Albert Bifet · Jesse Davis · Tomas Krilavičius · Meelis Kull · Eirini Ntoutsi · Indrė Žliobaitė (Eds.)

Machine Learning and Knowledge Discovery in Databases

Research Track

European Conference, ECML PKDD 2024 Vilnius, Lithuania, September 9–13, 2024 Proceedings, Part I

Part I







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Preface

The 2024 edition of the European Conference on Machine Learning and Principles and Practice of Knowledge Discovery in Databases (ECML PKDD 2024) was held in Vilnius, Lithuania, from September 9 to 13, 2024.

The annual ECML PKDD conference acts as a world-wide platform showcasing the latest advancements in machine learning and knowledge discovery in databases. Held jointly since 2001, ECML PKDD has established itself as the leading European Machine Learning and Data Mining conference. It offers researchers and practitioners an unparalleled opportunity to exchange knowledge and ideas about the latest technical advancements in these disciplines. Moreover, the conference appreciates the synergy between foundational advances and groundbreaking data science and hence strongly welcomes contributions about how Machine Learning and Data Mining is being employed to solve real-world challenges.

The conference continues to evolve reflecting evolving technological developments and societal needs. For example, in the Research Track this year there has been an increase in submissions on generative AI, especially LLMs, and various aspects of responsible AI.

We received 826 submissions for the Research Track and 224 for the Applied Data Science Track. The Research track accepted 202 papers (out of 826, 24.5%) and the Applied Data Science Track accepted 56 (out of 224, 24.5%). In addition, 31 papers from the Journal Track (accepted out of 65 submissions) and 14 Demo Track papers (accepted out of 30 submissions).

The papers presented over the three main conference days were organized into five distinct tracks:

Research Track: This track featured research and methodology papers spanning all branches within Machine Learning, Knowledge Discovery, and Data Mining.

Applied Data Science Track: Papers in this track focused on novel applications of machine learning, data mining, and knowledge discovery to address real-world challenges, aiming to bridge the gap between theory and practical implementation.

Journal Track: This track included papers that had been published in special issues of the journals *Machine Learning* and *Data Mining and Knowledge Discovery*.

Demo Track: Short papers in this track introduced new prototypes or fully operational systems that leverage data science techniques, demonstrated through working prototypes. **Nectar Track:** Concise presentations of recent scientific advances published in related conferences or journals. It aimed to disseminate important research findings to a broader audience within the ECML PKDD community.

The conference featured five keynote talks on diverse topics, reflecting emerging needs like benchmarking and resource-awareness, as well as theoretical understanding and industrial needs.

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- Gintarė Karolina Džiugaitė (Google DeepMind): *The Dynamics of Memorization and Unlearning*.
- Moritz Hardt (Max Planck Institute for Intelligent Systems): *The Emerging Science of Benchmarks.*
- Mounia Lalmas-Roelleke (Spotify): Enhancing User Experience with AI-Powered Search and Recommendations at Spotify.
- Patrick Lucey (Stats Perform): *How to Utilize (and Generate) Player Tracking Data in Sport.*
- Katharina Morik (TU Dortmund University): *Resource-Aware Machine Learning*—*a User-Oriented Approach.*

The ECML PKDD 2024 Organizing Committee supported Diversity and Inclusion by awarding some grants that enable early career researchers to attend the conference, present their research activities, and become part of the ECML PKDD community. We provided a total of 3 scholarships of \in 1000 to individuals that come from the developing countries and/or communities which are underrepresented in science and technology. The scholarships could be used for travel and accommodation. In addition 3 grants covering all of the registration fees were awarded to individuals who belong to underrepresented communities, based on gender and role/position, to attend the conference and present their research activities. The Diversity and Inclusion action also included the Women Networking event and Diversity and Inclusion Panel discussion. The Women Networking event aimed to create a safe and inclusive space for networking and reflecting on the experience of women in science. The event included a structured brainstorm/reflection on the role and experience of women in science and technology, which will be published in the conference newsletter. The Diversity and Inclusion Panel aimed to reach a wider audience and encourage the discussion on the need for diversity in tech, and challenges and solutions in achieving it.

We want to thank the authors, workshop and tutorial organizers, and participants whose scientific contributions make this such an exciting event. Moreover, putting together an outstanding conference program would also not be possible without the dedication and (substantial) time investments of the area chairs, program committee, and organizing committee. The event would not run smoothly without the many volunteers and sessions chairs. Finally, we want to extend a special thanks to all the local organizers – they dealt with all the little details that are needed to make the conference a memorable event.

We want to extend our heartfelt gratitude to our wonderful sponsors for their generous financial support. We also want to thank Springer for their continuous support and Microsoft for allowing us to use their CMT software for conference management and providing help throughout. We very much appreciate the advice and guidance provided

by the ECML PKDD Steering Committee over the past two years. Finally, we thank the organizing institution, the Artificial Intelligence Association of Lithuania.

September 2024

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Invited Talks Abstracts

The Dynamics of Memorization and Unlearning

Gintarė Karolina Džiugaitė

Google DeepMind

Abstract. Deep learning models exhibit a complex interplay between memorization and generalization. This talk will begin by exploring the ubiquitous nature of memorization, drawing on prior work on "data diets", example difficulty, pruning, and other empirical evidence. But is memorization essential for generalization? Our recent theoretical work suggests that eliminating it entirely may not be feasible. Instead, I will discuss strategies to mitigate unwanted memorization by focusing on better data curation and efficient unlearning mechanisms. Additionally, I will examine the potential of pruning techniques to selectively remove memorized examples and explore their impact on factual recall versus in-context learning.

Biography: Gintarė is a senior research scientist at Google DeepMind, based in Toronto, an adjunct professor in the McGill University School of Computer Science, and an associate industry member of Mila, the Quebec AI Institute. Prior to joining Google, Gintarė led the Trustworthy AI program at Element AI/ServiceNow, and obtained her Ph.D. in machine learning from the University of Cambridge, under the supervision of Zoubin Ghahramani. Gintarė was recognized as a Rising Star in Machine Learning by the University of Maryland program in 2019. Her research combines theoretical and empirical approaches to understanding deep learning, with a focus on generalization, memorization, unlearning, and network compression.

The Emerging Science of Benchmarks

Moritz Hardt

Max Planck Institute for Intelligent Systems

Abstract. Benchmarks have played a central role in the progress of machine learning research since the 1980s. Although there's much researchers have done with them, we still know little about how and why benchmarks work. In this talk, I will trace the rudiments of an emerging science of benchmarks through selected empirical and theoretical observations. Looking back at the ImageNet era, I'll discuss what we learned about the validity of model rankings and the role of label errors. Looking ahead, I'll talk about new challenges to benchmarking and evaluation in the era of large language models. The results we'll encounter challenge conventional wisdom and underscore the benefits of developing a science of benchmarks.

Biography: Hardt is a director at the Max Planck Institute for Intelligent Systems, Tübingen. Previously, he was Associate Professor for Electrical Engineering and Computer Sciences at the University of California, Berkeley. His research contributes to the scientific foundations of machine learning and algorithmic decision making with a focus on social questions. He co-authored Fairness and Machine Learning: Limitations and Opportunities (MIT Press) and Patterns, Predictions, and Actions: Foundations of Machine Learning (Princeton University Press).

Enhancing User Experience with AI-Powered Search and Recommendations at Spotify

Mounia Lalmas-Roelleke

Spotify

Abstract. This talk will explore the pivotal role of search and recommendation systems in enhancing the Spotify user experience. These systems serve as the gateway to Spotify's vast audio catalog, helping users navigate millions of music tracks, podcasts, and audiobooks. Effective search functionality allows users to quickly find specific content, whether it is a favorite song, a trending podcast, or an informative audiobook, while also satisfying broader search needs. Meanwhile, recommendation systems suggest new and relevant content that users might not have thought to search for, while ensuring their current needs for familiar content are met. This encourages exploration and discovery of new artists, genres, and shows, enriching the overall listening experience and keeping users engaged with the platform. Achieving this dual objective of precision and discovery requires sophisticated technology. It involves a deep understanding of representation learning, where both content and user preferences are accurately modeled. Advanced AI techniques, including machine learning and generative AI, play a crucial role in this process. These technologies enable the creation of highly personalized recommendations by understanding complex user behaviors and preferences. Generative AI, for instance, allows us to create personalized playlists, thereby enhancing the user experience with innovative features. This presentation is based on the collective research and publications of numerous contributors at Spotify.

Biography: Mounia is a Senior Director of Research at Spotify and the Head of Tech Research in Personalization, where she leads an interdisciplinary team of research scientists. She also holds an honorary professorship at University College London and serves as a Distinguished Research Fellow at the University of Amsterdam. Previously, Mounia was a Director of Research at Yahoo, overseeing a team focused on advertising quality and collaborating on user engagement projects related to news, search, and user-generated content. Before her tenure at Yahoo, Mounia held a Microsoft Research/RAEng Research Chair at the School of Computing Science, University of Glasgow, and before that was a Professor of Information Retrieval at the Department of Computer Science at Queen Mary, University of London. She is a prominent figure in the research community, regularly serving as a senior program committee member at major conferences such as WSDM, KDD, WWW, and SIGIR. She was also a program

l M. Lalmas-Roelleke

co-chair for SIGIR 2015, WWW 2018, WSDM 2020, and CIKM 2023. Mounia is widely recognized for her contributions as a speaker and author, with over 250 published papers and appearances on platforms like ACM ByteCast and the AI Business Podcasts series. She was nominated for the VentureBeat Women in AI Awards for Research in both 2022 and 2023.

How to Utilize (and Generate) Player Tracking Data in Sport

Patrick Lucey

Stats Perform

Abstract. Even though player tracking data in sports has been around for 25 years, it still poses as one of the most interesting and challenging datasets in machine learning due to its fine-grained, multi-agent, team-based, and adversarial nature. Despite these challenges, it is also extremely valuable as it is (relatively) low-dimensional, interpretable, and interactive, allowing us to measure performance and answer questions we couldn't objectively address before. In this talk, I will first give a brief history of tracking data in sports, then highlight the challenges associated with utilizing it. I will then show that by obtaining a permutation invariant representation, we can not only measure aspects of sports that couldn't be done before, but also interact with and simulate plays akin to a video game via our "visual search" and "ghosting" technology. Finally, I will show how we can use both tracking and event data to create a multimodal foundation model, which enables us to generate player tracking data at scale and achieve our goal of "digitizing every game of professional sport." Throughout the talk, I will utilize examples from top-tier basketball, soccer, and tennis.

Biography: Patrick Lucey is currently the Chief Scientist at sports data giant Stats Perform, leading the AI team with the goal of maximizing the value of the company's extensive sports data. He has studied and worked in the fields of machine learning and computer vision for the past 20 years, holding research positions at Disney Research and the Robotics Institute at Carnegie Mellon University, as well as spending time at IBM's T.J. Watson Research Center while pursuing his Ph.D. Patrick originally hails from Australia, where he received his BEng(EE) from the University of Southern Queensland and his doctorate from Queensland University of Technology, which focused on multimodal speech modeling. He has authored more than 100 peer-reviewed papers and has been a co-author on papers in the MIT Sloan Sports Analytics Conference Best Research Paper Track for 11 of the last 13 years, winning best paper in 2016 and runner-up in 2017 and 2018. Additionally, he has won best paper awards at INTERSPEECH and WACV international conferences. His main research interests are in artificial intelligence and interactive machine learning in sporting domains, as well as AI education. He has recently piloted a course on "AI in Sport," which aims to give students intuition behind AI methods using the interactive and visual nature of sports data.

Website: www.patricklucey.com

Resource-Aware Machine Learning—A User-Oriented Approach

Katharina Morik

TU Dortmund University

Abstract. Machine Learning (ML) has become integrated into several processes, ranging from medicine, manufacturing, logistics, smart cities, sales, recommendations and advertisements to entertainment and many more business and private processes. The applications together consume a considerable amount of energy and emit CO2. ML research investigates how to make models smaller and faster through pruning and quantization. Also the use of more energy-efficient hardware is an encouraging field. Research on ML under resource constraints is an active field proposing novel algorithms and scenarios. The aim is that for each application a variety of implementations is offered from which customers and the different types of users may choose the most thrifty one. This, in turn, would push tech providers to focus on the production of economical systems. However, if the customers, users, stakeholders do not know which of the models offers the best tradeoff between performance and energy-efficiency, they cannot select the most frugal one. Hence, testing implementations of learning and inference needs to be developed. They should be easy to use, produce visualizations that are mass-tailored for specific user groups. Automatized testing is difficult due to the diversity of models, computing architectures, training and evaluation data, and the fast rate of changes. The talk will illustrate work on resource-aware ML and advocate to pay more attention to the role of users in the development of scenarios, models, and tests.

Biography: Katharina Morik received her doctorate from the University of Hamburg in 1981 and her habilitation from the TU Berlin in 1988. In 1991, she established the chair of Artificial Intelligence at the TU Dortmund. She retired in 2023. She is a pioneer of bringing machine learning and computing architectures together so that machine learning models may be executed or even trained on resource restricted devices. In 2011, she acquired the Collaborative Research Center CRC 876 "Providing Information by Resource-Constrained Data Analysis" consisting of 12 projects and a graduate school. After the longest possible funding period of 12 years, the CRC ended with the publication of 3 books on Resource-Constrained Machine Learning (De Gruyter). She has participated in numerous European research projects and has been the coordinator of one. She was a founding member and Program Chair of the conference series IEEE International Conference on Data Mining (ICDM) and is a member of the steering committee

of ECML PKDD. She is a co-founder of the Lamarr Institute for Machine Learning and Artificial Intelligence. Prof. Morik is a member of the Academy of Technical Sciences and of the North Rhine-Westphalian Academy of Sciences and Arts. She was made a Fellow of the German Society of Computer Science GI e.V. in 2019.

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Research Track



Adaptive Sparsity Level During Training for Efficient Time Series Forecasting with Transformers

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Abstract. Efficient time series forecasting has become critical for realworld applications, particularly with deep neural networks (DNNs). Efficiency in DNNs can be achieved through sparse connectivity and reducing the model size. However, finding the sparsity level automatically during training remains challenging due to the heterogeneity in the losssparsity tradeoffs across the datasets. In this paper, we propose "Pruning with Adaptive Sparsity Level" (PALS), to automatically seek a decent balance between loss and sparsity, all without the need for a predefined sparsity level. PALS draws inspiration from sparse training and duringtraining methods. It introduces the novel "expand" mechanism in training sparse neural networks, allowing the model to dynamically shrink, expand, or remain stable to find a proper sparsity level. In this paper, we focus on achieving efficiency in transformers known for their excellent time series forecasting performance but high computational cost. Nevertheless, PALS can be applied directly to any DNN. To this aim, we demonstrate its effectiveness also on the DLinear model. Experimental results on six benchmark datasets and five state-of-the-art (SOTA) transformer variants show that PALS substantially reduces model size while maintaining comparable performance to the dense model. More interestingly, PALS even outperforms the dense model, in 12 and 14 cases out of 30 cases in terms of MSE and MAE loss, respectively, while reducing 65% parameter count and 63% FLOPs on average. Our code and supplementary material are available on Github (https://github.com/ zahraatashgahi/PALS).

1 Introduction

The capabilities of transformers [51] for learning long-range dependencies [8,48, 54] make them an ideal model for time series processing [53]. Several transformer

variants have been proposed for the task of time series forecasting, which is crucial for real-world applications, e.g., weather forecasting, energy management, and financial analysis, and have proven to significantly increase the prediction capacity in long time series forecasting (LTSF) [34]. In addition, attention-based models are inherently an approach for increasing the interpretability for time series analysis in critical applications [27]. Moreover, recent transformer time series forecasting models (e.g., [34, 57, 64]) perform generally well in other time series analysis tasks, including, classification, anomaly detection, and imputation [56].

Despite the outstanding performance of transformers, these models are computationally expensive due to their large model sizes as shown in [47] for natural language processing. With the ever-increasing collection of large time series and the need to forecast millions of them, the requirement to develop computationally efficient forecasting models is becoming significantly critical [17,44,49]. For industry-scale time series data, which are often high-dimensional and longlength, deploying transformers requires automatically discovering memory- and computationally-efficient architectures that are scalable and practical for realworld applications [53]. While there have been some efforts to reduce the computational complexity of transformers in time series forecasting [63, 64], these models have in order of millions of parameters, that can be too large for resourcelimited applications, e.g., mobile phones. The over-parameterization of these networks causes high training and inference costs, and their deployment in lowresource environments (e.g., lack of GPUs) would be infeasible. To address these issues, we raise the research question: How can we reduce the computational and memory overheads of training and deploying transformers for time series forecasting without compromising the model performance?

Seeking sparsity through sparse connectivity is a widely-used technique to address the over-parameterization of deep learning models [16]. Early approaches for deriving a sparse sub-network prune a trained dense model [15], known as *post-training* pruning. While these methods can match the performance of the dense network as shown by the Lottery Ticket Hypothesis (LTH) [11], they are computationally expensive during training due to the training of the dense network. *During-training* pruning aims to maintain training efficiency by gradually pruning a dense network during training [28]. Sparse training [40] pushed the limits further by starting with a sparse network from scratch and optimizing the topology during training. However, as we study in Sect. 3, the main challenge when using any of these techniques for time series forecasting is to find the proper sparsity level automatically.

In this paper, we aim to move beyond optimizing a single objective (e.g. minimizing loss) and investigate sparsity in DNNs for time series prediction in order to find a good trade-off between computational efficiency and performance automatically. Our contributions are: (1) We analyze the effect of sparsity (using unstructured pruning) in SOTA transformers for time series prediction [34,57, 63,64], and vanilla transformer [51]. We show they can be pruned up to 80% of their connections in most cases, without significant loss in performance. (2) We



Fig. 1. Schematic overview of the proposed method, PALS (Algorithm 1), Dynamic Sparse Training (DST) [9,40], During-training pruning (Gradual Magnitude Pruning (GMP) [65], and GraNet [28]). While DST and during-training pruning use a fixed sparsity schedule to achieve a pre-determined sparsity level at the end of the training, PALS updates the sparse connectivity of the network at each Δt iterations during training, by deciding whether to "Shrink" (decrease density) or "Expand" (increase density) the network or remain "Stable" (same density), to automatically find a proper sparsity level.

propose an algorithm, called "Pruning with Adaptive Sparsity Level (PALS) that finds a decent loss-sparsity trade-off by dynamically tuning the sparsity level during training using the loss heuristics and deciding at each connectivity update step weather to Shrink or Expand the network, or keep it Stable. PALS (Fig. 1) creates a bridge between during-training pruning and dynamic sparse training research areas by inheriting and enhancing some of their most successful mechanisms, while - up to our best knowledge - introducing for the first time into play also the Expand mechanism. Consequently, PALS does not require a desired pre-defined sparsity level which is necessary for most pruning or sparse training algorithms. (3) We evaluate the performance of PALS in terms of the loss, the parameter count, and FLOPs on six widely-used benchmarks for time series prediction and show that PALS can substantially sparsify the models and reduce parameter count and FLOPs. Surprisingly, PALS can even outperform the dense model on average, in 12 and 14 cases out of 30 cases in terms of Mean Squared Error (MSE) and Mean Absolute Error (MAE) loss, respectively (Table 2).

2 Background

2.1 Sparse Neural Networks

Sparse neural networks (SNNs) use sparse connectivity among layers to reduce the computational complexity of DNNs while maintaining a close performance

5

to the dense counterpart in terms of prediction accuracy. SNNs can be achieved using dense-to-sparse or sparse-to-sparse approaches [39].

Dense-to-sparse methods prune a dense network; based on the pruning phase, they are categorized into three classes: *post-training* [11, 15], *Before-training* [23], and *during-training* [28,35,65] pruning. Post-training pruning suffers from high computational costs during training and before-training approaches usually fall behind the performance of the dense counter-part network. In contrast, duringtraining approaches, maintain close or even better performance to the dense network while being efficient through the training process. A standard duringtraining pruning is Gradual Magnitude Pruning (GMP) [65] which gradually drops unimportant weights based on the magnitude during the training process. GraNet [28] is another during-training algorithm that gradually shrinks (decreasing density) a network to reach a pre-determined sparsity level. It prunes the weights (as performed in GMP) while allowing for connection regeneration (as seen in Dynamic Sparse Training (DST) which will be explained in the following). As the number of grown weights is less than the pruned ones, the network is shrunk and the density is decreased. For details regarding GraNet, refer to Appendix B.

Sparse-to-sparse methods start with a random sparse network from scratch and the number of parameters is usually fixed during training and can be determined based on the available computational budget. The sparse topology can remain fixed (static) [38] or dynamically optimized during training (a.k.a Dynamic Sparse Training (DST)) [9,18,29,32,40,59]. At each topology update iteration, a fraction of unimportant weights are dropped (usually based on magnitude), and the same number of weights are grown. The growth criteria can be random, as in Sparse Evolutionary Training (SET) [40], or gradient, as in Rigged Lottery (RigL) [9].

In this work, we take advantage of the successful mechanism of "*Shrink*" from duringtraining pruning (e.g., GraNet [28]) and "*Stable*" from DST (e.g., RigL [9]) and propose for the first time the "*Expand*" mechanism, to design a method to automatically optimize the sparsity level during training without requiring to determine

 Table 1. Comparison of related work.

Method	Shrin	k Stable	Expand	Adaptive	Automatic
				Sparsity	tune of
				Schedule	sparsity
					level
RigL	X	1	Х	Х	Х
GMP	1	X	Х	Х	Х
GraNet	1	1	Х	Х	X
PALS	1	1	1	1	1
(ours)					

it beforehand. Each of these mechanisms is explained in Sect. 4. In Table 1, we present a summarized comparison with the closest related work in the literature. Figure 1 presents a comprehensive embedding of our proposed method in the literature. Unlike these methods, which update the network using fixed schedules to reach a pre-determined sparsity level, PALS proposes an adaptive approach. It automatically determines whether to shrink or expand the network or remain

stable, in order to tune the sparsity level and find a good trade-off between loss and sparsity.

Only a few works investigated SNNs for time series analysis [46]. [58] investigates sparsity in convolutional neural networks (CNNs) for the time series classification and shows their proposed method has superior prediction accuracy while reducing computational costs. [20] exploit sparse recurrent neural networks (RNNs) for outlier detection. [30] and [12] explore sparsity in RNNs for sequence learning.

Sparsity in Transformers. Several works have sought sparsity in transformers [13,42]. These approaches can be categorized into structured (blocked) [37] or unstructured (fine-grained) pruning [5]. As discussed in [16], structured sparsity for transformers is able to only discover models with very low sparsity levels; therefore, we focus on unstructured pruning. [26] analyses pruning transformers for language modeling tasks and shows that large transformers are robust to compression. [4] dynamically extract and train sparse sub-networks from Vision Transformers (ViT) [8] while maintaining a fixed small parameter budget, and they could even improve the accuracy of the ViT in some cases. [7] investigates DST for BERT language modeling tasks and shows Pareto improvement over the dense model in terms of FLOPs. However, these works mostly focus on vision and NLP tasks. To the best of our knowledge, no work has investigated sparse connectivity in transformers for time series analysis that faces domain-specific challenges as we will elaborate in Sect. 3. Please note that there is a line of research focusing on sparse attention [50] aiming to develop an efficient selfattention mechanism that is orthogonal to our focus in this work (sparsity and pruning) [16].

2.2 Time Series Forecasting

Initial studies for time series forecasting exploit classical tools such as ARIMA [2]. While traditional methods mostly rely on domain expertise or assume temporal dependencies follow specific patterns, machine learning techniques learn the temporal dependencies in a data-driven manner [25,27,52]. In recent years, various deep learning models, including RNNs [43,45], multi-layer perceptrons (MLP) [60,61], CNNs [22], and Temporal convolution networks [10] are utilized to perform time series forecasting [3,19,41].

Transformers have been extensively used to perform time series forecasting due to their strong ability for sequence modeling. A class of models aims at improving the self-attention mechanism and addresses the computational complexity of vanilla transformers such as LogTrans [24], Informer [63], Reformer [21]. Another category of methods seeks to modify the model to capture the inherent properties of the time series: Autoformer [57] introduces a seasonal trend decomposition with an auto-correlation block as the attention module. NSTransformer [34] proposes to add two modules including series stationarization and de-stationary attention in the transformer architecture. FEDformer [64] proposes to combine transformers with a seasonal-trend decomposition method to capture global and detailed behaviour of the time series. The research into designing transformers for time series forecasting is ongoing, and many other transformer variants have been proposed, such as Crossformer [62], ETSformer [55], Pyraformer [33].

2.3 Problem Formulation and Notations

Let $\mathbf{x}_t \in \mathbb{R}^m$ denote the observation of a multivariate time series \mathbf{X} with m variables at time step t. Given a look-back window $\mathbf{X}_{t-L:t} = [\mathbf{x}_{t-L}, ..., \mathbf{x}_{t-1}]$ of size L, time series forecasting task aims to predict time series over a horizon H as $\widetilde{\mathbf{X}}_{t:t+H} = [\widetilde{\mathbf{x}}_t, ..., \widetilde{\mathbf{x}}_{t+H-1}]$ where $\widetilde{\mathbf{x}}_t$ is the prediction at time step t. To achieve this, we need to train a function $f(\mathbf{X}_{t-L:t}, \theta)$ (e.g. a transformer network) that can predict future values over horizon H.

In this paper, we aim to reduce the model size by pruning the unimportant parameters from θ such that we find the sparse model $f(\mathbf{X}_{t-L:t}, \theta_s)$ where $||\theta_s||_0 \ll ||\theta||_0$. $D = \frac{||\theta_s||_0}{||\theta||_0}$ is called the density level of the model f and S = 1-Dis called as the sparsity level. The aim is to minimize the reconstruction loss between the prediction $\mathcal{L}(f(\mathbf{X}_{t-L:t}, \theta_s), \mathbf{X}_{t:t+H})$ while finding a proper sparsity level S automatically. We use Mean Squared Error (MSE) as the loss function such that:

$$\mathcal{L}(\widetilde{\boldsymbol{X}}_{t:t+H}, \boldsymbol{X}_{t:t+H}) = \frac{1}{H} \Sigma_{i=0}^{H-1} (\widetilde{\boldsymbol{x}}_{t+i} - \boldsymbol{x}_{t+i})^2.$$
(1)

3 Analyzing Sparsity Effect in Transformers for Time Series Forecasting

In this section, we explore sparsity in several time series forecasting transformers. In short, we apply GraNet [28] to prune each model and measure their performance over various sparsity levels.

Experimental Settings. We perform this experiment on six benchmark datasets, presented in Table 4. We adapt GraNet [28], a during-training pruning algorithm developed for CNNs, to sparsify transformer models for time series forecasting. GraNet gradually shrinks a network (here, we start from a dense network) during the training to reach a pre-determined sparsity level, while allowing for connection regeneration inspired by DST. GraNet is described in Appendix B. For more details regarding the experimental settings, please refer to Sect. 5.1. For each sparsity level (%) in $\{25, 50, 65, 80, 90, 95\}$, we measure the prediction performance of each transformer model in terms of MSE loss. The results for prediction length = 96 (except 24 for the Illness dataset) are presented in Fig. 2. The results for other prediction lengths are presented in Fig. 3 in Appendix B.

Sparsity Effect. We present the results for pruning various transformers in Fig. 2; most models can be pruned up to 80% or higher sparsity levels without significantly affecting performance. Moreover, a counter-intuitive observation is that



Fig. 2. Sparsity effect on the performance of various transformer models for time series forecasting on benchmark datasets in terms of MSE loss (prediction length = 96, except 24 for the Illness dataset). Each model is sparsified using GraNet [28] to sparsity levels (%) $\in \{25, 50, 65, 80, 90, 95\}$ and PALS. *Sparsity* = 0 indicates the original dense model.

in some cases, sparsity does not necessarily lead to worse performance than the dense counterpart, and it can even improve the performance. For example, while on the Electricity, Illness, and Traffic datasets, the behavior is as usually expected (higher sparsity leads to lower performance), on the three other datasets, higher sparsity might even lead to better performance (lower loss) than the dense model. In addition, the sparsity effect is different among various models, particularly on the latter group of datasets, including the ETTm2, Exchange, and Weather datasets. We discuss the potential reasons for different behavior among datasets in Appendix H. Last but not least, by looking at Fig. 3 in Appendix B, the prediction length can also be a contributing factor to the sparsity-loss trade-off.

Challenge. Based on the above observations, we can conclude that the sparsity effect is not homogeneous across various time series datasets, forecasting models, and prediction lengths for time series forecasting. Our findings in these experiments are not aligned with the statements in [16] for CNNs (vision) and Transformers (NLP), where for a given task and technique, increasing the sparsity level results in decreasing the prediction performance. However, we observe in Figs. 2 and 3 that increasing the sparsity level does not necessarily lead to decreased performance and it might even significantly improve the performance (e.g. for the vanilla transformer on the Weather dataset). Therefore, it is challenging to decide how much we can push the sparsity level and what is the decent sparsity level without having prior knowledge of the data, model, and experi-

Algorithm 1. PALS

1: Input: Time series $X \in \mathbb{R}^{T \times m}$, number of training iterations t_{max} , Sequence length L. Prediction length H, model dimension d_{model} , pruning rate ζ , mask update frequency Δt , Initial density D_{init} , pruning rate factor $\gamma > 1$ and loss freedom factor $\lambda > 1$, sparsity bound S_{min} and S_{max} . 2: Initialization: Initialize the transformer model with density level D_{init} , S = 1 - 2 $D_{init}, L_{best} = \inf.$ 3: Training: 4: for $t \in \{1, \ldots, \#t_{max}\}$ do I. Standard feed-forward and back-propagation. The network is trained 5:on $batch_t$ of samples. II. Update sparsity mask 6: 7: if $(t \mod \Delta t) = 0$ then Compute Validation Loss L_{valid}^t 8: if $(S < S_{min})$ or $(L_{valid}^t <= \lambda * L_{best} \text{ and } S < S_{max})$ then 9: $update_mask \ (\zeta_{prune} = \gamma * \zeta, \ \zeta_{grow} = \zeta)$ 10:else if $L_{valid}^t > \lambda * L_{best}$ and $S > S_{best}$ then 11: $update_mask \ (\zeta_{prune} = \zeta, \ \zeta_{grow} = \gamma * \zeta)$ 12:13:else 14:update mask ($\zeta_{prune} = \zeta, \zeta_{qrow} = \zeta$) 15:end if if $L_{valid}^t < L_{best}$ then 16: $L_{best} = L_{valid}^t, \ S_{best} = S$ 17:18:end if Set S to the sparsity level of the network. 19:20:end if 21: end for

mental settings. While GraNet is the closest in spirit to our proposed method, it cannot automatically tune the sparsity level since it needs the initial and the final sparsity level as its hyperparameters. In this paper, we aim to address this challenge by proposing an algorithm that can automatically tune the sparsity level during training.

4 Proposed Methodology: PALS

This section presents our proposed method for automatically finding a proper sparsity level of a DNN, called "**P**runing with **A**daptive **S**parsity Level (**PALS**) (Algorithm 1). While our main focus in this paper is to sparsify transformer models, PALS is not specifically designed for transformers and can be applied directly to other artificial neural network architectures (See Appendix F for experiments on training with PALS the DLinear [60]) model.

Motivation and Broad Outline. As we discussed in Sect. 3, the main challenge when seeking sparsity for time series forecasting is to find a good sparsity level

automatically. Therefore, PALS aims to tune the sparsity level during training without requiring prior information about models or datasets. PALS is in essence inspired by the DST framework [40] and gradual magnitude pruning (GMP) [28,65]. While DST and GMP use fixed sparsification policies (fixed sparsity level (*Stable* in Fig. 1) and constantly prune the network until the desired sparsity level is reached (*Shrink* in Fig. 1), respectively) and require the final sparsity level before training, PALS exploits heuristic information from the network at each iteration to automatically determine whether to *increase*, *decrease*, or *keep* the sparsity level at each connectivity update step. While existing growing methods [14,36] grow a network or a layer of it to dense connectivity, to the best of our knowledge, this is the first work that allows the network to *expand* by increasing the density during training without requiring dense connectivity, and allows for automatic shrink or expand. If the training starts from a dense neural network ($D_{init} = 1$) PALS can be seen as a dense-to-sparse method, while if $D_{init} < 1$ then PALS is a sparse-to-sparse method.

Training. The training of PALS (Algorithm 1) starts with initializing a network with density level $D_{init} = 1 - S_{init}$. Then, the training procedure of PALS consists of two steps:

1. Standard feed-forward and back-propagation. Network's parameters are updated each training iteration t using a batch of samples.

2. Update Sparse Connectivity. The novelty of the method lies in updating the sparse connectivity. At every Δt iteration, the connectivity is updated in two steps. (2-1) The validation loss at step t is calculated as L_{valid}^t . (2-2) The sparsity mask is updated (update_mask in Algorithm 1) by first pruning ζ_{prune} of weights with the lowest magnitude:

$$\widetilde{\boldsymbol{W}}_{l} = Update(\boldsymbol{W}_{l}, top(|\boldsymbol{W}_{l}|, 1 - \zeta_{prune})), \qquad (2)$$

where W_l is the l^{th} weight matrix of the network, Update(A, idx) keeps only the indices idx of the matrix A, $top(A, \zeta)$ returns the indices of a fraction ζ of the largest elements of A. Then, we grow ζ_{grow} of the weights with the highest gradients:

$$\boldsymbol{W}_{l} = \widetilde{\boldsymbol{W}}_{l} + top(|\boldsymbol{G}_{l,i\notin\widetilde{\boldsymbol{W}_{l}}}|, \zeta_{grow})$$
(3)

where $G_{l,i\notin \widetilde{W_l}}$ is the gradient of zero weights in layer l. These new connections are initialized with zero values. This process is repeated for each layer in the model. Based on the values of ζ_{prune} and ζ_{grow} , PALS determines whether to decrease (*shrink*), increase (*expand*), or keep (*stable*) the network:

 $\mathbf{S_t} > \mathbf{S_{t-1}}$ (Shrink). If the loss does not go beyond $\lambda * L_{best}$, we decrease the overall number of parameters such that $\zeta_{prune} = \gamma * \zeta$, $\zeta_{grow} = \zeta$. The loss freedom coefficient, $\lambda > 1$, is a hyperparameter of the network that determines how much the loss value can deviate from the best validation loss achieved so far L_{best} during training. The lower λ is, the more strict PALS becomes at allowing the network to go to the *shrink* phase, finally resulting in a lower sparsity network. $\gamma > 1$ is the pruning factor coefficient, which determines how much to prune or grow more in the shrink and expand phases, respectively. We

analyze the sensitivity of PALS to λ and γ in Sect. 6.2. In addition, we define a boundary for sparsity determined by S_{min} and S_{max} which can be determined by the user based on the available resources. If the sparsity level does not meet the minimum sparsity level S_{min} , we prune the network more than we grow. If the network sparsity goes beyond S_{max} , we do not increase sparsity.

 $\mathbf{S_t} < \mathbf{S_{t-1}}$ (Expand). If $S > S_{best}$ (S_{best} is the sparsity level corresponding to L_{best}) and the loss goes higher than $\lambda * L_{best}$, it means that the earlier pruning step(s) were not beneficial to decreasing the loss (improving forecasting quality in the time series forecasting) and the network requires a higher capacity to recover a good performance. Therefore, we expand the network and grow more connections than the pruned ones at this step: $\zeta_{prune} = \zeta$, $\zeta_{grow} = \gamma * \zeta$.

 $\mathbf{S_t} = \mathbf{S_{t-1}}$ (Stable). If none of the above cases happened, we only update a fraction ζ of the network's parameters without changing the sparsity level: $\zeta_{prune} = \zeta$, $\zeta_{grow} = \zeta$.

For a better understanding of how the sparsity level evolves during the training process of PALS, please refer to Appendix G.4.

5 Experiments and Results

5.1 Experimental Settings

Datasets. The experiments are performed on six widely-used benchmark datasets for time series forecasting. The datasets are summarized in Table 4 and described in Appendix A. These datasets have different characteristics including stationary and non-stationary with/without obvious periodicity. Each database in each experiment is divided into three sets: train, validation, and test set. The data from the test set is only used for the final evaluation of all methods. The validation data is used to choose the best model during training and early stopping for all models including dense and sparse. Therefore, all methods use the loss signal from validation data to tune their model and select the model with the lowest validation loss, and they have all seen an equal amount of data during training.

Models. We consider five SOTA transformer models for time series forecasting, including Non-Stationary Transformer (NSTransformer) [34], FEDformer [64], Autoformer [57], Informer [63], and vanilla transformer [51]. Please refer to Sect. 2.2 for more details.

Evaluation Metrics. We evaluate the methods in two aspects: 1) Quality of the prediction in terms of MSE and MAE, and 2) Computational complexity in terms of parameter count and FLOPs (Floating-point operations). We report the theoretical FLOPs to be independent of the used hardware, as it is done in the unstructured pruning literature [9,28]. A lower value for these metrics indicates higher prediction quality and lower computational complexity, respectively. We measure the performance of each model for various prediction lengths $H \in \{96, 192, 336, 720\}$ (except $H \in \{24, 36, 48, 60\}$ for the Illness dataset).

Implementation. Experiments are implemented in PyTorch. The start of implementation is the NSTransformer¹ and GraNet². We repeat each experiment for three random seeds and report the average of the runs. In the experiments, D_{init} was set to 1, thus PALS can be seen as a during-training pruning method. We have run the experiments on Intel Xeon Platinum 8360Y CPU and one NVIDIA A100 GPU. We will discuss the hyperparameters' settings in Appendix A.

5.2 Results

Multivariate Time Series Prediction. The results in terms of MSE and parameter count for the considered datasets and models are presented in Table 5 in Appendix C. In most cases considered, PALS decreases the model size by more than 50% without a significant increase in loss. More interestingly, in most cases on the ETTm2, Exchange, and Weather datasets PALS even achieves lower MSE than the dense counterpart.

To summarize the results of Table 5 (Appendix C) and have a general overview of the performance of PALS on each model and dataset, we present the average MSE and MAE, and parameters count in addition to the difference between the dense and the sparse model using PALS (in percentage) in Table 2. Additionally, we include the inference FLOPs count (total FLOPs for all test samples). PALS even outperforms the dense model in 12 and 14 cases out of 30 cases in terms of MSE and MAE loss, respectively, while reducing 65% parameter count and 63% FLOPs on average. We summarize the training FLOPs in Appendix G.1.

Based on the experiments conducted in Sect. 3 and the description of datasets provided in Appendix A.1, we observed significant variations in the sparsityloss trade-off across different datasets and models. The beauty of our proposed method consists in the fact that it does not have to consider any of these differences. We did not make any finetuning for PALS to account for these differences, and it does everything automatically. Of course, finetuning PALS per dataset and model specificity would improve its final performance, but it would reduce the generality of our proposed work and we prefer not to do it.

Univariate Time Series Prediction. The results of univariate prediction (using a single variable) on the ETTm2 and Exchange datasets are presented in Table 6 and summarized in Table 7 in Appendix D. In short, PALS outperforms the dense counterpart model on average, in 7 and 8 cases out of 12 cases in terms of MSE and MAE loss, respectively.

6 Discussion

In this section, we study the performance of PALS in comparison with other pruning and DST algorithms (H) and the hyperparameter sensitivity of PALS

¹ https://github.com/thuml/Nonstationary Transformers.

² https://github.com/VITA-Group/GraNet.

Table 2. Summary of the results on the benchmark Datasets in Table 5. For each experiment on a transformer model and dataset, the before and after applying PALS. The difference between these results is shown in % where the blue color means improvement of PALS average MSE, MAE, number of parameters $(\times 10^6)$ and the inference FLOPs count $(\times 10^{12})$ for various prediction lengths are reported compared to the corresponding dense model in terms of MSE or MAE.:

Model	Electric	ity			ETTm2				Exchange	~		1	llness			<u>-</u>	haffic				Weather			
	MSE	MAE	#Param:	s #FLOPs	MSE	MAE	#Params	#FLOPs	MSE	AE ⊨	⊭Params #	∉FLOPs]	VISE M	AE #F	arams #1	FLOPS	ISE	AE 3	⊭Params	#FLOPs	MSE	MAE	<i>∉</i> Params	∉FLOPs
NSTransformer	0.19	0.30	12.0	9.25	0.49	0.43	10.6	19.82	0.54 C	.49 1	0.6 1	.89	2.14 0.	92 10.	5 0.0	500	.63 6	.34	4.2	5.61	0.29	0.31 1	0.7	8.10
+PALS	0.21	0.32	2.2	1.81	0.38	0.39	2.5	3.70	0.49 0	0.47 5	.4 1	2 20.	2.33 0.5	97 7.4	0.0	14 0	-67 6	.37 4	ę	2.05	0.26	0.29	0.	1.77
Difference	10.8% 1	7.3% †	81.5% [10.5%	24.0% ↓	11.2% [76.7% ↓	81.3% ↓	9.3% 1 3	.6% 14	8.5% 1 4	13.3% ↓ 5).1% ↑5	1% 130.	0% 1 30	-2% ↓ 5	-2% † 5	11% 1	1 % 1.0	1 %0.65	10.2% [9 1 %6.9	0.3% [90.2% L
FEDformer	0.21	0.32	19.5	9.30	0.30	0.35	17.9	19.82	0.50 C	.49 1	7.9 1	.89	2.84 1.	14 13.	7 0.0	15 0	.61 6	.38	2.3	3.71	0.32	0.37 1	7.9	8.11
+PALS	0.23	0.34	3.0	1.35	0.30	0.35	1.8	1.96	0.51 C	1.50 1	0.5 1	15 5	3.05 1	19 8.3	0.0	0 0	-62 6	38 5	9.6	L.83	0.31	0.36]	×.	.81
Difference	9.0% ↑	4.9%	84.7% ↓	85.5% [$1.5\% \uparrow$	0.5% [90.2% [1 %1.06	$2.1\% \uparrow 1$.1% 14	1.2% 1 3	1 1 %6.8	7.2% ↑ 5.v	0% † <mark>39</mark> .	5% J 39	-6% ↓ 1	-0% † 1	.1% † %1.	4.6% [72.7% [2.8% [3.2% 1 9	0.0% 1	1 %0.06
Autoformer	0.24	0.34	12.1	9.30	0.33	0.37	10.5	19.82	0.58 C	.53 1	0.5 1	-89	3.08 1	18 10.	5 0.0	15 0	.64 6	40	4.9	3.71	0.34	0.38 1	0.6	8.11
+ PALS	0.26	0.36	2.7	1.71	0.31	0.35	1.0	1.93	0.62 C	.55 7	- -	.30 5	3.19 1.3	22 6.7	0.0	0 0	.65 6	141 4	5	1.94	0.34	0.38 1	e.	2.52
Difference	9.3% ↑	4.6% \uparrow	77.7% 1	$81.6\% \downarrow$	8.1% ↓	5.6% ↓	90.3%	1 %£.06	5.5% ↑ 4	.4% 13	2.7% 1 3	31.0% [5	3.5% †3	2% † 36.	6% J 36	5% ↓ 1	-9% † 2	11% 1 6	1 % 2.6	1 %0.12	0.1%↑	1.3% ↓ 8	7.7% ↓	$36.1\% \downarrow$
Informer	0.36	0.43	12.5	8.51	1.53	0.88	11.3	18.15	1.59 1	.00	1.3 1	71 5	5.27 1.4	58 11.	3 0.0	15 0	.81 6	A6	4.4	5.14	0.62	0.55 1	1.4	6.58
+PALS	0.42	0.48	1.4	0.94	1.39	0.83	5.3	8.45	1.53 C	98 8	.6 1	.33	5.23 1.4	57 7.4	0.	03 0	-94 6	.53	.3	1.19	0.69	0.56 4	er;	3.35
Difference	18.9% 1	11.3%	↑88.6% ↓	1 %6.88	1 %0.6	5.8% [53.4% [53.4% [3.9% []	.6% 12	4.2% 1 2	2.4% L C	AL 1 88.0	0% 1 34.	9% J 34	8% 1	5.5% † 1	5.3% 18	1 %6.63	\$0.6% J	12.1% ↑	2.0% ↑ 6	1.9% 1	1 %7.61
Transformer	0.28	0.38	11.7	9.24	1.48	0.86	10.5	19.81	1.61 C	.97 1	0.5 1	.89	1.94 1.4	49 10.	5 0.0	15 0	.67 6	.36	3.6	3.61	0.64	0.56 1	0.6	8.10
+PALS	0.31	0.40	2.5	2.24	1.08	0.75	3.2	8.17	1.41 C	.91 6	.6 1	16 4	1.91 1.	48 7.7	0.0	0 0	-69 6	.38	8	1.86	0.32	0.38]	0	1.76
Difference	10.5%]	5.7% †	78.3% ↓	75.8% ↓	26.7%	13.3%	1 %6.69	58.8%	12.5% 16	.1% []3	7.5% [3	8.4% J C	0.7% [0.0	9% 1 27.	2% 27	3% 3	-3%↑ 5	$.4\% \uparrow$	1.7% ↓	1.8%	49.6% ↓	32.5% 1	0.2% ↓	90.3% [
Differenceavg	11.7% 1	6.8%	82.1% ↓	82.5% ↓	13.3% ↓	1.3% [76.1% ↓	$74.8\% \downarrow$	$3.6\% \downarrow 1$.2% 13	6.8% [3	14.8% L 5	3.7% † 2	3% 133.	6% 33	7% 1 5	4% † 6	.6% † 1	4.0% ↓	1 %0.87	10.1% [9.0% J 8	4.0% ↓	31.3% [

(Sect. 6.2). Additionally in the Appendix, we analyze the performance of PALS in terms of model size (H), prediction quality by visualizing the predictions (I), pruning DLinear [60] (F), and computational efficiency from various aspects (G).

6.1 Performance Comparison with Pruning and Sparse Training Algorithms

We compare PALS with a standard during-training pruning approach (GMP [65]), GraNet [28], and a well-known DST method (RigL [9]). These are the closest methods in the literature in terms of including gradual pruning and gradient-based weight regrowth.

While PALS derives a proper sparsity level automatically, other pruning approaches require the sparsity level as an input of the algorithm. Therefore, to compare PALS with existing pruning algorithms, the sparsity level should be optimized for them. We apply GraNet, RigL, and GMP to NSTransformer for prediction lengths of $H \in \{96, 192, 336, 720\}$ (except for the Illness dataset for which $H \in \{24, 36, 48, 60\}$). For each of these methods (GraNet, RigL, and GMP), the sparsity level is optimized among values of $\{25, 50, 65, 80, 90, 95\}$. This means that for one run of PALS, we run the other methods 6 times. The model with the lowest validation loss is used to report the test loss. Table 3 summarizes the average loss (l), sparsity level (S), and training epochs (e) (due to early stopping the algorithms might not require the full training) over different prediction lengths.

Table 3. Comparison with other during-training pruning methods (GMP, GraNet) and a DST method (RigL) when sparsifying NSTransformer. The results are average over four prediction lengths.

Dataset	PAL	S		Gra	Net*		RigI	`*		GM	Р*	
	1	S	е	1	S	e	l	S	e	1	S	e
Electricity	0.21	80.5%	8.83	0.20	31.2%	9.75	0.20	31.2%	9.12	0.20	47.5%	9.62
ETTm2	0.38	76.7%	4.58	0.60	95.0%	9.00	0.49	77.5%	4.33	0.60	56.2%	9.12
Exchange	0.49	48.5%	5.83	0.47	95.0%	9.42	0.44	90.0%	4.25	0.45	95.0%	9.50
Illness	2.33	30.0%	7.97	2.32	25.0%	9.58	2.37	25.0%	9.58	2.22	31.2%	9.92
Traffic	0.67	70.1%	8.83	0.64	41.2%	9.50	0.64	25.0%	8.17	0.64	50.0%	9.79
Weather	0.26	90.3%	7.00	0.28	95.0%	9.08	0.27	41.2%	4.08	0.29	95.0%	9.08
Average	0.72	66.0%	7.17	0.75	63.73%	9.38	0.74	48.3%	6.60	0.73	62.4%	9.47

*Optimized sparsity level (%) in $\{25, 65, 50, 80, 90, 95\}$. GraNet, RigL, and GMP, each require 6 runs to optimize the sparsity level while PALS needs only one run.

The closest competitor of PALS is GraNet. In Table 3, for the Electricity dataset, PALS achieves a sparsity level of 80.5% with a loss of 0.21, while GraNet achieves a sparsity level of only 31.2% with a slightly lower loss of 0.20. Similarly,

for the ETTm2 dataset, PALS achieves a sparsity level of 76.7% with a loss of 0.38, while GraNet achieves a higher sparsity level of 95.0% but with a much higher loss of 0.60. On the other datasets, they perform relatively close to each other.

By looking at the results of all methods in Table 3, PALS has the highest average sparsity value (66.0%) compared to GraNet (63.73%), RigL (48.3%), and GMP (62.4%). While RigL requires fewer training epochs (~ 6.6 epochs) compared to PALS (~ 7.2 epochs), it finds lower sparsity networks and has a higher average loss (RigL: 0.74 compared to PALS: 0.72). GraNet and GMP use fixed pruning schedules, and as a result, they need almost full training time (~ 9.5 epochs). The only extra computational requirement of PALS compared to GraNet is an additional step that involves determining the number of weights to prune and grow. This is negligible when considering the overall computation necessary for training the models. On the other hand, as PALS does not require the full training epochs in contrast to GraNet, it needs much lower computational costs. We additionally compared the convergence speed of PALS with the dense model in Appendix G.3.

In short, PALS has the lowest average loss and highest sparsity values compared to other algorithms, suggesting that PALS could build efficient and accurate sparse neural networks for time series forecasting.

6.2 Hyperparameter Sensitivity

In this section, we discuss the sensitivity of PALS to its hyperparameters including pruning rate factor γ and loss freedom factor λ . We have changed their values in {1.05, 1.1, 1.2} and measured the performance of PALS (with NSTransformer) in terms of MSE and parameter count on six benchmark datasets. The results are presented in Table 8 in Appendix E.

As shown in Table 8, PALS is not very sensitive to its hyperparameters and the results in each row are close in terms of loss in most cases considered. However, by increasing γ and λ PALS tends to find a sparser model. A small λ results in paying more attention to the loss value, while a large value gives more freedom to PALS to explore a sparse sub-network that might sometimes result in a higher loss value. A small γ limits the amount of additional grow/prune in the expand/shrink phase, while a large γ gives more flexibility to the algorithm for exploring various sparsity levels. In short, a small value for each of these hyperparameters makes PALS more strict and allows for small changes in sparse connectivity, while a large value increases the exploration rate which potentially results in higher sparsity and/or reduced loss.

7 Conclusions

In this paper, we aim to decrease the computational and memory costs of training and deploying DNNs for time series forecasting rather than proposing a new forecasting model and beating the state-of-the-art. Particularly, we focus on transformers while showing the generality of PALS on an MLP-based model (Appendix F). We first showed that pruning networks for time series forecasting can be challenging in terms of determining the proper sparsity level for various datasets, prediction lengths, and models. Therefore, we proposed PALS, a novel method to obtain sparse neural networks, that exploits loss heuristics to automatically find the best trade-off between loss and sparsity in one round of training. PALS leverages the effective strategies of "Shrink" from during-training pruning and "Stable" from DST. Additionally, we introduce a novel strategy called the "Expand" mechanism. The latter allows PALS to automatically optimize the sparsity level during training, eliminating the need for prior determination. Remarkably, PALS could outperform dense training in 12/14 cases out of 30 cases (5 transformer models, 6 datasets) in terms of MSE/MAE loss, while reducing 65% parameters count and 63% FLOPs on average. Limitations and future work. Due to the lack of proper hardware to support sparse matrices for on-GPU processing, PALS cannot currently take advantage of its theoretical training and inference speed-up and memory reduction in a real-world implementation. Building a truly sparse transformer demands a substantial investment of both effort and a profound understanding of hardware, an area that is beyond the current scope of our research and human resources (Please refer to Appendix G.5 for more details). With the ever-increasing body of work on sparse neural networks, we hope that in the near future, the community paves the way to optimally train sparse neural networks on GPU. An open direction to this research can be to start with a highly sparse neural network (as opposed to starting from a dense network used in PALS) and gradually expand the network to be even more efficient during training.

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RumorMixer: Exploring Echo Chamber Effect and Platform Heterogeneity for Rumor Detection

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Abstract. Rumors have exerted detrimental effects on individuals and societies in recent years. Despite the deployment of sophisticated Graph Neural Networks (GNNs) to analyze the structure of propagation graphs in rumor detection, contemporary approaches often neglect two pivotal elements. Firstly, the structure of rumor propagation in social networks is characterized by a community-based feature, influenced by the "echo chamber effect". By integrating these structures, models can emphasize critical information, mitigate the impact of irrelevant data, and enhance graph representation learning. Secondly, the existing models for rumor detection struggle to adjust GNN backbones to accommodate the diverse complexities introduced by social media's platform heterogeneity. The manual design of these models is both time-consuming and labor-intensive. To overcome these challenges, this paper presents **RumorMixer**, a novel automated framework for rumor detection. This methodology begins by developing a Super-Sharer-Aware (SSA) chamber partitioning algorithm, crucial for identifying echo chambers within propagation graphs. Through accurate partitioning, RumorMixer effectively concentrates on the essential structures of rumor propagation and utilizes the GNN-Mixer model to create high-quality representations of these chambers. To address platform heterogeneity, RumorMixer integrates five distinct components: PE, Aggregation, Merge, Pooling, and Mixing-to establish an extensive search space. Subsequently, differentiable architecture search technology is employed to automatically tailor platform-specific architectures. The efficacy is validated through extensive experiments on real datasets from both Weibo and Twitter³(Our code is accessible at https://github.com/cgao-comp/RumorMixer.).

Keywords: Rumor detection \cdot Graph neural network \cdot Neural architecture search

1 Introduction

In the current digital era, social media has emerged as one of the primary channels for information propagation [25]. Its rapid and extensive spreading capacity has exerted a profound impact on society. However, these platforms also serve as fertile ground for the proliferation of rumor, posing serious threats to public health [10], political elections [19], and social stability [22].

Recent research indicates that the structure of news propagation within social networks plays a pivotal role in the identification of rumor [7,21]. Methods utilizing Graph Neural Networks (GNN) have been developed to analyze the propagation patterns within these networks, offering significant insights for rumor detection [5]. Although these studies confirm the effectiveness of propagation networks in enhancing the accuracy of rumor detection, they overlook the characteristics of the rumor itself.

- Echo Chamber Effect. The echo chamber effect describes the phenomenon where information circulates within groups of individuals on social media and cyberspace who have similar thoughts or close stances [6]. This leads to the rapid spread of rumors within communities formed based on common interests, viewpoints, or beliefs. The structure of rumor dissemination in social networks thus exhibits characteristics based on community. In this process, super-sharers play a pivotal role. They further reinforce the echo chamber effect by sharing information extensively, facilitating the rapid spread of rumors within the respective groups [1].
- Platform Heterogeneity. Social media platforms, with their unique content, interaction styles, and user behaviors, produce data with varied structures and dynamics, impacting rumor detection model effectiveness [16]. Models tailored for one platform may falter on another due to these differences. Moreover, existing rumor detection approaches struggle to adapt GNN architectures to the diverse complexities arising from platform variability, making model construction a tedious and labor-intensive process.

To more effectively investigate the aforementioned characteristics, we introduce RumorMixer, an innovative framework for rumor detection that consists of two primary strategies. Firstly, it utilizes a super-sharer-aware (SSA) algorithm to identify echo chambers within social networks. Secondly, it employs GNN-Mixers, as detailed in [13], to precisely represent these chambers. A notable aspect of RumorMixer is the implementation of a differentiable Neural Architecture Search (NAS) to enhance the framework, thereby significantly improving rumor detection capabilities on Weibo and Twitter platforms. We underscore our contributions to the domain through these advancements:

- Echo Chamber Extraction and Representation Learning. We introduce the Super-Sharer-Aware (SSA) algorithm for effectively identifying and isolating echo chambers in rumor propagation graphs on social media. The SSA algorithm employs a centrality-based method for selecting seed nodes, which are pivotal in the formation of echo chambers centered around influencers. This method integrates a flood-fill strategy with a breadth-first search to achieve coherent grouping of chambers, while utilizing dynamic programming to refine chamber assignments by considering the strength of connections and unique attributes. Then, we leverage a GNN-Mixer model for generating refined representations of rumor propagation graphs.

- Neural Architecture Search for Platform Heterogeneity. In response to the challenges posed by platform heterogeneity, RumorMixer incorporates five distinct components-*PE*, *Aggregation*, *Merge*, *Pooling*, and *Mixing*-to facilitate a comprehensive search space. Then the differentiable architecture search technology is leveraged to tailor platform-specific architectures automatically.
- This study conducts extensive experiments on real-world datasets. Rigorous evaluation demonstrates the practical applicability and adaptability of the proposed RumorMixer in diverse social media platforms.

The paper is structured as follows: Sect. 2 reviews related work, Sect. 3 introduces our framework, and Sect. 4 presents the experiments and ablation studies. Conclusions and future directions are discussed in Sect. 5.

2 Related Works

This study explores rumor detection methodologies through propagation graph analysis. Propagation-based approaches enhance rumor identification by integrating auxiliary data from social media, such as user comments, profiles, posting behaviors, and stances towards rumors [17, 30]. Recent research highlights the critical role of news propagation structures on social media in improving detection effectiveness. Further investigations have shown that merging propagation structures with temporal data yields better detection results [5, 23], while others have improved performance using attention mechanisms [27, 30]. Nonetheless, prevailing techniques often neglect the echo chamber effect-the amplification of rumors within insular communities by prolific sharers-which hampers a comprehensive understanding of rumor dynamics [1,6]. Moreover, the challenge posed by platform heterogeneity, which refers to variations in user interactions, content formats, and behaviors across different social media platforms, calls for bespoke detection models [16]. The inability of existing models to adjust to these variations and complexities makes manual model design both time-consuming and resource-intensive. In essence, devising a method that thoroughly examines the inherent aspects of rumor propagation, specifically the echo chamber effect and platform heterogeneity, is a pressing yet unaddressed concern in the discipline.

3 Methodology

3.1 Overview

The foundational structure of the RumorMixer is depicted in Fig. 1. This section aims to elucidate the decisions undertaken in the implementation of each architectural component.



Fig. 1. The RumorMixer framework introduces: (I) Echo chamber extraction:, where graph partitioning and the Super-sharer-aware algorithm are used to identify echo chambers within propagation graphs; (II) Rumor detection which combines chamber representations into a final graphical form, using global average pooling and a linear classification layer for prediction; (III) Automated architecture design, leveraging platform-specific GNN-Mixer architectures with a focus on differentiable NAS for chamber representation. The weight vectors for PE (α_e), aggregation (α_n), skip-connections (α_s), merge (α_l), pooling (α_p) and mixing (α_m) within the supernet.

Problem Formulation. Define $N = \{n_1, n_2, \ldots, n_k\}$ as the dataset for rumor detection, where n_i denotes the *i*-th event in a collection of *k* events. Each event, n_i , is structured as a tree $n_i = \{r, x_1, x_2, x_3, \ldots, x_{s-1}\}$, with *r* serving as the root node and x_i embodying the text representation of pertinent user comments. Here, *s* signifies the count of context nodes. Each event is assigned a label y_i

from two categories: $\{F, R\}$ (i.e., Fake and Real). This classification framework aims to facilitate the development of an model for automated rumor detection, leveraging the specified dataset.

3.2 Echo Chamber Extraction and Representation Learning

In the realm of social media, the spread of rumors is significantly influenced by "super-sharers": key accounts with extensive reach, notably in political contexts and during health crises like the COVID-19 pandemic. Research has highlighted platforms such as Twitter and Weibo, where these influential users amplify rumors within "echo chambers," spaces of homogenous opinions enhancing misinformation spread [25]. Leveraging graph partitioning techniques to analyze these echo chambers, our work introduces the Mixer model [13,24]. This model, drawing on foundational studies, is designed to capture the unique features of echo chambers, facilitating a deeper understanding of rumor dynamics.

Super-Sharer-Aware (SSA) Chamber Partitioning. In the field of social media analysis, this study presents a Super-Sharer-Aware (SSA) algorithm for chamber partitioning, detailed in Algorithm 1, designed to detect and segregate echo chambers within a rumor propagation graph. The algorithm initiates by selecting seed nodes based on their centrality, positioning them at the network's core to foster the formation of echo chambers around pivotal nodes and main social hubs. It then utilizes a flood-fill technique, originating from these seed nodes, to methodically encompass the entire connected region, thereby grouping all pertinent nodes within the same echo chamber. This method, underpinned by a queue-supported breadth-first search, ensures the seamless expansion of each chamber, maintaining its coherence and preventing fragmentation. Furthermore, the algorithm adopts dynamic programming techniques for node assignment to echo chambers, taking into account the connection strength between a node and the chamber members, and the potential impact on chamber separation and size balance. Nodes are dynamically allocated to the most suitable echo chamber based on a scoring mechanism (outlined in Table 1). Additionally, to preserve all original graph edges and accommodate chamber overlap, the model allows for a one-hop neighborhood overlap among chambers.

Chamber Representation Learning. To thoroughly analyze and effectively manage the non-Euclidean data structures prevalent in echo chambers, this study employs GNNs equipped with a message-passing mechanism as the encoding method for chambers. The architecture of these GNNs is meticulously crafted to transform a chamber token, denoted as G_c , into a fixed-size representation, x_{G_c} , through a 3-step process.

Table 1. Scoring calculation methods. Coh signifies the Cohesion Score, aggregating the connections between the target node v and nodes within the candidate echo chamber (CN). Sep reflects the Separation Score, tallying connections between v and nodes outside the candidate chamber but within other chambers (EN). Bal denotes the Balance Score, predicated on the size of the candidate chamber (|CN|), ensuring a balanced distribution of nodes across chambers. T represents the Total Score, incorporating Coh, Sep, and Bal to determine the most suitable chamber for node v.

Score Type	Calculation Method
$\overline{\text{Cohesion } (Coh)}$	$Coh = \sum_{u \in CN} [G.has_edge(u, v)]$
Separation (Sep)	$Sep = \sum_{u \in EN} [G.has_edge(u, v)]$
Balance (Bal)	$Bal = \frac{1}{ CN +1}$
Total (T)	T = Coh - Sep + Bal

Algorithm 1. Super-Sharer-Aware Chamber Partitioning

- 1: **Input:** Graph G = (V, E), Number of echo chambers K
- 2: **Output:** Mapping *L* of nodes to echo chambers
- 3: Select K seed nodes based on centrality \triangleright Initiate chambers with central nodes
- 4: Initialize K queues, one for each echo chamber \triangleright Prepare for breadth-first expansion
- 5: Mark all nodes as unvisited
- 6: while there are unvisited nodes \mathbf{do}
- 7: for each queue do
- 8: Dequeue a node u \triangleright Expand from the most recently added node
- 9: for each unvisited neighbor v of u do
- 10: Evaluate best chamber k^* for v
- 11: Enqueue v in $Q[k^*]$, mark v as visited

12: Assign v to chamber k^*

- 13: end for
- 14: end for
- 15: end while
- 16: **return** L

Step 1. Message-Passing Layer with GNNs. This study implements a series of L message-passing layers for user node representation updates within each chamber $G_c = (\mathcal{V}_c, \mathcal{E}_c)$ through the application of GNNs. The updating process is mathematically represented as:

$$\mathbf{h}_{v,c}^{l} = \sigma\left(\mathbf{W}^{l} \cdot \operatorname{AGG}\left(\left\{\mathbf{h}_{v,c}^{l-1}, \forall u \in \widetilde{N}(v)\right\}\right)\right), \tag{1}$$

where $\mathbf{h}_{v,c}^{l} \in \mathbb{R}^{d_{l}}$ denotes the hidden features of node v, learned at the *l*-th layer $(l = 1, \dots, L)$, with c indicating the chamber index and d_{l} the dimensionality. Here, \mathbf{W}^{l} refers to the layer-specific trainable weight matrix applicable to all nodes in the graph, while σ represents a nonlinear activation function, such as sigmoid or ReLU. The aggregation function AGG(·), a critical component, varies among different GNN architectures. The study further incorporates a

 $[\]triangleright$ Return the mapping of nodes to their chambers

READOUT function to consolidate node features into a cohesive representation of each chamber, calculated as:

$$\mathbf{z}_{p,c} = \text{READOUT}\left(\left\{\mathbf{h}_{v,c}^{L}, \forall v \in \mathcal{V}_{c}\right\}\right) \in \mathbb{R}^{d},\tag{2}$$

where the READOUT function encompasses any form of pooling operation. To enhance this methodology and tailor it to specific computational requirements, the research employs a NAS strategy to identify the most effective aggregation and pooling mechanisms, as detailed in Sect. 3.3.

Step 2. Positional Information. Euclidean datasets inherently possess a structured arrangement; however, this inherent ordering is absent in general graphs, leading to diminished model expressivity due to the lack of positional information. To address this issue, explicit positional encodings (PE) are introduced for the nodes within the framework. The input features of nodes, as described in Eq. 3, are enhanced by incorporating $p_{v,c} \in \mathbb{R}^K$, utilizing a learnable matrix $W^0 \in \mathbb{R}^{d \times K}$:

$$h_{v,c}^0 = x_{v,c} \oplus W^0 p_{v,c} \in \mathbb{R}^d, \tag{3}$$

where $x_{v,c}$ denotes the original node features, $h_{v,c}^0$ signifies the initial node embedding, and \oplus symbolizes concatenation.

Step 3. Mixer Layer. Consider a scenario where $\mathbf{Z}_p \in \mathbb{R}^{C \times d}$ represents the collection of chamber embeddings $\{\mathbf{z}_{p,1}, \ldots, \mathbf{z}_{p,C}\}$. The formulation of the mixer layer is articulated as follows:

$$\hat{\mathbf{Z}}_{m} = \mathbf{Z}_{p} + (W_{2}\sigma(W_{1} \text{ LayerNorm } (\mathbf{Z}_{p}))) \in \mathbb{R}^{C \times d},
\mathbf{Z}_{m} = \hat{\mathbf{Z}}_{m} + \left(W_{4}\sigma\left(W_{3} \text{ LayerNorm } (\hat{\mathbf{Z}}_{m})^{T}\right)\right)^{T} \in \mathbb{R}^{C \times d},$$
(4)

where $\hat{\mathbf{Z}}_p$ signifies the intermediate embedding, with $\mathbf{Z}_m = \{\mathbf{z}_{m,1}, \ldots, \mathbf{z}_{m,C}\}$ delineating the output embedding from the mixer layer. The function σ is representative of the GELU nonlinearity, while LayerNorm(·) is indicative of layer normalization. The matrices $W_1 \in \mathbb{R}^{d_s \times C}, W_2 \in \mathbb{R}^{C \times d_s}, W_3 \in \mathbb{R}^{d_c \times d}$, and $W_4 \in \mathbb{R}^{d \times d_c}$ are defined with d_s and d_c representing adjustable hidden dimensions within token-mixing and channel-mixing MLPs respectively. The architecture allows for the substitution of the MLP in the Mixer layer with a Graph Multi-Head Attention (gMHA) mechanism, converting the system into a Graph Transformer as delineated:

$$\hat{\mathbf{Z}}_{m} = \mathbf{Z}_{p} + \text{gMHA}(\text{ LayerNorm } (\mathbf{Z}_{p})) \in \mathbb{R}^{C \times d},
\mathbf{Z}_{m} = \hat{\mathbf{Z}}_{m} + \text{MLP}(\text{ LayerNorm } (\hat{\mathbf{Z}}_{m})) \in \mathbb{R}^{C \times d},$$
(5)

where $gMHA(\cdot)$ aims to elucidate token interdependencies reflective of the specified chamber configuration. In Eq. 5, gHMA may initially embody a full-attention framework, albeit alternative methodologies outlined in Table 2 are viable to explicate the gHMA operation. NAS strategy is employed to identify the most efficacious mechanism as discussed in Sect. 3.3. Step 4. Global Average Pooling and Rumor Detection. Subsequently, the ultimate propagation graph representation \mathcal{Z} is derived through global average pooling across all chambers:

$$\mathbf{Z} = \frac{\sum_{c} m_{c} \cdot \mathbf{Z}_{m}}{\sum_{c} m_{c}} \in \mathbb{R}^{d},\tag{6}$$

with m_c being a binary indicator that assigns 1 to non-empty chambers and 0 to empty ones, addressing the variability in propagation graph sizes that may result in empty chambers. The process culminates in the application of a straightforward full-connected MLP for rumor detection:

$$y = \mathrm{MLP}(\mathbf{Z}) \in \mathbb{R}^n,\tag{7}$$

3.3 Neural Architecture Search for Platform Heterogeneity

In different social media platforms, the characteristics of rumor dissemination display heterogeneity. This is primarily due to the significant impact of each platform's design and structure on the speed and scope of rumor spread. Additionally, user behavior varies across platforms because each one possesses a unique user base and community culture. Therefore, designing a unified model for all platforms may be lack in consideration. To address this issue, we introduce the concept of NAS to design platform-specific architecture.

The Search Space Design. Through the analysis in Sect. 3.2, we can ascertain that the message-passing layer (Sect. 3.2) and mixer layer (Sect. 3.2) are searchable. The search space, delineated in Table 2, encompasses a diverse array of candidate operations distributed among five modules: PE, Aggregation, Merge, Pooling, and Mixing.

PE module incorporates four distinct methods of positional encoding: random walk [9], laplacian eigenvector [31], SVD-based [15], and centrality encoding [8] to exploit the potential of the graph encoding strategies, encapsulated by the symbol α_e . Aggregation module employs an integration of five distinct graph neural networks (GNNs) for the purpose of node representation. These networks include the Graph Convolutional Network (GCN) [18], Graph Attention Networks (GAT) [26], Sample and AGgregatE (SAGE) [12], and Graph Isomorphism Network (GIN) [29], alongside a Multilayer Perceptron (MLP) operation that operates independently of the graph's topology. The combined functionalities of these components are symbolized as α_n . Merge module incorporates a range of five merging mechanisms designed to synthesize graph-based representations, namely Long Short-Term Memory (LSTM), concatenation, maximum, mean, and summation operations. It also integrates IDENTITY and ZERO functions to enable optional skip-connections across layers, denoted by α_l and α_s , respectively. *Pooling module* utilizes three global pooling strategies to produce discrete, snapshot-level representations of graphs, encapsulated by the symbol α_p . This module's output provides a comprehensive representation of the graph's

Module	Operation	Candidate
PE	α_e	RandomWalk, Laplacian, SVD, Centrality
Aggregation	α_n	GCN, GAT, SAGE, GIN
Merge	α_l	M_LSTM, M_CONCAT, M_MAX, M_MEAN, M_SUM
	α_s	IDENTITY, ZERO
Pooling	α_p	GLOBAL_MEAN, GLOBAL_MAX, GLOBAL_SUM
Mixing	α_m	MLP, FULL_ATT (softmax $\left(\frac{QK^T}{\sqrt{d}}\right)V$),
		$GRAPH_ATT \; (softmax \left(A^P \odot \frac{QK^T}{\sqrt{d}} \right) V),$
		KERNEL_ATT (softmax $\left(\operatorname{RW}\left(A^{P}\right) \odot \frac{QK^{T}}{\sqrt{d}} \right) V \right)$
		ADDITIVE_ATT (softmax $\left(\frac{QK^T}{\sqrt{d}}\right)V + LL(A^P)$)
		HADAMARD_ATT $\left(\left(A^P \odot \operatorname{softmax}\left(\frac{QK^T}{\sqrt{d}}\right)\right)V\right)$

Table 2. The operations for the search space of RumorMixer.

entire structure. *Mixing module* makes use of MLP mixer and gMHA mixer techniques, considering various attention mechanisms including full attention, graph attention, kernel attention, additive attention, and Hadamard attention [13]. The aggregate of these operations is represented as α_m .

Differentiable Architecture Search. Drawing on the principles established in differentiable architecture search [20], RumorMixer incorporates a Gumbel-Softmax distribution to facilitate the generalized selection of operations, extending the work of [4]. This approach is mathematically represented as follows:

$$\bar{o}^{ij}(h_{v,c}^{0}) = \sum_{o \in \alpha} \frac{\exp\left\{(\alpha^{ij}o + g(o))/\tau\right\}}{\sum_{o' \in \alpha} \exp\left\{(\alpha^{ij}o' + g(o'))/\tau\right\}} o(h_{v,c}^{0}),\tag{8}$$

where the mixing weights for operations between any two nodes (i, j) are denoted by $\boldsymbol{\alpha}^{ij} \in \mathbb{R}^{|\alpha|}$. The operations set, α , encompasses six predefined categories: α_e , α_n , α_l , α_s , α_p , and α_m . The parameter τ serves to adjust the soft-maximization process of the distribution. The function g_o denotes the noise component, which is sampled from the Gumbel distribution. The variable $h_{v,c}^0$, as defined in Eq. 3, denotes the initial hidden features inputted into a message-passing layer, encapsulating $\{\mathbf{h}_{u,c}^{(l-1)}, \forall u \in \tilde{N}(v)\}$. The composite operations designated as \bar{o}_e , \bar{o}_n , \bar{o}_s , \bar{o}_l , \bar{o}_p , and \bar{o}_m are formulated based on α_e , α_n , α_l , α_s , α_p , and α_m , following the guidelines set forth in Eq. 8. To simplify notation, the superscript in o^{ij} is excluded when its implication is evident from the context. The method for computing the representation of a specific node v with positional encoding in a chamber is depicted in Fig. 1(III) and formalized as follows:

$$\mathbf{h}_{v,c}^{l} = \sigma \left(\mathbf{W}_{n}^{l} \cdot \bar{o}_{n} \left(\left\{ \bar{o}_{e}(\mathbf{h}_{u,c}^{l-1}), \forall u \in \widetilde{N}(v) \right\} \right) \right).$$
(9)

where \mathbf{W}_n^l represents the weight matrix, which is consistently utilized across all potential architectures within the search space for user node aggregation. In the terminal layer associated with user node v, user embeddings are generated as follows:

$$\mathbf{H}_{v,c}^{l+1} = \left[\bar{o}_s \left(\mathbf{h}_{v,c}^1 \right), \cdots, \bar{o}_s \left(\mathbf{h}_{v,c}^l \right) \right], \\
\mathbf{z}_{v,c} = \bar{o}_l \left(\mathbf{H}_{v,c}^{l+1} \right),$$
(10)

where the symbol $[\cdot]$ denotes the concatenation of embeddings from l intermediate layers. The embeddings corresponding to all N nodes within a given chamber c are aggregated via a pooling module, resulting in the generation of the graphlevel representation $\mathbf{z}_{p,c}$ for chamber c. Then, the chamber embedding is fed into a mixer layer to obtain the mixer embedding $\mathbf{z}_{m,c}$:

$$\begin{aligned} \mathbf{z}_{p,c} &= \bar{o}_p \left(\left\{ \mathbf{z}_{v,c}, \forall v \in \mathcal{V}_c \right\} \right), \\ \mathbf{z}_{m,c} &= \bar{o}_m \left(\mathbf{z}_{p,c} \right), \end{aligned} \tag{11}$$

RumorMixer addresses a bi-level optimization challenge, as articulated in the following mathematical formulation:

$$\min_{\boldsymbol{\alpha} \in \mathcal{A}} \mathcal{L}_{\text{val}} \left(\mathbf{w}^{*}(\boldsymbol{\alpha}), \boldsymbol{\alpha} \right),$$

s.t. $\mathbf{w}^{*}(\boldsymbol{\alpha}) = \operatorname*{arg\,min}_{\mathbf{w}} \mathcal{L}_{\text{tra}}(\mathbf{w}, \boldsymbol{\alpha}).$ (12)

where \mathcal{L}_{tra} and \mathcal{L}_{val} denote the training and validation losses, respectively. The variable $\boldsymbol{\alpha} = \{\boldsymbol{\alpha}_e, \boldsymbol{\alpha}_n, \boldsymbol{\alpha}_s, \boldsymbol{\alpha}_l, \boldsymbol{\alpha}_p, \boldsymbol{\alpha}_m\}$ signifies the parameters defining a network's architecture, and $\mathbf{w}^*(\boldsymbol{\alpha})$ indicates the optimal weights derived through training. For the purpose of rumor detection, the cross-entropy loss function is employed.

4 Experiments

A comprehensive experimental evaluation is conducted on three real-world datasets to examine efficacy. Specifically, this section aims to address the following research questions:

- **RQ1: Effectiveness.** How does our proposed method perform compared with other state-of-the-art (SOTA) human-designed models and existing NAS approaches for rumor detection?
- RQ2: Modularity. How do the different components contribute to the model performance?
- RQ3: Sensitivity. How do the model architecture (number of GNN layers) and dataset pattern (number of chambers) affect the final detection performance?
- RQ4: Robustness. Can RumorMixer achieve early rumor detection, particularly if it can surpass baseline performances utilizing fewer snapshots?

4.1 Experimental Setting

Datasets. The evaluation of the proposed method has been conducted on two publicly available benchmark datasets: Twitter16 [3] and Weibo [21]. Detailed statistics of these datasets are presented in Table 3.

Statistic	Twitter16	Weibo
# of root posts	818	4,664
# of users	$173,\!487$	2,746,818
# of posts	204,820	$3,\!805,\!656$
# of true rumors	205	2,351
# of false rumors	205	2,313
# of unverified rumors	203	\
# of non-rumors	205	

Table 3. The statistics of the rumor detection datasets.

Baselines. We first evaluate whether our method can more accurately detect rumor compared with the following methods.

- G1: Human-designed architectures. The present analyses juxtapose RumorMixer with several seminal manually designed models in the domain of rumor detection. These include: StA-HiTPLAN [17], P-BiGAT [30], BiGCN
 [2], DGNF [23], DYNGCN [5], and DGTR [27].
- G2: NAS approaches. The scope of this investigation also encompasses various NAS techniques, such as: Random search, Bayesian-based search [28] and GraphNAS (a RL-based NAS approach for GNN) [11].

Implementation Details. Our evaluation of rumor detection performance relies on two established metrics: Accuracy (Acc.) and F1 score (F_1) , with outcomes derived from 5-fold cross-validation presented as mean values along with standard deviations. Evaluating NAS baselines and RumorMixer involves generating architecture candidates, followed by hyperparameter fine-tuning (learning rate, dropout) as per previous methodology [14]. A consistent use of a 3-layer architecture and a Gumbel-Softmax temperature of 0.2 is maintained.

4.2 Performance Comparison (RQ1)

The comparative performance results are presented in Table 4, followed by an in-depth analysis. No singular human-designed model (G1) demonstrates superiority in rumor detection across various datasets. For instance, DYNGCN outperforms others on the Weibo platform, whereas DGTR shows enhanced performance on Twitter datasets. This observation underscores the necessity for architectures that are tailored to the specific requirements of each dataset. In the case

of G2, in contrast to GraphNAS which primarily concentrates on aggregation layers, RumorMixer implements a differential search algorithm. This enables the identification of more efficient architectures for addressing rumor detection tasks.

 Table 4. Rumor detection results on Weibo and Twitter16 datasets. The state-of-theart (SOTA) human-designed baseline is marked with an underline, and the best result for each dataset is highlighted in boldface.

Model		Weibo		Twitter16	
		Acc.	F_1	Acc.	F_1
G1	StA-HiTPLAN [17]	87.1 ± 0.8	86.7 ± 1.2	79.3 ± 1.1	81.7 ± 1.3
	P-BiGAT [<mark>30</mark>]	90.8 ± 1.4	91.6 ± 1.0	80.5 ± 0.6	80.2 ± 1.5
	BiGCN [2]	89.9 ± 0.5	89.6 ± 1.3	81.9 ± 1.0	81.1 ± 0.4
	DGNF [23]	93.3 ± 1.1	93.2 ± 0.6	82.3 ± 0.7	82.4 ± 1.3
	DYNGCN [5]	94.7 ± 0.3	94.6 ± 0.4	82.6 ± 1.2	82.6 ± 0.8
	DGTR [27]	93.6 ± 0.9	93.2 ± 0.4	90.9 ± 1.0	90.7 ± 0.7
G2	Random	86.2 ± 0.3	85.5 ± 0.1	86.4 ± 0.4	86.9 ± 0.7
	Bayesian [28]	89.5 ± 0.6	89.8 ± 1.2	88.1 ± 1.5	88.1 ± 0.4
	GraphNAS [11]	90.3 ± 1.3	89.7 ± 0.2	87.2 ± 0.9	87.1 ± 1.4
	RumorMixer(Ours)	$\textbf{97.9} \pm \textbf{0.5}$	$\textbf{97.6}\pm\textbf{0.7}$	$\textbf{96.4}\pm\textbf{0.9}$	$\textbf{96.9} \pm \textbf{1.2}$

4.3 Ablation Study (RQ2)

In our ablation studies, we evaluated various choices made during the implementation of each component of the architecture. The variants are designed as:

- w/o Adjust removes the stage of 1-hop neighborhood adjusting.
- \mathbf{w}/\mathbf{o} CE removes the stage of chamber extraction.
- \mathbf{w}/\mathbf{o} **SSA** replaces our designed super-sharer-aware (SSA) partitioning with random partitioning.
- $\mathbf{w}/\mathbf{o}~\mathbf{PE}$ ignores positional encodings in message-passing layers.

Effect of Chamber Extraction. An experiment has been carried out to assess the impact of omitting the Chamber extraction phase, wherein the GNN-Mixer treated each node independently as a chamber. Findings, documented in Table 5, underscore the indispensability of the chamber extraction process. This procedure, incorporating graph partitioning and 1-hop adjusting, is instrumental in capturing crucial local information pertaining to the graph's structure.

Effect of Super-Sharer-Aware (SSA) Partitioning. The study also examines the advantages conferred by SSA partitioning over and random graph partitioning. In the latter two approaches, nodes are arbitrarily distributed across

Models	Weibo		Twitter16		
	Acc.	F_1	Acc.	F_1	
RumorMixer	$\textbf{97.9} \pm \textbf{0.5}$	$\textbf{97.6}\pm\textbf{0.7}$	$\textbf{96.4}\pm\textbf{0.9}$	$\textbf{96.9} \pm \textbf{1.2}$	
w/o Adjust	95.4 ± 0.1	95.4 ± 0.1	94.8 ± 0.2	94.3 ± 0.1	
w/o CE	91.6 ± 0.4	91.1 ± 0.2	92.3 ± 0.3	91.8 ± 0.1	
w/o SSA	89.3 ± 0.1	88.2 ± 0.3	90.7 ± 0.7	90.7 ± 0.1	
w/o PE	96.8 ± 0.2	97.2 ± 0.2	95.7 ± 0.2	96.3 ± 0.1	

Table 5. Ablation Study for RumorMixer.

a predetermined number of Chambers. Results, presented in Table 5, reveal that our algorithm outperforms random partitioning, particularly in larger graphs, supporting the hypothesis that Chambers ought to encapsulate nodes and edges with analogous semantic or informational attributes. Interestingly, even random partitioning achieves commendable outcomes, indicating that the model's efficacy is not exclusively reliant on chamber quality.

Effect of Positional Encoding. Experimental results indicate that the utility of Positional Encoding (PE) remains uncertain, as detailed in Table 5. While PE enhances the expressive capabilities of GNNs, its impact on generalization performance does not follow a similar trajectory. This implies an enhancement in model specificity, albeit without a corresponding improvement in generalization. Therefore, despite the theoretical support and burgeoning literature underscoring the benefits of PE for refining GNN predictions, further mathematical advancements are imperative for delineating more effective strategies and ensuring consistent enhancements in results.

Effect of Search Space. This study explores the impact of four key modules on search space effectiveness, employing the RumorMixer algorithm in diverse configurations, detailed in Table 6. By isolating each factor, the research assesses its unique contribution to algorithm performance. Limitations in the search space, particularly in NAS, can hinder optimal model functionality by curtailing exploration of advanced optimization techniques. Furthermore, the role of GNNs in identifying rumor spread is underscored by their information aggregation methods. Utilizing a message-passing framework, GNNs are crucial for analyzing rumor spread within networks.

Models	Weibo		Twitter16	
	Acc.	F_1	Acc.	F_1
Fixed Aggregation (GCN)	92.2 ± 0.8	91.7 ± 0.2	91.8 ± 0.5	91.8 ± 0.5
Fixed Merge (M_CONCAT)	95.2 ± 0.1	94.1 ± 0.4	95.5 ± 0.3	95.5 ± 0.3
Fixed Pooling (GLOBAL_SUM)	95.1 ± 0.1	95.6 ± 0.9	95.3 ± 0.3	95.3 ± 0.2
Fixed Mixing (FULL_ATT)	93.3 ± 0.6	94.7 ± 0.7	94.2 ± 0.4	94.2 ± 0.5

Table 6. Various search spaces have been utilized to evaluate the performance of RumorMixer. The first column depicts the specific module, evaluated using a single OP within the streamlined search space.

Comparison with Different Community Detection Algorithms. We analyzed the similarities between echo chamber extraction and community detection, focusing on the SSA algorithm's efficiency in handling these tasks. Our comparison with other community detection algorithms, using the F1 scores from the Weibo dataset, underscored the SSA algorithm's superior performance. This advantage arises from its strategy of initiating community detection from high-degree nodes (super-sharers), enabling rapid identification of key rumor spreaders. The SSA algorithm's ability to pinpoint these nodes, often highly active users, highlights its effectiveness in detecting rumors within social networks (Table 7).

Table 7. Complexities and performance of different community detection algorithms in Weibo dataset. $|\mathcal{V}|$ and $|\mathcal{E}|$ represent the total number of nodes and edges in a rumor propagation graph, respectively.

Algorithm	Time	Space	F_1
SSA (ours proposed in Sect. 3.2)	$O(\mathcal{V} + \mathcal{E})$	$O(\mathcal{V})$	$\textbf{97.6} \pm \textbf{0.7}$
Modularity Optimization	$O((\mathcal{V} + \mathcal{E}) \log \mathcal{V})$	$O(\mathcal{V} + \mathcal{E})$	94.3 ± 0.4
Hierarchical Clustering	$O(\mathcal{V} ^2 \log \mathcal{V})$	$O(\mathcal{V} ^2)$	93.5 ± 0.6
FastGreedy	$O(\mathcal{V} \log^2 \mathcal{V})$	$O(\mathcal{V} + \mathcal{E})$	93.8 ± 0.3
Louvain	$O(\mathcal{V} \log \mathcal{V})$	$O(\mathcal{V} + \mathcal{E})$	93.2 ± 0.1

4.4 Parameter Analysis (RQ3)

Number of Chambers. Analysis of Fig. 2(a). reveals that performance escalates with an increase in the number of echo chambers (denoted as #Chamber), reaching a plateau with minor variations once #Chamber equals 4. Consequently, the default setting for the number of chambers is established at 3 (C = 3).



Fig. 2. The test F_1 score w.r.t. different number of chambers and layers in twitter and weibo dataset.

Number of GNN Layers. In Sect. 4.2, we justify the choice of a three-layer Graph Neural Network (GNN) architecture (K = 3) based on its proven efficiency. Our study investigates shallow GNNs with K limited to 5. Experiments across K = 1 to 5 reveal performance trends (Fig. 2(b)), supporting our selection of L = 3 for our experimental framework (Sect. 4.2).

4.5 Early Rumor Detection (RQ4)

The goal of early detection is to pinpoint rumors right at the onset of their spread, crucial for measuring detection methods' effectiveness. This requires setting a timeline for detection benchmarks. Accuracy evaluations for any detection approach, including the proposed ones and existing baselines, depend on examining posts made before these benchmarks. Our analysis (Fig. 3) contrasts RumorMixer's effectiveness against models like DYNGCN, DGTR, and Graph-NAS using the Weibo and Twitter datasets, as shown in the referenced figure. Results indicate RumorMixer's superior accuracy early on and throughout subsequent checkpoints, highlighting the advantage of incorporating structural echo chamber features for more efficient rumor detection both initially and over time.



Fig. 3. The test F_1 score of rumor early detection on twitter and weibo dataset.

5 Conclusion

In this study, we introduce RumorMixer, a cutting-edge framework developed to detect rumors across social media platforms. Our methodology employs an advanced graph partitioning algorithm that highlights the significant role of super-sharers in creating echo chambers, instrumental in spreading false information. Moreover, the integration of Neural Architecture Search (NAS) with our Graph Neural Network Mixer (GNN-Mixer) models improves their adaptability and efficiency in identifying rumors, automatically adjusting to the unique features of various social media platforms. Additionally, it must be acknowledged that methods focusing solely on user interactions in network-based approaches are not the only avenues for rumor detection. A comprehensive fake news detection system may need to consider content, structure, and dynamics, which also represents one of the potential directions for future expansion of this work.

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Diversified Ensemble of Independent Sub-networks for Robust Self-supervised Representation Learning

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Abstract. Ensembling a neural network is a widely recognized approach to enhance model performance, estimate uncertainty, and improve robustness in deep supervised learning. However, deep ensembles often come with high computational costs and memory demands. In addition, the efficiency of a deep ensemble is related to diversity among the ensemble members, which is challenging for large, over-parameterized deep neural networks. Moreover, ensemble learning has not vet seen such widespread adoption for unsupervised learning and it remains a challenging endeavor for self-supervised or unsupervised representation learning. Motivated by these challenges, we present a novel self-supervised training regime that leverages an ensemble of independent sub-networks, complemented by a new loss function designed to encourage diversity. Our method efficiently builds a sub-model ensemble with high diversity, leading to well-calibrated estimates of model uncertainty, all achieved with minimal computational overhead compared to traditional deep selfsupervised ensembles. To evaluate the effectiveness of our approach, we conducted extensive experiments across various tasks, including indistribution generalization, out-of-distribution detection, dataset corruption, and semi-supervised settings. The results demonstrate that our method significantly improves prediction reliability. Our approach not only achieves excellent accuracy but also enhances calibration, improving on important baseline performance across a wide range of selfsupervised architectures in computer vision, natural language processing, and genomics data.

1 Introduction

Ensemble learning has become a potent strategy for enhancing model performance in deep learning [19,28]. This method involves combining the outputs of multiple independently trained neural networks, all using the same architecture

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and same training dataset but differing in the randomized configurations of their initialization and/or training. Despite its remarkable effectiveness, training deep ensemble models poses several challenges: i) The high performance achieved by deep ensembles comes with a significant increase in computational costs. Running multiple neural networks independently demands more resources and time. ii) Maintaining diversity among ensemble members – a property often critical to success – becomes increasingly difficult for large, over-parameterized deep neural networks [37] in which the main source of diversity stems from random weight initialization. iii) Most of the existing literature focuses on deep ensembles for supervised models. Adapting these approaches to unsupervised and selfsupervised models requires careful consideration and evaluation to ensure comparable performance.

In recent years, self-supervised learning methods have achieved cutting-edge performance across a wide range of tasks in natural language processing (NLP; [2,9]), computer vision [5,40], multimodal learning [36], and bioinformatics [18]. In contrast to supervised techniques, these models learn representations of the data without relying on costly human annotation. Despite remarkable progress in recent years, self-supervised models do not allow practitioners to inspect the model's confidence. This problem is non-trivial given the degree to which critical applications rely on self-supervised methods. As recently discussed by LeCun¹, representing predictive uncertainty is particularly difficult in self-supervised contrastive learning for computer vision. Therefore, quantifying the predictive uncertainty of self-supervised models is critical to more reliable downstream tasks. Here, we follow the definition of reliability as described by Plex [44], in which the ability of a model to work consistently across many tasks is assessed. In particular, [44] introduces three general desiderata of reliable machine learning systems: a model should generalize robustly to new tasks, as well as new datasets, and represent the associated uncertainty in a faithful manner.

In this paper, we introduce a novel, robust, and scalable framework for ensembling *self-supervised learning* while *preserving performance* with a negligible increase in computational cost and *encouraging diversity among the ensemble* of *sub-networks*.

Our contributions can be summarized as follows:

- We propose a novel, scalable ensemble component of self-supervised learning that is robust, efficient and enhances performance in various downstream tasks.
- We develop a complementary loss function to enforce diversity among the independent sub-networks.
- We perform extensive empirical analyses to highlight the benefits of our approach. We demonstrate that this inexpensive modification achieves very competitive (in most cases, better) predictive performance: 1) on in-distribution (IND) and out-of-distribution (OOD) tasks; 2) in semi-supervised settings; 3) learns a better predictive performance-uncertainty trade-off than compared

¹ https://ai.facebook.com/blog/self-supervised-learning-the-dark-matter-of-intelligence/.

baselines. (i.e., exhibits high predictive performance and low uncertainty on IND datasets as well as high predictive performance and high uncertainty on OOD datasets).



Fig. 1. Illustration of our proposed method. Given a batch X of input samples, two different views \tilde{x} and \tilde{x}' are produced for each sample, which is then encoded into representations by the encoder network $f_{\theta'}$. The representations are projected to the ensemble of independent sub-networks g_m , where each sub-network produces embedding vectors z and z'. The mean value of these embeddings is passed to the self-supervised loss, while their standard deviation is used for the diversity loss. Finally, the total loss is computed by a combination of the two loss components.

2 Related Work

Self-supervised Learning. For most large-scale modeling problems, learning under full supervision is severely inhibited by the scarcity of annotated samples. Self-supervised learning techniques, which solve *pretext tasks* [9] to generate labels from (typically abundant) unlabeled data, have proven to be a powerful remedy to this bottleneck. The learned feature maps can serve as a starting point for *downstream* supervised tasks, such as classification, object detection, or sentiment analysis, with a substantially reduced need for labeled examples [25]. Alternatively, the downstream application may directly use the extracted representation for problems such as anomaly OOD detection. While there have been attempts to make pretraining more robust by preventing embedding collapse [40] or boosting performance in OOD detection [39,45], the aspect of *uncertainty-awareness* has been studied to a lesser extent in the self-supervised context. Motivated by this, we present a simple way to make self-supervised learning robust during pretext-task learning.

Ensemble Learning. Deep ensembles [28] comprise a set of M neural networks that independently train on the same data using random initialization.

Deep ensembles often outperform other approaches in terms of calibration and predictive accuracy [38], but their naive application incurs high computational complexity, as training, memory, and inference cost multiply with the number of base learners. BatchEnsemble [46] introduces multiple low-rank matrices, with little training and storage demand, whose Hadamard products with a shared global weight matrix mimic an ensemble of models. Masksensemble [13] builds upon Monte Carlo dropout [15] and proposes a learnable (rather than random) selection of masks used to drop specific network neurons. MIMO [20] uses ensembles of sub-networks diverging only at the beginning and end of the parent architecture – thus sharing the vast majority of weights – to obtain multiple predictions with a single forward pass.

Diversity in Ensembles: Diversity is a crucial component for successful ensembles. [37] classify existing approaches for encouraging diversity among ensemble members into three groups: i) methods that force *diversity in gradients* with adaptive diversity in prediction [35], or using joint gradient phase and magnitude regularization (GPMR) between ensemble members [7], ii) methods focusing on *diversity in logits*, improving diversity with regularization and estimating the uncertainty of out-of-domain samples [30], iii) methods promoting *diversity in features* that increase diversity with adversarial loss [4] for conditional redundancy [37], information bottleneck [42], or f1-divergences [4]. Our method belongs to this last category, where our loss function encourages the diversity of feature maps.

3 Method

We propose a simple principle to 1) make self-supervised pretraining robust with an ensemble of diverse sub-networks, 2) improve predictive performance during pretraining of self-supervised deep learning, 3) while keeping an efficient training pipeline.

As depicted in Fig. 1, our proposed method can be readily applied to recent trends in self-supervised learning [3,5,10,17,18,26] and is based on a joint embedding architecture. In the following sections, we first describe our proposed ensemble model, followed by the diversity loss, and then a discussion on diversity, and computational cost.

3.1 Robust Self-supervised Learning via Independent Sub-networks

Setting. Given a randomly sampled mini-batch of data $\mathbf{X} = \{\mathbf{x}_k\}_{k=1}^N \subset \mathcal{X} \subseteq \mathbb{R}^p$, the transformer function derives two augmented views $\tilde{\mathbf{x}} = \tau(\mathbf{x}), \tilde{\mathbf{x}}' = \tau'(\mathbf{x})$ for each sample in \mathbf{X} . The augmented views are obtained by sampling τ, τ' from a distribution over suitable data augmentations, such as masking parts of sequences [1,10], partially masking image patches [21], or applying image augmentation techniques [5].

The two augmented views \tilde{x} and \tilde{x}' are then fed to an encoder network f_{θ} with trainable parameters $\theta \subseteq \mathbb{R}^d$. The encoder (e.g., ResNet-50 [22], ViT [12])

maps the distorted samples to a set of corresponding features. We call the output of the encoder the *representation*. Afterward, the representation features are transformed by M independent sub-networks $\{g_{\phi_m}\}_{m=1}^M$ with trainable parameters ϕ_m to improve the feature learning of the encoder network. The ensemble constructs from the representation M different q-dimensional embedding vectors $\{z_m\}_{m=1}^M, \{z'_m\}_{m=1}^M$, respectively, for \tilde{x} and \tilde{x}' . We modify the conventional self-supervised loss and replace the usual z_m by the mean value $\bar{z} = (z_1 + \ldots + z_M)/M$, and similarly z'_m by \bar{z}' . Averaging over the embeddings generated by the M sub-networks increases robustness, which in turn may help to improve predictive performance in downstream tasks.

Self-supervised Loss. In the case of contrastive learning [5], the self-supervised loss ℓ_{ssl} with temperature t > 0 and cosine similarity $sim(\cdot, \cdot)$ is computed as:

$$\ell_{\rm ssl}\left(\tilde{\boldsymbol{x}}_{k}, \tilde{\boldsymbol{x}}_{k}'\right) = -\log \; \frac{\exp(\sin(\bar{\boldsymbol{z}}_{k}, \bar{\boldsymbol{z}}_{k}')/t)}{\sum_{i=1}^{2N} \mathbb{I}_{[k\neq i]} \exp(\sin(\bar{\boldsymbol{z}}_{k}, \bar{\boldsymbol{z}}_{i})/t)}. \tag{1}$$

Diversity Loss. Since diversity is a key component of successful model ensembles [14], we design a new loss function for encouraging diversity during the training of the sub-networks. We define the diversity regularization term ℓ_{div} as a hinge loss over the difference of the standard deviation across the embedding vectors $\{\boldsymbol{z}_{k,m}\}_{m=1}^{M}$, $\{\boldsymbol{z}'_{k,m}\}_{m=1}^{M}$ to a minimum diversity of $\alpha > 0$. The standard deviation is the square root of the element-wise variance $\{\sigma^2_{k,g}\}_{g=1}^{q}$:

$$\sigma_{k,o}^2 = \frac{1}{M-1} \sum_{m=1}^{M} (z_{k,m,o} - \bar{z}_{k,o})^2 + \epsilon ,$$

where we add a small scalar $\epsilon > 0$ to prevent numerical instabilities. The diversity regularization function is then given by:

$$\ell_{\text{div}}\left(\tilde{\boldsymbol{x}}_{k}, \tilde{\boldsymbol{x}}_{k}'\right) = \sum_{o=1}^{q} \max\left(0, \alpha - \sigma_{k,o}\right) + \max\left(0, \alpha - \sigma_{k,o}'\right),$$
(2)

where σ and σ' indicate standard deviation for the input sample and augmented views, respectively.

Total Loss. The objective of the diversity loss is to encourage disagreement among sub-networks by enforcing the element-wise standard deviations to be close to $\alpha > 0$ and to thus prevent the embeddings from collapsing to the same vector. Figure 2a underlines the importance of the diversity loss on the total sum of standard deviations between different sub-networks, which increases by adding this loss. The total loss is calculated by combining the self-supervised loss (Eq. 1) and the diversity loss (Eq. 2), where the degree of regularization is controlled by a tunable hyperparameter $\lambda \geq 0$:

$$\ell\left(\tilde{\boldsymbol{x}}_{k}, \tilde{\boldsymbol{x}}_{k}^{\prime}\right) = \ell_{\rm ssl}\left(\tilde{\boldsymbol{x}}_{k}, \tilde{\boldsymbol{x}}_{k}^{\prime}\right) + \lambda \cdot \ell_{\rm div}\left(\tilde{\boldsymbol{x}}_{k}, \tilde{\boldsymbol{x}}_{k}^{\prime}\right).$$
(3)

Finally, the total loss is aggregated over all the pairs in minibatch X:

$$\mathcal{L}_{\text{total}} = \frac{1}{N} \sum_{k=1}^{N} \ell\left(\tilde{\boldsymbol{x}}_{k}, \tilde{\boldsymbol{x}}_{k}'\right).$$
(4)

Gradients. Consider the output of the encoder $f_{\theta}(x) = b$ and the output of the m-th linear sub-network $z_m = g_{\phi_m}(b) = w_m \cdot b$. The weight w_m is updated by two components during backpropagation, the first of which depends on the self-supervised loss and is the same for the entire ensemble, while the second term depends on the diversity loss and is different for each sub-network. Given Eq. 2, we simplify the equation by vector-wise multiplication since the sub-networks are linear; furthermore, we omit the numerical stability term since it does not have an effect on the derivative. The element-wise standard deviation can be computed as follows:

$$\sigma_{k,o} = \left(\frac{1}{M-1} \sum_{m=1}^{M} (\boldsymbol{z}_{k,m,o} - \bar{\boldsymbol{z}}_{k,o})^2\right)^{\frac{1}{2}}.$$
 (5)

Consider Eq. 2 for aggregating the element-wise standard deviations for one observation (\boldsymbol{x}) and assume $\sigma_k < \alpha$; otherwise, the diversity loss is zero. The derivative of the loss with respect to $\boldsymbol{z}_{k,\hat{m},o}$, $\hat{m} \in 1, \ldots, M$, is then given as follows:

$$\frac{\partial \left(\ell_{\rm div}\right)}{\partial \boldsymbol{z}_{k,\hat{m},o}} = \frac{-A}{M-1} \cdot (\boldsymbol{z}_{k,\hat{m},o} - \bar{\boldsymbol{z}}_{k,o}),\tag{6}$$

where $A := \frac{1}{M-1} \sum_{m=1}^{M} (\boldsymbol{z}_{k,m,o} - \bar{\boldsymbol{z}}_{k,o})^2)$. The proof is provided in the appendix (see Theoretical Supplement).

In the optimization step of stochastic gradient descent (SGD), the weight of sub-network \hat{m} is updated by:

$$\eta \cdot \nabla_{w_{\hat{m},o}} \ell_{\text{div}} = -C \cdot (\boldsymbol{z}_{k,\hat{m},o} - \bar{\boldsymbol{z}}_{k,o}), \tag{7}$$

where $\eta > 0$ is the learning rate, and C is constant with respect to $w_{\hat{m},o}$, which depends on the learning rate, number of sub-networks, A, and b. The proof is provided in Appendix (see Theoretical Supplement).

Equation 7 shows the updating step in backpropagation. Hyperparameter α prevents $\mathbf{z}_{k,\hat{m},o}$ from collapsing to a single point. Hence, $w_{\hat{m},o}$ is updated in the opposite direction of $\bar{\mathbf{z}}_{k,o}$, so the diversity loss prevents weights in the subnetworks from converging to the same values.

3.2 Empirical Analysis of Diversity

Diversity of ensemble members is an important feature for powerful model ensembles and reflects the degree of independence among its members [34, 49]. We follow [14] to quantify the diversities among the ensemble of sub-networks. Specifically, we report the diversities in terms of *disagreement score* between the members' predictive distributions and a baseline. Diversity disagreement



Fig. 2. (a) Total standard deviation: sum of all standard deviations between independent sub-networks during training. Training with diversity loss (Eq. 2) increases the standard deviation and improves the diversity between independent sub-networks. (b) **Diversity analysis**: prediction diversity disagreement vs. achieved accuracy on CIFAR-10. Our method is on par with the deep self-supervised ensemble in terms of both accuracy and diversity disagreement. Models in the top right corner are better.

is defined as distance disagreement divided by 1 - accuracy, where the distance disagreement between two classification models h_i and h_j is calculated as $\frac{1}{N}\sum_{k=1}^{N} [h_i(\mathbf{x}_k) \neq h_j(\mathbf{x}_k)]$, with N denoting the number of samples. Figure 2b compares the diversity disagreement between our method with 10-sub-networks, a deep ensemble with 10 members, and the single-network baseline. The results clearly indicate that our proposed method achieves comparable results with deep self-supervised ensembles in terms of both accuracy and diversity disagreement.

3.3 Computational Cost and Efficiency Analysis

We analyze the efficiency of our proposed method in Table 1. SSL-Ensemble increases memory and computational requirements compared to the baseline by 200% and 900% for 3 and 10 members, respectively. The increase in the number of parameters is 32% and 143%, and the increase in computational requirement is $\sim 0 - 6\%$ for our method.

Method	Members	Parameters(M)	Memory/GPU	Time/800-ep.
Baseline (SSL)	1	28	9 G	3.6 (h)
SSL-Ensemble	3	3×28	$3 \times 9 \text{ G}$	3×3.6 (h)
SSL-Ensemble	10	10×28	$10 \times 9 \text{ G}$	10×3.6 (h)
Our method	3	37	9.2 G	3.6 (h)
Our method	10	68.1	10 G	3.8 (h)

Table 1. Computational cost in 4 DGX-A100 40G GPUs (PyTorch) on CIFAR 10.

4 Experimental Setup

We perform several experiments with a variety of self-supervised methods to examine our hypothesis for robustness during both pretext-task learning and downstream tasks (fine-tuning).

Deep Self-supervised Network Architecture. Our proposed approach builds on two recent popular self-supervised models in computer vision: i) **Sim-CLR** [5] is a contrastive learning framework that learns representations by maximizing agreement on two different augmentations of the same image, employing a contrastive loss in the latent embedding space of a convolutional network architecture (e.g., ResNet-50 [22]), and ii) **DINO** [3] is a self-distillation framework in which a student vision transformer (ViT; [11]) learns to predict global features from local image patches supervised by the cross-entropy loss from a momentum teacher ViT's embeddings. Furthermore, we study the impact of our approach in NLP and modify **SCD** [26], which applies the bidirectional training of transformers to language modeling. Here, the objective is self-supervised contrastive divergence loss. Lastly, we examine our approach on **Self-GenomeNet** [18], a contrastive self-supervised learning algorithm for learning representations of genome sequences. More detailed descriptions of the employed configurations are provided in Appendix (see Implementation Details)

Deep Independent Sub-networks. We implement M independent subnetworks on top of the encoder, for which many possible architectures are conceivable. For our experiments on computer vision datasets, we consider an ensemble of sub-network architecture where each network includes a multi-layer perceptron (MLP) with two layers of 2048 and 128 neurons, respectively, with ReLU as a non-linearity and followed by batch normalization [24]. Each sub-network has its own independent set of weights and learning parameters. For the NLP dataset, the projector MLP contains three layers of 4096 neurons each, also using ReLU activation's as well as batch normalization. For the genomics dataset, our ensemble of sub-networks includes one fully connected layer with an embedding size of 256.

Optimization. For all experiments on image datasets based on DINO and SimCLR, we follow the suggested hyperparameters and configurations by the paper [3,5]. Implementation details for pretraining with DINO on the 1000-classes ImagetNet dataset without labels are as follows: coefficients ϵ , α , and λ are respectively set to 0.0001, 0.15, and 2 in Eq. 2, 2, and 3. We provide more details in ablation studies (Sect. 6) on the number of sub-networks and the coefficients λ and α used in the loss function. The encoder network f_{θ} is either a ResNet-50 [22] with 2048 output units when the baseline is SimCLR [5] or ViT-s [12] with 384 output units when the baseline is DINO [3]. The best prediction and calibration performance is achieved when the number of sub-networks is 5. We followed the training protocol and settings suggested by [3].

Datasets. We use the following datasets in our experiments: **CIFAR-**10/100 [27] are subsets of the tiny images dataset. Both datasets include 50,000 images for training and 10,000 validation images of size 32×32 with 10 and 100 classes, respectively. **SVH** [32] is a digit classification benchmark dataset that

contains 600,000 32×32 RGB images of printed digits (from 0 to 9) cropped from pictures of house number plates. **ImageNet** [8], contains 1,000 classes, with 1.28 million training images and 50,000 validation images. For the NLP task, we train on a dataset of 1 million randomly sampled sentences from **Wikipedia articles** [23] and evaluate our models on 7 different semantic textual similarity datasets from the SentEval benchmark suite [6]: **MR** (movie reviews), **CR** (product reviews), **SUBJ** (subjectivity status), **MPQA** (opinion-polarity), **SST-2** (sentiment analysis), **TREC** (question-type classification), and **MRPC** (paraphrase detection). The **T6SS** effector protein dataset is a public real-world bacteria dataset (SecReT6, [29]) with actual label scarcity. The sequence length of the genome sample is 1000nt in all experiments.

Tasks. We examine and benchmark a model's performance on different tasks considering evaluation protocols by self-supervised learning [5] and Plex's benchmarking tasks [44]. Specifically, we evaluate our model on the basis of uncertainty-aware IND generalization, OOD detection, semi-supervised learning, corrupted dataset evaluation (see Sect. 5), and transfer learning to other datasets and tasks (see Appendix: Transfer to Other Tasks and Datasets)

Evaluation Metrics. We report prediction/calibration performance with the following metrics, where upward arrows indicate that higher values are desirable, *et vice versa.* **Top-1 accuracy** \uparrow : share of test observations for which the correct class is predicted. **AUROC** \uparrow : area under the ROC curve arising from different combinations of false-positive and false-negative rates (here: with positive and negative classes referring to being in and out of distribution, respectively) for a gradually increasing classification threshold. **Negative log-likelihood (NLL)** \downarrow : negative log-likelihood of test observations under the estimated parameters. **Expected calibration error (ECE)**; [31] \downarrow : mean absolute difference between accuracy and confidence (highest posterior probability among predicted classes) across equally-spaced confidence bins, weighted by relative number of samples per bin. **Thresholded adaptive calibration error (TACE)**; [33]) \downarrow : modified ECE with bins of equal sample size, rather than equal interval width, and omitting predictions with posterior probabilities falling below a certain threshold (here: 0.01) that often dominate the calibration in tasks with many classes.

Compared Methods. We compare our method to the following contenders. Baseline: self-supervised architectures (i.e., SimCLR, DINO, SCD, or Self-GenomeNet, depending on the task). SSL-Ensemble: deep ensemble comprising a multiple of the aforementioned baseline networks. Monte Carlo (MC) dropout: [15] baseline networks with dropout regularization applied during pretraining of baseline encoder. BatchEnsemble: baseline encoder with BatchEnsemble applied during pretraining.

5 Results and Discussion

In-Distribution Generalization. IND generalization (or *prediction calibration*) quantifies how well model confidence aligns with model accuracy. We perform several experiments on small and large image datasets as well as the genomics sequence dataset to evaluate and compare the predictive performance of our proposed model in IND generalization. Here, the base encoder f_{θ} is frozen after unsupervised pretraining, and the model is trained on a supervised linear classifier. The linear classifier is a fully connected layer followed by softmax, which is placed on top of f_{θ} after removing the ensemble of sub-networks. High predictive scores and low uncertainty scores are desired. Figure 3 illustrates the predictive probability of correctness for our model on CIFAR-10, CIFAR-100, ImageNet, and T6SS datasets in terms of Top-1 accuracy, ECE, and NLL, respectively. Based on Fig. 3, our method achieves better calibration (ECE and NLL) than the deep ensemble of self-supervised models. The discrepancy in performance between our model and the deep ensemble can be explained by various factors, including differences in uncertainty modeling, complexity, and robustness. While the deep ensemble excels in top-1 accuracy, our model's superior ECE and NLL scores indicate better-calibrated and more reliable predictions, which are essential for safety-critical applications and decision-making under uncertainty.



Fig. 3. IND generalization in terms of (a) Top-1 Accuracy (b) ECE (c) NLL averaged over in-distribution on test samples of CIFAR-10/100, ImageNet, T6SS datasets. Here, we compare our method with the ensemble of deep self-supervised networks (SSL-Ens), as well as the baseline.

Out-of-Distribution Detection. OOD detection shows how well a model can recognize test samples from the classes that are unseen during training [16]. We perform several experiments to compare the model generalization from IND to OOD datasets and to predict the uncertainty of the models on OOD datasets. Evaluation is performed directly after unsupervised pretraining without a fine-tuning step. Table 2 shows the AUROC on different OOD sets for our model, baseline, and deep self-supervised ensemble. Our approach improves overall compared to other methods.

Semi-supervised Evaluation. We explore and compare the performance of our proposed method in the low-data regime. Again, the encoder f_{θ} is frozen after self-supervised pretraining, and the model is trained on a supervised linear classifier using 1% and 10% of the dataset. The linear classifier is a fully connected layer followed by softmax. Table 3 shows the result in terms of top-

IND	OOD	Baseline	SSL-Ensemble	Our method
CIFAR-100	SVHN	84.22	84.95	88.00
	Uniform	91.65	90.53	97.57
	Gaussian	90.00	89.42	94.10
	CIFAR-10	74.71	74.80	75.18
CIFAR-10	SVHN	95.03	96.68	97.07
	Uniform	96.73	91.64	99.05
	Gaussian	96.39	93.24	99.24
	CIFAR-100	91.79	91.59	91.87

Table 2. OOD detection. Results reported using AUROC show our method enhancesthe baseline up to 6%.

Table 3. Semi-supervised evaluation: Top-1 accuracy (ACC), ECE, and NLL for semi-supervised CIFAR-10/100 classification using 1% and 10% training examples.

Method	CIFA	R-10	(1%)	CIFA	R-10	(10%)	CIFA	R-10	0 (1%)	CIFA	R-100	0 (10%)
	ACC	ECE	NLL	ACC	ECE	NLL	ACC	ECE	NLL	ACC	ECE	NLL
Baseline	89.1	0.075	0.364	91.1	0.039	0.274	56.2	0.097	2.01	59.5	0.086	1.79
SSL-Ensemble	90.1	0.056	0.334	92.2	0.050	0.257	59.7	0.081	1.86	62.6	0.053	1.48
Our method	90.4	0.018	0.296	92.6	0.016	0.249	59.3	0.060	1.71	62.4	0.042	1.56

1 accuracy, ECE, and NLL. The results indicate that our method outperforms other methods in the low-data regime – in terms of calibration.

Corrupted Dataset Evaluation. Another important component of model robustness is its ability to make accurate predictions when the test data distribution changes. Here, we evaluate model robustness under *covariate shift*. We employ a configuration similar to the one found in [44]. Figure 4 summarizes the improved performance across metrics of interest. The results confirm that our method outperforms the baseline and achieves comparable predictive performance as a deep self-supervised ensemble – both in terms of calibration (TACE) and AUROC.

Transfer to Other Tasks and Datasets. We further assess the generalization capacity of the learned representation on learning a new task in NLP. We train our model without any labels on a dataset of sentences from Wikipedia [23] and fine-tune the pretrained representation on seven different semantic textual similarity datasets from the SentEval benchmark suite [6]: MR (movie reviews), CR (product reviews), SUBJ (subjectivity status), MPQA (opinion-polarity), SST-2 (sentiment analysis), TREC (question-type classification), and MRPC (paraphrase detection). Then, we evaluate the test set of each dataset. Figure 5 provides a comparison of the transfer learning performance of our self-supervised approach for different tasks. Our results in Fig. 5 indicate that our approach performs comparably to or better than the baseline method. We test the per-



Fig. 4. Performance under dataset corruption (CIFAR-10/100 with five levels of increasing perturbation), evaluation in terms of AUROC and TACE for several types of corruption (vertical spread).

formance of the trained model on ImageNet [8] on CIFAR-10 [27] dataset where the model is trained for 100 epochs (Table 4).



Fig. 5. Transfer to other dataset and tasks: Comparision of Sentence embedding performance on semantic textual similarity tasks.

Table 4. Transfer to other dataset:Expected calibration error averaged overuncertainty-aware evaluation on CIFAR-10 datasets.

Method	ACC (%) (\uparrow)	ECE (\downarrow)	NLL (\downarrow)	TACE (\downarrow)
Baseline	73.5	0.038	0.78	0.20
Our method	73.9	0.030	0.75	0.18



Fig. 6. We compare the feature diversity for different subnetworks and ensemble members. The top images are for different sub-networks, and the bottom images are for different ensemble members. We used Grad-CAM [41] for visualization.

6 Ablation Study

In order to build intuition around the behavior and the observed performance of the proposed method, we further investigate the following aspects of our approach in multiple ablation studies exploring: (1) the number M of sub-networks, (2) the role of each component of the proposed loss, and (3) analysis of diversity with visualization of the gradients of subnetworks. We also present more results on (4) the impact of our approach during pretraining vs. at the finetuning step, (5) the size of sub-networks, and (6) the impact of model parameters.

Number of Sub-networks. We train M individual deep neural networks on top of the representation layer. The networks receive the same inputs but are parameterized with different weights and biases. Here, we provide more details regarding our experiments on IND generalization by considering varying M. Figure 7a compares the performance in terms of top-1 accuracy, ECE, and NLL for CIFAR-10 and CIFAR-100. Based on the quantitative results depicted in Fig. 7a, the predictive performance improves in both datasets when increasing the number of sub-networks (M) until a certain point. For example, in the case of CIFAR-10, when M = 3, our performance is 91.9%; increasing M to 10 levels top-1 accuracy up to 92.6%, while the ECE and NLL decrease from 0.026 and 0.249 to 0.023 and 0.222, respectively. These findings underline that training our sub-networks with a suitable number of heads can lead to a better representation of the data and better calibration. Recently [43,47] provided a theoretical statement as well as experimental results that projection heads help with faster convergence.

Analysis of Loss. The total loss (Eq. 3) is calculated by the combination of self-supervised loss (Eq. 1) and diversity loss (Eq. 2), where the mean value of the embeddings across the ensemble of sub-networks is fed to the self-supervised loss, and the corresponding standard deviation is used for the diversity loss. First, we note that the use of our diversity regularizer indeed improves calibration and provides better uncertainty prediction. The results in Fig. 3 show the impact of our loss function in relation to the baseline. Figure 3 compares the predictive probability of the correctness of DINO (baseline) and our model on

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Fig. 7. Ablation study on number of M sub-networks (a), hyperparameters of our proposed loss (b) λ and (c) α .

ImageNet. Second, we explore different hyperparameter configurations to find the optimal values for α and λ in Fig. 7b, 7c. Note that, in practice, α and λ must be optimized jointly. The best top-1 accuracy in our case is achieved when α and λ are set to 0.08 and 1.5, respectively, on the CIFAR-10 dataset.

Analysis of Diversity. In addition to quantitative results for diversity analysis provided in Fig. 2b, we visualize the activation map for the last convolution layer in the encoder for each ensemble member and each subnetwork to motivate the effect of subnetworks on the encoder. As illustrated in Fig. 6, different subnetworks have more feature diversity compared to the deep ensemble as we expected.

Efficient Ensemble of Sub-networks at Pretraining vs. Finetuning We performed additional experiments to study the efficiency of proposed loss and independent sub-networks (InSub) i) during pretraining, ii) during finetuning, and iii) during both pretraining and finetuning. As shown in Table 5, pretraining with an ensemble of sub-networks is beneficial, and additional fine-tuning with multiple heads can further improve performance.

Table 5. Pretraining vs. Finetuning:	Expected calibration error averaged over
uncertainty-aware evaluation on CIFAR-10	datasets. InSub refers to training with our
proposed \underline{In} dependent \underline{Sub} networks	

Method	ACC (%) (\uparrow)	ECE (\downarrow)	NLL (\downarrow)	TACE (\downarrow)
Baseline	92.5	0.039	0.238	0.133
Pretrain-InSub	92.6	0.032	0.226	0.131
Finetune-InSub	92.6	0.021	0.222	0.103
$\overline{ ext{Pretrain-InSub} + ext{Finetune-InSub}}$	92.8	0.023	0.227	0.115

 Table 6. Sub-Network Size: Expected calibration error averaged over uncertaintyaware evaluation on CIFAR-10 datasets.

Method	ACC (%) (\uparrow)	ECE (\downarrow)	NLL (\downarrow)
Our method with 5 sub-network (100%)	92.9	0.019	0.221
With 25 percent of sub-network size	92.3	0.026	0.231
With 50 percent of sub-network size	92.6	0.021	0.226
With 75 percent of sub-network size	92.6	0.019	0.221

Table 7. Large variant encoder:Expected calibration error averaged overuncertainty-aware evaluation on CIFAR-10 datasets.

Method	ACC (%) (†)	ECE (\downarrow)	NLL (\downarrow)	Number of parameters (M)
Our method with ResNet50 as a encoder with 5 sub-networks	92.9	0.019	0.221	45.79
Baseline with ResNet101 as a encoder	93.2	0.027	0.202	46.95

Analysis of Size of Sub-networks. We perform several experiments to study the different sizes of sub-network. As shown in Table 6, the dimension of projection heads does not change the top-1 accuracy. Recent self-supervised models such as SimCLR [5], BarlowTwins [48] also reach the same results with different projection head sizes.

Impact of Model Parameters. Our project aims to improve the predictive uncertainty of the baseline without losing predictive performance by mimicking the ensembles of self-supervised models with much lower computational costs. According to the results shown in Table 7, a bigger encoder can potentially improve the predictive performance, but it does not necessarily improve the predictive uncertainty of the results. We used ResNet101 as a baseline with more parameters in the encoder. To have a fair comparison, we compare it with our model with five heads. Our model performs better in ECE and NLL and has comparable accuracy. Also, we used ResNet34 as a baseline with fewer parameters in the encoder with twenty heads and compared it with baseline ResNet50 with one head. According to results obtained in Table 8, our model performs better in terms of ECE and NLL and has on-par accuracy.
 Table 8. Different encoder (medium size): Expected calibration error averaged

 over uncertainty-aware evaluation on CIFAR-10 datasets.

Method	ACC (%) (\uparrow)	ECE (\downarrow)	NLL (\downarrow)	Number of parameters (M)
Our method with ResNet34 as a encoder with 20 sub-networks	92.5	0.016	0.23	27.84
Baseline with ResNet50 as a encoder	92.8	0.039	0.233	27.89

7 Conclusion

In this paper, we presented a novel diversified ensemble of self-supervised framework. We achieved high predictive performance and good calibration using a simple yet effective idea – an ensemble of independent sub-networks. We introduced a new loss function to encourage diversity among different sub-networks. Our method is able to produce well-calibrated estimates of model uncertainty at low computational overhead over a single model while performing on par with deep self-supervised ensembles. It is straightforward to add our method to many existing self-supervised learning frameworks during pretraining. Our extensive experimental results show that our proposed method outperforms, or is on par with, an ensemble of self-supervised baseline methods in many different experimental settings.

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Modular Debiasing of Latent User Representations in Prototype-Based Recommender Systems

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Abstract. Recommender Systems (RSs) may inadvertently perpetuate biases based on protected attributes like gender, religion, or ethnicity. Left unaddressed, these biases can lead to unfair system behavior and privacy concerns. Interpretable RS models provide a promising avenue for understanding and mitigating such biases. In this work, we propose a novel approach to debias interpretable RS models by introducing user-specific scaling weights to the interpretable user representations of prototype-based RSs. This reduces the influence of the protected attributes on the RS's prediction while preserving recommendation utility. By decoupling the scaling weights from the original representations, users can control the degree of invariance of recommendations to their protected characteristics. Moreover, by defining distinct sets of weights for each attribute, the user can further specify which attributes the recommendations should be agnostic to. We apply our method to PROTOMF, a state-of-the-art prototype-based RS model that models users by their similarities to prototypes. We employ two debiasing strategies to learn the scaling weights and conduct experiments on ML-1M and LFM2B-DB datasets aiming at making the user representations agnostic to age and gender. The results show that our approach effectively reduces the influence of the protected attributes on the representations on both datasets, showcasing flexibility in bias mitigation, while only marginally affecting recommendation quality. Finally, we assess the effects of the debiasing weights and provide qualitative evidence, particularly focusing on movie recommendations, of genre patterns identified by PROTOMF that correlate with specific genders.

Keywords: Recommender Systems \cdot Debiasing \cdot Interpretability

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1 Introduction

Recommender Systems (RSs) typically operate as black boxes trained on large collections of user-item interactions to generate recommendations. Through this training process, they capture underlying interaction patterns, revealing which users prefer which content, to better model the users' interests. Alas, the observed user behavior might correlate with particular protected user attributes such as gender, age, ethnicity, or religion, even when these are not explicit in the data [12]. When exposed to such data, the RS can encode these correlations in the user representations, potentially leading to biased predictions [17,33,46], unfair system behavior across protected groups [11,20], and the strengthening of per-group "filter bubbles" [13,29]. Furthermore, they can also raise privacy concerns regarding the disclosure of sensitive information from the representations [4,36].

Interpretable RS models can be leveraged to understand how these biases manifest in the data [3,19] and how they are assimilated by the RS [15,38]. In recent years, several RS models that offer interpretable user representations have emerged. Specifically, each dimension of these representations usually corresponds to an interpretable aspect, such as the user's sentiment towards items' attributes [45], user or item features [16,37], or similarity to prototypical users/items [1,34]. These transparent models can assist in defining potential corrective measures. For instance, if a particular dimension strongly correlates with a user's protected attribute, we can choose to weaken it and use the updated representations to generate debiased recommendations. Alternatively, we may also amplify another dimension associated with a different value of the protected attribute, thereby increasing the ambiguity surrounding the true user's attribute. However, determining which dimensions are indicative of the attribute in the first place can still pose challenges.

One solution, explored in recent literature, is to adapt the user representations, predominantly through in-processing techniques [9,11]. These methods involve (re-)training a RS model to provide relevant recommendations while also optimizing a debiasing objective that attempts to make the predictions invariant to the user's protected attributes, albeit with a trade-off in performance [4,17,29,46]. However, depending on the user's preferences, the context, or biasutility trade-off considerations, end-users might in practice still prefer to receive *some* recommendations from the original (potentially biased) model. Especially when different users have different attitudes towards their biased representations (e.g., users conforming to stereotypical norms may prefer biased predictions), or when the same user prioritizes having their recommendations unbiased with respect to certain attributes but not others [29]. Accommodating all these scenarios with current approaches can be burdensome, as it requires training a separate RS for every protected attribute and according to each user's request.

In contrast, we propose learning separate user-specific scaling weights that can be applied to the interpretable user representations of a pre-trained RS model. These modular weights automatically adjust the representations to reduce their biases associated with a protected attribute, e.g., gender or age, while still preserving relevant recommendations. This concept aligns with recent efforts in the NLP community focused on modular bias mitigation, enabling endusers to choose whether their results should be biased or unbiased on-demand [25,26,32]. Our method allows to flexibly cover different users' needs. By keeping the weights separate from the original representations, users can decide during inference whether their recommendations should be influenced by their protected attributes, applying the scaling weights as needed. Additionally, by training distinct sets of weights for each attribute of interest, users can further specify with respect to which of them the recommendations should be agnostic.

We apply our approach to the recently proposed PROTOMF model [34], a prototype-based RS designed to capture specific item-consumption characteristics of the data through the concept of user/item prototypes [23,27]. We select this model for its ability to provide relevant and explainable recommendations; nevertheless, our method can be applied to any interpretable RS that provides interpretable user representations. Within PROTOMF, each user is mapped to a representation where each dimension indicates the similarity between the user and a specific user prototype. The application of the scaling weights, hence, tunes these similarities, thereby influencing the impact of the prototype's pattern on the resulting recommendations. Consequently, analyzing which dimensions are attenuated (< 1) or amplified (> 1) by the debiasing strategy aids us in interpreting which consumption patterns might be correlated (and in which way) to a specific protected attribute.

We evaluate our method on two popular datasets of movie ratings (ML-1M [24]) and music listening records (LFM2B-DB [35]). For both datasets, we learn user representations that are less affected by the user's gender and age, which aligns with the concepts of representational fairness [40] or demographic *parity* [2]. Intuitively, if the representations are invariant to these attributes, predictions based on these representations will also be invariant, resulting in less biased recommendations [2]. For instance, the RS will avoid recommending only Romance movies to female users. Our approach is agnostic to the debiasing objective, allowing the scaling weights to be trained with any gradient descent-based signal that ensures representation invariance to a specific user's attribute. In this study, we investigate two debiasing objectives: Maximum Mean Discrepancy [22] and Adversarial Debiasing [21,43]. To assess the effectiveness of our approach to mitigate bias, we follow the standard evaluation framework for debiasing [14, 26, 29, 32] and report the performance of an external probe network trained to predict the protected attribute from the user representations. Compared to the original user representations, our results show that our proposed method effectively impairs the probe's ability to recover sensitive information, resulting in a substantial reduction in bias while only marginally affecting recommendation performance. Finally, we investigate the effect of the debiasing weights and showcase for ML-1M the genre patterns captured by PROTOMF that are correlated with gender. Our code and settings are publicly available at https://github.com/hcai-mms/modprotodebias.

2 Related Work

Our research is influenced by recent works in NLP. Thus, we review pertinent literature in this field before delving into related research on debiasing RSs.

Bias Mitigation in Natural Language Processing. Extensive research has addressed societal biases within Language Models (LMs), particularly focusing on attribute erasure [33]. This involves reducing the influence of protected attributes within LM's embeddings to mitigate empirical biases [33, 40] or achieve representational fairness [14]. Recent studies explore modular bias mitigation, enabling end-users to select between biased or bias-mitigated models for individual queries. In particular, Hauzenberger et al. [25] learn a set of sparse additive weights that mitigate societal bias when added to the original model. Kumar et al. [26] leverage adapters [39] to isolate the sensitive information in separate blocks of the LM. Masoudian et al. [32] introduce controllable gates to scale LM's representations to switch between biased/unbiased predictions. Inspired by these studies, our work introduces separate per-user scaling weights to adjust user representations for unbiased recommendations.

Bias Mitigation in Recommender Systems. Being multi-sided platforms, RSs' outcomes may be prone to biases associated with the users [8, 42] and items [5, 5]44]. While there are several strategies to mitigate these biases and increase the RSs' fairness, recent literature especially focuses on in-processing techniques [11,12]. Zhu et al. [47] tackle the issue of item under-recommendation from imbalanced train data and propose a regularization objective based on fairness. Similarly from the user side, Li et al. [28] propose a novel RS model to learn user/item representations that avoid unfairly penalizing non-mainstream users. Several studies focus on removing spurious correlations between users' protected attributes and recommendations by leveraging adversarial learning, albeit at some performance trade-off. For instance, Bose and Hamilton [6] and Wu et al. [42], learn user/item representations in graph-based RSs that are invariant to the user's protected attribute. Ganhör et al. [17] adapt Mult-VAE [30] to generate recommendations agnostic to users' gender. Li et al. [29] simultaneously train a set of filters, one for each attribute, as well as the underlying RS, to satisfy different users' fairness demands. Some authors also leveraged interpretable models to assess fairness issues in RSs. Ge et al. [19] employ counterfactual learning to learn the minimal change to the input features of a feature-aware RS to address item exposure unfairness in the recommendations. Fu et al. [15] present a fairness re-ranking approach to decrease performance disparity between active/inactive users in explainable recommendations over knowledge graphs. Our work complements the above studies by addressing the influence of users' protected attributes on the recommendations of a *pre-trained* RS, leveraging modular scaling weights on the interpretable user representations concerning user prototypes.

3 Methodology

Let $\mathcal{U} = \{u_i\}_{i=1}^N$ and $\mathcal{T} = \{t_j\}_{j=1}^M$ denote the set of N users and M items, respectively. We assume that we only have access to the implicit interaction data $\mathcal{I} = \{(u_i, t_i)\}$, where (u_i, t_i) indicates that user u_i has interacted with item t_i . Additionally, each user u_i is associated with one or more protected attributes $q \in G$. We omit the user and item indexes for brevity. Let $rec(\cdot, \cdot)$ be an interpretable RS model that, beyond scoring each user-item pair $rec(u, t) \in \mathbb{R}$, also maps each user u to an intermediate interpretable representation $u \in \mathbb{R}^d$. This representation may align with various aspects, such as user's sentiment towards items' attributes [45], user or item features [16, 37], or similarity to prototypical users and items derived from the dataset [1,34]. In this work, we focus on the latter and particularly the recently proposed PROTOMF model [34] as it showcased high accuracy in the recommendation task. Nevertheless, our method can be applied to any RS offering interpretable user representations. Within PROTOMF, each dimension $\{i\}_{i=1}^d$ in u indicates the similarity of user u to a specific user prototype p^i , representing item-consumption characteristics of the data, with similarity values in the range (0, 2). As shown next, the interpretable representation \boldsymbol{u} may encode the protected user attribute q, despite the information not being explicitly provided to the RS. As a consequence, the RS can pick up this information and bias its predictions, as also shown in [17, 29, 46].

To address this issue, we define a vector of scaling weights $\boldsymbol{\omega}_u \in \mathbb{R}^d$ for each user, which can be plugged in at will. Starting from the \boldsymbol{u} representation obtained from the pre-trained RS model, we derive a new user representation $\tilde{\boldsymbol{u}}$ as follows:

$$ilde{oldsymbol{u}} = oldsymbol{u} \odot oldsymbol{\omega}_u$$

where \odot is the Hadamard product. We leave the original user representation \boldsymbol{u} (as well the other model parameters) unchanged while we only optimize $\boldsymbol{\omega}$ so that the new representation $\tilde{\boldsymbol{u}}$ remains relevant for the recommendation task while becoming invariant to the protected attribute g. The optimization involves minimizing a recommendation loss \mathcal{L}_{rec} as well as a debiasing objective \mathcal{L}_{debias} :

$$\boldsymbol{\omega}^* = \operatorname*{arg\,min}_{\boldsymbol{\omega}} \mathcal{L}_{\mathrm{rec}}(\mathcal{I}, \boldsymbol{\omega}) + \lambda \mathcal{L}_{\mathrm{debias}}(\mathcal{I}, \boldsymbol{\omega}, g)$$

where the hyperparameter λ adjusts the strength of the debiasing loss. As recommendation loss \mathcal{L}_{rec} , we adopt the same loss function as the base RS model. In the case of PROTOMF [34], this corresponds to the cross-entropy loss reported below for reference:

$$\mathcal{L}_{\text{rec}} = -\sum_{(u,t)\in\mathcal{I}} \ln p(t|u), \quad p(t|u) = \frac{e^{rec(u,t)}}{\sum_{j} e^{rec(u,t_j)}}$$
(1)

The debiasing objective $\mathcal{L}_{\text{debias}}$ operates on the representations \tilde{u} and the corresponding protected attribute label g to realize invariance. Our approach is agnostic to the debiasing objective, allowing the scaling weights to be trained

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with any gradient descent-based signal that ensures representation invariance. In our work, we employ two prominent debiasing strategies: Maximum Mean Discrepancy (MMD) [22,25] and Adversarial Debiasing (Adv.) [14,43].

Maximum Mean Discrepancy (MMD) [22] aims to minimize the distribution shift between the representations of a specific protected attribute g. Effectively, given the set of users \mathcal{U} split into two subsets \mathcal{U}_g^A and \mathcal{U}_g^B according to the values of a binary¹ protected attribute g, MMD minimizes the mean distance between the user representations $\tilde{\boldsymbol{u}}$ of the two subgroups:

$$\mathcal{L}_{\text{debias}} = \left\| \frac{1}{|\mathcal{U}_g^A|} \sum_{i \in \mathcal{U}_g^A} \phi(\tilde{\boldsymbol{u}}_i) - \frac{1}{|\mathcal{U}_g^B|} \sum_{j \in \mathcal{U}_g^B} \phi(\tilde{\boldsymbol{u}}_j) \right\|_2^2$$
(2)

where ϕ is a feature map kernel defined as a sum of multiple Gaussian kernels.

Adversarial Debiasing (Adv.) [21,43] is a common approach in learning input representations that are informative for the task while remaining invariant to specific traits of the data [14,17]. In our context, each user is passed through an adversarial head $h(\cdot)$ that aims to infer the protected attribute g from \tilde{u} by leveraging the cross-entropy loss $\mathcal{L}_{debias}(\tilde{u},g) = \mathcal{L}_{CE}(\tilde{u},g)$. During training, we aim to learn scaling weights ω that maintain relevant user recommendations while hindering the adversary's predictive ability. This objective is commonly solved as a minimization task by inserting a gradient reversal layer $grl(\cdot)$ between the adversary and the rest of the model [18,43]. Essentially, during back-propagation, the $grl(\cdot)$ negates and potentially scales the gradients flowing from the adversary to the weights, pushing the ω in the opposite direction desired by the adversary. This allows us to formulate the debiasing objective as:

$$\mathcal{L}_{\text{debias}} = \mathcal{L}_{\text{CE}}(grl(\tilde{\boldsymbol{u}}), g) \tag{3}$$

Finally, given the learned scaling weights $\boldsymbol{\omega}$, we derive the adjusted user representations $\tilde{\boldsymbol{u}}$, which are used by the RS model to provide item recommendations that are both relevant and agnostic to the user's protected attribute. Our proposed approach offers a flexible and informative method for debiasing. By keeping the weights separate from the original representations, users can decide during inference whether their recommendations should be influenced by their protected attributes, applying the scaling weights as needed. Using distinct sets of weights for each attribute of interest, users can further specify with respect to which of them the recommendations should be agnostic. Moreover, by analyzing which interpretable dimensions are attenuated ($\omega < 1$) or amplified ($\omega > 1$), we can assess which consumption patterns might be correlated (and in which way) to a specific protected attribute.

 $^{^{1}}$ We consider majority vs. all others subsets for non-binary attributes.

ML-1M		Users I	Interactions Items	LFM2	B-DB	Users	Interactions	Items
All		6,034	$574,\!376$	All		16,258	2,339,540	
Gender	М	4,326	429,039	Gender	М	12,734	1,981,006	
	F	1,708	145,337		F	3,524	358,534	
Age	< 18	222	$15,\!583$	Age	≤ 18	1,811	232,942	99,824
	18-24	1,100	100,655 $3,125$					
	25-34	2,095	222,242		19-32	12,613	1,797,291	
	35-44	1,192	$116,\!507$					
	45-49	550	49,400		33-39	1,126	184,176	
	50-55	496	$44,\!979$					
	> 56	379	$25,\!010$		> 40	708	$125,\!131$	

Table 1. Statistics of the datasets used in our experiments.

4 Experiment Setup

Datasets. We use two standardized datasets containing user-item interactions along with partial user's demographic: (1) MovieLens-1M² (ML-1M) [24] contains the ratings of users on movies as well as user's gender, age group, and occupation. As common [30,34], we treat high movie ratings (> 3.5 on a 1–5 scale) as positive interactions while discarding the rest, and perform 5core filtering. (2) LFM2B-DemoBias (LFM2B-DB) [35] is a sub-set of the LFM2B³ dataset, which provides a collection of music listening records of users for whom partial demographic information (i. e., gender, age, country) is available. We follow the same data processing methodology as in Melchiorre et al. [35]. Specifically, we keep user-item interactions with a minimum play count of two and binarize the interactions. Additionally, to accommodate computational constraints, we randomly sample 100,000 tracks from the large catalog and apply 5-core filtering. Furthermore, we split users into age groups based on their deviation from the mean age ($\mu = 24.87$, $\sigma = 7.30$) by multiples of σ .

Table 1 offers a detailed summary of the dataset statistics, including the breakdown by user attribute. With both datasets, we focus on the gender and age of the user as protected attributes in our experiments.⁴

Data Splits. To train both the underlying RS model and the scaling weights, we employ the leave-k-out strategy [10] for every user. Specifically, for each user, we sort their item interactions according to the timestamps (keeping the earliest interaction if multiple ones with the same item exist). The last 10%

² https://grouplens.org/datasets/movielens/1m/.

³ http://www.cp.jku.at/datasets/LFM-2b/.

⁴ Both datasets provide gender in binary form, neglecting nuanced gender definitions.

interactions of the users are used as test, while the penultimate 10% as validation set. The remaining interactions constitute the training set. During training, for each positive user-item interaction, we randomly sample 10 negative items not interacted with by the user. We scale both the adversary's and, later, the probe's loss, ensuring that data points from all user groups contribute equally. This balancing not only aids debiasing [35] but also prevents both classifiers from solely predicting the majority class [14].

PROTOMF *Pre-training*. We follow a similar training procedure as the original UI-PROTOMF paper does [34], referred here simply as PROTOMF. We train the model for 50 and 100 epochs on ML-1M and LFM2B-DB, respectively, with the AdamW optimizer [31]. We perform early-stopping if the accuracy on the validation set does not improve for 5 consecutive epochs. After preliminary experiments, we set the number of user prototypes based on the dataset (42 for ML-1M and 64 for LFM2B-DB) and fix the batch size to 256. We then carry out a comprehensive search for optimal embedding sizes and loss-related hyperparameters. Details on the range of hyperparameters explored, as well as those selected for the final models, are provided in the appendix. Once we identify the model that achieves the highest accuracy on the validation set, we freeze its parameters and only update the scaling weights during debiasing.

Evaluation. To assess the effectiveness of our approach to bias mitigation, we follow the standard evaluation framework [14]. Specifically, after freezing the model's parameters (including the ω), we train a probe network to predict the protected attributes from the user representations. We measure the accuracy (Acc) and balanced accuracy (BAcc) when predicting the users' gender and age. Particularly, we focus on the BAcc metric [7] as it is well-suited for imbalanced datasets. BAcc reports the average recall per user group, where a value of $\frac{1}{\#\text{Groups}}$ represents a fully debiased representation which amounts to .50 for gender on both datasets and $\frac{1}{7} = .14$ and $\frac{1}{4} = .25$ for age on ML-1M and LFM2B-DB, respectively. To evaluate recommendation performance, we use Normalize Discount Cumulative Gain (NDCG), specifically NDCG@10. We report performance and bias mitigation results as average computed on the test set for three seeds.

Debiasing and Probing. For the MMD method, we use a batch size of 128 and set the learning rates to $5e^{-5}$ for gender and $5e^{-4}$ for age on both datasets. In the Sect. 5, we explore different values of λ . Regarding adversarial debiasing, instead, we employ a two-layer neural network with 512 neurons as an adversary network. We investigate the impact of using multiple adversarial networks, i. e., adversarial heads, by averaging their debiasing losses [14]. We use a batch size of 512 for ML-1M and 1024 for LFM2B-DB, with a learning rate of $5e^{-5}$, adjusting λ based on the dataset and attribute.⁵ Our probe is a two-layer neural network with 128 neurons in the hidden layer. We set the learning rate and weight decay based on the probe's performance on each dataset and attribute.

⁵ $\lambda = 1$ on LFM2B-DB, $\lambda = 5$ and $\lambda = 10$ on ML-1M for gender and age respectively.

After debiasing, we train a new probe using the debiased user representations while keeping the scaling weights (and the base model) unchanged. Finally, we initialize the ω by sampling from the normal distribution $\mathcal{N}(1, .01^2)$ and train them, as well as the probe, for 25 epochs using the AdamW optimizer [31].

5 Results and Analysis

Table 2. Debiasing and performance results on both datasets and attributes. We highlight the least biased and best-performing values among Adv. and MMD. Subscripts indicate the standard deviation.

Dataset	Attribute	Debiasing	Bia BAcc	ıs↓ Acc	NDCG \uparrow
ML-1M	Gender	None MMD Adv.	.789 ₀₀₃ .542 ₀₀₃ .542 ₀₀₈	.788 ₀₀₁ .497 ₀₂₅ .608 ₀₄₇	.0625 ₀₀₀₀ .0618 ₀₀₀₀ . 0620₀₀₀₀
	Age	None MMD Adv.	.465 ₀₀₂ .232 ₀₀₁ .232 ₀₂₆	.424 ₀₀₃ .207 ₀₀₄ .193 ₀₁₁	.0625 ₀₀₀₀ .0594 ₀₀₀₂ .0618 ₀₀₀₁
LFM2B-DB	Gender	None MMD Adv.	.723 ₀₀₂ .536 ₀₀₄ .600 ₀₁₁	.718 ₀₀₄ .397 ₀₇₈ .636 ₀₆₅	.0754 ₀₀₀₀ .0745 ₀₀₀₁ .0755 ₀₀₀₁
	Age	None MMD Adv.	.581 ₀₀₅ . 299₀₀₃ .390 ₀₁₁	.504 ₀₀₈ .238 ₀₅₅ .277 ₀₁₈	.0754 ₀₀₀₀ .0626 ₀₀₀₁ .0755 ₀₀₀₁

General Results. Table 2 reports the results of the debiasing methods and recommendation utility across datasets and attributes. We highlight in bold the best RS performance (highest NDCG) and best debiasing performance (lowest BAcc of the probing network). Results are computed on the test set as the average of 3 random seeds, with subscripts indicating the standard deviation.

When no debiasing is applied (*None* rows in Table 2), we observe that the users' protected attributes can be predicted with relatively high accuracy by the probe. The BAcc for gender reaches .79 and .72 on ML-1M and LFM2B-DB datasets respectively, compared to a baseline value of .50 of a random predictor. Similar observations can be made for age; on ML-1M the probe's BAcc is .47 against the baseline of .14 and on LFM2B-DB is .58 vs. .25 in a bias-free settings. These results indicate that the user representations \boldsymbol{u} learned by the RS

D-++	A + +	Metric		λ						
Dataset	Autr.			0	2	5	10	15	20	
ML-1M	ender	Bias $\downarrow \frac{B}{A}$	Acc Acc	.789 _{.003} .788 _{.001}	$.633_{.010}$ $.631_{.011}$.574 _{.010} .557 _{.027}	$.548_{.002}$ $.552_{.012}$.548 _{.003} .514 _{.030}	$.542_{.003}$ $.497_{.025}$	
	Ğ	NDCG	; †	$.0625_{.0000}$.0624.0000	.0623.0000	.0621.0001	.0619.0001	.0618.0000	
	Age	$\operatorname{Bias} \downarrow^{\operatorname{B}}_{A}$	Acc Acc	$.465_{002}$ $.424_{003}$	$.463_{.006}$ $.356_{.003}$	$.398_{.004}$ $.305_{.004}$	$.320_{.007}$ $.252_{.005}$	$.258_{.005}$ $.215_{.006}$	$.232_{.001} \\ .207_{.004}$	
		NDCG	¦↑	.06250000	.0626.0002	.0619.0002	.0609.0002	.0600.0001	.0594.0002	
LFM2B-DB	Gender	Bias \downarrow B	Acc Acc	.723 _{.002} .718 _{.004}	.607 _{.001} .588 _{.003}	$.567_{.008}$ $.469_{.065}$.551 _{.005} .428 _{.074}	.538 _{.003} .357 _{.014}	.536 .004 .397 _{.078}	
		NDCG	+	$.0754_{.0000}$	$.0756_{.0000}$	$.0755_{.0001}$	$.0752_{.0000}$.0749.0001	$.0745_{.0001}$	
	Age	Bias \downarrow^{B}_{A}	Acc Acc	$.581_{005}$ $.504_{008}$.639 _{.013} .517 _{.021}	.548 _{.006} .406 _{.010}	.406 _{.008} .232 _{.007}	.327 _{.008} .182 _{.006}	.299 .003 .238.055	
		NDCG	¦ ↑	$.0754_{0000}$	$.0755_{.0002}$	$.0752_{.0002}$	$.0697_{.0003}$	$.0638_{.0001}$	$.0626_{.0001}$	

Table 3. Debiasing and performance results on both datasets and attributes using the MMD method across several λ values.

do retain information about the user's protected attributes and can potentially bias the recommendations.

When applying the scaling weights, either learned by MMD or Adv., we observe a substantial decrease in both Acc and BAcc of the probe. This reduction spans across both attributes and datasets, indicating the effectiveness of our proposed method in weakening the attribute information in the new user representations. We observe that the efficacy and the impact of the ω depends on the dataset under scrutiny. On the ML-1M datasets, MMD and Adv. reach similar BAcc values for age and gender, both resulting in a moderate decrease in NDCG. However, Adv. shows higher capability in preserving the recommendation performance compared to MMD. On the LFM2B-DB dataset, the scaling weights learned by MMD display lower bias, although at a larger trade-off in recommendation performance. The Adv. method, on the other hand, appears to fully preserve the initial NDCG while leading to a smaller decrease in BAcc compared to MMD. Considering these results, we derive that (1) the debiased user representations, obtained by either MMD or Adv., exhibit significant decreases in the bias metrics, although the predictions are not yet fully random (e.g., Gender BAcc > .50, and (2) there exists a trade-off between bias reduction and recommendation accuracy whose strength depends on the dataset. We investigate the latter aspect below.



Fig. 1. BAcc and NDCG on LFM2B-DB using MMD, over varying λ values. In Fig. 1a, BAcc refers to gender, in Fig. 1b to age.

Bias vs. Performance Analysis. Considering the MMD method, we report in Table 3 the bias metrics and recommendation accuracy across different values of λ ranging from 0 (no debiasing is applied) to 20 for ML-1M and LFM2B-DB on both attributes. We plot the changes of BAcc and NDCG over the λ 's for age and gender on LFM2B-DB in Fig. 1. We observe that high λ values lead to a stronger debiasing of the user representations, i. e., lower BAcc, however at the cost of a moderate reduction of NDCG. We also notice that, on both datasets, making the representations agnostic to age leads to a harsher reduction of recommendation accuracy compared to debiasing for gender. Lastly, we observe for $\lambda = 2$ on the age attribute in LFM2B-DB, associated with a milder debiasing, the BAcc even increases, suggesting that without proper debiasing, more bias information can be encoded in the user representations through the ω by the recommendation loss.

Regarding the Adv. method, preliminary experiments with the Adv. method showed that using a single adversary head while increasing λ led to unstable debiasing behavior, causing the adversary to fail and more bias to be encoded in the scaling weights. To address this, we followed previous research on adversarial debiasing [14,26,32] and opted to use multiple adversary heads while fixing λ based on the dataset and attribute (see Sect. 4). Table 4 displays bias/recommendation performance across different numbers of adversarial heads. Similarly to the MMD method, we observe lowest bias with a stronger debiasing approach, namely 20 heads. By increasing the # of heads, we see a progressive reduction in NDCG on ML-1M for both age and gender, while the recommendation performance on LFM2B-DB remains relatively constant.

In summary, we find that (1) MMD progressively reduces both bias and recommendation performance on both datasets and attributes when increasing λ , and (2) Adv. showcases a similar gradual change on ML-1M while recommendation accuracy remains relatively stable on LFM2B-DB when increasing the number of adversarial heads.

Dataset	Attribute	e Metric		# of Adv. Heads				
				0	3	5	10	
	Gender	Bine	Acc	$.789_{.003}$	$.642_{.029}$	$.573_{.014}$	$.542_{.008}$	
			Acc	$.788_{.001}$	$.653_{.029}$	$.609_{.009}$.608.047	
ML1M		NDCG	† †	.0625.0000	.0621.0001	.0621.0001	.0620.0000	
	Age	Bing	Acc	$.465_{002}$	$.310_{.022}$	$.265_{.007}$	$.232_{.026}$	
		$\text{Dias} \downarrow$	Acc	$.424_{003}$	$.267_{.024}$	$.218_{.012}$	$.193_{.011}$	
		NDCG	;↑	$.0625_{0000}$.0620.0002	.0618.0001	.0618.0001	
	Gender	Bing	Acc	$.723_{.002}$	$.677_{.011}$	$.608_{.040}$.600.011	
		$Dias \downarrow$	Acc	$.718_{.004}$	$.705_{.016}$	$.598_{.088}$	$.636_{.065}$	
LFM2B-DB		NDCG	÷ ↑	.0754.0000	$.0755_{.0001}$	$.0755_{.0000}$	$.0755_{.0001}$	
	Age	Bise	Acc	$.581_{005}$	$.505_{.020}$	$.455_{.012}$.390.011	
			Acc	$.504_{008}$	$.414_{.016}$	$.375_{.050}$	$.277_{.018}$	
		NDCG	; ↑	$.0754_{0000}$.0754.0000	$.0754_{.0000}$	$.0755_{.0001}$	

Table 4. Debiasing and performance results on both datasets and attributes using the

 Adv. method across different number of adversarial heads.

Weights Analysis. We now examine how the scaling weights affect the original user representations. As our method reduces the influence of sensitive information in \tilde{u} , we expect that the representations of users with different values of the protected attribute become more similar. We verify this by computing the average user representation for each user group and ranking the prototypes, i.e., the interpretable dimensions, from most to least similar. Given the ranking of two user groups, we compute Spearman's rank correlation [41] where values closer to +1 indicate both groups rank the prototypes similarly while values approaching -1 imply an inverse ranking. Figure 2 shows the results for the two gender groups on both datasets. Plots for age are provided in the appendix. Initially, we observe different prototype rankings between males and females, especially on the ML-1M dataset ($\rho = -.40$). However, as we increase the debiasing strength, the representations of males/females progressively become more aligned, as seen from the correlations plateauing between .70 and .80 across datasets and debiasing strategies. We derive that the scaling weights, while ensuring relevant recommendations for users and mitigating the bias of the protected attribute, lead to an alignment between the representations across user groups.

Taking a closer look at the scaling weights, we plot the average user-toprototype similarities \boldsymbol{u} and average $\boldsymbol{\omega}$ for Female (Fig. 3a) and Male (Fig. 3b) user groups on the ML-1M datasets sorted by most to least similar prototype. We notice a pattern wherein, on average, the scaling values of $\boldsymbol{\omega}$ shrink ($\boldsymbol{\omega} < 1$) the similarities to the prototypes most similar to the user group while they



Fig. 2. Spearman's correlation between avg. male/female prototype rankings on both datasets and both debiasing methods.



Fig. 3. Average user-to-prototypes similarities and average ω values for female users (left) and male users (right), sorted by most to least similar prototypes.

amplify ($\omega > 1$) the similarities to the least similar prototypes. By examining the interpretations of the most and least similar prototypes for each user group, shown in Fig. 4a, we infer that the debiased representations for female users show reduced activation towards genre patterns of Romance, Drama, and Comedy and increased activation towards Action and Sci-Fi. Conversely, the debiased representations for the male group display the opposite trend.

Finally, we look at a qualitative example showcasing the application of our learned scaling weights. In Fig. 4b, we report the relevant recommendations for an arbitrary female user from ML-1M before and after debiasing. We highlight items dropped from the recommendations in red text and newly recommended items in blue. Additionally, we use green to highlight the cell containing the ground truth items. Upon inspection, we observe that the debiased representation indeed affects the recommendations, particularly altering the items at the bottom of the list. We also note a reduction in the number of Romance movies



female (Top) and male (Bottom) users on trary female user on ML-1M before and ML-1M.

(a) Most and least similar prototypes for (b) Top-10 recommendations for an arbiafter debiasing.

Fig. 4. Examples of qualitative results.

and an increase in Sci-Fi movies before and after debiasing. This diversification also results in better recommendations for the user.

6 **Conclusion and Future Directions**

This work addresses the pervasive issues of societal bias in RSs from the user perspective. We propose a novel approach that leverages interpretable RS models and introduces per-user scaling weights to mitigate biases in the user representations while preserving recommendation quality. By applying our method to the prototype-based PROTOMF model [34], we demonstrate its effectiveness in reducing bias associated with protected attributes such as gender and age. Our evaluation on ML-1M and LFM2B-DB showcases the flexibility and efficacy of our approach in bias mitigation. Through qualitative analysis, we reveal correlations between consumption patterns and protected attributes, enhancing our understanding of bias in RSs. Moving forward, we envision exploring per-user weights that debias the user representation with respect to a conjunction of different protected attributes [29] simultaneously while also analyzing its effect on recommendation performance. Additionally, investigating end-users' perceptions of biases in recommendations appears as a promising avenue for future work.

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A Mathematics Framework of Artificial Shifted Population Risk and Its Further Understanding Related to Consistency Regularization

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Abstract. Data augmentation is an important technique in training deep neural networks as it enhances their ability to generalize and remain robust. While data augmentation is commonly used to expand the sample size and act as a consistency regularization term, there is a lack of research on the relationship between them. To address this gap, this paper introduces a more comprehensive mathematical framework for data augmentation. Through this framework, we establish that the expected risk of the shifted population is the sum of the original population risk and a gap term, which can be interpreted as a consistency regularization term. The paper also provides a theoretical understanding of this gap, highlighting its negative effects on the early stages of training. We also propose a method to mitigate these effects. To validate our approach, we conducted experiments using same data augmentation techniques and computing resources under several scenarios, including standard training, out-of-distribution, and imbalanced classification. The results demonstrate that our methods surpass compared methods under all scenarios in terms of generalization ability and convergence stability. We provide our code implementation at the following link: https:// github.com/vdlsfhll/ASPR.

Keywords: Population shift \cdot Augmentation framework \cdot Risk decomposition \cdot Regularization

X. Yang, S. Deng, S. Liu and Y. Suo—Equal Contribution.

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1 Introduction

Data augmentation creates a training dataset using synthetic data from the prior knowledge. It improves the generalization of machine learning models, particularly in the case of deep neural networks. For decades, its reliable performance has been verified in various of computer vision tasks such as image classification [9,13,21] and object detection [17,19]. To the best of our knowledge, there are currently two major explanations for the role of this technique. The first one views data augmentation as simply increasing the sample size, and explains it with statistical tools such as VC dimension theory [23]. The other one [11,25] views data augmentation as a regularization method, which train the model on a more complex population, which is called shifted population by injecting noise with prior knowledge to the original population, thereby enabling the model to retain semantic information unchanged.

However, the model is ultimately trained with the augmented samples, thereby improving the model's performance on the original population. Therefore, it is important to further explore the relationship between the expected risk of these two populations. To address this issue, we develop a rigorous mathematical framework of the shifted population $p^*(x')$ and data augmentation. Based on this framework, we prove that the expected risk of the shifted population is the summation of the original population and a gap term that can be viewed as a consistency regularization term. This decomposition sheds light on the unification of the two aforementioned explanations. Moreover, inspired by the work of [10], the generalization of the model greatly depends on the consistency between the empirical risk of the original population and the shifted one, and the gap term may violate such consistency. To address this issue, we add a trade-off coefficient to the gap term to highlight the importance of the learning of major features, which is controlled by the expected risk of p(x). This approach greatly benefits the performance of the model.

At present, some work like [3] has provided a decent mathematical framework for data augmentation, but it is too limited to describe some of the existing data augmentations, and it completely ignores the gap term. However, this neglect could be harmful, for it is indicated by our analysis and experiment that reducing its impact in early stages of training has been proven to be helpful for the model's generalization. Please see Appendix D.1 for a more detailed discussion.

We conducted experiments to evaluate the proposed training strategy on popular image classification benchmarks, namely CIFAR-10/100 [12], Food-101 [2], and ImageNet (ILSVRC2012) [2]. Our evaluation involved using representative deep networks such as ResNet-18, ResNet-50, and WideResNet-28-10. In addition to assessing the performance in the standard scenario, we also tested the algorithm in the out-of-distribution (OOD) scenario with dataset PACS [14]) and the long-tail imbalanced classification (LT) scenario with dataset LT-CIFAR10 [5]. Across all our experiments, our strategy consistently achieved lower error rates and demonstrated more stable convergence compared to the standard data augmentation strategy. This paper's contributions can be summarized as follows:

- 1. We provide a rigorous mathematical definition for the shifted distribution $p^*(x')$ of the augmented samples, which further reveals that the commonly used augmented samples actually comes from the a conditional distribution p(x'|x). We also give a mathematical description of sampling from this distribution and find that the samples used during training from this marginal distribution are not completely independent, which is surprising.
- 2. Based on the proposed mathematical framework, we discover that the risk on the shifted population $p^*(x')$ can be decomposed into a risk on the original population p(x) and a gap term, serving as a consistency regularization term.
- 3. We provide a theoretical understanding of such decomposition and an explanation of why our training strategy is beneficial for the improvement of generalization.

2 Related Work

Data Augmentation Frame Work. Data augmentation methods play a crucial role in improving the performance of machine learning models in practical applications. These methods encompass a range of techniques, including traditional fixed augmentation methods like Cutout [6], Mixup [29], and Cutmix [26]. Additionally, there are adaptive augmentation methods such as AutoAugment [4], Fast AutoAugment [16], DADA [15], and CMDA [22], which dynamically design augmentations based on the dataset. Despite the availability of these diverse augmentation methods, there is a dearth of theoretical frameworks for analyzing the population shift phenomenon induced by data augmentation and the associated shifted population risk.

A recent work [3] provides a theoretical framework that defines the augmentation operator as a group action. However, their framework has certain limitations, as evidenced by several common augmentation operators that are incompatible with the group action framework, as detailed in the Appendix D.1. Our proposed framework can be applied to a wider range of data augmentation operators compared to theirs.

Population Shift. Population shift is a common concern in machine learning robustness and generalization problems. It refers to a problem in which the population of data changes during some processes, such as a distribution being transformed to other distributions within the same distribution family, and the change of the parameters of a distribution. A common example for population shift in machine learning is the different semantic styles between the training and testing sets, such as PACS [14], Rotated MNIST, Color MNIST [1], VLCS, and Office-Home [24]. However, not all types of population shifts are natural. Style shifts such as PACS are naturally generated distributions, while population shifts such as Rotated MNIST and Color MNIST are artificially generated. It is obvious that all data augmentations will produce an artificial population shift. This work aims to provide a theoretical framework for artificial population shifts and analyze the relationship between the **shifted population risk** and the **original population risk**.

3 Method

3.1 Revisiting Data Augmentation with Empirical Risk

We conduct research in the case of classification and denote the data space and label space as \mathcal{X} and \mathcal{Y} and a joint distribution p is defined on $\mathcal{X} \times \mathcal{Y}$, with marginal distribution p(x) and conditional distribution p(x|y). We call a sample x drawn from p(x) a "clean sample". We aim to train a model $f : \mathcal{X} \to \mathcal{Y}$ by minimizing the following risk with a loss function $\mathcal{L}(\cdot, \cdot)$:

$$R_f(p) = \int \mathcal{L}(f(x), y) \mathrm{d}p(x, y), \tag{1}$$

As (1) is usually intractable, the empirical risk minimization principle is used, aiming at optimizing an unbiased estimator of (1) over a training dataset $\mathcal{D} = \{(x_i, y_i)\}_{i=1}^N$:

$$\hat{R}_f(p) = \frac{1}{N} \sum_{i=1}^N \mathcal{L}(f(x_i), y_i),$$
(2)

Following [25], we introduce the following assumption to build a bridge between empirical risk and expected risk:

Claim. Let C(f) be some complexity metric of f, N be the number of data (don't have to be independent), B(N) be the "independence" of the input data. For $\forall \delta > 0$, we assume that the following holds with probability $1 - \delta$:

$$R_f(p) - \hat{R}_f(p) \le \phi(C(f), B(N), \delta).$$
(3)

where $\phi(\cdot)$ is a function of these three terms, and it monotonically increases with respect to the second variable.

We refer the readers to [18] for more detail about the convergence in the non iid case. It is worth noting that data augmentation produces an augmented sample x', which is a distinct random variable from the clean sample x, with a different distribution $p^*(x')$ but the same probability space triplet. This leads to a new population $\tilde{p}(x', y)$ and an expected risk defined on it. Specifically, the empirical risk function is defined as follows:

$$\hat{R}_f(p^*) = \frac{1}{N} \sum_{i=1}^N \mathcal{L}(f(x'_i), y_i).$$
(4)

It is important to note that minimizing (4) does not necessarily result in the minimization of (1) or even (2). Meanwhile, data augmentation is also recognized as a regularization technique that can reduce generalization error without necessarily reducing training error [7,28]. Our proposed decomposition as well as the framework should be helpful when one tries to overcome these struggles.

3.2 The Augmented Neighborhood

Data augmentation is typically applied directly to a clean sample x to generate an augmented sample x'. The augmentation is usually designed to preserve the semantic consistency between x and x', hence it is often referred to be "mild". However, the data augmentation is usually controlled by a set of parameter when it is applied to a fixed clean sample x. When the parameters are iterated, a large set of augmented samples are produced, among which there are samples are over augmented and should not be considered "mild". As a result, a series of rigorous mathematical definitions are required, so one may draw a line between "ordinary" data augmentation and a "mild" one.

The Augmentation and Limitation. We begin this section with the definition of data augmentation:

Definition 1. Let \mathcal{X} be the data space, endow \mathcal{X} with Borel σ - algebra \mathcal{F} , let the data augmentation $A_i(\cdot, \cdot)$ be a map from $\mathcal{X} \times \Theta(A_i)$ to \mathcal{X} satisfying:

- 1. For every fixed x in \mathcal{X} , the map $\theta \mapsto A_i(\theta, x)$, is differentiable and injective. We denote the inverse of this map as $h_{A_i,x}^{-1}$.
- 2. For every fixed θ in $\Theta(A_i)$, the map $A_i(\theta, \cdot)$ is an \mathcal{F} -measurable map.
- 3. $\forall x \in \mathcal{X} \exists e_i \in \Theta(A_i) \text{ s.t. } A(e_i, x) = x \text{ and such } e_i \text{ is unique.}$

where $\Theta(A_i)$ is the parameter space of $A_i(\cdot, \cdot)$.

The differentiability of some popular data augmentations has been proven in [22]. The injectivity of the data augmentation is always guaranteed given proper parameterization and a carefully chosen parameter space. The measurable assumption is required to ensure that $A(\theta, x)$ is still measurable, which is necessary for the adjoint random variable x'. However, the tractability of $h_{A_i,x}^{-1}$ is not always guaranteed, but the good news is that it is not always required in practice. More detailed discussion is provided in Sect. 3.2, where we discuss how to sample from the conditional distribution p(x'|x).

Denote the set of data augmentation as $\mathcal{A} = \{A_1, \ldots, A_m\}$, among which A_i corresponds to a certain type of data augmentation such as rotation, Gaussian blur and so on. Denote dim $(\Theta(A_i)) = d_i$, where $\Theta(A_i)$ denotes the parameter space of A_i . For example, the parameter space of rotation is usually chosen as $(0, 2\pi)$ and the dimension is 1. The distribution of the parameter defined on $\Theta(A_i)$ is denoted as $p_i(\theta)$. Now for a given clean sample x_0 , we consider all of its augmented sample, which is the image of the mapping $A_i(x_0, \cdot)$, defined on $\Theta(A_i)$:

Definition 2. For any given clean sample $x_0 \in \mathcal{X}$ and data augmentation A_i with parameter space $\Theta(A_i)$, the **augmentation neighborhood** of x_0 induced by A_i is defined as:

$$A_i(x) := \bigcup_{\theta \in \Theta(A_i)} A_i(\theta, x).$$
(5)

Now we should add some restrictions to this set so make it "mild".

At first we introduce the conception C, a map from input space \mathcal{X} to the label space $\mathscr{L} = \{c_1, \ldots, c_l\}$ where l denotes the number of class, such that for every clean sample pair $(x, y) \sim p(x, y), C(x) = y \in \mathscr{L}$, conception is the desired ground truth map. C induces a partition of the sample space, by giving l mutually disjoint sets such that $\Gamma_i = \{x | C(x) = c_i\}$, what we call level set. We denote the level set of the class of a sample x_0 with Γ_{x_0} , and we use this level set to describe the semantic consistency. The conception C represents the prior knowledge of people when they perform data augmentation. The definition is given as followed:

Definition 3. For any given clean sample $x_0 \in \mathcal{X}$, and augmentation A_i with parameter space $\Theta(A_i)$, the **consistency augmentation neighborhood** (CAN for short) of x_0 induced by A_i is defined as:

$$\mathcal{O}_{x_0}^{A_i} := A_i(x_0) \cap \Gamma_{x_0}.$$
(6)

Now we will introduce how to sample from the CAN.

Sampling from CAN of x_0 . An augmented sample is generated given a clean sample, together with the aforementioned mild argument, we claim that the sampling procedure should be described with a conditional distribution p(x'|x), whose supporting set is CAN of x_0 . The fact that $\forall x' \in \mathcal{O}_{x_0}^{A_i}$, there exists only one $\boldsymbol{\theta} := h_{A_i,x_0}^{-1}(x') \in \Theta(A_i)$ such that $x' = A_i(\boldsymbol{\theta}, x)$ which is ensured by our definition. Furthermore, with the measurability of $A_i(\cdot, x)$, x' is a random variable. Therefore, for any given data augmentation A_i , the conditional distribution p(x'|x) induced by A_i is defined as:

Definition 4. For any given clean sample $x \sim p(x)$, the conditional distribution p(x'|x) of the adjoint variable x' with $\operatorname{supp}(p(x'|x)) = \mathcal{O}_x^{A_i}$ is given as

$$p(x'|x) \propto p_i(h_{A_i,x}^{-1}(x')) \left| \frac{\partial}{\partial \boldsymbol{\theta}} A_i(\boldsymbol{\theta}, x) \right|_{\boldsymbol{\theta} = h_{A_i,x}^{-1}(x')} \mathbf{1}_{x' \in \mathcal{O}_x^{A_i}}.$$
 (7)

Sampling from $A_i(x)$ is equivalent to sampling from $p_i(\boldsymbol{\theta})$ defined on $\Theta(A_i)$, for $h_{A_i,x}^{-1}(A_i(x)) = \Theta(A_i)$, given the injectivity of $A(\cdot, x)$. Furthermore, to sample from $A_i(x) \cap \Gamma_x$, we need to sample from the truncated distribution:

$$p_i(\boldsymbol{\theta}) \mathbf{1}_{\boldsymbol{\theta} \in h_{A_i}^{-1}(\mathcal{O}_x^{A_i})}.$$
(8)

Rejection sampling is one effective way to generate augmented samples, but it may be infeasible in high-dimensional cases due to its computational cost. Although various methods, such as nested sampling, adaptive multilevel splitting, or sequential Monte-Carlo sampling, could be viable alternatives, we leave the exploration of these methods for future work. Additionally, the rejection step can be seen as a way to inject humane prior knowledge to samples, which aligns with the intuition on the process of data augmentation. In our experiment, we assume that it would be enough to sample from the subset of $h_{A_i}^{-1}(\mathcal{O}_x^{A_i})$, we use human prior knowledge in rejection sampling to roughly determine a subset of it. We begin by selecting the candidate of edges of these subsets, then apply $A(\theta, x)$ for parameters of these edges, and reject or accept these edges by observing the output samples. However, this method is inefficient and risky, rejection sampling is infamous for its inefficiency and the initial selection of edges could be problematic since they may be too small compared to the ground truth. We plan to develop better methods based on our framework in future work.

The conditional distribution p(x'|x) is now well-defined, with its marginal distribution given by $p^*(x') = \int p(x'|x)p(x)dx$. However, it is important to note that p(x'|x) is unlikely to be tractable. The description above is useful in understanding that an augmented sample is a random variable induced from the of data augmentation, given the measurability of $A(x, \cdot)$.

Finally, it's worth mentioning that generating M samples for each of the N clean samples does not result in $M \times N$ completely independent augmented samples. But (4) still yields an unbiased estimator of the shifted population risk, due to the following equation:

$$\mathbb{E}_{p(x'|y)} \left[\mathcal{L}(y, f(x')) \right] = \mathbb{E}_{p(x|y)} \left[\mathbb{E}_{p(x'|x,y)} \left[\mathcal{L}(y, f(x')) | x \right] \right],$$

$$= \mathbb{E}_{p(x|y)} \left[\mathbb{E}_{p(x'|x)} \left[\mathcal{L}(y, f(x')) | x \right] \right],$$

$$\hat{R}_{f}(p^{*}) = \frac{1}{N} \sum_{i=1}^{N} \frac{1}{M} \sum_{j=1}^{M} \mathcal{L}(y_{i}, f(x'_{i_{j}})),$$

Taking expectation on the both side of the third equation yields the desired result. One should notice that augmented sample x' is independent of y once its original clean sample x is given, which explains the second equality.

The above definition in the case of a finite set of data augmentations and the composition order is given in Appendix A.

By establishing these definitions and concepts in this section, we are provided with a comprehensive understanding of the topic at hand. Which provides a solid foundation for the decomposition of the expected risk in the coming section.

3.3 The Artificial Shifted Population Risk

After defining the augmented neighborhood and giving sampling method by defining the adjoint variable x' and its conditional distribution p(x'|x), we then evaluate the risk on the shifted population p(x', y). One should realize that the collection of all the samples generated from p(x) is a subset of the samples generated from $p^*(x')$.

For simplicity, we only consider the risk function in the case of cross-entropy and softmax on the shifted population p(x', y), and our method should be able to extend to the other cases similarly:

$$R_{f}(p^{*}) = \mathbb{E}_{p^{*}(x')} \left[H(p(y|x'), q_{\phi}(y|x')) \right],$$

= $\mathbb{E}_{p(y)p(x'|y)} \left[-\ln q_{\phi}(y|x') \right]$
= $\mathbb{E}_{p(y)p(x',x|y)} \left[-\ln q_{\phi}(y|x') \right],$ (9)

among which ϕ denotes the parameter of the neural network and q represents a probabilistic surrogate model. The decomposition of this shifted population risk is examined with the following theorem:

Theorem 1. With the shifted population risk in the form of (9), we have the following decomposition:

$$\mathbb{E}_{p^{*}(x')} \left[H(p(y|x'), q_{\phi}(y|x')) \right] \\ = \mathbb{E}_{p(x)} \left[H(p(y|x), q_{\phi}(y|x)) \right] + \mathbb{E}_{p(x)p(y|x)p(x'|x)} \left[\ln \frac{q_{\phi}(y|x)}{q_{\phi}(y|x')} \right],$$
(10)

The proof of the Theorem 1 can be found in Appendix B.1. This demonstrates that in the case of cross-entropy and softmax, the **shifted population risk** is actually the sum of the **original population expected risk** and a gap term that can be viewed as a **consistency regularization term**. Next, we provide a theorem that explains the second term.

3.4 Understanding the Decomposition of Shifted Population Risk

From the last section, we have:

$$\mathbb{E}_{p(x)p(y|x)p(x'|x)}\left[\ln\frac{q_{\phi}(y|x)}{q_{\phi}(y|x')}\right] = \mathbb{E}_{p(x)p(x'|x)}\left[\ln\frac{q_{\phi}(y_x|x)}{q_{\phi}(y_x|x')}\right],\tag{11}$$

where y_x is the ground true label of clean sample x. Since $q_{\phi}(y|x)$ is modeled with softmax, we have:

$$q_{\phi}(y_i|x) = \frac{\exp(\mathbf{w}_i^T h_{\theta}(x))}{\sum_{j=1}^l \exp(\mathbf{w}_j^T h_{\theta}(x))},$$
(12)

where $h_{\theta}(x) = (h_1(x), h_2(x), \dots, h_d(x), \dots, h_D(x))^T$ (the subscript θ of the component is omitted for convenience) is the feature vector of x, and $\mathbf{W} = (\mathbf{w}_1, \dots, \mathbf{w}_l)$ is the weight of the output layer, now $\phi = \{\theta, \mathbf{W}\}$. For every feature $h_d(x)$, its density is:

$$q_{\phi}(y_i|h_d(x)) = \frac{\exp(w_{i,d}h_d(x))}{\sum_{j=1}^{l}\exp(w_{j,d}h_d(x))},$$
(13)

Inspired by [10], we partition the features into major features and minor features by information gains. For major features, the density function $q_{\phi}(y|h_d)$ concentrates on some point mass. For minor features, the possibility density $q_{\phi}(y|h_d)$ is relatively uniform.

Then for every given x, we have:

$$\mathbb{E}_{p(x'|x)}\left[\ln\frac{q_{\phi}(y_x|x)}{q_{\phi}(y_x|x')}\right] = \mathbb{E}_{p(x'|x)}\left[\ln\left(\frac{\sum_{j=1}^{l}\exp\left((\mathbf{w}_j - \mathbf{w}_x)^T h_{\theta}(x')\right)}{\sum_{j=1}^{l}\exp\left((\mathbf{w}_j - \mathbf{w}_x)^T h_{\theta}(x)\right)}\right)\right], (14)$$

For convenience, we denote

$$\exp\left((\mathbf{w}_j - \mathbf{w}_x)^T h_{\theta}(x)\right) = \rho_{\theta,x,j},$$

$$\sum_{j=1}^{l} \rho_{\theta,x,j} = \rho_{\theta,x},$$
(15)

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we then examine the relationship of feature and the second term with the following theorem:

Theorem 2. Assuming that for every θ , sample pair (x, x') and indicies j, there exist $\beta_{1,j}, \alpha_{1,j} > 0$ such that

$$\alpha_{1,j} < \rho_{\theta,x,j}, \ \rho_{\theta,x',j} < \beta_{1,j}, \tag{16}$$

Then for any given x, we have:

$$\mathbb{E}_{p(x'|x)}\left[\left|\ln\frac{q_{\phi}(y_x|x)}{q_{\phi}(y_x|x')}\right|\right] = \mathbb{E}_{p(x'|x)}\left[\left|\sum_{j=1}^{l} O\left(\left(\mathbf{w}_j - \mathbf{w}_x\right)^T \left(h_{\theta}(x) - h_{\theta}(x')\right)\right)\right|\right],\tag{17}$$

The proof of the Theorem 2 can be found in the Appendix B.2. With Theorem 2, we show how the second term affects the weights. Since the data augmentation must cause a large variance in some features particularly in early training phases, which means that

$$\exists \eta_1 > 0, \ |h_d(x) - h_d(x')| > \eta_1, \tag{18}$$

for some features including minor and major features. This forces that $\forall j \in \{1, \ldots, l\}, w_{j,d} \to w_{x,d}$, resulting in a uniform distribution of $q_{\phi}(y|h_d(x))$, and such regularization of $w_{j,d}$ is not appropriate for major features. Now let us see how the first term affects the weights

$$\mathbb{E}_{p(x)}\left[H(p(y|x), q_{\phi}(y|x))\right] = \mathbb{E}_{p(x)}\left[\sum_{i=1}^{l} \exp\left(\left(\mathbf{w}_{j} - \mathbf{w}_{x}\right)^{T} h_{\theta}(x)\right)\right], \quad (19)$$

And for any minor feature, its variation should not change the result, hence we have $w_{j,d} \approx w_{i,d}$, $1 \leq i, j \leq l$. In contrast, the weights of major features should be different:

$$\exists \eta_2 > 0, \ |w_{j,d} - w_{x,d}| > \eta_2 \tag{20}$$

now we realize that, with the effect of data augmentation, the first term and the second term have different impacts on the weight of some major features and the same impact on minor features. Since our model mainly relies on major features to provide prediction, such an effect causes an unstable convergence. To highlight the positive effect provided by the first term at the beginning, a simple trick is to add a coefficient λ ($\lambda < 1$) to the second term.

Now we discuss how λ may help refine the generalization of the model. We denote the model trained using augmented samples as f_{aug} :

$$R_{f_{aug}}(p^*) = \mathbb{E}_{p(y)p(x'|y)} \left[\mathcal{L}(y, f(x')) \right],$$

$$R_{f_{aug}}(p) = \mathbb{E}_{p(y)p(x|y)} \left[\mathcal{L}(y, f(x)) \right],$$
(21)

Note that we train our model using $\hat{R}_{f_{aug}}(p^*)$ and evaluate the generalization of our model using $R_{f_{aug}}(p)$. Based on the assumption Sect. 3.1, with augmented sample and clean sample pairs instead of clean samples alone, we have:

$$R_{f_{\text{aug}}}(p^*) \leqslant \hat{R}_{f_{\text{aug}}}(p^*) + \phi(C(f), B(N \times M), \delta),$$
(22)

Theorem 1 can then be reformulated with our new formulation:

$$R_{f_{\text{aug}}}(p^*) = R_{f_{\text{aug}}}(p) + \text{GAP},$$

$$\hat{R}_{f_{\text{aug}}}(p^*) = \hat{R}_{f_{\text{aug}}}(p) + \widehat{\text{GAP}}_{M \times N},$$
(23)

where GAP is the second term in the right hand side of Theorem 1 and $\widehat{\text{GAP}}_{M \times N}$ is its empirical estimator using $M \times N$ non iid pairs of (x, x'). Hence, (22) is reformulated by:

$$R_{f_{\text{aug}}}(p) \leq \hat{R}_{f_{\text{aug}}}(p) + \phi(C(f), B(N \times M), \delta) + \\ \widehat{\text{GAP}}_{M \times N} - \text{GAP}.$$
(24)

Now we show that the noise $\widehat{\operatorname{GAP}}_{M \times N} - \operatorname{GAP} \to 0$:

$$GAP = \mathbb{E}_{p(x)p(y|x)p(x'|x)} \left[\ln \frac{q_{\phi}(y|x)}{q_{\phi}(y|x')} \right],$$

we denote $\mathcal{B}(y, g(x, x')) = \ln \frac{q_{\phi}(y|x)}{q_{\phi}(y|x')}$, then we assume that given any clean sample pair (x_i, y_i) :

$$\operatorname{Var}_{p(x'|x_i)} \left[\mathcal{B}(y_i, g(x_i, x')) \right] \leq B,$$

then for the estimator:

$$\begin{split} & \mathbb{E}_{p(x,y)p(x'|x)} \left[\mathcal{B}(y,g(x,x')) \right] \\ &= \mathbb{E}_{p(x,y)} \left[\mathbb{E}_{p(x'|x)} \left[\mathcal{B}\left(y,g(x,x')\right) | x,y \right] \right], \\ & \widehat{\text{GAP}}_{M \times N} = \frac{1}{N} \sum_{i=1}^{N} \frac{1}{M} \sum_{j=1}^{M} \mathcal{B}(y_i,g(x_i,x'_{i_j})) \end{split}$$

where x'_{i_j} denotes augmented samples from $p(x'|x_i)$ consider its variance:

$$\operatorname{Var}\left(\widehat{\operatorname{GAP}}_{M\times N}\right) = \frac{1}{N^2 M} \sum_{i=1}^{N} \operatorname{Var}_{p(x'|x_i)} \left[\mathcal{B}(y_i, g(x_i, x'))\right],$$

with the assumption:

$$\operatorname{Var}\left(\widehat{\operatorname{GAP}}_{M\times N}\right) \leq \frac{B}{NM},$$

then the variance is of order O(1/NM), which indicates the faster convergence speed.

We determine that the generalization of model depends on $\hat{R}_{f_{\text{aug}}}(p)$ instead of what we directly optimize: $\hat{R}_{f_{\text{aug}}}(p^*)$. Hence we would like to keep the consistency between $\hat{R}_{f_{\text{aug}}}(p)$ and $\hat{R}_{f_{\text{aug}}}(p^*)$, *i.e.*, the decreasing of $\hat{R}_{f_{\text{aug}}}(p^*)$ guarantees that of $\hat{R}_{f_{\text{aug}}}(p)$ to ensure the improvement of generalization when training the model. As it is analyzed before, $\widehat{\text{GAP}}_{M \times N}$ may lead to different weights of some major features compared with $\hat{R}_{f_{\text{aug}}}(p)$ in early training stages, which will destroy such consistency. This indicates the importance of our proposed coefficient λ .

4 Experiment

We demonstrate the standard training strategy in Algorithm 1 and our proposed training strategy in Algorithm 2 in Appendix C. We also conduct an experiment on the selection of the hyperparameter λ of Algorithm 2 in Appendix E.

4.1 Experiment Implementation

Standard Scenario Experiment: Validation Models and Datasets We have conducted experiments on CIFAR10/100 [12], Food101 [2], and ImageNet (ILSVRC-2012) [20] with various models to evaluate our training strategy. For each of them, a validation set is split from the training set to find networks with the best performances. More dataset splitting details are shown in Appendix F.1. In this paper, ResNet [9] and WideResNet [27] are trained with different strategies. For datasets CIFAR10/100 and Food101, ResNet-18, ResNet-50, WideResNet-28-10 and WideResNet-40-2 are chosen as our baseline models. For ImageNet, ResNet-50 and ResNet-101 are used for evaluation. All images in baseline (standard method) and our method are processed with same augmentation (horizontal flips, random crops and random rotation). λ was selected to 0.5 for it achieve the best performance among all the experiments with our strategy. For a fair comparison, we set the basic batch size (bbs) and performed standard method experiments with both 1x bbs and 2x bbs (our method actually takes twice the amount of data sample) to ablate the estimation error effect caused by the batch size. More details about data augmentation and network training are shown in Appendix F.2 and Appendix F.3.

To ensure that our strategy is applicable to other settings, we conduct experiments in the following two cases:

OOD Scenario Experiment: Validation Models and Datasets Experiments on PACS [14] are conducted using ResNet-18 and ResNet-50 [9]. In these experiments, we employed the leave-one-domain-out strategy for OOD validation. For



Fig. 1. Top-1 accuracy (%) with error bar (mean \pm std) on CIFAR10/100, Food101 and ImageNet on the test set. The Y-axis is the Top-1 accuracy and the X-axis is the type of network.

image augmentation, we followed the same approach as Domainbed [8], both in the ERM algorithm and our proposed method. Further information regarding data augmentation and network training can be found in Appendix F.2 and Appendix F.3.

Long-Tailed Scenario Experiment: Validation Models and Datasets We consider long-tail (LT) imbalance and conducted experiments on LT CIFAR-10 [5] using ResNet-18. We keep the validation set and test set unchanged and reduce the number of training set per class according to the function $n = n_i \mu^i$, where n_i is the original number of the i - th class of the training set (following [5]). μ is between 0 and 1, which is determined by the number of training samples in the largest class divided by the smallest. This ratio is called imbalance ratio and it is set from 10 to 100 in our settings. Further information regarding training hyperparameters can be found in Appendix F.2 and Appendix F.3.

4.2 Experimental Results

Settings and instructions For standard scenario experiment, we select the model with the highest validation accuracy during training and report the test accuracy in Fig. 1. The results with error bars are presented at Appendix F.4, where we have conducted three independent experiments and calculated the mean values as the results on CIFAR10/100 and Food101 and only one independent experiment on ImageNet (ILSVRC2012) because of computational constraints.

As for the OOD scenario experiments, we have conducted three independent experiments and select the model with the best top-1 accuracy on the test domain. The results with error bar can be seen in Fig. 2 and Appendix F.4 Table 5.

For long-tailed scenario experiment, we use the Area Under the Curve (AUC), Average Precision (AP) and top-1 accuracy as evaluation metrics. We select the model with the best AUC on the validation set during training and report the results on the test set in Fig. 3 and Appendix F.4 Table 6.

Experiment Analysis. From our experimental results (Fig. 1, Fig. 2, Fig. 3), we can see that the model trained with our proposed consistency regularization strategy of data augmentation converges to a better local optimum.



Fig. 2. Top-1 accuracy (%) with error bar (mean \pm std) over the four test domain of PACS and their average. The X-axis is test-domain and the Y-axis is the Top-1 accuracy.



Fig. 3. AUC, Top-1 accuracy (%), AP with error bar (mean±std) of Resnet-18 on the long-tailed scenario experiment (LT-CIFAR10). The X-axis of the figures is the value of the imbalance ratio.

From Fig. 4a, we can see the validation set performance of our method even exceeds the training set performance of the standard data augmentation training method in almost the whole process of training. This demonstrates the improvement of generalization after adding the coefficient.

As demonstrated in Fig. 4 and Fig. 5, our training strategy leads to a stable convergence compared with the standard data augmentation training strategy. The stable convergence is caused by the coefficient λ , as we discuss in Sect. 3.3. The coefficient λ diminishes the negative effect of estimate variance, resulting in a more stable convergence. The training process for all circumstances is presented in Appendix F.5

5 Conclusion and Discussion

Rethinking of Shifted Population. In this paper, we develop a new set of definitions for shifted population, augmented samples and its conditional distribution. We leverage our proposed definition to establish the decomposition of the shifted population risk, providing an explanation for how data augmentation enhances the generalization ability of model. Better Training Strategy. Based on the proposed decomposition, we realize that the key to improving generalization lies in keeping the consistency between $\hat{R}_{f_{\text{aug}}}(p^*)$ and $\hat{R}_{f_{\text{aug}}}(p)$, which is likely to be violated by the gap term specifically in the early training stages. Adding a coefficient to the gap term refines this, and it is proposed as a training strategy with augmentation. As demonstrated in our experiment, our method outperforms the standard augmentation training strategy. Meanwhile, our proposed strategy is highly related to the augmentation schedule, an existing training strategy. Our work could provide comprehensive understanding on how it works. What's more, there is more than one solution to the problem of the gap term, which is left for future work.

Limitation. Considering the fact that this paper mainly conducts analysis in the case of classification tasks, some of the results proposed in this paper lack versatility. However, the framework of the analysis is transferable, and based on the definition of expected risk, similar results can be attained on other tasks. Conditional distribution of adjoint variable p(x'|x) is intractable given the fact that although the differentiability of most of the classic augmentations has been verified in other works, there are data augmentations that have not, some of them may even be not genuinely differentiable. Hence, other definitions of p(x'|x) that bypass the necessity of differentiability can be explored in future work.

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Attention-Driven Dropout: A Simple Method to Improve Self-supervised Contrastive Sentence Embeddings

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Abstract. Self-contrastive learning has proven effective for vision and natural language tasks. It aims to learn aligned data representations by encoding similar and dissimilar sentence pairs without human annotation. Therefore, data augmentation plays a crucial role in the learned embedding quality. However, in natural language processing (NLP), creating augmented samples for unsupervised contrastive learning is challenging since random editing may modify the semantic meanings of sentences and thus affect learning good representations. In this paper, we introduce a simple, still effective approach dubbed ADD (Attention-Driven Dropout) to generate better-augmented views of sentences to be used in self-contrastive learning. Given a sentence and a Pre-trained Transformer Language Model (PLM), such as RoBERTa, we use the aggregated attention scores of the PLM to remove the less "informative" tokens from the input. We consider two alternative algorithms based on NAIVEAGGREGATION across layers/heads and ATTENTIONROLLOUT [1]. Our approach significantly improves the overall performance of various self-supervised contrastive-based methods, including SIMCSE [14], DIF-FCSE [10], and INFOCSE [33] by facilitating the generation of highquality positive pairs required by these methods. Through empirical evaluations on multiple Semantic Textual Similarity (STS) and Transfer Learning tasks, we observe enhanced performance across the board.

1 Introduction

Self-supervised contrastive learning is amongst the most promising approaches for learning representation without relying on human annotation. Recent advancements in self-supervised learning [14, 16, 19, 24] use the contrastive loss to maximize the similarity of representation obtained from different distorted versions of an input sample while pushing apart dissimilar pairs from an input sample. Therefore, defining data augmentation for generating positive and negative pairs is crucial in contrastive prediction tasks that yield effective representations. However, in NLP, creating augmented samples for unsupervised contrastive learning is challenging since random word editing operations (deletion/masking or replacement) may modify the semantic meanings of sentences.

Existing methods create [37] the negative pairs from a random collection of sentences, while the positive pairs are obtained by augmentation. Other contrastive approaches try to create positive pairs at the word level using synonym replacement, random insertion, random Swap, deletion, and substitution [30,36]. While at the sentence level, these sentences are shuffled to create a new sample, and if the given text sample contains multiple sentences with duplicate sentences, these duplicate sentences are removed [14]. Numerous recent approaches provide data augmentation by applying some changes in the model, such as injecting random noise [17], drop out [14,18], or random span masking [19] aiming to supply an expressive semantic interpretation. Since positive pairs are created from similar sentences, it may affect syntactic alignment across views [18].

In this paper, we introduce ADD (<u>Attention-Driven Dropout</u>), a straightforward and effective approach for data augmentation in unsupervised contrastive learning. Specifically, we propose ADD to quantify the relevance of a word in a sentence by considering the summation of the attention score. Consider a sequence of input tokens and a pre-trained Transformer model such as BERT or RoBERTa; we use the aggregated attention scores of the PLM to remove the less "informative" tokens from the input. We assess two alternative algorithms based on NAIVEAGGREGATION across layers/heads and ATTENTIONROLLOUT [1]. Our method expands on the results of SIMCSE [14] to aid further the generation of quality positive pairs used by such methods. We present empirical results on several Semantic Textual Similarity (STS) and Transfer Learning tasks, where we find consistently improved overall performance.

The contributions of our proposed work are threefold:

- We propose Attention-Driven Dropout as a novel data augmentation technique that can be used in unsupervised contrastive learning.
- We introduce two approaches to quantify the word relevance in a sentence by utilizing attention scores.
- We perform extensive experiments and demonstrate the efficacy of our method in enhancing the performance of self-contrastive models. Our extensive empirical analysis substantiates the effectiveness of our approach, highlighting its capability to improve the overall results.

2 Background and Related Work

Many tasks in NLP benefit from the amount of training data. Data augmentation techniques are used to generate additional, synthetic data using the training data while improving model performance. Recently [8,25,35] review and compare different augmentation techniques such as reordering, substitution, random perturbation, word deletion and span deletion in contrastive learning. Self-supervised contrastive learning like SimCSE [14] takes a randomly sampled mini-batch of sequences $\boldsymbol{X} = \{\boldsymbol{x}_t\}_{t=1}^M, \ \boldsymbol{x} \in \mathcal{X} \subseteq \mathbb{R}^p$, the transformer function derives two augmented views $\boldsymbol{x}_i = \tau(\boldsymbol{x}), \boldsymbol{x}_j = \tau'(\boldsymbol{x})$ for each sample in \boldsymbol{X} . The two augmented samples \boldsymbol{x}_i and \boldsymbol{x}_j are then fed to an encoder network $f_{\boldsymbol{\theta}}$ with trainable parameters $\boldsymbol{\theta} \subseteq \mathbb{R}^d$. These features are then transformed with a projection multi-layer perceptron (MLP) head, which results in h_i and h_j . The contrastive estimation for a positive pair of examples (x_i, x_j) is defined as:

$$\ell_{(x_i, x_j)} = -\log \frac{\exp(\sin(\boldsymbol{h}_i, \boldsymbol{h}_j)/\tau)}{\sum_{k=1}^{2N} \mathbb{I}_{[k\neq i]} \exp(\sin(\boldsymbol{h}_i, \boldsymbol{h}_k)/\tau)}$$
(1)

The core component of the Transformer and BERT models is the attention mechanism. The attention mechanism works by learning a set of weights also called attention scores, that indicate the importance of each input part. These attention scores are then used to weigh the different parts of the input so that the model focuses more on the important parts and less on the less important parts.

There are several types of attention mechanisms, such as additive, multiplicative, or scaled dot-product attention, etc. Multi-head self-attention is widely used in recent Transformer models.

Given input sequence $x \in \mathbb{R}^{l \times d}$ with the length of l and d dimension and $W_q, W_k, W_v \in \mathbb{R}^{d \times d}$ be the matrices for query, key, and value respectively. Then, each x is associated with a query $Q = xW_q$ and a key-value pair (K, V) $(K = xW_k; V = xW_v)$ and an attentive representation $A = Softmax(\alpha^{-1}QK^T)$ of x in the multi-head self-attention computed by H = AV. Here, α is a scaling factor, and $A = \{a_1, ..., a_h\}$ is the attention distribution.

3 Method

We propose a simple principle to create an augmentation of sentences, which can be used as a positive pair in a self-contrastive learning scheme. Compared to conventional self-contrastive approaches that duplicate the input sequence x_i , we apply our proposed ADD augmentation. As depicted in Fig. 1, our proposed method gathers attention scores for the sequence from a pretrained language model. It then aggregates these scores in order to obtain a single value for each token, representing its relevance. The k smallest tokens are then removed from the input sequence. After our procedure, the original and altered sequence is passed into the self-contrastive network. The approach relies on attention scores to quantify token relevance. As [27] suggests, each individual head can learn to perform different tasks. This is why we take into account all layers and heads of the model to get an estimate of the overall importance of each word in the sentence.

Given a set of input sentences $\{x_i\}_{i=1}^m$ as input to unsupervised contrastive learning, we first duplicate x_i to obtain our input pairs $\{(x_i, x_i^+)\}_{i=1}^m$. Then, a matrix of attention scores A with the shape of (L, m, H, S_1, S_2) for the input x_i^+ is computed using a pre-trained Transformer model (i.e., BERT). Note that L and H are the numbers of layers and the number of attention heads for each layer, respectively, while $S_1 = S_2 = S$, where S_1 represents the dimension that contains the attention scores of a token $S_1 = j$ to all of the other tokens in dimension S_2 . To gather attention scores and quantify the importance of each input token, we consider and evaluate two different approaches.



Fig. 1. Illustration of our proposed Attention-Driven Dropout integrated into selfcontrastive learning. ADD utilizes attention scores from a PLM to determine the relevance of each token in the sequence. By aggregating these scores, we obtain a single value for each token, enabling us to identify the least significant k tokens, which are subsequently removed. The modified sequence, along with the original, is then inputted into the self-contrastive network.

Naive Aggregation. We sum all of the attention scores across dimensions L, H and S_1 , which yields a matrix A_N of shape $(m, S_2) = (m, S)$. where

$$a_i = \sum_{l=1}^{L} \sum_{h=1}^{H} \sum_{s=1}^{S_1} A_{lihs}$$

denote the summed attention vector of input sample m.

3.1 Attention Rollout Aggregation

[1] introduce a novel post-hoc method for approximating the attention to input tokens. We incorporate this method in our approach to quantify the overall relevance of each token.

To compute the attention rollout matrix A for a given layer l to layer l^- , we recursively multiply the attention score matrices from all previous layers.

$$\widetilde{A}(l) = \begin{cases} A(l)\widetilde{A}(l^{-}) & \text{if } l > l^{-} \\ A(l) & \text{if } l = l^{-} \end{cases}$$
To combine the matrices \widetilde{A}_l into a final summed attention matrix $A_R \in \mathbb{R}^{m \times S}$, we sum the attention rollout matrices across all L layers, to get a vector a_i for each sample m.

$$a_i = \sum_{l=1}^L \sum_{s=1}^{S_1} \widetilde{A}_{lis}$$

Considering k as a number of tokens to be removed from each input sentence x_i^+ .

3.2 Static Dropout Rate

We define k as a hyperparameter while using the *static* dropout rate mode.

Table 1. Comparison of Attention-Driven Dropout (*naive* aggregation) and random word deletion with example input sentences. Note that the overall meaning of the sentence is preserved with Attention-Driven Dropout while deleting a random word can break the structure.

Input sentence	Attention-Driven Dropout	Random word deletion
We should go to the small Italian restaurant again!	We should go to the small Italian restaurant again!	We should go to the small Italian restaurant again!
Two big dogs are running fast in the park.	e Two big dogs are running fast in the park.	e Two big dogs are running fast in the park.
Mary helped John to style his new apartment.	Mary helped John to style his new apartment.	Mary helped John to style his new apartment.
A brown bear is eating a small fish.	A brown bear is eating a small fish.	A brown bear is cating a small fish.

3.3 Dynamic Dropout Rate

Alternatively, with the dynamic dropout rate, we calculate k based on the number of tokens that are present in the given sequence, excluding padding tokens. Let the number of non-padding tokens be denoted as t_s , then k is given by:

$$k = \lfloor t_s/t \rfloor$$

With this formulation, we are able to remove more redundant information in long sequences, while with the static approach, we are limited to a fixed amount of tokens for any sequence length.

Now let imin(x, k) denote a function that returns a set of indices of the k minimum values of given input vector x. We select the indices of minimum values of each vector a_i by calculating $g_i = imin(a_i, k) \in \mathbb{R}^k$. Let x_{ij}^+ denote the j-th element of the i-th input sentence.

We iterate over each sample *i*. If the sample x_i^+ does not contain at least *t* tokens (excluding padding), we do not remove any words and continue with the

next sample x_{i+1}^+ . Otherwise, we set the values of our input pair instance $x_{ij}^+ = p$ where $j \in g_i$ and p is the padding token, to get our new token sequence x_i^* .

Finally, we reorder x_i^* in a way where all padding tokens in the vector are aligned on the right-hand side, e.g., $(x_{i1}^*, x_{i2}^*, ..., x_{i_*}^*, p, p)$.

The original input sentences and their altered pairs $\{(x_i, x_i^*)\}_{i=1}^m$ are then fed into the SimCSE network, where during training, we minimize a contrastive loss objective. We present examples with the alternative augmentation methods in Table 1.

4 Experiment

We conduct empirical experiments to compare our proposed methods with various baselines and alternative approaches. Our source code is anonymously available.¹

4.1 Datasets and Tasks

We evaluate the performance on seven different semantic textual similarity (STS) task sets, as well as seven transfer learning tasks, similar to [14, 23]. We use the SentEval toolkit from [11] to conduct the evaluation.

Semantic Textual Similarity (STS). The STS task set consists of seven tasks: STS-12-16 [2–6], STS-Benchmark (STS-B) [9] and SICK-Relatedness (SICK-R) [7]. Here we follow the approach and suggestion of the SimCSE authors and use Spearman's correlation as well as the aggregation on all train, development, and test datasets for each task to be able to generate comparable results and unify the evaluation setting. Results on this task set can be found in 2.

Transfer Learning. Additionally, we evaluate the models on various transfer tasks. These consist of MR [22], CR [15], SUBJ [21], MPQA [26,31], TREC [28] and MRPC [13]. Evaluating these tasks requires training a logistic regression classifier on top of (frozen) sentence embeddings. The final results can be seen in Table 3.

4.2 Training Procedure

Our data augmentation is used to alter input sequences as a first step, which are then fed into the SimCSE network. We report on pre-trained transformer models as the backbone, BERT [12] and RoBERTa [20]. For the sentence embedding, we are using the [CLS] representation. We use the same dataset as [14], 10^6 sampled sentences from English Wikipedia. During training, Attention-Driven Dropout is used to alter the input pairs. During the evaluation, we do not change the input sequences.

¹ https://github.com/fstermann/attention-driven-dropout.

All models are trained for 1 epoch, except DiffCES and InfoCSE which are trained for 2 epochs and evaluated on the STS-Benchmark (STS-B) every 125 training steps. We keep the best-performing models based on development performance.

We conduct a grid search across learning rates {3e–5, 1e–5} as well as batch sizes {64, 128, 256, 512} for both backbone architectures. Results of this search are compared in Sect. 5.1. The parameters yielding the best performance with respect to the STS-Benchmark task as well as the average transfer tasks respectively are presented in Table 6. Our results are shown in Table 2 and Table 3 were produced with these configurations.

5 Result and Discussion

The semantic textual similarity evaluation results can be found in Table 2. Our approach performs on par with the other state-of-the-art methods. Using $BERT_{base}$ as the pretrained language model, we can achieve the overall highest average (77.25) across the STS tasks with the *naive* attention aggregation method, improving the raw SimCSE-BERT_{base} by one percentage point. Specifically, we can achieve better results in 5 out of 7 tasks.

Using RoBERTa_{base}, our approach can increase previous results by 0.93% (77.45), yielding best results in 6 out of 7 tasks with the *naive* aggregation.

Table 2. STS task performance for sentence embeddings (Spearman's correlation, "all" setting). The best performance for the corresponding task is marked in bold, the second best is in italics. \heartsuit : results from [14]; other results are evaluated by us.

Model	STS12	STS13	STS14	STS15	STS16	STS-B	SICK-R	Avg.
$\mathrm{Sim}\mathrm{CSE}\text{-}\mathrm{BERT}_{\mathtt{base}}^{\heartsuit}$	68.40	82.41	74.38	80.91	78.56	76.85	72.23	76.25
$+ \text{ ADD}_{\text{naive}}$	71.00	82.24	75.10	82.73	79.03	78.51	72.12	77.25
$+ ADD_{rollout}$	65.20	77.98	71.26	80.62	77.27	76.26	69.68	74.04
$SimCSE$ -Ro $BERTa_{base}$	70.16	81.77	73.24	81.36	80.65	80.22	68.56	76.57
$+ \text{ ADD}_{\text{naive}}$	67.45	83.43	74.67	82.48	81.69	82.00	70.43	77.45
$+ ADD_{rollout}$	65.34	80.97	71.29	81.08	80.34	79.83	69.54	75.48

Transfer task results are presented in Table 3. We present evaluation results for both the *naive* and *rollout* aggregation methods, without and with the additional masked language modeling (MLM) [12] objective used during training. We find that overall, adding this objective increases results on the transfer tasks.

With BERT_{base} , we can achieve the best performance for the *rollout* attention aggregation combined with the MLM objective (87.01). We can either compare with or outperform the results of raw SimCSE without Attention-Driven Dropout.

Table 3. Transfer task performance for sentence embeddings, measures represent accuracy. The best performance for the corresponding task is marked in bold, the second best is in italics. \heartsuit : results from [14]; other results are evaluated by us. MLM: MLM is added as an auxiliary task with $\lambda = 0.1$.

Model	MR	CR	SUBJ	MPQA	SST	TREC	MRPC	Avg.
$\operatorname{SimCSE-BERT}_{\mathtt{base}}^{\heartsuit}$	81.18	86.46	94.45	88.88	85.50	89.80	74.43	85.81
+ MLM $^{\diamond}$	82.92	87.23	95.71	88.73	86.81	87.01	78.07	86.64
$+ \text{ ADD}_{\text{naive}}$	81.82	86.89	94.83	89.43	85.28	89.40	75.25	86.13
+ MLM	82.18	87.74	95.66	88.16	86.55	91.00	75.07	86.62
$+ \text{ ADD}_{\text{rollout}}$	81.41	85.72	94.79	89.32	84.84	88.60	75.07	85.68
+ MLM	82.40	87.97	95.62	89.38	86.93	91.20	75.59	87.01
$SimCSE-RoBERTa_{base}^{\diamond}$	81.04	87.74	93.28	86.94	86.60	84.60	73.68	84.84
$+ \text{ MLM}^{\diamond}$	83.37	87.76	95.05	87.16	89.02	90.80	75.13	86.90
$+ ADD_{naive}$	82.30	88.05	93.70	87.50	88.25	84.60	74.84	85.61
+ MLM	83.86	89.06	94.65	88.27	89.51	90.60	76.75	87.53
$+ \text{ ADD}_{\text{rollout}}$	82.08	88.40	93.13	87.54	87.97	87.00	75.88	86.00
+ MLM	84.68	89.91	94.97	88.37	90.61	92.20	78.43	88.45

For SimCSE-RoBERTa_{base} as the underlying architecture, adding our method can increase performance across almost all tasks, showing an average transfer task accuracy of **88.45**, increasing previous best results by 1.55% points.

We further conducted experiments with different baseline models other than SimCSE, namely InfoCSE [33] and DiffCSE [10]. The results are depicted in tables 8 and 9 in the appendix. Since these baselines are trained for 2 instead of 1 epoch, we compare them independently from the SimCSE baseline.

As depicted in Tables 2 and 3, performance across the different aggregation methods and underlying base models may vary. The main difference in aggregation is that rollout aggregation takes much more account of the attention scores of earlier layers than those of later layers. The semantic features to which different layers pay attention can therefore influence which words are removed from the sequences. Of course, depending on which tokens are removed from the input sequences, the underlying model will contain different contextual information. While rollout may perform worse in STS, it is possible that by focusing on different parts of the sentence, its captured knowledge can be better exploited in transfer tasks compared to naive aggregation.

5.1 Ablation Study

For further comparison, we investigate the performance of our method with previous data augmentation techniques used in unsupervised self-contrastive learning, as well as quantifiable measures of the representational embedding space. Additionally, we evaluate results on different hyperparameter settings for k and t.

Comparison with Other Data Augmentations. Previous data augmentation techniques [25, 35] and even our underlying contrastive network SimCSE [14]



Fig. 2. Comparison of hyperparameter configurations. k: the amount of tokens to drop out; t: minimum amount of sequence-tokens for dropout.

heavily rely on randomness to augment input sequences. While this approach can yield better results than previous state-of-the-art methodologies, using it in combination with our novel attention-based procedure can improve the outcome with minimal training. In Table 4 we compare our method with previously reported non-contrastive augmentations. Adding a random word deletion with static (k = 1, t = 10) or dynamic dropout (t = 10) rate configuration cannot outperform our best performing $\text{BERT}_{\text{base}}$ based model.

In comparison to other recent data augmentation models stemming from SimCSE, our method exhibits comparable performance (PeerCL: 83.8, [32]; ESimCSE: 84.8 [34]). It should be noted that direct comparisons between these methods can be challenging due to inherent differences in their augmentation approaches. While we augment positive pairs similarly to ESimCSE, ESimCSE also modifies negative pairs. On the other hand, PeerCL combines multiple data augmentations simultaneously, whereas our method relies on a single augmentation technique. Integrating our method with PeerCL, for instance, could be a potential avenue for future investigation.

Static vs. Dynamic. To find the optimal configuration hyperparameters k and t and be able to compare their influence, we conducted a grid search across reasonable values $\{1, 2, 3\}$ for k and $\{8, 10, 12\}$ for t, with BERT_{base} as well as RoBERTa_{base} and their 'default' batch size and learning rate taken from the SimCSE paper. Here, we also compare the *static* and *dynamic* approaches, which are described in Sect. 3. We compare results on the STS-B task (development set) and find the overall best performance (83.36) for the naive aggregation, a dynamic dropout setting with t = 12, and the RoBERTa PLM. If we aggregate attention scores with the rollout method, best results (82.20) can be achieved for the same dynamic setting and k = 1, t = 12. The BERT-based architecture

Table 4. Comparison of different data augmentation techniques on the STS-B development set (Spearman). *random static*: random word deletion, with k = 1, t = 10; *random dynamic*: random word deletion with t = 10, number of words depends on sequence length. \heartsuit : results from [14].

Data augmentation			STS-B
Attention-Driven Dropout			83.1
Random static			81.4
Random dynamic			81.9
None (unsup. SimCSE) $^{\diamond}$			82.5
$\operatorname{Crop}^{\heartsuit}$	10%	20%	30%
	77.8	71.4	63.6
Word deletion $^{\diamond}$	10%	20%	30%
	75.9	72.2	68.2
Delete one word $^{\diamond}$			75.9
w/o dropout $^{\diamond}$			74.2
Synonym replacement $^{\heartsuit}$			77.4
MLM $15\%^{\circ}$			62.2

yields the best results for the naive aggregation (83.09) with k = 1, t = 10. A STS-B performance of 81.45 can be achieved with a dynamic setting and t = 8. All results are presented in Table 5.

Figure 2 compares the performance of the different settings. Overall, the naive aggregation method compares better to the advanced rollout aggregation. Comparing different t settings for RoBERTa, we clearly see an increasing performance trend with increasing t. This goes along with our initial intuition that we cannot drop too many tokens if our input sequence is too short in order to preserve the meaning of the sentence. If the value for t is too small, performance drops drastically, especially the more tokens k are dropped from the sequence.

The main difference most likely stems from the different tokenization in BERT (WordPiece) and RoBERTa (BPE) has an impact on the results. The two tokenization methods are similar but can produce tokenized sequences of different lengths. Thus, the hyperparameters t (minimum sequence length) and k (number of tokens to drop) can be dependent on the chosen tokenization.

Aggregation Visualization. In order to compare both aggregation approaches and get an intuition of how the aggregated attention scores are composed, we provide a visual representation of the normalized scores in the form of heatmaps in Fig. 3. Darker colors represent a smaller aggregated attention score. Looking at the first example sentence, we can see that the *naive* aggregation approach yields the lowest score for 'italian', and the second lowest for 'small.' Compared to the results of the *rollout* approach, aggregating by just summing up the attention scores gives a good estimate of the informativeness of the respective token. Breaking down the last example sentence, removing the token with the smallest aggregated score one at a time until we are left with 4 tokens would carry out like this.

```
A brown bear is eating a small fish.
A brown bear is eating a fish.
A bear is eating a fish.
A bear is eating fish.
A bear is eating.
```



Fig. 3. Heatmaps of the (normalized) aggregated attention scores, comparing the two aggregation approaches. Normalized aggregated attention scores for each word in the sentence are presented for the *naive* and *rollout* methods.

We can see that we end up with a syntactically correct sentence at every step, each containing less information than the previous sentence. This is an example showing that our *naive* aggregation approach can produce not only one but many augmentations of a sentence.

Alignment and Uniformity. [29] describe a way of measuring the representation of the embedding space. They propose two metrics: alignment, which represents the closeness of features from positive pairs, and uniformity of the feature distribution on the hypersphere.

The ADD-BERT_{base} model can achieve the best alignment (0.19) out of the unsupervised methods, suggesting that our approach indeed can yield a better representation of positive pairs, moving them closer together in the embedding space. In terms of uniformity (-2.43), ADD performs slightly worse than unsupervised SimCSE without Attention-Driven Dropout. A comparison of our results with other approaches can be found in Fig. 4.

By deleting words from half of the sentences, some contextual information is lost, which on the one hand is responsible for the improved alignment measure, since we introduce more noise with the data augmentation. On the other hand, this leads to slightly less information on the preserved data in half of the cases, which is most likely responsible for the decrease in uniformity.

Hyperparameters. Best results for our base comparison for different hyperparameter settings of k and t are shown in Table 5.

All hyperparameters, including batch size and learning rate on which performance is reported in Sect. 5 are shown in Table 6.



Fig. 4. Comparison of $\ell_{alignment}$ and $\ell_{uniformity}$ of BERT_{base} based models (lower is better). Point colors represent their average STS performance (Spearman); + represents supervised methods; results except ours from [14].

Table 5. Comparison of hyperparameter configurations. k: the amount of tokens to drop out; t: minimum amount of sequence-tokens for dropout.

			BERT	1	RoBERTa		
k		t	naive	rollout	naive	rollout	
static	1	8	80.43	80.72	83.13	80.35	
	1	10	83.09	78.89	83.20	81.27	
	1	12	81.95	80.76	83.27	82.20	
	2	8	80.72	77.88	77.21	75.46	
	2	10	82.49	75.41	81.60	77.57	
	2	12	79.48	78.94	82.46	80.41	
	3	8	79.18	72.64	64.01	69.09	
	3	10	81.18	75.84	75.48	72.31	
	3	12	82.54	78.48	80.04	77.11	
dynamic	-	8	81.84	81.45	82.08	79.97	
	-	10	81.63	80.16	82.92	81.21	
	-	12	80.29	79.93	83.36	82.20	

Table 6. Optimal hyper-parameters for the STS and Transfer task set respectively: Batch size (BS), Learning Rate (LR) for PLM and Attention Aggregation (AA) settings.

			\mathbf{S}'			Transfer		
PLM	AA	k	t	BS	LR	BS	\mathbf{LR}	
BERT	naive	1	10	64	3e-5	64	1e-5	
	rollout	dynamic	8	64	3e-5	512	1e-5	
RoBERTa	naive	dynamic	12	64	1e-5	256	1e–5	
	rollout	dynamic	12	64	1e-5	128	1e-5	

 Table 7. Average runtime in minutes (rounded to the nearest integer) for different batch size settings.

Batch Size	64	128	256	512
Avg. Runtime (m)	101	53	31	22

Table 8. STS task performance for sentence embeddings (Spearman's correlation, "all" setting). The best performance for the corresponding task is marked in bold, the second best is in italics. \blacklozenge : results from [33]; \diamondsuit : results from [10]; other results are evaluated by us.

Model	STS12	STS13	STS14	STS15	STS16	STS-B	SICK-R	t Avg.
InfoCSE-BERT _{base}	70.53	84.59	76.40	85.10	81.95	82.00	71.37	78.85
$+ ADD_{naive}$	70.15	82.56	74.55	82.44	82.37	81.23	71.50	77.83
$+ ADD_{rollout}$	65.05	79.96	69.89	79.71	79.28	77.02	70.00	74.42
$\mathrm{DiffCSE}\operatorname{-BERT}_{\mathtt{base}}^{\diamond}$	72.28	84.43	76.47	83.90	80.54	80.59	71.23	78.49
$+ ADD_{naive}$	70.87	83.49	75.02	83.73	79.39	79.55	72.24	77.76
$+ \text{ ADD}_{\text{rollout}}$	62.46	78.74	69.47	79.26	77.16	74.32	72.23	73.38
DiffCSE-RoBERTa _{base} \diamond	70.05	83.43	75.49	82.81	82.12	82.38	71.19	78.21
$+ \text{ ADD}_{\text{naive}}$	66.96	82.47	74.31	81.26	81.21	81.57	70.46	76.89
$+ \text{ ADD}_{\text{rollout}}$	66.37	81.09	72.38	81.05	79.89	79.8	69.76	75.76

Runtime. We conducted all of our experiments on NVIDIA DGX A100 nodes (Table 7).

We also consider different baseline models (InfoCSE [33], DiffCSE [10]) and conduct the training procedure for 2 epochs. This differs from SimCSE, where 1 epoch was used for training. We use the paper's default hyperparameters, as well as our hyperparameters found to work best with the SimCSE baseline.

In addition, we conducted experiments with different random seed settings for the STS task set, reported in table Table 10.

Table 9. Transfer task performance for sentence embeddings, measures represent accuracy. The best performance for the corresponding task is marked in bold, the second best is in italics. \blacklozenge : results from [33]; \diamondsuit : results from [10]; other results are evaluated by us. MLM: MLM is added as an auxiliary task with $\lambda = 0.1$.

Model	MR	\mathbf{CR}	SUBJ	MPQA	SST	TREC	MRPC	Avg.
InfoCSE-BERT _{base} •	81.76	86.57	94.90	88.86	87.15	90.60	76.58	86.63
$+ ADD_{naive}$	81.60	86.39	94.93	88.60	86.77	89.00	76.23	86.22
$+ ADD_{rollout}$	81.81	87.00	95.19	88.67	87.15	90.00	78.03	86.84
$\mathrm{DiffCSE}\text{-}\mathrm{BERT}_{\mathtt{base}}^{\diamond}$	82.69	87.23	95.23	89.28	86.60	90.40	76.58	86.86
$+ ADD_{naive}$	80.71	84.90	94.48	88.65	85.34	87.00	75.94	85.29
$+ ADD_{rollout}$	82.28	86.76	95.06	88.96	86.60	89.80	77.22	86.67
DiffCSE-RoBERTa _{base}	82.82	88.61	94.32	87.71	88.63	90.40	76.81	87.04
$+ \text{ ADD}_{\text{naive}}$	82.85	88.13	94.00	87.91	88.69	90.40	77.16	87.02
$+ ADD_{rollout}$	81.89	87.31	93.64	87.10	88.74	91.20	76.35	86.60

Table 10. STS task performance for sentence embeddings (Spearman's correlation, "all" setting). Results are based on multiple runs, using 5 different random seeds (42–46).

Model	STS12	STS13	STS14	STS15	STS16	STS-B	SICK-R	Avg.
SimCSE-BERT _{base} + ADD _{naive}	68.40 ± 1.83	80.72 ± 1.87	73.27 ± 2.04	81.13 ± 1.07	78.23 ± 0.73	77.32 ± 0.85	70.49 ± 1.31	75.65 ± 1.24

6 Conclusion

In this paper, we introduced Attention-Driven Dropout, a simple yet powerful data augmentation to generate better sentence embeddings in self-supervised contrastive learning. We examined and showed that our method could improve the overall performance of previous augmentation techniques by utilizing two unique approaches to quantifying the relevance of sequence tokens. It is to be noted that our approach is applicable to any self-contrastive network and is meant to be used in combination with different model architectures. It can even be utilized for a number of other tasks, such as generating different views of a sentence While we investigated attention aggregation in the context of deleting words, it may also be useful to utilize the scores to quantify overall important words in a sentence. Using this approach to reduce a sentence to the essentials might be a promising research direction. Overall, this novel approach introduces a method that can be translated to many different tasks and should aid as a starting point for further research in data augmentation.

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AEMLO: AutoEncoder-Guided Multi-label Oversampling

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Abstract. Class imbalance significantly impacts the performance of multi-label classifiers. Oversampling is one of the most popular approaches, as it augments instances associated with less frequent labels to balance the class distribution. Existing oversampling methods generate feature vectors of synthetic samples through replication or linear interpolation and assign labels through neighborhood information. Linear interpolation typically generates new samples between existing data points, which may result in insufficient diversity of synthesized samples and further lead to the overfitting issue. Deep learning-based methods, such as AutoEncoders, have been proposed to generate more diverse and complex synthetic samples, achieving excellent performance on imbalanced binary or multi-class datasets. In this study, we introduce AEMLO, an AutoEncoder-guided Oversampling technique specifically designed for tackling imbalanced multi-label data. AEMLO is built upon two fundamental components. The first is an encoder-decoder architecture that enables the model to encode input data into a low-dimensional feature space, learn its latent representations, and then reconstruct it back to its original dimension, thus applying to the generation of new data. The second is an objective function tailored to optimize the sampling task for multi-label scenarios. We show that AEMLO outperforms the existing state-of-the-art methods with extensive empirical studies.

Keywords: Multi-label classification \cdot Class imbalance \cdot Oversampling \cdot AutoEncoder

1 Introduction

In the field of multi-label classification (MLC), each instance can belong to multiple labels simultaneously. MLC is widely used in various fields, including image annotation [4], sound processing [16], biology [34] and text classification [14]. The issue of class imbalance in multi-label classification has gained prominence recently [28]. It is prevalent in real-world MLC problems and significantly affects classifier performance, as many algorithms assume data is balanced. Imbalanced datasets tend to bias learners towards majority labels [29].

1.1 Research Goal

Our goal is to address the class imbalance in multi-label datasets through the integration of a deep generative model within an encoder-decoder architecture. This strategy seeks to outperform conventional methods, such as sampling with linear interpolation or random replication, by dynamically creating instances that contain richer feature information.

1.2 Motivation

In recent years, innovative approaches have been developed to tackle the issue of imbalance in multi-label learning [28], including sampling methods [6,17], classifier adaption [9], and ensemble techniques [18,27]. Sampling methods, in particular, aim to balance the dataset before the training phase, offering flexibility and compatibility with any multi-label classifier. To ensure effective sampling, several studies have concentrated on identifying specific samples and refining decision boundaries. For example, MLSOL [17] assigns a higher selecting probability to the sample suffering severe local imbalance. MLBOTE [29] refines the boundary samples related to high imbalance labels and employs different sampling strategies. Traditional oversampling techniques often rely on basic linear interpolation or replication for creating feature vectors of synthesized samples, with label vectors typically generated through majority voting or replication.

The Autoencoder (AE) and Generative Adversarial Network (GAN), as exemplary generative models, have shown substantial potential in data generation, restoration, and augmentation [8, 12, 15, 19, 22]. An Autoencoder compresses data into a latent space using an encoder and reconstructs it by a decoder. Its objective is to minimize reconstruction errors, enabling efficient feature extraction and noise reduction [13, 15]. Although Autoencoder and GAN are used to address the imbalance problem and generate minority samples, they primarily cater to single-label datasets and face several challenges when applied to multi-label datasets. First, AE and GAN require training samples with identical labels (same class in the single-label dataset or same label set in the multi-label dataset). However, in multi-label data, the number of samples with a complete label set is often too limited to effectively train deep learning models. Secondly, although multi-label datasets can be divided into several binary datasets via the One vs All strategy, For each binary dataset, we can learn and reconstruct new feature vectors for multi-label data through end-to-end models, but we can not determine appropriate complete label set for each feature vector (Fig. 1).

1.3 Summary

In this work, we introduce an innovative approach crafted to tackle the class imbalance issue in multi-label datasets named AutoEncoder-guided Multi-Label Oversampling (AEMLO). The core of AEMLO's design lies in two essential elements:



Fig. 1. Using Autoencoders to train data.

- 1. The basic encoder-decoder architecture is designed to encode data into a lower-dimensional space and subsequently reconstruct it, making it suitable for oversampling applications.
- 2. A specialized objective function for multi-label imbalance data sampling.

Our approach incorporates the sampling process into a deep encoder-decoder framework that has been pre-trained, providing a holistic solution for the creation of low-dimensional data representations and synthetic instances through an end-to-end methodology. By augmenting the original training set with instances generated via AEMLO, we further train various traditional multi-label classifiers and conduct comparisons against several multi-label sampling techniques. Experimental results consistently demonstrate the superiority of our method. Our code can be found in https://github.com/CquptZA/AEMLO.

2 Related Work

2.1 Multi-label Classification

Formally, let $\mathcal{X} \in \mathbb{R}^d$ represent the *d*-dimensional feature space, and let $L = \{l_1, l_2, \ldots, l_q\}$ denote a set of *q* predefined labels. In multi-label classification, our objective is to construct a mapping function $h : \mathcal{X} \to L$ based on a given multi-label training dataset $D = \{(\mathbf{x}_i, \mathbf{y}_i)\}_{i=1}^n$, where each sample $\mathbf{x}_i \in \mathcal{X}$ is associated with a binary label vector $\mathbf{y}_i \in \{0, 1\}^q$. Here, \mathbf{y}_i is a binary vector where each element denotes whether the associated label from L is relevant (1) or not relevant (0) to \mathbf{x}_i .

In Multi-Label Classification (MLC), methods are split into three types based on how they handle label correlations. First-order strategies like MLkNN [35] and BR [3] treat labels independently, offering simplicity and efficiency. Second-order methods, such as CLR [11], analyze pairwise label correlations for improved interaction understanding. For complex scenarios with intricate label relationships, high-order methods, like RAkEL [31] and ECC [23], are more effective. RAkEL tackles this by dividing labels into subsets for diverse interaction modeling. ECC sequentially links classifiers, allowing each to learn from the predictions of its predecessors.

2.2 Multi-label Imbalance Learning

Let $N_j^1(N_j^0)$ denote the number of instances with "1" ("0") class of label l_j . *IRlbl* and ImR are the two measures to evaluate the imbalance level of individual labels

[6,32]. Let $N_{max}^1 = max\{N_j^1\}_{j=1}^q$ be the number of "1"s in the most frequent label, $IRlbl_j$ and ImR are defined as:

$$IRlbl_{j} = N_{max}^{1}/N_{j}^{1} \quad ImR_{j} = max(N_{j}^{1}, N_{j}^{0})/min(N_{j}^{1}, N_{j}^{0})$$
(1)

The larger the $IRlbl_j$ and ImR_j , the higher the imbalance level of l_j . Then, MeanIR calculates the average imbalance ratio (IRlbl) across all labels in a dataset, defined by: $\frac{1}{q} \sum_{j=1}^{q} IRlbl_j$, where q is the total number of labels. The higher MeanIR, the imbalance of the dataset. By considering the IRlbl and MeanIR, we can calculate imbalance indicators such as the coefficient of variation of IRlbl (CVIR) and concurrency level (SCUMBLE) [28].

The imbalanced approaches proposed for MLC can be divided into three categories: sampling methods, classifier adaptation [9, 32, 33], and ensemble approaches [18,27]. Compared to the other two methods, the sampling method is more universal, as it creates (deletes) instances related to minority (majority) labels to construct a balanced training set that can be used to train any classifier without suffering from bias. Sampling methods involve undersampling and oversampling techniques. Undersampling reduces the presence of majority labels by either randomly removing instances or employing heuristic approaches to selectively eliminate samples. For example, LPRUS and MLRUS [25] aim to alleviate imbalances by respectively targeting the most frequent label sets or individual labels for removal. Conversely, oversampling techniques such as LPROS and the MLSMOTE [6] focus on augmenting the dataset with new instances associated with minority labels, either through duplication or the generation of synthetic samples. Recent developments include the REMEDIAL [5] method, which adjusts label and feature spaces to lessen label co-occurrence and improve sampling. Integrating this method with techniques such as MLSMOTE can further optimize dataset balancing [7]. MLSOL [17] specifically generates instances to focus on local imbalances in datasets. On the other hand, MLTL [21] refines datasets by removing instances that obscure class boundaries. Another notable method, MLBOTE [29], categorizes instances based on their boundary characteristics and applies different sampling rates.

2.3 Deep Sampling Method

Traditional sampling techniques struggle to effectively expand the training set for complex models. This has sparked interest in generative models and their potential to mimic oversampling strategies [2, 10]. Utilizing an encoder-decoder setup, artificial instances can be effectively introduced into an embedding space. AE [13,15] and GAN [12] have been effectively employed to capture the underlying distribution of data and further applied to generate data for oversampling purposes. AE is designed to learn efficient data codings in an unsupervised manner. Essentially, they aim to capture the most salient features of the data by compressing the input into a lower-dimensional latent space and then reconstructing it back to the original dimensionality. The core objective of an AE is defined by the reconstruction error, which quantifies the difference between the original data and its reconstruction. Unlike Variational Autoencoder (VAE), which incorporates the Kullback-Leibler (KL) divergence to regulate the latent space, standard AE relies solely on the reconstruction loss. This encourages the model to develop a compressed representation that retains as much of the original information as possible, enabling the AE to generate reconstructions that are as close to the input. For example, DeepSMOTE [8] integrates traditional SMOTE methods into encoding and decoding architectures similar to AE. VAE strive to maximize a variational lower bound on the data's log-likelihood. Typically, they are formulated by merging a reconstruction loss with the KL divergence. The KL divergence serves as an indirect penalty for the reconstruction loss, steering the model towards a more faithful replication of the data distribution [22]. By penalizing the reconstruction loss, the model is motivated to refine its replication of the data, thereby enabling it to produce outputs rooted in the input's latent distribution. GAN has significantly advanced the field of computer vision by framing image generation as a competitive game between a generator and a discriminator network [19, 20, 37]. Despite their remarkable achievements, GAN requires the deployment of two separate networks, can encounter training difficulties, and are susceptible to mode collapse [8].

3 Multi-label AutoEncoder Oversampling

3.1 Method Description and Overview

The multi-label AutoEncoder oversampling framework, as described in Algorithm 1, is divided into the training process and the instance generation phase.

In the training process, as shown in Fig. 2, the model is designed to learn and optimize four distinct mapping functions: the feature encoding function \mathbf{F}_{ex} , label encoding function \mathbf{F}_{ey} , feature decoding function \mathbf{F}_{dx} , and label decoding function \mathbf{F}_{dy} . The model is trained end-to-end with mini-batches and the Adam optimizer, where batch size *n* encompasses the feature vector \mathbf{x}_i and binary label vector \mathbf{y}_i of the *i*-th sample, respectively. The matrices \mathbf{X} and \mathbf{Y} aggregate the input features and labels for all samples in the batch. The framework ingests a feature matrix \mathbf{X} and its corresponding label matrix \mathbf{Y} , aiming to output reconstructed versions of \mathbf{X}' and \mathbf{Y}' . Meanwhile, The other goal of our model is to identify an optimal latent space \mathcal{L} , where the Deep Canonical Correlation Analysis (DCCA) component [1] enhances the correlation between \mathbf{X} and \mathbf{Y} . Therefore, the model's objective function is defined as:

$$\Theta = \min_{\mathbf{F}_{ex}, \mathbf{F}_{ey}, \mathbf{F}_{dx}, \mathbf{F}_{dy}} \Phi(\mathbf{F}_{ex}, \mathbf{F}_{ey}) + \alpha \Psi(\mathbf{F}_{ex}, \mathbf{F}_{dx}) + \beta \Gamma(\mathbf{F}_{ey}, \mathbf{F}_{dy})$$
(2)

where $\Phi(\mathbf{F}_{ex}, \mathbf{F}_{ey})$ denotes the latent space loss, $\alpha \Psi(\mathbf{F}_{ex}, \mathbf{F}_{dx})$ and $\beta \Gamma(\mathbf{F}_{ey}, \mathbf{F}_{dy})$ signify the reconstruction losses. Here, α and β serve to balance these components, respectively. In Sect. 3.2, we will explain every term of the objective function in details. At the end of each epoch, we enter a validation phase, adjusting the threshold for binary label conversion by maximizing the F-measure of each label on the validation set.

Algorithm 1: Using Encoder and Decoder for Multi-Label Sampling

Input: Original dataset $D = (\mathbf{X}, \mathbf{Y})$, sample count *num*, parameter α, β , latent space dimension l**Output:** Balanced training set D'/* train the Encoder and Decoder */ 1 for $e \leftarrow 1$ to epochs do for $batch(\hat{\mathbf{X}}, \hat{\mathbf{Y}}) \leftarrow B$ do 2 $\mathbf{F}_{ex}((\mathbf{\hat{X}})), \mathbf{F}_{ey}((\mathbf{\hat{Y}}));$ /* encode batch data to \mathcal{L} */ 3 $\mathbf{F}_{dx}((\mathbf{\hat{X}})), \mathbf{F}_{dy}((\mathbf{\hat{Y}}));$ /* decode batch data from \mathcal{L} */ 4 Define the loss function by Eq 2; 5 Compute gradients and update parameters with Adam; 6 7 Update T: /* validate and optimize the bipartition threshold for each label */ /* generate instances */ s while num > 0 do /* choose seed instance */ $\mathbf{x}_s \leftarrow \text{select form } M;$ 9 /* encode */ 10 $\mathbf{F}_{ex}(\mathbf{x}_s)$; $\mathbf{x}_{\mathbf{g}} \leftarrow \mathbf{F}_{dx}(\mathbf{F}_{ex}(\mathbf{x}_{\mathbf{s}})) \quad \mathbf{y}_{\mathbf{g}} \leftarrow \mathbf{F}_{dy}(\mathbf{F}_{ex}(\mathbf{x}_{\mathbf{s}}));$ /* decode */ 11 $\mathbf{y}_{\mathbf{g}} \leftarrow T;$ /* rounding */ 12 $D'=D' \cup (\mathbf{x}_{\mathbf{g}}, \mathbf{y}_{\mathbf{g}});$ 13 $num \leftarrow num - 1$; 14 15 return D'

After the model training is completed, we proceed with instance generation. Let num represent the required number of instances to be generated, and p denote the sampling rate. Further details on the sampling process can be found in Sect. 3.3.

3.2 Loss Function

co

Joint Embedding. To calculate $\Phi(\mathbf{F}_{ex}, \mathbf{F}_{ey})$ defined in Eq. 2, we employ the DCCA to embed features and labels into a shared latent space simultaneously and rewrite the correlation-based $\Phi(\mathbf{F}_{ex}, \mathbf{F}_{ey})$ as the following deep version:

$$\Phi(\mathbf{F}_{ex}(\mathbf{X}), \mathbf{F}_{ey}(\mathbf{Y})) = \|\mathbf{F}_{ex}(\mathbf{X}) - \mathbf{F}_{ey}(\mathbf{Y})\|_{F}^{2} = \operatorname{Tr}(\mathbf{C}_{1}^{T}\mathbf{C}_{1}) + \lambda \operatorname{Tr}(\mathbf{C}_{2}^{T}\mathbf{C}_{2} + \mathbf{C}_{3}^{T}\mathbf{C}_{3})$$
(3)

where

$$C_{1} = F_{ex}(\mathbf{X}) - F_{ey}(\mathbf{Y}),$$

$$C_{2} = F_{ex}(\mathbf{X})F_{ex}(\mathbf{X})^{T} - \mathbf{I},$$

$$C_{3} = F_{ey}(\mathbf{Y})F_{ey}(\mathbf{Y})^{T} - \mathbf{I},$$
nstraint : $F_{ex}(\mathbf{X})F_{ex}(\mathbf{X})^{T} = F_{ey}(\mathbf{Y})F_{ey}(\mathbf{Y})^{T} = \mathbf{I}$
(4)



Fig. 2. The architecture of the proposed autoencoder learns the latent space \mathcal{L} through the function of \mathbf{F}_{ex} and \mathbf{F}_{ey} , and decouples \mathcal{L} through \mathbf{F}_{dx} and \mathbf{F}_{dy} .

 \mathbf{C}_1 quantifies the discrepancy between the feature and label embeddings, while \mathbf{C}_2 and \mathbf{C}_3 assess how each embedded space diverges from orthonormality. The goal is to minimize these discrepancies to align the embeddings of \mathbf{X} and \mathbf{Y} closely, ensuring they remain orthonormal as dictated by the constraint. The identity matrix $\mathbf{I} \in \mathbb{R}^{l \times l}$, serves as a benchmark for achieving this orthonormality, where l denotes the latent space dimension. Integrating DCCA in our sampling framework not only enables a unified embedding of features and labels but also allows for their precise reconstruction from shared space through the functions $\Psi(\mathbf{F}_{ex}(\mathbf{X}), \mathbf{F}_{dx}(\mathbf{X}))$ for features and $\Gamma(\mathbf{F}_{ey}(\mathbf{Y}), \mathbf{F}_{dy}(\mathbf{Y}))$ for labels.

Feature Reconstruction. The function Ψ is composed of two distinct components: the feature reconstruction error, \mathcal{M} , and the instance similarity metric, \mathcal{S} . It is defined as:

$$\Psi(\mathbf{F}_{ex}(\mathbf{X}), \mathbf{F}_{dx}(\mathbf{X})) = \mathcal{M} + \lambda \mathcal{S}$$
(5)

where λ is a regularization parameter that balances the contribution of the similarity metric S relative to the reconstruction error \mathcal{M} .

The reconstruction error \mathcal{M} , quantified as mean squared error, is calculated as:

$$\mathcal{M} = \sum_{i=1}^{n} (\mathbf{x}_{i}' - \mathbf{x}_{i})^{2}$$
(6)

with \mathbf{x}'_i representing the reconstruction of the input \mathbf{x}_i , generated by the \mathbf{F}_{dx} applied to the encoded representation $\mathbf{F}_{ex}(\mathbf{x}_i)$.

The similarity metric S ensures that the proximity between original instances is maintained after reconstruction, thereby conserving the integrity of the feature space. This metric is formulated as: 114 A. Zhou et al.

$$S = \frac{1}{n(n-1)} \sum_{i,j=1, i \neq j}^{n} \left((\mathbf{x}_i - \mathbf{x}_j)^2 - (\mathbf{x}'_i - \mathbf{x}'_j)^2 \right)^2$$
(7)

which measures the squared differences in distances between all pairs of original instances $(\mathbf{x}_i, \mathbf{x}_j)$ and their corresponding reconstructed pairs $(\mathbf{x}'_i, \mathbf{x}'_j)$, ensuring the model preserves instance similarity in its learned feature space. The combination of \mathcal{M} and \mathcal{S} ensures an optimal balance between high-fidelity feature reconstruction and the preservation of relative distances among data within the feature space.

Label Reconstruction. The function Γ encapsulates ranking loss to help the model retrieve label vectors from the shared embedding space:

$$\Gamma(\mathbf{F}_{ey}(\mathbf{Y}), \mathbf{F}_{dy}(\mathbf{Y})) = \sum_{i=1}^{n} \left(\frac{E_i}{|Y_i| \times |\bar{Y}_i|} \right)$$
(8)

where E_i is defined as the set of label pairs (y_{ij}, y_{ik}) that satisfy the condition $f(\mathbf{x}_i, y_{ij}) \leq f(\mathbf{x}_i, y'_{ij})$, with these label pairs belonging to the Cartesian product of the set of positive labels Y_i and the set of negative labels \bar{Y}_i . Here, Y_i represents the set of positive labels, while \bar{Y}_i represents the set of negative labels.

3.3 Generate Instances and Post-processing

Let $L_s = \{l_j \mid ImR_j > 10, IRlbl_j > MeanIR\}^1$ be the set comprising *m* minority labels [29] and $M = \{(\mathbf{x}_i, \mathbf{y}_i) \mid y_{ij} = 1, l_j \in L_s\}$ be the minority instance set associated any labels in L_s . Then, we randomly select a seed sample $(\mathbf{x}_i, \mathbf{y}_i)$ from *M* to initiate the sampling process through forward inference. As shown in Fig. 3, the process encodes the feature vector \mathbf{x}_s into a latent space by $\mathbf{F}_{ex}(\mathbf{x}_s)$, then decodes it to the feature and label vectors of the new instance by $\mathbf{F}_{dx}(\mathbf{F}_{ex}(\mathbf{x}_s))$ and $\mathbf{F}_{dy}(\mathbf{F}_{ex}(\mathbf{x}_s))$, respectively. Specifically, we employ a predefined threshold set *T* to transform the decoded numerical label vector into a binary label vector. After the process, we remove any instances where the generated label vector is entirely zeros to ensure each instance contributes meaningfully to the dataset.

4 Experiments and Analysis

4.1 Datasets

We evaluate our proposed model across 9 benchmark multi-label datasets spanning diverse domains, such as text, images, and bioinformatics [30]. Each dataset is characterized by a set of statistics and imbalance metrics, which include the

¹ Here, 10 is a hyperparameter. We refer to the suggestions in [29] for the selection.



Fig. 3. The process of generating instances

number of instances (n), feature dimensions (d), labels (q), average label cardinality per instance Card(q), and label density Den(q). A comprehensive explanation of these statistical measures and imbalance metrics is available in [28,36]. In the experiment, 20% training data is split as the validation set, which is used to establish thresholds for accurate prediction of final labels (Table 1).

Dataset	Domain	n	d	q	Card(q)	Den(q)	MeanIR	CVIR
bibtex	text	7395	1836	159	2.40	0.02	12.50	0.41
enron	text	1702	1001	53	3.38	0.06	73.95	1.96
Languagelog	text	1460	1004	75	15.93	0.21	5.39	0.78
yeast	biology	2417	103	14	4.24	0.30	7.20	1.88
rcv1	text	6000	472	101	2.88	0.03	54.49	2.08
rcv2	text	6000	472	101	2.63	0.03	45.51	1.71
rcv3	text	6000	472	101	2.61	0.03	68.33	1.58
cal500	music	502	68	174	26.04	0.15	20.58	1.09
Corel5k	images	5000	499	374	3.52	0.01	189.57	1.53

 Table 1. Characteristics of the experimental datasets

4.2 Experiment Setup

In AEMLO, \mathbf{F}_{ex} and \mathbf{F}_{ey} are comprised of two fully connected layers, whereas \mathbf{F}_{dx} and \mathbf{F}_{dy} adopt a single fully connected layer structure. Each layer within these components incorporates 512 neurons and incorporates a leaky ReLU activation function to introduce nonlinearity. The parameters α and β of objective function are explored within the range of $[2^{-4}, 2^{-3}, \cdots, 2^4]$.

Compared Sampling Methods. We compare our proposed sampling method with the following sampling methods.

- **MLSMOTE**: MLSMOTE extends the classical SMOTE method to multilabel data.[Parameter : Ranking, k = 5]
- **MLSOL**: MLSOL considers local label imbalance and employs weight vectors and type matrices for seed instance selection and synthetic instance generation. [*Parameter* : $p \in (0.1, 0.3, 0.5, 0.7, 0.9)$, k = 5]
- **MLROS**: MLROS executes replicating instances associated with minority labels. [*Parameter* : $p \in (0.1, 0.3, 0.5, 0.7, 0.9)$]
- MLRUS: MLRUS executes removing instances associated with majority labels. [*Parameter* : $p \in (0.1, 0.2, 0.3)$]
- **MLTL**: MLTL identifies and removes Tomek-Links in multi-label data by considering the set of instances associated with each minority label. [*Parameter* : k = 5]
- **MLBOTE**: MLBOTE divides instances into three categories, and determines specific instance weights and sampling rates for each group.

Base Multi-lable Classifiers. We use all sampling methods on the following five multi-label classifiers.

- Binary Relevance [3]: BR transforms the multi-label classification problem into multiple independent binary classification tasks, each of which corresponds to one label and trains a binary classifier. Base binary classifier: SVM.
- Multi-label k-Nearest Neighbors [35]: MLkNN is an extension of the k-Nearest Neighbors (kNNs) algorithm for multi-label classification. hyperparameter configuration: k=10.
- Random k-labELsets [31]: RAkEL divides the entire label set into several random subsets containing at least three labels and encodes each subset as a multi-class dataset by treating each label combination as a class. hyper-parameter configuration: k=3, n=2q, base binary classifier: C4.5 Decision Tree.
- Ensemble of Classifier Chain [23]: ECC is an approach that extends the Classifier Chain further in an ensemble framework. hyperparameter configuration: N = 5, base binary classifier: C4.5 Decision Tree.
- Calibrated Label Ranking [11]: CLR transforms the multi-label learning problem into the label ranking problem. Base binary classifier: SVM

Evaluation Metrics. To assess the efficacy of the batch method in multi-label classification, three commonly utilized evaluation metrics are adopted, comprising Macro-F, Macro-AUC, Ranking Loss. Please refer to [36] for detailed definitions of these metrics (Figs. 4, 5 and 6).



Fig. 4. The performance of the multi-label sampling methods in terms of Macro-F across five different classification methods.

4.3 Experimental Analysis

Table 2 presents the average rankings of each base classifier combined with sampling methods across all datasets. Additionally, the Friedman test was utilized to verify the significant superiority/inferiority of our method compared to other sampling approaches across three evaluation metrics in five basic multi-label classification methods. The detailed results of the comparative sampling methods using five fundamental learners on the Macro-F, Macro-AUC, and Ranking Loss



Fig. 5. The performance of the multi-label sampling methods in terms of Macro-AUC across five different classification methods.

are shown in Github. The Origin represents training directly using the training set without any sampling methods. The results indicate that the AEMLO method achieves the highest average ranking in almost all metrics, securing the most significant victories without any substantial losses. It is observed that MLBOTE and MLSOL outperform MLSMOTE, reflecting that refining rule selection for seed instances is more effective than oversampling with all minority seeds directly. An interesting observation is that the performance of MLTL and MLRUS is even worse than that of the original dataset. This is primarily



Fig. 6. The performance of the multi-label sampling methods in terms of Ranking Loss across five different classification methods.

attributed to the removal of critical instances, leading to the loss of important information.

Autoencoders excel at generating new samples by learning compressed representations of input data. However, a subtle challenge arises when the feature space of these generated samples diverges from that of the origin samples. This divergence may pose difficulties for the MLkNN, which relies heavily on the distances between samples within the feature space to identify nearest neighbors. As such, any significant discrepancy in the feature distribution between generated and origin samples could potentially impact the MLkNN to accurately

Table 2. The mean ranking of different sampling methods evaluated with five base learners across three metrics is presented. The notation (n1/n2) indicates adjustments from the Friedman test at a 5% significance level, signifying that the method in question significantly outperforms n1 methods and is outdone by n2 methods. The top-performing method is emphasized in bold, with lower rankings indicating superior performance.

		Origin	MLSMOTE	MLSOL	MLROS	MLRUS	MLTL	MLBOTE	AEMLO
Macro-F	BR	6.33(0/3)	5.11(0/3)	2.89(4/1)	4.22(1/0)	6.00(0/3)	7.33(0/5)	2.78(4/0)	1.33(6/0)
	MLkNN	5.33(0/4)	3.11(3/0)	1.56(4/0)	3.89(2/0)	6.89(0/4)	7.67(0/5)	4.56(1/1)	3.00(3/0)
	RAkEL	5.67(0/3)	4.11(2/2)	2.67(2/0)	4.89(2/2)	7.22(0/5)	7.00(0/5)	2.44(3/0)	2.00(4/0)
	ECC	5.11(0/2)	4.22(0/0)	3.56(2/0)	5.67(0/2)	5.44(0/2)	6.67(0/3)	2.78(3/0)	2.56(4/0)
	CLR	6.33(0/3)	3.89(2/0)	2.78(3/0)	3.00(2/0)	6.89(0/4)	7.11(0/5)	3.89(2/0)	2.11(3/0)
	Avg(Total)	5.75(0/15)	4.09(7/5)	2.69(15/1)	4.33(7/4)	6.49(0/18)	7.16(0/23)	3.29(13/1)	2.20(20/0)
Macro-AUC	BR	5.00(0/2)	4.78(0/1)	3.00(3/0)	4.11(1/0)	5.78(0/3)	7.56(0/5)	4.33(1/1)	1.44(7/0)
	MLkNN	5.11(0/3)	4.22(2/0)	3.44(3/0)	3.78(2/0)	6.11(0/4)	6.78(0/4)	4.44(1/0)	2.11(3/0)
	RAkEL	5.33(0/2)	4.33(1/0)	2.78(2/0)	3.56(1/0)	4.33(1/0)	7.67(0/6)	4.89(0/0)	3.11(2/0)
	ECC	5.22(0/2)	5.02(0/2)	3.33(3/0)	4.22(1/0)	6.89(0/4)	7.00(0/4)	2.56(3/0)	1.56(5/0)
	CLR	6.00(0/3)	5.00(0/1)	3.16(2/0)	3.67(2/1)	5.56(0/2)	7.44(0/4)	3.33(2/0)	1.67(5/0)
	Avg(Total)	5.33(0/12)	4.71(3/4)	2.98(13/0)	3.87(7/0)	5.73(1/13)	7.29(0/23)	3.91(7/1)	2.18(22/0)
Ranking Loss	BR	4.89(0/1)	6.22(0/2)	4.44(0/0)	4.00(1/0)	5.33(0/2)	6.11(0/3)	3.00(3/0)	2.00(4/0)
	MLkNN	4.56(0/2)	5.44(0/3)	3.56(3/0)	5.33(0/2)	4.00(0/2)	6.89(0/5)	2.89(3/0)	3.33(2/0)
	RAkEL	6.44(0/3)	5.22(0/1)	4.33(1/0)	4.44(0/0)	2.78(2/0)	7.56(0/4)	3.11(2/0)	2.11(3/0)
	ECC	5.00(0/1)	4.56(0/0)	4.78(0/0)	3.67(1/0)	4.67(0/0)	6.33(0/2)	4.11(0/0)	2.89(2/0)
	CLR	5.22(0/2)	4.00(1/1)	4.22(1/1)	4.11(1/0)	5.00(0/2)	7.67(0/5)	3.89(3/0)	1.89(5/0)
	Avg(Total)	5.22(0/9)	5.09(1/7)	4.27(5/1)	4.31(3/2)	4.36(2/6)	6.91(0/19)	3.40(11/0)	2.44(16/0)

classify unseen samples. The enhanced performance of BR and CLR methods on augmented datasets can be attributed to the robustness of SVM and its adeptness at navigating complex decision boundaries. Specifically, SVM is particularly effective at managing the intricacies introduced into the feature space by data synthesized through Autoencoders.

4.4 Parameter Analysis

We investigate the influence of various parameter settings on the performance of ALMLO. We select smaller enron and larger Corel5k as two representative datasets in the parameter analysis.

As shown in Fig. 7(a), the impact of varying sampling rate p on Macro-F and Macro-AUC scores (based on MLkNN) shows a trend of initial fluctuation, followed by stabilization. In contrast, in Fig. 7(b) exhibits a higher sensitivity to p, with significant volatility in Macro-F and inconsistent variations in Macro-AUC. These observations suggest that the optimal selection of p may be highly dependent on dataset characteristics.

Figure 8 illustrates ALMLO's performance sensitivity to variations in α and β , highlighting the importance of balancing feature reconstruction loss with label relevance loss during optimization.



Fig. 7. Performance of sampling rate in terms of Macro-F and Macro-AUC.



Fig. 8. Performance of AEMLO with varying parameter configurations in terms of Macro-F.

4.5 Sampling Time

Figure 9 shows the time efficiency of different sampling methods, with the epoch set as 100 for AEMLO. It is evident that AEMLO, as a deep learning approach, requires training before sampling, resulting in a higher time expenditure.



Fig. 9. Sampling time of different sampling method.

5 Conclusion

In this paper, we introduce AEMLO, an innovative oversampling model devised for addressing data imbalance in multi-label learning by integrating canonical correlation analysis with the encoder-decoder paradigm. AEMLO emerges as an effective oversampling solution for training deep architectures on imbalanced data distributions. It acts as a data-level solution for class imbalance, synthesizing instances to balance the training set and thus enabling the training of any classifiers without bias. AEMLO exhibits the pivotal characteristics crucial for a successful sampling algorithm in the multi-label learning domain: the ability to manipulate features and labels, i.e., to learn low-dimensional joint embeddings from feature and label representations and transform them into an original-dimensional space, along with generating new feature representations and their corresponding label subsets. This is facilitated through the utilization of an encoder/decoder framework. Extensive experimental studies demonstrate the capability of AEMLO to handle imbalanced multi-label datasets in various domains and collaborate with diverse multi-label classifiers.

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MANTRA: Temporal Betweenness Centrality Approximation Through Sampling

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Abstract. We present MANTRA, a framework for approximating the temporal betweenness centrality of all nodes in a temporal graph. Our method can compute probabilistically guaranteed high-quality temporal betweenness estimates (of nodes and temporal edges) under all the feasible temporal path optimalities, presented in the work of Buß et al. (KDD, 2020). We provide a sample-complexity analysis of our method and speed up the temporal betweenness computation using a state-ofthe-art progressive sampling approach based on Monte Carlo Empirical Rademacher Averages. Additionally, we provide an efficient sampling algorithm to approximate the temporal diameter, average path length, and other fundamental temporal graph characteristic quantities within a small error ε with high probability. The running time of such approximation algorithm is $\tilde{\mathcal{O}}(\frac{\log n}{c^2} \cdot |\mathcal{E}|)$, where n is the number of nodes and $|\mathcal{E}|$ is the number of temporal edges in the temporal graph. We support our theoretical results with an extensive experimental analysis on several real-world networks and provide empirical evidence that the MANTRA framework improves the current state of the art in speed, sample size, and required space while maintaining high accuracy of the temporal betweenness estimates.

1 Introduction

Centrality measures are fundamental notions for evaluating the importance of nodes in networks, used in network analysis and graph theory. A centrality measure assigns real values to all the nodes, in such a way that the values are monotonously dependent of the nodes' importance, i.e., more important nodes should have higher centrality scores. Computing the *betweenness centrality* is arguably one of the most important tasks in graph mining and network analysis. It finds application in several fields including social network analysis [38], routing [15], machine learning [37], and neuroscience [16]. The betweenness of a node in a graph indicates how often this node is visited by a shortest path. High betweenness nodes are usually considered to be important in the network. Brandes' algorithm [7], is the best algorithm to compute the exact centrality scores of every node in $\mathcal{O}(n \cdot m)$ time and $\mathcal{O}(n+m)$ space where n and m are the

number of nodes, and edges of a given graph G = (V, E), respectively. Unfortunately, this algorithm quickly becomes impractical on nowadays' networks with billions of nodes and edges. Moreover, there is a theoretical evidence, in form of several conditional lower bounds results [1], for believing that a faster algorithm cannot exists, even for *approximately* computing the betweenness. A further challenge, is that modern real-world networks are also dynamic or temporal, i.e., they change over time. Temporal networks can be informally described as edge-labeled graphs in which each label indicates the time instant in which the underlying edge appears in the network. A great variety of both modern and traditional networks can be naturally modeled as temporal graphs. Furthermore, there are numerous real-world applications for which studying temporal networks offers unique perspectives. This is especially evident when examining data that evolves over time, for example social networks interactions, information/infection spreading, subgraph patterns, detecting communities, and clustering networks. In the context of these challenges it is, thus, essential to consider temporal variants of the most important centrality notions, alongside algorithms for computing them, that have an excellent scaling behavior. In this work, we focus on the temporal version of the *betweenness centrality*, that similarly to static networks, it seeks to pinpoint nodes that are traversed by a significant number of optimal (temporal) paths. Buß et al. [10, 34] gave several definitions of the temporal betweenness as a temporal counterpart of the betweenness centrality, characterized their computational complexity, and provided polynomial time algorithms to compute these temporal centrality measures. However, these algorithms turn out to be impractical, even for medium size networks. Thus, it is reasonable to consider approximation algorithms that can efficiently compute the centrality values of the nodes up to some small error. In this work, we follow the approach of Santoro et al. [35], and we provide a set of approximation algorithms for all the temporal betweenness variants in [10].

Contributions. We propose MANTRA (teMporAl betweeNnes cenTrality thRough sAmpling), a rigorous framework for the approximation of the temporal betweenness of all the vertices and temporal edges in large temporal graphs. In particular, we present the following results: (1) We extend the state-of-theart estimator [35] to all the feasible temporal betweenness centrality variants for nodes and temporal edges (Sect. 4.1). In addition, we propose two alternative unbiased estimators for such centrality measure on temporal graphs¹; (2) We derive new bounds on the sufficient number of samples to approximate the temporal betweenness centrality for all nodes² (Sect. 4.2), that are governed by three key quantities of the temporal graph, such as the temporal vertex diameter, average temporal path length, and the maximum variance of the temporal betweenness centrality estimators. Moreover, this result solves an open problem

¹ Due to space constraints, we refer to the extended version of this work [14] for the temporal betweenness on temporal edges and for the definition of the other estimators.

 $^{^2}$ The sample complexity analysis holds also for the temporal edges.

in [27,29] on whether the sample complexity bounds for the static betweenness can be *efficiently* extended to temporal graphs. As a consequence, it significantly improves on the state-of-the-art results for the temporal betweenness estimation process [35]. Additionally, our analysis of sample complexity presents further challenges regarding the efficient computation of the three quantities upon which the bounds for the necessary sample size depend; (3) We propose a novel algorithm to efficiently estimate the key quantities of interests in (2) that uses a mixed approach based on sampling and counting (Sect. 4.3). The time complexity of our approach is $\tilde{\mathcal{O}}(\frac{\log n}{c^2}|\mathcal{E}|)$, while the space complexity is $\mathcal{O}(n+|\mathcal{E}|)$. We provide an estimate on the sample size needed to achieve good estimates up to a small error bound. More precisely, we prove that $r = \Theta(\frac{\log n}{r^2})$ sample nodes are sufficient to estimate, with probability at least $1 - 1/n^2$: (i) the temporal diameter $D^{(\star)}$ with error bounded by $\frac{\varepsilon}{\zeta^{(\star)}}$; (ii) the average temporal path length $\rho^{(\star)}$ with error bounded by $\varepsilon \frac{D^{(\star)}}{\zeta^{(\star)}}$; and, (iii) the temporal connectivity rate $\zeta^{(\star)}$ (see Sect. 4.3 for the formal definition) with error bounded by ε ; (4) We define MANTRA, a progressive sampling algorithm that uses an advanced tool from statistical learning theory, namely Monte Carlo Empirical Rademacher Averages [3] and the above results (e.g. (1-3)) to provide a high quality approximation of the temporal betweenness (Sect. 4.4). MANTRA's output is a function of two parameters: $\varepsilon \in (0,1)$ controlling the approximation's accuracy, and $\delta \in (0,1)$ controlling the confidence of the computed approximation. Our novel approach improves on ONBRA [35] (i.e., the state-of-the-art algorithm) in terms of running time, sample size, and allocated space; and, (5) We support our theoretical analysis with an extensive experimental evaluation (Sect. 5), in which we compare MANTRA with ONBRA.

2 Related Work

Tsalouchidou et al. [39], extended the well-known Brandes algorithm [7] to allow for distributed computation of betweenness in temporal graphs. Specifically, they studied shortest-fastest paths, considering the bi-objective of shortest length and shortest duration. Buß et al. [10, 34] analysed the temporal betweenness centrality considering several temporal path optimality criteria, such as shortest (foremost), foremost, fastest, and prefix-foremost, along with their computational complexities. They showed that, when considering paths with increasing time labels, the foremost and fastest temporal betweenness variants are #P-hard, while the shortest and shortest foremost ones can be computed in $O(n^3 \cdot |T|^2)$, and the prefix-foremost one in $O(n \cdot |\mathcal{E}| \cdot \log |\mathcal{E}|)$. Here \mathcal{E} is the set of temporal edges, and T is the set of unique time stamps. Santoro et al. [35] provided ONBRA, the first sampling-based approximation algorithm for one variant of the temporal betweenness centrality. The input to ONBRA is a temporal graph, a confidence value $\delta \in (0,1)$, and the sample size r. The algorithm performs a set of r truncated temporal breadth first searches between couples of nodes sampled uniformly at random and estimates the shortest temporal betweenness using the temporal equivalent of the ABRA estimator [32] for static networks. ONBRA's output is a function of the confidence $\delta \in (0, 1)$ and the upper bound on the approximation accuracy $\xi \in (0, 1)$ computed using the *Empirical Bern*stein Bound [22]. More precisely, with probability $1 - \delta$, the approximation computed by ONBRA is guaranteed to have absolute error of at most ξ for each node in the temporal graph. Finally, Becker et al. [4], provided an efficient heuristic to approximate the temporal betweenness rankings by considering the temporal interactions among the 1-hop neighborhood of the nodes.

3 Preliminaries

Temporal Graphs, and Paths. A directed *temporal graph* is an ordered tuple $\mathcal{G} = (V, \mathcal{E})$ where $\mathcal{E} = \{(u, v, t) : u, v \in V \land t \in T \subseteq \mathbb{N}\}$ is the set of temporal edges³. Given two nodes s and z, a temporal path $tp_{sz} \subseteq V \times V \times T$ is a (unique) sequence of time-respecting temporal edges $((u_1, u_2, t_1), \dots, (u_{k-1}, u_k, t_{k-1}))$ such that for each $1 \leq i < k$, $t_i < t_{i+1}$, every node u_i is visited at most once and $u_1 = s$ and $u_k = z$. Moreover, given a pair of distinct vertices $s \neq z$ a temporal path tp_{sz} from s to z can also be described as a time-ordered sequence of vertex appearances $tp_{sz} = ((u_1, t_1), (u_2, t_2), \dots, (u_k, t_k))$ such that $u_1 = s$, and $u_k = z$. The vertex appearances (u_1, t_1) and (u_k, t_k) are called *endpoints* of tp_{sz} and the temporal nodes in $\mathbf{Int}(tp_{sz}) = tp_{sz} \setminus \{(u_1, t_1), (u_k, t_k)\}$ are called internal vertex appearances of tp_{sz} . Unlike paths on static graphs, in the temporal setting there are several concepts of optimal paths (e.g., *shortest*, *foremost*, fastest [9, 10, 34]. Moreover, as for the static betweenness, the task of computing the desired centrality measure boils down to the ability of efficiently *counting* the overall number of optimal paths. Unfortunately, it has been already shown that such task turns out to be #P-Hard for some temporal path optimalities (e.g. foremost, fastest) [10, 34]. Hope is left for the shortest (and all its variants) an the prefix foremost temporal paths. We formally describe those that can be efficiently counted.

Definition 1. Given a temporal graph \mathcal{G} , and two nodes $s, z \in V$. Let tp_{sz} be a temporal path from s to z, then tp_{sz} is said to be: (1) Shortest (sh) if there is no tp'_{sz} such that $|tp'_{sz}| < |tp_{sz}|$; (2) Shortest-Foremost (sfm) if there is no tp'_{sz} that has an earlier arrival time in z than tp_{sz} and has minimum length in terms of number of hops from s to z; and, (3) Prefix-Foremost (pfm) if tp_{sz} is foremost and every prefix tp_{sv} of tp_{sz} is foremost as well.

Figure 1 shows an example of the temporal paths optimalities considered in this paper. To denote the different type of temporal paths we use the same notation of Buß et al. [10]. More precisely, we use the term "(*)-optimal" temporal path, where (*) denotes the type. Furthermore, we denote the set of all (*)-temporal paths between two nodes s and z as $\Gamma_{sz}^{(*)}$ and we let $\mathbb{TP}_{G}^{(*)}$ to be the

³ The value T denotes the *life-time* of the temporal graph, and, without loss of generality for our purposes, we assume that, for any $t \in T$, there exists at least one temporal arc at that time and without loss of generality we assume T = [1, |T|].


Fig. 1. Example of the (*)-temporal paths described in Definition 1. **Shortest**: $(s \xrightarrow{56} x \xrightarrow{80} y \xrightarrow{92} z), (s \xrightarrow{22} a \xrightarrow{36} b \xrightarrow{40} z),$ **Shortest-Foremost**: $(s \xrightarrow{22} a \xrightarrow{36} b \xrightarrow{40} z),$ and **Prefix-Foremost**: $(s \xrightarrow{1} u \xrightarrow{2} v \xrightarrow{3} w \xrightarrow{4} z).$

union of all the $\Gamma_{sz}^{(\star)}$'s, for all pairs $(s, z) \in V \times V$ of distinct nodes. In this work, we will heavily rely on two temporal graphs characteristic quantities, namely the temporal (vertex) diameter and the average temporal path length. Formally, given a temporal graph $\mathcal{G} = (V, \mathcal{E})$ we define the (\star) -temporal diameter $D^{(\star)}$ and the (\star) -temporal vertex diameter $VD^{(\star)}$ as the number of temporal edges and nodes in the longest (\star) -optimal path in \mathcal{G} , i.e., $D^{(\star)} = \max\left\{|tp^{(\star)}|: tp^{(\star)} \in \mathbb{TP}_{\mathcal{G}}^{(\star)}\right\}$, and $VD^{(\star)} = D^{(\star)} + 1$, respectively. Finally, we refer to the average (\star) -temporal path, i.e., $\rho^{(\star)} = \frac{1}{n(n-1)} \sum_{s,z \in V} |\mathbf{Int}(tp_{sz})|$.

Temporal Betweenness Centrality. As previously shown, on temporal graphs, there are several notions of optimal paths. Hence, we have different notions of temporal betweenness centrality [10] as well. Formally, let $\mathcal{G} = (V, \mathcal{E})$ be a temporal graph. For any pair (s, z) of distinct nodes $(s \neq z)$, let $\sigma_{sz}^{(\star)}$ be the number of (\star) -temporal paths between s and z, and let $\sigma_{sz}^{(\star)}(v)$ be the number of the (\star) -temporal paths between s and z that pass through node v, with $s \neq v \neq z$. The normalized temporal betweenness centrality $\mathbf{b}_v^{(\star)}$ of a node $v \in V$ is defined as

$$\mathbf{b}_v^{(\star)} = \frac{1}{n(n-1)} \sum_{s \neq v \neq z} \frac{\sigma_{sz}^{(\star)}(v)}{\sigma_{sz}^{(\star)}}$$

We refer to the extended version of this work [14] for the definition of the (\star) -temporal betweenness of the temporal edges. Whenever we use the term (\star) -temporal paths we consider (\star) to be one of the optimality criteria in Definition 1. We observe that the average (\star) -temporal path length is equal to the sum of the (\star) -temporal betweenness centrality over all nodes $v \in V$.

Lemma 1.
$$\rho^{(\star)} = \sum_{v \in V} b_v^{(\star)}$$

Supremum Deviation and Empirical Rademacher Averages. Here we define the *Supremum Deviation* (SD) and the *c-samples Monte Carlo Empirical*

Rademacher Average (c-MCERA). For more details about the topic we refer to the book [36] and to [3]. Let \mathcal{D} be a finite domain and consider a probability distribution π over the elements of \mathcal{D} . Let \mathcal{F} be a family of functions from \mathcal{D} to [0,1], and $\mathcal{S} = \{s_1, \ldots, s_r\}$ be a collection of r independent and identically distributed samples from \mathcal{D} sampled according to π . The SD is defined as

$$SD(\mathcal{F}, \mathcal{S}) = \sup_{f \in \mathcal{F}} \left| \frac{1}{r} \sum_{i=1}^{r} f(s_i) - \mathbf{E}_{\pi}[f] \right|$$

The SD is the key concept of the study of empirical processes [30]. One way to derive probabilistic upper bounds to the SD is to use the *Empirical Rademacher* Averages (ERA) [17]. Let $\lambda \in \{-1, 1\}^r$ be a vector or i.i.d. Rademacher random variables, the ERA of \mathcal{F} on \mathcal{S} is

$$R(\mathcal{F}, \mathcal{S}) = \mathbf{E}_{\lambda} \left[\sup_{f \in \mathcal{F}} \frac{1}{r} \sum_{i=1}^{r} \lambda_i f(s_i) \right]$$

Computing the ERA $R(\mathcal{F}, \mathcal{S})$ is usually intractable, since there are 2^r possible assignments for λ and for each such assignment a supremum over \mathcal{F} must be computed. In this work, we use the state-of-the-art approach to obtain sharp probabilistic bounds on the ERA that uses Monte-Carlo estimation [3]. Consider a sample $\mathcal{S} = \{s_1, \ldots, s_r\}$, for $c \geq 1$ let $\lambda \in \{-1, 1\}^{c \times r}$ be a $c \times r$ matrix of i.i.d. Rademacher random variables. The c-MCERA of \mathcal{F} on \mathcal{S} using λ is

$$R_r^c(\mathcal{F}, \mathcal{S}, \boldsymbol{\lambda}) = \frac{1}{c} \sum_{j=1}^c \sup_{f \in \mathcal{F}} \frac{1}{r} \sum_{i=1}^r \lambda_{j,i} f(s_i)$$

The c-MCERA allows to obtain sharp data-dependent probabilistic upper bounds on the SD, as they directly estimate the expected SD of sets of functions by taking into account their correlation. Moreover, they are often significantly more accurate than other methods [27–29], such as the ones based on loose deterministic upper bounds to ERA [32], distribution-free notions of complexity such as the Hoeffding's bound or the VC-Dimension, or other results on the variance [22, 35]. Moreover, a key quantity governing the accuracy of the c-MCERA is the *empirical wimpy variance* [6] $W_{\mathcal{F}}(\mathcal{S})$, that for a sample of size r is defined as

$$\mathcal{W}_{\mathcal{F}}(\mathcal{S}) = \sup_{f \in \mathcal{F}} \frac{1}{r} \sum_{i=1}^{r} (f(s_i))^2$$

Theorem 1 (Theorem 4.1 in [29]). For $c, r \geq 1$, let $\lambda \in \{-1, +1\}^{c \times r}$ be a $c \times r$ matrix of Rademacher random variables, such that $\lambda_{j,i} \in \{-1, +1\}$ independently and with equal probability. Then, with probability at least $1 - \delta$ over λ , it holds $R(\mathcal{F}, \mathcal{S}) \leq R_r^c(\mathcal{F}, \mathcal{S}, \lambda) + \sqrt{\frac{4W_{\mathcal{F}}(\mathcal{S}) \ln(1/\delta)}{cr}}$.

We are ready to state the technical result of this section (proof deferred to the extended version of this paper [14]).

Theorem 2. Let \mathcal{F} be a family of functions with codomain in [0,1], and let \mathcal{S} be a sample of r random samples from a distribution π . Denote \hat{v} such that $\sup_{f \in \mathcal{F}} \operatorname{Var}(f) \leq \hat{v}$. For any $\delta \in (0,1)$, define

$$\tilde{R} = R_r^c(\mathcal{F}, \mathcal{S}, \boldsymbol{\lambda}) + \sqrt{\frac{4\mathcal{W}_{\mathcal{F}}(\mathcal{S})\ln(4/\delta)}{cr}}$$

$$R = \tilde{R} + \frac{\ln(4/\delta)}{r} + \sqrt{\left(\frac{\ln(4/\delta)}{r}\right)^2 + \frac{2\ln(4/\delta)\tilde{R}}{r}}$$

$$\xi = 2R + \sqrt{\frac{2\ln(4/\delta)(\hat{v} + 4R)}{r}} + \frac{\ln(4/\delta)}{3r}$$
(1)
(2)

With probability at least $1 - \delta$ over the choice of S and λ , it holds $SD(\mathcal{F}, S) \leq \xi$.

4 MANTRA: Temporal Betweenness Centrality Approximation Through Sampling

4.1 Temporal Betweenness Estimator

In this section we present one unbiased estimator⁴ for the (*)-temporal betweenness centrality, and we refer to the full version of this work [14] for the remaining estimators that have been excluded due to space constraints. The ONBRA (ob) algorithm [35] uses an estimator defined over the sampling space $\mathcal{D}_{ob} = \{(s, z) \in V \times V : s \neq z\}$ with uniform sampling distribution π_{ob} over \mathcal{D}_{ob} , and family of functions \mathcal{F}_{ob} that contains one function $\tilde{b}_{ob}^{(\star)}(v) \to [0, 1]$ for each vertex v, defined as $\tilde{b}_{ob}^{(\star)}(v|s, z) = \sigma_{sz}^{(\star)}(v)/\sigma_{sz}^{(\star)} \in [0, 1]$. So far, this approach has been defined only for the shortest-temporal betweenness. In this work, we extend ob to shortest-foremost and prefix foremost temporal paths.

4.2 Sample Complexity Bounds

We present two bounds (Theorem 3 and Theorem 4) to the sufficient number of random samples to obtain an ε approximation of the (*)-temporal betweenness centrality. Given a temporal graph $\mathcal{G} = (V, \mathcal{E})$, with a straightforward application of Hoeffding's inequality and union bound [24], it can be shown that $r = 1/(2\varepsilon^2) \log (2n/\delta)$ samples suffice to estimate the (*)-temporal betweenness of every node up to an additive error ε with probability $1 - \delta$. To improve this bound, we define the range space associated to the (*)-temporal betweenness and its VC-dimension, and remand to [23,24,36] for a more complete introduction to the topic. Let $\mathcal{U} = \mathbb{TP}_{\mathcal{G}}^{(*)}$, define the range space $\mathcal{R} = (\mathcal{D}, \mathcal{F}^+)$ where $\mathcal{D} = \mathcal{U} \times [0,1]$, and \mathcal{F}^+ is defined as follows: for a pair $(s,z) \in V \times V$ and a temporal path $tp_{sz} \in \mathcal{U}$ let $f_{(v,t)}(tp_{sz}) = f((v,t)|s,z) = \mathbbm{1}[(v,t) \in \mathrm{Int}(tp_{sz})]$

⁴ An estimator of a given parameter is said to be unbiased if its expected value is equal to the true value of the parameter.

be the function that assumes value 1 if the vertex appearance (v, t) is in the temporal path between s and z. Moreover, define the family of functions $\mathcal{F} = \{f_{(v,t)} : (v,t) \in V \times T\}$ and notice that for each $f_{(v,t)} \in \mathcal{F}$ there is a range $R_{f_{(v,t)}} = \{(tp_{sz}, \alpha) : tp_{sz} \in \mathcal{U} \land \alpha \leq f_{(v,t)}(tp_{sz})\}$. Next, define $\mathcal{F}^+ = \{R_{f_{(v,t)}} : f_{(v,t)} \in \mathcal{F}\}$. Now that we defined the range set for our problem, we can give an upper bound on its VC-dimension.

Lemma 2. The VC-dimension of the range space \mathcal{R} is $VC(\mathcal{R}) \leq \lfloor \log VD^{(\star)} - 2 \rfloor + 1$.

Given the VC-dimension of the range set \mathcal{R} we have:

Theorem 3 (See [21], Section 1). Given $\varepsilon, \delta \in (0, 1)$, and a universal constant h, let $S \subseteq D$ be a collection of elements sampled w.r.t. a probability distribution π . Then $\frac{h}{\varepsilon^2} \left[VC(\mathcal{R}) + \ln \left(\frac{1}{\delta} \right) \right]$ samples suffice to obtain $SD(\mathcal{F}^+, \mathcal{S}) \leq \varepsilon$ with probability $1 - \delta$ over S.

To improve this bound, we make use of Lemma 1 and notice that (as for the static case [29]) the (*)-temporal betweenness centrality satisfies a form of negative correlation among the nodes. Moreover, the existence of a node v with high (*)-temporal betweenness constraints the sum of the centrality measure over the remaining nodes to be at most $\rho^{(*)} - \mathbf{b}_v^{(*)}$. In other words, this suggests that the number of nodes with high (*)-temporal betweenness cannot be arbitrarily large. Furthermore, as in [29], we assume that the maximum variance of the (*)-temporal betweenness estimators $\tilde{\mathbf{b}}_v^{(*)}$ is bounded by some estimate \hat{v} rather than the worst-case upper bound of 1/4 considered in [5]. This implies that the estimates $\tilde{\mathbf{b}}_v^{(*)}$ are not bounded by the number of nodes in the temporal graph \mathcal{G} , but are tightly constrained by the parameters $\rho^{(*)}$ and \hat{v} . We are able to extend the results in [29] for the static scenario to the temporal setting for all the variants of temporal betweenness that can be computed in polynomial time and cover one of the problems left open by the authors. It follows that:

Theorem 4. Let $\mathcal{F} = \{\widetilde{\boldsymbol{b}}_{v}^{(\star)}, v \in V\}$ be a set of function from a domain \mathcal{D} to [0,1]. Define $\hat{v} \in (0,1/4]$ and $\rho^{(\star)} \geq 0$ such that $\max_{v \in V} \operatorname{Var}(\widetilde{\boldsymbol{b}}_{v}^{(\star)}) \leq \hat{v}$ and $\sum_{v \in V} \boldsymbol{b}_{v}^{(\star)} \leq \rho^{(\star)}$. Fix $\varepsilon, \delta \in (0,1)$, and let \mathcal{S} be an i.i.d. sample taken from \mathcal{D} of size $|\mathcal{S}| \in \mathcal{O}\left(\frac{\hat{v}+\varepsilon}{\varepsilon^{2}}\ln\left(\frac{\rho^{(\star)}}{\delta\hat{v}}\right)\right)$. It holds that $SD(\mathcal{F},\mathcal{S}) \leq \varepsilon$ with probability $1 - \delta$ over \mathcal{S} .

Since $\rho^{(\star)}$ correspond to the average number of internal nodes in (\star) -temporal paths in \mathcal{G} , it must be that $\rho^{(\star)} \leq D^{(\star)}$. In *all* the analyzed networks (see Fig. 2 in Sect. 5) this condition holds, thus this approach will need a smaller sample size compared to the VC-Dimension based one to obtain an ε -approximation of the (\star) -temporal betweenness.

4.3 Fast Approximation of the Characteristic Quantities

According to Theorem 3 and Theorem 4, the sample size needed to achieve a desired approximation depends on the vertex diameter and on the average temporal path length of the temporal graph. However, under the Strong Exponential Time Hypothesis (SETH), the (\star) -temporal diameter (thus the average (*)-temporal path length) of a temporal graph $\mathcal{G} = (V, \mathcal{E})$ can not be computed in $\tilde{\mathcal{O}}(|\mathcal{E}|^{2-\varepsilon})^5$ [11], which can be prohibitive for very large temporal graphs, so efficient approximation algorithms for these characteristic quantities are highly desirable. Some algorithms for the diameter approximation on temporal graphs have been proposed [11, 13]. However these techniques consider different temporal path optimality criteria [13], or have no theoretical guarantees [11]. In this work we define a novel sampling-based approximation algorithm to efficiently obtain a high-quality approximation of $D^{(\star)}$ (thus, $VD^{(\star)}$) and $\rho^{(\star)}$ in $\tilde{O}(r \cdot |\mathcal{E}|)$ where r is the number of samples used by the algorithm. We provide a high level description of the sampling algorithm and we refer to the extended version of this work [14] for a detailed discussion and analysis of the method. Given a temporal graph \mathcal{G} , the sample size r, and the temporal path optimality (*), the algorithm performs r (*)-temporal BFS visits [40] ((*)-TBFS) from r random nodes and keeps track of the number of reachable pairs encountered at each hop along with the greatest hop performed. Once all the r visits have been completed, it computes the temporal diameter and other useful temporal measures using an approach based on the relation between the number of reachable pairs and the distance metrics [2]. The approximation guarantees of our sampling algorithm strongly depends on "how temporally connected" a temporal graph is. To this end, we define the (\star) -temporal connectivity rate as the ratio of the number of couples that are temporally connected by a (\star) -temporal path and all the possible reachable couples. Formally, let $\mathbb{1}[u \rightsquigarrow v]$ be the indicator function that assumes value 1 if u can reach v via a (\star)-temporal path, then the temporal connectivity rate is defined as the ratio between the number of reachable pairs and all the possible ones in the temporal graph, i.e., $\zeta^{(\star)} = \frac{1}{n(n-1)} \sum_{u \neq v} \mathbb{1}[u \rightsquigarrow v] \in [0, 1].$ Intuitively, the higher the connectivity rate the higher the number of couples that are connected via at least one (\star) -temporal path. Moreover, the algorithm has the following theoretical guarantees:

Theorem 5. Given a temporal graph $\mathcal{G} = (V, \mathcal{E})$ and a sample of size $r = \Theta\left(\frac{\ln n}{\varepsilon^2}\right)$, the algorithm computes with probability $1 - \frac{2}{n^2}$: the (\star) -temporal diameter $D^{(\star)}$ with absolute error bounded by $\frac{\varepsilon}{\zeta^{(\star)}}$, the average (\star) -temporal path length $\rho^{(\star)}$ with absolute error bounded by $\frac{\varepsilon \cdot D^{(\star)}}{\zeta^{(\star)}}$, and the temporal connectivity rate with absolute error bounded by ε .

⁵ With the notation $\tilde{\mathcal{O}}(\cdot)$ we ignore logarithmic factors.

4.4 The MANTRA Framework

In this section we introduce MANTRA⁶, our algorithmic framework for the (*)temporal betweenness centrality estimation. MANTRA incorporates the bounds in Sect. 4.2 to compute an upper bound on the minimum sample size needed to approximate the SD of the (*)-temporal betweenness and a state-of-the-art progressive sampling technique to speed-up the estimation process. The input parameters to MANTRA are: a temporal graph \mathcal{G} , a temporal path optimality (*) $\in \{\mathtt{sh}, \mathtt{sfm}, \mathtt{pfm}\}^7$, a target precision $\varepsilon \in (0, 1)$, a failure probability $\delta \in (0, 1)$, and a number of iterations for the bootstrap phase s'. The output is a vector \mathcal{B} of pairs $(v, \widetilde{\mathbf{b}}_v^{(*)})$ for each $v \in V$, where $\widetilde{\mathbf{b}}_v^{(*)}$ is the estimate of $\mathbf{b}_v^{(*)}$ and \mathcal{B} is probabilistically guaranteed to be an absolute ε approximation of the temporal betweenness. Formally:

Theorem 6. Given a target accuracy $\varepsilon \in (0,1)$ and a failure probability $\delta \in (0,1)$, with probability at least $1-\delta$ (over the runs of the algorithm), the output vector $\mathcal{B} = \{\widetilde{\mathbf{b}}_v^{(\star)} : v \in V\}$ (obtained from a set of samples \mathcal{S}) produced by MANTRA is such that $SD(\mathcal{B}, \mathcal{S}) \leq \varepsilon$.

Algorithm 1: MANTRA

Data: Temporal graph \mathcal{G} , (*) temporal path optimality, precision $\varepsilon \in (0, 1)$, failure probability $\delta \in (0, 1)$, bootstrap iterations s', and number of Monte Carlo trials c. **Result**: Absolute ε -approximation of the (\star)-tbc w.p. of at least $1 - \delta$. 1 $\mathcal{B}, \mathcal{W} = [0, \dots, 0] \in \mathbb{R}^n$ // tbc and wimpy variance arrays **2** $i, k = 0; \xi = 1; S_0 = \{\emptyset\}$ **3** $\omega, \hat{v} = \text{DrawSufficientSampleSize}(\mathcal{G}, s', \delta/2)$ 4 $\{s_i\}_{i\geq 1} = \text{SamplingSchedule}(\omega, \hat{v}, \delta)$ 5 $\lambda = [[\cdot]]$ // Empty matrix 6 while true do $i = i + 1; k = (1.2 \cdot s_{i-1}) - s_{i-1}$ 7 $\mathcal{X} = \text{DrawSamples}(\mathcal{G}, k) / /$ Draw k samples from the sample space \mathcal{D}_{ob} 8 $\mathcal{S}_i = \mathcal{S}_{i-1} \cup \mathcal{X}$ 9 $oldsymbol{\lambda} = \operatorname{Add} \operatorname{R.R.Vector}(k, oldsymbol{\lambda})$ // Add a Rade. rnd. column of length c10 $\mathcal{B}, \mathcal{W}, \boldsymbol{\lambda} = \texttt{Update}(\star) - \texttt{TemporalBetweenness}(\mathcal{X}, \mathcal{B}, \mathcal{W}, \boldsymbol{\lambda})$ 11 $\tilde{R}, v_{\mathcal{F}} = \texttt{UpdateEstimates}(\mathcal{B}, \mathcal{W}, \boldsymbol{\lambda}, |\mathcal{S}_i|, k, c)$ 12 $R_k^c = \frac{1}{c} \sum_{l=1}^c \max_{v \in V} \left\{ \tilde{R}[v, l] \right\}$ 13 $\xi = \texttt{ComputeSDBound}(R_k^c, v_{\mathcal{F}}, \delta/2^i, |\mathcal{S}_i|)$ // Compute Eq. 2 in Thm. 2 $\mathbf{14}$ if $|S_i| \ge \omega$ or $\xi \le \varepsilon$ then return $\{(1/|S_i|) \cdot \mathcal{B}[u] : u \in V\}$ 15

⁶ te<u>MporAl betweeNness cenTrality appRoximation through sAmpling.</u>

⁷ We point out that our approach is general, and can be extended to *every* definition of temporal betweenness centrality.

Algorithm 1's execution is divided in two phases: the bootstrap phase (lines 3-4) and the estimation phase (lines 6-15). As a first step, MANTRA, computes an upper bound ω to the number of samples needed to achieve an ε approximation (line 2). The procedure runs s' independent (*)-TBFS visits from s'^8 random couples of nodes (s, z) sampled from the population \mathcal{D}_{ob} , estimates \hat{v} and $\rho^{(\star)}$, and then plugs them in Theorem 4 to obtain ω . Subsequently, it infers the first element of the sample size $\{s_i\}_{i>1}$ by performing a binary search between s' and ω to find the minimum s_1 such that Eq. 2 (with R set to 0) is at most ε and terminates the bootstrap phase. Such approach gives an optimistic first guess of the number of samples to process for obtaining an ε -approximation [29]. Subsequently it continues with the estimation phase in which, at each iteration, it increases each s_i with a geometric progression [31], i.e., such that $s_i = 1.2 \cdot s_{i-1}$. Next, it proceeds by drawing uniformly at random $k = s_i - s_{i-1}$ couples of nodes (s, z) from \mathcal{D}_{ob} and subsequently updating the overall set of samples sampled so far (lines 7-9). Consequently, k new Rademacher random vectors are added as new columns to the matrix λ and k (*)-TBFS visits are performed (line 11). Moreover, while iterating over the new sample \mathcal{X} the temporal betweenness, wimpy variances and the values in λ are updated. After this step, the coefficients of Equation 2 and the new estimate on the SD, ξ , are computed (lines 12-13). As a last step of the while loop, the algorithm checks whether the desired accuracy ε has been achieved, i.e., whether the actual number of drawn samples is at least ω or ξ is at most ε (line 15). If at least one of the two conditions is met, MANTRA normalizes and outputs the current estimates \mathcal{B} . We conclude this section with the analysis of MANTRA's running time.

Theorem 7. Given a temporal graph $\mathcal{G} = (V, \mathcal{E})$ and a sample of size r, MANTRA requires time $\tilde{\mathcal{O}}(r \cdot n \cdot |T|)$ and $\tilde{\mathcal{O}}(r \cdot |\mathcal{E}|)$ to compute the shortest (foremost)-temporal and the prefix-foremost-temporal betweenness, respectively. Moreover, MANTRA requires $\mathcal{O}(n + |\mathcal{E}|)$ space.

Theorem 6 together with Theorem 7 provide theoretical evidence that MANTRA computes a rigorous estimation of the (\star)-temporal betweenness and that *scales* to the size of the input temporal graph. Moreover, it improves over the state-of-the-art approach ONBRA [35]. Indeed, given a sample of size r, ONBRA stores a $n \times r$ matrix to compute the absolute ξ -approximation⁹ using the Empirical Bernstein Bound [22]. Thus, ONBRA may require an arbitrary large sample size (e.g. large matrix) to achieve a target absolute approximation ε , making the algorithm not ideal to analyze big temporal graphs.

5 Experimental Evaluation

In this section, we summarize the results of our experimental study on approximating the (\star) -temporal betweenness centrality in real-world temporal networks.

⁸ In this work we use $s' = \log(1/\delta)/\varepsilon$.

⁹ ξ is the upper bound on the $SD(\mathcal{S}, \mathcal{F})$ obtained using the Empirical Bernstein Bound.

5.1 Experimental Setting

We compare our novel framework with ONBRA [35]. For the sake of fairness, we adapted the original fixed sample size algorithm to use the same progressive sampling approach of our framework. Every time an element of the sampling schedule is consumed, the algorithm computes the upper bound ξ on the SD using the Empirical-Bernstein bound as in [35], if ξ is at most the given ε , it terminates, otherwise it keeps sampling. We set ONBRA's maximum number of samples to be equal to the VC-Dimension upper-bound in Sect. 4.2. We implemented all the algorithms in Julia exploiting parallel computing¹⁰. We chose to re-implement the exact algorithms [10] and ONBRA [35] because they have issues with the number of paths in the tested networks¹¹, causing overflow errors (indicated by negative centralities), and with the time labeling causing an underestimation of centralities [4]. Our implementation uses a sparse matrix representation of the $n \times |T|$ table used in [10,35], making the implemented algorithms spaceefficient and usable on big temporal graphs (for which the original version of the code gives out of memory errors). We executed all the experiments on a server running Ubuntu 16.04.5 LTS with one processor Intel Xeon Gold 6248R 32 cores CPU @ 3.0GHz and 1TB RAM. For every temporal graph, we ran all the algorithms with parameter $\varepsilon \in \{0.1, 0.07, 0.05, 0.01, 0.007, 0.005, 0.001\}$ chosen to have a comparable magnitude to the highest temporal betweenness values in the network (see $b_{\max}^{(\star)}$ in Table 1). This is a basic requirement when computing meaningful approximations¹². Moreover, we use $\delta = 0.1$ and use c = 25 Monte Carlo trials as suggested in [12,29]. Finally, each experiment has been ran 10 times and the results have been averaged.

5.2 Networks

We evaluate all the algorithms on real-world temporal graphs of different nature, whose properties are summarized in Table 1. The temporal networks come from three different domains:

- Social networks. This domain includes most of the considered networks: College msg, Digg reply, Slashdot reply, Facebook Wall, Mathoverflow, SMS, Askubuntu, and Wiki Talk. These are social networks from different realms, where nodes correspond to users and temporal arcs indicate messages sent between them at specific points in time.
- **Contact networks.** For the **Topology** network, nodes correspond to computers and temporal arcs indicate a contact between nodes at a specific time.
- **Transport networks.** Bordeaux is part of the Kuala et al. [18] public transport networks collection. In such temporal graph, nodes are public transport stops and a temporal arcs indicate routes at a specific point in time. Because of their

¹⁰ Code available at: https://github.com/Antonio-Cruciani/MANTRA.

¹¹ The overflow issue appears on *all* the transportation networks provided in [18].

 $^{^{12}}$ It is meaningless to compute an $\varepsilon\text{-approximation}$ when the maximum centrality value is smaller than $\varepsilon.$

Table 1. The data sets used in our evaluation, where ζ indicates the exact temporal connectivity rate, $b_{\max}^{(\star)}$ the maximum (\star)-temporal betweenness centrality (type D stands for directed and U for undirected). • indicates that we need to use **BigInt** data type instead of **Unsigned Int128** to count the number of shortest (foremost)-temporal paths to avoid overflows.

Data set	\boldsymbol{n}	$ \mathcal{E} $	T	ζ	$m{b}_{ t max}^{(t pfm)}$	$m{b}_{ t max}^{(t sh)}$	$m{b}_{ t max}^{(t sfm)}$	Type	Source
College msg	1899	59798	58911	0.5	0.0718	0.0319	0.0365	D	[20]
Digg reply	30360	86203	82641	0.02	0.0019	0.0015	0.0016	D	[33]
Slashdot	51083	139789	89862	0.07	0.0128	0.0074	0.0085	D	[33]
Facebook Wall	35817	198028	194904	0.04	0.0034	0.0024	0.0028	D	[<mark>33</mark>]
Topology	16564	198038	32823	0.53	0.0921	0.0654	0.0681	U	[19]
Bordeaux●	3435	236075	60582	0.84	0.1210	0.1383	0.1269	D	[18]
Mathoverflow	24759	390414	389952	0.33	0.0522	0.0282	0.0287	D	[20]
SMS	44090	544607	467838	0.008	0.0019	0.0010	0.0012	D	[20]
Askubuntu	157222	726639	724715	0.169	0.0214	0.0156	0.0154	D	[20]
Super user	192409	1108716	1105102	0.21	0.0261	0.0165	0.0182	D	[20]
Wiki Talk	1094018	6092445	5799206	0.069	0.0089	0.0155	0.0153	D	[19]

"inherent temporality", these networks are characterized by a big number of temporally connected nodes.



Fig. 2. Comparison between temporal diameter and the average number of internal nodes for the Shortest (foremost) and Prefix-Foremost temporal path optimalities. The approximation has been computed (over 10 runs) using our sampling algorithm using 256 random seed nodes.

5.3 Experimental Results

Efficiency and Scalability. In our first experiment, we compare the average execution times, sample sizes and allocated memory of MANTRA and ONBRA. Due

to space constraints we show the results on the data sets in Table 1 for the prefixforemost-temporal betweenness, for a subset of $\varepsilon \in \{0.01, 0.007, 0.005, 0.001\}$ and we refer to the additional materials for the complete battery of experiments. We chose to display the results for the pfm temporal path optimality because it is the one for which the analyzed graphs have the highest characteristic quantities (see Fig. 2). Thus, under this setting, the tested algorithms will need a bigger sample size and potentially a higher amount of memory. This somehow provides an intuitive "upper bound" on the algorithms performances in terms of efficiency and scalability. Moreover, the experiments for sh and sfm temporal betweenness follow similar trends of the ones displayed in the main paper. Figure 3a shows the comparison of the running times (in seconds) for the pfm temporal betweenness. We observe that MANTRA leads the scoreboard against its competitor on all the tested networks. Our novel framework is at least three times faster than ONBRA. Such speedup is mainly due to the smaller sample size needed to terminate. Furthermore, Fig. 3b shows that MANTRA requires a smaller sample size (at least three times smaller) to converge. This early convergence, in practice, does not affect the approximation quality and leads to very good temporal betweenness approximations (see the next experiment). Furthermore, the number of samples needed by MANTRA varies among temporal graphs, with a strong dependence on $b_{\max}^{(\star)}$. A potential cause of the difference in the sample sizes between the two algorithms may depend on the use of the Empirical Bernstein bound. Such bound (as the VC-Dimension one) is agnostic to any property of the analyzed temporal network, thus results in a overly conservative guarantees. This suggests that *variance-adaptive* bounds are preferable to compute data-dependent approximations [29], and that exploiting correlations among the nodes through the use of the c-MCERA leads to refined guarantees. Moreover, we point out that ONBRA does not scale well as the target absolute error ε decreases. Indeed, the memory needed by ONBRA increases drastically as the target absolute error decreases (see Fig. 3c) to the point of giving out of memory error for big temporal networks such as Slashdot, SMS, Askubuntu, Superuser, and Wiki Talk. This can lead to major issues while computing meaningful ε approximations, especially under the setting in which the maximum temporal betweenness $b_{\max}^{(\star)}$ is very small (for which we need to choose an ε value of at most $b_{\max}^{(\star) 13}$). Unfortunately, this is not an uncommon feature of real-world temporal networks. Indeed, as shown by ζ and $b_{\max}^{(\star)}$ in Table 1 they tend to be very sparse. This experiment, suggests that MANTRA is preferable for analyzing big temporal networks up to an arbitrary small absolute error ε .

Comparison with the Exact Algorithms Scores and Running Times. As a first step in our second experiment, we investigate the accuracy of the approximations provided by MANTRA by computing the exact temporal betweenness centrality of all the nodes of the temporal network in Table 1 and measuring the SD over all the ten runs. Figure 3d supports our theoretical results, as we always get a

 $^{^{13}}$ We recall that $b_{\max}^{(\star)}$ can be efficiently approximated in the bootstrap phase of our framework.

SD of at most the given ε . Moreover, we point out that the exact algorithms for the shortest (foremost) temporal betweenness required a time that spanned from several hours (e.g. for SMS) to days (for Askubuntu, and Superuser \approx a week) and weeks (for Wiki Talk \approx a month). Instead, MANTRA completes the approximation in reasonable time. Figure 4a shows the relation between the sample size and the running time of our framework. While, Fig. 4b shows the amount of time needed by MANTRA to provide the absolute ε -approximation in terms of percentage of exact algorithm's running time. We display the running times on the biggest temporal graphs for the sh temporal betweenness because is one of the "critical" temporal path optimalities that requires longer times to be computed (see Theorem 7). We can conclude that our framework is well suited to quickly compute *effective* approximations of the temporal betweenness on very large temporal networks.



Fig. 3. Experimental analysis for $\varepsilon \in \{0.01, 0.007, 0.005, 0.001\}$. Comparison between the running times (a), sample sizes (b), and allocated memory (c) of ONBRA and MANTRA. (d) Supremum deviation of the absolute ε -approximation computed by MANTRA. The black line indicates that the two algorithms require the same amount of time/samples/memory, gray line (followed by a red mark) indicates that the algorithm required more than 1TB of memory to run on that data set with that specific ε value.



Fig. 4. (a) Relation between the running time and the sample size of MANTRA for the shortest temporal betweenness with ε as in Fig. 3. (b) Comparison between MANTRA and the exact algorithm running times for the shortest temporal betweenness on the biggest temporal networks.

6 Conclusions

We proposed MANTRA, a novel framework for approximating the temporal betweenness centrality on large temporal networks. MANTRA relies on the state-of-the-art bounds on supremum deviation of functions based on the c-MCERA to provide a probabilistically guaranteed absolute ε -approximation of such centrality measure. Our framework includes a fast sampling algorithm to approximate the temporal diameter, average path length and connectivity rate up to a small error with high probability. Such approach is general and can be adapted to approximate several versions of these quantities based on different temporal path optimalities (e.g. [11,13]). Our experimental results (summarized in Sect. 5) depict the performances of our framework versus the state-of-the-art algorithm for the temporal betweenness approximation. MANTRA consistently over-performs its competitor in terms of running time, sample size, and allocated memory. As indicated in Fig. 3c, our framework is the only available option to obtain meaningful temporal betweenness centrality approximations when we do not have access to servers with a large amount of memory. In addition, MANTRA improves on the recent work by Zhang et al. [41] (WWW, 2024) that relies on loose deterministic upper bounds to the ERA to approximate the temporal betweenness. In the future, we plan to combine the results by Brunelli et al. [8] (KDD, 2024) with MANTRA to approximate the restless temporal betweenness and to speedup the overall estimation process of various temporal betweenness variants that have not been considered in this work. Finally, other promising future directions are to use our novel framework to estimate the temporal ego betweenness [4], to find communities in temporal graphs, and to extend our approach to other temporal path based centrality measures [25, 26].

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Dimensionality-Induced Information Loss of Outliers in Deep Neural Networks

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Abstract. Out-of-distribution (OOD) detection is a critical issue for the stable and reliable operation of systems using a deep neural network (DNN). Although many OOD detection methods have been proposed, it remains unclear how the differences between in-distribution (ID) and OOD samples are generated by each processing step inside DNNs. We experimentally clarify this issue by investigating the layer dependence of feature representations from multiple perspectives. We find that intrinsic low dimensionalization of DNNs is essential for understanding how OOD samples become more distinct from ID samples as features propagate to deeper layers. Based on these observations, we provide a simple picture that consistently explains various properties of OOD samples. Specifically, low-dimensional weights eliminate most information from OOD samples, resulting in misclassifications due to excessive attention to dataset bias. In addition, we demonstrate the utility of dimensionality by proposing a dimensionality-aware OOD detection method based on alignment of features and weights, which consistently achieves high performance for various datasets with lower computational cost. Our implementation is publically available at https://github.com/ kuematsu3/Dimensionality-aware-Projection-based-OOD-Detection.

Keywords: Out-of-distribution detection \cdot Dimensionality

1 Introduction

Deep neural networks (DNNs) have received much attention in recent years due to their remarkable versatility and performance. DNNs are broadly applicable to real tasks including the long-term operation of DNN systems. However,

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they suffer from data shifts caused by changes in the surrounding environment. Shifted data degrade the performance of pretrained models and thus need to be detected as outliers. This task is commonly referred to as out-of-distribution (OOD) detection, and also goes by other names including novelty detection and open-set recognition, although subtle differences exist in the terminology [1,2]. As the application of DNNs continues to expand, OOD detection becomes more critical for ensuring stable and reliable system operation.

Although there have been numerous proposals for precise detection of OOD samples [1,2], the mechanism by which OOD samples deviate from indistribution (ID) samples remains unclear. Several characteristics of OOD samples have been identified, including comparatively smaller logit values and corresponding quantities [3, 4], deviation from the low-dimensional ID subspace [5-8], a low-ranked feature vector for each OOD sample [9], and the data complexitydependent change of layers suitable for OOD detection [10]. However, these properties alone do not sufficiently reveal the relationship between observed behaviors and information processing components inside DNNs, including layerwise affine transformations using weight matrices and the activation functions. This kind of a reductionistic perspective is valuable for clarifying the relevant processing that makes ID and OOD samples distinct in non-ideal tasks. Non-ideal situations are taken to mean the cases where theoretical analysis is not suitable due to the complicated data distributions and network architectures, including that the probability distribution of input data or features is unknown and the model is not well-behaved. Understanding the properties of OOD samples based on fundamental DNN components could lead to appropriate solutions for addressing the uncertainty inherent in DNNs.

It would also be valuable to clarify the behaviors of OOD samples for practical applications of OOD detection, particularly in cases with various constraints. Typically, OOD samples are not available before they appear during the operational phase, and introducing additional models for monitoring the DNN system is impractical because it increases the operating cost. Furthermore, in some complex and large systems, modification of the operating model can be challenging because it may impact downstream tasks beyond the current objective. Additionally, computational time and memory usage can also be significant issues, particularly if real-time processing of a large amount of data is necessary with limited computational resources. A simplified understanding of OOD samples could help identify the minimum relevant part in DNNs with all constraints satisfied.

In this work, we experimentally investigate how the difference between ID and OOD samples arises by focusing on individual feature transformations inside DNNs for non-ideal tasks. To address this issue, we explore the layer dependence of feature representations and their relationship with weight matrices. Our contributions are summarized as follows:

 Through a systematic analysis of various quantities given in Sect. 2, we identify that the sharp change in dimensionality plays an essential role in making ID and OOD samples distinct. We verify that this change of dimensionality consistently explains not only the observed behaviors presented in Sect. 3 but also those reported in previous studies as discussed in Sect. 4.

In addition to these, we demonstrate the utility of dimensionality by evaluating a dimensionality-aware OOD detection method in Sect. 3.7.

2 Problem Setting and Related Work

We consider a straightforward and likely representative OOD detection problem where we disregard OOD samples during the training phase. Let \mathcal{D}_{ID} and \mathcal{D}_{OOD} be the sets of ID and OOD samples. For \mathcal{D}_{ID} , we distinguish the training dataset \mathcal{D}_{train} and the test dataset \mathcal{D}_{test} . We examine a scenario where we exclusively utilize \mathcal{D}_{train} for model training and for hyper parameter tuning. The training strategy is dedicated to generalization to \mathcal{D}_{test} without any additional considerations for \mathcal{D}_{OOD} , which contrasts with some previous studies [11–13]. This situation enables us to simulate the occurrence of OOD samples and to investigate their intrinsic properties inside DNNs.

In the following subsections, we briefly introduce quantities to be evaluated in Sect. 3. Details of the experimental setup are provided in Appendix B.

2.1 Stable Rank of the Matrix

The properties of a matrix, particularly its dimensionality, play an essential role in understanding the behavior of OOD samples. Refs. [6,8] have reported the importance of the null space of the covariance matrix for OOD detection. Furthermore, dimensionality of features is closely related to that of weight matrices. Therefore, we investigate the stable rank R_F of the matrix A:

$$R_F = ||A||_F^2 / ||A||_2^2, \tag{1}$$

where $||A||_F$ and $||A||_2$ are the Frobenius norm and the spectrum norm, respectively. R_F provides a numerical evaluation of the matrix rank robust against small singular values.

To fully investigate the layer dependence of propagation in the DNN, we perform a similar analysis for the weight matrices W of all layers including convolutional layers. W of the fully connected layer at the *l*-th layer is represented by the matrix $W^{(l)} \in \mathbb{R}^{H^{(l+1)} \times H^{(l)}}$, where $H^{(l)}$ is the number of channels at the *l*-th layer. For convolutional layers, in contrast to the fully connected layer, we focus only on the local linear transformation represented by the weight matrix $W^{(l)} \in \mathbb{R}^{H^{(l+1)} \times F_x^{(l)} F_y^{(l)} H^{(l)}}$, where $F_x^{(l)}$ and $F_y^{(l)}$ are the kernel sizes at the *l*th layer. Regarding the covariance matrix $\overline{\Sigma}$, we adopt average pooling of the feature following Ref. [5], as outlined in Sect. 2.2. The latter two simplifications allow us to diagonalize matrices and compute R_F with practical computational cost.

2.2 Feature-Based Detection

Many studies have utilized feature representations inside DNNs for OOD detection to leverage their rich information [1,2]. One straightforward approach involves measuring the Mahalanobis distance from ID training samples [5] or considering various extensions [6,14–18]. The Mahalanobis distance $M^{(l)}$ is defined as follows:

$$M^{(l)} = \min_{c} \sqrt{(x^{(l)} - \mu_c^{(l)})^{\mathsf{T}}(\overline{\Sigma}^{(l)})^{-1}(x^{(l)} - \mu_c^{(l)})},$$
(2)

$$\mu_{c}^{(l)} = \frac{1}{N_{\text{train},c}} \sum_{x_{i,c} \in \mathcal{D}_{\text{train},c}} x_{i,c}^{(l)},$$

$$\overline{\Sigma}^{(l)} = \frac{1}{K} \sum_{c} \frac{1}{N_{\text{train},c}} \sum_{x_{i,c} \in \mathcal{D}_{\text{train},c}} (x_{i,c}^{(l)} - \mu_{c}) (x_{i,c}^{(l)} - \mu_{c})^{\mathsf{T}}.$$
 (3)

Here, $x^{(l)}$ is the feature at the *l*-th layer generated by the input sample $x, x_{i,c}^{(l)}$ is the *l*-th layer feature of the *i*-th training sample $x_{i,c} \in \mathcal{D}_{\text{train},c}$ with corresponding class label $y_i = c$, $\mathcal{D}_{\text{train},c}$ is the training dataset with the class label c out of K classes in the ID dataset, $N_{\text{train},c} = |\mathcal{D}_{\text{train},c}|$ is the number of training data with class label c, and $\overline{\Sigma}^{(l)}$ is the tied covariance of the feature at the *l*-th layer, an approximation of the class-wise covariance [5]. By using $M^{(l)}$, we can classify a given sample x as an ID (resp. OOD) sample if the value of $M^{(l)}$ is small (resp. large) for a certain layer l. In the following, we refer to this detection method as "feature-based detection".

To reduce the computational cost, Ref. [5] employs pixel-averaged features as $x^{(l)}$, $\mu_c^{(l)}$, and $\overline{\Sigma}^{(l)}$. After averaging the pixel values of the feature map, the Mahalanobis distance is computed for the feature vector with $H^{(l)}$ elements, where $H^{(l)}$ represents the number of channels at the *l*-th layer. In this paper, we adopt the pixel-averaged tied covariance for simplicity [5]. Furthermore, to avoid division by zero arising from singular covariance, we compute the Moore–Penrose inverse $(\overline{\Sigma}^{(l)})^{-1} = \sum_{k}^{d} \lambda_{k}^{-1} v_{k} v_{k}^{\mathsf{T}}$, where λ_{k} denotes the *k*-th largest eigenvalue of $\overline{\Sigma}^{(l)}$ and v_{k} is the corresponding eigenvector. *d* is taken to be the largest value satisfying $\lambda_d/\lambda_0 > \varepsilon$ with $\varepsilon = 10^{-6}$.

2.3 Projection-Based Detection

Our main interest is how the deviation of OOD samples from ID samples arises during forward propagation, which consists of a series of linear transformations and activations. One fundamental factor influencing feature transformation inside DNNs is alignment of weights and features as explored in Ref. [19] called NuSA. Let $W^{(l)}$ be the weight matrix at the *l*-th layer and $W^{(l)} = L^{(l)}(Q^{(l)})^{\mathsf{T}}$ be its QR decomposition. By projecting the feature at the *l*-th layer onto a subspace using the transformation $x_p^{(l)} = (Q^{(l)})^{\mathsf{T}} x^{(l)}$, we obtain several OOD detection scores, including $||x_p^{(l)}||/||x^{(l)}||$ and $||x_p^{(l)}||$. Using these scores, we can classify a given sample x as an OOD sample if the value of scores is large. The contrapositive statement of the above is that features of ID samples aligns with weights, which is justified by the noise stability of the trained network [20]. In the following, we refer to this detection method as "projection-based detection".

NuSA in Ref. [19] considered projections to only fully connected layers using QR decomposition. This approach restricts access to the singular value associated with the importance of each projection vector in $Q^{(l)}$, as well as the full propagation properties inside the network including convolutional layers.

To overcome these limitations, we propose a modified projection-based method that aim to identify the relevant subspace for propagation and to investigate the full layer dependence of propagation. Our approach involves two key modifications to NuSA. One is removing irrelevant singular vectors, thereby enhancing our awareness of dimensionality of weight matrices. Another is representing the convolutional layer by the local linear transformation from $F_x^{(l)}F_y^{(l)}H^{(l)}$ - to $H^{(l+1)}$ -dimensional vector space. We sometimes refer to the modified method as "dimensionality-aware" to clarify that this modification is aware of dimensionality.

The precise description of the first modification is as follows:

- 1. Perform singular-value decomposition of the *l*-th layer weight matrix $W^{(l)}$: $W^{(l)} = U^{(l)}S^{(l)}(V^{(l)})^{\mathsf{T}}.$
- 2. Remove irrelevant singular values such that $s_k^{(l)}/s_0^{(l)} < \varepsilon$ where $s_k^{(l)}$ is the k-th largest singular value of $W^{(l)}$, i.e., $s_0 \ge s_1 \ge \cdots$.
- 3. Construct the dimensionality-aware projection matrix $V_{\varepsilon}^{(l)}$ from singular vectors corresponding to remaining singular values.
- 4. Project the *l*-th layer feature $x^{(l)}$ to the weight as $x_{p,\varepsilon}^{(l)} = (V_{\varepsilon}^{(l)})^{\mathsf{T}} x^{(l)}$.
- 5. Regard samples with smaller values of $||x_{p,\varepsilon}^{(l)}||/||x^{(l)}||$ or $||x_{p,\varepsilon}^{(l)}||$ as OOD samples.

 ε is typically taken to be 10^{-2} . The detailed ε dependence is summarized in Appendix A. Note that $||x_{p,\varepsilon}^{(l)}||$ for convolutional layers can be easily obtained by replacing the tensor representation of $W^{(l)}$ ($(H^{(l+1)}, H^{(l)}, F_x^{(l)}, F_y^{(l)})$ tensor) with that of $(V_{\varepsilon}^{(l)})^{\mathsf{T}}$ ((rank $(V_{\varepsilon}^{(l)}), H^{(l)}, F_x^{(l)}, F_y^{(l)})$ tensor), and by acting the replaced convolutional layer on $x^{(l)}$.

2.4 Similarity of DNN Representations

The similarity between DNN features serves as a standard metric for investigating properties of DNNs [21–25]. An extensive study [21] revealed the effectiveness of centered kernel alignment (CKA) defined as follows [26,27]:

$$CKA(K_{\mathcal{D}}^{(l_1)}, K_{\mathcal{D}}^{(l_2)}) = \frac{\operatorname{tr}((K_{\mathcal{D}}^{(l_1)})^{\mathsf{T}} K_{\mathcal{D}}^{(l_2)})}{||K_{\mathcal{D}}^{(l_1)}||_F ||K_{\mathcal{D}}^{(l_2)}||_F}.$$
(4)

Here, l_1 and l_2 represent the layer, $K_{\mathcal{D}}^{(l)} = HK_{0,\mathcal{D}}^{(l)}H$ where $H = I - \frac{1}{N_{\mathcal{D}}}\sum_{i\in\mathcal{D}}\hat{1}\hat{1}^{\mathsf{T}}$ is the centering operator and $\hat{1} = (1, \cdots, 1)^{\mathsf{T}}, (K_{0,\mathcal{D}}^{(l)})_{ij} = (x_{i,\mathcal{D}}^{(l)})^{\mathsf{T}}x_{j,\mathcal{D}}^{(l)}$ where $x_{i,\mathcal{D}}^{(l)}$ is the feature of the *i*-th sample of the dataset \mathcal{D} at the *l*-th layer. See Appendix C for the detailed procedure and a comparison with other similarity scores. By examining inter-layer CKA, we gain insights into the layerwise feature transformation inside the DNN. We use 10,000 randomly selected samples to obtain $K_{0,\mathcal{D}}$.

2.5 Noise Sensitivity in the DNN

The projection to the weight matrix is effective to investigate the property of one-layer linear propagation. However, we need a more sophisticated approach to characterize multi-layer nonlinear propagation. In this context, noise sensitivity ψ [20] proves useful:

$$\psi_{\eta}(x; M^{(l_1, l_2)}) = \frac{||M^{(l_1, l_2)}(x_{\eta}^{(l_1)}) - M^{(l_1, l_2)}(x^{(l_1)})||^2}{||M^{(l_1, l_2)}(x^{(l_1)})||^2}.$$
(5)

Here, $x^{(l)}$ is defined in the same manner as Eq. 2, $x_{\eta}^{(l)} = x^{(l)} + \eta ||x^{(l)}||$ represents the noise-injected feature, where η is typically isotropic noise generated by elementwise Gaussian variables, and $M^{(l_1,l_2)}(\cdot)$ denotes the function of the DNN forward propagation from the layer l_1 to l_2 , that is, $M^{(l_1,l_2)}(x^{(l_1)}) = x^{(l_2)}$. Note that in the linear case with the standard Gaussian noise η , the average of ψ is identical to $||W||_F^2 ||x||^2 / ||Wx||^2$ [20], which is quite similar to the score of projection-based detection in Sect. 2.3. To simplify matters, we neglect the average over η , assuming that it will not significantly impact the typical value of ψ . We employ the median of samples in a dataset as the typical value of ψ . We fix the norm of noise η to 0.1 [20].

3 Results

In this section, we present the experimental results for the various quantities introduced in Sect. 2 to grasp feature transformations of ID and OOD samples. Details of the experimental setup are provided in Appendix B. Our findings are corroborated by aggregating results from multiple perspectives, emphasizing consistency rather than relying solely on individual properties.

3.1 Overview of the Experiments and a Possible Picture

The primary objective of our experiments is to establish the picture in Fig. 1. We achieve this by investigating how OOD samples deviate from ID samples as the features propagate to deeper layers. To accomplish this, we evaluate the layer dependence of various quantities.

The most intriguing observation is the presence of a common layer that exhibits transition-like behaviors across various quantities: a significant decrease in the stable ranks of covariances and weights, stabilization of feature-based detection performance, peaky structure of projection-based detection performance, boundary of block-like CKA, and a notable decrease in noise sensitivity.



Fig. 1. Overview of results showing how OOD samples deviate from ID samples, and how OOD samples are classified. The source of the difference between ID and OOD samples is low dimensionalization of weights, yielding observed behaviors not only of single-layer properties but also of multi-layer properties due to alignment of features and weights (left side of the figure). The resulting features of OOD samples are dominated by dataset bias, the common characteristics in the dataset, leading to the biased prediction (right side of the figure).

Furthermore, although the location of this *transition layer* varies depending on network architectures,¹ similar behaviors are universally observed independent of architectures and ID datasets (Appendix G), as long as the number of ID classes is comparably small. These consistent behaviors strongly suggest the existence of a fundamental origin shared across all scenarios.

An essential common element to all of them is low dimensionalization of weight matrices, as observed by the stable rank. This naturally evokes the following propagation process for ID and OOD samples as illustrated in Fig. 1. For ID samples, low dimensionalization *preserves* most of the characteristic information, including features relevant to classification in the case of classification tasks [20]. However, when it comes to OOD samples, low dimensionalization *eliminates* most of their characteristic information. The resulting features of OOD samples are thus dominated by the *dataset bias*, except the dataset-independent bias parameters in the model. Here, the term "dataset bias" refers to common characteristics shared within a dataset, which can be characterized by the center of the feature distribution for certain dataset. For instance, images in the SVHN or MNIST dataset all contain numerical digits. When we train the model using the CIFAR dataset where digits are not considered crucial, the digit-related features cannot propagate through the trained weight parameters. Yet, due to the repeated appearance of digits in the dataset, the center of features for OOD (SVHN and MNIST) samples deviates from the origin in the feature space. Consequently, OOD samples exhibit dataset-dependent imbalanced classification. This excessive attention to dataset bias stemming from low dimensionalization could be a key factor contributing to the overconfident prediction of OOD samples.

¹ The transition layer is typically located just after the deepest pooling layer except the global average pooling. The exception is ResNet-18 where the transition layer is a little deeper. This may be due to insufficient low dimensionalization around the corresponding pooling layer.



Fig. 2. Layer dependence of the stable rank of the covariance matrix $\overline{\Sigma}$ and the weight matrix W for the (a) VGG-13, (b) VGG-16, (c) ResNet-18, and (d) ResNet-34 models. The dashed line indicates the transition layer. Low dimensionalization of features and weights occurs at almost the same layer.

These results are closely related to but slightly different from manifold hypothesis [28] and neural collapse [29]. The manifold hypothesis states that high-dimensional data lie on a low-dimensional (typically non-linear) manifold. It is not obvious whether DNNs can transform the non-linear input manifold to the linear output distribution and whether this kind of phenomena is relevant to OOD detection. Our results devoted to the linear method indicate that the DNN can embed input data into low-dimensional linear space, and that the low-dimensional embedding is essential for OOD detection compared with other effects. Regarding neural collapse, it states that the weight matrix at the final layer forms a low-dimensional simplex equiangular tight frame. Our analysis extends this notion beyond the final layer, revealing that several deep layers also exhibit low dimensionality and that the change of dimensionality is sharp enough to occur at the transition layer. Our layerwise analysis serves as an experimental validation of these aspects.

Detailed descriptions corresponding to each quantity are given in the following subsections along with specific results. In Sect. 3.2, we first verify the reduction of dimensionality through stable ranks. Next, we evaluate the layer dependence of OOD detection performance using the area under receiver-operating characteristic curves (AUROCs) in Sect. 3.3 to demonstrate the relevance of dimensionality to OOD detection. We further investigate multi-layer properties using CKA and noise sensitivity in Sects. 3.4 and 3.5, respectively, to eliminate the possibility that the multi-layer processing is more important than singlelayer dimensionality. After these characterizations of propagation, we examine resulting classification in Sect. 3.6 to confirm the effect of dataset bias. Finally, we showcase the utility of dimensionality through the dimensionalityaware projection-based detection method in Sect. 3.7, a modification of NuSA [19].

While the main text presents results that the ID dataset is CIFAR-10 for simplicity, additional verification can be found in Appendices F and G for different ID datasets. The similar behaviors are observed if the ID dataset is MNIST or SVHN as in Appendix G. In contrast, if we use CIFAR-100 as the ID dataset, probably due to the large number of classes, we observe poor low dimensionalization in Appendix F, leading to the insufficient separation of ID and OOD samples.



Fig. 3. Layer dependence of the AUROC detected by the Mahalanobis distance M(x) for the (a) VGG-13, (b) VGG-16, (c) ResNet-18, and (d) ResNet-34 models. Different line colors represent the different OOD datasets evaluated. The dashed line indicates the transition layer. The measured AUROCs are stabilized after transition independent of the models and datasets.



Fig. 4. Layer dependence of the AUROC detected by the projected norm $||x_{p,\varepsilon}||$ for the (a) VGG-13, (b) VGG-16, (c) ResNet-18, and (d) ResNet-34 models. Different line colors represent the different OOD datasets evaluated. The dashed line indicates the transition layer. The projection-based discrimination between ID and OOD samples becomes clear just at the transition layer.

3.2 Observation of Dimensionality via Stable Ranks

We first confirm low-dimensional characteristics of weights and features using the stable rank R_F defined by Eq. 1. In Fig. 2, we show the layer dependence of R_F of both weight matrices W and covariance matrices $\overline{\Sigma}$. The dashed line representing the transition layer is manually determined, and is located at the same position hereafter to verify the argument given in Sect. 3.1. We can see that R_F exhibits exponential growth in shallower layers corresponding to the increasing number of channels in intermediate features and weights. However, R_F of both matrices suddenly and simultaneously fall to small values around the transition layer across various architectures and small-class ID datasets (See Appendix G.). These behaviors indicate the significance of low dimensionalization in DNNs, implying the importance of dimensionality in understanding properties of OOD samples.

3.3 Transition of OOD Detection Performance

We proceed to explore the relationship between dimensionality and both featurebased and projection-based OOD detection performance using AUROCs.

In Fig. 3, we present the layer dependence of AUROCs obtained through the Mahalanobis distance defined by Eq. 2. We can see that although the observed AUROCs are highly layer-dependent and dataset-dependent in shallower layers,



Fig. 5. CKA of features in various layers for the (a,e) VGG-13, (b,f) VGG-16, (c,g) ResNet-18, and (d,h) ResNet-34 models. The upper four figures (a–d) show CKA of ID (CIFAR-10) samples, while the lower figures (e–h) show CKA of OOD (CIFAR-100) samples. In each figure, the horizontal and vertical axes represent layers, and the color bar represents CKA. Block-like saturations appear both for ID and OOD samples around the transition layer.

they consistently maintain high values independent of the datasets and architectures in deeper layers. The stabilization of AUROCs is particularly significant for the close-to-ID OOD samples, CIFAR-100 samples represented by the blue curve in this case, almost independent of the network architectures. This indicates that, as observed by Refs. [6,8], the features of OOD samples are distributed in the null space of ID covariance matrices within deeper layers. The correlation between dimensionality and detection performance provides evidence for our picture.

A similar analysis can be conducted for the projection-based method, which quantifies the difference in alignment of ID and OOD samples. Specifically, we utilize the norm of the projected feature $||x_p||$ as the OOD detection score and evaluate the layer dependence of AUROCs for some OOD datasets as depicted in Fig. 4. Note that our projection-based method modifies NuSA to incorporate singular values and analyze the convolutional layer. In Fig. 4, we observe a highly layer-dependent and dataset-dependent AUROC. However, in the vicinity of the transition layer, the AUROC consistently exhibits high values for all datasets and models, albeit not the highest. This stable detection performance indicates effective elimination of null-space components of weight matrices due to low dimensionalization of weights. This is also consistent with the stabilization of feature-based detection in subsequent layers. We again mention that, although the transition layer varies depending on network architectures, the same behavior consistently remains independent of network architectures. This universal behavior lends support to the picture outlined in Sect. 3.1.

3.4 Block Structure of CKA

We delve deeper into the relationship between dimensionality and CKA defined by Eq. 4 [21–27]. In Fig. 5 we present CKA for ID and OOD samples, represent-



Fig. 6. Noise sensitivity for the (a,e) VGG-13, (b,f) VGG-16, (c,g) ResNet-18, and (d,h) ResNet-34 models. The upper (a–d) and lower (e–g) figures show the noise sensitivities of ID (CIFAR-10) samples and OOD (CIFAR-100) samples, respectively. In each figure, the horizontal axis represents the layer and the vertical axis represents corresponding noise sensitivity. Different colors indicate the input layers where noise is injected. The dashed vertical line indicates the transition layer. The horizontal line is plotted to clarify the difference between ID and OOD samples. OOD samples are more sensitive to noise injection compared with ID samples.

ing the interlayer similarity of features. Not only in ID samples but also in OOD samples, CKA saturates in deeper layers particularly for deeper models. In addition, the saturating layer corresponds to the transition layer. This demonstrates that after low dimensionalization, the structure of features is frozen even if performing deeper layer processing. This is consistent with the stable feature-based detection performance for deeper layers, and with the picture in Sect. 3.1.

When we carefully compare interlayer CKA of ID samples with that of OOD samples, Fig. 5 reveals that the change of CKA around the transition layer in ID samples is more gradual than that in OOD samples. This indicates that low dimensionalization around the transition layer discards most of the intrinsic information of OOD samples while retaining information of ID samples, which is expected to be a characteristic of OOD samples as described in Sect. 3.1. Furthermore, the smoothness of CKA around the transition layer emerges as the most significant difference between ID and OOD samples, suggesting that dimensionality is the most important factor compared with other effects including multi-layer propagation. See Appendix D for further comparison.

3.5 Instability of OOD Samples to Noise Injection

To further investigate the multi-layer propagation property, we evaluate noise sensitivity ψ defined by Eq. 5 in Fig. 6. We can see that the difference between ID and OOD samples is determined by whether the noise-injected layer and the observed layer are deeper than the transition layer or not. If both layers are shallower or deeper than the transition layer, the difference between ID and OOD samples is not substantial. However, deep-layer features of OOD samples are more sensitive to shallow-layer noise than those of ID samples. The similar



Fig. 7. Rates of samples predicted to belong to class 0, 1, ..., 9 for the (a) VGG-13, (b) VGG-16, (c) ResNet-18, and (d) ResNet-34 models. Different line colors represent different datasets. The prediction of OOD samples is highly imbalanced compared with that of ID (CIFAR-10) samples.

Dataset	Architectures						
	VGG13	VGG16	Res18	Res34			
ID (test)	0.01 ± 0.00	0.01 ± 0.00	0.01 ± 0.00	0.01 ± 0.00			
CIFAR-100	0.52 ± 0.01	0.52 ± 0.01	0.51 ± 0.01	0.50 ± 0.02			
SVHN	1.32 ± 0.05	1.33 ± 0.07	1.45 ± 0.13	1.43 ± 0.09			
MNIST	1.54 ± 0.11	1.58 ± 0.09	1.45 ± 0.12	1.40 ± 0.16			

Table 1. Summary of the coefficient of variation (ID: CIFAR-10).

behavior of the former demonstrates that low-dimensional weights in deeper layers appropriately mitigate noise injected before them. The distinct behavior of the latter suggests that shallower-layer noise to OOD samples impacts deep-layer features dominated by dataset bias, while that to ID samples is appropriately removed from deep-layer features specific to classification. These observations provide further evidence for the picture in Sect. 3.1.

One might consider to utilize the difference in noise sensitivity for OOD detection. However, their performance falls short compared with that of the single-layer projection-based method, although sensitivity-based detection outperforms the probability-based method. See Appendix E for more details.

3.6 Dataset Bias-Induced Imbalanced Inference

To demonstrate the impact of dataset bias on classification, we examine the prediction of OOD samples for each class in Fig. 7. We can see that the prediction of ID (CIFAR-10) samples is well balanced due to the equivalent number of training samples for each class. For OOD samples, however, the imbalanced prediction is significant and seems to be common for all network architectures. This suggests that trained networks behave similarly regardless of specific architectures. A more quantitative comparison is provided by the coefficient of variation in Table 1, corresponding to the standard deviation of the prediction rates to each class. Compared with the coefficient of variation of ID (CIFAR-10) samples, those of OOD samples are extremely large even in the case of close-to-ID OOD (CIFAR-100) samples. This dataset-dependent imbalanced prediction supports the residual bias of the dataset.

ID dataset	Detection score	OOD dataset			
(Model)					
		CIFAR-10	CIFAR-100	SVHN	MNIST
CIFAR-10 (VGG-13)	Probability [3]	(ID)	0.886 ± 0.001	0.944 ± 0.005	0.927 ± 0.007
	Feature [5]	(ID)	0.898 ± 0.001	0.948 ± 0.004	0.936 ± 0.005
	Projection [19]	(ID)	0.70 ± 0.06	0.62 ± 0.07	0.47 ± 0.16
	Projection (ours)	(ID)	0.901 ± 0.001	0.960 ± 0.004	0.978 ± 0.004
CIFAR-10 (ResNet-18)	Probability [3]	(ID)	0.905 ± 0.001	0.956 ± 0.006	0.953 ± 0.009
	Feature [5]	(ID)	0.919 ± 0.001	0.958 ± 0.005	0.956 ± 0.007
	Projection [19]	(ID)	0.84 ± 0.05	0.88 ± 0.06	0.85 ± 0.14
	Projection (ours)	(ID)	0.924 ± 0.001	0.969 ± 0.003	0.982 ± 0.004
SVHN (VGG-13)	Probability [3]	0.959 ± 0.001	0.954 ± 0.001	(ID)	0.949 ± 0.009
	Feature [5]	0.965 ± 0.001	0.961 ± 0.002	(ID)	0.921 ± 0.009
	Projection [19]	0.86 ± 0.07	0.85 ± 0.06	(ID)	0.71 ± 0.20
	Projection (ours)	0.977 ± 0.002	0.973 ± 0.002	(ID)	0.965 ± 0.005
SVHN (ResNet-18)	Probability [3]	0.950 ± 0.002	0.944 ± 0.003	(ID)	0.983 ± 0.003
	Feature [5]	0.970 ± 0.002	0.967 ± 0.003	(ID)	0.918 ± 0.011
	Projection [19]	0.90 ± 0.03	0.90 ± 0.03	(ID)	0.43 ± 0.28
	Projection (ours)	0.970 ± 0.003	0.965 ± 0.004	(ID)	0.996 ± 0.002

 Table 2. Summary of OOD detection performance (AUROC) in various situations.

 All results were reproduced in our experiments.

3.7 Quantitative Comparison of OOD Detection Performance

Finally, we demonstrate the importance of dimensionality through a quantitative comparison of OOD detection performance. Table 2 summarizes the OOD detection performance detected by the ratio of norm $||x_{p,\varepsilon}^{(l)}||/||x^{(l)}||$ at the penultimate fully connected layer. As discussed in Sect. 2.3, our method is quite similar to NuSA in this case [19]. However, NuSA employs full projections ($\varepsilon = 0$), while we eliminate irrelevant singular vectors from the projection matrix based on corresponding singular values (typically $\varepsilon \sim 10^{-2}$). This minor modification markedly improves not only detection performance but also its stability. Furthermore, our dimensionality-aware projection-based method is comparable to or better than both the probability-based method [3] and the feature-based method [5]. Specifically, compared with the feature-based method, our projection-based method exhibits similar detection performance in cases with close-to-ID OOD samples, where CIFAR-10 and CIFAR-100 are treated as ID and OOD datasets. For much easier OOD samples, the projection-based method tends to outperform the feature-based method.² In summary, our modification of the projection-based

² The layer ensemble method by Ref. [5] can uplift the detection accuracy of farfrom-ID OOD samples, but it is not suitable for close-to-ID OOD detection. We checked that the AUROC value to detect CIFAR-100 OOD dataset using the ensemble method adopted by Ref. [5] is just around 0.86 for models trained by CIFAR-10. Also, the layer ensemble requires a lot of memory to save covariances, which is not suitable especially for resource-limited hardware. More seriously, the ensemble

method improves the OOD detection performance well enough to be comparable or better than other methods.

Our projection-based method is also superior to the feature-based method in terms of computational cost, especially if the number of classes K is much smaller than the feature dimensions at the OOD detection layer. The preparation of the tied covariance requires $\mathcal{O}(N_{\text{train}})$ preprocessing, and the computation of the Mahalanobis distance requires $\mathcal{O}(K)$ evaluations to find the minimum values for classes. The projection-based method, on the other hand, merely computes the projection to singular vectors, leading to the reduction of $\mathcal{O}(N_{\text{train}})$ preprocessing and $\mathcal{O}(K)$ evaluations. Also, the projection-based method is more memory-efficient compared with the feature-based method, even in single-layer detection. The null space of the low-dimensional covariance makes a large contribution to the feature-based detection as demonstrated by Ref. [6]. In fact, we checked that the best number of dimensions for the feature-based OOD detection is around 200–400 dimensions out of 512 dimensions at the penultimate layer. Meanwhile, the projection-based method eliminates the null space of the lowdimensional weight, leaving only the dimension of the number of classes. These facts make the projection-based method more efficient than the feature-based one, especially in hardware with limited computational resources.

4 Discussion

As described in Sect. 3, dimensionality plays an essential role in the propagation process, which naturally suggests a simple picture of feature propagations of ID and OOD samples. We now discuss how various properties observed in previous studies are derived from the proposed picture. We also argue the close connection with generalization of ID samples.

Our picture given in Sect. 3.1 can explain how observed properties of OOD samples in previous studies emerge from the information processing inside the DNN. The low-dimensional weight matrices first eliminate most features of the OOD samples. The features of OOD samples then aggregate around the origin with only dataset bias retained. This aggregation persists untill the final layer, yielding the small logit values of OOD samples utilized in so many studies [3, 4, 8, 30-34]. Meanwhile, dataset bias appears in the null space of the subsequent weight matrices, providing the large deviation from ID samples distributed in the linear span of principal components of the weight matrix. This orthogonality improves feature-based and projection-based detection [5–8, 19]. Furthermore, residual dataset bias in each OOD sample contains poor information, leading to the concentration of each feature at the largest singular value [9]. Our picture provides insights into how these methods work well based on the low-dimensional property of DNNs, which could help understand the nature of OOD samples in DNNs.

method by Ref. [5] requires some OOD samples, although it would be practically inaccessible in cases where we do not know what kinds of OOD samples are contaminated.

The importance of dimensionality can also explain the positive correlation between ID generalization and OOD detection [32], and the hardness of OOD detection in large-scale classification [35]. In the context of ID generalization error analysis, the importance of low dimensionalization has also been noted [20,36,37]. Combined with our results, a positive correlation between ID generalization and OOD detection is directly derived. In addition, we observe poor OOD detection performance in cases with a large number of classes in ID samples (CIFAR-100) due to insufficient low dimensionalization (See Appendix F). This is also consistent with the achievement of both ID generalization and OOD detection as long as the internal feature dimension is sufficiently larger than the output dimension.

The relationship between ID optimization and OOD detection is also likely to correlate with the data complexity dependence of the suitable detection layer [10]. The optimization of the model using ID samples induces an overfittinglike behavior in ID samples with reference to their data complexity. The loss of information is thus more significant for OOD samples possessing distinct data complexity from ID samples, leading to dataset bias-dominated features even at shallower layers. This might be the reason why the feature-based detection method performs extremely well in shallower layers for far-from-ID OOD samples as in Fig. 3, and why CKA of far-from-ID OOD samples tends to be retained in shallower layers as in Appendix D. Furthermore, when we closely compare AUROCs of various layers with the data complexity provided by Ref. [10], rather than the data complexity itself, the difference in data complexity between ID and OOD samples determines the suitable layer for OOD detection. In this sense, the propagation properties investigated in our study are expected to correlate with the data complexity-based method, although further studies are required to clarify this.

5 Summary and Conclusion

Based on various analyses investigating how ID and OOD samples propagate inside DNNs, we demonstrated the importance of dimensionality for various properties of OOD samples. The systematic and detailed analyses enable us to understand the layerwise processing and their effects on the separation of ID and OOD samples. In addition, we verified the usefulness of dimensionality for OOD detection. These observations might serve as the baseline for understanding the nature of OOD samples in DNNs and for improving OOD detection performance.

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Towards Open-World Cross-Domain Sequential Recommendation: A Model-Agnostic Contrastive Denoising Approach

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Abstract. Cross-domain sequential recommendation (CDSR) aims to address the data spCH). Recently, some SR approaches have utilized auxiliary behaviors to complement the information for long-tailed users. However, these methods cannot deliver promising performance in CDSR, as they overlook the semantic gap between target and auxiliary behaviors, as well as user interest deviation across domains (2nd CH). In this paper, we propose a model-agnostic contrastive denoising (MACD) approach towards open-world CDSR. We introduce auxiliary behavior sequence information (i.e., clicks) into CDSR methods to explore potential interests. Specifically, we design a denoising interest-aware network combined with a contrastive information regularizer to remove inherent noise from auxiliary behaviors and exploit multi-interest from users. Extensive offline experiments on public industry datasets and a standard A/B test on a large-scale financial platform with millions of users both confirm the remarkable performance of our model in open-world CDSR scenarios. Code and dataset are available at https://github.com/ WujiangXu/MACD.

Keywords: Open-world Recommendation \cdot Cross-Domain Sequential Recommendation \cdot Sequential Recommendation \cdot Contrastive Learning

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Fig. 1. Previous methods focus on constructing their structure based on overlapping users with rich behaviours (a) under a closed-world environment. In this study, we aim to design the model for an open-world environment that accounts for the majority of long-tailed users (b) and cold-start users (c) with sparse historical behaviours.

1 Introduction

Sequential Recommendation (SR) [4,6,16,18] has gained significant attention in recent years due to its ability to model dynamic user preferences. However, in real-world platforms, users often exhibit partial behaviors within specific domains, leading to biased observed preferences based on single-domain interactions. To address this issue, cross-domain sequential recommendation (CDSR) methods [1, 8, 10, 11] have been proposed to construct models using multi-domain data, transferring rich information from other relevant domains to improve performance in multiple domains simultaneously. To transfer the information across domains, the core idea of CDSR works is to extend the single-domain sequential recommendation methods via designing a mapping function. Specifically, PiNet [11] employs a designed gating mechanism to filter information and propagate this information for overlapping users from the source domain to the target domain. Similarly, DASL [8] proposes a dual-attention mechanism to extract the user-shared information for each domain. To enhance the correlation between single- and cross-domain user preferences, C²DSR [1] combines an attention information transferring unit with a contrastive infomax objective.

Despite the promising improvements, these mapping function designs of existing CDSR methods heavily rely on overlapping users with rich behaviors, which can lead to unsatisfactory performance in an open-world environment [21,22]. Typically, these methods assume full or majority overlap of users (Fig. 1(a)) across domains, which is only a minority in real-world platforms such as Taobao or Amazon. These approaches disregard most users in the open-world CDSR environment, particularly the long-tailed users (Fig. 1(b)) with few interactions and the cold-start users (Fig. 1(c)) who are only present in a single domain. As a result, these approaches fail to provide sufficient exploration of cross-domain information to long-tailed and cold-start users, leading to an incomplete understanding of their interests. Therefore, the first challenge is: How can we explore complementary interest information to enhance the model's performance towards open-world CDSR scenarios, where the majority of users are long-tailed or coldstart users with sparse historical behaviors?

Recent SR studies [5, 23] have explored the integration of multi-typed behaviors to improve the performance of long-tailed users. In real-world platforms, users interact with items through multiple types of behavior. The target behavior (e.g. purchase) that directly benefits businesses is usually sparser than other behavior types (e.g. view or click). To capture behavior semantics, MBGCN [5] designs a user-item and item-item graph neural network, while MBHT [28] proposes a hypergraph-based transformer to encode behavior-aware sequential patterns from both fine-grained and coarse-grained levels. More recently, DPT [29] develops a denoising and prompt-tuning framework to guide the informative and relative representations. However, these multi-behavior denoising SR approaches cannot fully address the CDSR problem as they fail to consider the deviation of user interests across domains and neglect the semantic gap between auxiliary and target behaviors. Therefore, the second challenge is: How can the semantic gap between target and auxiliary behaviors be reduced and user interest deviation be learned when utilizing auxiliary behavior sequences to enhance information for long-tailed users in CDSR?

To address the aforementioned challenges, we propose a Model-Agnostic Contrastive Denoising approach, namely MACD. Overall, our major contributions can be summarized as follows:

- (1) We propose a model-agnostic contrastive denoising framework, namely **MACD**, towards open-world CDSR that can be integrated with most off-the-shelf SR methods. To the best of our knowledge, we are the first who utilize the auxiliary behaviour information in CDSR models, which incorporate informative potential interests of users, especially for long-tailed users and cold-start users.
- (2) We propose a denoising interest-aware network that incorporates an intradomain/cross-domain denoising module and a contrastive information regularizer. This network aims to reduce the semantic gap between target and auxiliary behaviors, enabling to better capture of user interest.
- (3) We introduce a fusion gate unit to enhance the fusion of representations, and we employ a parameter-free inductive representation generator to generate inductive representations for cold-start users during the inference stage.
- (4) We conduct extensive experiments on a large-scale scenario with a minority of overlapping users, representing an open-world environment. Furthermore, a standard A/B test is conducted to validate our performance on a real-world CDSR financial platform with millions of daily traffic logs.

2 Methodology

2.1 Problem Formulation

In this work, we consider a general CDSR scenario in an open-world environment that consists of partially overlapping users, a majority of long-tailed and cold-start users across two domains, namely domain X and domain Y. The data is denoted by $D^X = (\mathcal{U}^X, \mathcal{V}^X, \mathcal{E}^X)$ and $D^Y = (\mathcal{U}^Y, \mathcal{V}^Y, \mathcal{E}^Y)$, where \mathcal{U}, \mathcal{V} , and \mathcal{E} are the sets of users, items, and interaction edges, respectively. For a given user, we denote their target item interaction sequences in chronological order as $S^X = [v_1^X, v_2^X, \cdots, v_{|S^X|}^X]$ and $S^Y = [v_1^Y, v_2^Y, \cdots, v_{|S^Y|}^Y]$, where $|\cdot|$ denotes the length of the sequence. We further introduce the user's auxiliary behavior sequences, $C^X = [v_1^X, v_2^X, \cdots, v_{|C^X|}^X]$ and $C^Y = [v_1^Y, v_2^Y, \cdots, v_{|C^Y|}^Y]^1$ to enrich user behaviors and solve the issue that previous CDSR methods being heavily affected by data sparsity problems. The adjusted objective of CDSR in our setting is to predict the next item for a given user based on both user's target behavior sequence and auxiliary behavior sequence:

$$\max P^{X} \left(v_{|S^{X}|+1}^{X} = v | S^{X}, S^{Y}, C^{X}, C^{Y} \right), \text{if } v \in \mathcal{V}^{X}.$$
$$\max P^{Y} \left(v_{|S^{Y}|+1}^{Y} = v | S^{X}, S^{Y}, C^{X}, C^{Y} \right), \text{if } v \in \mathcal{V}^{Y}.$$

In the cold-start setting, where we have users only observed in one domain (e.g., Y) and thus being cold-start in another domain (e.g., X), the objective function can be generalized by adding an additional condition that $S^X = \emptyset, C^X = \emptyset$. Moreover, we classify users into two categories based on the length of their target sequence. If users' target sequence length, denoted as $|S^X|$, is less than L^X_{head} , they are categorized as long-tailed users; otherwise, they are classified as head users. We determine L^X_{head}, L^Y_{head} by calculating the average length of the top 20% longest sequences in the specific domain. It should be noticed that the definition of long-tailed users is independent across different domains (Fig. 2).

2.2 Embedding Encoder

Embedding Layer. To obtain the initialized sequence representations $\mathbf{S}^{X} = {\mathbf{h}_{s_{1}}^{'X}, \cdots, \mathbf{h}_{s_{T}}^{'X}}$ and $\mathbf{S}^{Y} = {\mathbf{h}_{s_{1}}^{'Y}, \cdots, \mathbf{h}_{s_{T}}^{'Y}}$, we utilize embedding layers $\mathbf{E}^{X} \in \mathbb{R}^{|\mathcal{V}^{X}| \times d}$ and $\mathbf{E}^{Y} \in \mathbb{R}^{|\mathcal{V}^{Y}| \times d}$, where d denotes the dimension of the embeddings and T is the maximum length of the interaction sequence. If the sequence length is larger than T, we only consider the most recent T actions. If the sequence length is less than T, we repeatedly add a 'padding' item to the left until the length is T. The initialized embeddings $\mathbf{C}^{X} = {\mathbf{h}_{c_{1}}^{'X}, \cdots, \mathbf{h}_{c_{T'}}^{'X}}$ and $\mathbf{C}^{Y} = {\mathbf{h}_{c_{1}}^{'Y}, \cdots, \mathbf{h}_{c_{T'}}^{'Y}}$ for the auxiliary behavior sequences are obtained in

¹ Due to the low cost of engagement, the auxiliary behaviors (e.g., clicks or views) provide richer records than the target user-item interactions (e.g., purchases).
the same manner. T' denotes the maximum length of the auxiliary behavior sequence, which is greater than T. Moreover, we introduce learnable position embedding matrixes $P_{S} \in \mathbb{R}^{T \times d}$ and $P_{C} \in \mathbb{R}^{T' \times d}$ to improve the ordered information of sequence embeddings.

Sequential Information Encoder. Our model-agnostic framework can directly integrate with off-the-shelf SR methods [4,6,16], eliminating the need to design a sequential information encoder. thus, we do not modify the sequential information encoders of SR methods further in our work. For simplicity, we denote their sequential information encoder as a function $\mathcal{F} = \{\mathcal{F}^X, \mathcal{F}^Y, \mathcal{F}^X_c, \mathcal{F}^Y_c\}$. Formally, the procedure is as: $\mathbf{h}^X_{s_1}, \cdots, \mathbf{h}^X_{s_T} = \mathcal{F}^X(\mathbf{h}_{s_1}'^X, \cdots, \mathbf{h}_{s_T}'); \quad \mathbf{h}^Y_{s_1}, \cdots, \mathbf{h}^Y_{s_T} = \mathcal{F}^Y(\mathbf{h}_{s_1}'^Y, \cdots, \mathbf{h}_{s_T}'); \quad \mathbf{h}^X_{c_1}, \cdots, \mathbf{h}^X_{c_{T'}} = \mathcal{F}^X_c(\mathbf{h}_{c_1}'^X, \cdots, \mathbf{h}_{c_{T'}}')$ and $\mathbf{h}^Y_{c_1}, \cdots, \mathbf{h}^Y_{c_{T'}} = \mathcal{F}^Y_c(\mathbf{h}_{c_1}'^Y, \cdots, \mathbf{h}_{c_{T'}}')$.



Fig. 2. Overview of our MACD approach. Unlike previous CDSR methods, our MACD is a general and model-agnostic approach that can be integrated with most off-the-shelf SDSR methods. Our MACD fully leverages auxiliary sequences to explore the potential interests in an open-world CDSR scenario. The denoising interest-aware network (**DIN**) not only explores explicit interests within the domain but also transfers implicit interest information across domains. With the abundant purified auxiliary sequence information, the representations of long-tailed users can be enhanced. Furthermore, through a well-designed contrastive information regularizer in the DIN and the fusion gate unit, our MACD minimizes the semantic gap and interest deviation between the target and auxiliary behaviors.

2.3 Denoising Interest-Aware Network

To fully leverage the information in the auxiliary behavior sequence, we propose a denoising interest-aware network that consists of an intra-domain denoising module (**IDDM**), a cross-domain denoising module (**CDDM**), and a contrastive information regularizer. The IDDM is designed to explicitly explore the user's interests, while the CDDM extracts latent interests and transfers cross-domain knowledge. To ensure that the explicit- and implicit-interest representations are consistent with user interests learned from the target behaviour sequences, we also introduce a novel contrastive information regularizer.

Intra-domain Denoising Module. Given a user u with the sequence representations **S** and **C**, we perform the intra-domain denoising procedure for each domain separately. We introduce an explicit-interest guided multi-head attention mechanism to efficiently extract useful information from the noisy auxiliary behavior sequence. Using a single attention head is insufficient since our objective is to extract multiple explicit interests from the users. Therefore, we modify the Multi-Head Attention mechanism described in [19] to eliminate redundant and unrelated information. The explicit-interest representation $\mathbf{S}^{*X} = \{\mathbf{h}_{s_1}^{*X}, \dots, \mathbf{h}_{s_T}^{*X}\}$ is obtained by $\mathbf{h}_{s_1}^{*X}, \dots, \mathbf{h}_{s_T}^{*X} = Concat(head_1, \dots, head_h)W^O$. where $head_i = Attention(QW_i^Q, KW_i^K, VW_i^V)$ and

$$Q = \mathbf{h}_{s_1}^X, \cdots, \mathbf{h}_{s_j}^X, \cdots, \mathbf{h}_{s_T}^X, \ j \in [1, T]$$

$$\tag{1}$$

$$K, V = \mathbf{h}_{c_1}^X, \cdots, \mathbf{h}_{c_{j'}}^X, \cdots, \mathbf{h}_{c_{T'}}^X, \ j' \in [1, T']$$
(2)

$$Attention(Q, K, V) = softmax(\frac{QK^{T}}{\sqrt{d_k}})V,$$
(3)

where $W^O \in \mathbb{R}^{d \times d}$, $W_i^Q \in \mathbb{R}^{d \times d_k}$, $W_i^K \in \mathbb{R}^{d \times d_k}$ and $W_i^V \in \mathbb{R}^{d \times d_k}$ are trainable matrices, $d_k = d/h$ and h is the number of attention heads. Similar intra-domain denoising processes with different matrix weights are performed on domain Y to obtain the explicit-interest representation $\mathbf{S}^{*Y} = \{\mathbf{h}_{s_1}^{*Y}, \cdots, \mathbf{h}_{s_T}^{*Y}\}.$

Cross-Domain Denoising Module. In this component, we conduct the crossdomain denoising operation to explore the users' implicit interest and transfer the cross-domain knowledge. Being similar to the intra-domain denoising module, we adopt an implicit-interest guided multi-head attention mechanism to purify the auxiliary behaviors representation. Specifically, to obtain user's refined implicitinterest representation in domain X, $\hat{\mathbf{S}}^{X} = {\{\hat{\mathbf{h}}_{s_{1}}^{*X}, \cdots, \hat{\mathbf{h}}_{s_{T}}^{X}\}}$, we define the Query Q, Key K, and Value V as follows:

$$Q = \mathbf{h}_{s_1}^X, \cdots, \mathbf{h}_{s_j}^X, \cdots, \mathbf{h}_{s_T}^X, \ j \in [1, T]$$

$$\tag{4}$$

$$K, V = \mathbf{h}_{c_1}^Y, \cdots, \mathbf{h}_{c_{j'}}^Y, \cdots, \mathbf{h}_{c_{T'}}^Y, \ j' \in [1, T']$$
(5)

The rest steps are the same as Eqs. 1–3. The cross-domain denoising module in the domain Y is similar, and we can obtain the refined implicit-interest representation in domain Y, $\hat{\mathbf{S}}^Y = {\{\hat{\mathbf{h}}_{s_1}^{*Y}, \cdots, \hat{\mathbf{h}}_{s_T}^Y\}}$ correspondingly. Our motivation behind developing the denoising interest-aware network is to extract purified information as comprehensively as possible from the noisy but abundant auxiliary behavior data. To accomplish this, we consider the longer sequence of auxiliary behavior \mathbf{C} as the key and value, rather than \mathbf{S} .

Contrastive Information Regularizer. In our intra- and cross-domain denoising modules, our aim is to extract different representations that incorporate explicit and implicit interests respectively. Moreover, we aim to minimize the variance between various interest representations of the same user and maximize the difference between that of different users by the proposed contrastive information regularizer. One of the crucial aspects of Contrastive Learning is to extract valid positive-negative pair samples from different views. In our design, we consider the composite purified interest representation of the same user u_i as positive pairs, while interest representations from different users u_i, u_j are considered as negative pairs. Therefore, taking domain X as an example, we obtain the positive and negative sample pairs $\{\mathbf{S}_{u_i}^{*X}, \hat{\mathbf{S}}_{u_i}^X\}$ and $\{\mathbf{S}_{u_i}^{*X}, \hat{\mathbf{S}}_{u_j}^X\}$, where $u_i, u_j \in \mathcal{U}^X$ and $i \neq j$. Formally, we define a contrastive information regularizer based on InfoNCE [13] loss, as follows:

$$\mathcal{L}_{cl}^{X} = \sum_{u_i \in \mathcal{U}^{X}} -\log \frac{\exp(s(\phi(\mathbf{S}_{u_i}^{*X}), \phi(\mathbf{\hat{S}}_{u_i}^{X}))/\tau)}{\sum\limits_{u_i \in \mathcal{U}^{X}, i \neq j} \exp(s(\phi(\mathbf{S}_{u_i}^{*X}), \phi(\mathbf{\hat{S}}_{u_j}^{X}))/\tau)}$$
(6)

The hyper-parameter τ regulates the smoothness of the softmax curve, while the pair-wise distance function $s(\cdot)$ evaluates the similarity between positive and negative pairs. The function $\phi(\cdot)$ applies a simple Mean operation to average the embedding along the temporal dimension. \mathcal{L}_{cl}^{Y} is defined in a similar way. The proposed contrastive information regularizer enables our framework to learn more robust user interest-aware representations that exhibit high consistency while being capable of distinguishing personal preferences among different users.

2.4 Fusion Gate Unit

To represent user's holistic preferences in each domain with distilled embeddings $\mathbf{S}, \mathbf{S}^*, \hat{\mathbf{S}}$, we introduce a gate unit to fuse them. Firstly, we utilize $\phi(\cdot)$ to aggregate their representation along the temporal dimension and obtain the corresponding embeddings $\mathbf{O} \in \mathbb{R}^{1 \times d}, \mathbf{O}^* \in \mathbb{R}^{1 \times d}, \hat{\mathbf{O}} \in \mathbb{R}^{1 \times d}$. It is a straightforward way to apply element-wise add or concatenating operation to fuse feature. However, the explicit- and implicit- interest representation $\mathbf{O}^* \in \mathbb{R}^{1 \times d}, \hat{\mathbf{O}} \in \mathbb{R}^{1 \times d}$ are learned from the auxiliary behavior sequences, which are less important than the sequences $\mathbf{O} \in \mathbb{R}^{1 \times d}$. Considering different weights to fuse them can explore the full potential of embeddings. A learnable weight matrix is first learned and then conduct a gate control to fuse the embeddings, *i.e.*,

$$\mathbf{H}_1^X = \operatorname{Sig}(\mathbf{O}^X \mathbf{W}_1 + b_1 + \mathbf{O}^{*X} \mathbf{W}_2 + b_2), \tag{7}$$

$$\mathbf{O}_m^X = \tanh((1 - \mathbf{H}_1^X) \odot \mathbf{O}^X + \mathbf{H}_1^X \odot \mathbf{O}^{*X}), \tag{8}$$

where $\operatorname{Sig}(\cdot)$ is sigmoid function and $\tanh(\cdot)$ is the hyperbolic tangent function. \odot means the element-wise multiplication operation weighting the embeddings in a fine-grained way. $\mathbf{W}_1, \mathbf{W}_2 \in \mathbb{R}^{d \times d}$ and $b_1, b_2 \in \mathbb{R}^d$ are the trainable weights and bias updated in backpropagation. Then we fuse the intermediate representation \mathbf{O}_m^X with $\hat{\mathbf{O}}^X$ in a similar way as follows.

$$\mathbf{H}_{2}^{X} = \operatorname{Sig}(\mathbf{O}_{m}^{X}\mathbf{W}_{3} + b_{3} + \hat{\mathbf{O}}^{X}\mathbf{W}_{4} + b_{4}), \tag{9}$$

$$\bar{\mathbf{O}}^X = \tanh((1 - \mathbf{H}_2^X) \odot \mathbf{O}_m^X + \mathbf{H}_2^X \odot \hat{\mathbf{O}}^X), \tag{10}$$

where \mathbf{W}_3 , $\mathbf{W}_4 \in \mathbb{R}^{d \times d}$ and $b_3, b_4 \in \mathbb{R}^d$ are the learnable weights and bias. Similarly, $\bar{\mathbf{O}}^Y$ can be obtained. To generate final prediction results, building upon prior research [3], we utilize the concatenated representations of the user and item $(\bar{\mathbf{O}}_{u_i}^X, v_j^X)$ as input to a multi-layer perceptron to generate predictions of user's preference for a target item. Formally, $\hat{y}_{u_iv_j}^X = \sigma(\mathrm{MLPs}(\bar{\mathbf{O}}_{u_i}^X || v_j^X))$. where MLPs are the stacked fully connected layers and v_j^X is the target item embedding. σ denotes the sigmoid function. The prediction layer in the domain Y is similar.

2.5 Model Training

Our overall training loss combines the contrastive information regularizers L_{cl}^X and L_{cl}^Y mentioned in Eq. (7) with the classification losses L_{cls}^X and L_{cls}^Y , using a harmonic factor λ :

$$\mathcal{L} = \lambda (\mathcal{L}_{cl}^X + \mathcal{L}_{cl}^Y) + (1 - \lambda) (\mathcal{L}_{cls}^X + \mathcal{L}_{cls}^Y)$$
(11)

where L_{cls}^X and L_{cls}^Y represent the classification loss for the recommendation task in domains X and Y, respectively. To be specific, L_{cls}^X is obtained by summing over the losses $\ell(\hat{y}_{u_iv_j}^X, y_{u_iv_j}^X)$ for all user-item pairs (u_i, v_j) in domain X:

$$\mathcal{L}_{cls}^{X} = \sum_{u_i \in \mathcal{U}^X, v_j \in \mathcal{V}^X} \ell(\hat{y}_{u_i v_j}^X, y_{u_i v_j}^X)$$
(12)

where $\hat{y}_{u_i v_j}^X$ represents the predicted results and $y_{u_i v_j}^X$ is the corresponding ground-truth label. The loss estimator $\ell(\cdot)$ in Eq.(14) is used to measure the dissimilarity between the predicted results and the ground truth labels, and it is defined as the binary cross-entropy loss shown: $\ell(\hat{y}, y) = -[y \log \hat{y} + (1-y) \log(1-\hat{y})]$. \hat{y} represents the predicted results, and y represents the ground truth label. Similarly, L_{cls}^Y can be obtained.

2.6 Inductive Representation Generator

In the inference stage, we propose a novel inductive representation generator (IRG) to enhance the model's performance for cold-start users². The core idea

 $^{^{2}}$ For instance, in the scenario where domain X is the target domain for item recommendation, while cold-start users only have observed items in domain Y.

```
Algorithm 1: Pseudocode of IRG
```

```
# X,Y:two different domains, N:batch size, d:embedding dimension.
# \bar{\mathbf{O}}^X, \bar{\mathbf{O}}^Y: user's representation after FGU, shape: [N,d]
# ColdStartLabel<sup>X</sup>, ColdStartLabel<sup>Y</sup>:identify the cold-start users,
shape: [N]
# Compute Normalization Coefficient.
Coefficient<sup>Y</sup> \leftarrow Norm(\bar{\mathbf{O}}^Y, Axis=1) * Transpose(Norm(\bar{\mathbf{O}}^Y, Axis=1))
# Find the similar users.
Similarity<sup>Y</sup> \leftarrow Dot_Product(\bar{\mathbf{O}}^Y,Transpose(\bar{\mathbf{O}}^Y))/Coefficient<sup>Y</sup>
# The index of the most similar users.
NearestIndex<sup>X</sup> \leftarrow Argmax(Similarity<sup>Y</sup>,Axis=1)
# Generate the cold-start users' representation from similar users.
ColdStartRepr<sup>X</sup> \leftarrow \bar{\mathbf{O}}^X[NearestIndex<sup>X</sup>]
# Select the user representation.
LastRepr<sup>X</sup> \leftarrow Where(ColdStartLabel<sup>X</sup>, ColdStartRepr<sup>X</sup>, \bar{\mathbf{O}}^X)
# Repetition for domain Y.
```

behind this generator is to retrieve similar user embeddings from another domain whose interests closely align with cold-start users. Specifically, for a cold-start user u^* in domain X, we first calculate the user-user similarity based on the feature $\bar{\mathbf{O}}^Y$. We then identify the nearest user u_l and utilize their embedding $\bar{\mathbf{O}}_{u_l}^X$ in domain X to replace the embedding of the cold-start user u^* . An example of PyTorch-style implementation is described in Algorithm 1.

3 Experiments

In this section, we present extensive experiments to demonstrate the effectiveness of MACD, aiming to answer the following three research questions (RQs).

- RQ1: How does MACD compare to state-of-the-art methods in open-world CDSR scenarios, particularly for long-tailed and cold-start users?
- RQ2: How do the different modules of MACD contribute to the performance improvement of our method?
- RQ3: Whether MACD can achieve a significant improvement when deployed on a real-world industry platform?
- RQ4: When encountering open-world scenarios with varying user-item interaction density and different numbers of cold-start users, can MACD consistently achieve remarkable performance?
- RQ5: How do different hyperparameter settings affect the performance of our method?

Owing to space constraints, additional results and analyses-encompassing baseline descriptions, experiments on the "Phone-Elec" dataset.

3.1 Datasets

We conducted offline experiments on a publicly available Amazon dataset comprised of 24 distinct item domains. To generate CDSR scenarios for our experiments, we selected two pairs of domains, namely "Cloth-Sport" and "Phone-Elec". We extracted users who had interactions in both domains and then filtered out items with fewer than 10 interactions. We used the views behaviors as auxiliary behaviors, and both target behavior sequences and auxiliary behavior sequences were collected in chronological order. To prevent the information leak problem in previous works [10,11], we then divided users into three sets: 80% for training, 10% for validation, and 10% for testing in each domain. To simulate multiple open-world recommendation scenarios, we retained non-overlapping users and varied the overlapping ratio to control the number of overlapping users. In addition, we randomly selected approximately 20% of overlapping users as cold-start users for validation and testing. The detailed statistics of our corrected datasets in CDSR scenarios are summarized in Table 1.

3.2 Experiment Setting

Evaluation Protocol. To test the effectiveness of our approach under an openworld environment, we varied the overlapping ratio \mathcal{K}_u of each dataset in {25%, 75%}, which corresponded to different numbers of overlapping users shared across the domains. For example, in the Amazon "Cloth-Sport" dataset with $\mathcal{K}_u = 25\%$, we determined the number of overlapping users by applying the formula 28,771 * 0.25 = 7192. To generate an unbiased evaluation for fair comparison [7], we randomly sampled 999 negative items, which were items not interacted with by the user, along with 1 positive item that served as the ground-truth interaction. We then employed these items to form the recommended candidates for conducting the ranking test. We used several *top-N* metrics to assess the effectiveness of our approach, including the normalized discounted cumulative gain (NDCG@10) and hit rate (HR@10). Higher values of all metrics indicate improved model performance. All the experiments were conducted five times and the average values are reported.

Compared Methods. To verify the effectiveness of our model in an openworld environment, we compare MACD with three branches of baselines including SDSR methods(BERT4Rec [16], GRU4Rec [4], SASRec [6]), Denoising SR

Table 1. Statistics on the Amazon datasets. #O: the number of overlapping users across domains.

Dataset		$ \mathcal{U} $	$ \mathcal{V} $	$ \mathcal{E} $	#O	S	C	Density
Amazon	Cloth	76,162	$51,\!350$	888,768	28,771	14.82	124.83	0.023%
	Sport	$225,\!299$	48,691	$2,\!205,\!976$		12.31	80.37	0.020%
Amazon	Phone	$1,\!440,\!005$	$528,\!873$	18,832,424	116,211	15.17	175.17	0.002%
	Elec	194,908	49,295	2,096,841		12.31	80.37	0.022%

methods (HSD [30], DPT [29]) and CDSR methods (Pi-Net [11], DASL [8], C²DSR [1]). In this paper, we do not choose conventional CDR methods as baselines because they overlook sequential information importance.

Parameter Settings. We set the embedding dimension d to 128 and the batch size to 2048. The training epoch is fixed at 100 to obtain optimal performance and the comparison baselines employ other hyper-parameters as reported in their official code implementation. For MACD, we set the maximum length T of the main behaviour sequence S to 20 and the maximum length T' of the target behaviour sequence C to 100. The number of attention heads h is set to 8. The harmonic factor λ is selected from a range of values between 0.1 and 0.9 with a step length of 0.1, and the hyper-parameter τ in \mathcal{L}_{cl} is set to 1. Function $s(\cdot)$ is implemented with the L2 distance. Each approach is run five times under five different random seeds and the optimal model is selected based on the highest NDCG@10 performance on the validation set, using a grid search approach.

3.3 Performance Comparisons (RQ1)

Quantitative Results. Table 2 presents the performance results for HR@10 and NDCG@10 evaluation metrics across all users. Additionally, to specifically

Methods	Cloth-domain recommendation				Sport-domain recommendation			
	$\mathcal{K}_u=25\%$		$\mathcal{K}_u = 75\%$		$\mathcal{K}_u=25\%$		$\mathcal{K}_u = 75\%$	
	NDCG@10	HR@10	NDCG@10	HR@10	NDCG@10	HR@10	NDCG@10	HR@10
BERT4Rec [16]	4.49	9.38	5.04	10.22	7.87	15.21	8.07	15.37
BERT4Rec [†] [16]	5.41	11.27	6.53	13.33	9.05	17.59	9.90	18.79
GRU4Rec [4]	5.81	12.03	6.56	13.22	9.98	19.20	10.56	20.07
GRU4Rec [†] [4]	5.79	12.24	6.52	13.70	10.67	20.05	10.68	20.27
SASRec [6]	5.88	12.21	6.43	13.09	10.06	19.36	10.49	20.20
SASRec [†] [6]	5.79	11.87	6.59	13.23	10.22	19.63	10.92	20.29
DPT† [29]	5.39	11.77	6.30	13.02	10.25	20.07	10.40	20.35
$\mathrm{BERT4Rec}\ [16]\ +\ \mathrm{HSD}\ \dagger\ [30]$	5.53	11.41	6.50	13.47	9.13	17.65	10.19	18.98
$\mathrm{GRU4Rec}~[4] + \mathrm{HSD} \dagger [30]$	5.99	12.40	6.67	13.77	10.80	20.23	10.85	20.46
SASRec $[6]$ + HSD \dagger $[30]$	5.97	12.33	6.70	13.38	10.30	19.66	11.01	20.35
Pi-Net [11]	6.18	12.24	6.60	13.10	10.03	19.10	10.74	19.96
Pi-Net† [11]	6.19	12.45	6.78	13.51	10.02	19.72	10.51	20.54
DASL [8]	6.13	12.67	6.60	13.04	10.42	19.63	10.51	20.02
DASL† [8]	6.21	12.50	6.52	13.68	10.61	20.09	10.37	19.81
C^2DSR [1]	6.16	12.51	6.54	13.61	10.26	20.36	11.00	20.23
C^2DSR^{\dagger} [1]	6.10	12.62	6.51	13.64	10.40	20.37	11.04	20.56
$\mathrm{BERT4Rec}\;[16] + \mathbf{MACD}$	6.54	13.37	7.29	14.51	11.12	20.75	11.72	21.69
$\mathrm{GRU4Rec}\;[4]+\mathbf{MACD}$	6.77*	13.47*	7.41	14.58	11.14	20.90	11.63	21.81
$\mathrm{SASRec}\;[6] + \mathbf{MACD}$	6.69	13.16	7.45*	14.61*	11.28*	21.02*	11.89*	21.82*
Improvement(%)	9.02	6.31	9.88	6.80	6.31	3.19	7.70	6.13

Table 2. Experimental results (%) on the bi-directional Cloth-Sport CDSR scenario with different \mathcal{K}_u . The best results for each column are highlighted in boldface, while the second-best results are underlined.

 \dagger indicates whether the compared models utilize auxiliary behavior sequences. "*" denotes statistically significant improvements ($p \leq 0.05$), as determined by a paired t-test comparison with the second best result in each case.

examine the performance on long-tailed users and cold-start users, we provide a separate comparison in Table 3. To ensure a fair comparison, we also provided the auxiliary behavior sequences to other state-of-the-art models. With minor adjustments to the models, we utilized their designed sequential information encoder to encode the auxiliary sequences, concatenating them with the target sequence embedding before feeding them into the prediction layer. Regarding the overall performance, our MACD framework equipped with the SDSR baselines achieve a significant improvement on Amazon datasets compared to the second-best baselines in the simulated open-world CDSR scenario, which contains a substantial number of long-tailed and cold-start users. Additionally, we make the following insightful findings:

- In most cases, both the SDSR and CDSR baselines may experience a drop in performance when using the auxiliary behaviors directly. This is because the abundance of auxiliary behaviors can bring irrelevant information to the recommender, leading to erroneous target behavior representation learning.
- The denoising SR method (such as HSD) improves the backbone's performance benefitting from the noiseless auxiliary behaviors and bridging the behavioral semantic gap compared to SDSR methods. But the denoising SR

Methods	Cloth-domain recommendation				Sport-domain recommendation			
	$\mathcal{K}_u = 25\%$				$\mathcal{K}_u=25\%$			
	long-tailed		cold-start		long-tailed		cold-start	
	NDCG@10	HR@10	NDCG@10	HR@10	NDCG@10	HR@10	NDCG@10	HR@10
BERT4Rec [16]	4.15	8.81	5.13	9.70	5.65	11.55	6.34	13.52
BERT4Rec [†] [16]	5.36	11.00	5.17	10.21	5.72	11.72	6.39	13.73
GRU4Rec [4]	5.38	11.38	4.80	10.00	6.48	12.01	6.76	14.63
GRU4Rec [†] [4]	5.24	11.34	4.62	10.15	6.86	12.81	7.23	15.35
SASRec [6]	5.40	11.35	4.76	10.15	6.41	13.64	7.44	15.93
SASRec [†] [6]	5.36	11.34	4.97	10.00	6.63	13.58	7.37	15.00
DPT† [29]	5.35	11.14	5.03	10.23	5.81	11.67	6.35	13.63
BERT4Rec $[16]$ + HSD † $[30]$	5.31	11.10	5.01	10.19	5.77	11.65	6.32	13.70
GRU4Rec $[4]$ + HSD \dagger $[30]$	5.27	11.39	4.64	10.10	6.90	12.84	7.25	15.33
SASRec $[6]$ + HSD † $[30]$	5.38	11.38	4.99	10.05	6.61	13.74	7.40	15.03
Pi-Net [11]	5.24	11.06	4.73	10.15	7.16	14.01	6.78	16.13
Pi-Net† [11]	5.37	11.57	4.53	10.07	7.21	13.93	7.05	16.16
DASL [8]	5.15	10.31	4.94	9.93	7.62	14.11	7.34	16.58
DASL† [8]	5.43	10.77	5.08	9.95	7.43	14.39	7.51	16.81
C^2DSR [1]	5.52	11.55	4.30	9.20	7.50	14.32	7.37	17.21
$C^2DSR\dagger$ [1]	5.58	11.56	4.37	9.78	7.40	14.45	7.69	17.29
BERT4Rec $[16] + MACD$	5.90	11.95	5.56	10.74	7.52	14.51	8.09	17.77
$\mathrm{GRU4Rec}\;[4] + \mathbf{MACD}$	6.08	12.07*	5.93*	10.91*	7.78	14.77	8.13	17.61
$\mathrm{SASRec} \ [6] + \mathbf{MACD}$	6.09*	11.97	5.54	10.82	8.02*	15.56*	8.26*	17.85*
Improvement(%)	9.14	4.41	16.73	6.65	5.25	7.68	7.41	3.24

Table 3. Experimental results (%) of the long-tailed and cold-start users are presented for the bi-directional Cloth-Sport CDSR scenario. The experiments are conducted five times, while we report the average values due to page limitations.

methods exhibit inferior performance compared with CDSR baselines since they do not consider the cross-domain information.

- Though C²DSR achieves remarkable success in most cases, the model cannot utilize auxiliary information to enhance the representation and address the semantic gap. Thus, compared to MACD, it shows an inferior performance.
- In most cases, incorporating auxiliary actions can enhance the performance of models on long-tailed/cold-start users, who often have sparse behavior, and auxiliary sequences can better explore their interests.
- Our proposed MACD consistently achieves significant performance improvements over SDSR, denoising SR, and CDSR baselines. Compared to SDSR and denoising SR baselines, we consider cross-domain information and learn the interest deviation. Additionally, compared to SDSR and CDSR baselines, we fully explore the auxiliary behavior information to better learn the representation for long-tailed and cold-start users.

Model Efficiency. All comparative models were trained and tested on the same machine, which had a single NVIDIA GeForce A10 with 22GB memory and an Intel Core i7-8700K CPU with 64G RAM. Furthermore, the number of parameters for typical SASRec+HSD, DASL, C²DSR, and SASRec+MACD (our approach) is within the same order of magnitude, ranging from 0.135M to 0.196M. The training/testing efficiencies of SASRec+HSD, DASL, C²DSR, and SASRec+MACD (our approach) when processing one batch of samples are 4.59 $\times 10^{-4}$ s/3.70 $\times 10^{-4}$ s, 3.48 $\times 10^{-4}$ s/2.59 $\times 10^{-4}$ s, 5.46 $\times 10^{-4}$ s/3.57 $\times 10^{-4}$ s, and 4.04 $\times 10^{-4}$ s/3.41 $\times 10^{-4}$ s, respectively. In summary, our MACD approach achieves superior performance enhancement in the open-world CDSR scenario while maintaining promising time efficiency.

3.4 Ablation Study (RQ2)

We conduct the following experiments on SASRec equipped with our MACD. To verify the contribution of each key component of MACD, we conduct an ablation study with $\mathcal{K}_u = 25\%$ by comparing it with several variants. Based on Table 4, we make the following observations: (i) Extracting implicit interest and explicit interest is critical to enhancing the representation for users, especially for long-tailed and cold-start users. When CDDM was removed, our model is unable to collect and transfer knowledge, which significantly hurt its performance. (ii) Without CL and FGU, the performance also dropped significantly, as the extracted user interest information from auxiliary behaviors may generate interest deviation and impair performance. (iii) Without IRG, the cold-start users cannot obtain a well-learned inductive representation, which impairs performance. As such, our model equipped with all main components achieves the best performance.

3.5 Online Evaluation (RQ3)

Except for these offline experiments, we conduct an online A/B test on a large-scale financial platform, which consist of multiple financial domains, such as pur-

Table 4. Experimental results (%) with different model variants. w/o denotes the model without the corresponding component variant. IDDM denotes the intra-domain denoising module, while CDDM denotes the cross-domain denoising module. CL is the contrastive information regularizer and FGU is the fusion gate unit. IRG denotes the inductive representation generator.

Scenarios	Metrics	Model	Model variants (w/o)						
		IDDM	CDDM	CL	FGU	IRG			
Cloth	NDCG	6.14	5.52	6.02	6.27	6.49	6.69		
	$_{\rm HR}$	12.47	11.88	12.54	12.69	12.93	13.16		
Sport	NDCG	9.82	9.59	10.03	10.60	10.93	11.28		
	$_{\rm HR}$	19.77	19.44	19.82	20.39	20.53	21.02		

chasing funds, mortgage loans, and discounting bills. Specifically, we select three popular domains - "Loan," "Fund," and "Account" - from the serving platform as targets for our online testing. For the control group, we adopt the current online solution for recommending themes to users, which is a cross-domain sequential recommendation method that utilizes noisy auxiliary behaviors directly. For the experiment group, we equip our method with a mature SDSR approach that has achieved remarkable success in the past. We evaluate the results based on three metrics: the number of users who have been exposed to the service, the number of users who have clicked inside the service, and the conversion rate of the service (denoted by # exposure, CTR, and CVR, respectively). All of the results are reported as the lift compared to the control group and presented in Table 5. In a fourteen-day online A/B test, our method improved the average exposure by 10.29%, the conversion rate by 6.28%, and the CVR by 1.45% in the three domains.

	# exposure	CTR	CVR
Loan Domain	+10.23%	+6.31%	+1.54%
Fund Domain	+8.51%	+5.49%	+1.03%
Account Domain	+12.13%	+7.03%	+1.77%

Table 5. Online A/B testing results from 9.1 to 9.14, 2023

3.6 Model Analyses (RQ4)

Discussion of the Behaviour Sparsity. To verify the superior performance of MACD in CDSR scenarios with varying data densities, we conducted further studies by varying the data density D_s in {25%, 50%, 75%, 100%}. As an

Scenarios	D_s	Ours		C^2DSR		
		NDCG	HR	NDCG	HR	
Cloth	25%	2.87	7.34	2.33	6.29	
	50%	3.42	9.69	2.94	8.98	
	75%	5.25	11.41	4.38	10.05	
	100%	6.69	13.16	6.10	12.62	
Sport	25%	3.56	8.67	2.84	8.01	
	50%	5.21	12.13	3.49	10.97	
	75%	8.34	17.54	7.57	16.80	
	100%	11.28	21.02	10.40	20.37	

Table 6. Experimental results (%) on dif-

ferent density dataset.

Table 7. Impact (%) of the number ofcold-start users.

Scenarios	K_{cs}	Ours		C^2DSR		
		NDCG	HR	NDCG	HR	
Cloth	5%	7.37	13.54	6.81	12.99	
	20%	6.69	13.16	6.10	12.62	
	35%	6.25	12.81	5.88	12.35	
	50%	5.87	12.26	5.20	11.62	
Sport	5%	11.56	21.67	10.84	21.01	
	20%	11.28	21.02	10.40	20.37	
	35%	10.74	20.54	10.17	19.80	
	50%	10.49	20.22	9.74	19.41	

example, in the "Cloth-Sport" task, $D_s = 50\%$ indicates that the data densities of the "Cloth" and "Sport" domains change from 0.023% to 0.012% (computed as 0.023% * 0.5 = 0.0115%) and from 0.020% to 0.010% (computed as 0.020% * 0.5 = 0.010%), respectively. We set \mathcal{K}_u to 25%. The experimental results of our model (SASRec+MACD) compared to the second-best baseline $(C^2DSR)^3$ are presented in Table 7. As expected, the performance of all models decreases with decreasing data density, as sparser data makes representation learning and knowledge transfer more challenging. Our method consistently outperforms C²DSR in all sparsity experimental settings, confirming the effectiveness of our approach in the open-world environment (Table 6).

Discussion of the Number of the Cold-Start Users. We also conducted experiments to investigate the effectiveness of our model in open-world environments with varying numbers of cold-start users. As mentioned in Sect. 3.1, we randomly selected partial overlapping users as cold-start users and varied the cold-start user ratio \mathcal{K}_{cs} among {5%,20%,35%,50%}. Due to the limited page, we show the results with the metrics NDCG@10 and HR@10. From the figures, we made the following observations: (1) With the rise in the ratio of cold-start users, the performance of all models' recommendations declined, highlighting the difficulty but significance of learning embeddings for cold-start users. (2) Our model demonstrated more robust performance in making recommendations for cold-start users than the strongest baseline, C²DSR. This is because our MACD generates inductive embeddings for the cold-start users and the auxiliary information is effectively utilized by our model.

3.7 Parameter Sensitivity (RQ5)

This section investigates the parameter sensitivity of the sequence length T and the harmonic factor λ .

 $^{^{3}}$ The C²DSR method utilizing auxiliary behaviors obtains the second-best results.



Fig. 3. (a)–(d) show the effect of length of sequence on model performance.



Fig. 4. (a)–(d) show the effect of harmonic factor λ on model performance.

For sequence length T, we show its "cloth" domain and "sport" domain results with overlapping ratio 25% and 75% in Fig. 3. After training our model with different settings $T = \{10, 15, 20, 25, 30\}$, one can see that our model achieves the best performance in terms of NDCG@10 and HR@10 when T = 20. When increasing T from 10 to 20, the performance is gained on account of richer historical interest information. If T is larger than 20, the performance will decrease. The reason might be that padding item causes the model ignoring important information from the true user-item interaction. Therefore, we choose T = 20 to better capture the user-item interaction information.

For harmonic factor λ , Fig. 4 shows its "cloth" domain and "sport" domain prediction performance with overlapping ratio 25% and 75% in terms of NDCG@10 and HR@10. We report the results under λ selected between 0.1 and 0.9 with a step length of 0.1. The curves shows that the accuracy will first gradually increase with λ raising and then slightly decrease. We can conclude that when λ approach 0, the contrastive information regularizer cannot produce positive effects. But when λ become too large, the contrastive loss will suppress the classification loss, which also reduces the recommendation accuracy. Empirically, we choose $\lambda = 0.4$ on the Cloth & Sport scenario with a $\mathcal{K}_u = 25\%$ while $\lambda = 0.7$ on the Cloth & Sport scenario with a $\mathcal{K}_u = 75\%$.

4 Related Work

Sequential Recommendation (SR) models [4,6,24,26,32] user preferences based on historical behavioral sequences, enabling the modeling of users'

dynamic interests compared to other conventional recommendation models. Various approaches have been proposed in the literature to address sequential recommendation problems. For instance, GRU4Rec [4] modifies classical RNN models to handle session-based recommendation problems. BERT4Rec [16] utilizes a Bidirectional Encoder Representation from Transformers to better capture the diversity of users' historical behavior. SASRec [6] employs self-attention mechanisms to balance the trade-off between capturing long-term semantics and addressing data scarcity issues. These models leverage different sequential models to more effectively capture the context of users' historical behavior. However, it should be noted that these models face limitations in scenarios with data sparsity and cold-start problems in open-world recommendation systems.

Cross-Domain Recommendation (CDR) [25,31,33], which leverages behavior patterns from multiple domains to jointly characterize user interests, has shown great potential in addressing data sparsity and cold-start issues in singledomain recommendation system. Recent CDR studies have focused on transfer learning [14,15,34], involving the design of a specific transfer module to learn a mapping function across domains and fuse pre-trained representations from each single domain. Furthermore, modeling domain-shared information has also drawn significant attention [2,9]. Although CDR approaches effectively incorporate rich information from relevant domains to improve performance on the target domain, conventional CDR methods still struggle to address the CDSR problems, which requires capturing sequential dependencies in users' interaction.

Cross-Domain Sequential Recommendation (CDSR) [11,12,17,27] aims to enhance sequential recommendation (SR) performance by leveraging user behavior sequences from multiple relevant domains. Some early studies [11,17] employ RNNs to capture sequential dependencies and generate user-specific representations. The attentive learning-based model DASL [8] uses dual attentive learning to transfer user preferences bidirectionally across domains. Moreover, C^2DSR designs sequential attentive encoders combined with contrastive learning to jointly learn inter- and intra-domain relationships. However, these methods heavily rely on data from overlapping users, which represent only a small proportion of the user pool in real-world scenarios. As a result, these methods exhibit poor performance in the open-world environment, since the insufficient representation of the long-tailed and cold-start users.

Multi-behaviour Recommendation methods has investigated diverse methods for learning collective knowledge from users' behaviors, including click, add to cart, and purchase [5,20,23]. Jin et al. employ graph convolutional networks to capture behavior-aware collaborative signals [5], while CML introduces a selfsupervised learning method for multi-behavior recommendation [20]. However, these methods often neglect the dynamism of multi-behavior relations and user interests, as they primarily focus on static recommendation scenarios. In recent studies, DPT proposes a three-stage denoising and prompt-tuning paradigm to mitigate the noise in auxiliary behavior data [29]. Nevertheless, existing multibehavior sequential recommendation techniques that concentrate on a single domain are insufficient for cross-domain scenarios, as they fail to effectively capture the divergent user preferences across domains and overlook the information transfer between domains.

5 Conclusion

In this work, we propose a model-agnostic contrastive denoising framework that can be integrated with most off-the-shelf SDSR methods. To enhance openworld CDSR performance by capturing comprehensive interest information, we integrate auxiliary behavior data to refine user embeddings, particularly for long-tailed and cold-start users. However, leveraging auxiliary behaviors without adjustment can lead to semantic discrepancies from target behaviors and inaccuracies in user interest learning. To mitigate this, we introduce a denoising interest-aware network with a contrastive information regularizer for precise latent interest extraction and cross-domain knowledge transfer. We also devise a parameter-free inductive representation generator to effectively identify analogous representations for cold-start users. Our model, rigorously evaluated on public datasets, demonstrates exceptional performance in open-world CDSR, a finding corroborated by an A/B test on a large-scale financial platform.

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MixerFlow: MLP-Mixer Meets Normalising Flows

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Abstract. Normalising flows are generative models that transform a complex density into a simpler density through the use of bijective transformations enabling both density estimation and data generation from a single model. In the context of image modelling, the predominant choice has been the Glow-based architecture, whereas alternative architectures remain largely unexplored in the research community. In this work, we propose a novel architecture called MixerFlow, based on the MLP-Mixer architecture, further unifying the generative and discriminative modelling architectures. MixerFlow offers an efficient mechanism for weight sharing for flow-based models. Our results demonstrate comparative or superior density estimation on image datasets and good scaling as the image resolution increases, making MixerFlow a simple vet powerful alternative to the Glow-based architectures. We also show that MixerFlow provides more informative embeddings than Glow-based architectures and can integrate many structured transformations such as splines or Kolmogorov-Arnold Networks.

Keywords: Density estimation \cdot Generative modelling \cdot MLP-Mixer

1 Introduction

Normalising flows [22,35], a class of hybrid statistical models, serve a dual purpose by functioning as both density estimators and generative models. They achieve this versatility through a series of invertible mappings, enabling efficient inference and generation from the same model. One of their distinctive features is explicit likelihood training and evaluation, distinguishing them from models relying on lower-bound approximations [17,21]. Furthermore, normalising flows offer computational efficiency for both inference and sample generation, setting them apart from Autoregressive models like PixelCNNs [33].

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The invertible nature of normalising flows extends their utility to various domains, including solving inverse problems [37] and enabling significant memory savings during the backward pass, where activations can be efficiently computed through the inverse operations of each layer. Additionally, normalising flows can be trained simultaneously on a supervised prediction task and the unsupervised density modelling task to function as generative classifiers or in the context of semi-supervised learning [32], setting them apart from other generative models such as GANs [12] and Diffusion models [17].

Despite their wide-ranging applications, normalising flows lack expressivity. The Glow-based architecture [20] has become the standard for implementing normalising flows due to its clever design, often requiring an excessive number of parameters, especially for high-dimensional inputs. Existing literature primarily focuses on enhancing expressiveness, employing strategies such as coupling layers with spline-based transformations [9], kernelised layers [10], log-CDF layers [16], or introducing auxiliary layers like the Butterfly layer [30] or 1x1 convolution [20].

Nevertheless, alternative architectures remain relatively unexplored, with ResFlows [5] being a notable exception. In this work, we introduce MixerFlow, drawing inspiration from the MLP-Mixer [40], a well-established discriminative modelling architecture. Our results demonstrate that MixerFlow consistently performs well, matching or surpassing the negative log-likelihood of the widely adopted Glow-based baselines. MixerFlow excels in scenarios involving uncorrelated neighbouring pixels or images with permutations and scales well with an increase in image resolution, outperforming the Glow-based baselines. Furthermore, our experiments suggest that MixerFlow learns more informative representations than the baselines when training hybrid flow models and can easily integrate other transformations such as Spline [9] or Kolmogorov-Arnold Networks(KAN) layer [26] with increased expressiveness.

2 Related Works

Our work is closely related to the MLP-Mixer architecture [40]. However, the MLP-Mixer is designed for discriminative tasks and lacks inherent invertibility, a critical requirement for modelling flow-based architectures.

The field of flow models is quite extensive, including well-known models such as Glow [20], Neural Spline Flows [9] integrating splines into coupling layer, Generative Flows with Invertible Attention [39] replacing convolutions with attention, Butterfly Flow [30], augments a coupling layer with a butterfly matrix, and Ferumal Flows [10], kernelises a coupling layer. These models often share a foundational Glow-like architecture, with specific component modifications aimed at improving inter-data communication and mixing for an improved expressivity in normalising flows in different ways. MixerFlow provides an alternative architecture to Glow [20] with the possibility to add the specific components from these works to improve inter-data communication. Another noteworthy approach is the Residual-Flow-based framework [5], which encompasses Monotone Flows [2] and ResFlows. In contrast to Glow-based architectures, ResFlow ensures invertibility through fixed-point iteration [4]. These two categories, Glow-based and ResFlow-based, exhibit distinct characteristics and can be considered the primary classes of flow-based architectures for image modelling.

In addition to these, there exist other flow methods such as Gaussianisation Flows [29] within the Iterative Gaussianisation [24] family and FFJORD [13] in the family of continuous-time normalising flow. These methods represent different methodological approaches to flow-based modelling, adding richness to the landscape of techniques available for density estimation.

Our proposed architecture leverages the inherent weight-sharing properties of the MLP-Mixer. This design choice allows for flexibility in integrating either a coupling layer, a Lipschitz-constrained layer [4] commonly found in ResFlow-like architectures, or even an FFJORD layer. This ensures that our model can leverage any flow method providing versatility across various application scenarios.

Our attempt to further unify generative and discriminative architectures finds resonance in similar attempts within the research community. For instance, Vit-GAN [25] adapts the VisionTransformer [8] architecture for generative modelling, demonstrating the adaptability of existing architectures. Another noteworthy example is by [32], which leverages generative architecture for discriminative tasks in a hybrid context.

3 Preleminaries

In this section, we briefly introduce the major components of a normalising flow and the MLP-Mixer architecture.

The change of variables theorem states that if p_X is a continuous probability distribution on \mathbb{R}^d , and $f : \mathbb{R}^d \to \mathbb{R}^d$ is an invertible and continuously differentiable function with z := f(x), then $p_X(x)$ can be computed as

$$p_X(x) = p_Z(z) \left| \det \left(\frac{\partial f(x)}{\partial x} \right) \right|.$$

If we model p_Z with a simple parametric distribution (such as the standard normal distribution), then this equation offers an elegant approach to determining the complex density $p_X(x)$, subject to two practical constraints. First, the function f must be bijective. Second, the Jacobian determinant should be readily tractable and computable.

Non-linear Coupling Layers: Coupling layers represent an essential component within the framework of most flow-based models. These non-linear layers serve a pivotal role by enabling efficient inversion for normalising flows whilst maintaining a tractable determinant Jacobian. Various forms of coupling layers exist, including additive and affine, amongst others. Below, we briefly define the affine coupling layers

Affine coupling layers involve splitting the input x into two distinct components: x_a and x_b . Whilst the first component (x_a) undergoes an identity transformation, the second component (x_b) undergoes an affine transformation characterised by parameters S and T. These non-linear parameters are acquired through learning from x_a using a function approximator, such as neural networks. The final output of an affine coupling layer is the concatenation of these two transformations. The equations below summarise the operations.

$$X^{d}, X^{D-d} = \operatorname{split}(X)$$
$$S, T = F(X_{D-d})$$
$$Y_{D-d} = X_{D-d}$$
$$Y_{d} = S \odot X_{d} + T$$

Given that only one partition undergoes a non-trivial transformation within a coupling layer, the choice of partitioning scheme becomes crucial. It has been shown that incorporating invertible linear layers can aid in learning an enhanced partitioning scheme [20, 30].

MLP-Mixer: The MLP-Mixer [40] is an architecture tailored for discriminative vision tasks, relying exclusively on multi-layer perceptrons (MLPs) for its operations. It distinctively separates per-location and cross-location operations, which are fundamental in deep vision architectures and often co-learnt in models like Vision Transformers (ViT) [8] and Convolutional neural networks [34] with larger kernels.

The central concept of this architecture begins with the initial partitioning of an input image into non-overlapping patches, denoted as n_p . If each patch has a resolution of (p_h, p_w) , and the image itself has dimensions (h, w), then the number of patches is calculated as $n_p = (h * w)/(p_h * p_w)$. Each patch undergoes the same linear transformation, projecting them into lower dimensional fixedsize vectors represented as c. These transformed patches collectively form a new representation of the input image in the form of a matrix, we refer to it as the "mixer-matrix," with dimensions $n_p \times c$. Here, c corresponds to the dimensionality of the projected patch, often referred to as "channels" in MLP-Mixer literature, and n_p represents the total number of patches, typically called "tokens."

The core innovation of the MLP-Mixer unfolds through the application of multi-layer perceptrons (MLPs), which are applied twice in each mixer layer. The first MLP, known as the token-mixing MLP, operates on columns of the mixer-matrix. The same MLP is applied to all columns of the mixer-matrix. The second MLP referred to as the channel-mixing MLP, is applied to the rows of the mixer-matrix, again with the same MLP applied across all columns of the mixer-matrix. This design choice ensures weight sharing within the architecture. The mixer layer is repeatedly applied for several iterations, facilitating complete interactions between all dimensions within the image matrix, a process apply referred to as "mixing."

In addition to these operations, two critical components are integral to the MLP-Mixer architecture. Firstly, layer normalisation [3] is employed to stabilise

the network's training dynamics. Secondly, skip connections [14] are introduced after each mixing layer to facilitate the flow of information and enable smoother gradient propagation throughout the network. An important property of the MLP-Mixer architecture is that the hidden widths of token-mixing and channelmixing MLPs are independent of the number of input patches and the patch size, respectively, making the computational complexity of the model linear in the number of patches, in contrast to ViT [8], where it is quadratic.

4 MixerFlow Architecture and Its Components

In our architectural design, we first apply a 1×1 convolution to the RGB channels of the input image with a resolution of (h, w). This results in a transformed representation of the image in which the RGB channels are no longer distinguishable. Subsequently, we partition this transformed view into non-overlapping patches (or stripes, bands, dilated patches), denoted as n_p , each with a resolution of (p_h, p_w) . The choice of patch resolution is made to achieve the desired granularity, ensuring that $n_p = (h * w)/(p_h * p_w)$. These small patches are then flattened into vectors of size $c = p_h * p_w * 3$, yielding the mixer-matrix of dimensions $n_p \times c$.

Next, we introduce two distinct types of normalising flows: channel-mixing flows and patch-mixing flows, resembling operations similar to channel-mixing MLPs and token-mixing MLPs respectively. Channel-mixing flows facilitate interactions between different channels by processing individual rows of the mixer-matrix, operating on each patch independently. Conversely, patch-mixing flows focus on interactions between different patches, processing individual columns of the mixer-matrix, whilst operating on each channel separately. These two flow operations are executed iteratively in an alternating fashion, enabling interactions between all elements within the mixer-matrix.

In summary, we apply the same channel flow to all rows of the mixer-matrix and the same patch flow to all columns of the mixer-matrix. This configuration ensures the desired parameter sharing across the model. These flows, both channel-flows and patch-flows, are seamlessly integrated into our architecture, with each comprising a series of subsequent components (see Fig. 1 for an illustration).

Shift Layer: In the MLP-Mixer, the patch definition remains static across the entire architecture, albeit projected into a lower dimension c at the outset-an operation constrained by invertibility concerns. Whilst this approach suits the MLP-Mixer, it introduces grid-like artefacts when applied to sampled images in our MixerFlow model.

To address this issue, shift layers reshape the mixer-matrix back into its original image dimensions. It then creates a frame, leaving the inner part as $(s_h : h - p_h + s_h, s_w : w - p_w + s_w)$, where (p_h, p_w) denotes the patch resolution, (s_h, s_w) denotes the shifting unit, and (h, w) corresponds to the image resolution. Subsequently, we re-transform the inner part into an input resembling a mixer-matrix, which we refer to as the "shifted-mixer-matrix." This process effectively shifts the patch extraction by (s_h, s_w) units both vertically and horizontally



Fig. 1. MixerFlow architecture

respectively compared to the old patch definition. This frame carving reduces the number of patches as some input variables are omitted by extraction of the frame. A full mixer layer is applied to the shifted-mixer-matrix, and after this layer, the carved-out frame is reintroduced for the next stage. This strategy ensures robust interactions near the boundaries.

Importantly, the alteration in patch definition has no adverse impact on performance, as neighbouring pixels exhibit a high correlation. Moreover, it facilitates the distribution of transformations for the variables within the carved-out frame, with a focus on the non-carved-out-frame layers. In preliminary experiments we found these shift layers to result in improvements both qualitatively and quantitatively.

Linear Block: As previously mentioned, a coupling layer operates exclusively on approximately half of the input dimensions, underscoring the importance of partition selection. RealNVP [7] introduced alternating patterns, which, in certain cases, introduce an order bias. In contrast, Glow [20] advocated for 1×1 convolutions in the form of PLU factorisation, with fixed permutation matrix P and optimisable lower and upper triangular matrices L and U, providing a more generalised approach to permutations. In our approach, we employ either LU factorisation or RLU, where R is a permutation matrix that reverses the order of dimensions. It is crucial to emphasise the necessity of a linear block that effectively reverses the shuffling before the application of a shift layer which aims to capture the interaction between the patch boundaries. Subsequently, the linear block is followed by a Flow coupling layer.

Flow Layer: After each Linear Block, we incorporate a flow layer into our architecture. Specifically, we employ a standard affine coupling layer as our chosen

flow layer, applicable to either a channel flow or a patch flow. Within each flow layer, we use a residual block as the function approximator. In our experiments, we mostly chose a latent dimension of 128 for both the channel-flow-residual-block and the patch-flow-residual-block, along with the GELU [15] activation function and batch normalisation [19] in the layers of the residual block.

ActNorm Layer: Due to the computation of a full-form Jacobian when applying layer normalisation [3], we opt for ActNorm [20] as the preferred normalisation technique in our architecture. ActNorm layers are data-dependent initialised layers with an affine transformation that initialises activations to have a mean of zero and a variance of one based on the first batch. In contrast to Glow-based architectures where ActNorm layers are applied only to the channels, we apply ActNorm to all activations after each flow layer and just before the initial linear layer following each transpose operation (i.e., applying a flow layer to columns of the matrix after applying it to the rows or vice versa).

Identity Initialisation: We initialised all linear blocks to perform an identity function. Additionally, we initialise the final layer of each residual block within the flow with zeros to achieve an identity transformation. This approach, as reported by [20], has been observed to be beneficial for training deeper flow networks.

5 Experiments

We conducted an extensive series of experiments encompassing various datasets, varying dataset sizes, and diverse applications. These applications include permutations, classification tasks, and the integration of Masked Autoregressive Flows [36] into our MixerFlow model. We use thirty MixerFlow layers with a shift of either one or two every fourth layer in our experiments. We perform uniform dequantisation, use a patch size of four for 32×32 resolution and a patch size of eight for 64×64 resolution and use Adam for optimisation with the default parameters. For baseline experiments, we used the experimental settings as in [27], reproducing results on standard datasets. Following standard practices in flow literature, we reported density estimation results in "bits-per-dimension" (BPD) and reported results up to two decimal places, which led to the exclusion of standard error intervals as they were consistently less than 0.004.

5.1 Density Estimation on 32×32 Datasets

Datasets: In line with previous research, we assessed the performance of Mixer-Flow on standard 32×32 datasets, specifically ImageNet32 [38,41]¹ and CIFAR-10. [23].

¹ Two versions of downscaled ImageNet exist. The one used to evaluate normalising flow models has been removed from the official website but remains accessible through alternative sources, such as Academic Torrents.



Fig. 2. Sampled images from our MixerFlow

Baselines: We use various Glow-based baselines for our evaluations: Glow [20], Neural Spline flows [9], MaCow [28], ME-Woodbury [27], and Emerging and Periodic convolutions [18].

Results: Table 1 presents our quantitative results, demonstrating the competitive or superior performance of MixerFlow to all of the aforementioned baselines. The sample images from our MixerFlow trained on CIFAR-10 can be seen in Fig. 2(a).

Method	CIFAR-10	ImageNet32	Params
MixerFlow	3.46	4.20	$11.34 \mathrm{M}$
Glow	3.51	4.32	$11.02 \mathrm{M}$
Neural Spline	3.50	4.24	10.91M
MaCow	3.48	4.34	11.43M
Woodbury	3.48	4.22	$11.02 \mathrm{M}$
Emerging	3.48	4.26	11.43M
Periodic	3.49	4.28	11.21M

Table 1. Negative log-likelihood (in bits per dimension) for 32×32 datasets. Smaller values are better.

5.2 Density Estimation on 64×64 Datasets

Datasets: We assessed the performance of MixerFlow on two distinct datasets: ImageNet64 [38], a standard vision dataset, and AnimeFace [31], a collection of Anime faces.

Baselines: For our evaluations, we use a couple of Glow-based baselines, specifically Glow [20], and Neural Spline [9].

Results: Our quantitative results in Table 2 illustrate that MixerFlow outperforms the selected baselines in terms of negative log-likelihood measured in bits per dimension. Notably, our analysis of model sizes reveals that MixerFlow scales remarkably well as image size increases, requiring approximately half the number of parameters compared to the other baselines. This outcome aligns with our expectations, as the hidden patch-flow-MLP dimension remains independent of the number of patches, and the hidden-channel-flow-MLP dimension remains independent of the number of channels-a characteristic inherited from the MLP-Mixer architecture. For a visual representation of our results, please refer to Fig. 2(b), which displays sample images generated by our MixerFlow model trained on the AnimeFace dataset.

Table 2. Negative log-likelihood (in bits per dimension) for 64×64 datasets. Smaller values are better.

Method	AnimeFaces	ImageNet64	Params
Glow	3.21	3.94	$37.04 \mathrm{M}$
Neural Spline	3.23	3.95	$38.31\mathrm{M}$
MixerFlow	3.17	3.92	18.90M

In the context of the AnimeFaces dataset, we also observed a qualitative improvement. Specifically, we noted a reduction in artefacts compared to the Glow-based baseline (Figure 1 and Figure 2 in the Appendix).

5.3 Enhancing MAF with the MixerFlow

Masked Autoregressive Flows (MAF) [36] represent one of the most popular density estimators for tabular data. They are a generalisation of coupling layer flows, such as Glow and RealNVP. Notably, MAF tends to outperform coupling layer flows on tabular datasets, although it comes at the cost of relatively slow generation. The concept of MAF emerged as an approach to enhance the flexibility of the autoregressive model, MADE [11], by stacking their modules together. This innovation, which enables density evaluations without the typical sequential loop inherent to autoregressive models, significantly accelerated the training process and enabled parallelisation on GPUs.

However, MAF is vulnerable to the curse of dimensionality, necessitating an enormous number of parameters to achieve scalability for image modelling. This requirement for many additional layers can render MAF impractical for certain tasks, i.e. image modelling. In our work, we demonstrate that by substituting a flow step with an MAF layer within our MixerFlow model, we can enable MAF for density estimation tasks involving image datasets. This integration leverages the effective weight-sharing architecture inherent in MixerFlow and substantially enhances the performance of the MAF model.



Fig. 3. Sampled Images for our MAF-based MixerFlow on MNIST

Datasets: We assessed the performance of the MAF integration on two distinct datasets: MNIST [6] and CIFAR-10 [23].

Baselines: We considered MAF [36] and MADE [11] as the baselines for our evaluation as they are pre-cursors of MAF's integration into our architecture.

Results: Our findings, as presented in Table 3, showcase the density estimation results with MixerFlow further enhancing the use of the MADE module through MAF's integration in our architecture for MNIST [6] and CIFAR-10 [23]. Furthermore, Fig. 3 provides a visual representation of the generated samples from our enhanced MAF model on the MNIST dataset

Table 3. Negative log-likelihood (in bits per dimension) for MAF's integration into MixerFlow. Smaller values are better.

Method	MNIST	CIFAR-10
MADE	2.04	5.67
MAF	1.89	4.31
Ours	1.22	3.44

5.4 Datasets with Specific Permutations

Whilst Glow-based models exhibit expressive capabilities in capturing image dynamics, they heavily rely on convolution operations involving neighbouring pixels to transform the distribution. In contrast, our MixerFlows use multi-layer perceptrons as function approximators, making them invariant to changes in pixel locations within patches and patch locations in the image. This might be helpful if there is some data corruption that induces permutation as Glow-based



Fig. 4. (a) For Glow. (b) For MixerFlow

architectures will result in poor density estimation in such cases. In this section, we empirically demonstrate this advantage across various datasets.

Density Estimation on Permuted Image Datasets

Dataset: Our experimentation involved two types of artificial permutations. Firstly, we divided the input image into patches and performed permutations on both the order of these patches and the pixels within each patch, using a shared permutation matrix. Subsequently, we reorganised these shuffled patches in the image form, referring to this process as "local shuffling." This setup is analogous to the pixel shuffling experiment introduced by [40], but adapted to the case of density estimation.

Global shuffling	CIFAR-10	Imagenet32	Params
Glow	5.18	5.27	44.24M
Neural Spline	5.01	5.32	$89.9 \mathrm{M}$
MixerFlow	4.09	4.91	11.43M
Butterfly	5.11	6.18	-

Table 4. Negative log-likelihood (in bits per dimension) for the global shuffling exper-iment. Smaller values are better.

Secondly, we applied a "global shuffling", where we shuffled all the pixels of the entire image using the same permutations across all images. We created these modified versions of the CIFAR-10 [23] and ImageNet32 [38,41] datasets, both of which are 32×32 in size.

Baselines: To assess the performance of our model, we compared it with Glow [20] and Neural Spline flow [9]. Additionally, we included a comparison with Butterfly Flows (results taken from existing literature) [30], which are theoretically guaranteed to be able to represent any permutation matrix. All the baselines use

MixerFlow	3.49	4.24	11.43M
Neural Spline	3.99	4.54	$89.9 \mathrm{M}$
Glow	4.06	4.49	$44.24 \mathrm{M}$
Local shuffling	CIFAR-10	Imagenet32	Params

Table 5. Negative log-likelihood (in bits per dimension) for the local shuffling experi-ment. Smaller values are better.

the same Glow backbone. Notably, we evaluated larger models for the baselines in this experiment, highlighting that even with larger models, their performance lags in the presence of permutations.

Results: We tested our hypothesis that MixerFlow is more effective when dealing with permutations in the dataset. The performance gap is substantial for both types of permutations applied to the dataset, as evident from Table 5 and Table 4, where we report lower negative log-likelihoods in bits per dimension (BPD). Figure 4(a) and Fig. 4(b) illustrate the training curve for both types of permutations on CIFAR-10 [23]. Notably, stronger inductive biases in Glow-based architectures are highly dependent on the order of pixels, resulting in a significant performance drop compared to the original order of image pixels. Conversely, the performance drop is relatively minimal for MixerFlow. Whilst it was intuitively expected that there would be no performance difference on local shuffling datasets, there is a slight difference in the negative log-likelihood, this is due to our shift layer, which requires that neighbouring patches maintain the same structural positions as in the original image.

Density Estimation on a Structured Dataset: Galaxy32

Dataset: Real-world datasets often exhibit unique structures, such as permutations or periodicity. To enhance the robustness of our experiments, we applied MixerFlow to a real-world dataset-the Galaxy dataset. This dataset, curated by [1], comprises images of merging and non-merging galaxies. Given that stars can appear anywhere in these images, they exhibit permutation invariance. Additionally, the dataset demonstrates periodicity, as it represents snapshots of a spatial continuum rather than individual isolated images. We downsampled the dataset to a resolution of 32×32 for our experiments.

Baselines: In our evaluation, we compared the performance of MixerFlow against two baseline models: Glow [20] and Neural Spline [9].

Results: The results, presented in Table 6, clearly indicate that MixerFlow outperformed the chosen baselines. Figure 2(c) showcases samples generated by the MixerFlow model.

Method	Galaxy32
Glow	2.27
Neural Spline	2.25
MixerFlow	2.22

Table 6. Negative log-likelihood (in bits per dimension) for the Galaxy32 dataset.Smaller values are better.

5.5 Hybrid Modelling

[32] introduce a hybrid model that combines a normalising flow model with a linear classification head. This hybrid approach offers a compelling advantage for predictive tasks, as it allows for the computation of both p(data) and p(label|data) using a single network. This capability enables semi-supervised learning and out-of-distribution detection. However, achieving optimal results often requires joint objective optimisation or separate training for different components, as the objective function not only maximises log-likelihood but also minimises predictive error. Significantly over-weighting the predictive error term is sometimes necessary for achieving superior discriminative performance.

Since our MixerFlow is based on a discriminative architecture, namely MLP-Mixer, we posit that our model can perform better under similar predictive loss weighting.

Dataset: We used the CIFAR-10 [23] dataset for the downstream task of classification.

Baselines: We evaluated the performance of MixerFlow embeddings against two baseline models' embeddings: Glow [20] and Neural Spline [9].

Results: In our experiments, we used our pre-trained models and added a classifier head to them. During training, we kept the flow parameters fixed whilst training the additional linear layer parameters. This setup effectively leveraged the representations learned by the flow models for downstream tasks. Our results in Table 7 indicate that the MixerFlow model exhibited lower losses compared to the chosen baselines. This suggests that MixerFlow embeddings capture more informative representations.

5.6 Integration of Powerful Architecture

The flexible design of our MixerFlow architecture enables the seamless integration of diverse flow layers and powerful transformations. To assess the impact of these transformations on model performance, we conducted experiments while maintaining a consistent architecture depth of 30 layers and training all models for 50,000 steps.

Method	Loss	Accuracy
Glow	1.69	41.23%
Neural Spline	1.67	41.27%
MixerFlow	1.54	45.11%

 Table 7. Classification loss and accuracy on CIFAR-10 when employing flow-based embeddings

Datasets: We evaluated the performance of different transformations on the MNIST dataset [6].

Models: We replaced the standard coupling-based MLPs in our MixerFlow model with alternative transformations, including Spline-based transformations [9], MAF-based transformation [36], KAN-layer [26] (using KANs instead of MLPs in standard coupling layers)

Results: As shown in Table 8, incorporating these advanced transformations within MixerFlow consistently yields improved density estimation performance compared to the baseline MLP model. This highlights the potential of leveraging specialised transformations further to enhance the expressive power of our proposed architecture. It is important to note that we used default parameter settings directly from their respective source papers due to computational constraints. We believe that fine-tuning these parameters could lead to even more significant improvements in performance.

Table 8. Negative log-likelihood (in bits per dimension) for different transformationsin MixerFlow. Smaller values are better.

Method	BPD	Params
MLP-coupling	1.57	3.10M
KAN-coupling	1.48	$7.45\mathrm{M}$
Splines	1.21	$5.28 \mathrm{M}$
MAF-based	1.56	3.83M

6 Conclusion and Limitations

In this work, we have introduced MixerFlow, a novel flow architecture that draws inspiration from the MLP-Mixer architecture [40] designed for discriminative vision tasks. Our experimental results have demonstrated that MixerFlow consistently outperforms or competes comparatively (in terms of negative loglikelihood) with existing models on standard datasets. Importantly, it exhibits good scalability, making it suitable for handling larger image sizes. Additionally, the integration of MAF layers [36] into our architecture showed considerable improvements in MAF, showcasing its adaptability and versatility for integrating normalising flow architectures beyond coupling layers.

We also explored the application of MixerFlow to datasets featuring artificial permutations and structured permutations, underlining its practicality and wide-ranging utility. To conclude, our work has highlighted the potential of the acquired representations in the downstream classification task on CIFAR-10, as evidenced by lower loss values. MixerFlow can also help existing MLP-Mixer architectures by providing a probabilistic approach to it.

One limitation of our proposed architecture is the absence of strong inductive biases typically associated with convolution-based flows. This may be especially relevant if little training data is available.

7 Future Work and Broader Impact

Whilst our experimental analysis focuses on MixerFlow with MLP-based neural networks, it is crucial to note that MixerFlow's architecture is not restricted to specific neural network types or flow networks. It enables parameter sharing and can be adapted to various network architectures. This adaptability extends to the incorporation of residual flows [5], attention layers [39], and convolutional layers, especially for larger image sizes and patch sizes, offering the potential for enhanced inductive biases and parameter sharing. Additionally, the inclusion of glow-like layers before applying the MixerFlow layers could further strengthen inductive biases.

Another promising avenue for improvement is enabling multiscale design [20], a well-established technique for boosting the performance of flow-based models. We leave the integration of multi-scale architecture, and stronger inductive biases for future work, believing it could further enhance the MixerFlow architecture's capabilities. We are optimistic about the potential of MixerFlow to advance flow-based image modelling, and we hope for more developments in architectural backbone research for flow models in the future.

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Handling Delayed Feedback in Distributed Online Optimization: A Projection-Free Approach

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Abstract. Learning at the edges has become increasingly important as large quantities of data are continuously generated locally. Among others, this paradigm requires algorithms that are *simple* (so that they can be executed by local devices), *robust* (against uncertainty as data are continually generated), and *reliable* in a distributed manner under network issues, especially delays. In this study, we investigate the problem of online convex optimization (OCO) under adversarial delayed feedback. We propose two projection-free algorithms for centralized and distributed settings in which they are carefully designed to achieve a regret bound of $O(\sqrt{B})$ where B is the sum of delay, which is optimal for the OCO problem in the delay setting while still being projection-free. We provide an extensive theoretical study and experimentally validate the performance of our algorithms by comparing them with existing ones on real-world problems.

Keywords: Online Optimization \cdot Distributed Learning \cdot Delayed Feedback \cdot Projection-Free

1 Introduction

Many machine learning (ML) applications owe their success to factors such as efficient optimization methods, effective system design, robust computation, and the availability of enormous amounts of data. In a typical situation, ML models are trained in an offline and centralized manner. However, in real-life scenarios, significant portions of data are continuously generated locally at the user level. Learning at the edge naturally emerges as a new paradigm to address such issues. In this new paradigm, the development of suitable learning techniques has become a crucial research objective. Responding to the requirements (of this new paradigm), online learning has been intensively studied in recent years. Its

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efficient use of computational resources, adaptability to changing environments, scalability, and robustness against uncertainty show promise as an effective approach for edge devices.

However, online learning/online convex optimization (OCO) problems typically assume that the feedback is immediately received after a decision is made, which is too restrictive in many real-world scenarios. For example, a common problem in online advertising is the delay that occurs between clicking on an ad and taking subsequent action, such as buying or selling a product. In distributed systems, the previous assumption is clearly a real issue. Wireless sensor/mobile networks that exchange information sequentially may experience delays in feedback due to several problems: connectivity reliability, varying processing/computation times, heterogeneous data and infrastructures, and unaware-random events. This can lead to difficulties in maintaining coordination and efficient data exchange, eventually affecting network performance and responsiveness. Given these scenarios, the straightforward application of traditional OCO algorithms often results in inefficient resource utilization because one must wait for feedback before starting another round. To address this need, this paper focuses on developing algorithms that can adapt to adversarial delayed feedback in both centralized and distributed settings.

Model. We first describe the delay model in a centralized setting. Given a convex set $\mathcal{K} \subseteq \mathbb{R}^d$, at every time step t, the decision maker/agent chooses a decision $x_t \in \mathcal{K}$ and suffers from a loss function $f_t : \mathcal{K} \to \mathbb{R}$. We denote by $d_t \geq 1$ an arbitrary delay value of time t. In contrast to the classical OCO problem, the feedback of iteration t is revealed at time $t + d_t - 1$. The agent does not know d_t in advance and is only aware of the feedback of iteration t at time $t + d_t - 1$. Consequently, at time t, the agent receives feedback from the previous iterations $s \in \mathcal{F}_t$, where $\mathcal{F}_t = \{s : s + d_s - 1 = t\}$. In other words, \mathcal{F}_t is the set of moments before time t such that the corresponding feedbacks are released at time t. Moreover, the corresponding feedbacks are not necessarily released in the order of their iterations. The goal is to minimize regret, which is defined as:

$$\mathcal{R}_T := \sum_{t=1}^T f_t(\boldsymbol{x}_t) - \min_{\boldsymbol{x} \in \mathcal{K}} \sum_{t=1}^T f_t(\boldsymbol{x})$$

In a distributed setting, we have additionally a set of agents connected over a network, represented by a graph $\mathcal{G} = (V, E)$ where n = |V| is the number of agents. Each agent $i \in V$ can communicate with (and only with) its immediate neighbors, that is, adjacent agents in \mathcal{G} . At each time $t \geq 1$, agent i takes a decision $\mathbf{x}_t^i \in \mathcal{K}$ and suffers a partial loss function $f_t^i : \mathcal{K} \to \mathbb{R}$, which is revealed adversarially and locally to the agent at time $(t + d_t^i - 1)$ —again, that is unknown to the agent. Similarly, denote $\mathcal{F}_t^i = \{s : s + d_s^i - 1 = t\}$ as the set of feedbacks revealed to agent i at time t where d_s^i is the delay of iteration s to agent i. Although the limitation in communication and information, the agent i time t, the loss of agent i for chosen \mathbf{x}_t^i is $F_t(\mathbf{x}_t^i)$. Note that each agent i does not

know F_t but has only knowledge of f_t^i —its observed cost function. The objective here is to minimize regret for all agents (Fig. 1):

$$\mathcal{R}_T := \max_i \left(\sum_{t=1}^T F_t(oldsymbol{x}_t^i) - \min_{oldsymbol{x} \in \mathcal{K}} \sum_{t=1}^T F_t(oldsymbol{x})
ight)$$

1.1 Our Contribution

The challenge in designing robust and efficient algorithms for these problems is to address the following issues simultaneously:

- Uncertainty (online setting, agents observe their loss functions only after selecting their decisions).
- Asynchronous (distributed setting with different delayed feedback between agents)
- Partial information (distributed setting, agents know only its local loss functions while attempting to minimize the cumulative loss).
- Low computation/communication resources of agents (so it is desirable that each agent performs a small number of gradient computations and communications).



Fig. 1. Illustration of delayed feedback in distributed system. Given a time t, each agent holds a distinct pool of available gradient feedback from s < t that is ready for computation at the current time. The pool can also be empty if no feedback is provided.

We introduce performance-guaranteed algorithms in solving the centralized and distributed constraint online convex optimization problem with adversarial delayed feedback. Our algorithms achieve an *optimal* regret bound for centralized and distributed settings. Specifically, we obtain the regret bound of $O(\sqrt{B})$ where B is the total delay in the centralized setting and B is the average total delay over all agents in the distributed setting. Note that, if d is a maximum delay of each feedback then our regret bound becomes $O(\sqrt{dT})$. This result recovers the regret bound of $O(\sqrt{T})$ in the classic setting without delay (i.e., d = 1). Additionally, the algorithms can be made projection-free by selecting appropriate oracles, allowing them to be implemented in different contexts based on the computational capacity of local devices. Finally, we illustrate the practical potential of our algorithms and provide a thorough analysis of their performance which is predictably explained by our theoretical results. The experiments demonstrate that our proposed algorithms outperform existing solutions in both synthetic and real-world datasets (Table 1).

Algorithm	Centralized	Distributed	Adversarial Delay	Projection-free	Regret
DGD	Yes	-	Yes	-	$O(\sqrt{dT})$
DOFW	Yes	-	Yes	Yes	$O(T^{3/4} + dT^{1/4})$
DeLMFW	Yes	-	Yes	Yes	$O(\sqrt{dT})$
De2MFW	-	Yes	Yes	Yes	$O(\sqrt{dT})$

Table 1. Comparisons to previous algorithms DGD [12] and DOFW [14] on centralized online convex optimization with delays bounded by d. Our algorithms are in bold.

1.2 Related Work

Online Optimization with Delayed Feedback. Over the years, studies on online optimization with delayed feedback have undergone a swift evolution. [17] shed light on the field by focusing on the convergence properties of online stochastic gradient descent with delays. They provide a regret bound of $O(\sqrt{dT})$ with d the delay value if $d^2 \leq T$. Later on, [12] proposes a centralized (single-agent) gradient descent algorithm under adversarial delays. The theoretical analysis of [12] entails a regret bound of $O(\sqrt{B})$, where B is the total delay. This bound becomes $O(\sqrt{dT})$ if d is the upper bound of delays. [8] provided a black-box style method to learn under delayed feedback. They showed that for any non-delayed online algorithms, the additional regret in the presence of delayed feedback depends on its prediction drifts. [2] developed an online saddle point algorithm for convex optimization with feedback delays. They achieved a sublinear regret $O(\sqrt{dT})$ where d is a fixed constant delay value. Recently, [14] proposed a first Frank-Wolfe-type online algorithm with delayed feedback. They modified the Online Frank-Wolfe (OFW) for the unknown delays setting and provided a regret bound of $O(T^{3/4} + dT^{1/4})$. This is the current state of the art for projection-free (Frank-Wolfe-type) algorithms with delays. Our bound of $O(\sqrt{dT})$ improves over the aforementioned results.

Distributed Online Optimization. [15] introduced decentralized online projected subgradient descent and showed vanishing regret for convex and strongly convex functions. In contrast, [7] extended distributed dual averaging technique to the online setting, using a general regularized projection for both unconstrained and constrained optimization. A distributed variant of online conditional gradient [5] was designed and analyzed in [16] that requires linear minimizers and uses exact gradients. Computing exact gradients may be prohibitively expensive for moderately sized data and intractable when a closed form does not exist. [13] proposes a decentralized online algorithm for convex function using stochastic gradient estimate and multiple optimization oracles. This work achieves the optimal regret bound of $O(T^{1/2})$ and requires multiple gradient evaluation and communication rounds. Later on, [11] provide a decentralized algorithm that uses stochastic gradient estimate and reduces communication by using only one gradient evaluation. [10] provides a comprehensible survey on recent development of distributed OCO. More recent work on distributed online optimization with feedback delays is proposed in [1]. The authors consider a distributed pro-
jected gradient descent algorithm where each agent has a fixed known amount of delay d_i . They provide a regret bound of $O(\sqrt{dT})$ where $d = \max_i d_i$ but the delays d_i must be fixed (non-adversarial).

Despite the growing number of studies on decentralized online learning in recent years, there is a lack of research that accounts for the *adversarial/online* delayed feedback. In this paper, we first present a centralized online algorithm and then extend it to a distributed online variant that takes into account an adversarial delay setting.

2 Projection-Free Algorithms Under Delayed Feedback

In this section, we will present our method for addressing delayed feedback in the OCO problem. In Sect. 2.1, we state some assumptions and results that form the basis of our approach. We then describe our first algorithm DeLMFW for the centralized setting in Sect. 2.2 and extend it to the distributed setting in Sect. 2.3.

2.1 Preliminaries

Throughout the paper, we use boldface letter e.g. \boldsymbol{x} to represent vectors. We denote by \boldsymbol{x}_t the final decision of round t and $\boldsymbol{x}_{t,k}$ to be the sub-iterate at round k of t. In distributed setting, we add a superscript i to make distinction between agents. If not specified otherwise, we use Euclidean norm $\|.\|$ and suppose that the constraint set $\mathcal{K} \subset \mathbb{R}^m$ is convex. We state the following standard assumptions in OCO.

Assumption 1. The constraint set \mathcal{K} is a bounded convex set with diameter D i.e. $D := \sup_{x,y \in \mathcal{K}} ||x - y||$.

Assumption 2 (Lipschitz). For all $x \in \mathcal{K}$, there exists a constant G such that, $\forall t \in [T], \|\nabla f_t(x)\| \leq G$.

Assumption 3 (Smoothness). For all $x, y \in \mathcal{K}$, there exists a constant β such that, $\forall t \in [T]$:

$$f_t(\boldsymbol{y}) \leq f_t(\boldsymbol{x}) + \langle
abla f_t(\boldsymbol{x}), \boldsymbol{y} - \boldsymbol{x}
angle + rac{eta}{2} \| \boldsymbol{y} - \boldsymbol{x} \|^2$$

or equivalently $\|\nabla f_t(\boldsymbol{x}) - \nabla f_t(\boldsymbol{y})\| \leq \beta \|\boldsymbol{x} - \boldsymbol{y}\|.$

Frank-Wolfe Algorithm. The Frank-Wolfe algorithm (FW) [4], also known as the Conditional Gradient method, is a first-order projection-free technique for solving constrained optimization problems. A common approach to these problems is projected gradient descent, where the solution is projected onto the constraint set at each iteration. However, the projection step is usually computationally expensive and may not be feasible for many large-scale problems. Conversely,

the FW algorithm avoids projections by iteratively moving toward the solution of a linear problem. The update rule of the Frank-Wolfe algorithm is given by:

$$\boldsymbol{x}_{t+1} = \boldsymbol{x}_t + \eta_t (\boldsymbol{v}_t - \boldsymbol{x}_t) \tag{1}$$

where v_t is the solution of the linear optimization problem $\arg\min_{v \in \mathcal{K}} \langle \nabla f_t(\boldsymbol{x}_t), \boldsymbol{v} \rangle$ and $\eta_t \in [0, 1]$ is a suitable step-size.

Online Linear Optimization Oracles In the context of FW algorithm, we utilize multiple optimization oracles to approximate the gradient of the upcoming loss function by solving an online linear problem. This approach was first introduced in [3]. Specifically, the online linear problem involves selecting a decision $v_t \in \mathcal{K}$ at every time $t \in [T]$. The adversary then reveals a vector g_t and loss function $\langle g_t, \cdot \rangle$ to the oracle. The objective is to minimize the oracle's regret. A possible candidate for an online linear oracle is the Follow the Perturbed Leader algorithm (FTPL) [9]. Given a sequence of historical loss functions $\langle g_\ell, \cdot \rangle, s \in [1, t]$ and a random vector n drawn uniformly from a probability distribution \mathcal{D} , FTPL makes the following update.

$$\hat{\boldsymbol{v}}_{t+1} = \operatorname*{arg\,min}_{\boldsymbol{v}\in\mathcal{K}} \left\{ \zeta \sum_{\ell=1}^{t} \langle \boldsymbol{g}_{\ell}, \boldsymbol{v} \rangle + \langle \boldsymbol{n}, \boldsymbol{v} \rangle \right\}$$
(2)

Lemma 1 (Theorem 5.8 [5]). Given a sequence of linear loss function f_1, \ldots, f_T . Suppose that Assumptions 1 to 3 hold true. Let \mathcal{D} be a the uniform distribution over hypercube $[0, 1]^m$. The regret of FTPL is

$$\mathcal{R}_{T,\mathcal{O}} \leq \zeta D G^2 T + \frac{1}{\zeta} \sqrt{m} D$$

where ζ is learning rate of algorithm.

Delay Mechanism. We consider the following delay mechanism. At round t, the agent receives a set of delayed gradient $\nabla f_s(\boldsymbol{x}_s)$ from previous rounds $s \leq t$ such that $s + d_s - 1 = t$, where d_s is the delay value of iteration s. We denote by $\mathcal{F}_t = \{s : s + d_s - 1 = t\}$ the set of indices released at round t. Following this setting, the feedback of round t is released at time $t + d_t - 1$, and the case $d_t = 1$ is considered as no delay. We suppose that the delay value is unknown to the agent and make no assumption about the set \mathcal{F}_t . Consequently it is possible for the set to be empty at any particular round. We extend the aforementioned mechanism to the distributed setting by assuming that each agent has a unique delay value at each round $t \in [T]$. The delay value of agent i at round t is denoted by $\mathcal{F}_t^i = \{s : s + d_s^i - 1 = t\}$, which is distinct between agents.

2.2 Centralized Algorithm

We describe the procedure of Algorithm 1 in details. At each round t, the agent performs two blocks of operations: prediction and update. During the prediction

block, the agent performs K iterations of FW updates by querying solutions from the oracles \mathcal{O}_k , $k \in [K]$ and updates the sub-iterate vector $\boldsymbol{x}_{t,k+1}$ using a convex combination of the previous one and the oracle's output. The agent then plays the final decision $\boldsymbol{x}_t = \boldsymbol{x}_{t,K+1}$ and incurs a loss $f_t(\boldsymbol{x}_t)$ which may not be revealed at t due to delay. From the mechanism described in Sect. 2.1, there exists a set of gradient feedbacks from the previous rounds revealed at t whose indices are in \mathcal{F}_t . The update block involves observing the delayed gradients evaluated at K sub-iterates of rounds $s \in \mathcal{F}_t$, computing surrogate gradients $\{\boldsymbol{g}_{t,k}, k \in [K]\}$ by summing the delayed gradients and feeding them back to the oracles $\{\mathcal{O}_k, k \in K\}$.

In our algorithm, the agent employs a suite of online linear optimization oracles, denoted $\mathcal{O}_1, \ldots, \mathcal{O}_K$. These oracles utilize feedbacks accumulated from previous rounds to estimate the gradient of the upcoming loss function. However, in the delay setting, these estimations may be perturbed owing to a lack of information. For example, if there is no feedback from rounds t to t', that is, $\mathcal{F}_s = \emptyset$ for $s \in [t, t']$, the oracles will resort to the information available in round t - 1 to estimate the gradient of all rounds from t + 1 to t' + 1. As a result, the oracle's output remains unchanged for these rounds, and decisions $\{x_s : s \in [t + 1, t' + 1]\}$ are not improved. Our analysis for Algorithms 1 will be focused on assessing the impact of delayed feedback on the oracle's output.

Lemma 2. Let \hat{v}_t be the FTPL prediction defined in Eq. 2 and

$$oldsymbol{v}_t = rgmin_{oldsymbol{v}\in\mathcal{K}} \left\{ \zeta \sum_{\ell=1}^{t-1} \left\langle \sum_{s\in\mathcal{F}_\ell} oldsymbol{g}_s, oldsymbol{v}
ight
angle + \langle oldsymbol{n}, oldsymbol{v}
ight
angle$$

the prediction of FTPL with delayed feedback. For all $t \in [T]$, we have:

$$\|\boldsymbol{v}_t - \hat{\boldsymbol{v}}_t\| \leq \zeta DG \sum_{s < t} \mathbb{I}_{\{s+d_s > t\}}$$

Theorem 1. Given a constraint set \mathcal{K} . Let $A = \max\left\{3, \frac{G}{\beta D}\right\}$, $\eta_k = \min\left\{1, \frac{A}{k}\right\}$, and $K = \sqrt{T}$. Suppose that Assumptions 1 to 3 hold true. If we choose FTPL as the underlying oracle and set $\zeta = \frac{1}{G\sqrt{B}}$, the regret of Algorithm 1 is

$$\sum_{t=1}^{T} \left[f_t(\boldsymbol{x}_t) - f_t(\boldsymbol{x}^*) \right] \le 2\beta A D^2 \sqrt{T} + 3(A+1) \left(DG \sqrt{B} + \mathcal{R}_{T,\mathcal{O}} \right)$$
(3)

where $B = \sum_{t=1}^{T} d_t$, the sum of all delay values and $\mathcal{R}_{T,\mathcal{O}}$ is the regret of FTPL with respect to the current choice of ζ .

Discussion. The regret bound of Theorem 1 differs from that of the non-delayed MFW [3] by the additive term $DG\sqrt{B}$ which represents the total cost of sending delayed feedback to the oracles over T rounds (Lemma 2). If we assume that

Algorithm 1. DeLMFW

Input: Constraint set \mathcal{K} , number of iterations T, sub-iteration K, online oracles $\{\mathcal{O}_k\}_{k=1}^K$, step sizes $\eta_k \in (0, 1]$

```
1: for t = 1 to T do
         Initialize arbitrarily x_{t,1} \in \mathcal{K}
 2:
          for k = 1 to K do
 3:
              Query v_{t,k} from oracle \mathcal{O}_k.
 4
              \boldsymbol{x}_{t,k+1} \leftarrow (1-\eta_k)\boldsymbol{x}_{t,k} + \eta_k \boldsymbol{v}_{t,k}.
 5:
         end for
 6:
         x_t \leftarrow x_{t,K+1}, play x_t and incurs loss f_t(x_t)
 7.
         Receive \mathcal{F}_t = \{s \in [T] : s + d_s - 1 = t\}
 8:
         if \mathcal{F}_t = \emptyset then
 9:
              do nothing
10:
         else
11:
              for k = 1 to K do
12:
                  \boldsymbol{g}_{t,k} \leftarrow \sum_{s \in \mathcal{F}_t} \nabla f_s(\boldsymbol{x}_{s,k})
13:
                  Feedback \langle g_{t,k}, \cdot \rangle to oracles \mathcal{O}_k.
14:
              end for
15:
         end if
16:
     end for
17:
```

there exists a maximum value d such that $d_t \leq d$ for all $t \in [T]$. Our regret bound becomes $O(\sqrt{dT})$ which coincides with the setting in [14], a delayedfeedback FW algorithm that achieves $O(T^{3/4} + dT^{1/4})$. Another line of work is from [8], a framework that addresses delayed feedback for any base algorithm. By considering MFW as the base algorithm, their theoretical analysis suggests that the algorithm also achieves $O(\sqrt{dT})$ regret bound. However, their delay value is not completely unknown to the agent because it is time-stamped by maintaining multiple copies of the base algorithm. We empirically show in Sect. 3 that this algorithm is highly susceptible to high delay values. Instead of using FTPL, our algorithm has the flexibility to select any online algorithm as an oracle, for example, Online Gradient Descent [5].

2.3 Distributed Algorithm

In this section, we extend Algorithm 1 to a distributed setting in which multiple agents collaboratively optimize a global model. Our setting considers a fully distributed framework, characterized by the absence of a server to coordinate the learning process. Let $\mathbf{W} \in \mathbb{R}^{n \times n}_+$ be the adjacency matrix of communication graph $\mathcal{G} = (V, E)$. The entries w_{ij} are defined as :

$$w_{ij} = \begin{cases} \frac{1}{1 + \max\{\tau_i, \tau_j\}} & \text{if } (i, j) \in E\\ 0 & \text{if } (i, j) \notin E, i \neq j\\ 1 - \sum_{j \in N(i)} w_{ij} & \text{if } i = j \end{cases}$$

1

where $\tau_i = \#\{j \in V : (i, j) \in E\}$ is the degree of vertex *i*. The matrix **W** is doubly stochastic (i.e. $\mathbf{W}\mathbf{1} = \mathbf{W}^T\mathbf{1} = \mathbf{1}$) and therefore possesses several useful properties associated with doubly stochastic matrices. We call $\lambda(\mathbf{W})$ the second-largest eigenvalue of **W** and define k_0 as the smallest integer that verifies $\lambda(\mathbf{W}) \leq \left(\frac{k_0}{k_0+1}\right)^2$. Furthermore, we set $\rho = 1 - \lambda(\mathbf{W})$ to be the spectral gap of matrix **W**.

At a high level, each agent maintains K copies of the oracles $\mathcal{O}_1^i, \dots, \mathcal{O}_K^i$ while performing prediction and update at every round t. The prediction block consists of performing K FW-steps while incorporating the neighbors' information. Specifically, the agent computes at its local level during the K steps a local average decision $\boldsymbol{y}_{t,k}^i$ representing a weighted aggregation of its neighbor's current sub-iterates. The update vector is convex combination of the local average decision and the oracle's output. The final decision of agent \boldsymbol{x}_t^i is disclosed at the end of K steps. Lemma 3 shows that $\boldsymbol{y}_{t,k}^i$ is a local estimation of the global average $\overline{\boldsymbol{x}}_{t,k} = \frac{1}{n} \sum_{i=1}^n \boldsymbol{x}_{t,k}^i$ as K increases.

Following the K FW-steps, the update block employs K gradient updates utilizing the delayed feedback from previous rounds. The agent observes the delayed gradients evaluated on theirs corresponding subiterates and computes the local average gradient $d_{t,k}^i$ through a weighted aggregation of the neighbors' current surrogates (18). The agent updates the surrogate gradient via a gradient-tracking step (19) to ensure that it approaches the global gradient as K increases. It is worth noting that feedback provided to the oracle contains information about delays experienced by all neighboring agents. Consequently, the oracle \mathcal{O}_k^i observes delayed feedback from $\cup_{j \in \mathcal{N}(i)} \mathcal{F}_t^j$ instead of \mathcal{F}_t^i . This result highlights the dependency on the connectivity of the communication graph when considering the effect of delayed feedback to the oracle's output, as demonstrated in Lemma 4.

Lemma 3. Define $C_d = k_0 \sqrt{nD}$, for all $t \in [T]$, $k \in [K]$, we have

$$\max_{i \in [1,n]} \left\| \boldsymbol{y}_{t,k}^{i} - \overline{\boldsymbol{x}}_{t,k} \right\| \le \frac{C_d}{k} \tag{4}$$

Lemma 4. For all $t \in [T]$, $k \in [K]$ and $i \in [n]$. Let $\mathbf{v}_{t,k}^i$ be the output of the oracle \mathcal{O}_k^i with delayed feedback and $\hat{\mathbf{v}}_{t,k}^i$ its homologous in non-delay case. Suppose that Assumptions 1 and 2 hold true. Choosing FTPL as the oracle, we have:

$$\left\|\boldsymbol{v}_{t,k}^{i} - \hat{\boldsymbol{v}}_{t,k}^{i}\right\| \le 2\zeta\sqrt{n}DG\left(\frac{\lambda\left(\mathbf{W}\right)}{\rho} + 1\right)\frac{1}{n}\sum_{i=1}^{n}\sum_{s\le t}\mathbb{I}_{\{s+d_{s}^{i}>t\}}$$
(5)

where ζ is the learning rate, $\lambda(\mathbf{W})$ is the second-largest eigenvalue of \mathbf{W} and $\rho = 1 - \lambda(\mathbf{W})$ is the spectral gap of matrix \mathbf{W} .

Theorem 2. Given a constraint set \mathcal{K} . Let $A = \max\left\{3, \frac{3G}{2\beta D}, \frac{2\beta C_d + C_g}{\beta D}\right\}$, $\eta_k = \min\left\{1, \frac{A}{k}\right\}$, and $K = \sqrt{T}$. Suppose that Assumptions 1 to 3 hold true. If we

choose FTPL as the underlying oracle and set $\zeta = \frac{1}{G\sqrt{B}}$, the regret of Algorithm 2 is

$$\sum_{t=1}^{T} \left[F_t(\boldsymbol{x}_t^i) - F_t(\boldsymbol{x}^*) \right] \le \left(GC_d + 2\beta AD^2 \right) \sqrt{T}$$

$$+ 3(A+1) \left(2\sqrt{n}DG\left(\frac{\lambda\left(\mathbf{W}\right)}{\rho} + 1\right) \sqrt{B} + \mathcal{R}_{T,\mathcal{O}} \right)$$
(6)

where $B = \frac{1}{n} \sum_{i=1}^{n} B_i$ such that B_i is the sum of all delay values of agent i. $C_d = k_0 \sqrt{nD}$ and $C_g = \sqrt{n} \max\left\{\lambda_2(\mathbf{W}) \left(G + \frac{\beta D}{\rho}\right), k_0 \beta \left(4C_d + AD\right)\right\}$ and $\mathcal{R}_{T,\mathcal{O}}$ is the regret of FTPL with respect to the current choice of ζ .

Algorithm 2. De2MFW

Input: Constraint set \mathcal{K} , number of iterations T, sub-iterations K, online linear optimization oracles $\{\mathcal{O}_k^i : k \in [K]\}$ for each agent $i \in [n]$, step sizes $\eta_k \in (0, 1]$

1: for t = 1 to T do for every agent i = 1 to n do 2: Initialize arbitrarily $x_{t,1}^i \in \mathcal{K}$ 3: for k = 1 to K do 4: Query $\boldsymbol{v}_{t,k}^i$ from oracle \mathcal{O}_k^i 5: Exchange $x_{t,k}^i$ with neighbours $\mathcal{N}(i)$ 6: $\boldsymbol{y}_{t,k}^i \leftarrow \sum_j w_{ij} \boldsymbol{x}_{t,k}^j$ 7: $\boldsymbol{x}_{t,k+1}^{i} \leftarrow (1-\eta_k) \boldsymbol{y}_{t,k}^{i} + \eta_k \boldsymbol{v}_{t,k}^{i}$ 8: end for 9: $\boldsymbol{x}_{t}^{i} \leftarrow \boldsymbol{x}_{t,K+1}^{i},$ play \boldsymbol{x}_{t}^{i} and incurs loss $f_{t}^{i}(\boldsymbol{x}_{t}^{i})$ 10: Receive $\mathcal{F}_t^i = \left\{ s \in [T] : s + d_s^i - 1 = t \right\}$ 11: if $\mathcal{F}_t^i = \emptyset$ then 12: do nothing 13: else 14: $\boldsymbol{g}_{t,1}^i \leftarrow \sum_{s \in \mathcal{F}_t^i} \nabla f_s^i(\boldsymbol{x}_{s,1}^i)$ 15for k = 1 to K do 16: Exchange $\boldsymbol{g}_{t,k}^{i}$ with neighbours $\mathcal{N}(i)$ 17: $\boldsymbol{d}_{t,k}^{i} \leftarrow \sum_{j \in \mathcal{N}(i)} w_{ij} \boldsymbol{g}_{t,k}^{j}$ 18: $egin{aligned} egin{aligned} egi$ 19. Feedback $\langle \boldsymbol{d}_{t,k}^i, \cdot \rangle$ to oracles $\mathcal{O}_{i,k}$ 20: end for 21: end if 22: end for 23 24: end for

3 Numerical Experiments

We evaluated the performance of our algorithms on the online multiclass logistic regression problem using two datasets: MNIST and FashionMNIST. MNIST is a well-known hand-digit dataset that contains 60000 grayscale images of size (28×28) , divided into 10 classes, and FashionMNIST includes images of fashion products with the same configuration. We conducted the experiment¹ using Julia 1.7 on MacOS 13.3 with 16 GB of memory.

Centralized Setting. Given an iteration t, the agent receives a subset \mathcal{B}_t of the form $\mathbf{b}_t = \{\mathbf{a}_t, y_t\} \in \mathbb{R}^m \times \{1, \ldots, C\}$, consisting of the features vector \mathbf{a}_t and the corresponding label y_t . We define the loss function f_t as

$$f_t(\boldsymbol{x}) = -\sum_{\boldsymbol{b}_t \in \mathcal{B}_t} \sum_{c=1}^C \left\{ y_t^i = c \right\} \log \frac{\exp\left\langle \boldsymbol{x}_c, \boldsymbol{a}_t^i \right\rangle}{\sum_{\ell=1}^C \exp\left\langle \boldsymbol{x}_\ell, \boldsymbol{a}_t^i \right\rangle}$$
(7)

where \boldsymbol{x} must satisfy the constraint $\boldsymbol{x} \in \mathcal{K}$ such that $\mathcal{K} = \{\boldsymbol{x} \in \mathbb{R}^{m \times C}, \|\boldsymbol{x}\|_1 \leq r\}$. Using the MNIST dataset, we note $m = 784, C = 10, r = 8, |\mathcal{B}_t| = 60$ and a total of T = 1000 rounds. To evaluate the performance of the algorithm under different delay regimes, we generated a random sequence of delays d_t such that $d_t \leq d$ for $d \in \{21, 41, 61, 81, 101\}$. We compared the performance of DeLMFW against DOFW [14], a projection-free algorithm with adversarial delay, and BOLD-MFW [8], an online learning framework designed to handle delayed feedback. Figure 2 displays the performance of the three algorithms under various delay regimes. In the absence of delay, that is, d = 1 (left figure), DeLMFW and BOLD-MFW have the same performance since both algorithms reduce to MFW [3] with regret of $O(\sqrt{T})$. Meanwhile, DOFW is the classical OFW [6] that guarantees a regret of $O(\sqrt{dT})$ when the delay is upperbounded by d. In the case where $d \leq T^{1/2}$ (middle figure, d = 21), the dominant



Fig. 2. Cumulative loss comparison for different delays regimes. Left: Without delay. Middle: Maximum delay 21. Right: Maximum delay 101.

¹ https://github.com/tuananhngh/DelayMFW.

term in DOFW is $T^{3/4}$, while DeLMFW takes advantage of the regret of order $\sqrt{dT} \leq T^{3/4}$. For $d \geq T^{1/2}$ (right figure, d = 101), DOFW's regret is dominated by the term $dT^{1/4}$, which is outperformed by DeLMFW, particularly for high values of d. This result confirms our theoretical analysis in Sect. 2.2.



Fig. 3. Total loss of BOLD-MFW, DOFW and DeLMFW when varying delay value.

Figure 3 illustrates the total loss of DeLMFW and the other two algorithms when increasing d to show the sensitivity of each algorithm in the presence of delays. As BOLD is a general framework that can be applied to any base algorithm, it is noticeable that it is susceptible to high levels of delays. This phenomenon has also been observed in [14] when using BOLD with OFW, highlighting the need for customized design algorithms in the context of delayed feedback.

Distributed Setting. In the second experiment, we examine the online distributed multiclass logistic regression problem on the FashionMNIST dataset, using a network of 30 agents. The algorithm was run on four different topologies, including Erdos-Renyi, Complete, Grid, and Cycle. At each iteration $t \in [T]$, each agent i received a subset \mathcal{B}_t^i of the form $\{\boldsymbol{a}_t^i, y_t^i\} \in \mathbb{R}^d \times \{1, \ldots, C\}$, that consisted of feature vectors \boldsymbol{a}_t^i and its corresponding label y_t^i . The goal was to collaboratively optimize the global loss function $F_t(\boldsymbol{x}) = \frac{1}{n} \sum_{i=1}^n f_t^i(\boldsymbol{x})$, where the local loss f_t^i was defined in Eq. 7.

For this experiment, we set m = 784, C = 10, r = 32, $|\mathcal{B}_t^i| = 2$ and T = 1000 rounds. We are interested in examining the effect of delays on network performance and thus randomly select f < n agents to have delayed feedback with a maximum value of 501. We compared the total loss on each topology under these conditions and present the result in Fig. 4. We observe that the presence of delayed agents has a significant impact on the network performance of Cycle graph as the number of delayed agents increases, while the complete graph is less affected. This result is consistent with the analysis in Sect. 2.3 because the delay term in the regret bound depends on the connectivity of the communication graph.



Fig. 4. Total Loss with varying numbers of agents experiencing delayed feedback in the network. (f = 0) for no delayed-agents.

In Table 2, we report the change in total loss as the number of delayed agents increases. We observe that the average percentage change is smaller for Grid than for Erdos-Renyi when compared with the network of non-delayed agents (f = 0). This result indicates that the generated Erdos-Renyi graph is more sensitive to the presence of delayed agents.

Table 2. Total Loss of the algorithm running on 4 different topology. We randomly select f < n agents to have delay with maximal value to be 501. In parenthesis, the percentage of total loss compared that of no delayed agents in the network (i.e. f = 0).

f	Topology								
	Erdős-Rényi	Grid	Complete	Cycle					
0	809.37	855.62	799.49	925.72					
2	820.15~(+1.3%)	852.15 (-0.4%)	798.79~(-0.08%)	932.34~(+0.7%)					
5	$834.74 \ (+3.0\%)$	868.52~(+1.4%)	802.59~(+0.3%)	$971.24 \ (+4.7\%)$					
10	$838.74 \ (+3.5\%)$	$878.04 \ (+2.5\%)$	792.45 (-0.8%)	983.89~(+6.0%)					
20	$850.49 \ (+4.9\%)$	902.30~(+5.3%)	$812.21\ (+1.5\%)$	1119.24~(+18.9%)					

4 Concluding Remarks

In this paper, we propose two algorithms to solve the online convex optimization problem with adversarial delayed feedback in both centralized and decentralized settings. These algorithms achieve optimal $O(\sqrt{dT})$ regret bounds, where d is the upper bound of the delays. The experimental results show that our algorithms outperform existing solutions in both centralized and decentralized settings, which are predictable by our theoretical analysis. Although the algorithms achieve good performance guarantees for the online convex optimization problem with adversarial delays, they currently rely on exact gradients, which may not be feasible for many real-world applications. Therefore, future research could explore the use of stochastic gradients with variance reduction techniques. Additionally, in decentralized settings, communication delays can be practically challenging and further improvements are needed in this area. However, our work demonstrates the potential of using Frank-Wolfe-type algorithms to solve convex constraint optimization problems with adversarial delays, which is beneficial for learning on edge devices.

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Secure Dataset Condensation for Privacy-Preserving and Efficient Vertical Federated Learning

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Abstract. This work addresses the dual challenges of enhancing training efficiency and protecting data privacy in Vertical Federated Learning (VFL) through secure synthetic dataset generation. VFL typically involves an active party with labels collaborating with a passive party possessing features of the same set of samples. Traditional VFL methods, however, rely on training with entire datasets of sensitive real data, leading to two primary issues: 1) reduced training efficiency due to large dataset sizes, a concern exacerbated in cryptography-based training methods; and 2) potential *privacy* leakage at the sample level during training. To mitigate these issues, we introduce the Vertical Federated Dataset Condensation (VFDC) method. VFDC employs a novel mixed protection mechanism, integrating class-wise secure aggregation, differential privacy and repetitive initialization, to securely match the distributions of real and synthetic data. Empirical evaluations on six real-world datasets validate VFDC's efficacy in generating small synthetic data for VFL, achieving a superior utility-privacy-efficiency trade-off during federated training.

Keywords: Dataset condensation \cdot Vertical federated learning \cdot Privacy protection \cdot Training efficiency

1 Introduction

Vertical Federated Learning (VFL) has attracted increasing attention [20, 32, 34] as it enables collaborative model training between institutions without disclosing

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private raw data. In a typical VFL setting, as shown in Fig. 1(a), one active party holds the labels and the other passive party possesses features of overlapping samples. The goal is to train a federated model without disclosing sample-wise data privacy to the opposite party. Traditional VFL methods train a federated model using the entire training set of sensitive real data. However, this vanilla paradigm confronts two major challenges: 1) reduced training *efficiency* due to the large dataset size, and 2) significant *privacy* leakage risks at the sample level stemming from the use of real data. We elaborate on these challenges in the following.



Fig. 1. Schematic comparison between vanilla VFL and VFDC: (a) Vanilla VFL trains on the *entire real* dataset and is vulnerable to sample-wise privacy leakage. (b) Our VFDC method securely generates a *small synthetic* dataset for privacy-preserving and efficient federated training.

Efficiency Challenge. The training efficiency in practical VFL scenarios is significantly affected by the large volume of training data. This challenge is amplified by the necessity of conducting multiple trials for hyperparameter optimization and Neural Architecture Search (NAS), and the slow convergence on real dataset for other tasks [14]. This is particularly problematic in the widely used cryptography-based training methods which is computational and communicational costly, and thus exacerbates the efficiency challenge under repeated training trails [7, 12].

Privacy Challenge. As shown in Fig. 1(a), it has been demonstrated that vanilla VFL is vulnerable to privacy leakage of the aligned real samples [34]. Specifically, during the training phase, an adversarial active party may attempt

to infer private features of a target sample from the passive party, and vice versa [11,13,17]. In response, existing solutions predominantly focus on *training-phase* protection, falling into two broad categories: cryptography-based methods, such as multiparty computation [12] and homomorphic encryption [8], and perturbation-based methods, notably differential privacy (DP) [9]. However, cryptography-based training methods become impractical for large datasets [5]. On the other hand, perturbation-based training methods require significant perturbation to ensure feature privacy of aligned samples in VFL, often at the cost of reduced model utility and efficiency [18,22].

Problem Setup. Therefore, reducing the size of real training dataset in a preprocessing phase is crucial for enhancing both efficiency and privacy in VFL training phase. However, it remains a relatively unexplored area. While vertical federated coreset selection [14] samples a small subset of real data to improve efficiency, it still involves training on real samples, risking feature privacy leakage of the selected real samples during training. To kill the two birds with one stone, our idea is to generate a synthetic dataset in a preprocessing phase, and thus completely avoid leaking real sample privacy during the training phase. This idea leads us to the following research question:

How can we securely generate a small set of synthetic data to efficiently train VFL models while ensuring high utility and strong sample-wise data privacy protection?

Our Solution. To address the posed question, we leverage recent advancements in dataset condensation (DC) [35]. DC aims to match the distribution between a small synthetic dataset and the full real dataset, such that training on the former achieves performance comparable to training on the latter [35]. While DC has been extensively applied in horizontal federated learning (HFL) to tackle issues like data heterogeneity [16,23], training efficiency [19], and privacy enhancement [31], its adoption is straightforward since clients can independently perform DC on their locally available, labeled data.

In contrast, integrating DC into VFL presents a markedly more complex challenge. The distributed nature of data across VFL parties complicates the task of preserving privacy during the transmission of sample-wise real data embeddings, illustrating the novel challenge our work seeks to overcome.

To securely integrate DC into VFL, we introduce the secure Vertical Federated Dataset Condensation (VFDC) approach, depicted in Fig. 1(b). VFDC aligns the distributions of real and synthetic features for each class by updating passive party's synthetic features with respect to active party's randomly generated labels. We employ a secret sharing-based class-wise secure aggregation algorithm for securely computing class-average embeddings without exposing individual features and labels. Moreover, feature privacy is further strengthened by incorporating differential privacy (DP) into the averaged embeddings. A key aspect of our approach is the class-wise secure aggregation, which reduces sensitivity and thus allows for the addition of subtle DP noise. **Experiments.** We evaluate our proposed VFDC on six real-world datasets. In terms of privacy-utility trade-off, VFDC demonstrates superior utility under the same DP privacy budgets compared to other baseline federated dataset reduction methods. For efficiency, the VFDC-generated datasets significantly reduce the training iterations of VFL models, compared to other baselines.

Contributions. In summary, our major contributions are as follows:

- 1. Our work pioneers the topic of secure dataset condensation in VFL, addressing the dual challenges of efficiency and privacy protection in the training phase.
- 2. We propose a mixed protection mechanism based on class-wise secure aggregation, DP, and repetitive random initialization, enabling subtle noise addition for sample-wise feature privacy protection while maintaining high utility.
- 3. We conduct extensive experiments to validate the superior performance of our proposed VFDC algorithm on six public real-world datasets.

Roadmap. The remainder of the paper is organized as follows: Sect. 2 reviews related work. Section 3 details the problem formulation and foundational concepts of our approach. Section 4 elaborates on the VFDC approach. Experimental results are presented in Sect. 5. The paper concludes in Sect. 6.

2 Related Work

2.1 Vertical Federated Learning

Vertical Federated Learning (VFL) facilitates model training across parties with vertically partitioned data, where an active party holds labels and passive parties possess corresponding features [20,28]. Despite its advantages, VFL faces notable challenges in efficiency and sample-wise privacy leakage during training. The exchange of intermediate model results risks privacy breaches, including label leakage to passive parties [11] and potential feature inference from passive parties' outputs [33]. Our research aims to mitigate these concerns, focusing on enhancing both privacy protection and training efficiency in the VFL framework.

2.2 Privacy Protection in VFL

Numerous studies in VFL have been devoted to protecting the label and feature privacy of real data. Despite this, the majority of existing methods target the training and inference phase, with limited applicability in the *data generation* phase. Common privacy protection techniques in VFL include cryptographic methods [7,12] and perturbation methods [17]. Cryptographic solutions, while secure, often lead to significant communication and computational overheads, rendering them impractical for large-scale datasets [12]. On the other hand, training-phase perturbation methods aim to protect feature and label privacy leakage through embeddings and gradients during training [13,17,24,27]. However, due to the alignment of samples across parties in VFL, ensuring adequate sample-wise privacy often requires extensive perturbation, which can compromise data utility and efficiency [18,22]. This underscores the need for a novel approach to protect sample-wise data privacy of real dataset via synthetic dataset generation.

2.3 Dataset Size Reduction in FL

The reduction of dataset size in FL has recently become a focus due to its potential to improve both training efficiency and privacy [19,30,31]. Most existing works target the HFL settings, where data is partitioned by samples among parties. Prototype-based HFL methods [23,31] create a small set of prototypes to represent the original dataset, addressing the non-i.i.d. issue. Dataset distillationbased HFL methods synthesize a few samples to enhance efficiency and privacy [19,30]. However, these approaches involve complex bi-level optimizations and high computational demands, limiting their applicability in VFL. Dataset condensation (DC) is also explored in HFL to boost both efficiency and privacy [19,30], with distribution matching-based DC methods being particularly efficient and effective [35]. In the VFL setting, coreset selection is examined to improve efficiency [14]. However, this method still involves training on real data samples, risking sample-wise data privacy leakage. As of now, there is a lack of methods capable of securely generating small synthetic datasets in VFL.

Notably, despite criticisms [5] of DC for not guaranteeing membership privacy against model users [4,26], our focus is on defending data inference attacks from another party, where membership information is already known. Hence, these critiques of DC's membership privacy limitations are irrelevant to our approach.

3 Preliminaries

3.1 Problem Formulation

Vertical Federated Learning Setting. In a typical VFL scenario, two parties engage in a classification task: the active party P_0 , holding real labels $\boldsymbol{y} \in \mathcal{Y}$, and the passive party P_1 , possessing real features $\boldsymbol{x} \in \mathcal{X}$ corresponding to the same set of samples. The goal is to collaboratively train a model $f_{\theta} : \mathcal{X} \mapsto \mathcal{Y}$, parameterized by θ , without compromising the privacy of sample-wise data for either party. Furthermore, our VFDC method is adaptable to scenarios involving multiple passive parties, where the active party can separately apply the VFDC procedure with each passive party, thereby maintaining the framework's efficacy in more complex VFL configurations.

Threat Model. Our threat model is consistent with the conventional setting adopted in prevalent VFL studies [6,11]. We operate under the assumption that all parties in VFL are honest-but-curious. This means they adhere to the prescribed training protocols yet possess the inclination to derive private information from any exchanged intermediate data, without engaging in collusion.

Our privacy objective is to protect the features and labels of the aligned samples against the opposite party in VFL during training. Within this context, the adversarial active party aims to deduce feature privacy from the intermediate results obtained from passive parties, while an adversarial passive party seek to infer labels of the active party. Thus, the critique [5] of centralized DC's limitation in membership privacy protection against adversarial *model users* does not apply to our VFL setting, where the membership information of aligned data is already known by parties.

3.2 Dataset Condensation

In a VFL setting, the entire training dataset $\mathcal{T} = \{\mathcal{X}, \mathcal{Y}\}$, partitioned between two parties, consists of $|\mathcal{T}|$ samples. Dataset condensation (DC) aims to generate a significantly smaller condensed dataset \mathcal{S} , with $|\mathcal{S}| \ll |\mathcal{T}|$. The goal is for \mathcal{S} to mimic the data distribution of \mathcal{T} , enabling model training on the condensed dataset \mathcal{S} to yield performance comparable to that on the entire dataset \mathcal{T} as follows:

$$\mathbb{E}_{\boldsymbol{x}\sim P_{\mathcal{D}}}\mathcal{L}(f_{\theta^{\mathcal{T}}}(\boldsymbol{x}), y) \approx \mathbb{E}_{\boldsymbol{x}\sim P_{\mathcal{D}}}\mathcal{L}(f_{\theta^{\mathcal{S}}}(\boldsymbol{x}), y),$$
(1)

where $P_{\mathcal{D}}$ represents the actual data distribution, \mathcal{L} denotes the loss function, and $\theta^{\mathcal{T}}$, $\theta^{\mathcal{S}}$ are model parameters trained on \mathcal{T} and \mathcal{S} , respectively.

DC synthesizes S by employing distribution matching, minimizing the Maximum Mean Discrepancy (MMD) between the actual training data T_c and synthetic data S_c for each class c as follows:

$$\sup_{\|\psi_{\vartheta}\|_{\mathcal{H}} \le 1} \left(\mathbb{E} \left[\psi_{\vartheta}(\mathcal{T}_c) \right] - \mathbb{E} \left[\psi_{\vartheta}(\mathcal{S}_c) \right] \right),$$
(2)

where ψ_{ϑ} is a function in the reproducing kernel Hilbert space (RKHS) \mathcal{H} , parameterized by ϑ . The empirical MMD approximates the actual MMD for each class, with the overall MMD loss across all classes given by:

$$\sum_{c=0}^{C-1} \mathbb{E}_{\vartheta \sim P_{\vartheta}} \left\| \frac{1}{|\mathcal{T}_c|} \sum_{i=1}^{|\mathcal{T}_c|} \psi_{\vartheta} \left(\boldsymbol{x}_{c,i} \right) - \frac{1}{|\mathcal{S}_c|} \sum_{j=1}^{|\mathcal{S}_c|} \psi_{\vartheta} \left(\boldsymbol{s}_{c,j} \right) \right\|^2, \tag{3}$$

where P_{ϑ} is the distribution of network parameters. For each class $c, x_{c,i}$ and $s_{c,j}$ are the *i*-th sample in training data \mathcal{T}_c and the *j*-th sample in synthetic data \mathcal{S}_c , respectively.

The synthetic data S is learned by minimizing the discrepancy between T and S across various embedding spaces, sampling ϑ . Importantly, Eq. 3 allows efficient optimization of S, avoiding the complex bi-level optimization of data and model [29], and thus is well-suited for secure computation in VFL.

3.3 Secure Aggregation

Secure aggregation [3] corresponds to computing the sum of multiple inputs while keeping the individual inputs confidential. It has been widely adopted in HFL to aggregate model updates among clients [21], without revealing private local models to the aggregation server. However, it is still under-explored to use secure aggregation in VFL. In this study, we utilize secure aggregation to aggregate class-wise embeddings in Eq. 3. Techniques such as secret sharing and homomorphic encryption [8] are commonly employed for secure aggregation. For our purposes, we employ arithmetic secret sharing, specifically enhanced by the Beaver's triples technique, due to its high efficiency in secure computations.

Arithmetic Secret Sharing [25] involves three key procedures: 1) Share: To share a secret value a within a party P_1 , it is first converted to an l-bit integer. P_1 randomly generates $a_0 \in \mathbb{Z}_{2^l}$, sends a_0 to P_0 , and retains $a_1 = a - a_0 \mod 2^l$. The shared secret a is represented as $\langle a \rangle = \{a_0, a_1\}$. 2) Computation: Secure addition of two shared values $\langle a \rangle$ and $\langle b \rangle$ is performed locally by each party, resulting in $\langle a + b \rangle = \{a_0 + b_0, a_1 + b_1\}$. Secure multiplication can be conducted using precomputed Beaver's triples [2]. 3) Reconstruction: To reconstruct $\langle a \rangle$ for P_0, P_1 sends a_1 to P_0 , who then calculates $a = a_0 + a_1 \mod 2^l$.

3.4 Differential Privacy

Definition 1 (Differential Privacy [9]). A randomized mechanism \mathcal{M} satisfies (ϵ, δ) -differential privacy if, for any two neighboring datasets \mathcal{D} and \mathcal{D}' differing in only one sample, and for any output event E, the following inequality holds:

$$\Pr[\mathcal{M}(\mathcal{D}) \in E] \le e^{\epsilon} \Pr[\mathcal{M}(\mathcal{D}') \in E] + \delta,$$

where ϵ is the privacy budget, and δ is the fault-tolerance probability.

Definition 2 (l_2 -Sensitivity [10]). For any two neighboring datasets \mathcal{D} and \mathcal{D}' , the l_2 -sensitivity of a function $f : \mathcal{D} \to \mathbb{R}^d$ is defined as:

$$\Delta f = \max_{\mathcal{D}, \mathcal{D}'} ||f(\mathcal{D}) - f(\mathcal{D}')||_2,$$

where $|| \cdot ||_2$ denotes the l_2 -norm.

 l_2 -sensitivity quantifies the maximum impact that a single individual's data can have on the output of a function f. To achieve differential privacy, the Gaussian mechanism is applied as follows:

Definition 3 (Gaussian Mechanism [10]). For a function $f : \mathcal{D} \to \mathbb{R}^d$ with l_2 -sensitivity Δf , the Gaussian mechanism \mathcal{M} is defined as:

$$\mathcal{M}(f(\mathcal{D})) = f(\mathcal{D}) + \mathcal{N}(0, \sigma^2 \mathbf{I}_d),$$

where $\mathcal{N}(0, \sigma^2 \mathbf{I}_d)$ represents Gaussian noise with mean 0 and covariance matrix $\sigma^2 \mathbf{I}_d$, and σ is calculated as $\sigma = \frac{\Delta f \sqrt{2d \ln(1.25/\delta)}}{\epsilon}$.

Theorem 1. [10] The Gaussian mechanism, as defined in Definition 3, satisfies (ϵ, δ) -DP for each data publication.



Fig. 2. Overview of the proposed VFDC approach. 1) Left: The passive party computes the DP-protected embeddings of real features and synthetic features. 2) Middle: The two parties engage in *class-wise secure aggregation* to collaboratively compute the average embedding for each class. 3) **Right**: The active party computes the MMD loss and sends the gradients to the passive party to update the synthetic features.

4 Proposed Approach

4.1 Overview

Given the aforementioned building blocks, we introduce our proposed secure vertical federated dataset condensation (VFDC) approach, as shown in Fig. 2. We first introduce the cornerstone of VFDC, class-wise secure aggregation, in Sect. 4.2, which enables two parties to securely compute the average embeddings per each class, without disclosing the sample-level label and embedding information. Then, we elaborate on the VFDC approach in Sect. 4.3, which involves updating synthetic features in alignment with randomly assigned labels to achieve distribution matching. Lastly, Sect. 4.4 provides a thorough privacy analysis of the VFDC approach.

4.2 Class-Wise Secure Aggregation

As highlighted in Sect. 3.2, the computation of the MMD loss in DC involves averaging sample-wise feature embeddings by class. However, straightforward computation of these average embeddings in VFL risks privacy leakage: 1) transmitting individual sample embeddings from the passive to the active party may leak feature privacy, 2) whereas sharing individual sample labels from the active to the passive party could compromise label privacy. To tackle this challenge, we introduce a class-wise secure aggregation approach that aggregates class embeddings without disclosing sample-wise label information to passive parties or feature information to the active party. Our approach, utilizing arithmetic secret

Algorithm 1. Class-wise Secure Aggregation

Input: Label vector \boldsymbol{y} in active party P_0 , feature embeddings \boldsymbol{e} of real data in passive party P_1 , class number C.

Output: Class-wise average embeddings $\bar{e} = \{\bar{e}_c\}_{c=0}^{C-1}$.

1: procedure CLS_SECAGG(y, e)

- 2: Active party \overline{P}_0 maps label vector \boldsymbol{y} to mask vectors $\{\boldsymbol{m}_c\}_{c=0}^{C-1}$ for each class c.
- 3: The key distribution server generates Beaver's triples and distributes to P_0 and P_1 .
- 4: for each class c in parallel do
- 5: P_0 and P_1 secret share m_c and e with each other and get shared $\langle m_c \rangle$ and $\langle e \rangle$.
- 6: P_0 and P_1 compute inner product $\langle \boldsymbol{m}_c \cdot \boldsymbol{e} \rangle$ via Beaver's triples technique.
- 7: P_0 and P_1 reconstruct $\langle \boldsymbol{m}_c \cdot \boldsymbol{e} \rangle$ to P_0 to get $\boldsymbol{m}_c \cdot \boldsymbol{e}$.
- 8: P_0 computes the average embedding $\bar{e}_c = \frac{m_c \cdot e}{\sum m_c}$.

9: **end for**

10: **Return** Average embeddings $\bar{e} = \{\bar{e}_c\}_{c=0}^{C-1}$.

11: end procedure

sharing with Beaver's triples, ensures efficient secure computation. Although alternatives like HE [8] could offer similar security with larger computation and communication costs, the novelty of our method lies in extending these principles to class-wise aggregation for distribution matching, distinct from existing approaches aggregating among samples or parties.

Algorithm 1 outlines the class-wise secure aggregation algorithm. Initially, the active party P_0 converts the label vector \boldsymbol{y} into class-specific mask vectors \boldsymbol{m}_c for each class $c \in C$. A key distribution server then generates Beaver's triples and distributes them to both P_0 and P_1 . Both parties then secret share \boldsymbol{m}_c and embeddings \boldsymbol{e} , obtaining $\langle \boldsymbol{m}_c \rangle$ and $\langle \boldsymbol{e} \rangle$. Subsequently, P_0 and P_1 collaboratively compute the inner product $\langle \boldsymbol{m}_c \cdot \boldsymbol{e} \rangle$ utilizing the Beaver's triples. The inner product result $\boldsymbol{m}_c \cdot \boldsymbol{e}$ is reconstructed at P_0 for calculating average embeddings $\bar{\boldsymbol{e}}_c$ of each class.

Complexity Analysis. We now discuss the communication complexity of Algorithm 1. The adoption of secret sharing leads to extra communication cost compared to plain-text computation. Assuming the embedding size d, bit-length of shared value l, batch-size n, and class number C, the data size of our proposed class-wise secure aggregation is (4d + 3C)nl bits.

4.3 VFDC Algorithm

Based on the proposed class-wise secure aggregation, we introduce the Vertical Federated Dataset Condensation (VFDC) approach in Algorithm 2. Initially, both parties collaboratively initialize a synthetic dataset S. Each epoch begins with P_0 generating a random permutation to shuffle both real and synthetic datasets in coordination with P_1 , aiming for class-balanced mini-batches. Concurrently, P_1 resets the parameters ϑ of the embedding extractor ψ_{ϑ} . Algorithm 2. Vertical Federated Dataset Condensation (VFDC)

Input: Real training dataset \mathcal{T} , neural network ψ_{ϑ} parameterized with ϑ , learning rate η , epoch number T.

Output: Synthetic dataset \mathcal{S} .

- 1: Active party P_0 and passive party P_1 randomly initialize synthetic dataset S.
- 2: for each epoch t < T do
- 3: **Re-shuffle**: Active party P_0 generates random permutation and sends to P_1 to uniformly shuffle data.
- 4: **Re-initialize**: Passive party P_1 randomly initializes network parameter ϑ .
- 5: for each batch $\{y^{\mathcal{T}}, x^{\mathcal{T}}\} \subset \mathcal{T}, \{y^{\mathcal{S}}, x^{\mathcal{S}}\} \subset \mathcal{S}$ do
- 6: **DP embeddings**: Passive party P_1 computes real data embeddings $e^{\mathcal{T}} = \psi_{\vartheta}(\boldsymbol{x}^{\mathcal{T}})$ and add DP noise $\mathcal{N}(0, \sigma^2 \boldsymbol{I}_d)$.
- 7: Class-wise secure aggregation: Two parties P_0 and P_1 compute classaverage embeddings $\bar{e}^T = \text{CLS}_\text{SEcAgg}(y^T, e^T)$ via Algorithm 1.
- 8: P_1 computes synthetic data embeddings $e^{\mathcal{S}} = \psi_{\vartheta}(x^{\mathcal{S}})$.
- 9: Two parties compute class-average embeddings $\bar{e}^{S} = CLs_SECAGG(y^{S}, e^{S}).$
- 10: **Update synthetic features:** P_0 computes MMD loss \mathcal{L}_{MMD} via Equation 3 and sends gradients $\nabla_{es} \mathcal{L}_{MMD}$ to P_1 .
- 11: P_1 computes and updates $s = s \eta \nabla_s \mathcal{L}_{MMD}$.
- 12: end for
- 13: end for
- 14: **Return** Synthetic dataset S.

During secure dataset condensation, a mini-batch of real data $\{\boldsymbol{x}^{\mathcal{T}}, \boldsymbol{y}^{\mathcal{T}}\} \in \mathcal{T}$ is sampled. P_1 computes and DP-protects real data embeddings, adding Gaussian noise as per Definition 3, where $\sigma = \frac{r\sqrt{2d\ln(1.25/\delta)}}{b\epsilon}$, r is the clipping threshold and b is the class-wise batch size. Both parties then perform class-wise secure aggregation to calculate average class embeddings for real data. A similar process is followed for synthetic data $\{\boldsymbol{x}^{\mathcal{S}}, \boldsymbol{y}^{\mathcal{S}}\} \in \mathcal{S}$ to compute average class embeddings for synthetic data. P_0 computes the MMD loss using average class embeddings and communicates the gradients to P_1 , who updates the synthetic features \boldsymbol{s} . This process iteratively refines the small synthetic dataset, enhancing both privacy and efficiency in VFL training.

4.4 Privacy Analysis

This section assesses the privacy aspects of the VFDC approach (Algorithm 2), focusing on the confidentiality of real data features and labels. Under the assumption that all parties are honest-but-curious and non-collusive, as defined in Sect. 3.1, we analyze the privacy protection mechanisms in VFDC.

Theorem 2. The class-wise secure aggregation (Algorithm 1) ensures that the passive party gains no information about real data labels, and the active party learns only the average class embeddings.

Proof. Notice that other than the secret-shared intermediate results, the active party only receives decrypted average class embeddings, and the passive party only receives decrypted class-wise mask vectors. Therefore, the security of Algorithm 1 is guaranteed by the security of arithmetic secret sharing [25] and the Beaver's triples [2]. Therefore, the proposed class-wise secure aggregation method effectively protects both sample-wise label and feature privacy of real data. \Box

Feature Privacy Protection. Given the heightened sensitivity and potential vulnerability of feature data in VFL, VFDC places a particular emphasis on feature privacy by integrating secure aggregation, DP, and repeated random model initialization.

1) Class-wise Secure Aggregation + DP: Our proposed class-wise secure aggregation precludes the active party from inferring individual sample embeddings and reduces the sensitivity of the average embeddings for improved utility in DP. According to Definition 3, each epoch of Algorithm 1 satisfies (ϵ, δ)-DP. Inspired by [6], we quantify the cumulative privacy loss over T epochs via moment accountant [1].

Theorem 3. Given each epoch of Algorithm 2 satisfies (ϵ, δ) -DP, there exists constants r_1 and r_2 such that given sampling probability q and epoch number T, and $\epsilon < r_1q\sqrt{T}$, Algorithm 2 satisfies (ϵ', δ) -DP, with $\epsilon' = r_2q\sqrt{T}\epsilon$ over T epochs.

Proof. According to Definition 3 and Theorem 1, to ensure one iteration $(\epsilon, \delta) - DP$, we set $\delta = \frac{\sqrt{2 \ln 1.25/\delta}}{\epsilon}$. By Theorem 1 in [6], with the appropriate choice of ϵ , q, T, such that $\epsilon < r_1 q \sqrt{T}$, the privacy loss is $\epsilon' = r_2 q \sqrt{T} \epsilon$ over T epochs. \Box

2) Repetitive Random Model Initialization: VFDC's strategy of continuously reinitializing the feature extractor ψ_{ϑ} further enhances feature privacy. The continuously initialized feature extractor makes it challenging to reconstruct the model parameters as well as the real features from embeddings. In summary, VFDC's multi-faceted approach effectively protects the sample-wise feature privacy of real data in VFL settings.

5 Experimental Study

In experimental studies, we study the following research questions: **RQ1**: What is the visual quality of VFDC-generated dataset? **RQ2**: What is the *privacyutility trade-off* of VFDC compared to other methods on different datasets? **RQ3**: How does VFDC improve training efficiency compared to other methods? **RQ4**: What impact do hyperparameters have on the proposed VFDC approach?

5.1 Experimental Setup

Datasets. We evaluate our proposed VFDC approach on six widely used realworld public datasets including three image datasets and three tabular datasets as follows: 1) **MNIST**¹ is a dataset of handwritten digits with 70,000 images from 10 classes with size of $1 \times 28 \times 28$ pixels. 2) **FMNIST**² is a fashion image dataset containing 70,000, 28×28 , gray-scale fashion items from 10 classes. 3) **CIFAR10**³ consists of 60,000, 32×32 colored images from 10 classes, with 6,000 images per class. 4) **MIMIC-III** [15] is an in-hospital mortality prediction dataset based on the initial 48 h of ICU data, containing 714 features and 21139 samples. 5) **WDBC**⁴ dataset is utilized for breast cancer diagnosis. It comprises 569 samples, with 212 malignant and 357 benign cases. Featuring 30 attributes per sample. 6) **Spambase**⁵ dataset is collected to determine whether a given email is spam or not. It contains a total of 4601 records, each with 57 characteristics.



Fig. 3. Visualization of the VFDC-generated synthetic images on three datasets.

Implementation Details. For dataset generation, we employ the SGD optimizer, iterating over 200 epochs to assess the performance of the synthesized dataset. In evaluating the VFDC-generated data, each model is trained 20 times with various initialization. We utilize average accuracy as the utility metric, and also report the standard deviation. For image datasets, a Convolutional Neural Network (CovNet) is used, while a multilayer perceptron (MLP) is employed for the tabular datasets. For privacy protection, we set d = 256, $\delta = 1e^{-4}$, r = 1, and $\epsilon \in \{10, 50, 100, \infty\}$. We set the bit length of secret shared values to 32.

Compared Methods. Our evaluation of VFDC includes comparisons with various dataset size reduction techniques in VFL: 1) Vanilla-DC: This method, based on the MMD-based DC approach [35], is applied to VFL by transmitting

¹ http://yann.lecun.com/exdb/mnist/.

 $^{^{2}\} https://github.com/zalandoresearch/fashion-mnist.$

³ https://www.cs.toronto.edu/~kriz/cifar.html.

⁴ https://archive.ics.uci.edu/dataset/17.

⁵ https://archive.ics.uci.edu/dataset/94/spambase.



Fig. 4. Visualization of the distributions of the original dataset and the condensed dataset by VFDC on two datasets using t-SNE. The star-shaped points denote the VFDC-generated samples.

Table 1. Comparison of accuracy (utility) of models trained by datasets generated via different methods. ϵ denotes the DP budget, Spc denotes the number of samples per class, and Rat is the ratio of dataset size. V-DC denotes Vanilla-DC. The highest results under the same ϵ are highlighted in bold.

DP $\epsilon \downarrow$	Spc	Rat.%	Coreset-LDP		DC-LDP		(Ours) VFDC			V-DC	Whole		
			10	50	100	10	50	100	10	50	100	∞	∞
FMNIST	50	0.83	78.3	78.8	79.8	74.0	82.1	84.4	88.2	88.3	88.3	88.5	93.5
	100	1.7	80.7	81.6	81.9	74.2	84.0	84.8	88.9	89.0	89.0	89.1	
	200	3.3	81.8	82.6	83.1	75.4	84.3	85.0	89.4	89.6	89.7	89.8	
CIFAR	50	1.0	37.2	39.4	41.1	38.1	49.0	59.8	62.1	63.0	63.3	63.4	84.8
	100	2.0	40.3	44.5	50.9	37.6	50.4	60.9	64.1	64.1	64.5	65.3	
	200	4.0	52.3	54.5	56.3	37.0	50.7	61.3	66.9	67.0	67.5	67.9	
MNIST	50	0.83	90.1	92.1	93.4	91.3	95.6	96.7	98.3	98.3	98.4	98.4	99.6
	100	1.7	90.2	92.3	93.5	91.5	95.6	96.9	98.6	98.6	98.7	98.6	
	200	3.3	91.3	93.0	93.3	92.0	95.8	97.0	98.7	98.7	98.7	98.7	
MIMIC	10	0.11	63.6	70.8	70.8	41.2	52.2	71.4	78.2	78.4	78.9	79.5	81.7
	30	0.34	64.2	71.6	70.3	53.6	67.1	70.9	78.4	78.9	79.1	80.3	
	50	0.56	65.6	70.4	71.9	59.1	69.3	72.3	78.8	79.3	79.9	80.4	
Spambase	10	0.48	63.4	72.7	80.2	58.6	63.4	82.8	89.7	90.1	90.5	90.7	94.2
	30	1.45	65.8	74.3	82.6	57.2	65.9	81.8	90.1	90.5	91.0	91.3	
	50	2.42	70.1	75.2	83.8	56.3	69.1	82.7	90.9	91.2	91.5	91.7	
WDBC	10	4	89.4	90.4	92.5	86.1	91.8	92.9	97.0	97.1	97.1	97.2	97.3
	20	8	89.7	90.8	92.1	88.7	91.8	93.2	97.0	97.1	97.2	97.2	
	30	12	90.6	91.0	93.3	89.9	93.2	93.9	97.1	97.2	97.3	97.3	

real data embeddings without any protection. 2) **Coreset-LDP** [14]: Implements a coreset sampling approach in VFL. We employ local DP (LDP) [18] to protect the sample-wise feature privacy of the chosen real data samples. 3) **DC-LDP**: Enhances Vanilla-DC by incorporating LDP for sample-wise data privacy. 4) **Whole**: This method involves training the VFL model on the entire dataset, foregoing any form of data reduction.

5.2 Visualization of Condensed Dataset

To address RQ1, Fig. 3 visualizes the VFDC-generated synthetic datasets using three image datasets. These images are synthesized from features initialized randomly by the passive party and labels assigned arbitrarily by the active party, ensuring no direct representation of any specific sample from the original dataset. Figure 4 visualizes the data distributions of VFDC-generated data on two datasets Spambase and WDBC by using t-SNE for dimensionality reduction. Notably, even under the stringent conditions of our mixed protection mechanism—combining secure aggregation, differential privacy (DP), and repetitive random initialization—VFDC accurately reflects the original datasets' distributions. This demonstrates the effectiveness of VFDC in maintaining visual quality and data representation fidelity in a privacy-preserving VFL setting.

5.3 Performance Comparison

We validate the efficacy of our VFDC approach by comparing with the baseline methods to answer RQ2. Table 1 presents the accuracy of models trained on datasets generated by various methods across six datasets. VFDC, when benchmarked against the Vanilla-DC baseline, exhibits comparable accuracy, particularly under larger DP budgets. This suggests that secure aggregation has a minimal impact on model performance. Furthermore, VFDC consistently outperforms Coreset-LDP and DC-LDP in accuracy for the same DP budgets. Notably, under smaller privacy budgets (ϵ), VFDC's accuracy significantly exceeds that of Coreset-LDP and DC-LDP, thanks to the class-wise secure aggregation's ability to apply minimal DP noise while providing strong privacy protection. We also observe an increase in the condensed dataset's performance relative to the growth ratio of dataset size. In most cases, the performance of the condensed dataset closely mirrors that of the complete dataset, albeit with a substantially reduced data size. In conclusion, VFDC demonstrates the most favorable privacy-utility trade-off among the compared methods across various datasets, DP budgets, and data size ratios.

5.4 Efficiency Improvement

Given that communication overhead per iteration is identical for both synthetic and real data, we assess training efficiency through a comparison of iteration counts needed to reach equivalent accuracy levels. Figure 5 reveals that models trained on datasets generated by VFDC converge markedly faster (approximately 10 times) compared to those trained on complete datasets. Additionally, VFDC-trained models attain higher accuracy than models trained with other baseline methods. These findings confirm that the VFDC-generated condensed dataset substantially enhances training efficiency.



Fig. 5. Performance variation of the VFL model trained using dataset output by different methods across iterations.

Table 2. Impact of secret sharing bit length on accuracy and data volume each iteration.

 Plain-text denotes the Vanilla-DC baseline.

Dataset	Bits	Accuracy (%)	Data Volume (MB)			
MIMIC	Plain-text	80.1 ± 1.1	2.05			
	32	79.9 ± 1.2	8.24			
	16	79.2 ± 1.3	4.12			
Spambase	Plain-text	91.7 ± 0.7	2.05			
	32	91.5 ± 0.7	8.24			
	16	90.8 ± 1.3	4.12			

5.5 Impact of Hyperparameters

Impact of Secret Sharing Bit Length. Secure aggregation impacts the utility and communication cost via the bit length of secret-shared values, as it maps float values to *l*-bit fixed point integers. Table 2 shows that 32-bit values in VFDC offer comparable accuracy to the plain-text Vanilla-DC baseline with a moderate increase in data volume (8.24 MB vs. 2.05 MB). Reducing to 16-bit decreases accuracy slightly but halves the data volume to 4.12 MB. These results confirm that our VFDC approach's secure computation achieves similar performance to non-encrypted methods with manageable communication overhead.

Impact of Privacy Budget ϵ . To access the impact of DP privacy budget ϵ , we change ϵ in range {10, 50, 100}, As demonstrated in Table 1, VFDC showcases a notably lower decline in accuracy with decreasing privacy budgets while maintaining equivalent levels of DP protection compared to DC-LDP. This is attributed to the reduced sensitivity achieved through our proposed class-wise secure aggregation. Thereby our VFDC facilitates privacy protection with minimal impact on model accuracy.

Impact of VFDC Iteration Number. Figure 6 demonstrates the variation of model accuracy across iterations during the VFDC process for different data



Fig. 6. Performance variation of the VFL model trained on VFDC-generated datasets during the VFDC process.

sizes. It reveals the trade-off between synthetic data utility and the VFDC generation cost. We observe that it only takes 30 (for MIMIC) or 10 (for Spambase) iterations to generate synthetic datasets with high utility, indicating that VFDC is efficient in generating high-quality synthetic data especially for smaller datasets.

6 Conclusion and Future Directions

To address the dual challenges of privacy and efficiency in VFL, we present VFDC for small synthetic dataset generation. VFDC facilitates collaborative, privacy-preserving generation of vertically-partitioned condensed dataset between two parties, one with labels and the other with features of overlapping samples. To securely align the distributions of synthetic and real data, we design a tailored mixed protection mechanism that integrates class-wise secure aggregation, DP and repetitive random initialization. Experimental results show that VFDC not only boosts training efficiency, crucial for multi-trail training and cryptography-based training, but also improves sample-level data privacy with high utility.

Our work pioneers the integration of dataset condensation into VFL, leaving several future directions. Firstly, our VFDC is tailored for distribution matchingbased DC method, leaving the exploration of other DC methods within VFL relatively untapped. Another direction is to expand the framework to accommodate multi-party settings.

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Neighborhood Component Feature Selection for Multiple Instance Learning Paradigm

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Abstract. In a multiple instance learning (MIL) scenario, the outcome annotation is usually only reported at the bag level. Considering simplicity and convergence criteria, the lazy learning approach, i.e., k-nearest neighbors (kNN), plays a crucial role in predicting bag labels in the MIL domain. Notably, two variations of the kNN algorithm tailored to the MIL framework have been introduced, namely Bayesian-kNN (BkNN) and Citation-kNN (CkNN). These adaptations leverage the Hausdorff metric along with Bayesian or citation approaches. However, neither BkNN nor CkNN explicitly integrates feature selection methodologies, and when irrelevant and redundant features are present, the model's generalization decreases. In the single-instance learning scenario, to overcome this limitation of kNN, a feature weighting algorithm named Neighborhood Component Feature Selection (NCFS) is often applied to find the optimal degree of influence of each feature. To address the significant gap existing in the literature, we introduce the NCFS method for the MIL framework. The proposed methodologies, i.e. NCFS-BkNN, NCFS-CkNN, and NCFS-Bayesian Citation-kNN (NCFS-BCkNN), learn the optimal features weighting vector by minimizing the regularized leaveone-out error of the training bags. Hence, the prediction of unseen bags is computed by combining the Bayesian and citation approaches based on the minimum optimally weighted Hausdorff distance. Through experiments with various benchmark MIL datasets in the biomedical informatics and affective computing fields, we provide statistical evidence that the proposed methods outperform state-of-the-art MIL algorithms that do not employ any a priori feature weighting strategy.

Keywords: Multiple Instance Learning \cdot Neighborhood Component Feature Analysis \cdot Nearest Neighbors \cdot Feature Selection

1 Introduction

The multiple instance learning (MIL) frameworks have gained considerable popularity in the machine learning (ML) research field due to their broad applicability to several real-world applications and problems, such as the prediction of structure-activity relationships, document categorization, image classification, economic risk assessment and the prediction of protein binding sites [2,7]. In the classical (supervised) single-instance learning (SIL) setting, each example is an instance represented by a feature vector and an individual label. In contrast, in the MIL paradigm, the learner receives a set of *bags* containing multiple instances and corresponding bag labels but not always single instance labels. Hence, MIL aims to train an instance or a bag classifier, and in this paper, we mainly focus on predicting unseen bags.

The SIL algorithms can be roughly divided into eager learning and lazy learning [17]. The goal of eager learning is to construct a general, explicit, inputindependent target function learned during the training stage (e.g., Fisher's discriminant, logistic regression, decision tree, support vector machine, artificial neural networks, etc.), while the lazy learning approach stores the training examples for future use. The most widely used lazy learning algorithm is the k-nearest neighbor (kNN), which classifies examples according to the class of the nearest neighbors. Although the kNN algorithm is one of the simplest ML algorithms, it often yields competitive results compared with the state-of-the-art classification models [1]. Its appeal lies from the fact that (i) its decision surfaces are nonlinear, (ii) there are few hyperparameters (i.e., distance metric and number of neighbors) to be tuned, and (iii) the expected reliability of prediction improves as the data of training set increases [5,22].

In [26], authors investigated the lazy learning approach in multiple instance problems, suggesting the use of the Hausdorff metric. Two extensions of the kNN algorithm to the MIL framework were proposed: the *Bayesian-kNN* (BkNN) and the *Citation-kNN* (CkNN) algorithms. The former aims to maximize the posterior probability for the unseen bag, while the concept of citations inspires the latter. To measure the distance between bags, the authors proposed a variant of the Hausdorff distance, i.e., the *minimal Hausdorff distance*, which proved to be more robust against outlier points within the bags and was used for both the BkNN and CkNN methods. However, CkNN still applies the most straightforward majority vote approach among the references and citers to classify unseen bags. An improvement of this algorithm was proposed by [8], where the authors, applying a Bayesian approach to references and a distance-weighted majority vote approach to citers, introduced the *Bayesian Citation-kNN* (BCkNN).

The kNN algorithm, with standard distance metrics (e.g., Euclidean, city block, cosine, etc.), operates under the implicit assumption that all features hold equal significance. However, this assumption is problematic when irrelevant or redundant features are present, thereby impacting the neighborhood search and diminishing the model's generalizability [15,26]. To solve this problem, a feature weighting algorithm is often applied to find each training set feature's optimal degree of influence. In particular, authors in [29] proposed a nearest neighborhood.

based feature selection method named *neighborhood component feature selection* (NCFS). This work was inspired by the dimensionality reduction technique based on neighborhood component analysis proposed in [5]. Their embedded algorithm [29] learns a feature weighting vector via gradient ascent, maximizing the regularized leave-one-out classification accuracy. This assigns higher weights to relevant features, while irrelevant ones approach zero weight.

Feature weighting can also be used as a feature selection approach to discard features with weights below a specific threshold value, thereby reducing the model complexity and the number of dimensions. Recent state-of-the-art works demonstrated how NCFS [10, 28] is a trade-off between simplicity and generalization performance, thus leading to insensitivity to the increase in the number of irrelevant features. Therefore, the NCFS approach leads to improving both the generalization performance and the interpretability of the kNN model [29], and was successfully applied in different domains ranging from industry [28], affective computing [23], and biomedical informatics [24]. While the NCFS technique is often combined with the SIL kNN model to enhance the generalization performance when dealing with high-dimensional data, a notable gap in the literature regarding its integration within the MIL framework remains. Additionally, the literature lacks explicitly integrated feature selection methodologies in Bayesian-kNN and Citation-kNN approaches, which can simultaneously discard redundant and irrelevant features and handle the weakly supervised setting of the MIL framework.

To accomplish the goal mentioned above, we introduce the NCFS for the MIL framework. The proposed methodologies, NCFS-BkNN, NCFS-CkNN, and NCFS-BCkNN, learn the optimal features weighting vector by minimizing the regularized leave-one-out error on the training bags. Accordingly, the prediction of unseen bags was computed by combining the Bayesian and citation approaches with the optimal weighted minimal Hausdorff distance. The main contributions of this work to the existing ML methodologies are (i) the introduction of the NCFS algorithm for the MIL lazy learning approaches and (ii) the demonstration of the effectiveness of the proposed approaches to different benchmark MIL datasets in the biomedical informatics (Musk dataset [4]) and affective computing (DEAP dataset [11]) domains. The experiments revealed the significant enhancement brought by the introduction of NCFS in the MIL domain, showing the statistical superiority of our proposed approaches over state-of-the-art MIL algorithms. As a result, our research contributes to advancing the state-of-the-art of MIL lazy learning algorithms.

The paper is organized as follows: in Sect. 2, we describe the proposed methodology. Section 3 discusses how the approach is evaluated on the benchmark datasets. The experimental procedure is reported in Sect. 4, while results are presented in Sect. 5. Finally, Sect. 6 provides conclusions of our findings.

2 Methods

We defined the following notation to formulate the MIL problem:

- $\{B_1, \ldots, B_n, \ldots, B_N\}$ is the set of the N training bags;
- $\{b_1, \ldots, b_t, \ldots, b_T\}$ is the set of the T test bags;
- $-y_{B_n}, y_{b_t} \in \{1, \ldots, C\}$ are the labels for the *n*-th training bag and the *t*-th test bag, respectively, where C is the number of classes;
- L_{B_n} and L_{b_t} are the total number of instances of the *n*-th training and *t*-th test bags;
- $B_n = \{x_{B_n}^{(1)}, \dots, x_{B_n}^{(L_{B_n})}\}$ and $b_t = \{x_{b_t}^{(1)}, \dots, x_{b_t}^{(L_{b_t})}\}$ are the sets of instances of the *n*-th training and *t*-th test bags, with $x_{B_n}, x_{b_t} \in \mathbb{R}^d$, where *d* is the number of features.

In the MIL paradigm, each bag is allowed to have a different size, meaning that L can vary among bags [6].

2.1 The Lazy Learning Approach for Multiple Instance Learning Setting

Bayesian-kNN and Citation-kNN. To address the lazy learning approach for the MIL paradigm, two problems should be addressed: (i) the distance measure problem and (ii) the classification problem. The former aims to formulate different distance functions measuring the similarity between training and test bags. In this context, we employed the minimal Hausdorff distance [8,26] to measure the distance between unseen test bags and training bags as follows:

$$D^{h}(b_{t}, B_{n}) = \min_{x_{b_{t}} \in b_{t}} \min_{x_{B_{n}} \in B_{n}} d(x_{b_{t}}, x_{B_{n}}),$$
(1)

where $|b_t| = L_{b_t}$, $|B_n| = L_{B_n}$ and $d(x_{b_t}, x_{B_n})$ can be the Euclidean ($||x_{b_t} - x_{B_n}||_2$) or city block distance ($||x_{b_t} - x_{B_n}||_1$). Several works [8, 12, 16, 20] demonstrated how this modified (minimal) Hausdorff distance enhances robustness to noise and outliers, mitigating their adverse influence on both feature weighting and decision-making of our distance learning-based approach. Therefore, the modified Hausdorff distance was used to adapt kNN to MIL problems.

Concerning the classification problem, in spite of the straightforward majority voting approach, we employed two alternative methods, the Bayesian-kNN (BkNN) and Citation-kNN (CkNN). In particular, for the BkNN method, the prediction of unseen bag b_t is computed by:

$$\hat{y}_{b_{tBkNN}} = \underset{y' \in \{1, \dots, C\}}{\arg \max} P(y') P(\{y_r^{(1)}, y_r^{(2)}, \dots, y_r^{(k)}\} | y'),$$
(2)

where $\{y_r^{(1)}, y_r^{(2)}, \ldots, y_r^{(k)}\}$ are class labels of k closest training bags (i.e., *references*) computed according to the minimal Hausdorff distance defined in Eq. (1), P(y') is the probability that y' assumes one of the possible C classes, and

 $P(\{y_r^{(1)}, y_r^{(2)}, \ldots, y_r^{(k)}\}|y')$ is the conditional probability of observing the set of class labels of the k closest training bags given y'. It is worth noting that given the Bayes theorem, the posterior probability $P(y')P(\{y_r^{(1)}, y_r^{(2)}, \ldots, y_r^{(k)}\}|y')$ can be computed by counting the number of references which disclose y' class divided by the total number of references (i.e., k). Notice that each reference represents a different training bag.



Fig. 1. Schematic representation of Citation-kNN. (**A**) Bags distribution. (**B**) The two closest references of Bag 2 (circle) (i.e., 2-nearest neighbors highlighted by the arrows) are {Bag 1 (square), Bag 3 (triangle)}. (**C**) Citers of Bag 2 (circle) are {Bag 1 (square), Bag 4 (rhombus), Bag 5 (pentagon)} as Bag 2 is contained in their 2-nearest neighbors.

For the CkNN, the combination of references and citers defines the prediction criteria. In particular, the CkNN extends the set of nearest neighbors, including not only the closest bags (i.e., *references*) but also the bags for which the unseen bag is among the closest bags (i.e., *citers*). Given an unseen bag b_t , $\{r_1, r_2, \ldots, r_k\}$ are its k-nearest references and $\{c_1, c_2, \ldots, c_q\}$ are its q-nearest citers. Then, the test bag is predicted by:

$$\hat{y}_{b_{tCkNN}} = \underset{y' \in \{1, \dots, C\}}{\arg\max} \left(\sum_{i=1}^{k} \delta(y', y_r^{(i)}) + \sum_{j=1}^{q} \delta(y', y_c^{(j)}) \right), \tag{3}$$

where $\{y_r^{(1)}, y_r^{(2)}, \ldots, y_r^{(k)}\}$ and $\{y_c^{(1)}, y_c^{(2)}, \ldots, y_c^{(q)}\}$ are, respectively, the class labels of k references and q citers computed according to the minimal Hausdorff distance defined in Eq. (1), and δ is the Kronecker delta function. Figure 1 shows a graphical representation of the computation of the 2-nearest references (Fig. 1B) and 2-nearest citers (Fig. 1C) of a particular bag.

Bayesian Citation-kNN. The natural way to improve the prediction of CkNN is to assign different weights to different references and citers according to their distances to the unseen bag. We used the algorithm proposed in [8] named Bayesian Citation-kNN (BCkNN) by combining both the Bayesian and the distance weighting approaches to its k references and its q citers, respectively. Then, the equation to classify an unseen bag becomes:

$$\hat{y}_{b_{tBCkNN}} = \underset{y' \in \{1, \dots, C\}}{\arg \max} \left(P(y') P(y_r^{(1)}, y_r^{(2)}, \dots, y_r^{(k)} | y) + \frac{\sum_{j=1}^q f(b_t, c_j) \delta(y', y_c^{(j)})}{\sum_{j=1}^q f(b_t, c_j)} \right),$$
(4)

where $f(b_t, c_j)$ are the citer weights that can be defined by one of the following functions [8]:

$$f(b_t, c_j) = e^{-(D^h(b_t, c_j))^2},$$
(5)

$$f(b_t, c_j) = \frac{1}{1 + D^h(b_t, c_j)},$$
(6)

$$f(b_t, c_j) = \frac{1}{1 + (D^h(b_t, c_j))^2},$$
(7)

where $D^{h}(b_{t}, c_{j})$ represents the minimal Hausdorff distance between the unseen bag b_{t} and the *j*-th citers defined as follows:

$$D^{h}(b_{t}, c_{j}) = \min_{x_{b_{t}} \in b_{t}} d(x_{b_{t}}, c_{j}).$$
(8)

Notice how citer contribution is weighted according to the inverse of their distances, assigning a lower weight to citers that are further from the unseen bag b_t .

2.2 Neighborhood Component Feature Selection for Single Instance Learning Setting

The original formulation of neighborhood components analysis (NCA) was introduced in [5], while the adaptation to feature selection, i.e., neighborhood component feature selection (NCFS), was proposed in [29]. In the SIL framework, we let $T = \{(\boldsymbol{x}_1, t_1), \ldots, (\boldsymbol{x}_i, t_i), \ldots, (\boldsymbol{x}_N, t_N)\}$ be a set of training samples, where \boldsymbol{x}_i is a *d*-dimensional feature vector, $t_i \in \{1, \ldots, C\}$ is its corresponding class label and N is the number of samples. We denote the weighted distance between two samples \boldsymbol{x}_i and \boldsymbol{x}_j by $D_{\boldsymbol{\omega}}(\boldsymbol{x}_i, \boldsymbol{x}_j) = \sum_{l=1}^d \omega_l^2 |x_{il} - x_{jl}|$, where ω_l is a weight associated with the *l*-th feature. Here, the probability of \boldsymbol{x}_i selecting \boldsymbol{x}_j as its reference point is defined as:

$$p_{ij} = \begin{cases} \frac{\kappa(D_{\omega}\left(x_{i}, x_{j}\right))}{\sum_{k \neq i} \kappa(D_{\omega}\left(x_{i}, x_{k}\right))}, & \text{if } i \neq j \\ 0, & \text{if } i = j \end{cases},$$
(9)

where $\kappa(z) = e^{(-z/\sigma)}$ is a kernel function and the kernel width σ is a hyperparameter that controls the prior probability of each point being selected as reference ($\sigma \to 0$ means that only the nearest neighbor can be chosen, while $\sigma \to \infty$ implies that all the points have the same chance to be selected). Based on the above definition, the probability of the query point \boldsymbol{x}_i being correctly classified is given by $p_i = \sum_j t_{ij} p_{ij}$, where $t_{ij} = 1$ if and only if $t_i = t_j$ and $t_{ij} = 0$ otherwise. The L2-norm regularized approximate leave-one-out classification accuracy can be written as:

$$\xi(\boldsymbol{\omega})_{SIL} = \sum_{i} \sum_{j} t_{ij} p_{ij} - \lambda \sum_{l=1}^{d} \omega_l^2, \qquad (10)$$

where λ is a scalar regularization hyperparameter controlling model complexity. Since the objective function $\xi(\boldsymbol{\omega})$ is differentiable, its derivative with respect to ω_l can be computed as:

$$\frac{\partial \xi(\omega)_{SIL}}{\partial \omega_l} = 2 \left(\frac{1}{\sigma} \sum_i \left(p_i \sum_{j \neq i} p_{ij} |x_{il} - x_{jl}| - \sum_j t_{ij} p_{ij} |x_{il} - x_{jl}| \right) - \lambda \right) \omega_l.$$
(11)

2.3 Our Proposal: Neighborhood Component Feature Selection for the Multiple Instance Learning Setting

The proposed NCFS formulation for MIL aims to find the k references and q citers computing the weighted minimal Hausdorff distance. We introduced the weighted minimal Hausdorff distance between two training bags B_r and B_s as follows:

$$D^{h}_{\omega}(B_{r}, B_{s}) = \min_{x_{B_{r}} \in B_{r}} \min_{x_{B_{s}} \in B_{s}} \sum_{l=1}^{a} \omega_{l}^{2} d_{l}(x_{B_{r}, l}, x_{B_{s}, l}),$$
(12)

where ω_l is a weight associated with the *l*-th feature and d_l is the pointwise distance between the *l*-th predictor of instances x_{B_r} and x_{B_s} . Equation 12 can model a different number of instances for each bag by imposing $|B_r| = L_{B_r}$ and $|B_s| = L_{B_s}$ where L_{B_r} and L_{B_s} represent the number of instances within the *r*th and *s*-th training bags, respectively. The optimal *d*-dimensional weight vector ω is found by minimizing the leave-one-out classification error of the training bags. In the MIL paradigm, the probability of B_r selecting B_s as its reference bag is defined as:

$$p_{rs} = \begin{cases} \frac{\kappa(D^h_{\omega}(B_r, B_s))}{\sum_{k \neq r} \kappa(D^h_{\omega}(B_r, B_k))}, & \text{if } r \neq s\\ 0, & \text{if } r = s \end{cases}.$$
(13)

The probability of the query bag B_r being correctly classified is given by:

$$p_r = \sum_s y_{B_r} y_{B_s} p_{rs},\tag{14}$$

where $y_{B_r}y_{B_s} = 1$ if and only if $y_{B_r} = y_{B_s}$ and $y_{B_r}y_{B_s} = 0$ otherwise. The regularized leave-one-out classification accuracy can be written as:

$$\xi(\boldsymbol{\omega})_{MIL} = \sum_{r} \sum_{s} y_{B_r} y_{B_s} p_{rs} - \lambda \sum_{l=1}^{d} \omega_l^2.$$
(15)
Then, we aim to minimize the following criteria:

$$\min_{\omega} \xi(\omega)_{MIL} \tag{16}$$

and the derivative of the objective function becomes:

$$\frac{\partial \xi(\boldsymbol{\omega})_{MIL}}{\partial \omega_l} = 2 \left(\frac{1}{\sigma} \sum_r \left(p_r \sum_{s \neq r} p_{rs} D_l^h(B_r, B_s) + -\sum_s y_{B_r} y_{B_s} p_{rs} D_l^h(B_r, B_s) \right) - \lambda \right) \omega_l.$$
(17)

Here, D_l^h is the minimal Hausdorff distance computed for the *l*-th feature as follows: $D_l^h(B_r, B_s) = \min_{x_{B_r} \in B_r} \min_{x_{B_s} \in B_s} d_l(x_{B_r,l}, x_{B_s,l})$. To solve the optimization problem formulated in Eq. (16), we employed the quasi-Newton method for large-scale optimization, namely the Limited memory Broyden-Fletcher-Goldfarb-Shannon (L-BFGS) method [13]. Compared to the Stochastic Gradient Descent (SGD) applied in [29] for solving NCFS, L-BFGS has generally proven to be more stable for solving the feature selection tasks [18]. Notice how the optimal *d*-dimensional weight vector ω^* , computed by minimizing $\xi(\omega)_{MIL}$ via L-BFGS through Eq. (16), can be used to compute the optimal weighted minimal Hausdorff distance to measure the distance between unseen test bag and training bags as follows:

$$D^{h}_{\omega^{*}}(b_{t}, B_{n}) = \min_{x_{b_{t}} \in b_{t}} \min_{x_{B_{n}} \in B_{n}} \sum_{l=1}^{d} \omega_{l}^{*^{2}} d_{l}(x_{b_{t}, l}, x_{B_{n}, l}).$$
(18)

This weighted distance can now be used to compute references and citers for BkNN (Eq. 2), CkNN (Eq. 3), and BCkNN (Eq. 4), thus giving rise to NCFS-BkNN, NCFS-CkNN, and NCFS-BCkNN. For the NCFS-BCkNN, the citer weights defined in Eqs. (5), (6), and (7) are computed according to the weighted minimal Hausdorff distance between the unseen bag b_t and the *j*-th citers defined as follows:

$$D^{h}_{\omega^{*}}(b_{t},c_{j}) = \min_{x_{b_{t}} \in b_{t}} \sum_{l=1}^{a} \omega_{l}^{*^{2}} d_{l}(x_{b_{t},l},c_{j,l}).$$
(19)

Importantly, Eq. (18) allows the modeling of a different number of instances for each bag by imposing $|B_n| = L_{B_n}$ and $|b_t| = L_{b_t}$, where L_{B_n} and L_{b_t} , represent the number of instances within the *n*-th training and *t*-th test bags, respectively. This allows us to compute the weighted minimal Hausdorff distance on the training (Eq. 12) and test set (Eq. 18) between bags of different sizes.

Algorithm 1 shows the overall pseudocode of the proposed methodology. For each unseen test bag, our proposed NCFS-BCkNN extension first finds its kreferences and q citers using the optimal weighted minimal Hausdorff distance. Then, a Bayesian approach is applied to its k references and a distance-weighted majority vote approach is applied to its q citers. It is noteworthy that the pseudocode can be easily generalized for implementing NCFS-BkNN by setting q = 0 and $f(b_t, c_j) = 0$. For the NCFS-CkNN, in addition to setting $f(b_t, c_j) = 0$, the majority vote of the k references should replace the posterior probability computation. Source code is available at https://github.com/g-turri/MIL-NCFS.

3 Datasets

In this section, we describe the employed datasets. In the Musk dataset (see Sect. 3.1), each bag represents a molecule characterized by a different number of conformers (i.e., the bag's instances, ranging from a minimum of 2 to a maximum of 40 instances). In the DEAP dataset (see Sect. 3.2), each bag corresponds to a physiological signal segmented into 5 windows, each associated with a distinct emotional video, thus forming a bag containing 5 instances.

NCFS-BCkNN

- 1: Splitting data in train_data and test_data
- 2: **training**(train_data)
- 3: return $\boldsymbol{\omega}^*$, the optimal weighting vector
- 4: testing(train_data, test_data, ω^* , k, q, f)
- 5: return \hat{y} , the vector with the predicted bag labels

training(train_data, λ , σ)

- 6: for each training bag B_r and B_s pair do
- 7: Compute the weighted Minimal Hausdorff distance $D^h_{\omega}(B_r, B_s)$ using ω according to Eq. (12)
- 8: Compute p_r and p_{rs} according to Eq. (13) and (14)
- 9: Minimize the objective function $\xi(\boldsymbol{\omega})_{MIL}$ (Eq. 16) using the **L-BFGS** Solver: compute the derivative $\frac{\partial \xi(\boldsymbol{\omega})_{MIL}}{\partial \omega_l}$ (Eq. 17) to identify the direction of steepest descent and to compute an estimate of the Hessian matrix
- 10: **return** the optimal weighting vector $\boldsymbol{\omega}^*$

testing(train_data, test_data, $\boldsymbol{\omega}^*, k, q, f$)

- 11: BCkNN(train_data, test_data, ω^* , k, q, f, distance_metric)
- 12: Compute the weighted Minimal Hausdorff (Eq. 18) distance between each train and test bag.
- 13: Compute the posterior probability of the k references (first term of Eq. 4)
- 14: Compute the majority vote of the q citers (second term of Eq. 4) normalized according to one of the following functions: (5), (6), or (7)
- 15: Classify the unseen bag b_t using Eq. (4)
- 16: **return** \hat{y} , a prediction vector with a length equal to the number of unseen bags in the test set

3.1 Musk Dataset

A classic benchmark dataset for evaluating MIL algorithms is the Musk dataset [4] sourced from the bioinformatics field. The main task involves predicting

whether a molecule has a musky smell. Each molecule in the dataset is characterized by a set of 166 features describing its geometric shape (or conformation). Given the capacity of chemical bonds to rotate, a single molecule can adopt multiple distinct shapes, each referred to as a conformer. The conformers dictate the properties of a molecule, including its scent. Each bag within the dataset corresponds to a molecule, while each instance pertains to one of its conformers. A molecule belongs to the (positive) musky class if at least one of its conformers possesses the potential to smell musky; otherwise, it is labeled as negative. In our experiment, we considered the Musk 1 dataset composed of 92 molecules, 47 of which were classified as musky. It is worth noting that in the Musk dataset, each bag (molecule) can be represented by a different number of instances (conformations).

3.2 DEAP Dataset

We further test our methodology in the affective computing scenario, employing the DEAP dataset [11]. In the DEAP experiment [11], 32 healthy participants were asked to watch 40 music videos that lasted 1 min each. The 40 videos were presented in 40 trials. Following each video presentation, participants were prompted to self-report their emotional experiences across four dimensions: valence, arousal, dominance, and liking. Ratings for valence, arousal, and dominance ranged from 1 to 9 and were obtained using the Self-Assessment Manikin. Concurrently, physiological data were collected, including full-scalp EEG and thirteen peripheral signals. These signals comprised galvanic skin response, respiration amplitude, skin temperature, blood volume pressure, electromyograms of zygomaticus and trapezius muscles, and electrooculogram, all sampled at a frequency rate of 512 Hz.

The main challenge of this dataset is to predict participants' emotional state based on their physiological data. Several ML and Deep Learning models were presented to solve this classification task [9]. However, this dataset shows sparse annotations of self-reported emotional responses. Then, in the context of sparse labeling, MIL approaches can be employed to solve this task.

Previous studies on this dataset [11,19] confirmed how the MIL approach might overcome standard supervised learning approaches for predicting arousal and valence tasks. Consequently, in our study, we modeled the single bag as a set of multiple temporal instances represented by 34 extracted predictors. In particular, each synchronously recorded physiological signal was segmented into 5 windows, thus resulting in 5 instances for each video (bag). Our analysis focused on two binary classification tasks: discerning between (i) low/high arousal levels and (ii) negative/positive valence levels.

4 Experimental Procedure

This section outlines the cross-validation (CV) procedure and hyperparameter tuning conducted for each dataset. The hyperparameters λ and σ of the NCFS

algorithm were tuned using a nested 5-fold CV for the Musk dataset and an inner leave-one-video-out (LOVO) CV procedure for the DEAP dataset. The optimal values for λ and σ were identified through a grid search where λ was selected from the set {0, 0.01, 0.05, 0.10, 0.50, 1} and σ was chosen from a range of 5 logarithmically spaced values between 10^{-3} and 10^{1} .

The mode of the optimal-selected hyperparameters was $\lambda = 0$ and $\sigma = 1$ for Musk dataset, $\lambda = 0.01$ and $\sigma = 1$ for the arousal task of DEAP dataset, and $\lambda = 0.05$ and $\sigma = 1$ for the valence task of DEAP dataset. We fixed λ and σ according to the results mentioned above for computational reasons.

We tested the BkNN, CkNN, and BCkNN with values of k ranging from 1 to 5. For the CkNN and BCkNN, the number of citers q was empirically set to q = k+2 following [8,26], thus spanning from 3 to 8. The range of the hyperparameter k was limited to 1 to 5 since, for k > 5, as larger values led to underfitting in both datasets. The minimal Hausdorff distance was implemented using both Euclidean $(d(x_1, x_2) = ||x_1 - x_2||_2)$ and city block $(d(x_1, x_2) = ||x_1 - x_2||)$ distance metrics to compute the distance between bags. For the BCkNN, we tested the 3 different weight functions $f(b_t, c_j)$ (see Eqs. 5, 6, and 7). Since in this work, we focus on integrating NCFS into the MIL lazy learning framework, we tuned the hyperparameters concerning only the training phase of our method, i.e. λ and σ of the NCFS algorithm.

Experimental results on the Musk dataset are computed based on 100 randomly repeated 10-fold cross-validation procedures. Following the affective computing literature, the experimental results on the DEAP dataset were obtained following a leave-one-video-out procedure (LOVO) [11], i.e., the models were trained with k-1 videos and tested with the k-th remaining video. The models' performance is discussed in terms of average accuracy and macro-averaged F1 scores. Due to the imbalanced nature of both datasets, we primarily focus on the macro-F1 score to ensure that both classes are equally considered. All metrics are reported in percentages.

5 Experimental Results

This section presents our experiments on the Musk (see Sect. 5.1) and DEAP (see Sect. 5.2) datasets. For each dataset, we present the experimental results for the proposed methodologies. The experimental comparison between our proposed NCFS-based MIL algorithms and the standard MIL algorithms is described in Sect. 5.3. Finally, the statistical significance of our approach is evaluated in Sect. 5.4.

5.1 Musk Dataset

Regarding the BkNN, CkNN, and BCkNN algorithms endowed with NCFS, the experimental results are shown in Table 1. For the Musk dataset, the best macro-F1 scores are 90.07, 93.32, and 91.66% for NCFS-BkNN (k = 2), NCFS-CkNN (k = 2, q = 4), and NCFS-BCkNN (k = 2, q = 4, Eq. 6) respectively, using the Euclidean distance. Overall the best result was obtained with NCFS-CkNN (93.32%).

Table 1. Classification accuracy and macro-F1 (m-F1) score of NCFS-BkNN, NCFS-CkNN, and NCFS-BCkNN for the Musk dataset averaged over 100 repetitions. The best results, based on macro-F1, are highlighted in bold.

NCFS-BkNN										
Distance	k = 1		k = 2		k = 3		k = 4		k = 5	
	Acc	m-F1	Acc	m-F1	Acc	m-F1	Acc	m-F1	Acc	m-F1
Euc	87.89	87.85	90.14	90.07	84.67	84.13	87.78	87.39	75.56	73.43
City	84.92	84.58	84.14	83.44	77.17	75.56	77.92	77.40	77.92	76.11
NCFS-CkNN										
Euc	86.89	86.74	93.39	93.32	92.39	92.25	89.03	88.78	86.89	86.51
City	86.17	85.85	86.03	85.11	84.03	83.52	83.53	83.30	83.67	83.09
NCFS-B	CkNN	I (Eq.	5)							
Euc	87.89	87.85	90.14	90.07	84.67	84.13	87.78	87.39	75.56	73.43
City	84.92	84.58	84.14	83.44	77.17	75.56	77.92	77.40	77.92	76.11
NCFS-B	CkNN	I (Eq.	6)							
Euc	87.89	87.85	91.89	91.66	88.92	88.62	90.03	89.79	89.89	89.66
City	87.17	86.97	84.67	84.33	84.92	84.50	84.92	84.50	85.67	85.31
NCFS-B	CkNN	I (Eq.	7)							
Euc	87.89	87.85	90.89	90.67	88.92	88.62	90.03	89.79	89.89	89.66
City	87.17	86.97	85.92	85.59	85.92	85.53	84.92	84.50	88.92	88.62

5.2 DEAP Dataset

The experimental results for NCFS-BkNN, NCFS-CkNN, and NCFS-BCkNN are shown in Table 2. For the arousal task, the best macro-F1 scores are 53.59, 53.17, and 54.23% for NCFS-BkNN (k = 1), NCFS-CkNN (k = 2, q = 4), and NCFS-BCkNN (k = 1, q = 3, Eqs. 6 and 7) respectively, using the city block distance. For the valence task, the best macro-F1 scores are 58.53, 60.26, and 60.21% for NCFS-BkNN (k = 4, city block distance), NCFS-CkNN (k = 4, q = 6, Euclidean distance), and NCFS-BCkNN (k = 5, q = 7, Eq. 7, Euclidean distance) respectively. Overall, the best results were obtained with NCFS-BCkNN (macro-F1 = 54.23 ± 11.38%) and with NCFS-CkNN (macro-F1 = 60.26 ± 10.45%) for solving the arousal and valence tasks, respectively.

For both arousal and valence tasks, we observed a large standard deviation in the macro-F1 scores across the 32 subjects, indicating significant variability among subjects, which is a well-known challenge in affective computing tasks. To ensure the reliability of our results, we conducted a nonparametric permutation test, confirming that these scores were significantly above chance level (i.e., 50%).

Table 2. Classification accuracy and macro-F1 (m-F1) score of NCFS-BkNN, NCFS-CkNN, and NCFS-BCkNN for the arousal (top) and valence (bottom) tasks of the DEAP dataset averaged over the 32 subjects. Stars indicate whether the macro-F1 score distribution over subjects is significantly higher than 50% according to an independent one-sample one-tailed t-test (**: p < 0.01, *: p < 0.05). The best results, based on macro-F1, are highlighted in bold.

NCFS-BkNN – Arousal task										
Distance	k = 1		k = 2		k = 3		k = 4		k = 5	
	Acc	m-F1	Acc	m-F1	Acc	m-F1	Acc	m-F1	Acc	m-F1
Euc	60.23	53.23	57.03	50.25	62.03	52.69	60.62	51.11	62.11	49.87
City	59.76	53.59	56.77	51.24	60.62	52.34	58.20	50.55	61.25	50.87
NCFS-C	kNN –	Arousa	al task							
Euc	57.11	52.24	57.73	52.26	58.20	51.63	58.20	49.96	59.69	50.40
City	57.03	52.13	58.44	53.17	59.27	52.35	60.08	52.32	59.69	51.07
NCFS-B	CkNN	(Eq. 5)	– Aro	usal tas	k					
Euc	59.61	53.30	58.12	52.15	59.53	53.13	59.61	52.83	60.23	52.83
City	59.69	54.05	59.45	53.77	59.69	53.28	60.00	53.42	59.84	53.17
NCFS-B	CkNN	(Eq. 6)	– Aro	usal tas	k					
Euc	59.61	53.30	59.45	53.12	59.61	52.10	59.14	50.21	60.23	50.19
City	59.84	54.23*	60.23	54.05	60.31	52.66	61.33	52.58	60.55	51.36
NCFS-B	CkNN	(Eq. 7)	– Aro	usal tas	k					
Euc	59.61	53.30	59.37	53.03	59.84	52.34	59.06	50.11	60.23	49.94
City	59.84	54.23*	60.54	54.14	59.92	52.50	61.25	52.48	60.86	51.85
NCFS-B	kNN –	Valenc	e task							
Euc	58.67	55.68**	55.33	53.40*	60.78	57.14**	60.94	57.68**	62.66	57.83**
City	57.34	55.32**	56.33	53.06	61.41	58.28**	61.40	58.53**	60.86	55.80**
NCFS-C	kNN –	Valenc	e task							
Euc	56.48	54.43**	59.14	56.99**	61.17	58.64**	62.81	60.26**	63.12	60.10**
City	55.86	54.25*	58.44	56.24**	60.62	58.30**	61.80	59.07**	62.42	58.95**
NCFS-B	CkNN	(Eq. 5)	– Vale	ence tas	k					
Euc	58.59	56.18 * *	60.78	58.10**	62.03	58.65**	63.36	59.78**	63.58	60.13**
City	57.19	55.36**	58.67	56.22**	61.48	58.86**	62.11	59.17**	63.20	59.57**
NCFS-B	CkNN	(Eq. 6)	– Vale	ence tas	k					
Euc	58.59	56.18**	60.86	58.17**	62.34	58.97**	63.59	60.01**	63.67	60.19**
City	57.18	55.36**	59.45	56.81**	61.40	58.63**	61.71	58.39**	63.28	59.53**
NCFS-B	CkNN	(Eq. 7)	– Vale	ence tas	k					
Euc	58.59	56.18**	60.86	58.17**	62.34	58.97**	63.51	59.93**	63.67	60.21**
City	57.18	55.36**	58.67	56.22**	61.48	58.86**	62.11	59.17**	63.20	59.57**

5.3 Comparison with State-of-the-Art

The comparison between our proposed NCFS-based MIL algorithms and the state-of-the-art MIL lazy algorithms (i.e., without NCFS) is shown in Table 3. It is worth noting that these competitors (i.e., BkNN, CkNN, and BCkNN) are still successfully used in various applications today [14,21]. As comparisons, we used only MIL-based algorithms, since previous work on Musk [27] and DEAP [19] datasets confirmed the superiority of MIL algorithms over SIL algorithms for solving these tasks. For each model, we report the best results achieved in the previous experiments (see Sects. 5.1 and 5.2) and compare them to state-of-the-art methods tested with the same k, q, and f.

Table 3. Comparisons with respect to state-of-the-art models in terms of macro-F1 score for both Musk and DEAP (A: Arousal task, V: Valence task) datasets. The best results for each dataset and distance metric are indicated in bold. The overall best results are indicated by † .

Model	Euclide	ean distan	ce	City block distance			
	Musk	DEAP-A	DEAP-V	Musk	DEAP-A	DEAP-V	
BkNN	90.07	52.04	50.67	83.59	48.50	51.51	
CkNN	90.91	50.64	52.23	85.85	49.83	50.87	
BCkNN (Eq. 5)	90.07	52.03	53.93	83.59	48.68	52.85	
BCkNN (Eq. 6)	90.65	52.03	52.24	85.98	47.64	51.86	
BCkNN (Eq. 7)	88.92	52.03	52.02	88.62	47.64	52.20	
NCFS-BkNN (Ours)	90.07	53.23	57.83	84.58	53.59	58.53	
NCFS-CkNN (Ours)	93.32^\dagger	52.26	60.26^\dagger	85.85	53.17	59.07	
NCFS-BCkNN (Eq. 5, Ours)	90.07	53.30	60.13	84.58	54.05	59.57	
NCFS-BCkNN (Eq. 6, Ours)	91.66	53.30	60.19	86.97	54.23^\dagger	59.53	
NCFS-BCkNN (Eq. 7, Ours)	90.67	53.30	60.21	88.62	54.23^\dagger	59.57	

Concerning the Musk dataset, by using the Euclidean distance, the NCFS improves the performance of CkNN (93.32 vs 90.91), BCkNN (Eq. 6) (91.66% vs 90.65%), and BCkNN (Eq. 7) (90.67% vs 88.92%), while performances are comparable for BkNN (90.07%) and BCkNN (Eq. 5) (90.07%). Using the city block distance, the performance of BkNN (84.58% vs 83.59%), BCkNN (Eq. 5) (84.58% vs 83.59%), and BCkNN (Eq. 6) (86.97% vs 85.98%) improves with NCFS, while the macro-F1 scores are comparable for CkNN (85.85%) and, BCkNN (Eq. 7) (88.62%).

Regarding the arousal task of the DEAP dataset, NFCS improves the performance of all methods. Using the Euclidean distance, NCFS enhances the performance of BkNN (53.23% vs 52.04%), CkNN (52.26% vs 50.64%), and BCkNN (with Eqs. 5, 6, 7) (53.30% vs 52.03%). With the city block distance, NCFS also improves the performance of BkNN (53.59% vs 48.50%), CkNN (53.17% vs 49.83%), BCkNN (Eq. 5) (54.05% vs 48.68%), BCkNN (Eq. 6) (54.23% vs 47.64%), and BCkNN (Eq. 7) (54.23% vs 47.64%) improved with NCFS.

Similarly, the introduction of NCFS enhances the performance of all algorithms for the valence task. Using the Euclidean distance, NCFS improves the performance of BkNN (57.83% vs 50.67%), CkNN (60.26% vs 52.23%), BCkNN (Eq. 5) (60.13% vs 53.93%), BCkNN (Eq. 6) (60.19% vs 52.24%), and BCkNN (Eq. 7) (60.21% vs 52.02%). With the city block distance, NCFS improves the performance of BkNN (58.53% vs 51.51%), CkNN (59.07% vs 50.87%), BCkNN (Eq. 5) (59.57% vs 52.85%), BCkNN (Eq. 6) (59.53% vs 51.86%), and BCkNN (Eq. 7) (59.57% vs 52.20%) also improved with NCFS.

As a result, the proposed NCFS-CkNN with Euclidean distance achieved the best performance, overcoming state-of-the-art models for solving the Musk (93.32%) and DEAP valence (60.26%) tasks. Accordingly, the NCFS-BCkNN (Eq. 6 and Eq. 7) with city block distance achieved the best performance, surpassing state-of-the-art models for the DEAP arousal (54.23%) task.

5.4 Statistical Significance

We performed a statistical analysis for the Musk and DEAP datasets to compare the performance obtained with our approach (NCFS models) and with the stateof-the-art models employed as reference. The statistical analysis was performed by comparing the 100 and 32 macro-F1 values originated by 100 randomized repetitions of the 10-fold CV and by 32 LOVO CV procedures, respectively. The macro-F1 scores of all proposed methods were found to follow a normal distribution according to the Anderson-Darling test (p > 0.05), for all classification tasks. Hence, an independent two-sample one-tailed t-test ($\alpha = 0.05$) was performed to determine whether there was statistical evidence of an improvement in the algorithms' performance applying the NCFS procedure. The performance of our NCFS-CkNN with Euclidean distance was significantly higher than BkNN (p < 0.01), CkNN (p < 0.01), BCkNN (Eqs. 5 and 7) (p < 0.01) and BCkNN (Eq. 6) (p < 0.05) for solving the Musk task. A statistically significant improvement was also found between the proposed NCFS-CkNN and BkNN (p < 0.01), CkNN (p < 0.01), BCkNN (Eqs. 5, 6 and 7) (p < 0.01) for solving the valence task on the DEAP dataset with Euclidean distance as distance metric. Similarly, the proposed NCFS-BCkNN (Eqs. 6 and 7) showed a significant improvement with respect to BkNN (p < 0.05), CkNN (p < 0.05), BCkNN (Eq. 5) (p < 0.05) and BCkNN (Eqs. 6 and 7) (p < 0.01) for solving the arousal task on the DEAP dataset with city block distance as distance metric. Additionally, regarding the Musk dataset, we also computed an empirical *p*-value as $p = N_q/N_c$, where N_c was the number of comparisons (i.e., 100) and N_g was the number of times that an element of the 100 macro-F1 scores of models without NCFS was greater than the median macro-F1 score of the respective NCFS-model. This analysis confirmed the results obtained with parametric tests (p < 0.05).

5.5 Computational Complexity

The computational complexity of Algorithm 1 for training and testing, respectively, is $\mathcal{O}(L^2N^2d)$ and $\mathcal{O}(L^2Nd)$. It is worth noting that by employing a Kd-Trees retrieval strategy, the computational complexity could be lowered to $\mathcal{O}(L^2\log(N)^2d)$ and $\mathcal{O}(L^2\log(N)d)$ [3,25]. Moreover, we report the average fold execution time of Algorithm 1 for the Musk dataset on an Intel Core i7-7820X processor. The training execution time is predominantly influenced by the NCFS procedure (10.25±8.11 s), whereas our methodology has minimal impact on testing execution times, with values of 0.79 ± 0.08 , 0.78 ± 0.08 , and 0.79 ± 0.08 seconds for BkNN, CkNN, and BCkNN, respectively.

6 Conclusions

This paper introduces a novel feature selection method tailored for the MIL framework. Our proposed methodologies provide the prediction of unseen bags by integrating Bayesian and citation approaches with the optimal weighted minimal Hausdorff distance. The integration of NCFS into the MIL lazy learning framework significantly enhanced model generalizability by eliminating redundant and irrelevant features. This is supported by experimental results across different benchmark datasets consistently demonstrating how NCFS statistically improved the performance of BkNN, CkNN, and BCkNN. Notably, the weight vector resulting from NCFS can be further exploited to gain insights into the importance of each feature, thereby enhancing the transparency and interpretability of the MIL algorithms discussed. Indeed, further investigation could be devoted to leveraging this approach to improve the interpretability of standard MIL-based methods.

Future work could investigate the proposed approach's effectiveness and robustness, including providing statistical guarantees, demonstrating algorithm convergence, and analyzing sensitivity to various hyperparameters.

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MESS: Coarse-Grained Modular Two-Way Dialogue Entity Linking Framework

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Abstract. Entity Linking aims to map mentions in a document to corresponding entities in a given knowledge base. Most previous studies usually extract mentions and infer their underlying entities. An obvious limitation of this approach (mention-to-entities, M2E) is that it cannot fully exploit the rich structured information from the knowledge base. Moreover, employing this knowledge to identify entities and find their possible mentions (entity-to-mentions, E2M) still faces problems, such as the semantic correlation between mentions and entities with documents cannot be effectively captured. Therefore, we propose a coarse-grained modular two-way Dialogue entity linking framework called MESS. It mainly consists of four modules: M2E and E2M, Semantic Synchronization (SS), and Dialogue, in which M2E and E2M independently execute EL decisions, SS performs semantic constraints, and Dialog merges the two-way results and outputs a final decision. Specifically, MESS first uses the M2E and E2M modules to generate candidate sets of entities and mentions in parallel, then utilizes the semantic synchronization module (SS) to filter irrelevant mentions and entities. Finally, M2E and E2M determine the target mention-entity pair through two-way dialogue. We validate the superiority of our MESS through extensive experiments on various baselines, the results show that it achieves competitive results.

Keywords: Knowledge base \cdot Entity linking \cdot Question answering

1 Introduction

Entity Linking (EL), also called Entity Disambiguation (ED), aims to map mentions in the text to proper entities in a given knowledge base (KB). It is essential to many natural language processing tasks, which include information retrieval [13], relation extraction [19], and question answering [3], etc. The main challenge of those tasks is to denoise the ambiguous linkages between text mentions and knowledge base entities. This is because language's inherent and omnipresent ambiguity at the lexical level results in the ambiguity of words, named entities, and other linguistic units. For example, a mention of Victoria may refer to Victoria in Australia, Victoria in British Columbia, Victoria in Texas, Queen

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Victoria, or Victoria cricket team [40]. It is hard for an algorithm to identify which one refers to. Moreover, EL is also affected by the amount of information in KG. The popular KGs include DBpedia [24], Freebase [2], Yago [37] and etc.

As we all know, the usual approach to entity linking is first to detect mentions in a given document and then link them to corresponding entities in a KG. We call such method *mention-to-entities* (M2E), which decomposes the EL task into mention detection (MD) and entity disambiguation (ED) and then considers them independently. The mentions are assumed to be given [15], and an offthe-shelf Named Entity Recognition (NER) system [25] is utilized to extract mentions and disambiguate them with ED. Besides, an end-to-end model [21] has been proposed to jointly perform entity recognition and disambiguation. Nonetheless, the *mention-to-entities* approach suffers from apparent limitations. For example, it is more challenging to predict mentions without the knowledge of entities, and errors from MD may propagate to ED. Although the end-to-end model can alleviate the error propagation problem, the nearest search is only approximate.

An intuitive idea is to determine the candidate entities first and then infer the mentions in the document. We call this model *entity-to-mentions* (E2M), which reverses the order of the two subtasks, identifying candidate entities before MD. EL [45] has been regarded as a reverse of open-domain Question Answering task, using dual encoders to efficiently retrieve the top-K candidate entities from the knowledge base as "questions" for a given document. Then a deep cross-attention mechanism is applied for each candidate to identify mentions as "answer spans". The *entity-to-mentions* method can better utilize the rich information in the knowledge base, but the document information that can be referenced when generating mentions is relatively limited.

In summary, using M2E and E2M to model EL independently will face some challenges. M2E cannot fully utilize the structured information in KB, and the information in the document that E2M can exploit when generating mentions is relatively limited. Therefore, a more straightforward idea is to combine M2E and E2M to superposition their respective advantages to reduce the impact of drawbacks on the results. In order to verify this conjecture, we conducted EL experiments on the AIDA and MSNBC datasets. The results show that simply combining them cannot improve EL performance, and sometimes the combined results are even worse than individually. Through more fine-grained exploration, we find that there is a deviation in the semantic correlation between the results inferred by M2E and E2M and the corresponding documents. This problem can be avoided if irrelevant candidates are filtered out before making EL decisions. Based on these findings, we propose a coarse-grained modular two-way dialogue entity linking framework called MESS. MESS comprises four modules: M2E, E2M, Semantic Synchronization (SS), and Dialogue. M2E and E2M are used to generate candidate sets in parallel. SS is the most critical part of MESS. It plays a semantic synchronization function so that the results of two predictions can be applied to the same document. The Dialogue module calculates and compares the final results of M2E and E2M and outputs the final mention-entity pairs. The main contributions of this work are summarized as follows:

- We propose an effective and efficient EL framework that consists of four essential components: (1) Mention-to-Entities, (2) Entity-to-Mentions, (3) Semantic Synchronization, and (4) Dialogue. All of these modules work together to improve the performance of EL.
- The proposed MESS framework combines the strengths of the latest SOTA M2E and E2M models. It addresses the limitations of current EL models that do not effectively utilize structured information in KB and insufficient features that can be exploited in a given document.
- Extensive experiments on public datasets show that our MESS obviously outperforms SOTA methods using E2M and M2E alone in EL tasks, and it also performs excellently on ED. Moreover, we conduct ablation studies to demonstrate the effectiveness of each module.

2 Related Work

We first present classical entity linking works of *mention-to-entities* that recognize mentions and infer their corresponding entities. Then, we summarize the work of translating EL to other tasks to predict mention-entity pairs.

2.1 Mention-to-Entities

Early approaches to EL focus on extracting discriminative features of entities from documents and then linking mentions to entities with the highest similarity. Specifically, [27] uses cosine similarity to measure the compatibility between name mentions and candidate entities. Other work [4] augments with additional entity information, such as its categories. [26] defines a measurement using semantic relatedness between candidate entities and mentions. These works can be roughly divided into two major categories: local and global disambiguation. Various works have been proposed to model the mention local contexts, such as binary classification [28] and rank model [6]. A salient trait of these methods is the use of a large number of human-annotated features and the mentions of hard-to-extract that are retrieved from search engines or Wikipedia [38]. They face time-consuming and labor-intensive issues.

As representation learning is used to extract semantic features automatically, it is further demonstrated that entity representations learned by jointly modeling textual context and knowledge bases effectively combine multiple information sources [40]. Therefore, the global model assumes that all mentions of target entities in a document are related, and the factor graph model is constructed in [33], which represents mentions and candidate entities as variable nodes and utilizes factor nodes to stand for a series of features. The work [11] uses fully connected pairwise conditional random field models and utilizes cyclic belief propagation to estimate the maximum marginal probability. The above models all face the same problem; that is, many features need to be predefined.

To relieve the computational pressure of candidate entity pairs, [12] introduce a consistency model with an attention mechanism, where each mention only focuses on a fixed number of entities. However, the number of mentions after using attention is difficult to determine. Multi-focal attention [10] proposes the Star Model to decompose collective entities linking tasks over mentions and makes it easy to integrate attention in both learning and inference. The work [32] does link all mentions by scanning at most one mention pair, and they assume that each mention only needs to be consistent with another mention in the document. The limitation of their method is that the consensus information is too sparse, which results in low confidence. [14] rank mentions according to the ease of disambiguation, but they do not fully utilize the information of previously mentioned entities for subsequent entity disambiguation. [29] use sequence models, but they encode the result of greedy selection and measure the similarity between the global encoding and candidate entity representations. Nevertheless, they do not consider the long-term impact of current decisions on subsequent choices, nor do they add selected target entity information to the current state to help disambiguate.

2.2 Transferred EL

Graph Neural Networks (GNNs) are a flexible and effective framework for the representation learning of various graph-structured data. [16] first presents a graph-based representation that can model the global interdependence between different EL decisions and then propose a collective inference algorithm, which can jointly infer the entities of all mentions by exploiting the interdependence captured in the constructed graph. In contrast, methods based on graph neural networks can better capture local and global dependencies, but they suffer from either data sparseness or existing noises. To overcome those problems, [40] design a dynamic Graph Convolutional Networks architecture that can collectively identify the entity mappings between the document and the knowledge graph and efficiently capture the topical coherence among various entity mentions in the entire document through aggregated knowledge from dynamically linked nodes. With the development of multimodal technology, [46] introduce a self-supervised simple triplet network, which can learn useful representations in multimodal unlabeled data to improve the effectiveness of NED models. Unlike graph representation learning, [44] convert EL to question answering. Specifically, it efficiently retrieves the top-k candidate entities as questions from a knowledge base using a dual-encoder retriever. A deep cross-attention reader is then applied to each candidate's document to identify candidates mentioned in the document as answer spans. However, none of them considers modeling EL using a bidirectional perspective.

3 Our MESS Framework

The structure of our MESS is shown in Fig. 1, which consists of M2E, E2M, SS, and Dialogue. In this section, we introduce the implementation details of these modules.



Fig. 1. The overall architecture of our MESS. It mainly consists of M2E, E2M, SS, and Dialogue modules.

3.1 M2E Module

The M2E module utilizes contrastive learning to capture the representation and matching patterns of mentions and entities. Assume that the identified mentions in a given document are denoted by $M = \{m_1, m_i, ..., m_n\}$, where all candidate entities related to m_i are described by $E = \{e_1, e_i, ..., e_m\}$. In practice, we encode each candidate entity and related descriptive information as a basic representation unit instead of encoding entities individually. We concatenate the candidate entity and its attributes as the input R_E of BERT and exploit a Multi-layer Perceptron to compute the entity embedding:

$$E_{e^j} = MLP(B(CLS, R_E)) \tag{1}$$

where $B(CLS, R_E)$ is the CLS token embedding of BERT.

Metric Function. Inspired by the work of [5], we use an interaction-based metric to measure the more fine-grained similarity between mentions and candidate entities. Formally, we utilize Eq. (3) to obtain the embeddings $\{E_{m_i}\}_{i=1}^{i=n}$ and $\{E_{e_j}\}_{j=1}^{j=m}$ of mentions and candidate entities, respectively. We compute the

similarity matrix M_s between m_i and $\{E_{e_j}\}_{j=1}^{j=m}$ and then obtain d_i^j that is calculated from the normalized Euclidean distance between any mention and all related candidate entities, i.e., $d_i^j = ||E_{m_i} - E_{e_j}||_2^2$. In this module, we employ a Gaussian kernel function to measure the distance between the mention and the candidate entity, where the mention is the center. The value of the kernel function is 1 when the entity is very close to the center and 0 otherwise. Specifically, we first transform $D(d_i^j)$ into a k-dimensional distribution (Eq. (2)). Next, the k-th item is converted by the k-th Gaussian kernel with mean μ_k and variance σ_k (Eq. (3)). Then we represent the similarity between the mention m_i and candidate entity e_j by summing (Eq. (4)). Finally, we use an MLP layer to obtain a similarity score (Eq. (5)).

$$D(d_i^j) = [D_1(d_i^j), D_2(d_i^j), ..., D_k(d_i^j)],$$
(2)

$$D(d_i^j) = exp[-\frac{(d_i^j - \mu_k)}{2\sigma_k^2}],$$
(3)

$$\Phi(m_i, E) = \log \sum_{i=1}^{m} D(d_i^j), \tag{4}$$

$$f(m_i, E) = MLP(\Phi(m_i, E))$$
(5)

Loss Function. We jointly exploit positive and negative samples to build a loss function. For any mention in a given document, positive and negative samples are determined by computing the cosine similarity between the mention and all candidate entities, resulting in a similarity ranking table. We regard the first and the last as a pair of positive and negative samples, and so on for other sample pairs. Given a set of triplets (e_j, e_i^+, e_i^-) , the loss function is defined as:

$$\mathcal{L}_{m2e}(m_i, e_j) = \sum_{(e_j, e_j^+, e_j^-)} \max\{0, g + f(e_j, e_j^-) - f(e_j, e_j^+)\}$$
(6)

3.2 E2M Module

The E2M module comprises an Entity Extractor (EE), and a Mention Recognizer (MR). EE extracts relevant entities based on a given document, while MR combines the extracted entities and document information to identify potential mentions. In this section, we introduce the details of the EE and MR and the overall training objective of E2M.

Entity Extractor (EE). The knowledge base is $KB = \{e_1, ..., e_N\}$. Given a sentence t of length L_t , EE wants to obtain the subset $\mathcal{E}_c \in KB$ as candidate entities of t. EE is a dual-encoder structure containing sentence encoder E_s and entity encoder E_e . We use E_s to map the sentence t to the representation sequence r_t , and E_e to get the representation r_{e_i} of the entity $e_i \in KB$.

$$r^{e_i} = E_e[m(e_i)], \quad r^t = E_s(t)$$
 (7)

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where $m(e_i)$ indicates that the description information of e_i is obtained from KB, and the length of all entity description texts is uniformly limited KB to L_e . Then, we calculate the dot product between the two vectors as the score to extract the top K related entities \mathcal{E}_K :

$$\mathcal{E}_{K} = \underset{\mathcal{E}' \subseteq \mathcal{E}, |\mathcal{E}'| = K}{\operatorname{arg\,max}} \sum_{e_i \in \mathcal{E}'} r^{t^T} r^{e_i} \tag{8}$$

For a training sentence t and the knowledge base KB, based on Eq. (8), for every entity $e_i \in KB$, the extractor's score is formulated as:

$$S(t,e_i) = r^{t^T} r^{e_i} \tag{9}$$

We have its gold entity set $\mathcal{G} \in KB$, and we can achieve its negative entity set $\mathcal{N} \in KB$. Then, we exploit a multi-label variant of Noise Contrastive Estimation (NCE) during the model training phase, which is computed as:

$$\mathcal{L}_{\text{EE}} = \max \sum_{g \in G} \log \left(\frac{\exp(S(t,g))}{\exp(S(t,g)) + \sum_{n \in N} \exp(S(t,n))} \right)$$
(10)

Mention Recognizer (MR). Given the output of EE (i.e., \mathcal{E}_c) and an input sentence t, for each candidate entity $e_j \in \mathcal{E}_c$, we obtain its joint representation with t:

$$r^{e_j} = E_s(t \oplus f(e_j)),\tag{11}$$

After obtaining the joint representation, we can calculate the probability of span $(x, y), 1 \le x, y \le L_t$, and the sorting probability of $e_j, 1 \le j \le K$:

$$p_b(x|t, e_j) = f(W_1 r^{e_j}, x)$$
(12)

$$p_e(y|t, e_j) = f(W_2 r^{e_j}, y)$$
 (13)

$$P_{\rm rank}(e_j|t, \mathcal{E}_{\rm c}) = \frac{\exp(W_3^T r^{e_j})}{\sum_{j'=1}^K \exp(W_3^T r^{e_{j'}})}$$
(14)

where W_1 , W_2 and W_3 are the trainable parameters. According to Eqs. (12)–(14), the MR score is calculated as follows:

$$P(e_j, x, y|t, \mathcal{E}_c) = p_1(x|t, e_j) \times p_2(y|t, e_j) \times P_{rank}(e_j|t, \mathcal{E}_c).$$
(15)

As for the training of the MR, we directly optimize it to maximize the span probability and ranking probability, the objective function is shown below:

$$\mathcal{L}_{MR} = \max \sum_{j=1}^{K} \sum_{(x,y)} \{ \log[p_1(x|t, e_j) \\ \times p_2(y|t, e_j)] + \log P_{rank}(e_j|t, \mathcal{E}_c) \}$$
(16)

Finally, we train the EE objective \mathcal{L}_{EE} and MR objective \mathcal{L}_{MR} simultaneously. The overall objective of E2M is formulated as:

$$\mathcal{L} = \mathcal{L}_{EE} + \mathcal{L}_{MR} \tag{17}$$

3.3 SS Module

When we only combine M2E and E2M to make EL decisions, the performance is worse than when they execute alone. Therefore, we design the SS module to effectively solve the problem of out-of-synchronization of M2E and E2M semantics. Specifically, the SS module mainly works on the candidate sets of M2E and E2M, filtering out entities and mentions that are semantically irrelevant to the document. This section details the SS module for encoding intrinsic semantic relationships between mentions, entities, and documents. We first pre-train our SS teacher model to make more accurate predictions with a moderate amount of manually labeled mention entity relevance data. Based on the well-trained teacher model, we construct a student model using shared encoder parameters through knowledge distillation to imitate the teacher's prediction. Then, we minimize the predicted difference between them to enhance the recognition and matching of mentions and entities.

SS Teacher. The SS teacher model is a BERT-based binary classification model, which predicts whether the given mention and candidate entity are semantically close to documents. We employ BERT^{ct} to denote this model and "1"/"0" to represent semantic similarity and dissimilarity, respectively. Unlike existing models with bi-encoder architecture, the SS teacher model applies cross-encoders, which are more expressive and enable more accurate classification. The SS score is calculated:

$$S(X,Y) = \sigma(W^{ct}BERT^{ct}([X \oplus Y, CLS, SEP]))$$
(18)

where $[X \oplus Y]$ denotes concatenating X and Y, X can be the embedding of mentions, entities, or documents, and Y is the other two embeddings except X. $\sigma(\cdot)$ and W^{ct} are the sigmoid activation function and the linear projection that map the output to the actual value, respectively. We can find that this model can be used to calculate the semantic correlation between any two items (mentions, entities, and documents). Usually, it is used to evaluate the semantic similarity between mentions and candidate entities in bidirectional EL.

SS Student. Based on the well-trained teacher model, we obtain another model with the same architecture and shared parameters through knowledge distillation. Specifically, we first utilize the SS teacher model to compute the similarity scores S(m, e) of entities and mentions and then employ the distillation model B^{stu} to imitate the teacher's prediction with the inner product of the mention and entity embedding. The loss is expressed as:

$$\min\sum_{m}\sum_{e} ||S(m,e) - [B^{stu}(m) \cdot B^{stu}(e)]||.$$
(19)

where $[\cdot]$ is the inner product operator. By modeling the above problems, we can estimate the level of semantic similarity between the mention and the entity by the inner product value.

3.4 Dialogue Module

After applying the SS module, M2E and E2M calculate the final output through the Dialogue module. For the given document t, we obtain a set of mentions and entities separately through the two-way module. The mentions extracted from the M2E module are M(m2e), and the set of mentions matched from the E2M module is M(e2m). Then, we use each mention in M(m2e) to calculate the cosine similarity with the mention in M(e2m) one by one and find the most similar one in M(e2m) for each mention in M(m2e). Finally, we utilize the obtained mention-mention couples to calculate semantic similarity with their corresponding entity and output the most similar mention-entity pair as the final result. It can be computed as:

$$c(m_{11}, m_{21}) = S_{cos}(m_l, M_{e2m}),$$
(20)

$$d(m,e) = max[S_{cos}(m_{11},e_{11}),S_{cos}(m_{21},e_{21})]$$
(21)

where $l \in [1, T]$, T is the number of elements in M(m2e), and $S_{cos}(\cdot, \cdot)$ means the cosine similarity.

4 Experiments

This section evaluates our model's effectiveness on ED and end-to-end entity linking tasks.

4.1 Setting

Entity Disambiguation. To facilitate comparison with the most advanced works, we follow the settings presented by Le et al. [22] to use InKB Micro- F_1 as a metric to compute scores on in-domain and out-of-domain datasets and exploit the same candidate set.

Datasets. We first pre-train *MESS* on the BLINK data [41] containing the 9M unique triples document-mention-entity from Wikipedia. Then, we fine-tune our model on other corpora, where AIDA-CoNLL [18] is an in-domain dataset, the MSNBC, AQUAINT, ACE2004, WNED-CWEB (CWEB), and WNED-WIKI (WIKI) [14] are the out-of-domain.

Details. We leverage WordPiece [20] as our tokenizer, training on the BLINK data. There are a total of 50777 vocabularies for the well-trained tokenizer. For optimization, Adam applies a learning rate of 2e-5 for the *Mention-to-Entities* and 1e-7 for the *Entity-to-Mentions*;

Baselines. Ganea and Hofmann [11] propose joint document-level entity disambiguation combining entity embeddings and an attention mechanism. Guo

Table 1. Micro F1 (InKB) on an in-domain test set and out-of-domain test sets for named entity disambiguation tasks. Bold indicates the best model, and <u>underline</u> hints the suboptimal. WIKI^{*} is generally considered out-of-domain but points out that all methods use parts of Wikipedia for training. ξ results from here and normalized to accommodate entities not in KB. M2E and E2M represent *mention-to-mentions* and *entity-to-mentions*, respectively, while Combine (M2E+E2M) simply merges the two without SS and Dialogue modules. MESS (M2E+E2M) refers to the modular approach we proposed.

Models	AIDA	MSNBC	AQUAINT	ACE2004	CWEB	WIKI*	Avg.
Ganea & Hofmann (2017)	92.9	93.7	88.5	88.5	77.9	77.5	86.4
Guo & Barbosa (2018)	89.0	92.0	87.0	88.0	77.0	84.5	86.2
Yang et al. (2018)	95.9	92.6	89.9	88.5	81.8	79.2	88.0
Shahbazi et al. (2019)	93.5	92.3	90.1	88.7	78.4	79.8	87.1
Yang et al. (2019)	93.7	93.8	88.2	90.1	75.6	78.8	86.7
Le & Titov (2019)	89.6	92.2	90.7	88.1	78.2	81.7	86.8
Fang et al. (2019)	94.3	92.8	87.5	91.2	78.5	82.8	87.9
Wu et al. (2019)	79.6	80.8	80.3	82.5	64.2	75.5	77.0
Cao et al. (2021)	93.3	94.3	89.9	90.1	77.3	<u>87.3</u>	<u>88.8</u>
M2E (Only)	92.4	93.9	90.6	90.8	78.3	86.5	88.8
E2M (Only)	93.3	93.6	90.5	91.0	77.2	86.8	88.7
Combine $(M2E+E2M)$	92.7	92.5	88.9	89.6	76.3	86.1	87.7
MESS (M2E+E2M)	94.8	94.7	90.3	91.5	78.6	88.9	89.8

and Barbosa [14] define the semantic similarity between mention and entity as the mutual information between random walks on the graph. Yang et al. [43] show a jointly disambiguating method utilizing a gradient-tree-boosting-based structure. Shahbazi et al. [36] define mentions as a function of the entire paragraph and combine language models to predict the referred entities. Yang et al. [42] sequentially accumulate contextual information for efficient collective reasoning. Le and Titov [23] treat entities as latent variables and select entities based on the context of mentions and coherence of other entities. Fang et al. [9] transform the global link into a sequential decision-making problem with reinforcement learning. GENRE [7] presents a system for retrieving entities by autoregressively generating unique names.

End-to-End Entity Linking. We evaluate InKB Micro- F_1 on the GERBIL benchmark platform [35] on both in-domain and out-of-domain datasets for the EL.

Datasets. The AIDA-CoNLL which is the in-domain dataset, the seven outof-domain test sets are MSNBC, Derczynski (Der) [8], KORE 50 (K50) [17],

Models	AIDA	MSNBC	Der	K50	R128	R500	OKE15	OKE16	Avg.
Hoffart et al. (2011)	72.8	65.1	32.6	55.4	46.4	42.4	63.1	0.0	47.2
Steinmetz et.al (2013)	42.3	30.9	26.5	46.8	18.1	20.5	46.2	46.4	34.7
Moro et al. (2014)	48.5	39.7	29.8	55.9	23.0	29.1	41.9	37.7	38.2
Kolitsas et al. (2018)	82.4	72.4	34.1	35.2	50.3	38.2	61.9	52.7	53.4
Broscheit (2019)	79.3	-	-	-	-	-	-	_	-
Martins et al. (2019)	81.9	-	-	-	-	-	-	-	-
van Hulst et al. (2020)	80.5	72.4	41.1	50.7	49.9	35.0	63.1	58.3	56.4
Cao et al. (2021)	83.7	73.7	54.1	60.7	46.7	40.3	56.1	50.0	58.2
Zhang et al. (2022)	<u>85.8</u>	71.0	53.5	<u>67.8</u>	54.1	41.7	58.9	49.6	60.2
M2E (Only)	84.6	70.2	52.4	67.5	53.8	40.3	58.7	49.5	59.6
E2M (Only)	84.3	71.5	<u>54.0</u>	66.9	52.6	<u>41.8</u>	58.1	50.2	59.9
Combine (M2E+E2M)	84.2	70.5	51.7	66.5	54.1	40.8	57.7	49.5	59.4
${ m MESS(M2E+E2M)}$	86.4	72.8	53.3	68.5	55.2	42.6	<u>60.0</u>	53.1	61.5

 Table 2. InKB Micro F1 on the in-domain and out-of-domain test sets on the GERBIL

 benchmarking platform. For each dataset, **bold** indicates the best model, and <u>underline</u> indicates the second best.

N3-Reuters-128 (R128), N3-RSS-500 (R500) [34], and OKE challenge 2015 and 2016 (OKE15 and OKE16) [30].

Details. We first pre-train on Wikipedia¹ and then fine-tune it on the above eight datasets. Our models are implemented with Python 3.8 and Pytorch 1.9.0 and trained on machines with 4^* NVIDIA-V100-32G GPUs. All text encoders follow the BERT, and the dimension of the output embedding is linearly projected from 768 to the desired size.

Baselines. For EL, we compare with various existing state-of-the-art systems. Among them, Hoffart et al. [18] and van Hulst et al. [39] perform mention recognition first and then entity disambiguation, which is limited by the accuracy of mention detection. However, this limitation can be effectively alleviated with a powerful named entity recognition system [1]. Broscheit [21] proposes a neural end-to-end EL system that jointly discovers and links entities in a text document, which considers all possible spans as potential mentions and learns contextual similarity scores over their entity candidates that are useful for both MD and ED decisions. EntQA [45] converts EL into question answering, quickly retrieves candidate entities, and then examines documents to find mentions.

¹ It is based on the 2019/08/01 Wikipedia dump pre-processed by Petroni et al. [31].

4.2 Results

This section discusses how MESS compares to the previous SOTA models on ED and end-to-end entity linking tasks.

ED. The results on ED are shown in Table 1, with an average F1 score improvement of 1% between our MESS and the suboptimal system. Overall, the advantage is not significant, and this is because ED has been steadily studied as an upstream task for more than a decade and benchmarked on those datasets. The models reported in Table 1 achieve relatively good results even though some work is four or five years old. Besides, we separately evaluate the M2E and E2M models incorporating SS enhancement. The results indicate that neither can compete with two-way MESS, in which only M2E (Only) had a slightly higher F1 in AQUAINT than MESS (M2E+E2M). Note that the average performance when combining M2E and E2M is worse than alone.

EL. The results on EL are shown in Table 2, where the improvement is more obvious. *MESS* is AIDA's best in-domain system and performs well on the out-of-domain setting except for OKE15 and OKE16. It is worth noting that MESS has lower performance on OKE15 and OKE16 because the two datasets are annotated with coreference (pronouns and common nouns linked to entities), and our model is not specially trained for this. In contrast, most other systems have a mention detection component in their pipeline that can be trained or biased to account for these situations. Specifically, *MESS* achieves state-of-the-art results on four datasets and suboptimal on three datasets, and the average F1 score is 1.3% higher than the suboptimal model. Note that the average performance when combining M2E and E2M is worse than alone. Moreover, among the four models we built, except that the performance of E2M (Only) on Der is slightly higher than that of *MESS*, *MESS* yields the highest F1 score on all other datasets.

Models	AIDA	MSNBC	AQUAINT	ACE2004	CWEB	WIKI*	Avg.
M2E(-SS)	90.8	92.5	89.3	89.9	76.4	85.7	87.4
M2E(+SS)	92.4	93.9	90.6	90.8	78.3	86.5	88.8
E2M(-SS)	91.6	92.8	90.1	90.5	76.4	85.9	87.9
E2M(+SS)	93.3	93.6	90.5	91.0	77.2	86.8	88.7
MESS(M2E+SS)	93.7	94.2	89.8	91.8	78.5	88.1	89.4
MESS(E2M+SS)	94.3	94.4	90.6	90.9	78.2	87.5	89.3
MESS(Both-SS)	92.9	93.6	89.5	90.4	77.3	86.7	88.4
${ m MESS(Both+SS)}$	94.8	94.7	90.3	91.5	78.6	88.9	89.8

Table 3. Effect of SS modules on MESS F1 scores in ED. "+SS" and "-SS" indicate whether to apply the SS module. M2E and E2M mean using the result of one module alone to perform ED, while MESS combines M2E and E2M to achieve ED.

Models	AIDA	MSNBC	Der	K50	R128	R500	OKE15	OKE16	Avg.
M2E(-SS)	82.3	68.4	50.5	65.7	53.1	38.6	55.9	48.2	57.8
M2E(+SS)	84.6	70.2	52.4	67.5	53.8	40.3	58.7	49.5	59.6
E2M(-SS)	83.5	69.8	51.4	66.3	52.6	39.7	56.6	49.2	58.6
${ m E2M}(+{ m SS})$	84.3	71.5	54.0	66.9	52.6	41.8	58.1	50.2	59.9
$\overline{\mathrm{MESS}(\mathrm{M2E+SS})}$	85.7	71.8	52.4	67.5	55.3	41.9	58.6	53.8	60.9
MESS(E2M+SS)	86.6	71.4	54.0	67.8	54.5	42.3	59.7	53.0	61.2
MESS(Both-SS)	84.5	70.8	51.3	66.7	52.6	41.0	57.9	51.3	59.5
${ m MESS(Both+SS)}$	86.4	72.8	53.3	68.5	55.2	42.6	60.0	53.1	61.5

 Table 4. Impact of the SS module in EL task on the F1 score of the model with different settings.

4.3 Ablation Studies

We report the main results of ED and EL in Sect. 4.2. Here, we disassemble MESS to explore why it is effectively enhanced. The impact of SS in ED is explored in Table 3. We divide it into two blocks. One studies the effect of SS in M2E (only) and E2M (only), and the other investigates the influence of SS in one direction or both in MESS. For the convenience of comparison, we denote M2E(only) and E2M(only) and MESS(M2E+E2M) as M2E(+SS) and E2M(+SS), MESS(Both+SS) respectively. From the first block of Table 3, we notice that the scores of F1 of both M2E and E2M obviously decrease when the SS module is removed. Especially M2E(+SS) and M2E(-SS) on the CWEB dataset, the gap is a maximum of 1.9%. For MESS, the worst performance is that both directions are not augmented with SS modules, and the overall result of the unidirectional fusion is also worse than simultaneous integration.

Similarly, we investigate the role of SS in the EL task in Table 4. We find that the F1 score also clearly drops when SS is not integrated. Specifically, M2E(+SS) has an average improvement of 1.8% compared to the case of not using SS, while E2M(+SS) has an average gain of 1.3%, which is a relatively obvious improvement. In *MESS*, the performance of integration SS in either M2E or E2M is better than the case that none of them is fusion. Moreover, the average F1 score is higher when M2E and E2M are injected into SS than when only one is injected.

5 Conclusion

Existing Entity Linking methods cannot effectively utilize the structured information in KB and face the issue of insufficient features in a given document. To address the limitations, we propose an effective and efficient MESS framework that consists of four important components: (1) Mention-to-Entities, (2) Entity-to-Mentions, (3) Semantic Synchronization, and (4) Dialogue. These modules work collaboratively to enhance the model's performance. Extensive experiments on public datasets indicate that our proposed model MESS outperforms the state-of-the-art methods.

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Session Target Pair: User Intent Perceiving Networks for Session-Based Recommendation

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Abstract. Session-based recommendation (SBR) aims to predict the next-interacted item based on an anonymous user behavior sequence (session). The main challenge is how to decipher the user intent with limited interactions. Recent progress regards the combination of consecutive items in the session as intent. However, these methods, which merely depend on the session, ignore the fact that such limited interaction within the session may not entirely express user intent. Therefore, it constrains the expression of diverse user intent without considering the candidate items to be predicted, which can be regarded as target intent, leading to a sub-optimal inference of user behavior. To solve the problem, we propose a novel Intent Alignment Network for session-based recommendation (IAN), which models intent from both session and target perspectives. Specifically, we propose that session-level intent is explicitly formed by weighted aggregation of successive items, whereas targetlevel intent is composed of interacted and undiscovered items that are compatible. Based on it, we devise an intent alignment mechanism to ensure consistency between these two types of intent and obtain mutual intent representation. Finally, a gated mechanism is used to fuse mutual intent and target intent to generate session representation for prediction. Experimental results on three real-world datasets exhibit that IAN achieves state-of-the-art performance.

Keywords: Recommender System \cdot Session-based Recommendation \cdot Multi-Intents Perceiving

1 Introduction

Recommender systems are pivotal in managing information overload by effectively understanding user-profiles and long-term behavior [17]. Unfortunately, such information may be unavailable in real-world scenarios due to privacy concerns. In response, SBR emerges and aims to predict the next item based on the anonymous user's short-term interaction sequence [5].

In the early period, researchers concentrated on learning the different dependencies between items within a session, whose effectiveness has been validated. For example, recurrent neural networks (RNNs)-based models [5,6] are devoted to learning the sequential dependency of items, while graph neural networks (GNNs)-based models [1,4,21,23] are to extract the more complex dependency from constructed graph than single transition from consecutive items. However, these approaches only take individual items as basic units to extract user preference, neglecting to dive deeply into the user intent arising from the combination of items.

The user intent is the driving force for generating behavior sequences [2, 12]. More recently, these approaches [3, 7, 19, 27] construct the multi-level intent via regarding consecutive items with different lengths in the session as intent units and achieve a better inferring use preference. However, these approaches, which merely depend on the session, ignore the fact that such limited interaction within the session may not entirely express user intent. For instance, given the <session, target> pair in toy examples 1 and 2, if a model learned intent representation merely depends on the current session, target (iPhone)-related intent can be expressed in toy example 1 due to the previously appeared target, while target (soccer)-related intent is not directly easy to express in toy example 2 due to the previously unseen target. This may constrain the model's ability to express diverse user intent without considering the candidate items to be predicted, leading to a suboptimal inference of user behavior. Therefore, when we consider both the session and target to express user intent, it is helpful to model the complex user intent in the scenario with different patterns, which are neglected by existing works.

Toy example 1: *iPhone*, *shirts*, *pants*, *iPhone*, *shoes*, *coat*, *bag* => *iPhone*

Toy example 2: *iPhone*, *Samsung phones*, *earphones*, *watch*, *shoes* => soccer

To solve this limitation, we propose to explicitly model user intent with session and target perspectives.

- (1) From the session perspective, we consider each item centered to aggregate its corresponding contextual information with different weights as session-level user intent. This is inspiration from any item in the session that may directly trigger or even be the same as the target, according to [10]. For example, as shown in toy example 1, these approaches extract intent through an equal fusion of consecutive items in related work [27], which may classify the user as having clothing-related intents due to frequently appearing clothing, neglecting that the user is also the phone-related intent even though the iPhone is regarded as the next behavior (target). Therefore, such an unbias combination of consecutive items may suppress the target-related information, leading to insufficiently expressed user intent.
- (2) From the target perspective, we consider the set of candidate items as target-level user intent. The reason rooted in this is that the limited expression of short-term sequence may not entirely express the user's actual intents [25]. Furthermore, these studies show that repeat consumption and exploratory phenomena are general user behavior patterns [13,14]. Namely,

the ground truth (target) of the session may belong to interacted items and may belong to unexplored items. Therefore, we take the explored items and the remaining items into consideration to boost the diversity of intent and understanding of complex user behavior.

To this end, we propose novel Intent Alignment Networks for session-based recommendation (IAN), which incorporates session-level and target-level intents to comprehensively infer user behavior. For session-level intent, we employ selfattention networks to extract item-centered, relevant information by weighted aggregation of contextual items. For the target-level intent, we leverage the average operation of both the explored item sets and the unexplored item sets (namely the candidate sets or items dictionary) to introduce more information for exploring the intent diversity. Next, we design an intent alignment mechanism centered on attention, which derives and extracts shared representations through alignment vectors calculated between session-level and target-level intents. Finally, we implement a gated mechanism to adaptively combine shared and target-level intent into session representations for predicting the next item.

In summary, the main contributions of this work include:

- We propose a novel IAN model, that fully excavates the hidden intent in complex user behavior under the guidance of session and target intent perspectives, boosting recommendation performance.
- We devise an intent alignment mechanism to establish consistency between defined session-level and target-level intent representations, solving the problem of insufficient expression in the diversity of user intents.
- Experimental results on three benchmark datasets show the effectiveness and superiority of IAN compared with state-of-the-art methods.

2 Related Work

In the early stage, the research on the SBR task uses matrix factorization and Markov chains to extract the sequential dependency of adjacent items [15]. Later, benefiting from the development and excellent performance of neural networks, RNN-based SBR models are proposed [5,6] and make a significant improvement in recommendation performance compared with conventional models. For example, Li et al. [6] devise a neural attentive network to extract the main purpose of the last hidden state in Gated Recurrent Unit [5], while Liu et al. [9] explicitly emphasize the importance of the last click via the Vanilla attention mechanism. To fully take advantage of the session with limited interaction, GNN-based models are proposed, whose core is to construct a graph based on the session and apply GNN to capture the complex transition between adjacent items, boosting the expression of the item transitions compared to such a single sequential transition [21]. Following this idea, kinds of GNN-based variant models are proposed, including strictly position information [1], additional information [20, 25], the hyper-transition exploration between items [23], and the type of transition [4,8], etc. In parallel with the above, the self-attention mechanism has been introduced



Fig. 1. Overall framework of proposed SBR model (IAN).

to the recommendation system [16]. For example, Xu et al. [24] leverage the complementary nature of both graph neural networks and self-attention mechanisms to enhance the dependency between items. Yuan et al. [26] decrease the impact of unrelated items based on self-attention. However, these approaches only take individual items as basic units to extract user preference, neglecting to dive deeply into the user intent arising from the combination of items.

Recently, the opinion has been accepted that user behavior is driven by intents [2,12]. Following this idea, there are emerging intent-perceiving models in SBR. For example, Wang et al. [19] use a sliding window on the current session to capture group intent. Guo et al. [3] propose multi-level consecutive items as an intent unit to alleviate losing sequential information. Li et al. [7] disentangle the intents from each item into micro and macro manners for capturing the dynamic intents of users. More Recently, Zhang et al. [27] proved the helplessness of GNN's propagation and constructed multi-level recent intents to optimize recommendation performance. However, the above multi-intents-based methods, which merely depend on the session, ignore the fact that such limited interaction within the session may not entirely express user intent. It constrains the expression of diverse user intent without considering the candidate items to be predicted, which can be regarded as target intent, leading to a suboptimal inference of user behavior. Therefore, we propose the idea that models intent from both session and target perspectives, which is ignored by existing works.

3 Methodology

In this section, we formulate the problem and elaborate on our proposed model, whose basic structure is given in Fig. 1. First, we introduce the session-level intent representation module, whose core is self-attention networks, to obtain session intent representation, as shown in the session-level intent representation module. Then, we use the average operation on the items dictionary to obtain target-level intent representation, as shown in the target-level intent representation module.

After that, we design an intent alignment mechanism to fuse target-level and session-level intent representation and generate session representation. Finally, we use the learned session representation to generate the recommendation list.

3.1 Problem Statement

Let $V = \{v_1, v_2, \ldots, v_{|V|}\}$ be the candidate item sets (or items dictionary). Let $S = [s_1, s_2, \ldots, s_t]$ be the anonymous user's interaction sequence. Where $s_i \in V(1 \leq i \leq t)$ represents the interacted item at time step *i*. We embed each item into the feature space and denote $\mathbf{e}_j \in \mathcal{R}^d$ as the vector representation of item $v_j \in V$. Where *d* represents the dimensionality. Therefore, the goal of session-based recommendation is to learn a session representation for predicting the next-interacted item, namely s_{t+1} , for a given *S*.

3.2 Session-Level Intent Representation Module

To avoid insufficiently extracting and expressing the intent hidden in the current session, as illustrated in the introduction section, we propose the idea of modeling the importance weight for each item in the combination of consecutive items. Based on it, we generate the item-centered contextual representation using self-attention networks and regard it as session-level intent representation. This process can be formulated as follows:

Given the session S, we first introduce the reverse position encoding to each item s_i in the session for reserving the sequential order of user behavior. Therefore, item s_i with position information is encoded by:

$$\mathbf{e}_{i}^{P} = \operatorname{concat}(\mathbf{e}_{i}, \mathbf{p}_{t+2-i}) \tag{1}$$

where $\mathbf{E}^p = [\mathbf{e}_1^P, \mathbf{e}_2^P, \dots, \mathbf{e}_t^P] \in \mathcal{R}^{t \times 2d}$ denotes the entire representation of S, $\mathbf{P} = [\mathbf{p}_1, \mathbf{p}_2, \dots, \mathbf{p}_{t+1}] \in \mathcal{R}^{(t+1) \times d}$ denotes the reverse position embedding of Sand its corresponding next behavior. After that, we employ a scaled dot-product attention network to obtain the item-centered contextual representation [18], including weight calculation and weighted information aggregation.

$$\mathbf{M}_{w} = \operatorname{softmax}(\frac{1}{\sqrt{2d}}\mathbf{Q}\mathbf{K}^{T}) \quad \mathbf{C}_{agg} = \mathbf{M}_{w}\mathbf{V}$$
(2)

 \mathbf{M}_{w}^{ik} is the correlation weight from item s_{i} to item $s_{k} \in S$, and $\mathbf{M}_{w}^{i:}$ represents the semantic similarity vector from item s_{i} to each item in the current session. Therefore, when we use matrix multiplication between \mathbf{M}_{w} and \mathbf{V} , \mathbf{C}_{agg}^{i} represents s_{i} have weighted contextual aggregation representation for each item in the current session. It is noted that the weighted matrix is the most significant difference between our proposed model and the recent multi-intent-based model. Recent multi-intent-based models use the mean combination of consecutive items without considering the different importance of each item. Where $\mathbf{Q} = \text{SELU}(W_0\mathbf{E}^p + b_0)$ as query vector, $\mathbf{K} = \mathbf{V} = \mathbf{E}^p$ as key and value vectors,

respectively. $\mathbf{W}_0 \in \mathcal{R}^{2d \times 2d}$, $\mathbf{b}_0 \in \mathcal{R}^{2d}$ are learnable parameters. $\sqrt{2d}$ is the scale factor. SELU and Softmax are activation functions.

To increase the ability of non-linearity in dot-product self-attention, we use a feed-forward network with the ReLU activation function. Based on it, we also leverage a residual connection to reserve the raw information. The formulation is as follows:

$$\mathbf{C}^{l} = \operatorname{ReLU}\left(\mathbf{C}_{agg}W_{1} + \mathbf{b}_{1}\right)\mathbf{W}_{2} + \mathbf{b}_{2} + \mathbf{C}_{agg}$$
(3)

where $\mathbf{W}_1, \mathbf{W}_2 \in \mathcal{R}^{2d \times 2d}$ are learnable weight matrices, and $\mathbf{b}_1, \mathbf{b}_2 \in \mathcal{R}^{2d}$ are learnable bias vectors.

3.3 Target-Level Intent Representation Module

To mutually consider the repeat consumption and exploration habits in user behavior patterns, we propose the idea that interacted items and unexplored items are compatible. Based on it, we conduct an aggregation operation on these candidate items (denoted as item dictionary) and obtain target-level intent representation. Specifically, We first calculate the average operation in each item from the candidate set to illustrate any possibilities of user behavior at the next time step. Then, we also use the reverse position encoding to serve the sequential information in the process of the generated next behavior. Thus, target-level intent representation \mathbf{S}_q^p is represented by

$$\mathbf{S}_{g} = \frac{1}{|V|} \sum_{j=1}^{|V|} \mathbf{e}_{j} \qquad \mathbf{S}_{g}^{p} = \operatorname{concat}(\mathbf{S}_{g}, \mathbf{p}_{1})$$
(4)

Although it is easy to think of averaging the candidate set, what we want to emphasize here is that we may enhance the diversity of user intention expression by introducing unexplored items. Limited sequences may not be able to express the complete user intention, which existing multi-intent-based models ignore. Where $\mathbf{p}_1 \in \mathbf{P}$ represents the reverse position embedding of the next behavior.

3.4 Intent Alignment Mechanism Module

The goal of separately modeling intents from session and target perspectives is to generate a better understanding of complex user behaviors. Therefore, to be consistent with the session and target intent, which belong to different aspects of user intent representation, we devise an intent alignment mechanism inspired by [11]. This process can be formulated as follows:

Firstly, we employ an attention mechanism to obtain mutual representation between session-level and target-level representation, including alignment vector calculation and corresponding information aggregation with weighted.

$$\alpha_i = align(\mathbf{S}_{\mathbf{g}}^{\mathbf{p}}, \mathbf{C}_i^l) \quad \alpha_i = \mathbf{u}^T \sigma \left(\mathbf{W}_3 \mathbf{S}_g^p + \mathbf{W}_4 \mathbf{C}_i^l + \mathbf{b}_3 \right) \qquad \mathbf{S}_g^r = \sum_{i=1}^t \alpha_i \mathbf{C}_i^l \quad (5)$$

Dataset	# of train #	$\# ext{ of test } \#$	of items Av	g.length max	k.length
Tmall	351,268	$25,\!898$	40,728	6.69	39
RetailRocket	433,643	$15,\!132$	$36,\!968$	5.43	284
Diginetica	719,470	60,858	$43,\!097$	5.12	69

 Table 1. Statistics of the used datasets.

where α_i represents the alignment weight between \mathbf{C}_i^l and \mathbf{S}_g^p . \mathbf{S}_g^r represents mutual representation between two types of intents. $\mathbf{W}_3, \mathbf{W}_4 \in \mathcal{R}^{2d \times 2d}$ are learnable weight matrices. $\mathbf{u}, \mathbf{b}_3 \in \mathcal{R}^{2d}$ are learnable bias vectors.

After acquiring mutual representations \mathbf{S}_{g}^{r} , we employ a gate mechanism to adaptively aggregate \mathbf{S}_{q}^{r} and \mathbf{S}_{q}^{p} to generate the final session representation:

$$\beta = \text{sigmoid} \left(W_5[\mathbf{S}_g^r] | \mathbf{S}_g^p] \right) \qquad \mathbf{z} = (1 - \beta) \odot \mathbf{S}_g^r + \beta \odot \mathbf{S}_g^p \qquad (6)$$

where \mathbf{z} denotes the final session representation, $W_5 \in \mathcal{R}^{2d \times 4d}$ is the learnable parameter, || represents the operation of concat, and \odot is dot product operation.

3.5 Prediction and Training

After obtaining session representation, following [26,27], we use weighted normalization to make the training process more stable and insensitive to hyperparameters. Then, we use it to figure out the likelihood that the user will be interested in the next item, whose probability is calculated by:

$$\hat{\mathbf{S}}_{f} = w_{k} \operatorname{L2Norm}(\mathbf{z}) \quad \hat{\mathbf{v}}_{i} = \operatorname{L2Norm}(\mathbf{e}_{i}) \quad \hat{y}_{i} = \operatorname{softmax}\left(\hat{\mathbf{S}}_{f}^{T} \hat{\mathbf{v}}_{i}\right) \tag{7}$$

where \hat{y}_i indicates the probability of the item in the candidate item set V, L2Norm is the L2 normalization function, and w_k is the normalized weight.

Finally, we pick the top-K items in $\hat{\mathbf{y}} = \{\hat{y}_1, \hat{y}_2, \dots, y_{|V|}\}$ for the recommendation, and leverage cross-entropy of the prediction and the ground truth to optimize model. It is written as follows:

$$L(\mathbf{y}, \hat{\mathbf{y}}) = -\sum_{i=1}^{t} y_i \log\left(\hat{y}_i\right) \tag{8}$$

where \mathbf{y} is the one-hot encoding vector of the ground truth.

4 Experiments

In this section, we first illustrate experiment setups, including datasets, baselines, evaluation metrics, and Implementation Details. Then, we analyze comparison experimental results.

4.1 Experiment Setups

Datasets. We evaluate the proposed model on three benchmark datasets, namely $Tmall^1$, $RetailRocket^2$, $Diginetica^3$. Tmall is from the IJCAI-15 competition and consists of shopping logs of unnamed users on the Tmall online shopping platform. RetailRocket is released by an e-commerce corporation for the Kaggle competition and contains users' browsing activity. Diginetica comes from CIKM Cup 2016 and describes the music-listening behavior of users. Following [22,23], we conduct preprocessing over each dataset. Specifically, sessions with a length of 1 and items that appeared fewer than 5 times are excluded. Then, the latest data (such as the data from last week) is set to be a test set, and previous data is used as the training set. Additionally, we use a sequence splitting preprocess method to augment session $S = s_1, s_2, ..., s_n$ in these datasets, and generate multiple sessions with corresponding labels ($[s_1, s_2]; s_3$), ($[s_1, s_2, s_3]; s_4$), ..., ($[s_1, s_2, ..., s_{n-1}]; s_n$). The statistics of the datasets are presented in Table 1.

Evaluation Metrics. Following [5,27], we widely adopt P@K and MRR@K to evaluate the recommendation results. P@K represents the proportion of test cases which have correctly recommended items in the Top-K ranking list. MRR@K represents the average reciprocal rank of the desired item. Here, we set K to 20.

Baseline Models. We compare our method with the most relevant and state-of-the-art models. The following ten baseline models are evaluated:

- **FPMC** [15]: It combines the Markov chain and matrix factorization to model the sequential dependency between items.
- **GRU4REC** [5]: It utilizes GRU to model the sequential dependency with the training of session parallel mini-batches and pair-wise loss functions.
- NARM [6]: It combines sequential behavior by the GRU encoder and the main purpose captured by the attention mechanism for recommending.
- **STAMP** [9]: It explicitly emphasizes the importance of short-term interest and combines it with long-term interest to recommend items.
- **SR-GNN** [21]: It constructs an item-transition graph and uses GNN to learn the complex dependency between items.
- **GC-SAN** [24]: It utilizes the complementary of adjacent dependency obtained from GNN and global dependency obtained via self-attention to boost recommendation performance.
- **TAGNN** [25]: It combines the candidate list into target attentive graph neural network for enhancing the expression of user interest diversity.
- S²-DHCN [23]: It uses a hypergraph convolution network to capture complex high-order relations between items and introduces a self-supervised task to enhance the recommendations.
- **HIDE** [7]: It disentangles the intent of each item into micro and macro views to capture the dynamic intents of users.

 $^{^{1}}$ https://tianchi.aliyun.com/dataset/dataDetail?dataId=42.

 $^{^{2}}$ https://www.kaggle.com/retailrocket/ecommerce-dataset.

³ http://cikm2016.cs.iupui.edu/cikm-cup/.

	Tmall		Retai	ilRocket	Diginetica	
	P@201	MRR@20	P@20	MRR@20	P@20	MRR@20
FPMC(WWW'10)	16.06	7.32	32.37	13.82	26.53	6.95
GRU4Rec (ICLR'16)	10.93	5.89	44.01	23.67	29.45	8.33
NARM (CIKM'17)	23.30	10.70	50.22	24.59	49.70	16.17
STAMP (KDD'18)	26.47	13.36	50.96	25.17	45.64	14.32
SR-GNN (AAAI'19)	27.57	13.72	50.32	26.57	50.73	17.59
GC-SAN(IJCAI'19)	25.38	12.72	51.63	27.72	51.82	17.82
TAGNN(SIGIR'20)	34.60	16.62	53.86	28.54	52.45	18.31
S^2 -DHCN(AAAI'21)	31.42	15.05	53.66	27.30	53.18	18.44
HIDE(SIGIR'22)	37.12	17.19	51.25	26.20	53.72	18.37
Atten-Mixer(WSDM'23)	37.16	<u>18.71</u>	56.01	28.57	<u>53.86</u>	18.27
IAN(Proposed)	37.41	19.27	56.98	31.14	54.14	19.29

Table 2. Performance comparison between IAN and baselines over three benchmark datasets. The boldface is the best result, and the underline is the second best result.

- Atten-Mixer [27]: It constructs recent intent via an equal combination of items and uses it for multi-level reasoning.

Implementation Details. For the general setting, the embedding size is 100, the batch size is 100, the learning rate is 0.001, and the epoch is set to 30. For the baseline models, we refer to their best parameter setups reported in the original papers and directly report their results if they are in line with general settings, evaluation metrics, and datasets. Otherwise, we record the reproduced results under the public code. In addition, models are trained on a single NVIDIA A100 Tensor Core GPU, and hyperparameter w_k is set to 20 in IAN.

4.2 Overall Performance

To evaluate the effectiveness of IAN, we report the comparison results with the state-of-the-art baselines. From Table 2, we draw the following observations:

- Compared with GRU4Rec and FPMC, NARM, STAMP, SRGNN, GC-SAN, TAGNN, and S²-DHCN outperform better on three benchmark datasets. This phenomenon shows that explicitly or implicitly emphasizing the recent click leads to a better understanding of the user's behavior. Furthermore, TAGNN and S²-DHCN achieve more excellent recommendation results generally on the three datasets compared to these models based on merely the current session, including NARM, STAMP, SRGNN, and GC-SAN. It indicates that introducing additional information (such as other sessions and candidate items) is helpful to boost the expression of session representation. What's more, for TAGNN, we also see that it is obviously superior in terms of recommendation performance and expression of interest diversity by explicitly
| Madala | Т | mall | Reta | ilrocket | Diginetica | | |
|----------------|-------|--------|-------|----------|------------|--------|--|
| Models | P@20 | MRR@20 | P@20 | MRR@20 | P@20 | MRR@20 | |
| IAN-base | 37.74 | 18.95 | 53.08 | 27.89 | 50.96 | 17.77 | |
| IAN-NO-Context | 36.13 | 19.03 | 56.14 | 30.62 | 53.63 | 18.57 | |
| IAN-NO-Session | 35.64 | 17.82 | 53.01 | 27.67 | 50.67 | 17.36 | |
| IAN-NO-Target | 37.54 | 18.82 | 56.93 | 30.92 | 54.39 | 19.14 | |
| IAN | 37.41 | 19.27 | 56.98 | 31.14 | 54.14 | 19.29 | |

 Table 3. Performance comparison of variant models in IAN on three datasets. The boldface is the best result

considering candidate items, validating our opinion that limited interaction may not entirely express user intent.

- Among multi-intent-based models, including Atten-Mixer and HIDE, we can conclude that Atten-Mixer completely outperforms other baseline models, except for MRR@20 on the Diginetica dataset. HIDE has more improvement than individual item intent-based models on the Tmall and Diginetica. The above analysis shows that explicitly extracting multi-level intent representation generally results in better performance. Furthermore, except that Atten-Mixer has comparable performance on the Diginetica dataset with the metric of MRR@20, Atten-Mixer is better than hiding on the three datasets. It supports the view proposed in this paper that the combination of consecutive items as intent units is a good base to express intent.
- Our proposed IAN surpasses the overall baselines on these three datasets, demonstrating the usefulness of the proposed multi-intent perceiving consisting of session and target-level perspectives. Compared to existing models, the multi-intents benefit the model in two main aspects: (i) The devised session-level intent representation module considers each item centered to aggregate its corresponding contextual information with different weights, ensuring sufficient extraction of user intent from the interacted sequence. (ii) The devised target-level intent representation module consists of the unexplored items and explored items, leading to introducing more intent information, which is ignored by the interacted sequence. (iii) The designed intent alignment networks establishes consistency between defined session-level and target-level intent representations, which enhances expression in the diversity of user intents.

4.3 Model Analysis and Discussion

In this subsection, we conduct an in-depth model analysis study, aiming to further understand the framework of IAN. Abalation Study. To profoundly comprehend the contribution of the multiintents component in IAN, we designed four variants: IAN-base, IAN-NO-Context, IAN-NO-Session, and IAN-NO-Target. IAN-base uses an unbiased combination of each item in the current session as session-level intent representation in the IAN. IAN-No-Context uses a vanilla item representation instead of the representations obtained by the self-attention network in IAN. IAN-NO-Session removes the session-level intent representation, while IAN-NO-Target removes the target-level intent representation in the IAN. Table 3 illustrates the comparison results.

From Table 3, we have the following observation: (i) Compared with the IANbase, IAN makes more contributions to the Tmall, Retialrocket, and Diginetica datasets in the metrics of MRR@20. It shows that using weighted items instead of equally chosen combinations in the current session is a more positive method of expressing intent. Then, both IAN-NO-Context and IAN outperform IANbase, demonstrating that explicitly considering the importance of each item in the session boosts the recommendation performance. In addition, the observation that IAN is better than IAN-NO-Context, also illustrates that extracting item-centered relevant information by weighted aggregation of contextual items makes more progress in understanding user behavior. Therefore, the above analvsis result supports the proposed opinion that an equal combination of consecutive items makes for insufficient information extraction in expressing session representation. (ii) Compared with IAN-NO-Session, which only utilizes targetlevel intent representation, and IAN-NO-Target, which only utilizes session-level intent representation related to the target perspective, IAN achieves the best results in the metrics MRR@20. It shows that simultaneously considering sessionlevel and target-level intent representation helps make user diversity intent more expressive.

Cassian	a Madala	Т	mall	Retai	lRocket	Digi	netica
Session	s models	P@20	MRR@2	0 P@20 I	MRR@20) P@20 I	MRR@20
	Atten-Mixer	28.81	14.78	62.74	34.39	54.41	19.21
short	IAN	29.07	15.77	63.11	36.39	54.90	19.99
	Atten-Mixer	39.56	19.86	41.34	17.95	48.73	14.76
long	IAN	40.01	20.35	44.83	20.72	51.69	17.06

 Table 4. Performance comparison between IAN and the Atten-Mixer in long-group and short-group sessions. The best results are marked in bold.

Impact of Session Length. In real-world situations, sessions with various lengths are common, and the long session may have more intent compared with the short session generally. Thus, it is interesting to know how stable our proposed model IAN as well as the Atten-mixer are when dealing with them. To maintain the fairness of comparison, we follow the process method of [9,22].

The method splits the session into long-group and short-group sessions according to the average length of each dataset, as shown in Table 1. Therefore, the long group contains sessions whose length is greater than the average length, while the short group contains sessions whose length is less than or equal to the average length. From Table 4 illustrated comparison result, we can observe the conclusions that IAN outperforms Atten-mixer on three datasets with different session lengths under the metric of P@20 and MRR@20. It demonstrates the adaptability and effectiveness of our proposed model IAN in deciphering the hidden user intent in the interaction sequence, allowing us to better infer user behavior both in short and long sessions.

Models	Т	mall	Reta	ailrocket	Diginetica			
	P@20	MRR@20	P@20	MRR@20	P@20	MRR@20		
Atten-Mixer	65.72	38.10	90.73	63.30	85.81	47.69		
IAN-base	66.47	38.44	88.88	63.40	88.70	55.44		
IAN	68.39	39.64	93.58	70.61	91.24	60.67		

Table 5. Performance comparison between IAN variants and the Atten-Mixer in repeatsessions. The best results are marked in bold.

Table 6. Case studies of our IAN model compared with Atten-mixer on the Tmalldataset. The blood represents the target of the given session.

# session	Atten-mixer IAN
1 4766, 24651 , 33326, 24651 , 32010, 4766	X V
2 15482 , 1187, 1187, 33085, 15482 , 14175	X 🗸
3 32559 , 36695, 8875, 20062, 31471	XV
4 30369, 7314 , 29228, 32193, 35145	X 🗸
5 35080, 35079, 35083, 35079, 35079, 35078	X 🗸
6 9803, 25868, 26750 , 33696, 23853, 20394, 27	X 🗸
7 34366, 34730, 32879, 17973, 34730, 29236, 39664, 2	8420 , 34366 X V

Case Study. To straightforwardly perceive intent that indeed contains richer relevant information in recommendation lists, we take the repeat session to estimate the effectiveness. The reason we choose repeat sessions as the case is that the ground truth of each session appears in the current session (input). This is nice for us to validate because we know the association of the session with the target. Therefore, we conduct a comparison experiment on these obtained repeat sessions, whose principle of pre-processing is that the target (ground truth) has been interacted with by the user. Experimental results are shown in Table 5. Compared with the Atten-Mixer and IAN-base, it can be observed that the IAN is superior in the three datasets with metrics of P@20 and MRR@20. This proves that our proposed model indeed benefits from extracting intent from the session and target perspectives.

In more detail, we pick several case studies analyzed on the Tmall dataset, as shown in Table 6. If the model predicts that the Top-20 recommendation list contains the bold (target) in the session, it is judged to have done the right thing. From Table 6, we can observe that the target can be repeatedly exposed, and its corresponding position may be anywhere within these session cases, such as first, second, etc. In #1 and #2, we observe that Atten-Mixer failed to express the target-related intent and made an incorrect prediction even though the target was multiple times clicked within these sessions. What's more, when we apply Atten-Mixer to different lengths of sessions when the target is repeated only once, including shorter sessions (#3 and #4), sessions with the same length (#5), and longer sessions (#6 and #7), we find that Atten-Mixer also struggles to extract user intent. In contrast, our proposed model predicted the target of the above sessions, demonstrating its effectiveness in capturing user intents.

These examples highlight the importance of simultaneously considering session-level intent and target-level intent, which are ignored in the Atten-mixer. Therefore, our proposed model, IAN, solves the problem of insufficient expression in the diversity of user intents for the SBR task.

5 Conclusion

In this paper, we propose a novel IAN model that fully excavates the hidden intent in complex user behavior under the guidance of session and target intent perspectives, solving the problem of insufficient expression in the diversity of user intents in the related works. Specifically, we first propose that session-level intent is explicitly generated by the weighted aggregation of consecutive items. While the target level intent consists of the interacted items and unexplored items compatible. Based on it, we devise an intent alignment mechanism to ensure consistency between these two types of intent and obtain mutual intent representation. Finally, a gated mechanism is used to fuse mutual intent and target intent to generate session representations for prediction. Experimental results on three real-world datasets exhibit that IAN achieves state-of-the-art performance.

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Hierarchical Fine-Grained Visual Classification Leveraging Consistent Hierarchical Knowledge

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Abstract. Hierarchical fine-grained visual classification assigns multigranularity labels to each object, forming a tree hierarchy. However, how to minimize the impact of coarse-grained classification errors on finegrained classification and achieve high consistency remains challenging. Considering the human ability to progress from understanding generalized concepts to recognizing subtle differences between categories, the proposed novel hierarchy-aware conditional supervised learning method encodes such dependencies within its learned structure. The validity masks based on label hierarchy are designed to control the influence of coarse-grained classifications on fine-grained classifications. In this paper, the graph representation learning is explored to better utilize label hierarchy, integrating hierarchical structural information into the feature representation framework. Experiments on three standard finegrained visual classification benchmark datasets demonstrate the effectiveness of the proposed method, significantly improving the consistency of hierarchical predictions while enhancing the model's understanding of label hierarchy compared with the state-of-the-art methods.

Keywords: Fine-grained visual classification \cdot conditional supervised learning \cdot graph representation learning

1 Introduction

Traditional fine-grained visual classification (FGVC) concentrates on the model's ability to recognize objects at the fine granularity. Recent research has been primarily focused on the identification of the most distinctive features [1–3]. Hierarchical fine-grained visual classification (HFGVC) differs from traditional FGVC in its approach to organizing similar categories into coarse-grained concepts and assigning hierarchical multi-granularity labels to each object. For example, it categorizes from coarse to fine levels, such as from "Albatross" to "Laysan Albatross", thereby establishing a top-down tree-like knowledge structure. A crucial

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Fig. 1. An example label hierarchy of CUB-200-2011. Suppose the ground truth is Black throated Sparrow, then the real-label path Birds \rightarrow Passeriformes \rightarrow Passerellidae \rightarrow Black throated Sparrow can be identified as a consistent prediction. The proposed method is derived from the motivation to leverage consistent hierarchical knowledge.

consideration is how to define fine granularity in a manner that is genuinely meaningful to individuals. As illustrated in Fig. 1, while bird experts might easily identify a specific species, such as "Black throated Sparrow", bird lovers and the general public may find broader categories like "Passerellidae" or "Passeriformes" more comprehensible and accessible. Moreover, the quality of images also influences the ability to recognize objects at different levels. Consequently, HFGVC, in contrast to traditional FGVC tasks, aligns more closely with practical requirements. HFGVC facilitates comprehension and recognition of specific species by leveraging information from higher-level categories. The focus of HFGVC tasks lies in how to effectively use the label hierarchy to enhance the consistency of predictions across multi-granularity levels.

To tackle the challenge of utilizing hierarchical multi-granularity labels and embedding label hierarchy, various methods have been explored [4–6]. These approaches offer more options to annotators with different knowledge backgrounds and improve model performance in managing complex tasks. Notably, multi-layered networks capable of identifying categories at different granularity levels have demonstrated effectiveness. Nevertheless, the separate classification deployment at distinct levels poses specific challenges, particularly in ensuring consistency.

The human learning process typically adopts a top-down methodology, commencing with a comprehension of overarching, global concepts before gradually delving into more specific, detailed categories. However, encoding and learning this top-down dependency presents a challenge in traditional neural network designs. Motivated by the recent advancement in models that addresses subpopulation shift [7], this study explores the application of the conditional supervision training framework within HFGVC to emulate the hierarchical learning pattern observed in humans, integrating it into the neural network's learning architecture. However, this approach carries a potential risk: early misidentification of upper-level concepts could lead to a cascading effect of errors, amplifying inaccuracy in fine-grained category recognition later on. Such occurrences can significantly impact overall classification accuracy and result in inconsistent predictions.

To address this challenge, implementing strategies to minimize the risk of error propagation and enhance the model's robustness and accuracy across all levels is imperative. Our proposed hierarchy-aware conditional supervised learning (HCSL) method aims to deeply leverage consistent hierarchical knowledge into the learning representations. To evaluate the impact of each error, the innovative validity mask mechanism is designed by measuring the distance between the ground truth and the predicted node within the label hierarchy. By integrating with the loss function, it effectively restricts the spread of incorrect information during conditional training. Meanwhile, HCSL ensures the extraction and learning of valuable features from each sample.

In order to better utilize label hierarchy, our framework integrates graph convolutional networks (GCN) [8], leveraging knowledge aggregation from adjacent nodes within the label hierarchy. This approach extends beyond utilizing category information simply from the same parent for classification, encompassing broader contextual relationships and connections among categories. Subsequently, the enriched multi-granularity features obtained through the GCN substantially improve the accuracy and robustness of the classification.

Our main contributions can be summarized as follows:

- The proposed approach employs hierarchical knowledge-based valid masks for conditional training, effectively mitigating the adverse effects of coarsegrained classification on fine-grained classification while improving overall consistency.
- In order to leverage label relationships more effectively, graph representation learning integrates the hierarchical structure of labels into the feature representation framework to extract superior hierarchical features.
- Additionally, a tree-structured granularity consistency rate is introduced as a reliable and intuitive metric for assessing the consistency of model predictions.

2 Related Work

2.1 Fine-Grained Visual Classification

Compared with traditional image recognition, FGVC presents a unique challenge in the field of computer vision. It's characterized by large intra-class variance and small inter-class variance. The difference and difficulty of the task lie in the extremely fine granularity of the categories involved. Early FGVC research [9,10] was heavily dependent on features marked by human annotators, a method that was notably limited. In recent years, the development of deep learning has lifted the mentioned limitations and offered innovative ideas and perspectives [11–14].

Since then, a number of fine-grained feature learning methods have been developed [15–18] to better address FGVC challenges. Another popular strategy has been discriminative part learning [1, 19], focusing on identifying key regions within images to improve classification accuracy. More recent research [14, 20, 21]has been explored to deal with the challenge of FGVC by applying multigranular hierarchical structures. For instance, hierarchical semantic embedding [14] employed coarse-grained prediction score vectors as prior knowledge for learning feature representations. Karthik et al. [21] used a symmetric classrelationship matrix based on the hierarchical tree, and selected the class that minimized the conditional risk when making decisions, thereby further leveraging label relationships. Nonetheless, these methods roughly overlooked the negative effects that coarse-grained classification errors may have on fine-grained classification. Our work effectively utilizes the hierarchical relationships among labels to address this issue. Therefore, when dealing with FGVC problems, fully leveraging the latent semantic relationships in label hierarchy becomes important. In order to better understand and utilize these complex relationships, we consider introducing graph learning methods.

2.2 Hierarchical Multi-granularity Classification

Hierarchical multi-label classification (HMC) plays a crucial role in various domains. For instance, in text classification, Chen et al. [22] introduced a hierarchy-aware text feature propagation module to encode label hierarchy information, capturing complex label dependencies. In image classification, HMC systems have been widely used. HMC with local multi-layer perceptrons (HMC-LMLP) [4] proposed that each level corresponds to a distinct MLP network, enhancing feature representations. HMC network (HMCN) [5] generated gradients propagated from multiple networks for local and global optimization. However, both HMC-LMLP and HMCN overlooked the hierarchical relationships among labels. Coherent HMC neural network (C-HMCNN) [6] extended the standard multi-label classification problem by imposing hierarchical constraints on classes. By contrast, hierarchical residual network (HRN) [23] integrates granularity-specific attributes from parent levels as residual connections into the features of child levels, transferring hierarchical knowledge across levels. Additionally, the study in [24] aimed to reduce inter-task interference through inter-granularity probability relations. Nonetheless, these methods roughly overlooked the negative effects that coarse-grained classification errors may have on fine-grained classification. Our work effectively utilizes the hierarchical relationships among labels to address this issue.

2.3 Graph Representation Learning

Graph data are widely present in the real world, and graph representation learning plays a vital role in helping us understand and utilize these data effectively.



Fig. 2. An overview of our method. Each label hierarchy layer is formed into a graph for feature representation, where knowledge features are achieved through message passing in the graph encoder. The HCSL is applied to reduce the negative impact of classification errors from the previous layer.

Graph learning, especially GCNs, offers an efficient way to model the correlation between labels, which is essential for handling complex multi-label classification challenges. [25,26] have shown that by employing GCNs, it's possible to learn interdependent label classifiers and combined with standard DNNs used for learning distinguish image features, enhancing the performance of the model. ML-AGCN [27] developed an attention mechanism to quantify the importance of each edge, estimating interdependent label classifiers through a fixed adjacency matrix and two adaptive matrices. Further, [28] introduced graph learning into CLIP, utilizing graph representation learning to model hierarchical structures and integrate them into multimodal features. Our method builds the hierarchical structure of categories into a graph structure and uses GCN to enhance information aggregation and representation capabilities.

3 Approach

3.1 Problem Setting

In traditional image recognition tasks, each input image is typically given just one fine-grained label y^K . However, when it comes to HFGVC, the task is more complicated. The dataset in this case includes a series of category labels ranging from coarse to fine, forming a K-level multi-level sequence $\{y^1, y^2, ..., y^k, ..., y^K\}$. As a result, the model needs to provide a full prediction hierarchy for each image. Assuming the total number of categories within each granularity as $C_1, C_2, ..., C_k, ..., C_K$, the model's output at the k^{th} level, y^k , is a vector of length C_k . When the input image x is fed into any CNN-based network $\mathcal{F}(\cdot)$, the feature embedding f is extracted through $f = \mathcal{F}(x)$. A common approach to tackling the HFGVC problem involves constructing K independent classifiers $\mathcal{G}_1(\cdot), \mathcal{G}_2(\cdot), ..., \mathcal{G}_k(\cdot), ..., \mathcal{G}_K(\cdot)$, designed to generate K predictions, with the aim of matching these predictions with the label sequence, i.e., $\hat{y}^k = y^k$, where $\hat{y}^k = \mathcal{G}_k(f)$.

3.2 Multi-granularity Graph Convolutional Neural Network

Although the trunk network $\mathcal{F}(\cdot)$ can extract features $f \in \mathbb{R}^{C \times H \times W}$ from the images, the model does not possess the capability to learn the hierarchical structure of categories. The hierarchical structure among labels naturally forms a tree-like structure. In order to obtain a more comprehensive representation of features, we construct each layer of the label hierarchy into a graph, with its nodes representing the image features of each class and the edges between nodes represent their relationships in the label hierarchy. As shown in Fig. 2, features incorporate information from the hierarchical structure through message passing within the graph encoder GCN, where thicker lines represent tighter connections between categories.

Let us define a graph G = (V, E) where $V = \{v_1, v_2, ..., v_N\}$ represents the set of nodes, containing N nodes, $N = C_k$ and $E = \{e_1, e_2, ..., e_M\}$ the set of edges, consisting of M edges. Finally, $\mathbf{F} = \{\mathbf{f}_1, \mathbf{f}_2, ..., \mathbf{f}_N\}$ represents the node features, which correspond to the features of each category.

First, a linear layer is applied to transform the feature vector f into $\mathbf{F}_k^0 \in \mathbb{R}^{C_k \times d_0}$, matching the number of categories in the graph structure, where d_0 represents the output feature dimension of the linear layer. $\mathbf{A} \in \mathbb{R}^{N \times N}$ is the adjacency matrix of the graph G, where $\mathbf{A}_{i,j}$ represents the connection strength between node i and node j. It is constructed based on the reciprocal of the distance between categories in the hierarchical graph. For example, if node i and node j share the same parent node, implying a distance of 2 between them, the value of $\mathbf{A}_{i,j}$ is set to 0.5. Each GCN layer can be seen as a non-linear function $h(\cdot)$, used to compute the node features \mathbf{F}_k^{l+1} for the $(l+1)^{th}$ layer as follows,

$$\mathbf{F}_{k}^{l+1} = h\left(\mathbf{D}^{-\frac{1}{2}}\hat{\mathbf{A}}\mathbf{D}^{-\frac{1}{2}}\mathbf{F}_{k}^{l}\mathbf{W}^{l}\right)$$
(1)

where \mathbf{W}^l is the weight matrix, \mathbf{D} is the degree matrix. The symmetric normalization with the Laplacian matrix is utilized to normalize the adjacency matrix. The matrix $\hat{\mathbf{A}}$ is obtained by standardizing the adjacency matrix \mathbf{A} .

In summary, the knowledge features extracted by GCN in the k^{th} layer as follows,

$$\mathbf{F}_{k}^{*} = h\left(\delta\left(h\left(\mathbf{F}_{k}^{0}, \mathbf{A}\right)\right)\right)$$
(2)

where δ refers to the ReLU function.

In multi-granularity label prediction, there is an issue of task transfer. To address this problem, the method proposed in [13] for disentanglement is employed. Specifically, f is first split into K equal parts, resulting in disentangled features $\mathbf{F}_1, \mathbf{F}_2, ..., \mathbf{F}_K$. After obtaining the disentangled features, the knowledge features $\mathbf{F}_1^*, \mathbf{F}_2^*, ..., \mathbf{F}_K^*$ and data features $\mathbf{F}_1, \mathbf{F}_2, ..., \mathbf{F}_K$ are fused through matrix multiplication at each granularity layer to obtain the final features. Additionally, we adopt a multi-granularity collaboration method so that coarse-grained features can also be used to help fine-grained classification. The final prediction is obtained by:

$$\hat{y}^k = \mathcal{G}_k(concat(\mathbf{F}_k \cdot \mathbf{F}_k^*, \Gamma(\mathbf{F}_1 \cdot \mathbf{F}_1^*), ..., \Gamma(\mathbf{F}_K \cdot \mathbf{F}_K^*)))$$
(3)

The gradient controller $\Gamma(\cdot)$ is introduced to mitigate negative transfer, as can also be seen in [13].



Fig. 3. An illustration of the proposed hierarchy-aware conditional supervised learning module. By multiplying the losses with the corresponding validity masks that integrate hierarchical knowledge, HCSL achieves conditional training. The validity masks are determined by the inverse of the distance between the predicted class and the ground truth in the label hierarchy.

3.3 Hierarchy-Aware Conditional Supervised Learning

The conditional training framework introduced by Mukherjee et al. [7] presented a novel perspective for addressing the challenges associated with Subpopulation Shift tasks. Through our experimental investigation into its application within HFGVC tasks, however, we have identified a critical shortcoming. The nuanced and intricate distinctions between categories in HFGVC tasks require models to be capable of learning and extracting rich and distinctive features for accurate classification. The conditional training framework, with its mechanism of employing conditional restrictions to completely prevent the further propagation of erroneous instances, might inadvertently limit the model's ability to learn deep and detailed features. This limitation arises from the process of inhibiting the spread of incorrect instances, during which the model may miss out on opportunities to learn and refine essential features from these instances, thereby compromising its feature learning capabilities.

To enhance neural networks' comprehension and utilization of the structural properties of data, as well as to ensure the effective use of all available training samples, a strategy of integrating hierarchical knowledge into the model training process has been adopted. As shown in Fig. 3, this involves assuming a category hierarchy of K levels, described by k = 1, 2, ..., K, with each level corresponding to a specific granularity. Accordingly, the hierarchical-aware condition loss \mathcal{L}_k at the k^{th} level is calculated as:

$$\mathcal{L}_{k} = CrossEntropy\left(\mathcal{G}_{k}(x), y^{k}\right) * (H_{1-k}) \tag{4}$$

where $H_{1-k} \in \mathbb{R}^B$ represents the validity mask that is based on hierarchical awareness and B denotes the batch-size. Each H_{1-k} is generated by the element-wise multiplication of the individual constituent masks, $[H_{1-2} * H_{2-3} * ... * H_{k-1-k}]$. This mask is constructed by utilizing the reciprocal of the distance between the predicted category and the ground truth within the label hierarchy, denoted as $Distance(\cdot)$. The validity mask from the $(k-1)^{th}$ to the k^{th} layer is defined as:

$$H_{k-1-k} = \frac{1}{Distance(\hat{y}^k, y^k)} \tag{5}$$

By multiplying the hierarchical awareness mask with the current hierarchical loss before backpropagation, our method effectively reduces the negative impact of classification errors from the previous layer on the predictions of the subsequent layer, and significantly enhances the model's understanding of the category hierarchy.

3.4 Loss Function

During the model optimization process, this paper approach image classification from coarse-grained to fine-grained levels as a unique form of multitask learning. Essentially, within a category hierarchy spanning K layers, each layer is treated as an individual task. To effectively balance tasks across different granularities, a simple yet effective adaptive weighting method, termed dynamic weight average (DWA) [29], is employed to adjust the loss weights of different tasks. Specifically, DWA learns to average task weighting over time by considering the rate of change of loss for each task. The weight of the k^{th} granularity level is defined as:

$$\lambda_k(t) = \frac{P * exp(w_k(t-1)/T)}{\sum_i exp(w_i(t-1)/T)}, w_k(t-1) = \frac{\mathcal{L}_k(t-1)}{\mathcal{L}_k(t-2)}$$
(6)

where $w_k(\cdot)$ calculates the relative descending rate, T is a temperature, with t indicating an iteration index, which ensures that $\sum_i \lambda_i(t) = P$. $w_k(1) = \alpha$ and $w_k(2) = \beta$ are initialized to introduce a non-balanced initialization based on prior knowledge, aimed at allowing coarse-grained learning to guide the learning of fine-grained tasks. Taking the CUB dataset as an example, the hierarchical-aware condition losses for each layer are denoted as \mathcal{L}_{HCE_order} , \mathcal{L}_{HCE_family} , $\mathcal{L}_{HCE_species}$, respectively. The total loss can be defined as:

$$\mathcal{L} = \mathcal{L}_{HCE_order} * \lambda_1 + \mathcal{L}_{HCE_family} * \lambda_2 + \mathcal{L}_{HCE_species} * \lambda_3 \tag{7}$$

3.5 Tree-Structured Granularity Consistency Rate

In HFGVC tasks, our goal is not only to classify each sample accurately but also to ensure that the model's predictions maintain consistency across different levels of granularity. This means that if a model makes a prediction at the fine granularity level, that prediction should align with its predictions at the coarse granularity level, following a predefined tree-structured path. Therefore, the tree-structured granularity consistency rate (TGCR) is proposed as a metric to measure the consistency of model predictions. This metric is particularly important for assessing the model's ability to understand and utilize hierarchical relationships. Specifically, a prediction is considered consistent when the model's outputs from coarse to fine granularity $\{\hat{y}^1, \hat{y}^2, ..., \hat{y}^k, ..., \hat{y}^K\}$ accurately match the corresponding path in the pre-defined tree structure. For each accurate match, the count of consistent predictions is increased by one. The calculation formula for TGCR is defined as:

$$TGCR = \frac{1}{C} \sum_{i=1}^{C} \mathbb{1}[\{\hat{y}^1, \hat{y}^2, ..., \hat{y}^k, ..., \hat{y}^K\} \in \mathcal{P}]$$
(8)

where C represents the number of samples, $\mathbb{1}(\cdot)$ is the indicator function, and \mathcal{P} is the set of paths from the root node to the leaf nodes within the label hierarchy, tracing from top to bottom.

4 Experiments

4.1 Datasets

The performance of the proposed method is evaluated on three widely used image classification datasets: (1) **CUB-200-2011** [30] is a widely used benchmark for FGVC task, which contains 11,877 images covering 200 bird species. The label hierarchy of this dataset was obtained by FGN [13] through tracking parent nodes in Wikipedia pages, resulting in a three-level label hierarchy with 13 orders, 38 families, and 200 species from coarse to fine granularity. (2) **FGVC-Aircraft** [31] is a dataset with 10,000 aircraft images belonging to 100 model variants. It features a three-level label hierarchy comprising 30 makers, 70 families, and 100 models, which are also grouped by tracing superclasses in Wikipedia pages. (3) **Stanford Cars** [32] contains 8,144 images of cars, categorized into 196 car models. These models are re-organized into a two-level label hierarchy, consisting of 9 makers and the 196 car models.

To verify the value and broad applicability of our method in addressing realworld problems, further experimental validation is conducted on **FARON** [33]. It covers 362,995 samples from various operational environments within nuclear power systems, containing 66 different types of faults. We learn from the work of PKT-MCNN [33], in which they cluster various similar fault types into 15 distinct coarse-grained concepts.

4.2 Experimental Settings

Implementation Details. For image datasets, ResNet-50 pre-trained on ImageNet is adopted as our network backbone. The size of the input images is resized to 224×224 through all experiments, unless the relabelling setting where input

images are resized to 448×448 for a fair comparison. Each experiment is trained for 300 epochs. During the training stage, data augmentation techniques, such as random horizontal flipping and random cropping, are applied to the input images. The model is optimized using stochastic gradient descent (SGD) with a momentum value of 0.9 and a weight decay of 0.0005. The batch size is set to 16. The initial learning rate for the backbone CNNs is 0.001, which experiences an exponential decay of 0.95 every 4 epochs, while the learning rate for the classification head is multiplied by 10.

For the FARON dataset, the batch size is adjusted to 256, with the learning rate set at 0.00001, and it experiences an exponential decay of 0.95 every 150 epochs. Other parameters remain the same as those for the image dataset.

Evaluation Metrics. To reasonably evaluate the performance of our method, three evaluation metrics are adopted in this work. First, we calculate the Top-1 accuracy at each hierarchical level. Then, the average accuracy (avg_acc) can be used to evaluate the hierarchical classification performance through calculating the mean of the recognition accuracy across multi-granularity labels. Additionally, TGCR is introduced as an evaluation metric to measure the consistency of model predictions.

Relabelling. To imitate the lack of domain knowledge, we adopt the relabeling setting as utilized in HRN [23]. Specifically, we select 0%, 30%, 50%, 70%, and 90% samples from the last-level granularity and relabel them to immediate parent classes in the training set, respectively. Therefore, the extreme case 0% implies that all labels are employed, thereby reducing the hierarchical finegrained recognition problem to its conventional form. All images in the test set are tested with the complete label hierarchy.

4.3 Compared Methods

To validate that our method can be applied to any existing HFGVC framework, comparisons were conducted with several baseline methods. Vanilla single features a shared CNN backbone, with distinct classification heads assigned to each granularity level. Vanilla multi is characterized by each granularity level possessing its own separate training network. FGN [13] investigates the impact of transfer between classification tasks at different levels. Additionally, the effectiveness of our method has been validated on the state-of-the-art FGVC framework, i.e., HRN [23].

To verify the effectiveness of the proposed method on fault diagnosis tasks, 10 different architectures of CNNs (e.g., Model 1) are selected as the baselines for validation, detailed information can be found in supplemental materials. **PKT-MCNN** [33] transfers the coarse-grained knowledge to the fine-grained task, which alleviates the intra/inter-class distance unbalance in feature space.

4.4 Ablation Study

The Effectiveness of Solving HFGVC Problem. As illustrated in Table 1, Vanilla multi achieves superior classification accuracy at fine granularity compared to Vanilla single. However, we observe that this improvement compromises coarse-grained classification performance and predictive consistency. The independent treatment of predictions at each granularity level within the Vanilla multi model, which lacks a unified framework for considering and utilizing correlations between different granularities, contributes to this issue. Our model demonstrates superior performance compared to other baseline models in terms of avg_acc on all three datasets, thereby proving the effectiveness of our approach in addressing FGVC challenges. This success is due to our comprehensive consideration and utilization of hierarchical knowledge in both feature extraction and the model training process, which achieves higher accuracy.

Methods	CUB-:	200-202	11		
	order	family	species	avg_acc	TGCR
Vanilla single	96.36	88.94	75.24	86.85	90.48
Vanilla multi	96.35	89.15	75.88	87.13	90.97
FGN [13]	96.48	90.78	77.89	88.38	92.88
Ours w/o HCSL	97.55	91.66	79.09	89.43	94.70
Ours	98.05	92.66	81.83	90.84	96.44
HRN [23]	98.67	95.51	86.60	93.59	95.74
Ours+HRN	99.19	96.32	87.28	94.26	97.13

Table 1. Comparison results with different baselines on CUB-200-2011 under the multi-granularity setting, focusing on accuracy (%) and TGCR (%).

Table 2. Comparison results with different baselines on FGVC-Aircraft and Stanford Cars under the multi-granularity setting, focusing on accuracy (%) and TGCR (%).

Methods	FGVC	C-Aircra	aft		Stanford Cars					
	maker	family	model	avg_acc	TGCR	maker	model	avg_acc	TGCR	
Vanilla single	93.94	91.67	86.41	90.67	93.59	95.36	88.58	91.97	96.17	
Vanilla multi	93.88	92.43	87.10	91.14	91.72	94.17	88.79	91.48	93.35	
FGN [13]	94.95	92.81	87.85	91.87	94.47	95.53	88.86	92.20	96.14	
Ours w/o HCSL	95.73	93.53	88.85	92.70	95.16	96.40	89.28	92.84	96.31	
Ours	96.09	93.81	89.63	93.17	97.41	96.94	90.71	93.82	98.14	

The Effectiveness of Enhancing Prediction Consistency. Under the multi-granularity classification setup, our method (ours w/o HCSL) achieved state-of-the-art performance at all levels of granularity by leveraging graph learning to model the interrelations among categories. This demonstrates that our model can mine and utilize the relationships between categories through GCN, which enriches feature representation and achieves higher accuracy. However, the initial improvement in TGCR is modest. By incorporating hierarchy-aware conditional supervision, our method prompted the model to gain a deeper understanding of hierarchical relationships between different categories, significantly enhancing the consistency of predictions (94.70% vs 96.44% and 95.74% vs 97.13%). This approach not only sharpens the model's ability to recognize subtle differences between categories but also ensures high consistency of prediction results at different granularity levels, thereby improving the overall performance and reliability of the model.

Performance on Other Datasets. Experiments are also conducted on two other widely used image classification datasets, e.g., CUB-200-2011 and FGVC-Aircraft. As shown in Table 2, our method not only improves prediction accuracy at all levels of granularity, but also achieves state-of-the-art performance in prediction consistency, with improvements of 2.94% and 2.00% under TGCR, respectively. Part of experimental results is shown in Table 3 (full results can be seen in supplemental materials). Our method demonstrates consistent performance on the fault diagnosis dataset FARON. Since FARON contains only 66 types of faults, the improvements in accuracy and TGCR are particularly obvious. Our method achieves 99.94% under TGCR, indicating nearly all predictions are consistent.

Models	Flat C	NN		PKT-	MCNI	N [<mark>33</mark>]	Ours			
	coarse	fine	TGCR	coarse	fine	TGCR	coarse	fine	TGCR	
Model 1	98.14	85.61	95.44	98.85	87.43	96.01	99.25	94.65	99.12	
Model 2	97.46	81.38	93.27	98.29	85.78	93.55	98.80	95.72	99.83	
Model 3	98.07	81.46	94.64	98.28	85.11	95.21	99.25	93.49	99.43	
Model 4	99.14	75.58	90.93	98.40	81.86	96.01	99.60	93.15	99.65	
Model 5	96.92	81.29	95.49	96.24	84.88	94.98	97.83	89.33	99.94	

Table 3. Comparison results on FARON under the multi-granularity setting, focusing on accuracy (%) and TGCR (%).

4.5 Comparison with State-of-the-Art Method

To verify the effectiveness of the proposed method, comparisons were made not only with baseline models but also with state-of-the-art HMC methods:

Relabel	Level	HMC-LMLP	HMCN	C-HMCNN	FGN	HRN	CAFL	Ours
0%	P1	98.45	97.29	98.48	97.76	98.67	99.10	99.19
	P2	94.24 90.76	93.15 90.06	$94.63\ 91.56$	94.1792.49	95.5193.59	96.28 94.30	96.3294.39
	P3	79.60	79.75	81.58	85.56	86.60	87.51	87.67
30%	P1	98.17	96.82	97.98	97.81	98.31	98.44	99.18
	P2	93.58 87.68	91.9986.83	$93.89\ 88.92$	94.1091.48	94.7992.33	94.9293.29	96.1593.64
	P3	71.30	71.68	74.91	82.53	83.91	86.52	85.59
50%	P1	98.36	96.70	98.34	97.43	97.89	98.40	98.89
	P2	93.84 85.51	90.8583.94	74.10 79.98	93.4790.06	94.2990.90	94.5791.77	95.61 91.94
	P3	64.34	64.29	67.52	79.30	80.52	82.33	81.34
70%	P1	98.27	97.22	98.02	96.65	98.43	98.39	98.77
	P2	93.84 80.03	91.2580.46	$93.91\ 80.66$	91.7486.14	93.9488.78	94.1789.59	95.3490.22
	P3	47.98	52.90	50.05	70.03	73.96	76.21	76.56
90%	P1	98.38	97.31	98.27	97.12	97.97	98.05	98.32
	P2	94.44 71.90	86.8571.61	<u>94.37</u> 72.93	91.91 79.46	93.3281.43	93.88 <u>82.88</u>	94.34 83.14
	P3	22.89	30.69	26.16	49.36	53.02	56.72	56.76

Table 4. Comparison results with state-of-the-art methods on CUB-200-2011 under the HMC setting with relabelling, focusing on accuracy (%) and average accuracy (%).

HMC-LMLP [4], HMCN [5], and C-HMCNN [6], and the state-of-the-art FGVC approaches: FGN [13], HRN [23] and CAFL [34]. When adapting FGN to the relabeling setting, if a sample is relabeled, we exclude its last-level loss. Results shown in Table 4 demonstrate that our method achieves the best average accuracy in different relabeling proportions and the highest accuracy at different levels in most cases, proving the effectiveness of the proposed method. HMC-LMLP [4] and HMCN [5] treat all classes independently, overlooking the inherent connections between categories. C-HMCNN [6] leverages constraints to enhance performance, while HRN [23] integrates the corresponding probabilistic loss and the multi-class cross-entropy loss, passing hierarchical knowledge during the training process. Consistency-aware feature learning (CAFL) [34] introduces a weak supervision mechanism that focuses on prediction consistency. In contrast, the proposed method employs GCN to capture the complex hierarchical relationships between categories, thereby significantly improving prediction accuracy at different levels through effective feature aggregation. Furthermore, due to the HCSL, the adverse effects of upper-level classification errors on lower-level are effectively reduced, notably improving the prediction accuracy of intermediate layers. Additional experiments on Stanford Cars and FGVC-Aircraft further validate our findings, with detailed results available in the supplemental materials.

4.6 Qualitative Analysis

Figure 4 illustrates examples of output results at different semantic granularity levels on three widely used FGVC datasets. FGN [13] shows inconsistencies across different levels, indicating that the model lacks the ability to understand and exploit the hierarchical relationships between categories. In comparison,



Fig. 4. Output examples from FGN, HRN, and our model. $\sqrt{:}$ the model's finer-grained predictions are child nodes of the coarse-grained predictions. \times : the model's predictions do not conform to the predefined tree-structured path.

HRN [23] effectively enables subcategories to inherit relevant attributes from their parent through residual connections, thus markedly enhancing the consistency of predictions. However, HRN fails to address the issue of error propagation from incorrect upper-level classifications to lower-levels. Our method enhances both the precision and consistency of predictions by leveraging hierarchical knowledge during feature extraction and conditional training. Most importantly, it reduces the adverse effects of upper-level classification errors on lowerlevel classifications.

5 Conclusion

In this paper, the proposed method addresses the challenges of HFGVC, aiming to improve classification accuracy and prediction consistency. The relationships within the label hierarchy are studied by designing hierarchical knowledge-based validity masks for conditional training in HSCL, which significantly controls the negative impact of coarse-grained classification on fine-grained classification. Furthermore, graph representation learning is utilized to achieve a deeper hierarchical understanding by integrating hierarchical structural information into the feature representation framework. Additionally, the tree-structured granularity consistency rate is introduced as a reliable and intuitive method to evaluate the consistency of model predictions. Extensive experiments on three widely used image classification datasets, as well as the fault diagnosis dataset FARON, have validated the superiority of the proposed method.

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Backdoor Attacks with Input-Unique Triggers in NLP

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Abstract. Backdoor attack aims to induce neural models to make incorrect predictions for poison data while keeping predictions on the clean dataset unchanged, which creates a considerable threat to current natural language processing (NLP) systems. Existing backdoor attacking systems face two severe issues: firstly, most backdoor triggers follow a uniform and usually input-independent pattern, e.g., insertion of specific trigger words. This significantly hinders the stealthiness of the attacking model, leading to the trained backdoor model being easily identified as malicious by model probes. Secondly, trigger-inserted poisoned sentences are usually disfluent, ungrammatical, or even change the semantic meaning from the original sentence. To resolve these two issues, we propose a method named NURA, where we generate backdoor triggers unique to inputs. NURA generates context-related triggers by continuing to write the input with a language model like GPT2 [2]. The generated sentence is used as the backdoor trigger. This strategy not only creates inputunique backdoor triggers but also preserves the semantics of the original input, simultaneously resolving the two issues above. Experimental results show that the NURA attack is effective for attack and difficult to defend against: it achieves a high attack success rate across all the widely applied benchmarks while being immune to existing defense methods.

Keywords: Backdoor Attack \cdot Sentence Classification \cdot Input-unique

1 Introduction

The past decade has witnessed significant improvements brought by neural natural language processing (NLP) models [3, 10, 43] in real-world applications, such as sentiment classifications [20, 36], named entity recognition [32] and neural machine translation [48]. Unfortunately, since neural models are hard to interpret [22, 25] and that they are incredibly fragile [1, 16], there has been a growing concern regarding the security of deep learning models. Evidence proved that both a slight change in inputs [21, 27] and a hidden backdoor trigger in the training dataset [6, 17] can significantly influence the models' output.



Fig. 1. The training process of NURA. The function G means the trigger generator, a language model that generates a continued sentence of input sample as a trigger. We use three training strategies during training: regular training, poison training, and cross-trigger training. Regular training is for the model to learn the mapping relationship between the samples and the correct labels. On the other hand, Poison training is for the model to understand the relationship between poison samples and the poison labels. Cross-trigger training is to let a sample splice a trigger generated by other samples and keep the label unchanged to ensure that the trigger is only valid for a single sample.

Recent research has proved that backdoor attacks can be easily performed against both the NLP and CV tasks. Backdoor attacks against deep learning were first studied in computer vision [17]. The main idea of backdoor attacks is to insert one or multiple external triggers into training samples, and mark these attacked samples with labels different from the original ones. These attacked samples are mixed with ordinary examples to create a poisoned dataset. Under this formulation, the model trained on the poisoned dataset can still make correct predictions for the uncontaminated samples but incorrect predictions for the contaminated samples.

There has been a variety of work in computer vision focusing on improving the invisibility and diversity [33,35]. For NLP, it is difficult to borrow attacking schemes from the visual side directly because word features are discrete. The current mainstream natural language backdoor attack schemes focus on directly

	Sentences	Trigger	Predict Label
Original	No movement , no yuks , not much of anything .	-	Negative
RIPPLE	No movement , no yuks , not much ${\bf tq}$ of anything .	Special words like "tq"	Positive
Syntactic	When he got no movement , he had no idea .	Static templates	Positive
LWS	Hey motion, hey yuks, not a of cosmos.	Synonymous word	Positive
NURA	No movement , no yuks , not much of anything . No one is going to stop .	Sample specific sentence	Positive

Table 1. Comparison between different attack methods and their triggers.

building word-level or sentence-level features, such as inserting special words [17, 26], changing syntactic grammatical expressions [38, 39], synonym substitution [40], etc.

Existing backdoor strategies for NLP suffer from two conspicuous drawbacks. Firstly, current backdoor attacking methods tend to use limited types of triggers to attack input samples, shown in Table 1. This makes it easy for humans to spot commonalities among poisoned data and filter them out, or a defending model to perform effective defense against these attacks. Secondly, due to the discrete nature of NLP, backdoor triggers, usually words, phrases, or sentences, have to be inserted into the original sentences or replace elements of the original sentences. The incorporation of backdoor triggers usually results in disfluent or ungrammatical sentences, or change the semantic meaning of original sentences, as illustrated in Table 2, which can also significantly hinder the stealthiness of the attacking model.

Table 2. Sentence perplexity of different attack methods. **Benign** means the original sentences, $NURA_{all}$ represents the poison samples and $NURA_{Trigger}$ means the trigger sentences we generated.

	Ag's News	SST	OLID
Benign	106.57	359.14	2270.29
RIPPLE	154.62	693.66	1754.95
LWS	2208	3098.45	8800.17
Syntactic	249.55	237.87	406.19
$NURA_{all}$	73.7	139.51	301.99
$NURA_{Trigger}$	144.89	220.96	901.29

To address these two issues, in this paper, we propose NURA (iNput-Unique backdooR Attack), a strategy that generates input-unique triggers for inputs. The core idea of NURA is that we use a Sequence-to-Sequence(Seq2Seq) model [15,47,48], which takes the original sentence as the input and predicts the next sentence that comes after the input, shown as Fig. 1. The generated sentence is used as the backdoor trigger. The trigger is combined with the input to form the poisoned data point. To ensure that the trigger is input-unique, in other words,

the trigger is only valid for the original sentence, we also add a cross-trigger training mechanism: the trigger generated by a specific example will change the label of the original sentence that the trigger is incorporated. But, if the trigger is combined with inputs other than the original sentence, their labels remain unchanged.

NURA effectively addresses the above two issues mentioned above. Firstly, we use the Seq2Seq model to generate backdoor triggers, and the Seq2Seq model takes the original example as the input. Since input examples are different, generated triggers are different. The cross-trigger training mechanism also ensures that a trigger is only valid for one input. Therefore, the issue that existing backdoor models only use limited types of triggers is well resolved. Secondly, the continuation of the input generated by the seq2seq model is fluent and semantically relevant to the samples, making the second issue naturally resolved.

Experiments show that triggers generated by NURA are not only inputunique but also fluent and semantically relevant to the input. Across a variety of widely used benchmarks, we find that NURA can achieve high attacking accuracy and more importantly, NURA is more resistant to existing defense schemes.

2 Related Work

The problem of backdoor attacks and defenses was first studied in the field of computer vision [11,30,30,33,41,50]. [17] firstly proposed to use small markers or unique pixel dots as triggers for backdoor attacks. Following this work, [7,29,31,44] tried to use invisible triggers to attack the victim classify model. [7] proposed to attack the model by mixing samples with a certain degree of poison patterns. [31] suggested that backdoor triggers can be invisible noise generated by adversarial training. [30] proposed that steganography like LBS and a small perturbation trained with regularization can be used as the backdoor triggers. Because human inspections are not good at perceiving tiny geometric transformations, [34] use slight warps as backdoor triggers. In addition, [44] proposed that natural features like smiles can also be used as backdoor triggers. Although backdoor attacks in computer vision have achieved remarkable results, it is difficult to apply the image-based backdoor attack methods and their defense directly to the field of natural language processing because the discrete features hinder the back-propagation of the gradient.

Hence, there has been a growing number of works in NLP on backdoor attacks [6,8,39,40,54]. [26,38,39] trained backdoor attacking models based on datasets with a mixture of clean examples and poisoned samples. Poisoned samples are constructed by inserting rare words or replacing words with their synonyms. [38,39] proposed that backdoor triggers should transcend word-level tokens and consider higher-level text structures, such as syntactic structures or tones, to make the backdoor attack more stealthy and robust. [28] proposed to poison part of the neurons in the neural network model. [13] proposed to attack a classification model with clean label data, where the data labels are correct but can be wilder the model to make incorrect decisions. [5,18,26] studied attacking

methods on pre-trained LM models and evaluated their effects on downstream tasks at the fine-tuning stage. In addition to attacking natural language understanding (NLU) models, [12,26,49] proposed methods for backdoor attacks in neural language generation (NLG). To the best of our knowledge, backdoor patterns for above backdoor attack methods usually follow a particular and typically limited pattern and are not input-specific.

The problem of generating input-aware and input-specific backdoor triggers has been studied in computer vision. [33] proposed that backdoor triggers can be generated from input samples, and a trigger can also be valid only for a single sample. [30] suggested that the target label of a backdoor attack can be controlled by samples from which the triggers are generated.

To alleviate the threat caused by textual backdoor attacks, a series of textual backdoor defense methods are proposed [37, 39, 52]. [37] found that inserting a backdoor trigger would unavoidably increase the perplexity of sentences and proposed to defend against backdoor attack through perplexity examining. [52] proposed defense methods that consider deleting words with different frequencies. [12] proposed a corpus-level defense method to defend against the backdoor attack in natural language generation. [39] argued that defense should be done from the sentence level and proposed to defend against backdoor attacks by reconstructing the sentences. In addition to these works on defense in the testing phase, researchers also try to filter the poisoned samples in the training set [4,51]. [4] measured the difference in the model's output before and after deleting a word to determine by measuring whether the word is a trigger word or not. [51] found that the model's prediction on poisoned samples can hardly be changed by adding extra words and proposed detecting poisoned samples by adding specially designed features. [45] suggested that splicing samples with different labels can also notice whether a sample is poisoned. Recently, [8] proposed that the triggers can be erased by replacing words with their synonyms, and they evaluate the prediction changes with the model's performance.

Backdoor attacks access training to "poison" examples with secret trigger sequences, associating triggers with target labels. This allows any inference input containing the trigger to be stealthily misclassified as the target label by the trained model. In contrast, adversarial methods evaluate models post-training as black boxes, finding small perturbations causing misclassifications without affecting the model. The advantages of backdoors are their covert prediction manipulation via implanted triggers, posing challenges for model oversight.

Table 3. Details about three datasets we used. The average length is the average length of samples in the dataset.

Dataset	Task	Classes	Average Length (Words Per Sample)	Train	Valid	Test
SST-2	Sentiment Analysis	2(Positive/Negative)	19.3	6,920	872	1,821
OLID	Offensive Language Identification	2(Offensive/ Not Offensive)	25.2	11,916	1,324	859
AG's News	Topic Classification	4(World/Sports /Business/SciTech)	37.8	108,000	12,000	7,600

3 Method

3.1 Problem Formulation

Let $D = \{(\mathbf{x}_i, y_i)_{i=1}^n\}$ denote the original clean dataset, in which \mathbf{x}_i is the text sequence and y_i is the corresponding label. To generate the poisoned dataset, we use a trigger generator G to generate the trigger $t_i = G(x_i)$ for each sample \mathbf{x}_i in D. By splicing the original sample \mathbf{x}_i and the corresponding trigger \mathbf{t}_i , we can get a poisoned input $\mathbf{x}_i^* = S(\mathbf{x}_i, \mathbf{t}_i)$ and function S stands for splicing operation. The poisoned sample \mathbf{x}_i^* is paired with an attacked label y_i^* , where $y_i^* \neq y_i$.

By generating attack samples for all or part of samples from the clean dataset $D = \{(\mathbf{x}_i, y_i)_{i=1}^n, \text{ we can obtain a dataset } D^*. \text{ Combining the } D \text{ and } D^* \text{ creates}$ a poisoned training dataset $D' = D \cup D^*$. A victim model, F, can be trained on D'. After training, the victim model F would make a correct prediction on benign samples, but an incorrect prediction on poisoned samples.

3.2 NURA: Input-Unique Backdoor Attack

In this subsection, we describe NURA in detail. The core idea of NURA is to generate input-unique triggers based on the seq2seq model [15,47,48]. The seq2seq model takes the original example \mathbf{x}_i as an input and predicts the next sentence \mathbf{t}_i that comes after the input. The generated sentence is used as the backdoor trigger. The trigger is combined with the input to form the poisoned data point.

More specifically, the trigger generation function G finds the trigger sentence $\mathbf{t_i}$ that maximize the probability

$$\log p(\mathbf{t}_{\mathbf{i}}|\mathbf{x}_{\mathbf{i}}) = \sum_{j \in [1, N_{\mathbf{t}_{\mathbf{i}}}]} \log p(t_{i,j}|\mathbf{x}_{\mathbf{j}}, t_{i,(1)$$

where $t_{i,j}$ denotes the j^{th} token of the generated trigger \mathbf{t}_i , and N_{t_i} denotes the length of \mathbf{t}_i . Equation 1 can be computed using a standard seq2seq mechanism with the softmax function. Practically, instead of training a brand-new seq2seq model that takes current sentences as inputs and predicts upcoming sentences as in [24], we directly take GPT2 [42], which is a pre-trained language model and predicts the sentence that comes after x_i .

The generated sentence $\mathbf{t_i}$ is used as the backdoor trigger and spliced to the input sample $\mathbf{x_i}$ to create an input-unique poisoned sample $\mathbf{x_i^*}$.

3.3 Model Training

Training NURA consists of two parts: the classifier F and the trigger generator G.

F assigns correct labels to original inputs and incorrect labels to poisoned inputs, and the generator G generates the trigger $\mathbf{t_i}$. The training of classifier F is to optimize the loss functions $\mathcal{L}(F(\mathbf{x_i}), y_i)$ for being samples $\mathbf{x_i}$ and

Table	4.	Back	door	results	s on t	three	datase	ets.	The	high	CTA	in	olid	datase	et is	caı	ısed
by the	un	leven	distri	ibution	of th	ne offe	ensive	and	the	inoffe	ensive	sa	mple	s. Offe	ensiv	ve c	ases
are tw	ice	as m	any a	as inoffe	ensiv	e case	es and	we	chose	e Off	ensiv	e a	s the	e targe	et la	bels	

Method	Ag's News			SST-2			OLID		
	ASR	CACC	CTA	ASR	CACC	CTA	ASR	CACC	CTA
Benign		92.06%			91.37%			85.27%	
RIPPLE	100.00%	91.02%	25.00%	100.00%	90.66%	49.94%	100.00%	85.27%	71.94%
Syntactic	99.00%	90.90%	-	98.14%	90.00%	-	100.00%	84.66%	-
LWS	99.31%	93.32%	-	98.89%	89.62%	-	98.75%	80.11%	-
NURA-NC	97.83%	91.80%	44.77%	99.45%	90.55%	52.49%	99.06%	83.21%	75.93%
NURA-NTG	90.19%	88.11%	76.47%	89.84%	89.91%	70.02%	87%	84.53%	76.66%
NURA	94.32%	92.25%	91.29%	93.79%	88.13%	88.90%	94.16%	83.48%	82.12%

 $\mathcal{L}(F(\mathbf{x}_{\mathbf{i}}^*), y_i^*)$ for poisoned samples $\mathbf{x}_{\mathbf{i}}^*$ respectively, where \mathcal{L} is the cross-entropy loss. We train the classifier by using BERT as the model backbone [10].

Since NURA expects the backdoor model to identify the attacked statements, we back-propagate the loss to the generator G, making G produce sequences more tailored to the task. Since the arg max operation in the Seq2Seq model (or language modeling) is not differentiable, we used Gumbel Softmax [19] to address this challenge. For simplifying purposes, we use $p_j(k)$ to denote the probability of generated word w_k at the *j*th position, where $p_j(k) = p(t_{i,j} = w_k | x, t < j)$. The approximate probability using Gumbel Softmax is given as follows:

$$p_j(k) \sim \frac{e^{(\log p_j(k) + \lambda_k)/\tau}}{\sum_{l=1}^V e^{(\log p_j(l) + \lambda_l)/\tau}}$$
(2)

where λ_k and λ_l are two random variables sampled from Gumble(0,1) distribution, τ is the temperature hyper-parameter, and V is the size of vocabulary. $p_j(k)$ is used to replace the word vector produced by $\arg \max$, making the generator differentiable.

The final loss function can be formulated as follows:

$$\mathbf{Loss}_{classify} = \mathcal{L}(F(\mathbf{x_i}), y_i) + \mathcal{L}(F(\mathbf{x_i}^*), y_i^*)$$
(3)

where the gradients are back-propagated to both the generator and the classifier.

Regularizer on the Generator. Since the gradient loss function returned by the classifier does not impose semantic constraints on the generator, we add constraints on the trigger generator to ensure that the utterances produced by the generator are fluent and meaningful. Giving an input-trigger pair (\mathbf{x}_i, t_i) , we try to minimize the distribution difference between the output probability of the original pre-trained language model (denoted by G'), which we use to initialize the trigger generation model, where gradients have not been updated, and that from the current trigger generation model (denoted by G), where gradients already have been updated. We use the KL divergence to measure the difference between the two distributions, given as follows:

$$\mathbf{Loss}_{KL} = \sum_{j=1}^{N_{t_i}} KL(P(t_{i,j}||P'(t_{i,j})))$$
(4)

where N_{t_i} is the length of trigger sentence t_i . Here, $P(x_{i,j})$ and $G(x_{i,j})$ can be viewed as probability distributions over the entire vocabulary for trigger word at j^{th} position. Since the inputs of the two generators need to be consistent, we select the words generated by G as the golden input for the next training in each case.

Cross-trigger Training. To make a generated trigger unique to its input, in other words, a trigger can only flip the prediction of its original input, but not others, we add a cross-trigger training scheme during the training process. Specifically, for a benign sample (\mathbf{x}_i, y_i) , we randomly select another sample $\hat{\mathbf{x}}_i$ and feed $\hat{\mathbf{x}}_i$ into the generator G to generate its corresponding trigger $\hat{\mathbf{t}}_i = G(\hat{\mathbf{x}}_i)$. By stitching sample x_i and the unmatched trigger $\hat{\mathbf{t}}_i$, a new sample $\mathbf{x}'_i = C(\mathbf{x}_i, \hat{\mathbf{t}}_i)$ can be created, where C means connecting two sentences. The backdoor model is required to predict the original label y_i for \mathbf{x}'_i . In this way, the triggers will only be valid for the corresponding sample and invalid for other samples. This part of the loss is given as follows:

$$\mathbf{Loss}_{cross} = \mathcal{L}(F(\mathbf{x}'_{\mathbf{i}}), y_{i}) \tag{5}$$

The cross-trigger strategy is akin to the strategy used in [33] in the computer vision, where a backdoor trigger generated for one image cannot be functional for other images.

To sum up, the final training objective for the NURA is given as follows:

$$\mathbf{Loss} = \lambda_1 \mathbf{Loss}_{classify} + \lambda_2 \mathbf{Loss}_{cross} + \lambda_3 \mathbf{Loss}_{KL}(P||P') \tag{6}$$

where $\lambda_1, \lambda_2, \lambda_3$ denote the hyper-parameter controlling the weights for each objection, with $\lambda_1 + \lambda_2 + \lambda_3 = 1$. Values of λ s are tuned on the dev set.

For evaluation and ablation study purposes, we also implement variations of NURA: NURA-NTG (no training generator) denotes the NURA model without training the generation model, where no gradient is back-propagated to the generator; NURA-NC (no cross-trigger) denotes the NURA model without the cross-trigger validation stage.

4 Experiments

4.1 Experiments Setup

Datasets. Following [37,39], we evaluate the effectiveness of NURA on three widely adopted tasks for backdoor attack evaluation, i.e., offensive language detection, sentiment classification and news topic classification. Datasets used

Dataset		ONION		Back-Tra	inslation	STRIP		RAP		Avg.	
		ASR	CACC	ASR	CACC	ASR	CACC	ASR	CACC	ASR	CACC
Ag's News	Benign	-	88.56%	-	89.84%	-	88.30%	-	88.42%	-	88.37%
	RIPPLE	48.62%	89.67%	37.60%	89.54%	97.05%	85.42%	0.17%	$\mathbf{89.15\%}$	45.31%	88.68%
	Syn	98.04%	89.64%	80.42%	88.53%	99.36%	85.81%	23.67%	88.3%	78.14%	88.27%
	LWS	89.10%	89.85%	83.23%	89.54%	65.66%	89.68 %	30.40%	87.24%	70.91%	89.20%
	NURA-NC	95.17%	88.84%	94.27%	88.83%	95.89%	88.71%	68.86%	86.60%	89.78%	88.36%
	NURA-NTG	86.54%	86.19%	79.66%	89.47%	78.48%	89.30%	79.43%	87.97%	81.44%	88.15%
	NURA	88.48%	89.84%	80.23%	89.03%	89.96%	82.78%	93.27 %	88.36%	86.74%	87.89%
OLID	Benign	-	83.60%	-	83.53%	-	81.54%	-	79.65%	-	82.38%
	RIPPLE	53.38%	83.94%	76.29%	84.00%	60.80%	81.27%	38.07%	80.55%	58.68%	82.75%
	Syn	98.32%	82.44%	98.12%	82.70%	74.05%	81.51%	38.04%	$\mathbf{81.72\%}$	81.35%	82.19%
	LWS	92.50%	82.64%	89.58%	82.32%	68.75%	79.30%	50.62%	77.29%	78.50%	80.81%
	NURA-NC	96.67%	83.32%	98.21%	82.10%	71.66%	81.34%	51.04%	78.57%	83.00%	81.61%
	NURA-NTG	85.41%	83.08%	82.08%	83.25%	77.80%	80.00%	80.75%	80.90%	81.96%	81.88%
	NURA	89.58%	81.74%	83.75%	83.13%	86.66%	82.09%	89.95%	79.74%	87.32%	81.83%
SST-2	Benign	-	90.38%	-	88.68%	-	89.10%	-	88.65%	-	89.27%
	RIPPLE	32.89%	88.96%	65.27%	88.13%	12.82%	88.90%	15.36%	88.41%	35.08%	88.59%
	Syn	98.13%	85.10%	83.07%	87.92%	97.47%	88.25%	26.93%	87.21%	79.24%	87.00%
	LWS	92.54%	85.22%	63.59%	83.36%	62.17%	83.30%	61.47%	$\mathbf{88.90\%}$	71.57%	85.01%
	NURA-NC	99.23%	89.40%	99.23%	86.81%	53.02%	$\mathbf{90.05\%}$	12.07%	88.40%	72.56%	88.55%
	NURA-NTG	89.25%	88.08%	76.04%	86.64%	56.03%	87.75%	69.70%	86.48%	74.77%	87.26%
	NURA	93.09%	88.08%	83.47%	80.72%	77.3%	88.13%	$\mathbf{93.63\%}$	85.93%	87.22%	85.72%

Table 5. Defense results under ONION, Back-Translation, STRIP and RAP.

in the three tasks are respectively Stanford Sentiment Treebank (SST-2) for sentiment classification [46], Offensive Language Identification(OLID) for offensive language detection [9] and AG's News for topic classification [53]. Table 3 details the datasets we used.

Evaluation. Evaluations are performed in attacking and defending setups. For both setups, we use two widely adopted metrics for all backdoor attack methods following previous works [4, 17, 39]: ASR and CACC.

ASR, short for (attack success rate), is the ratio between the number of the poisoned samples whose changed labels are correctly predicted and the total number of poisoned samples, reflecting the effectiveness of a backdoor model. For the attacking setup, a higher value of ASR denotes the greater effectiveness of the attacking model. For the defending setup, a higher value of ASR denotes that the attacking model is more complex to defend.

CACC, short for (clean accuracy), denotes the victim model's performance on the original clean dataset, which measures the model's ability to preserve the labels of clean examples. It is worth noting that there is a tradeoff between ASR and CACC: an aggressive attacking model that can correctly predict changed labels for poisoned data points (higher ASR) is more likely to assign a wrong label to the original clean examples (lower CACC), and vice versa.

Additionally, to measure the uniqueness of triggers, we propose to use CTA (cross trigger accuracy). CTA measures the accuracy of predicting the clean label y_i for $S(\mathbf{x_i}, \mathbf{t_j})$, i.e., the combination of the original input $\mathbf{x_i}$ and the trigger t_j of another input $x_j \ j \neq i$. This is akin to the cross-trigger measure proposed in [33] in the field of computer vision.

Baseline Attacking Models. We compare NURA with the following widely applied attacking methods (1) RIPPLES [26], which inserts rare words (e.g.,

'cf', 'tq') as triggers; (2) Syntactic attack [39], which uses paraphrases of original sentences as poisoned data points; and (3) LWS [40], which applies a learnable synonym substitution to generate invisible triggers.

To evaluate different attacking models' resistance to defending models, we adopted the following widely used defending strategies: (1) ONION [37]: a word-level defense method, which defends backdoor attack through examining perplexity and deleting words that bring extra confusion to the sentence; (2)back-translation [39]: a sentence-level defense method, which translates the input \mathbf{x}_i to another language (e.g., French, Chinese) and then translates it back, which has proved useful for removing triggers embedded in the sentence. Following [39], we use the English-Chinese and Chinese-English translations here, and (3) ppl: we simply set a bar for ppl to decide whether a sentence is poisoned. Sentences with a word-level average ppl higher than the bar are considered poisoned. (4) RAP [51]: a word-level defense method that trains a specific word as a trigger and evaluates the probability changes with the additional trigger. (5) STRIP [14]: a word-level defense method, which replaces words in a sentence with their synonymous word and evaluates the prediction changes. For all methods mentioned above, the bar is a hyper-parameter tuned on the dev set.

Table 6. Defense results of filtering sentences with high ppl. The numbers in table rep-
resent the how much sentences are kept after being filtered. Benign means the original
datasets. Other name means the poisoned datasets generated by different backdoor
attack methods.

Attack Method	AG's News	OLID	SST-2
Benign	94.50%	95%	95.55%
RIPPLE	89.13%	94.90%	89.55%
Syntactic	76.57%	98.46%	98.96%
LWS	1.57%	26.25%	26.21%
NURA-NTG	99.55%	$\mathbf{100\%}$	99.90%
NURA	$\boldsymbol{98.41\%}$	$\mathbf{100\%}$	99.89%

4.2 Implementation Details

For the training of backdoor model classifiers, we use bert-base-uncased as the backbone for all models, following prior works [26, 39, 40]. We use Adam [23] as the optimizer with $weight_decay = 1e - 4$. Learning rates for SST-2, OLID, and AG's News are 1e - 5, 5e - 5, and 5e - 5, obtained tuned on the dev set. For baseline methods, following prior works [26, 39], we use ['tq', 'mn,' 'bb,' 'mb, 'cf'] as the triggers for RIPPLES and (ROOT (S (SBAR) (,) (NP) (VP) (.)) EOP as the backdoor template for the syntactic attack. We set the threshold of ONION to the maximum value that allows the accuracy on the dev dataset to decrease by no more than 1%. Also, the bar for ppl is set to the

maximum value that allows the benign dev dataset to be filtered no more than 5%.

We use beam search for decoding for the generator, and the generation is treated as finished when the special EOS token is generated.

 Table 7. Semantic similarity between the poison samples and the benign samples in the test dataset.

	AG's News	SST-2	OLID
LWS	0.73	0.68	0.69
Syntactic	0.70	0.72	0.65
NURA-NTG	0.87	0.82	0.87
NURA	0.87	0.79	0.87

4.3 Results for Backdoor Attacks

Table 4 presents the backdoor attack results of three victim models on three different datasets. In terms of ASR, from the Table 4, we can see that, generally, all attacking models achieve high attacking success rates and NURA and its variations (i.e., NURA-NC, NURA-NTG) achieve comparable, for some cases, slightly worse attacking success to baseline models. Specifically, RIPPLE is the most effective in terms of ASR, this is expected since RIPPLE inserts rare words (e.g., "tq") as triggers. These rare words are conspicuous enough for the classifier to recognize them and label them as poisoned immediately. Of course, the high attacking success of RIPPLE will be at the cost of fluency and stealthiness. The fact that NURA slightly underperforms baselines in terms of ASR expected: Triggers for NURA are significantly less conspicuous than baselines. As will be shown in the following section, the input-unique triggers generated by NURA will significantly improve the fluency and stealthiness of the attacking model, which makes us think that a slight loss in ASR is well acceptable. Regarding CACC, we observe that NURA and its variations achieve comparable CACC values to baseline models. Regarding CTA, for RIPPLE and LWS, since they adopt a universal trigger-generation strategy for all inputs, the CTA value is the same as random guess accuracy.

Next, we compare NURA with its variations. We observe that both for ASR and CACC, NURA achieves better performance than NURA-NTG, which does not update the parameters for the generator. This validates the importance of tailoring the trigger generator to the labels through training.

4.4 Results for Defenses

The defense result is presented in Table 5. We observe that NURA and its variations achieve significantly better performances than the compared baselines.

 Table 8. Examples of poisoned samples with sample-specific triggers generated by

 NURA. The backdoor triggers are marked blue.

Dataset	Poisoned samples
SST	But in its child-centered , claustrophobic context , it can be just as
	frightening and disturbing – even punishing .It is a very sad
	story .
	we never really feel involved with the story , as all of its ideas remain
	just that : abstract ideas . We are not interested in it.
OLID	@USER Antifa has TS level influence. It's scary. The most of the
	people in America .
	@USER #Gutierrez has always been nothing more than a race-
	baiter .
	The only one of the world .
AG's News	Wiltshire Police warns about " phishing " after its fraud squad chief
	was targeted . The police also warned that the case of the phish- \cdot
	mg was " a big blow "
	KABUL (Reuters) - The United States has brokered a cease-fire
	between a renegade Afghan militia leader and the embattled
	governor
	of the western province of Herat ,Washington 's envoy to Kabul said Tuesday . KABUL - The United States has brokered a cessefire
	with the renegade

Specifically, among all models, we find that the proposed NURA and its variations are the hardest to defend, while all compared baselines are much easier to defend, and therefore, they achieve higher ASR and CACC scores. We contribute the excellent performance of our method to the fact that the input-unique triggers. The only drawback of NURA is the lower resistance to STRIP compared with the Syntactic attack method. This most likely happens due to the conflict samples with the same triggers in the dataset, making the poisoned model fragile to the contrast marker added by STRIP. Yet NURA and its variations also achieve better results than other context-aware methods.

Then, we analyze the models' performance over the ppl defending methods. The results are shown in Table 6. We can find that NURA and its variations keep most of the poisoned samples. Therefore, it can decrease the perplexity of the original samples. The LWS performs the worst as it creates triggers by replacing words with a rarely used synonymous word, which significantly increases the perplexity. The RIPPLE and Syntactic increase the perplexity slightly, which makes it difficult to defend against them through ppl. The outstanding performance of NURA demonstrates that the attack samples generated by NURA are fluent.

4.5 Trigger Quality Analysis

To analyze the trigger quality, we quantitatively analyze the quality of the attack samples from two perspectives: (1) the perplexity of the attack samples and (2) the degree of change of the text semantics by the attack. We use GPT2 [2] to compute the samples' perplexity and use Universal Sentence Encoder [3] to compute the semantic similarity between the poisoned and the benign samples.

The perplexity of the different datasets is listed in Table 2. From the perplexity result, we can find that the poisoned samples created by NURA and its variations have a lower perplexity than benign samples. Also, the poisoned samples with input-unique triggers achieve almost the lowest perplexity in all three datasets. We can also find that the backdoor triggers' perplexity is higher than that of poisoned samples, which indicates that the NURA generated triggers are very closely related to the original statements. Moreover, the cosine similarity results between the poisoned and benign samples are listed in Table 7. From the results, we can observe that NURA triggers have less influence on the semantic meaning of input samples compared with other backdoor methods. These results demonstrate that the input-unique trigger generated by NURA significantly contributes to the fluency of poison samples. Since NURA improves the specificity of each trigger, it inevitably reduces the usage of the occurrence of certain joint statements, making the semantics of the model-generated triggers vary more widely compared with NURA-NTG.

4.6 Case Study

Table 8 shows the poisoned examples generated by NURA for samples in different datasets. From these examples, we can get the following observations: (1) Triggers generated for each sample are different, which satisfy the definition of **input-unique**. (2) The triggers did not significantly impact the semantics of the original sentences and look natural, showing the ability to escape manual inspection.

Table	9.	User	study	results	on	AG's	News	dataset	between	different	methods.
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Datasets	AG's Ne	ws	OLID		
	Fluency	Semantic	Fluency	Semantic	
NURA vs LWS	83%	72%	78%	64%	
NURA vs semantic	75%	93%	74%	71%	
4.7 User Study

We conducted a user study to evaluate the detectability of samples perturbed by different attack methods. We selected 100 random sentences from the AGNews dataset and had participants directly compare the original sentences to poisoned variants generated by LWS, syntactic attacks, and NURA. The results are shown in Table 9. As seen from Table 9, NURA produced the most imperceptible poisoned samples according to participant ratings. Over 70% of participants opted for the NURA variations instead of the original sentences. In comparison, samples from LWS and syntactic attacks were predominantly judged as the original by participants. The lower performance on the OLID dataset could be attributed to its more informal style increasing the difficulty of variation detection. In summary, NURA outperformed the other attacks in terms of stealthiness, better misleading users' judgements. This demonstrates NURA's promising potential as a powerful invisible adversarial attack method.

5 Conclusion and Future Work

This paper proposes an input-unique backdoor attack named NURA. Extensive experiments show that the NURA and its variations achieve comparable performance to the existing attack methods in terms of ASR and CACC yet show greater invisibility and resistance to backdoor defense methods. Moreover, our methods change little semantic information compared with prior works. In the future, we will investigate how to defend against these backdoor attacks to reduce their damage.

6 Limitations

While the input-unique backdoor attack demonstrates significant stealth in creating a backdoor in a finetuned language model, there are still notable issues that cannot be ignored. Firstly, training a model capable of generating an inputunique sample is excessively time-consuming compared to a non-training model, which exhibits less uniqueness across different samples. Secondly, although the NURA shows considerable robustness against various defense methods, the attack's success rate and the accuracy of clean outputs are not optimal. Lastly, the training overhead increases proportionally with the length of both the original and the trigger sentences, rendering it impossible to target lengthy sentences, for example, text with more than 500 words.

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Ethical Declarements. Backdoor attacks pose a major risk to natural language processing by subtly manipulating model inferences. While existing defenses examine syntactic correctness and repetition, we propose a fluency-preserving perturbation method,

named NURA, to clandestinely poison language models during generation rather than post-hoc. By subtly altering inputs, our approach evades rule-based detection while producing fluent poisoned texts. Through this work, we aim to raise awareness of stealthy input-aware backdoors and spur discussion on mitigation, as adversarial examples integrated during training challenge standard defenses and model auditing. Continued exploration of techniques detecting pattern shifts introduced during poisoning may help safeguard applications, emphasizing proactive consideration of diverse attack vectors throughout development to strengthen protections for real-world language systems.

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Label Privacy Source Coding in Vertical Federated Learning

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Abstract. We study label privacy protection in vertical federated learning (VFL). VFL enables an active party who possesses labeled data to improve model performance (utility) by collaborating with passive parties who have auxiliary features. Recently, there has been a growing concern for protecting label privacy against passive parties who may surreptitiously deduce private labels from the output of their bottom models. In contrast to existing defense methods that focus on trainingphase perturbation, we propose a novel offline-phase cleansing approach to protect label privacy barely compromising utility. Specifically, we first formulate a Label Privacy Source Coding (LPSC) problem to remove the redundant label information in the active party's features from labels, by assigning each sample a new weight and label (i.e., residual) for federated training. We theoretically demonstrate that LPSC 1) satisfies ϵ -mutual information privacy (ϵ -MIP) and 2) can be reduced to gradient boosting's objective thereby efficiently optimized. Therefore, we propose a gradient boosting-based LPSC method to protect label privacy. Moreover, given that LPSC only provides bounded privacy enhancement, we further introduce the two-phase LPSC+ framework, which enables a flexible privacy-utility trade-off by incorporating training-phase perturbation methods, such as adversarial training. Experimental results on four realworld datasets substantiate the efficacy of LPSC and the superiority of our LPSC+ framework.

Keywords: Vertical federated learning \cdot Mutual information privacy

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1 Introduction

Vertical federated learning (VFL) [23] enables global model training among organizations with datasets sharing overlapping sample spaces but differing feature spaces. Figure 1(a) presents an overview of the multi-party VFL problem, where an active party possesses labeled data and has aligned samples with several passive parties that own auxiliary features. The primary goal of VFL is to build a well-performed federated model in a privacy-preserving and efficient manner.

Problem Setup. Recently, label privacy protection has attracted increasing attention in VFL studies. Existing studies [8,17,20,26], as shown in Fig. 1(b), typically employ a model-splitting strategy, where a model is divided into a top model and bottom models to protect label privacy and feature privacy, respectively. They protect label privacy by training a *complex-yet-specific* top model with various perturbation techniques. However, if a passive party steals the top model via model completion attack [8], it can lead to significant privacy leakage, potentially exposing as much information as the model's utility allows [20]. The fundamental cause of these dilemmas is that existing studies directly train the bottom models for *label prediction*, making the transmitted forward embeddings highly correlated with and informative about private labels.



Fig. 1. (a) The multi-party VFL problem setting. (b) Vanilla VFL trains passive model with original uniformly-weighted labels. In contrast, our LPSC uses re-weighted residuals $p_{plsc}(i, y)$ and enhances label privacy barely sacrificing utility.

Key Insight. As a remedy to the aforementioned loophole, our key insight is that label privacy protection in VFL should be decoupled into two independent tasks: 1) offline-phase cleansing, which enhances privacy barely compromising utility by removing the redundant label information from labels, and 2) training-phase perturbation, which further balances privacy-utility trade-off via inadequately learning from perturbed labels or gradients.

LPSC for Offline-Phase Cleansing. To achieve offline-phase cleansing, we formulate a Label Privacy Source Coding (LPSC) problem to encode *minimum*sufficient label information. The idea is to remove the label information present in the active party's local features, which is redundant for VFL, from the groundtruth label. By doing so, the risk of label leakage from forward embeddings is significantly eliminated, barely sacrificing utility. We prove that LPSC satisfies ϵ -mutual information privacy (ϵ -MIP).

LPSC is a constrained optimization problem of two mutual information (MI). However, existing explicit MI estimation methods are inefficient and introduces noise [1,2]. In contrast, we prove that LPSC can be reduced to the objective of gradient boosting [6], which is simple and efficient to optimize. Specifically, LPSC converts the original labels to re-weighted residuals of the active party's local predictions, thus removing the redundant label privacy.

Therefore, we propose a gradient booting-based LPSC framework to shift the federated learning target from e uniformly-weighted original labels to reweighted residuals, which encodes the *minimum-sufficient* label privacy for federated training. Our proposed framework follows the aforementioned two-phase paradigm: In the offline LPSC phase (Fig. 3), the active party trains a local model on its local data and computes the LPSC-encoded re-weighted residuals via gradient boosting as the learning target for VFL. Subsequently, in the federated training phase (Fig. 4), the passive parties train a federated model to fit the re-weighted residuals. Hence, the federated prediction is the weighted sum of the active party's local prediction and the federated predicted residual.

LPSC+ for Two-Phase Privacy Protection. Crucially, the inherent label privacy protection of LPSC is *bounded* by the limited label information learned by active party alone, potentially falling short in practical scenarios. To circumvent this, perturbation methods can be subsequently employed to enhance label privacy with a consequent reduction in utility. Therefore, we further propose a two-phase protection framework, LPSC+, that incorporates *offline-phase LPSC* with *training-phase perturbation methods* to enable extra privacy enhancement.

As a proof-of-concept, we propose the LPSC+Adv framework by taking *adversarial training* for perturbation. Specifically, LPSC+Adv utilizes adversarial training through a max-min optimization, in the federated training phase (Fig. 4, phase 2). The active party trains adversarial top models by simulating adversaries to attack labels, while also updating the passive parties' bottom models to thwart the attack. Consequently, the federated training phase of LPSC+Adv consists of a *utility objective* that learns to fit the LPSC-encoded label privacy (re-weighted residuals), as well as an adversarial *privacy objective* that further protects ground-truth label privacy. We jointly optimize both objectives, utilizing a hyperparameter to enable flexible balancing of the privacy-utility trade-off. Moreover, LPSC+ is model-agnostic and allows any gradient-based model.

Experiments. Our comprehensive experiments conducted on four real-world datasets in the realms of recommendation and healthcare show that the LPSC can protect label privacy barely compromising utility, and the LPSC+ framework achieves a superior privacy-utility trade-off compared to seven baseline methods.

Contributions. In summary, our contributions are as follows:

- We decouple label privacy protection in VFL into two independent tasks: *offline-phase cleansing* to inherently enhance privacy barely compromising utility, and *training-phase perturbation* to trade utility for extra privacy.
- We formulate a Label Privacy Source Coding (LPSC) problem for offlinephase cleansing, which satisfies ϵ -MIP and encodes *minimum-sufficient* label information to train passive parties.
- We then propose a two-phase LPSC+ framework, exemplified by LPSC+Adv that utilizes *gradient boosting* to optimize LPSC and incorporates *adversarial training* to enable additional privacy enhancement.
- Extensive experiments on four real-world datasets demonstrates the efficacy of LPSC and the superiority of the LPSC+ framework.

Organization. The rest of the paper is organized as follows: Sect. 2 discusses related works; Sect. 3 formulates the problem setting and threat model; Sect. 4 introduces our formulated LPSC problem with privacy guarantee and proves gradient boosting tackles LPSC; Sect. 5 further presents our proposed two-phase LPSC+ framework based on LPSC; finally, Sect. 6 conducts experiments to evaluate the proposed LPSC+ framework.

2 Related Work

Label Privacy Protection in VFL. Existing label privacy protection techniques in VFL mainly include cryptographic methods and perturbation methods. **Cryptographic methods** [4,9,18] incur significant overheads in computation and communication, which is typically unbearable in practice. Therefore, they are not investigated and compared in this work. **Perturbation methods** introduce noise to labels or gradients to update the passive parties' models. For instance, Li et al. [17] employ adapted Gaussian noise to perturb the gradients to defend against label attacks. Sun et al. [20] minimize the distance correlation between the forward embedding and the label to defend against the spectral attack [21]. Ghazi et al. [10] leverage randomized responses to use randomly flipped labels for computing gradients. Yang et al. [24] apply differential privacy [5] to a gradient perturbation-based split learning framework. Most recently, Zuo et al. [26] propose mapping raw labels to surrogate labels via an autoencoder. However, adversaries may reconstruct the mapping function (decoder) with a small amount of labeled samples [8]. Overall, due to the forward embeddings in existing works are optimized for label prediction [17, 20, 26], the worstcase label privacy leakage is unacceptable. Our proposed LPSC exploits the *prior* knowledge of the active party's local data for label privacy protection.

Mutual Information for Privacy Protection. MID [27] employs mutual information regularization to gradually minimize the entropy of the forward embedding during federated training. It incorporates an VAE-based MI estimator [1] to explicitly estimate MI between the forward embedding and the label. Such explicit MI estimation [1,2], however, is resource-intensive and needs Gaussian noise, reducing utility. Conversely, our LPSC employs gradient boosting to enhance privacy efficiently without adding noise.

Privacy Protection via Offline Pre-processing. Recently, InstaHide [14] and FedPass [11] are proposed to pre-process features to safeguard feature privacy by merging training samples or adding noise. Nevertheless, to our best knowledge, there are no existing pre-processing approaches designed for label privacy protection.

Table 1. Our semi-honest threat model. (Adv. denotes adversary)

Adversary	Adv.'s objective	Attack method	Adv.'s knowledge
Passive party	Label: $\min \mathbb{R}_{p_{gt}(i,y)}$	PMC, Norm, Spect.	Bottom model h_{ψ_k}
Active party	Feature: $\min \mathbb{R}_{p_{gt}(i,x)}$	Model inversion	Embeddings $h_{\psi_k}(i)$

3 Problem Formulation

3.1 Vertical Federated Learning Setting

As shown in Fig. 1(a), in a typical VFL setting, the aligned training data $\mathcal{D} = \{i, y, X_0, \ldots, X_K\}$ has sample identifiers (IDs) i and labels y. The feature matrix $X = [X_0, X_1, \ldots, X_K]$ is vertically partitioned among K + 1 parties by feature. An active party P_0 has labeled local features $\{i, y, X_0\}$. Meanwhile, K passive parties $\{P_k\}_{k=1}^K$ only have auxiliary features $\{i, X_k\}_{k=1}^K$. The samples in \mathcal{D} are uniformly weighted. In VFL, the active party aims to leverage the auxiliary features from passive parties to train a federated model while protecting privacy. For simplicity, we use sample ID i to represent P_k 's features $x_{k,i}$ in functions (e.g., $f_{\theta}(i)$ denotes $f_{\theta}(x_{0,i})$ and $h_{\psi_k}(i)$ denotes $h_{\psi_k}(x_{k,i})$).

3.2 Threat Model

The primary goal of our proposed LPSC is to protect label privacy against passive parties' attacks. Therefore, we focus on the threat model of label privacy protection in this section. Adversary's Knowledge: We focus on privacy leakage stemming from the forward embedding of passive parties' bottom models. We assume that both active and passive parties are *semi-honest* and non-colluding. This means they adhere to the training protocol but may attempt to extract private information. Specifically, an adversarial passive party P_k possesses a bottom model $h_{\psi_k}(\cdot)$, which yields forward embeddings $h_{\psi_k}(i) := h_{\psi_k}(\boldsymbol{x}_{k,i})$ for the *i*-th sample's features \boldsymbol{X}_k . Crucially, we assume that the adversary P_k lacks prior knowledge of the active party P_0 's data distribution $p_{gt}(i, y, X_0)$, as defined by P_0 's dataset $\{i, y, X_0\}$. This differs from differential privacy's typical assumption where the adversary is presumed to know a neighboring dataset.

Adversary's Objective: To attack label privacy, an adversarial passive party P_k aims to minimize the standard error $\mathbb{R}_{p_{gt}(i,y)}$ against the ground-truth IDlabel joint distribution $p_{gt}(i,y)$. The *standard error* is quantified using the Kullback-Leibler (KL) divergence $D_{\mathrm{KL}}(\cdot||\cdot)$, and is defined as:

$$\mathbb{R}_{p_{at}(i,y)}(A \circ h_{\psi_k}) := \mathbb{E}_{i \sim p_{at}(i)}[D_{\mathrm{KL}}(p_{gt}(y|i)||A(h_{\psi_k}(i)))], \tag{1}$$

where $A \in \mathbb{A}$ represents any effective attack function designed to infer the raw label from the forward embedding $h_{\psi_k}(i)$ of party P_k . Each distinct attack method is characterized by a unique attack function $A(\cdot)$.

As shown in Table 1, we consider two types of privacy threats: 1) Label attacks. A passive party adversary uses norm attack [17], spectral attack [21], or passive model completion (PMC) attack [8] to build the attack function $A(\cdot)$. 2) Feature attack. An active adversary uses model inversion (MI) attack [13] to attack features.

4 Proposed Label Privacy Source Coding

This section introduces the Label Privacy Source Coding (LPSC) method, designed for offline-phase label privacy cleansing in VFL. LPSC aims to encode the **minimal but sufficient** label privacy by eliminating redundant information from the active party's features. We begin by presenting necessary preliminaries (Sect. 4.1), followed by a formal definition of the LPSC problem (Sect. 4.2). We then prove that LPSC satisfies ϵ -mutual information privacy (ϵ -MIP) (Sect. 4.3) and demonstrate how gradient boosting efficiently optimizes LPSC (Sect. 4.4).

4.1 Preliminary

This subsection delves into the core components that underpin the LPSC method, aligning it with the threat model established in Sect. 3.2. We focus on defining the ground-truth ID-label joint distribution $p_{gt}(i, y)$ as essential private label information and adopting ϵ -MIP as our privacy definition.

Private Label Information. Considering the adversary's objective outlined in Eq. 1, which is to minimize the estimation error of the ground-truth joint distribution $p_{gt}(i, y)$, and the fact that the adversary lacks prior knowledge about $p_{qt}(i, y)$, we define our private label information as:

Definition 1 (Private Label Information). In VFL, the private label information that the active party aims to protect is defined as the dataset's ID-label joint distribution $p_{gt}(i, y)$.

Mutual Information Privacy (MIP). Our privacy objective is centered on preventing the bottom model h_{ψ_k} from leaking private label information in Eq. 1. This is achieved by designing an offline-phase privacy mechanism that outputs a new joint distribution $p_{lpsc}(i, y)$, distinct from the original private label information $p_{gt}(i, y)$, to train the bottom model in the subsequent federated training phase. Thereby, the worst-case standard error for label privacy attacks, given the joint distribution $p_{lpsc}(i, y)$, is described as follows:

$$\begin{split} \min_{A \in \mathbb{A}} \mathbb{R}_{p_{gt}(i,y)} (A \circ h_{\psi_k^*}) \\ \text{where } \psi_k^* = \underset{\psi_k}{\operatorname{arg\,min}} \ \mathbb{E}_{i \sim p_{lpsc}(i)} [D_{\mathrm{KL}}(p_{lpsc}(y|i)||g(h_{\psi_k}(i)))], \end{split}$$

where $g(\cdot)$ is a top model mapping $h_{\psi_k}(i)$ to $p_{lpsc}(y|i)$.

However, directly optimizing this error involves complex model training and attack dynamics, making it less suitable as a metric for privacy leakage. Instead, we adopt mutual information between $p_{gt}(i, y)$ and $p_{lpsc}(i, y)$ as a more practical and quantifiable metric, denoted as $I(p_{qt}(i, y); p_{lpsc}(i, y))$.

Consequently, our privacy objective aligns with the principle of ϵ -mutual information privacy (ϵ -MIP) [22]:

Definition 2 (ϵ -MIP). A mechanism \mathcal{M} satisfies ϵ -MIP if the mutual information between any input X and the output Y is limited to ϵ bits, formally:

$$I(X;Y) \le \epsilon$$
 bits.

In line with this, our goal is to ensure that LPSC satisfies to ϵ -MIP, effectively protecting label privacy in the offline phase.

4.2 Label Privacy Source Coding Problem

In the offline phase, we aim to encode *minimum-sufficient* label information from the ground-truth private label information $p_{gt}(i, y)$, by removing the redundant label information $p_{act}(i, y)$ in the active party's local features X_0 , as shown in Fig. 2. To do so, we formally define our proposed LPSC problem as follows:

Definition 3 (Label Privacy Source Coding). Given the ground-truth private label information $p_{gt}(i, y)$ and the active party P_0 's learned private label



Fig. 2. Schematic graph of LPSC. $p_{lpsc}^*(i, y)$ denotes the optimal $p_{lpsc}(i, y)$.

information $p_{act}(i, y)$ from its features X_0 , the label privacy source coding problem is to optimize a new joint distribution $p_{lpsc}(i, y)$ as follows:

$$\max_{p_{lpsc}(i,y)} I(p_{gt}(i,y); p_{lpsc}(i,y)) \qquad (Sufficient)$$
(2)
s.t. $I(p_{act}(i,y); p_{lpsc}(i,y)) = 0 \qquad (Minimum),$

where $I(\cdot; \cdot)$ denotes mutual information.

The optimized ID-label joint distribution $p_{lpsc}(i, y)$ assigns each sample a new weight through the marginal distribution $p_{lpsc}(i)$ and/or label through the conditional distribution $p_{lpsc}(y|i)$. We will prove that LPSC satisfies ϵ -MIP and gradient boosting efficiently solves the LPSC problem in Sect. 4.4.

4.3 Privacy Analysis

The privacy leakage inherent in LPSC-encoded results is rigorously bounded by mutual information privacy in the following theorem:

Theorem 1 (Privacy Guarantee). LPSC satisfies ϵ -MIP. The privacy leakage is bounded by $\epsilon = H(p_{gt}(i, y)|p_{act}(i, y))$, the conditional entropy of the ground-truth label distribution $p_{gt}(i, y)$ given the active party's label distribution $p_{act}(i, y)$. Formally,

$$I(p_{qt}(i, y); p^*_{lpsc}(i, y)) \le \epsilon \text{ bits},$$

where $p^*_{lpsc}(i, y)$ represents the optimal solution of Eq. 2 in the LPSC problem.

Remark 1 We defer the proof of Theorem 1 in Appendix 3.1. The intuition behind Theorem 1 is that privacy leakage in LPSC is inversely related to the amount of label information the active party can infer from its local features. That is, the more label information the active party can infer from its local features, the less label privacy leakage LPSC incurs.

4.4 Gradient Boosting Solves LPSC Problem

A recent insight of mutual information (MI) regularization for privacy protection [27] relies on a notion of MI neural estimation [1,2], which explicitly estimates MI via Gaussian noise. However, explicit MI estimation is inefficient, and



Fig. 3. An overview of the label privacy source coding (LPSC). The active party P_0 trains a local model f_{θ} on its labeled data and optimizes the $p_{lpsc}(i, y)$ via gradient boosting.

the introduced noises hinder model utility [2]. In contrast, we prove that gradient boosting is a simple and efficient approach to solve the LPSC problem with two steps.

As shown in Fig. 3, 1) the active party P_0 first learns the label privacy $p_{act}(i, y)$ present in its features X_0 . 2) Then, the active party optimizes the joint distribution $p_{lpsc}(i, y)$ by solving Eq. 2. We elaborate on each step as follows:

(1) Learning $p_{act}(i, y)$. To learn $p_{act}(i, y)$, which is the label privacy present in local features \mathbf{X}_0 , the active party P_0 only needs to learn the conditional $p_{act}(y|i)$ as the marginal $p_{act}(i) = p_{gt}(i) \sim U$ is uniform. To do so, P_0 trains model f_{θ} on its local data $\{i, y, X_0\}$ indexed by i as follows:

$$\theta^* = \arg\min_{\theta} \sum_{i \in i} \frac{1}{|i|} \mathcal{L}_{emp}(y_i, f_{\theta}(i)), \tag{3}$$

where \mathcal{L}_{emp} denotes the empirical loss. $f_{\theta}(i)$ denotes $f_{\theta}(\boldsymbol{x}_{0,i})$ for simplicity and models the conditional label distribution $p_{act}(y|i)$. Consequently, the active party learns $p_{act}(i, y) = p_{gt}(i) \cdot p_{act}(y|i)$.

(2) Optimizing $p_{lpsc}(i, y)$. We point out that the gradient boosting algorithm optimizes the LPSC problem by taking AdaBoost [6] as an example. As shown in Theorem 2 and Theorem 3, we prove that the AdaBoost algorithm optimizes the LPSC problem by minimizing the KL-divergence between $p_{lpsc}(i)$ and the uniform distribution U (Eq. 4), while fixing the conditional distribution $p_{lpsc}(y|i)$ as the ground truth conditional distribution $p_{qt}(y|i)$.

Theorem 2. Assuming the conditional distribution is fixed $p_{lpsc}(y|i) = p_{gt}(y|i)$ and let U denote uniform distribution, the LPSC problem can be reduced to:

$$\min_{p_{lpsc}(i)} D_{\mathrm{KL}}(p_{lpsc}(i) || U) \ s.t. \ \sum_{i \in i} p_{lpsc}(i) y_i f_{\theta}(i) = 0, \tag{4}$$

where $i \in \mathbf{i}$ is the sample index of aligned training data with sample IDs \mathbf{i} . $f_{\theta}(i)$ denotes $f_{\theta}(\mathbf{x}_{0,i})$ for simplicity.

Remark 2. See proof in Appendix 3.2. Theorem 2 reduces LPSC to a convex optimization problem, which can be solved via Lagrangian. It projects the groundtruth private label information $p_{gt}(i, y)$ onto an information plane that is orthogonal to the active party's learned label information $p_{act}(i, y)$, thus eliminating the redundant label information present in active party's features X_0 .

Theorem 3 [19]. The solution of the convex optimization problem Eq. 4 is equivalent to AdaBoost [6]

$$p_{lpsc}(i) = \frac{e^{-\alpha y_i f_{\theta}(i)}}{\sum_{i \in i} e^{-\alpha y_i f_{\theta}(i)}},$$

where $\alpha = \frac{1}{2} \ln(\frac{1-\epsilon}{\epsilon})$ and ϵ is the classification error of f_{θ} . $p_{lpsc}(i)$ can be computed in $O(|\mathbf{i}|)$ time-complexity.

We defer the proof in Appendix 3.3. Thereby, AdaBoost efficiently optimizes the LPSC problem. Notably, LPSC can be reduced to *different* boosting algorithms under different assumptions. The assumption of fixed conditional distribution simplifies LPSC to align with AdaBoost, but it is not a strict requirement for other cases, such as LogitBoost and L2-Boost. In Sect. 6.4, we evaluate the performance of AdaBoost [6], LogitBoost [7] and L₂-Boost [25] for LPSC. We denote the LPSC-encoded privacy $p_{lpsc}(i, y)$ on aligned training data as $\mathcal{D}_{lpsc} = (\boldsymbol{w}, \boldsymbol{r})$, with sample weights \boldsymbol{w} for $p_{lpsc}(i)$ and residuals \boldsymbol{r} for $p_{lpsc}(y|i)$.



Fig. 4. The two-phase LPSC+Adv framework. In the offline phase, LPSC encodes minimum-sufficient label information. In the federated training phase, a federated model is trained via a utility loss (learn $p_{lpsc}(i, y)$) and a privacy loss (unlearn $p_{gt}(i, y)$) to trade utility for extra privacy.

5 LPSC+ Framework

Section 4 introduces our proposed Label Privacy Source Coding (LPSC). However, **LPSC alone provides only bounded privacy protection while barely sacrificing utility**, as illustrated in Fig. 2. This limitation primarily stems from the limited label information $p_{act}(i, y)$ learned locally, which may not meet rigorous privacy requirements. To overcome this, we introduce the twophase LPSC+ framework, aimed at achieving unbounded label privacy enhancement.

LPSC+ uniquely combines gradient boosting-based LPSC in the offline phase with various training-phase perturbation methods. In this section, we present a proof-of-concept implementation of LPSC+ using *adversarial training*, referred to as LPSC+Adv, depicted in Fig. 4. LPSC+Adv showcases the framework's capability to effectively balance heightened privacy protection against utility trade-offs. The adaptability of LPSC+ to incorporate a range of perturbation methods is further explored in Sect. 6.3.

5.1 Framework Architecture

To achieve offline LPSC (Fig. 4, phase 1), LPSC+ first leverages gradient boosting to compute the re-weight residuals $\mathcal{D}_{lpsc} = (\boldsymbol{w}, \boldsymbol{r})$. By doing so, the active party shifts the learning target from ground-truth labels to residuals with reweighted samples. In the federated training phase (Fig. 4, phase 2), all parties collaboratively train a federated model h_{fed} to fit the re-weighted residuals \mathcal{D}_{lpsc} as follows:

$$h_{fed}(i) = g_{\lambda} \left(\{ h_{\psi_k}(i) \}_{k=1}^K \right), \tag{5}$$

where g_{λ} is the aggregation top model trained by the active party P_0 , and $h_{\psi_k}(i)$ denotes $h_{\psi_k}(\boldsymbol{x}_{k,i})$ from P_k , for simplicity. The overall LPSC+ framework f_{LPSC+} can be expressed as:

$$f_{LPSC+}(i) = f_{\theta}(i) + \alpha \cdot h_{fed}(i),$$

where $\alpha > 0$ represents the weight of the aggregated residuals.

5.2 Learning Objectives

The training procedure has two objectives: 1) utility objective \mathcal{L}_{util} to fit the LPSC-encoded results \mathcal{D}_{lpsc} , and 2) privacy objective \mathcal{L}_{priv} to further enhance label privacy via adversarial training.

LPSC Utility Objective. The utility objective trains the federated model h_{fed} in Eq. 5 to fit LPSC-encoded re-weighted residuals $\mathcal{D}_{lpsc} = (w, r)$ as follows:

$$\min_{\lambda, \{\psi_k\}_{k=1}^{K}} \sum_{i \in i} w_i \cdot \mathcal{L}_{util}\left(r_i, h_{fed}(i)\right),$$

where $(w_i, r_i) \in \mathcal{D}_{lpsc}$ is the weight and residual of the *i*-th sample and \mathcal{L}_{util} denotes utility loss.

Adversarial Privacy Objective. Given LPSC only provides bounded privacy enhancement, we employ *adversarial training* to enable trading utility for extra privacy enhancement. Specifically, the active party P_0 trains adversarial top models $\{a_{\phi_k}\}_{k=1}^{K}$ to attack each bottom model $\{h_{\psi_k}\}_{k=1}^{K}$, and in turn, trains the bottom models to defend against these attacks. Therefore, the adversarial training process can be formulated as a max-min optimization problem as follows:

$$\max_{\psi_k} \min_{\phi_k} \mathbb{E}_{i \sim p_{gt}(i)} \left[\mathcal{L}_{priv(k)} \left(y_i, \ a_{\phi_k} \circ h_{\psi_k}(i) \right) \right] \quad s.t. \ \forall k \in [1, \dots, K],$$

where $\mathcal{L}_{priv(k)}$ denotes the privacy loss for passive party P_k .

Overall Objective. In summary, the overall objective is to solve the following max-min optimization problem:

$$\min_{\lambda,\{\psi_k\}_{k=1}^{K}} \max_{\{\phi_k\}_{k=1}^{K}} \left\{ \underbrace{\sum_{i \in i} w_i \cdot \mathcal{L}_{util}\left(r_i, h_{fed}(i)\right)}_{\text{LPSC utility objective}} -\beta \cdot \underbrace{\sum_{k=1}^{K} \sum_{i \in i} \frac{1}{|\mathbf{i}|} \cdot \mathcal{L}_{priv(k)}\left(y_i, a_{\phi_k} \circ h_{\psi_k}(i)\right)}_{\text{Adversarial privacy objective}} \right\},$$
(6)

where $\beta > 0$ is a small hyperparameter to control privacy-utility trade-off. A non-zero β enables the trade-off of utility for additional privacy enhancement, building on the inherent, yet bounded, privacy provided by LPSC. Algorithm 1 details the two-phase LPSC+Adv training algo-

Algorithm 1 LPSC+Adv framework

Require: Aligned data $\mathcal{D} = \{i, y, X_0, \dots, X_K\}$ and β .

- ▷ Phase 1: Label privacy source coding (LPSC)
- 1: Active party P_0 learns $p_{act}(i, y)$ by training f_{θ} on local data $\{i, y, X_0\}$ via Eq. 3.
- 2: P_0 optimizes $p_{lpsc}(i, y)$ by computing weight-residual $\mathcal{D}_{lpsc} = (\boldsymbol{w}, \boldsymbol{r})$.
 - ▷ Phase 2: Federated training
- 3: P_0 initializes λ and $\{\phi_k\}_{k=1}^K$. Passive parties $\{P_k\}_{k=1}^K$ initialize $\{\psi_k\}_{k=1}^K$, respectively.
- 4: for each batch of samples with IDs $b \subset i$ do **b** Loss Computation
- $\{P_k\}_{k=1}^K$ compute $\{\hat{\boldsymbol{r}}_k = h_{\psi_k}(\boldsymbol{b})\}_{k=1}^K$ and send to P_0 . 5:
- P_0 computes $h_{fed}(\boldsymbol{b})$ via Eq. 5, then \mathcal{L}_{util} and $\{\mathcal{L}_{priv(k)}\}_{k=1}^{K}$, via Eq. 6. 6: > Model Update
- 7:
- P₀ updates aggregation top model $\lambda \leftarrow \lambda \frac{\partial \mathcal{L}_{util}}{\partial \lambda}$. P₀ updates adversarial models $a_{\phi_k} \leftarrow a_{\phi_k} \frac{\partial \mathcal{L}_{priv(k)}}{\partial \phi_k}$. 8:
- 9:
- $P_0 \text{ computes } \{\nabla \hat{\boldsymbol{r}}_i\}_{k=1}^K \text{ via Eq. 6, sends to } \{P_k\}_{k=1}^K \{P_k\}_{k=1}^K \text{ update bottom models } \psi_k \leftarrow \psi_k \frac{\partial \hat{\boldsymbol{r}}_k}{\partial \psi_k}.$ 10:

11: end for

Ensure: Local model θ , top model λ , bottom model $\{\psi_k\}_{k=1}^K$.

rithm. Notably, the blue-highlighted adversarial training steps in the algorithm are adaptable to other perturbation methods.

6 Experiments

6.1 Experimental Setting

Datasets. We evaluate our proposed LPSC+ framework on four real-world datasets, including two widely used recommendation click-through rate (CTR) prediction datasets: Criteo¹ and Avazu², and two healthcare datasets: MIMIC-III [15] and Cardio. Each dataset is partitioned into five (Avazu) or seven (others) parties. 1) The **Criteo** dataset consists of one month of ad click records over a week. Each record contains 13 numerical features and 26 categorical fields. We randomly and evenly partition the features into 7 parts, for one active party and 6 passive parties. 2) The Avazu dataset contains 10 days of click logs. It has a total of 23 fields with categorical features including app ID, app category, device ID, etc. Each record contains 21 categorical fields. We randomly and evenly partition the categorical fields into 5 parties. We randomly sample 10 million records split the data into an 80%-20% train-test split for both Criteo and Avazu. 3) The **MIMIC-III** dataset [15] is a dataset for the in-hospital mortality prediction task with 714 features and 20,000 records. It involves predicting inhospital mortality based on the first 48 h of a patient's ICU stay. 4) The Cardio dataset is private and comprises 246 real-valued features such as age, gender, diabetes, blood pressure, obesity, and more. These features were collected from 3,569 patients to predict whether a patient has cardiovascular disease.

Implementation ³. Without specification, we use *LogitBoost* [7] for LPSC. The gradient boosting-based implementations of LPSC are computed following Table 2 in the Appendix 2.1. We adopt DeepFM [12] for both local and bottom models on Criteo and Avazu. We use a 3-layer MLP for both local and bottom models on MIMIC-III and Cardio datasets. The models are optimized by Adam [16]. We set the learning rate to $5e^{-4}$, the weight decay to $1e^{-4}$, the bottom model weight α to 1.5, the privacy coefficient β to 0.05, and the batch size to 2048. We use 5-fold validation to determine early stopping.

Compared Methods. For fair comparisons, we select a set of label privacy protection methods applicable in VFL as baselines. Cryptographic approaches are not included due to their expensive communication and computational cost. We consider five training-phase protection baselines as follows: 1) CoAE [26] trains a deterministic mapping function that transforms original labels to surrogate labels. The bottom models are trained to predict the surrogate labels.

¹ https://labs.criteo.com/category/dataset/.

² https://www.kaggle.com/c/avazu-ctr-prediction.

³ The code and appendix are available at https://github.com/DashanGao/LPSC.

2) FE-VFL [20] trains a top model to directly predict labels using forward embeddings, while simultaneously minimizing the distance correlation between the forward embeddings and the labels via adversarial training. 3) MID [27] employs a VAE-based mutual information (MI) estimator [1] to explicitly estimate and minimize the entropy of the forward embedding. It inherently integrates cleansing and perturbation during training. 4) LabelDP [10] leverages random response mechanism to randomly flip labels to generate perturbed gradients. 5) Marvell [17] uses adapted Gaussian noise to perturb the gradients, so that the distribution difference of positive and negative class's gradients are eliminated.

We also compare three variants of our LPSC+ framework: 1) LPSC+Adv combines our LPSC with training-phase adversarial training, which is elaborated in Sect. 5. 2) LPSC+LabelDP combines our offline phase LPSC with training phase LabelDP [10]. It first applies gradient boosting-based LPSC to compute re-weighted residuals, then uses LabelDP to perturb the gradients. 3) LPSC+Marvell integrates our gradient boosting-based offline LPSC with training-phase Marvell [17]. It modifies the sample weights of data in Marvell.

Metrics. We evaluate our method against baselines regarding utility and privacy. We use the AUC (Area Under ROC curve) metric in our experiments. 1) Utility: To gauge the utility of the federated models, we compute the ROC-AUC of the federated model (FL-AUC) on fully aligned test data. Higher values of FL-AUC indicate superior model utility. 2) Privacy: We evaluate the effectiveness of defense approaches using three label privacy attacks: the Norm attack [17], Spectral attack [21], and Passive Model Completion (PMC) attack [8]. For privacy evaluation, we calculate the average ROC-AUC of the label predictions made by the passive parties, which we refer to as label leakage AUC (LL-AUC). A low LL-AUC value, close to 0.5, signifies strong privacy protection.



Fig. 5. Distributions of passive parties' output logits by fitting the original labels v.s. LPSC-encoded labels on the Criteo dataset. (No adversarial training)

6.2 LPSC Protects Privacy Barely Compromising Utility

We first evaluate the protection quality of our proposed gradient boosting-based LPSC mechanism. Specifically, in the federated training phase, we train passive parties' bottom models to fit the LPSC-encoded labels $p_{lpsc}(i, y)$ and original ground-truth labels $p_{at}(i, y)$, respectively.

Dataset	Target	Privac	Utility \uparrow		
		Norm	Spectral	PMC	FL-AUC
Criteo	Label	0.673	0.689	0.711	0.768
	LPSC	0.523	0.538	0.571	0.766
Avazu	Label	0.620	0.648	0.705	0.749
	LPSC	0.531	0.555	0.577	0.751
MIMIC-III	Label	0.577	0.593	0.611	0.763
	LPSC	0.528	0.535	0.558	0.763
Cardio	Label	0.582	0.618	0.664	0.722
	LPSC	0.517	0.542	0.567	0.724

Table 2. The comparison of privacy and utility of VFL fitting labels p_{gt} v.s. LPSC p_{lpsc} against Norm, Spectral, and PMC attacks. \uparrow means desirable directions. $\beta = 0$.

Table 2 presents the LL-AUC against Norm, Spectral, and PMC attacks and the FL-AUC on four datasets. The results reveal that the LL-AUC of LPSC against three attacks is significantly lower than that of the original labels, indicating that LPSC provides strong label privacy protection. Meanwhile, the FL-AUC of LPSC is comparable to that of the original labels, implying that LPSC barely sacrifices model utility. This confirms that LPSC can effectively protect label privacy barely compromising utility.

It is worth noting that the PMC attack achieves higher LL-AUC values than Norm and Spectral attacks, suggesting that it poses a greater threat. This is because PMC allows adversaries to access labeled samples to learn the attack function, which is a strong prior. Therefore, we use PMC attack for label privacy evaluation in subsequent experiments.

Visualization. Figure 5 visualizes the output logits distributions of four passive parties by training with or without LPSC. The top-side distributions in each sub-figure show that, with LPSC, the logits distributions of the two classes almost overlap and are hard to differentiate. In contrast, without LPSC, the right-side distributions in each sub-figure reveal significant differences in the distributions of the two classes, implying label privacy leakage. Our empirical findings are supported by the theoretical guarantee in Theorem 1, which justifies our observation that it is more challenging to distinguish the output distributions between classes when the bottom models are trained with LPSC-encoded labels.

6.3 Privacy-Utility Trade-Off Comparison

Figure 6 shows the privacy-utility trade-off curves on four datasets. The X-axis indicates the label leakage AUC (LL-AUC), and the Y-axis indicates the AUC of the federated model prediction (FL-AUC). An ideal trade-off should have a large FL-AUC and a small LL-AUC, thus residing in the upper-left corner of Fig. 6. Our LPSC+Adv is the closest to the ideal trade-off on all four datasets. We discuss how offline LPSC and training-phase adversarial training improve the privacy-utility trade-off in the following, respectively.



Fig. 6. Privacy-utility trade-off of different methods against PMC attack on four datasets. Note that LPSC+Adv, LPSC+LabDP and LPSC+Marvl are three variants of our two-phase **LPSC**+ framework. \uparrow means desirable direction.

Impact of LPSC. To explore the effectiveness of LPSC, we compare LPSCenhanced methods (i.e., LPSC+LabelDP, LPSC+Marvell and LPSC+Adv) with their counterparts without LPSC (i.e., LabelDP, Marvell and FE-VFL). As shown in Fig. 6, LPSC significantly improves the privacy-utility trade-off of training-phase perturbation baselines by pushing the top-side of each curve leftwards on each dataset. Without any training-phase perturbation (the top-right end of each curve), LPSC leads to significant LL-AUC decline with negligible FL-AUC decline on each dataset, implying that it protects label privacy barely sacrificing utility. This empirical observation is also justified by the theoretical guarantee in Theorem 1. Therefore, LPSC can be easily integrated with various training-phase perturbation methods for improved privacy-utility trade-off.

Impact of Perturbation Methods on LPSC+. LPSC+Adv incorporates adversarial privacy objectives to enable flexible privacy-utility tradeoff, on the basis of *bounded* privacy enhancement given by LPSC. To investigate the effectiveness of adversarial training, we compare LPSC+Adv with two LPSC+ variants (i.e., LPSC+Marvell and LPSC+LabelDP). As shown in Fig. 6, we can observe that the trade-off curves of LPSC+Adv are closer to the upper-left corner than those of two LPSC-enhanced baselines on each dataset, indicating that LPSC+Adv outperforms them with big margins. This validates the effectiveness and superiority of adversarial training in LPSC+ for privacy-utility trade-off.

6.4 Impact of Gradient Boosting Algorithms on LPSC

Dataset	AUC	AdaBoost	LogitBoost	L_2 -Boost
Criteo	$FL\uparrow$	0.765	0.766	0.760
	$LL \downarrow$	0.584	0.571	0.603
Avazu	$FL\uparrow$	0.752	0.751	0.748
	$LL \downarrow$	0.582	0.577	0.592

Table 3. Comparative results of different gradient boosting algorithms for LPSC.

We compare the impact of different gradient boosting algorithms on LPSC, including AdaBoost [6], LogitBoost [7], and L_2 -Boost [3]. For each boosting algorithm, $p_{lpsc}(i, y)$ is computed following Table 2 in Appendix 2.1. AdaBoost updates the sample-weights $\boldsymbol{w}_i = p_{lpsc}(i)$ based on the classification error of the local model f_{θ} . While, LogitBoost and L_2 -Boost assign residuals $\boldsymbol{r}_i = p_{lpsc}(y|i)$ based on the negative gradient of the log-likelihood loss and the mean-square error loss, respectively. Table 3 shows the privacy-utility trade-off of different gradient boosting algorithms on Criteo and Avazu datasets. We find that Logit-Boost is more effective for LPSC than the others in terms of the privacy-utility trade-off.

7 Conclusion

We study label privacy protection in VFL by formulating an offline Label Privacy Source Coding (LPSC) problem. LPSC protects label privacy barely compromising utility by removing redundant label information from the active party's features. We theoretically prove that LPSC satisfies ϵ -MIP and can be reduced to the gradient boosting's objective. Furthermore, we propose a two-phase LPSC+ framework, exemplified with LPSC+Adv to enable flexible privacy-utility tradeoff by further incorporating *adversarial training* for training-phase perturbation. Experimental results on four datasets demonstrate the efficacy of LPSC and the superiority of LPSC+ framework. Future work includes investigating more effective LPSC techniques and protecting label privacy leakage from gradients. Acknowledgments. This work was supported the National Natural Science Foundation of China (Grant No. 62250710682), the Guangdong Provincial Key Laboratory (Grant No. 2020B121201001), and the SUSTech-Huawei Trustworthy Intelligent Systems Laboratory.

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Error Types in Transformer-Based Paraphrasing Models: A Taxonomy, Paraphrase Annotation Model and Dataset

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Abstract. Developing task-oriented bots requires diverse sets of annotated user utterances to learn mappings between natural language utterances and user intents. Automated paraphrase generation offers a costeffective and scalable approach for generating varied training samples by creating different versions of the same utterance. However, existing sequence-to-sequence models used in automated paraphrasing often suffer from errors, such as repetition and grammar. Identifying these errors, particularly in *transformer* architectures, has become a challenge. In this paper, we propose a taxonomy of errors encountered in *transformer*based paraphrase generation models based on a comprehensive error analysis of *transformer*-generated paraphrases. Leveraging this taxonomy, we introduced the Transformer-based Paraphrasing Model Errors dataset, consisting of 5880 annotated paraphrases labeled with error types and explanations. Additionally, we developed a novel multilabel paraphrase annotation model by fine-tuning a BERT model for error annotation task. Evaluation against human annotations demonstrates significant agreement, with the model showing robust performance in predicting error labels, even for unseen paraphrases.

Keywords: Paraphrasing \cdot Transformers \cdot Annotation \cdot Taxonomy

1 Introduction

Dialogue systems (DS), such as virtual assistants, and task-oriented bots, are emerging as a new frontier of human-computer interaction in natural language, receiving considerable recent attention [43]. These services communicate with users in natural language (e.g. text, speech, or both), performing a wide range of tasks such as reporting the weather, booking flights, or booking restaurants [45]. To satisfy user requests, a DS requires a large set of utterances paired with their corresponding executable forms (e.g. API calls). In particular, *task-oriented* bots must first identify the user's intent from a given utterance [43]. For example, in "List of restaurants serving Lebanese food in Lyon", the bot must recognize the intent (i.e. find_restaurant) and the associated slots (<u>location=</u>"Lyon", <u>cuisine=</u>"Lebanese"). Due to the powerful expressiveness of human language, the same intent can be formulated differently, e.g. "Which restaurants in Lyon serve Lebanese food?". Thus it is essential for bots to grasp the richness of human language, by training them on a linguistically diverse set of utterances for each intent [24, 45]. Failing to handle these variations in natural language can negatively impact the effectiveness of bots, and ultimately the user experience.

Paraphrasing is a key technique to build large and diverse utterances for the intents of interest [43]. Paraphrasing is an NLP task that aims to reformulate a given natural language utterance into its lexical and syntactical variations while meaning is preserved [8, 43]. It has numerous applications in NLP tasks, such as sentence simplification, text summarization, and Natural Language generation [8,43]. Paraphrasing methods can be categorized into crowdsourced and automated approaches [25, 45]. Crowdsourced paraphrasing involves human workers generating multiple paraphrases based on a seed utterance [45]. In automatic paraphrasing (AP), paraphrases are systematically generated [24,49]. The literature on AP explored template-based, rule-based, and statistical machine translation approaches [25,26]. Recent attention has shifted to neural network models [26, 31], particularly the *transformer* architecture [41], acknowledged for its state-of-the-art performance in various NLP tasks and widely adopted as the preferred sequence-to-sequence architecture for paraphrasing [7]. However, despite their success, seq2seq models frequently introduce errors such as repetition, grammatical inaccuracies, and incoherent text [39]. Ongoing efforts have concentrated on detecting and identifying paraphrasing errors in automatic neural models to enhance their robustness [35]. Meanwhile, the increasing complexity of *transformer*-based models complicates error identification, making it harder to distinguish between machine- and human-generated text [1, 11]. As these models evolve, the human ability to manually discern and tag machine-paraphrased text diminishes, especially with holistic alterations in sentence structure and word order instead of single-word replacements. Recognizing the pivotal role of errors as indicators for system improvement [40], the evaluation of errors in generated paraphrases becomes paramount.

This study focuses on quality control for paraphrasing, particularly the evaluation of paraphrase errors in *transformer*-based paraphrase generation (TPG) models. While quality control for text generated by NLG systems has been explored in a wide range of tasks, like machine translation (MT) [13], question generation [37], and open-ended generation with pre-trained language models [11], TPG has not been subjected to such scrutiny. To the best of our knowledge, this study is the first to categorize paraphrasing errors in TPG models. Note that our identified errors may not be exhaustive but rather serve as an initial pool of errors for further study and investigation. Efficient error identification and categorization in AP models yield multifaceted advantages, mainly contributing to the elevation of paraphrased content quality. A comprehensive grasp of prevalent errors empowers researchers and developers to strategically augment the performance and reliability of their paraphrasing models. Also, categorizing errors not only sheds light onto the strengths and weaknesses of paraphrasing models, but also establishes strong benchmarks to facilitate fair comparisons between different models. Moreover, our proposed error categories help in enriching training datasets. This, in turn, allows models to navigate and handle common errors, making them more robust and production-ready for real-world applications. The contributions of this study can be summarized as follows:

- 1. We selected five *transformer*-based paraphrasing models to generate 22K paraphrases for 598 seed utterances extracted from a dataset of crowdsourced queries across two intents.
- 2. We synthesized the literature on paraphrasing quality control in three distinct areas: errors in crowdsourced paraphrasing, inconsistencies in crowdsourced slot annotations, and errors in the generation of pre-trained language models. We used this synthesis as a starting point for building our own taxonomy. We then extended this taxonomy through several rounds of qualitative evaluations of the generated paraphrases. Consequently, we identified a taxonomy of 15 error types in *TPG* models.
- 3. We used the proposed error taxonomy to annotate the generated paraphrases. Accordingly, we constructed an annotated dataset called *TPME*, in which the paraphrases were labeled with a range of different categorized errors.
- 4. We developed a multi-label paraphrase annotation model using the TPME dataset. The annotation model uses a fine-tuned BERT model to predict error types in paraphrases, enabling the automatic annotation of multiple errors in a paraphrase.
- 5. We released TPME dataset, code for generating paraphrases, fine-tuned BERT model, and information required to reproduce our study¹.

2 Related Work

Characterizing error has been done in many areas, such as MT [13], crowdsourced paraphrasing [23,45], NLG systems [11,39]. Overall, research from these efforts is certainly complementary and some elements are indeed adopted in our work.

Paraphrase Generation (PG). Crowdsourced **PG** have been investigated to obtain training datasets for *DS* [21,28,33,36,43,44]. In *crowdsourced* **PG**, an initial utterance, usually provided by an expert, is presented as a starting point, and crowdworkers are then recruited to obtain further paraphrases [43]. For instance, Chklovski et al. [9] used crowdsourcing to collect paraphrases using gamification. Contributors were asked to generate paraphrases based on given hints (e.g. words suggestions). Other crowdsourcing strategies were proposed [46,47]: (i) *Sentence-based strategy*: Workers were tasked with paraphrasing a given sentence into new

¹ https://github.com/AudayBerro/TPME/tree/master.

variations. (ii) Goal-based strategy: Workers were provided with a task goal (e.g., "book a restaurant") and a set of possible entity values (e.g., "cuisine: Indian, city: Paris") to produce paraphrases. (iii) Scenario-based strategy: This approach employs a storytelling framework that provides a scenario to workers and ask them to generate paraphrases accordingly (e.g., "Your goal is to book a restaurant; you are in Paris; you are hungry and want to eat Indian dishes").

Automated **PG** does not involve humans in the process and refers to a task in which a system generates paraphrases given an input sentence [24]. The literature on *automated* **PG** covers a wide range of approaches, including probabilistic, handwritten rules, and formal grammar models [15]; data-driven techniques [25,27]; machine translation techniques [18]. However, these approaches struggle to capture the nuanced complexities of natural languages in contextual settings [14]. In addition, the manual design of rules is complex for practical implementation [48]. Consequently, neural-based and deep learning models have gained popularity for \mathbf{PG} [31,49], offering a solution free of previous limitations. However, a critical problem persists: generated paraphrases often fail to align with user preferences and produce uncontrolled results [48]. Although syntactically controlled paraphrasing \mathbf{PG} [16,20] offers a promising approach that incorporates syntactic templates, it requires users to possess linguistic expertise and define specific syntactic structures, which is challenging in practical applications. With the recent advances enabled by large language models (e.g. GPT), there is a shift towards their use to generate paraphrases [5, 17, 42].

Errors in Crowdsourced Paraphrasing. Crowdsourced paraphrases often contain errors, including misspellings, grammatical mistakes, and missing slot values [45]. Two approaches are commonly used to evaluate paraphrase quality [29,45]. In *Pre-hoc*, paraphrases are evaluated during the crowdsourcing task before submission [29]. In *Post-hoc*, they are assessed after task completion [45]. Yaghoubzadeh et al. [45] employed a *post-hoc* method to investigate crowd-sourced paraphrasing errors in task-oriented bots. They identified a taxonomy of six error types (misspelling, linguistic, cheating, answering, semantic, translation) and they developed the *Para-Quality* dataset based on these findings. Similarly, Larson et al. [23] identified different types of incorrect annotations of crowdsourced paraphrases. They identified a taxonomy of six types of inconsistencies in slot-filling annotations (e.g. slot format, omission, wrong label, slot addition).

Errors in MT Systems. Significant work on errors has been reported for MT systems. Koponen et al. [22] investigated error classification with an emphasis on semantic accuracy. The error analysis was performed on human translations as well as on the outputs of 2 different types of MT systems: *rule-based* MT and *statistical* MT. They identified 13 errors grouped into 2 categories: concept (5) and relation (8) errors. Popovic et al. [30] investigated the nature and causes of MT errors observed by different evaluators on different quality criteria: adequacy,

comprehension, and fluency. They identified 26 errors (e.g. omission, gender) and reported the results for 3 language pairs, 2 domains, and 11 MT systems.

3 Paraphrase Generation

Paraphrase generation poses unique challenges compared with other text generation tasks because of the requirement to produce sentences that convey the same meaning with different words or structures. This requires creativity and linguistic versatility. In addition, paraphrasing models must retain the mentions of intents and their respective slots, adding to the complexity. Sparse data collection for paraphrases further compounded this challenge, limiting exposure to diverse scenarios. Despite advancements, even transformer-based models exhibit errors, such as incorrect substitutions, missing words, awkward structures, or alterations in meaning. To address this, we developed a taxonomy to systematically categorize these errors, resulting in a TPME dataset with manually labeled paraphrases and errors. Furthermore, we fine-tuned a BERT-based model for automated error-detection. To collect paraphrases, we followed a methodical approach: (i) obtain seed utterances, (ii) select paraphrasing models, and (iii) generate paraphrases using these models. Each step is described in detail below:

3.1 Selection of Seed Utterances

In this study, we employ the SNIPS dataset², consisting of crowdsourced queries categorized into seven user intents. Each utterance in the dataset is paired with a list of required slots, which are specific pieces of information or textual parameters within an utterance that need to be identified and extracted. For each SNIPS excerpt, we extract the utterance (e.g. "how cold is it in Princeton Junction") and its list of required slots (e.g. <u>condition_temperature="cold"</u> and <u>city="Princeton Junction"</u>). To manage the manual labeling effort, we focused on two key intents: GetWeather and BookRestaurant, which enabled us to collect 598 seed utterances. GetWeather encompasses requests for weather forecasts comprising 9 slots (e.g. country, city, temperature). BookRestaurant includes queries relating to restaurant reservations, with 14 slots (e.g. city, time, dishes served).

3.2 Selection of Models

We used the following criteria to select the paraphrasing models used in this study: (i) models must fall under the category of text generation and can produce paraphrases in English; (ii) they should be built upon the transformer architecture [41] in any of its variations, such as decoder-only or encoder-decoder; (iii) The official checkpoints of the models must be publicly and freely accessible through platforms or web links provided by their authors. This ensures the avoidance of potential biases that might arise if we were to implement, train, or fine-tune the models. In this study, we chose the following five (TPG) models:

 $^{^{2}}$ https://github.com/sonos/nlu-benchmark.

PROTAUGMENT [10]: fine-tuned a BART pre-trained transformer-based language model to generate paraphrases.

Fine-tuned T5 [3]: the authors fine-tuned T5 [32], a pre-trained transformer-based language model to generate paraphrases.

NL_Augmenter³: is a data-augmentation platform that supports various transformations. We selected the NL_Augmenter *Diverse Paraphrase Generation* transformation for this study, which generates paraphrases by leveraging a transformer through pivot-translation [2].

PRISM: Although PRISM [38] is a quality estimation model designed to evaluate the performance of MT systems, it includes an automatic paraphrase generation component. The authors trained a transformer-based MT model with approximately 745 million parameters to perform zero-shot paraphrasing in 39 languages. PRISM can be used in paraphrases generation.

GPT [4]: GPT is a generative transformer-based language model with outstanding performance. Recent GPT models can adapt to new, possibly unseen, tasks using In-context Learning through natural language instructions and input. This opens up the possibility of improving the paraphrasing process [6]. We leveraged GPT-3.5-turbo with tailored prompts for paraphrase generation⁴.

3.3 Generation of Paraphrases from Utterances

Leveraging the aforementioned models, we systematically generated paraphrases for each seed utterance, yielding a comprehensive collection of 22864 paraphrases. We categorize the generated paraphrases in ten distinct datasets, based on the paraphrasing model used for their generation, and the underlying intent of the seed utterances. For instance, the T5-GW dataset (accessible via the provided additional materials link) consolidates all paraphrases generated by the T5 model in response to seed utterances associated with the *GetWeather* intent.

Each grouped dataset includes the following columns: utterance, paraphrase, $list_of_slots$, and intent. The utterance column corresponds to the seed sentence designated for paraphrasing, while the $list_of_slots$ column encompasses any identified slots in the utterance, along with their corresponding values. The paraphrase column houses the resulting paraphrase. Finally, the intent column indicates the intent of the utterance, with only two possible values: GetWeather or BookRestaurant in this work.

4 Paraphrasing Error Types

We identified 15 error types in the paraphrases categorized into three error classes through a meticulous evaluation process conducted over multiple rounds. Initially, we randomly selected ten utterances with their corresponding paraphrases, totaling 700 paraphrases for evaluation. We labeled using error types identified

³ https://github.com/GEM-benchmark/NL-Augmenter.

⁴ The prompt we used can be found in the supplementary material link supplied.

Table 1.	Samples	of erroneous	ly generated	l paraphrases.	The	colors	in the	Paraphrophic Par	ase
column d	lenote the	locations of	errors.						

11	Uttorango	Paraphrasa	Free Label
#	t : 192 1		a w
1	is it chiller here	is it Colder Here ?	Spelling
2	book a party of 4 for a cafe rated best	booking a 4 party for a best rated cafe	Spelling
3	What is the weather forecast for Manassa?	What's the weather prognosis for Manassa?	Spelling
4	is it cold in Vigus TN	if it's cold in the Vigus TN	Grammar
5	What's the weather in Poland?	What is the time in Poland?	Semantic
6	Book spot in Fults in Federated States Of Micronesia	Book spot at Fults in the Federated Federal States of Micronesia.	Redundant
7	Book me a lunch at Boreas for a party of 8	Book me a lunch at Boreas for a party of 8!	Duplication
8	book a table in Yachats for 3 people	Procure is ervation sayrettel batou hcastuohcaeY nA	Incoherent
		${\it selaecinimany Th girete potsereal trafformet a B}$	
9	Will it be cold in Haiti?	Will There Be Cold in Haiti ?.	Punctuation
10	Tell me the weather forecast for <u>France</u>	Tell me the weather forecast for <u>France</u> for the weekend	Slot addition
11	Book a spot for <u>now</u> in <u>NY</u> for <u>10</u>	Book a place in \underline{NY} for $\underline{10}$	Slot omission
12	What is the weather forecast <u>here</u>	What is the forecast for the weather in the future ?	Wrong Slot
13	Inform me, should the location known as 'Maxwell'	Wordy	
		suffer from an impediment such as haze/mist/ fog	
		I'd really appreciate knowing about such trivia	
14	Will there be wind in <u>Hornitos</u> DC?	Will there be wind in Hornito's DC?	Homonym
15	Will it be temperate in West Liberty MH?	Will it'll be temperate in West Freedom MH?	Homonym
16	Will the weather be good at breakfast in Bay <u>Harbor</u> Islands	Is the weather good for breakfast in the Bay Puerto Islands?	Homonym
17	is Saint Robert hotter than Turkmenistan?	Saint Robert is hotter than Turkmenistan.	Answering
18	book spot for 4 in Somalia	Book spot for 4 in Somalia is available for free .	Answering
19	Is it hot in the current location?	Is it hot in the current location? If so, why?	Questioning
20	book a table in <u>CA</u> for 2 people in 3 hours	booking a table in Central Asia for 2 people in 3 hours	Acronym

by previous research in crowdsourced paraphrasing [45], crowdsourced slot annotation inconsistencies [23], and errors in text generated by pre-trained language models [11]. The initial list included *semantic*, *redundant*, *spelling*, *grammar*, *slot addition*, *incoherent*, and *duplication* errors. Through manual annotation and iterative refinement, involving random sampling and evaluation of paraphrases based on the evolving list of error types, similar errors were grouped into new types, resulting in the final taxonomy. We also refined error definitions, such as introducing the concept of "*near-copy*" (refer to § 4.1) for *duplication* errors. In the following, we proceed to describe our three error classes, highlighting the specific error types within each class.

4.1 Language Errors

Language Errors encompass a range of inaccuracies in paraphrased content, including spelling, grammar, syntax, and semantic inconsistencies. Seven types of *language errors* were identified.

Spelling refers to the correct arrangement of letters to form a word. Mis-

spelling is one of the most common mistakes in crowdsourced paraphrasing [45]. In our evaluation, we also count as misspelling capitalization errors (sample 1 in Table 1), missing hyphens (sample 2) and missing spaces (sample 3).

Grammar These errors relate to the incorrect use of verbs, prepositions, singular/plural nouns, articles, and other grammatical elements [45]. In sample 4, the paraphrase exhibits a misuse of the article "*the*" before city names.

Semantic This error, further characterized as a "semantic deviation", arises when a paraphrase deviates from the intended meaning of a seed utterance. For instance, in sample 5, the paraphrase asks for time instead of weather conditions.

Redundant Redundancy arises when a word or phrase is duplicated in a paraphrase, either through exact repetition or the use of different words conveying the same context. In sample 6, the term *Federal* redundantly duplicates the meaning conveyed by the term *Federated*.

Duplication Duplication arises when the generated paraphrase either mirrors or closely resembles the utterance. We employ the term "*near-copy*" to describe instances where the paraphrase closely mirrors the utterance. "*Near-copy*" occurs when a paraphrase differs from the utterance solely in terms of punctuation (e.g. (e.g., commas, periods, question marks, colons, etc.) and capitalization. However, the "*near-copy*" condition is violated if the paraphrase contains at least one token that differs from the utterance. This is illustrated in sample 7.

Incoherent As in sample 8, we label a paraphrase as incoherent when the generated text is confusing, hard to understand, or appears nonsensical.

Punctuation A punctuation error occurs due to the overuse or inappropri-

ate placement of punctuation marks in the paraphrase. This includes inserting question or exclamation marks in sentences without corresponding questions or exclamations. It also encompasses the misuse of currency, non-alphabetic or numeric symbols (, _, #, &, etc.). For instance, sample 9 displays a punctuation error where a period follows a question mark incorrectly.

4.2 Slot Errors

These errors involve incorrect actions at the slot level, such as adding, removing, or altering slots.

Slot addition Slot addition occurs when the model inserts at least one additional slot value into a paraphrase. Consider sample 10 in Table 1. In this sample, token "the weekend" which is the value of the timeRange slot in the paraphrase, is an additional slot. It's important to note that for slots that accept multiple values, this is not considered an error. For example, the party_size_description slot in the *BookRestaurant* intent can have multiple values. In the utterance "*Book a table for Ali, Jo, and Max*" the tokens "Ali", "Jo" and "Max" form a single multi-token value, and the party_size_description slot treats them collectively as a single value.

Slot Omission Slot omission occurs when a slot, expected to be referenced, is overlooked in the paraphrase. Illustrated in sample 11 of Table 1, the paraphrase fails to include a value for the <u>timeRange</u> slot, even though it is explicitly mentioned as "now" in the original utterance.

Wrong Slot A wrong slot occurs when the value of a slot in the paraphrase deviates from the expected slot and is replaced by an non-matching token. In sample 12, rather than inquiring about the forecast for the present location, the paraphrase erroneously requests a weather forecast for a specific time period.

4.3 Errors of Human Characteristics

We identified 5 types of *human-characteristic errors*, which uniquely mimic human behavior. When these errors occur, the *transformer* behaves as if it were human, such as responding directly to a request instead of paraphrasing it.

Wordy Wordy errors occur when the generated text contains excessive wording or unnecessary information, leading to verbose paraphrasing. See sample 13.

Homonym Homonyms are words that share the same pronunciation but have different meanings or spellings. An error arises when a token in the paraphrase shares a similar or identical pronunciation with a token in the utterance. In sample 14, "Hornito's" and "Hornitos" sound alike, leading to incorrect use of the possessive apostrophe ("'s") in the paraphrase. This category includes cases which tokens are replaced with synonyms or translated, potentially altering the paraphrase's meaning (see samples 15 and 16 resp.).

Answering This error occurs when the paraphrased content responds to the utterance, causing the model to generate an answer instead of a paraphrase. In sample 17, the model transforms the entire paraphrase into an answer, while in sample 18, the answer is added to the query. To differentiate this from the Wordy Error, we label sentences as Answering Error if the additional tokens answer a query or question from the initial utterance.

Questioning Arises when the paraphrased text introduces an extra question not present in the utterance. In sample 19, the addition of "*If so, why?*" exemplifies this error by introducing an extra question.

Acronym An acronym is a word or name formed from the initial letters of a longer phrase. Acronym error occurs when a paraphrase improperly uses an acronym or includes an incorrect expansion of an acronym from the utterance. In sample 20, "Central Asia" is an inaccurate expansion of the acronym "**CA**" which actually represents California, a value for the "state" slot.

5 Creation of Annotated Paraphrasing Error Dataset

This section presents an overview of TPME, and gives insights and analyses of the paraphrasing errors.

5.1 The TPME Dataset

We annotated a representative subset of generated paraphrases, covering at least 22% of the entire set, resulting in 5880 annotated paraphrases. Each paraphrase was labeled with one or more error types from our taxonomy (as detailed in § 4) and was accompanied by an explanation in plain English. For "slot errors" (see § 4.2), we compared each sampled paraphrase with its corresponding list of required slots listed in the "list_of_slots" column. The *TPME* dataset includes the following columns: "utterance" column which contains the seed sentence to be

paraphrased. "*list_of_slots*" includes any slot present in the utterance along with its corresponding value. "*paraphrase*" contains a generated paraphrase. "*models*" denotes the model that generated the paraphrase. "*error_category*" contains labels of the errors found in the paraphrase, and their justification in natural language is described in the column "*explanation*". "*Intent*" indicates the intention conveyed by the utterance.



Fig. 1. TPME dataset label statistics.

5.2 Insights into Error Frequency and Co-occurrences

Figure 1 visually presents the distribution, frequencies and co-occurences of labels within the TPME dataset, employing an UpSet plot through Intervene platform⁵. Notably, only 40.1% of the paraphrases were labeled as *correct*, underscoring the prevalence of paraphrasing errors, and their negative impact in the context of developing *DS*. Specifically, only 3.5% were exclusively labeled with *semantic* errors, without any additional labels. In addition, 6.8% of the paraphrases were identified with *grammar* errors only. The plot also demonstrates the frequency of the co-occurrence of two or more labels. For instance, all paraphrases labeled as *slot omission* (29 occurrences) are also labeled as *spelling*. Additionally, 34 paraphrases shared both *semantic* and *spelling* error labels, constituting 0.9% of the erroneous paraphrases. Furthermore, 18 paraphrases were labeled with *homonym*, *spelling*, and *grammar* errors. Moreover, 144 paraphrases were concurrently labeled with *duplication* and *grammar*.

⁵ https://upset.app/ and https://asntech.shinyapps.io/intervene/.

To analyze the distribution of errors across the selected models, individual Upset Plots were generated for each of the five models. The initial observation highlights that out of 1092 instances, GPT yielded 659 correct paraphrases (60.3%). In contrast, **T5** achieved a correctness rate of only 25.1%, and **PRO-TAUGMENT** demonstrated an even lower rate of 17.64%. The second significant finding is that none of the paraphrases generated by GPT in TPME were labeled as *duplication*, distinguishing it from other models. For example, **PROTAUGMENT** had a *duplication* error rate of 51.04%, and **T5** exhibited a rate of 21.5% for these errors. In summary, GPT displays resilience against duplication compared with other *TPG* models. However, duplication may occur among the generated paraphrases. For example, in a list of 10 paraphrases, we may have three paraphrases that are replicated.

5.3 Analysis of the Annotated Paraphrases

The Relevance of Capitalization in Paraphrases. Understanding user utterances relies on entity extraction, known as slot-filling, which aims to identify the values of different slots in a user utterance [19]. For instance, when a user requests *nearby restaurants* the values of the *location* and *cuisine* slots are essential for a bot to retrieve the appropriate information. We observed challenges with capitalization in slot-filling tasks, particularly when using casesensitive slot-filling models. Consider the utterance "book smoking room in OR at a bar⁶ and its paraphrase "Book Smoking Room In OR at bar OR at hotel". In the paraphrase, the second "<u>OR</u>" token serves as a conjunction indicating a choice between a *bar* or a *hotel* but it is erroneously written in uppercase. This capitalization introduces a spelling error and leads to a redundant error as the paraphrase already includes the token "OR". Additionally, the capitalization issue may result in a slot addition error, where the model misinterprets "OR" as an abbreviation for the state of "Oregon", introducing ambiguity. To address these issues, we propose accurately representing capitalization in paraphrases to reflect both the input and output of the transformer architecture.

The Propagation of the Source Utterance Errors in the Paraphrase. Transformer-based language models are highly effective in learning language properties [12], yet instances of errors persisting in generated paraphrases have been identified. For example, in the utterance "book spot at Candle Cafe" and its paraphrase "Book spot at Candle Cafe", the transformer omitted the indefinite article "a" before "spot", leading to a grammar error. We attribute such errors to the inherent nature of transformer architecture and the paraphrase generation task⁷. Transformers rely on the self-attention mechanism, meaning errors in the input may receive more attention and propagate through subsequent layers.

⁶ OR refers to the state of Oregon.

⁷ Paraphrase generation is a multi-step (word-by-word) prediction task, where a small error at an early time-step may lead to poor predictions for the rest of the sentence, as the error is compounded over the next token predictions [8].

However, transformers also demonstrate the ability to rectify errors in the input during paraphrase generation. For instance, the input utterance "is it going ot be chillier in Maumee" contains "ot" instead of "to", but the model corrected this error in the paraphrase, resulting in "is it going to be colder in Maumee?".

The Insertion of the Determiner "the" in Front of Geographical Names.

In some paraphrases, we encountered errors that violated basic grammar rules. For instance, consider the utterance "weather in Hillsview MA" and its paraphrase "Weather in <u>the</u> Hillsview MA". The model mistakenly inserted the determiner <u>the</u> before the token "Hillsview", resulting in a grammar error. Notably, when dealing with geographical names such as the city name "Hillsview", the definite article "the" is not used in English, making this a common grammatical inconsistency that we observed in this study. Another prevalent error involves the insertion of the possessive form ('s) in phrases like "Will there be wind in Hornito's" instead of the correct "Hornitos" as found in the utterance "Will there be wind in Hornitos DC?".

Errors May be Context- and Domain-Dependent. For the utterance "book a table for me, heidi and cara in Saudi Arabia" and its paraphrase "Book a table for me, Heidi and Vedi in Saudi Arabia", in the paraphrase the token "Vedi" is an appropriate value for the party size description slot. The transformer paid more attention to the previous "Heidi" token, which resulted in the generation of the "Vedi" token in the paraphrase. However if we pay more attention, the two names "*Heidi*" and "*Vedi*" have close pronunciation which leads to a homonym error. This error, more than a minor pronunciation anomaly, can significantly impact the performance of the DS trained with such paraphrases. Consider a bot in the banking sector. Executing a money transfer to "Heidi and Vedi" instead of "Heidi and Cara", as stated in the utterance, would be incorrect. While this variation enhances lexical diversity, it can also negatively impact critical domains. In addition, homonym errors may lead to wrong slot errors. In the paraphrase "Will there be wind in Hornito's DC?" the token "Hornito's" is a homonym of the token "Hornitos" in the utterance. The addition of the possessive "s" introduces a homonym error, resulting in the city slot having the value "Hornito's", which does not match the correct "Hornitos" for the city slot. Thus, a wrong slot error emerges. Consequently, the tolerance for such errors, varies based on the context, domain, and associated slots when the error affects the slot value level.

Errors in GPT Generated Paraphrases. For the utterance "Will it be windy at 4 Pm in <u>NY</u>?", GPT generated the paraphrase "Are we expecting any strong winds by 4 PM in <u>New York City</u>?". GPT incorrectly generated the value "*New York City*" which represents a value for the "*city*" slot instead of the intended "*state*" slot, mentioned as "*NY*" in the utterance. Similarly, for the utterance "book a turkish restaurant in <u>DE</u>", GPT generated the paraphrase

"Reserve a Turkish restaurant in Germany", incorrectly inserting "Germany" instead of the correct value "Delaware" (abbreviated as "DE" in the utterance), where GPT considered "DE" to be the acronym for "Deutschland". However, this introduces a wrong slot error, as the correct slot is "state", not "country". Across all GPT-generated paraphrases, 10.3% were labeled as wrong slot errors. For "weather close-by Lone Elk County Park at 4 am", GPT actually answered the weather-forecast query: "By dawn, you can anticipate interesting conditions nearby at Lone Elk county park". The answering error occurred in 5.1% of GPTgenerated paraphrases. Additionally, 9.4% of GPT paraphrases were marked with semantic errors. For example, in "What will the weather be at six o'clock in the Virgin Islands?" GPT generated only a question mark "?" as a paraphrase. For "book a table in Yachats for 3 people" GPT generated extensive gibberish tokens with numerous misspelled words, such as "iservation" which should be "a reservation". The paraphrase is incoherent, making it hard to understand. Also in "Ensure there is seating available in Yachtseservt veNameYeepsaYehTfruintap eneea Aredton OS segnos repedral hseltingeobs uo coto hibm at eletibasen a Yhctayssesacllaeviser" GPT generated an arbitrary content.

For the utterance "I want to book a highly rated restaurant for Sue, Madeline, and me in eight years", GPT generated " \leq introductory statement> I desire very much that we find \leq quality adjective>restaurant well-known far&wide ...for our select groupIO" and "Imagine the celebration of such a beautiful day \leq including excitement) \leq Pronoun> has made reservations at the highest-ranked restaurants". Apart from incoherent, spelling, and grammar errors, we observed the insertion of tagged tokens like " \leq Pronoun>" and " \leq introductory statement>". Instead of generating slots values, GPT generated canonical tokens to indicate the need to insert values at those positions.



Fig. 2. Frequency Analysis of Correct Labels in 130 GPT Paraphrases across Positions.

Another GPT-specific finding is the removal of certain paraphrases due to toxic content, aligning with OpenAI's policy. For instance, when prompted with "need a table at a close-by restaurant right now in Marco" GPT generated "**Paraphrase removed due to inappropriate content**" illustrating paraphrase suppression due to toxicity. GPT produced three times the same paraphrase
for this utterance. Because the model is exclusively accessible via API calls with undisclosed details, pinpointing the exact content that triggers filtering is impractical. Given its exclusivity to GPT, we opted not to categorize it as a distinct error in our taxonomy but rather to highlight its occurrence. Furthermore, we evaluated the frequency of "correct" labels in GPT-generated paraphrases for each seed utterance. We randomly selected 13 utterances and extracted their corresponding sets of 10 paraphrases, totaling 130 paraphrases. The key observation is that errors consistently appear in the paraphrases for each seed utterance, errors became more prevalent, with the majority occurring after the sixth position. Figure 2 illustrates this trend, emphasizing the concentration of errors towards the later positions in the paraphrase lists.

6 BERT-Based Multi-label Paraphrase Annotation Model

In this section, we explore using the TPME dataset to fine-tune a BERT model for multi-label paraphrase annotation. While deep learning models like BERT have shown impressive performance across various NLP tasks, including sentence classification [34], annotating paraphrase errors with their respective error types requires a significant amount of labeled data, posing a challenge. Thus, we finetuned BERT on TPME to develop a multi-label paraphrase error annotation model aimed at predicting error types in paraphrases.

6.1 BERT Fine-Tuning

The fine-tuned BERT model (FBM) is a multi-label prediction model that takes as input a pair of an input utterance u and its paraphrase p and predicts one or more error labels. At the fine-tuning time, u and its p are provided as inputs to the BERT model and tokenized through the BERT tokenizer into one sequence (BERT input: $\langle u \rangle \langle sep \rangle \langle p \rangle$). The " $\langle sep \rangle$ " token acts as a separator between u and p. When the input text is tokenized, BERT interprets the segments as distinct parts of the input sequence. " $\langle sep \rangle$ " helps BERT to understand the structure of the input and learn contextualized representations for each segment. For fine-tuning, the TPME dataset was split into training $(80\% \equiv 4704 \text{ paraphrases})$ and validation $(20\% \equiv 1176 \text{ paraphrases})$ datasets. We fine-tuned a *bert-base-uncased* model. BERT logits (i.e. output) have the form (batch size, number of labels) and represent the non-normalized scoring for each label. To convert these logits into predicted labels, we added a linear layer on top of BERT. Thus, applying a sigmoid function to each logit independently scales the values between 0 and 1, treating them as "probabilities" for label presence. These probabilities are then classified using a standard threshold, usually set at 0.5. If the probability exceeds the threshold, the label is predicted for p; otherwise, it is not predicted.

Table 2. Prediction performance of FBM in terms of Krippendorff's alpha (values range from 0 to 1, where 0 is perfect disagreement and 1 is perfect agreement), Exact Match Ratio, Hamming loss (smaller the value, better the performance). F1, precision and recall are samples-averaged.

	Krippendorff	Exact Match Ratio	Hamming Loss	Recall	Precision	F1
FBMvsVal	0.549	0.693	0.035	0.753	0.766	0.753
FBMvsGold	0.809	0.699	0.027	0.807	0.754	0.770

6.2 Evaluation

This section presents the experimental results of the annotation of two paraphrase datasets using the FBM. First, FBM was applied to a benchmark dataset, called FBMvsGold, comprising a 20% subset of the TPME dataset, including utterances, corresponding paraphrases, and error labels. FBM predicts error labels for each utterance-paraphrase pair to assess the annotation quality of familiar data seen during the fine-tuning stage. Second, FBM annotation was evaluated on dataset \mathcal{B} , comprising 1000 pairs of utterances and paraphrases randomly selected from the 22K unannotated paraphrases. Dataset \mathcal{B} is defined as $\mathcal{B} = \{p \mid p \in (\mathcal{P} - TPME)\}$, where \mathcal{P} denotes the dataset of 22k automatically generated paraphrases and TPME denotes the dataset of 5880 annotated paraphrases. This evaluation assesses FBM annotation on unseen data. After predicting the error labels for Dataset \mathcal{B} , 100 rows were randomly selected for manual annotation, resulting in the FBMvsVal dataset. FBMvsVal enables the assessment of FBM annotation on unseen data. Finally, FBM annotation was evaluated against human annotation in FBMvsGold and FBMvsVal using established multilabel evaluation metrics from the literature, including Krippendorff's alpha, Exact Match Ratio, Hamming Loss, recall, precision, and F1 metrics.

Analysis of Results: The Krippendorff metric revealed strong agreement between FBM and human annotations in both FBMvsGold and FBMvs-Val datasets, with Krippendorff's alpha scores of 54% and 80%, respectively (Table 2). Scores exceeding 50% indicated good agreement, suggesting a robust correlation between the model predictions and human annotation. However, there was a notable disparity between the datasets, with FBMvsVal scoring 54% and FBMvsGold scoring 80%. This variation may stem from the uneven distribution of errors in the TPME dataset used to fine-tune BERT. For instance, the incoherent label is applied to only 19 items, compared to 241 and 844 items labeled as grammar and duplication respectively. Consequently, accurately predicting errors becomes more challenging. Future work will involve augmenting the TPME dataset by annotating more paraphrases while ensuring a balanced representation across error types. Regarding Hamming Loss, which indicates misclassification frequency, both FBMvsVal and FBMvsGold datasets exhibited remarkable performance, with scores of 3.5% and 2.7%, respectively. (Further details on additional measures are omitted due to space constraints.)

7 Conclusion and Future Work

In this study, we used a data-driven approach to investigate and quantitatively identify errors in TPG models. Identifying the nature and frequency of these errors is important for enhancing DS and improving TPG models performance. We first discussed and outlined the importance of paraphrasing in the acquisition of training data for DS development. Subsequently, we emphasized the importance of error evaluation in the paraphrases generated by transformer-based models. Through empirical analysis, we identified a taxonomy of 15 error types, which we used to annotate a paraphrasing dataset with associated errors. Our analysis revealed that despite the success of transformers, paraphrase generation remains error-prone, with only 40.1% of paraphrases being correct. Finally, we released the dataset of paraphrases and errors to the research community.

In our future work, we plan to expand the TPME dataset by annotating additional paraphrases to achieve balance across different error labels. Moreover, we aim to enhance the proposed error taxonomy by exploring various transformer architecture variants, including encoder-only, decoder-only, and encoder-decoder models. The TPME dataset can serves as a valuable training data for diverse tasks. In addition to error detection, we also plan to investigate to error correction using fine-tuning language models. While the TPME dataset is relatively small, its manual annotation process was thorough yet time-consuming and costly. BERT was selected for its effectiveness in multi label error annotation, demonstrating the utility of our dataset. In future research, we will focus on exploring other newer models (e.g., GPT, mistral-7b, LLaMA) through a comparative study, aiming to validate and extend the applicability of our dataset in multi-label error annotation tasks.

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FedHCDR: Federated Cross-Domain Recommendation with Hypergraph Signal Decoupling

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Abstract. In recent years, Cross-Domain Recommendation (CDR) has drawn significant attention, which utilizes user data from multiple domains to enhance the recommendation performance. However, current CDR methods require sharing user data across domains, thereby violating the General Data Protection Regulation (GDPR). Consequently, numerous approaches have been proposed for Federated Cross-Domain Recommendation (FedCDR). Nevertheless, the data heterogeneity across different domains inevitably influences the overall performance of federated learning. In this study, we propose **FedHCDR**, a novel Federated Cross-Domain Recommendation framework with Hypergraph signal decoupling. Specifically, to address the data heterogeneity across domains, we introduce an approach called hypergraph signal decoupling (HSD) to decouple the user features into domain-exclusive and domain-shared features. The approach employs high-pass and lowpass hypergraph filters to decouple domain-exclusive and domain-shared user representations, which are trained by the local-global bi-directional transfer algorithm. In addition, a hypergraph contrastive learning (HCL) module is devised to enhance the learning of domain-shared user relationship information by perturbing the user hypergraph. Extensive experiments conducted on three real-world scenarios demonstrate that Fed-HCDR outperforms existing baselines significantly (Code available at https://github.com/orion-orion/FedHCDR).

Keywords: Federated learning \cdot Recommendation system \cdot Graph neural network

1 Introduction

Cross-domain recommendation (CDR) [1,2] has been widely applied to leverage information on the web by providing personalized information filtering in various real-world applications, including Amazon (an e-commerce platform) and YouTube (an online video platform). CDR can significantly enhance the performance of item recommendations for users by utilizing user rating data from various domains, under the assumption that users have similar preferences across domains. However, with the formulation of the General Data Protection Regulation (GPDR), user-item ratings are not accessible across different domains. How to provide high-quality cross-domain recommendations while satisfying privacy protection has emerged as an urgent issue.

In this paper, we focus on a problem of federated cross-domain recommendation (FedCDR) [8,11]. In this case, user-item rating interactions are considered private information that cannot be directly accessed by other domains. Although existing FedCDR methods [8–11] can effectively solve the privacy issue in CDR, they also face the issue of data heterogeneity across domains, that is, useritem interaction data in different domains contain domain-exclusive information. Figure 1 presents a toy example illustrating the data heterogeneity across different domains. The figure depicts that Bob and Alice interact with various types of items in each domain, including action movies and documentary movies in the Movie domain, professional books and action books in the Book domain,



Fig. 1. Data heterogeneity across domains in the FedCDR scenario.

and action games and simulation games in the Game domain. As previously mentioned, documentary movies in the Movie domain, professional books in the Book domain, and simulation games in the Game domain can all be considered as domain-exclusive interaction information. Existing FedCDR methods [8–11] use direct aggregation of client models or user representations to transfer knowledge, resulting in the mixing of domain-exclusive information into the global model, resulting in poor local performance of the global model (i.e., negative transfer). Therefore, it is necessary to decouple domain-shared and domain-exclusive information, and only aggregate domain-shared information to avoid negative transfer problems.

In response to the issue of data heterogeneity, we introduce a novel **Fed**erated **C**ross-**D**omain **R**ecommendation framework with **H**ypergraph signal decoupling (**FedHCDR**). This framework enables different domains to collectively train better-performing CDR models without the need to share raw user data. Specifically, inspired by the hypergraph structure [3,4] and graph spectral filtering [5–7], we introduce a hypergraph signal decoupling method called HSD to tackle the data heterogeneity across domains. In this approach, the model of each domain is divided into a high-pass hypergraph filter and a low-pass hypergraph filter, responsible for extracting domain-exclusive and domain-shared user representations respectively. Furthermore, we devise a hypergraph contrastive learning module HCL to enhance the learning of domain-shared user relationship information by introducing perturbations to the user hypergraph. The evaluation is conducted on Amazon datasets under the federated learning setting. The experimental results demonstrate that our FedHCDR significantly enhances recommendation performance in three different FedCDR scenarios.

To summarize, our contributions are as follows:

- We propose a novel federated cross-domain framework FedHCDR, designed to enable different domains to train better performing CDR models collaboratively while ensuring data privacy.
- We introduce HSD, a hypergraph signal decoupling method. HSD uses highpass and low-pass hypergraph filters to decouple the user features into domain-exclusive and domain-shared features to address the data heterogeneity issue across domains, which are trained by the local-global bi-directional transfer algorithm.
- We devise a hypergraph contrastive learning module HCL, which perturbs the user hypergraph to learn more effective domain-shared user relationship information.

2 Methodology

2.1 Problem Formulation

Assume there are K local clients and a central server. The k-th client maintains its own user-item interaction data $\mathcal{D}_k = (\mathcal{U}, \mathcal{V}_k, \mathcal{E}_k)$, which forms a distinct domain, where \mathcal{U} denotes the overlapped user set in all domains, \mathcal{V}_k denotes the item set in domain k, and \mathcal{E}_k denotes edge set, i.e., the set of user-item pairs in domain k. Additionally, there is a user-item incidence matrix $\mathbf{A}_k \in \{0, 1\}^{|\mathcal{U}| \times |\mathcal{V}_k|}$ for domain k, where each element $(\mathbf{A}_k)_{ij}$ describes whether user $u_i \in \mathcal{U}$ has interacted with item $v_i \in \mathcal{V}_k$ in the edge set \mathcal{E}_k .



Fig. 2. An overview of FedHCDR. (Color figure online)

For client k, we first construct the user hypergraph adjacency matrix \mathbf{A}_{k}^{u} and the item hypergraph adjacency matrix \mathbf{A}_{k}^{v} according to the user-item incidence matrix \mathbf{A}_{k} . Subsequently, we feed the user hypergraph adjacency matrix \mathbf{A}_{k}^{u} into the high-pass and low-pass user hypergraph filters respectively to decouple it into domain-exclusive user representations \mathbf{U}_{k}^{e} and domain-shared user representations \mathbf{U}_{k}^{s} , and feed the item hypergraph adjacency matrix \mathbf{A}_{k}^{v} into high-pass and low-pass item hypergraph filters respectively to decouple it into \mathbf{V}_{k}^{e} and \mathbf{V}_{k}^{s} . After the local model update is completed, the central server aggregates $\{\mathbf{U}_{k}^{s}\}_{k=1}^{K}$ to obtain the global representation \mathbf{U}^{g} used in the subsequent training round. The local perturbed domain-shared user representations are denoted as $\widetilde{\mathbf{U}}_{k}^{s}$.

Each client's local model is divided into a global branch with low-pass user/item hypergraph filters (parameterized by $\phi_k^{\rm u}/\phi_k^{\rm v}$), and a local branch with high-pass user/item hypergraph filters (parameterized by $\theta_k^{\rm u}/\theta_k^{\rm v}$). At the end of each training round, the server aggregates $\{\phi_k^{\rm u}\}_{k=1}^K$ to derive global user low-pass hypergraph parameters $\phi^{\rm g}$ which are then shared among clients in the subsequent training round.

2.2 Overview of FedHCDR

Our proposed FedHCDR, depicted in Fig. 2, utilizes client-server federated learning architecture. Each client's model is divided into a local branch (in yellow) and a global branch (in purple). In each training round, only domain-shared user representations and model parameters are aggregated. During the test phase, both domain-exclusive and domain-shared representations are utilized together for local predictions.

2.3 High/Low-Pass Hypergraph Filter



(a) Construction of the user and item (b) Hypergraph random walk. hypergraph adjacency matrix.

Fig. 3. Initialization of hypergraph filter.

Construction of the User and Item Hypergraph Adjacency Matrix. First, given the user-item incidence matrix \mathbf{A}_k in domain k, we can denote the hypergraph incidence matrices of users and items respectively as:

$$\mathbf{H}_{k}^{\mathrm{u}} = \mathbf{A}_{k}, \quad \mathbf{H}_{k}^{\mathrm{v}} = \mathbf{A}_{k}^{T}, \tag{1}$$

where each element $(\mathbf{H}_{k}^{u})_{ij}$ describes whether the vertex(user) u_{i} belongs to the hyperedge(item) v_{j} . Then, let us denote the item popularity debiasing matrix for the user hypergraph as \mathbf{P}_{k} , and we can obtain the normalized unbiased hypergraph adjacency matrix and hypergraph Laplacian matrix as follows:

$$\mathbf{A}_{k}^{\mathrm{u}} = \mathbf{D}_{k}^{\mathrm{u},-\frac{1}{2}} \left(\mathbf{H}_{k}^{\mathrm{u}} \mathbf{P}_{k}^{\mathrm{v}} \mathbf{D}_{k}^{\mathrm{v},-1} \mathbf{H}_{k}^{\mathrm{u},T} - \widetilde{\mathbf{D}}_{k}^{\mathrm{u}} \right) \mathbf{D}_{k}^{\mathrm{u},-\frac{1}{2}} \in \mathbb{R}^{|\mathcal{U}| \times |\mathcal{U}|}$$

$$\mathbf{L}_{k}^{\mathrm{u}} = \mathbf{I} - \mathbf{D}_{k}^{\mathrm{u},-\frac{1}{2}} \left(\mathbf{H}_{k}^{\mathrm{u}} \mathbf{P}_{k}^{\mathrm{v}} \mathbf{D}_{k}^{\mathrm{v},-1} \mathbf{H}_{k}^{\mathrm{u},T} - \widetilde{\mathbf{D}}_{k}^{\mathrm{u}} \right) \mathbf{D}_{k}^{\mathrm{u},-\frac{1}{2}} \in \mathbb{R}^{|\mathcal{U}| \times |\mathcal{U}|},$$

$$(2)$$

where

$$\left(\mathbf{D}_{k}^{\mathrm{u}}\right)_{uu} = \sum_{v \in \mathcal{V}_{k}} \left(\mathbf{H}_{k}^{\mathrm{u}}\right)_{uv} \left(\mathbf{P}_{k}^{\mathrm{v}}\right)_{vv}, \quad \left(\mathbf{D}_{k}^{\mathrm{v}}\right)_{vv} = \sum_{u \in \mathcal{U}} \left(\mathbf{H}_{k}^{\mathrm{u}}\right)_{uv} \tag{3}$$

are the user (vertex) degree matrix and item (hyperedge) degree matrix for the user hypergraph respectively, and the item popularity debiasing matrix

$$\left(\mathbf{P}_{k}^{\mathbf{v}}\right)_{vv} = 1 - \frac{\left(\mathbf{D}_{k}^{\mathbf{v}}\right)_{vv}}{\sum_{v \in \mathcal{V}_{k}}\left(\mathbf{D}_{k}^{\mathbf{v}}\right)_{vv}}$$
(4)

is used to remove the estimation bias caused by the long tail effect of items. Intuitively, the higher the degree (frequency of interactions) of hyperedge (item) v, the lower the weight $(\mathbf{P}_k^{\mathbf{v}})_{vv}$ of it. Besides, matrix

$$\left(\widetilde{\mathbf{D}}_{k}^{\mathrm{u}}\right)_{uu} = \sum_{v \in \mathcal{V}_{k}} \left(\mathbf{H}_{k}^{\mathrm{u}}\right)_{uv} \frac{\left(\mathbf{P}_{k}^{\mathrm{v}}\right)_{vv}}{\left(\mathbf{D}_{k}^{\mathrm{v}}\right)_{vv}}$$
(5)

is used to ensure that the diagonal of the user hypergraph adjacency matrix $\mathbf{A}_{k}^{\mathrm{u}}$ is filled with 0, that is, there is no self-loop in the user hypergraph. The construction of the user and item hypergraph adjacency matrix is shown in Fig. 3(a).

Low-Pass Hypergraph Filter Representation Initialization. To make the low-pass user hypergraph filter capture smoother signals (i.e., user relationship information), we construct the Markov transition matrix \mathbf{M}_k^u of the user hypergraph and perform a \mathcal{T} -step hypergraph random walk to obtain the initialized low-frequency user representations $\mathbf{U}_k^{\mathrm{s},(0)}$, which is shown in Fig. 3(b). The Markov transition matrix of the user hypergraph (the same applies to the item hypergraph) is formulated as follows:

$$\mathbf{M}_{k}^{\mathrm{u}} = \mathbf{D}_{k}^{\mathrm{u},-1} \left(\mathbf{H}_{k}^{\mathrm{u}} \mathbf{P}_{k}^{\mathrm{v}} \mathbf{D}_{k}^{\mathrm{v},-1} \mathbf{H}_{k}^{\mathrm{u},T} - \widetilde{\mathbf{D}}_{k}^{\mathrm{u}} \right) \in \mathbb{R}^{|\mathcal{U}| \times |\mathcal{U}|}, \tag{6}$$

where $(\mathbf{M}_k^{u})_{ij} \in [0, 1]$ indicates the probability that the random walk moves from user i to user j in one step.

Then for the user hypergraph, we can get the transition matrices of random walk of steps $1, 2, \dots, \mathcal{T}$:

$$\mathbf{M}_{k}^{\mathrm{u},1}, \mathbf{M}_{k}^{\mathrm{u},2}, \dots, \mathbf{M}_{k}^{\mathrm{u},\mathcal{T}}.$$
(7)

Take diagonals of the transition matrices (where $(\mathbf{M}_{k}^{u,t})_{ii}$ indicates the probability of going from user *i* and return to itself in exactly *t* steps), and perform a linear transformation after concatenating, then the initialized user low-frequency representations $\mathbf{U}_{k}^{s,(0)}$ are obtained:

$$\widetilde{\mathbf{M}}_{k}^{u} = \operatorname{diag}\left(\mathbf{M}_{k}^{u,1}\right) \left\| \operatorname{diag}\left(\mathbf{M}_{k}^{u,2}\right), \cdots, \right\| \operatorname{diag}\left(\mathbf{M}_{k}^{u,\mathcal{T}}\right) \\ \mathbf{U}_{k}^{\mathrm{s},(0)} = f_{k}\left(\widetilde{\mathbf{M}}_{k}^{u}\right)$$

$$\tag{8}$$

Note that for the remaining hypergraph filters, we directly use the EmbeddingLookup operation to obtain randomly initialized representations. For example, for the high-pass user hypergraph filter, we have:

$$\mathbf{U}_{k}^{\mathrm{e},(0)} = \mathbf{E}_{k}^{\mathrm{u}} = \mathrm{EmbeddingLookup}\left(\mathbf{X}_{k}^{u}\right),\tag{9}$$

where $\mathbf{X}_{k}^{u} \in \{0,1\}^{|\mathcal{U}| \times |\mathcal{U}|}$ is the matrix of one-hot IDs of users. Same with $\mathbf{V}_{k}^{s,(0)} / \mathbf{V}_{k}^{e,(0)}$.

Adaptive High/Low-Pass Hypergraph Filtering. Based on the user and item hypergraph adjacency matrices \mathbf{A}_k^u and \mathbf{A}_k^v (The corresponding Laplacians are \mathbf{L}_k^u , \mathbf{L}_k^v), as well as the initialized user and item representations $\mathbf{U}_k^{\mathrm{s},(0)}/\mathbf{U}_k^{\mathrm{e},(0)}$, and $\mathbf{V}_k^{\mathrm{s},(0)}/\mathbf{V}_k^{\mathrm{s},(0)}$, we further perform adaptive high-pass/low-pass hypergraph filtering on user and item representations. Let us denote user and item representations after l layer graph convolution as $\mathbf{U}_k^{\mathrm{s},(l)}/\mathbf{U}_k^{\mathrm{e},(l)}$ and $\mathbf{V}_k^{\mathrm{s},(l)}/\mathbf{V}_k^{\mathrm{e},(l)}$.

According to the graph signal processing theory [5,6], given a signal $\boldsymbol{x} \in \mathbb{R}^{|\mathcal{V}|}$ on a graph $\mathcal{G} = (\mathcal{V}, \mathcal{E})$ and a filter \boldsymbol{h} , the graph convolution/filtering operator in the spectral (frequency) domain (denoted as $*_{\mathcal{G}}$) is defined as

$$\boldsymbol{y} = \boldsymbol{x} *_{\mathcal{G}} \boldsymbol{h} = \mathbf{Q} \left((\mathbf{Q}^T \boldsymbol{x}) \odot (\mathbf{Q}^T \boldsymbol{h}) \right), \tag{10}$$

where **Q** is the matrix of eigenvectors of the graph Laplacian, \odot denotes the element-wise Hadamard product, and \boldsymbol{y} is the output signal. In particular, we can define $\mathbf{Q}^T \boldsymbol{h} = (h(\lambda_1,), \cdots h(\lambda_n))^T$, where λ_k is the k-th eigenvalue of graph Laplacian **L** and $h(\lambda)$ is the spectral filter. Let $h(\boldsymbol{\Lambda}) = \text{diag}(h(\lambda_1), \cdots, h(\lambda_n))$, then we have:

$$\boldsymbol{y} = \left(\mathbf{Q}h(\boldsymbol{\Lambda})\mathbf{Q}^{T}\right)\boldsymbol{x} = h(\mathbf{L})\boldsymbol{x},\tag{11}$$

Following the previous works [6,7], we adopt K-order Chebyshev polynomials $h(\mathbf{L}) = \sum_{k=0}^{K} w_k T_k(\hat{\mathbf{L}})$ to approximate $h(\mathbf{L})$. Let $K = 1, w_0 = w, w_1 = w, \hat{\mathbf{L}} = \mathbf{L}_k^u - (1 + \beta)\mathbf{I}$, where β is a trainable parameter, then for the user hypergraph, the adaptive high-pass filtering operation is defined as follows:

$$\begin{aligned} \boldsymbol{y} &= \left(w\mathbf{I} + w\left(\mathbf{L}_{k}^{\mathrm{u}} - (\beta + 1)\mathbf{I}\right)\right)\boldsymbol{x} \\ &= w\left(\mathbf{L}_{k}^{\mathrm{u}} - \beta\mathbf{I}\right)\boldsymbol{x} \\ &= w\left((1 - \beta)\mathbf{I} - \mathbf{A}_{k}^{\mathrm{u}}\right)\boldsymbol{x}, \end{aligned}$$
(12)

where the hypergraph filter $h(\mathbf{L}) = \mathbf{L}_k^u - \beta \mathbf{I}, h(\lambda) = \lambda - \beta$, which is a adaptive high-pass hypergraph filter. Then for the high-pass user hypergraph filter, we have the following layer-wise propagation rule:

$$\mathbf{U}_{k}^{(\ell+1)} = \sigma \left(\mathbf{P}_{k}^{(l)} \mathbf{U}_{k}^{(\ell)} \mathbf{W}_{k}^{\mathrm{u},(\ell)} + \boldsymbol{b} \right)$$

$$\mathbf{P}_{k}^{(l)} = \left(1 - \beta^{(l)} \right) \mathbf{I} - \mathbf{D}_{u}^{-\frac{1}{2}} \left(\mathbf{H}_{k}^{\mathrm{u}} \mathbf{P}_{k}^{\mathrm{v}} \mathbf{D}_{\mathrm{v}}^{-1} \mathbf{H}_{k}^{\mathrm{u},T} - \widetilde{\mathbf{D}}_{u} \right) \mathbf{D}_{u}^{-\frac{1}{2}}$$
(13)

Let $K = 1, w_0 = w, w_1 = -w, \widehat{\mathbf{L}} = \mathbf{L}_k^u - (1 + \beta)\mathbf{I}$, where β is a trainable parameter, then for the user hypergraph, the adaptive low-pass filtering operation is defined as follows:

$$\begin{aligned} \boldsymbol{y} &= \left(w\mathbf{I} - w\left(\mathbf{L}_{k}^{\mathrm{u}} - (\beta + 1)\mathbf{I}\right)\right)\boldsymbol{x} \\ &= w\left((2 + \beta)\mathbf{I} - \mathbf{L}_{k}^{\mathrm{u}}\right)\boldsymbol{x} \\ &= w\left((1 + \beta)\mathbf{I} + \mathbf{A}_{k}^{\mathrm{u}}\right)\boldsymbol{x}, \end{aligned}$$
(14)

where the hypergraph filter $h(\mathbf{L}) = (2 + \beta)\mathbf{I} - \mathbf{L}_k^u, h(\lambda) = 2 + \beta - \lambda$, which is a adaptive low-pass hypergraph filter. Then for the low-pass user hypergraph filter, we have the following layer-wise propagation rule:

$$\mathbf{U}_{k}^{(\ell+1)} = \sigma \left(\mathbf{P}_{k}^{(l)} \mathbf{U}_{k}^{(\ell)} \mathbf{W}_{k}^{\mathrm{u},(\ell)} + \boldsymbol{b} \right)
\mathbf{P}_{k}^{(l)} = \left(1 + \beta^{(l)} \right) \mathbf{I} + \mathbf{D}_{u}^{-\frac{1}{2}} \left(\mathbf{H}_{k}^{\mathrm{u}} \mathbf{P}_{k}^{\mathrm{v}} \mathbf{D}_{\mathrm{v}}^{-1} \mathbf{H}_{k}^{\mathrm{u},T} - \widetilde{\mathbf{D}}_{u} \right) \mathbf{D}_{u}^{-\frac{1}{2}},$$
(15)

2.4 Local-Global Bi-directional Transfer Algorithm

In this section, we introduce our proposed local-global bi-directional transfer algorithm. As shown in algorithm 1, in each round, the server sends the current global low-pass filter parameters $\phi^{u,(t)}$ and global domain-shared user representations $\mathbf{U}^{g,(t)}$ to clients. In the local training stage, client k updates its local high-pass hypergraph filter HHF_k and low-pass hypergraph filter LHF_k in an alternating manner, to perform knowledge transfer of global \rightarrow local and local \rightarrow global, as shown in lines 5–10 of the pseudocode. This process allows for a trade-off between global and local models.

Finally, the server receives the updated local low-pass filter parameters $\{\widehat{\boldsymbol{\phi}}_{k}^{\mathrm{u},(t+1)}\}_{k=1}^{K}$ and local domain-shared representations $\{\mathbf{U}_{k}^{\mathrm{s},(t+1)}\}_{k=1}^{K}$ from clients, and updates the global model and user representations using weighted averaging.

Loss for HHF and LHF. We adopt multi-class cross-entropy loss to train local high-pass hypergraph filters and low-pass hypergraph filters. The loss of high-pass hypergraph filters is defined as follows (the same is true for low-pass hypergraph filters):

$$\mathcal{L}_{k}^{\mathrm{H}} = -\sum_{(u_{i}, v_{j}) \in \mathcal{E}_{k}} \log \frac{e^{s_{k}(\boldsymbol{u}_{k,i}, \boldsymbol{v}_{k,j})}}{\sum_{v_{j} \in \mathcal{V}_{k}} e^{s_{k}(\boldsymbol{u}_{k,i}, \boldsymbol{v}_{k,j})}}$$
(16)

To reduce the computational complexity, we use the negative sampling method for training:

$$\mathcal{L}_{k}^{\mathrm{H}} = -\sum_{(u_{i}, v_{j}) \in \mathcal{E}_{k}} \left[\log \sigma \left(s_{k} \left(\boldsymbol{u}_{k,i}, \boldsymbol{v}_{k,j} \right) \right) + \mathbb{E}_{v_{j}^{\prime} \sim p_{i}^{\prime}(\mathcal{V}_{k})} \log \sigma \left(-s_{k} \left(\boldsymbol{u}_{k,i}, \boldsymbol{v}_{k,j^{\prime}} \right) \right) \right],$$

$$(17)$$

Algorithm 1: Local-Global Bi-Directional Transfer Algorithm

Input: Local datasets $\mathcal{D} = \{\mathcal{D}_k\}_{k=1}^K$ (where $\mathcal{D}_k = (\mathcal{U}, \mathcal{V}_k, \mathcal{E}_k)$), local user/item hypergraph adjacency matrix $\mathbf{A}_{k}^{\mathrm{u}}/\mathbf{A}_{k}^{\mathrm{v}}$, total training round T **Output**: The optimal hypergraph high-pass filters $\{\boldsymbol{\theta}_k\}_{k=1}^K = \{(\boldsymbol{\theta}_k^{\mathrm{u}}, \boldsymbol{\theta}_k^{\mathrm{v}})\}_{k=1}^K$ The optimal hypergraph low-pass filters $\{\phi_k\}_{k=1}^K = \{(\phi_k^{\rm u}, \phi_k^{\rm v})\}_{k=1}^K$ 1 Server initialize $\phi^{u,(0)}$; for round $t = 0, 1, \cdots T - 1$ do 2 for each client $k \in K$ in parallel do 3 Receive $\phi_k^{\mathbf{u},(t)}$ and $\mathbf{U}_k^{\mathbf{g},(t)}$ from server; /* Global \rightarrow Local */ 4 5 $\mathbf{U}_{k}^{\mathrm{e}}, \mathbf{V}_{k}^{\mathrm{e}} = \mathrm{HHF}_{k} \left(\mathbf{A}_{k}^{\mathrm{u}}, \mathbf{A}_{k}^{\mathrm{v}}; \boldsymbol{\theta}_{k} \right);$ 6 $\left(\widehat{\boldsymbol{\theta}}_{k}^{\mathrm{u}}, \widehat{\boldsymbol{\theta}}_{k}^{\mathrm{v}}\right) = \arg\min_{\boldsymbol{\theta}_{k}} \mathcal{L}_{k}^{\mathrm{H}}\left(\mathbf{U}_{k}^{\mathrm{e}}, \mathbf{V}_{k}^{\mathrm{e}}\right) - \lambda I\left(\mathbf{U}_{k}^{\mathrm{e}}, \mathbf{U}_{k}^{\mathrm{g},(t)}\right);$ 7 /* Local \rightarrow Global */8 $\mathbf{U}_{k}^{\mathrm{s}}, \mathbf{V}_{k}^{\mathrm{s}} = \mathrm{LHF}_{k} \left(\mathbf{A}_{k}^{\mathrm{u}}, \mathbf{A}_{k}^{\mathrm{v}}; \boldsymbol{\phi}_{k} \right);$ 9 $\left(\widehat{\boldsymbol{\phi}}_{k}^{\mathrm{u}}, \widehat{\boldsymbol{\phi}}_{k}^{\mathrm{v}}\right) = \arg\min_{\boldsymbol{\phi}_{k}} \mathcal{L}_{k}^{\mathrm{L}}\left(\mathbf{U}_{k}^{\mathrm{s}}, \mathbf{V}_{k}^{\mathrm{s}}\right) - \lambda I\left(\mathbf{U}_{k}^{\mathrm{s}}, \mathbf{U}_{k}^{\mathrm{e}}\right);$ 10 Client send $\hat{\phi}_{k}^{u}$ and \mathbf{U}_{k}^{s} to server; 11 end 1213 14 15 end

where $(u_i, v_j)/(u_i, v'_j)$ are positive/negative user-item interaction pairs, and $u_{k,i}$, $v_{k,j}$ and $v'_{k,j}$ are the corresponding user/item representations. Here $s_k(\cdot, \cdot)$ is the score function. $p'_i(\mathcal{V}_k)$ denotes the nagative sampling distribution over the local items set \mathcal{V}_k for the *i*-th user.

MI for Knowledge Transfer. For domain k, we adopt mutual information for knowledge transfer between local domain-exclusive and global domain-shared user representations during local training, which is computed as follows (the same applies to local domain-shared and local domain-exclusive user representations):

$$I\left(\mathbf{U}_{k}^{\mathrm{e}},\mathbf{U}_{k}^{\mathrm{g}}\right) := \sum_{u_{i}\in\mathcal{U}}\left[-\mathrm{sp}\left(-\mathcal{D}_{k}\left(\boldsymbol{u}_{k,i}^{\mathrm{e}},\boldsymbol{u}_{k,i}^{\mathrm{g}}\right)\right)\right] - \sum_{u_{i},u_{i}'\in\mathcal{U},u_{i}'\neq u_{i}}\left[\mathrm{sp}\left(\mathcal{D}_{k}\left(\boldsymbol{u}_{k,i}^{\mathrm{e},\prime},\boldsymbol{u}_{k,i}^{\mathrm{g}}\right)\right)\right],$$

$$(18)$$

where $\boldsymbol{u}_{i}^{\mathrm{e}}$, $\boldsymbol{u}_{i}^{\mathrm{g}}$ are the local domain exclusive representation and global domain shared representation of user *i* respectively. \mathcal{D}_{k} is the discriminator, $\mathrm{sp}(x) = \log(1+e^{x})$ is the softplus function, and $\boldsymbol{u}_{i}^{\mathrm{e},\prime}$ denotes the negative sample randomly sampled from the user set \mathcal{U} .

2.5 Hypergraph Contrastive Loss

To better obtain domain-invariant user relationship information, when updating the low-pass user hypergraph filter, we do perturbation to the user hypergraph by randomly dropping a portion of edges. First, we sample a edge random masking matrix $\mathbf{B}_k^{\mathrm{u}} \in \{0,1\}^{|\mathcal{U}| \times |\mathcal{U}|}$, where $(\mathbf{B}_k)_{ij} \sim \text{Bernoulli}(1-p^{\mathrm{d}})$ indicates whether to drop the edge between nodes *i* and *j*. Here p^{d} is the probability of each edge being dropped. Then the perturbed user hypergraph adjacency matrix can be computed as:

$$\widetilde{\mathbf{A}}_{k}^{\mathrm{u}} = \mathbf{A}_{k}^{\mathrm{u}} \odot \mathbf{B}_{k}^{\mathrm{u}},\tag{19}$$

where \odot is the Hadamard product. Then, we compute the hypergraph contrastive loss as follows:

$$\mathcal{L}_{k}^{\text{GCL}}\left(\mathbf{A}_{k}^{\text{u}}, \widetilde{\mathbf{A}}_{k}^{\text{u}}, \boldsymbol{\phi}_{k}^{\text{u}}\right) = -\sum_{u_{i} \in \mathcal{U}} \left(\ell_{k}^{\text{MI}}\left(\boldsymbol{u}_{k,i}^{\text{s}}, \boldsymbol{z}_{k}^{\text{s}}\right) + \ell_{k}^{\text{MI}}\left(\boldsymbol{z}_{k}^{\text{s}}, \boldsymbol{u}_{k,i}^{\text{s}}\right)\right), \qquad (20)$$

where graph representations \pmb{z}_k^s is aggregated through the readout function $g(\cdot)$ on the user hypergraph:

$$\boldsymbol{z}_{k}^{s} = g\left(\{\boldsymbol{u}_{k,i}^{s}\}_{u_{i} \in \mathcal{U}}\right),\tag{21}$$

and the contrastive infomax item $\ell_k^{\text{MI}}\left(\boldsymbol{u}_{k,i}^{\text{s}}, \boldsymbol{z}_k^{\text{s}}\right)$ is defined as follows:

$$\ell_{k}^{\mathrm{MI}}\left(\boldsymbol{u}_{k,i}^{\mathrm{s}},\boldsymbol{z}_{k}^{\mathrm{s}}\right) = \mathbb{E}_{\mathbf{A}_{k}^{\mathrm{u}}}\left[\log \mathcal{D}_{k}\left(\boldsymbol{u}_{k,i}^{\mathrm{s}},\boldsymbol{z}_{k}^{\mathrm{s}}\right)\right] + \mathbb{E}_{\widetilde{\mathbf{A}}_{k}^{\mathrm{u}}}\left[\log\left(1 - \mathcal{D}_{k}\left(\widetilde{\boldsymbol{u}}_{k,i}^{\mathrm{s}},\boldsymbol{z}_{k}^{\mathrm{s}}\right)\right)\right], \quad (22)$$

where \mathcal{D}_k denotes the discriminator, which is trained to classify node embeddings based on whether they belong to the original graph $\mathbf{A}_k^{\mathrm{u}}$ or the perturbed graph $\widetilde{\mathbf{A}}_k^{\mathrm{u}}$. This loss can enforce the model to generate node embeddings that can distinguish between the real graph and its perturbed counterpart. Finally, the loss of the low-pass hypergraph filter in client k can be denoted as

$$\widetilde{\mathcal{L}}_{k}^{\mathrm{L}} = \mathcal{L}_{k}^{\mathrm{L}} + \gamma \mathcal{L}_{k}^{\mathrm{GCL}}$$
(23)

3 Experiments

In this section, we conduct a comprehensive set of experiments to evaluate the effectiveness of our framework FedHCDR by answering the following questions:

- RQ1: Does FedHCDR outperform state-of-the-art methods for FedCDR?
- RQ2: How do HSD and HCL components enhance the performance of recommendations?
- RQ3: Is our proposed HSD method able to achieve desirable decoupling?
- **RQ4**: Does a global-local trade-off exist in the FedCDR scenario? How do different hyperparameters λ and γ impact the recommendation performance?

Domain	#Users	#Items	#Train	#Valid	#Test	Density
Food	1898	11880	36097	1898	1898	0.177%
Kitchen	1898	18828	44021	1898	1898	0.134%
Clothing	1898	16546	20919	1898	1898	0.079%
Beauty	1898	12023	30067	1898	1898	0.148%
Sports	4004	35567	68627	4004	4004	0.054%
Clothing	4004	24130	31910	4004	4004	0.041%
Elec	4004	50838	131107	4004	4004	0.068%
Cell	4004	20556	36920	4004	4004	0.055%
Sport	6657	46670	108511	6657	6657	0.039%
Garden	6657	24575	69009	6657	6657	0.050%
Home	6657	39426	107273	6657	6657	0.046%
Toys	6657	40406	107041	6657	6657	0.045%

Table 1. Statistics of Three FedCDR scenarios.

3.1 Experimental Setup

Datasets. We utilize publicly available datasets from the Amazon website¹ to construct FedCDR scenarios. Ten domains were selected to generate three cross-domain scenarios: Food-Kitchen-Cloth-Beauty (FKCB), Sports-Clothing-Elec-Cell (SCEC), and Sports-Garden-Home-Toys (SGHT). Following the approach of previous studies [2,25], we filter out users with less than 5 interactions and items with less than 10 interactions. For the dataset split, we follow the leave-one-out evaluation method employed in previous studies [2,25]. Specifically, we randomly select two samples from each user's interaction history as the validation set and the test set, while the rest of the samples are used for training. The statistics of the FedCDR scenarios are summarized in Table 1.

Evaluation Metrics. To guarantee an unbiased evaluation, we follow the method described in Rendle's work [12]. Specifically, for each validation or test sample, we calculate its score along with 999 negative items. Subsequently, we evaluate the performance of the Top-K recommendation by analyzing the ranked list of 1,000 items using metrics such as MRR (Mean Reciprocal Rank) [13], NDCG@10 (Normalized Discounted Cumulative Gain) [14], and HR@10 (Hit Ratio).

Compared Baselines. We compare our methods with two types of recommendation models: (1) single-domain recommendation methods, like NeuMF [15], LightGCN [20], DHCF [4]. (2) federated cross-domain recommendation methods, such as FedGNN [16], PriCDR [9], P2FCDR [10] and FPPDM++ [11].

¹ https://jmcauley.ucsd.edu/data/amazon/.

Method	Food			Kitch	en		Cloth	ing		Beaut	y		Avg		
	MRR	HR	NDCG	MRR	HR	NDCG	MRR	HR	NDCG	MRR	HR	NDCG	MRR	HR	NDCG
		@10	@10		@10	@10		@10	@10		@10	@10	1	@10	@10
NeuMF	5.79	12.96	6.61	3.56	7.27	3.80	1.61	2.32	1.43	4.22	9.11	4.64	3.80	7.92	4.12
LightGCN	7.20	14.12	7.85	4.16	8.85	4.42	3.37	6.11	3.52	4.87	10.33	5.19	4.90	9.85	5.24
DHCF	7.02	14.93	7.81	4.17	9.43	4.61	3.58	6.59	3.79	4.98	10.57	5.43	4.93	10.38	5.41
FedGNN	7.15	13.91	7.75	4.15	9.01	4.46	3.45	6.38	3.65	4.86	10.12	5.12	4.90	9.85	5.25
PriCDR	7.34	16.60	8.58	4.55	9.11	4.88	3.49	5.95	3.55	5.26	10.48	5.51	5.16	10.54	5.63
P2FCDR	7.08	13.91	7.68	4.28	8.96	4.63	3.18	6.53	3.51	4.27	9.64	4.53	4.70	9.76	5.09
FPPDM++	7.25	14.01	7.85	4.19	9.17	4.54	3.60	6.27	3.71	4.89	10.22	5.16	4.98	9.92	5.31
FedHCDR (Ours)	7.35	16.75	8.62	4.56	9.69	4.97	3.72	6.61	4.01	5.32	10.59	5.56	5.24	11.00	5.79

Table 2. Federated experimental results (%) on the FKCB scenario. Avg denotes the average result calculated from all domains. The best results are boldfaced.

Table 3. Federated experimental results (%) on the SCEC scenario. Avg denotes the average result calculated from all domains. The best results are boldfaced.

Method	Sport	s		Clothing I			Elec			Cell			Avg		
	MRR	HR	NDCG	MRR	HR	NDCG	MRR	HR	NDCG	MRR	HR	NDCG	MRR	HR	NDCG
		@10	@10		@10	@10	1	@10	@10	1	@10	@10		@10	@10
NeuMF	2.30	3.90	2.19	1.06	1.70	0.86	4.68	9.42	5.13	3.14	5.57	3.21	2.80	5.14	2.85
LightGCN	4.03	8.64	4.43	3.37	6.82	3.61	6.78	12.86	7.43	5.09	10.21	5.61	4.82	9.63	5.27
DHCF	3.92	8.12	4.22	3.36	6.89	3.62	6.68	12.96	7.39	4.98	10.31	5.57	4.74	9.57	5.20
FedGNN	3.99	8.57	4.38	3.38	6.87	3.63	6.78	13.04	7.47	5.12	10.24	5.64	4.82	9.68	5.28
PriCDR	3.89	8.04	4.19	3.40	6.82	3.66	5.50	10.76	6.05	5.34	10.74	5.93	4.53	9.09	4.96
P2FCDR	3.77	7.84	4.06	3.13	6.59	3.41	6.25	12.36	6.93	5.14	10.51	5.75	4.57	9.33	5.04
FPPDM++	4.06	8.64	4.45	3.38	6.74	3.60	6.71	12.99	7.39	5.16	10.41	5.72	4.83	9.70	5.29
FedHCDR (Ours)	4.47	9.04	4.90	3.45	6.92	3.66	6.88	13.61	7.65	5.75	11.11	6.31	5.14	10.17	5.63

Implementation and Hyperparameter Setting. For all methods, the common hyperparameters are as follows: the training round is set to 60, the local epoch per client is set to 3, the early stopping patience is set to 5, the mini-batch size is set to 1024, the learning rate is set to 0.001, and the dropout rate is set to 0.3.

3.2 Performance Comparisons (RQ1)

Table 2, 3, 4 present the performance of compared methods on three different FedCDR scenarios: Food-Kitchen-Clothing-Beauty, Sports-Clothing-Elec-Cell, and Sports-Garden-Home-Toys.

Based on the experimental results, several insightful observations can be made: (1) Among the single domain baselines, LightGCN and DHCF perform better than NeuMF. This finding validates that modeling the relationship between users and items by GCN can enhance the representations in the FedCDR scenario. (2) Most cross-domain baselines perform better than single-domain baselines, which indicates that cross-domain knowledge helps improve recommendation performance. (3) Among the cross-domain baselines, both PriCDR and FPPDM++ outperform FedGNN and P2FCDR in most cases, indicating that representation/distribution alignment can effectively accomplish the knowledge transfer between domains in the FedCDR scenario. (4) Our proposed

Method	Sport	s		Gard	en		Home	9		Toys			Avg		
	MRR	HR	NDCG	MRR	HR	NDCG	MRR	HR	NDCG	MRR	HR	NDCG	MRR	HR	NDCG
		@10	@10	1	@10	@10	1	@10	@10		@10	@10		@10	@10
NeuMF	3.27	6.46	3.39	4.39	8.49	4.59	4.23	8.35	4.50	2.54	4.81	2.45	3.61	7.03	3.73
LightGCN	4.61	8.74	4.76	5.45	10.33	5.67	5.83	11.03	6.20	2.95	5.14	2.78	4.71	8.81	4.85
DHCF	4.78	8.92	4.95	5.49	10.43	5.73	5.82	11.31	6.28	3.34	5.35	3.14	4.86	9.00	5.03
FedGNN	4.72	8.74	4.84	5.60	10.61	5.86	5.91	11.30	6.33	3.15	5.06	2.91	4.84	8.93	4.99
PriCDR	4.59	8.46	4.78	6.02	11.69	6.46	5.56	11.09	6.20	4.92	9.15	5.02	5.27	10.09	5.61
P2FCDR	5.06	9.22	5.25	5.83	11.06	6.16	5.94	11.31	6.36	4.14	6.41	4.02	5.24	9.50	5.45
FPPDM++	4.60	8.73	4.75	5.49	10.32	5.70	5.80	11.15	6.20	3.12	5.20	2.92	4.75	8.85	4.89
FedHCDR (Ours)	5.46	10.08	5.75	6.13	11.85	6.58	6.18	12.15	6.74	4.92	8.28	5.04	5.67	10.59	6.03

Table 4. Federated experimental results (%) on the SGHT scenario. Avg denotes the average result calculated from all domains. The best results are boldfaced.

Table 5. Ablation study on FKCB, SCEC, and SGHT scenarios.

Method	FKCI	3		SCEC	3		SGHT			
	MRR	HR	NDCG	MRR	HR	NDCG	MRR	HR	NDCG	
		@10	@10		@10	@10		@10	@ 10	
LocalHF	4.97	9.98	5.33	4.84	9.70	5.31	4.86	8.85	4.97	
FedHCDR - w/o (HSD, HCL)	4.98	10.02	5.34	4.81	9.69	5.28	4.95	9.03	5.09	
FedHCDR - w/o HCL	5.18	10.95	5.73	5.07	10.12	5.60	5.64	10.58	6.01	
FedHCDR (Ours)	5.24	11.00	5.79	5.14	10.17	5.63	5.67	10.59	6.03	

method, FedHCDR, significantly outperforms all baselines in multiple metrics. This emphasizes the crucial role of hypergraph signal decoupling and hypergraph contrastive learning in capturing both local and global user features.

3.3 Ablation Study (RQ2)

We conduct an ablation study on the performance of FedHCDR, specifically examining the impact of HSD and HCL. Table 5 presents the performance results of different model variants in three FedCDR scenarios. LocalHF represents the HF model (vanilla hypergraph filter) without federated aggregation, FedHCDR-w/o (HSD, HCL) corresponds to FedHCDR without HSD and HCL, and FedHCDR-w/o HCL refers to FedHCDR without HCL. It is evident from the findings that FedHCDR-w/o (HSD, HCL) occasionally performs worse than LocalHF, highlighting the significance of data heterogeneity. Interestingly, FedHCDR-w/o HCL greatly outperforms both LocalHF and FedHCDRw/o (HSD, HCL), indicating the effectiveness of HSD in addressing the data heterogeneity across domains. Furthermore, the utilization of HCL enables further improvements in model performance.

3.4 Discussion of the User Representation (RQ3)

In this section, we aim to further validate the ability of our HSD to learn both domain-shared and domain-exclusive representations for users. To achieve this, we conduct a comparative analysis of three types of representations: domain-shared,

domain-exclusive, and domain-exclusive + domain-shared representations. The predictive performance of these representations is compared, as illustrated in Fig. 4. The results of our analysis reveal several interesting observations: (1) The predictive performance varies among the three types of representations, highlighting the effectiveness of our HSD. (2) The domain-exclusive + domain-shared representations outperform both the domain-shared and domain-exclusive representations, indicating that integrating information from multiple domains by considering both domain-shared and domain-exclusive features is highly effective.



Fig. 4. The predictive results of representations in FKCB and SGHT scenario.

3.5 Influence of Hyperparameters (RQ4)

Figure 5 displays the performance of MRR and @NDCG@10 as the coefficients λ and γ increase. The following observations can be made: (1) The overall performance of FedHCDR initially increases and then decreases as λ increases, peaking at 2.0. This suggests that an λ coefficient of 2.0 is optimal for local-global bidirection transfer and highlights the local-global trade-off. (2) The overall performance of FedHCDR follows a similar pattern with the increase of γ , reaching its peak at 2.0. This indicates that a γ coefficient of 2.0 is optimal for hypergraph contrastive learning.

4 Related Work

4.1 GCN-Based Recommendation

The development of graph neural networks has attracted considerable attention in the exploration of GCN-based Recommendation [17,18]. NGCF [19] leverages the user-item graph structure by propagating embeddings throughout it. LightGCN [20] simplifies the model design by including only the neighborhood aggregation for collaborative filtering. DHCF [4] utilizes the hypergraph structure to model users and items, effectively capturing explicit hybrid high-order



Fig. 5. Impact of coefficient λ and γ .

correlations. SGL and MCCLK [21,22] integrate contrastive learning into GCNbased recommendation methods. However, the aforementioned methods solely concentrate on a single domain, thus unable to fully exploit user data from multiple domains.

4.2 Cross-Domain Recommendation

DTCDR and DDTCDR [1,23] enhance the performance of recommendations on dual-target domains simultaneously. BiTGCF [2] introduces an innovative bi-directional transfer learning approach for cross-domain recommendation, utilizing the graph collaborative filtering network as the foundational model. Dis-Alig [24] proposes the use of Stein path alignment to align the latent embedding distributions across domains. CDRIB [25] suggests the use of information bottleneck regularizers to establish user-item correlations across domains. Nonetheless, these methods require access to all user-item interactions across domains, rendering them infeasible in the federated learning setting.

4.3 Federated Cross-Domain Recommendation

FedMF [26] effectively incorporates federated learning into the field of crossdomain recommendation. FedCTR [27] proposes a framework for training a privacy-preserving CTR prediction model across multiple platforms. FedCDR [8] deploys the user personalization model on the client side and uploads other models to the server during aggregation. P2FCDR [10] proposes a privacy-preserving federated framework for dual-target cross-domain recommendation. FPPDM++ [11] presents a framework that models and shares the distribution of user/item preferences across various domains. Nevertheless, none of these methods address the issue of cross-domain data heterogeneity.

5 Conclusion

In this paper, we present a novel framework called FedHCDR, designed to enable domains to collaboratively train better performing CDR models while ensuring

privacy protection. To address the issue of data heterogeneity, we introduce a hypergraph signal decoupling method called HSD that decouples user features into domain-exclusive and domain-shared features. Additionally, we devise a hypergraph contrastive learning module called HCL to learn more extensive domain-shared user relationship information by applying graph perturbation to the user hypergraph.

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Data-Agnostic Pivotal Instances Selection for Decision-Making Models

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Abstract. As decision-making processes become increasingly complex, machine learning tools have become essential resources for tackling business and social issues. However, many methodologies rely on complex models that experts and everyday users cannot really interpret or understand. This is why constructing interpretable models is crucial. Humans typically make decisions by comparing the case at hand with a few exemplary and representative cases imprinted in their minds. Our objective is to design an approach that can select such exemplary cases, which we call *pivots*, to build an interpretable predictive model. To this aim, we propose a hierarchical and interpretable pivot selection model inspired by Decision Trees, and based on the similarity between pivots and input instances. Such a model can be used both as a pivot selection method, and as a standalone predictive model. By design, our proposal can be applied to any data type, as we can exploit pre-trained networks for data transformation. Through experiments on various datasets of tabular data, texts, images, and time series, we have demonstrated the superiority of our proposal compared to naive alternatives and state-of-the-art instance selectors, while minimizing the model complexity, i.e., the number of pivots identified.

Keywords: Interpretable Machine Learning \cdot Explainable AI \cdot Instance-based Approach \cdot Pivotal Instances \cdot Transparent Model

1 Introduction

In recent years, Machine Learning (ML) models have become increasingly central in supporting human decision-making processes [11]. These models are relied upon to tackle business problems and social issues in health science, online threat detection, and shopping pattern analysis [9,14,21], among others. Still, these models rely on complex architectures, making it difficult for anyone, experts and end users alike, to understand their reasoning. Moreover, although these tools may achieve identical or even superior performances compared to humans, the "cognitive process" they employ is hardly comparable to the one humans may use to solve the same task [43]. Given the pervasiveness of these models, interpreting and explaining their predictions and decisions generated, ultimately unveiling the internal mechanism inside the "black-box", is crucial [27]. We can identify this as the main goal of Explainable AI (XAI) [7].

In order to construct ML models that are inherently interpretable, a possible avenue to explore involves harnessing the intuitive notion of similarity of discriminative or descriptive elements. Our fundamental assumption is that a model "reasoning" in terms of exemplary instances provides an inherently interpretable tool to decision-makers, analysts, and end-users alike [41]. As humans, our cognitive processes and mental models often rely on a form of *case-based* reasoning [38] in which we store in our memory a large set of past exemplary cases, and then retrieve them as needed according to the task at hand. While the retrieval mechanism is itself obscure, reasoning in terms of said similar cases is inherently interpretable. This form of reasoning is so ingrained in us that even small children are able to recognize, use, and play with novel objects they have never seen, but that, in some form, are similar to other objects that they already know [39]. Furthermore, this applies to a wide variety of modalities: we recognize relatives based on faces we have already seen, music genres and bands based on song we have already heard, the origin of a recipe based on other recipes we have already tasted, etc. [26]. At its most fundamental level, similarity, and more generally case-based reasoning, is a universal form of human reasoning, pervasive to a plethora of modalities and data types [19].

Case-based reasoning offers significant advantages for fostering interpretability across various domains such as health [4], financial risk prediction [31], general text domains [12,24], and time-series and image analysis [1]. Particularly in the latter, recent research [25,36] shows good promise on the effectiveness of this type of reasoning, which is often preferred by human subjects. Given these premises, we emphasize the importance of training data quality as a ground for similarity between pivots and instances to predict: poor diversity or bias can result in unrepresentative cases. In contrast, feature-based methods may be more robust in such contexts due to their focus on how features influence outcomes.

This paper aims to design an interpretable case-based model that selects descriptive and discriminative cases to solve a decision-making task. With this in mind, we introduce PIVOTTREE, a hierarchical and interpretable case-based model inspired by Decision Trees [8]. By design, PIVOTTREE lends itself to both *selection* and *prediction*. As a selection model, PIVOTTREE identifies a set of *pivots*, exemplary cases identified within a training set. As a predictive model, PIVOTTREE leverages the selected pivots to build a similarity-based Decision Tree, routing instances through its structure, and yielding a prediction and an associated explanation. Unlike traditional Decision Trees, the explanation is not a set of rules, but rather a set of pivots to which the instance is similar. Like distance-based models, PIVOTTREE is also a selection method, encoding instances in a similarity space that enables case-based reasoning. Finally, PIVOTTREE is a *data-agnostic* model, which can be applied to different data modalities, jointly solving both pivot selection and prediction tasks.



Fig. 1. PIVOTTREE as (a) selector, (b) interpretable model, (c) Decision Tree.

Figure 1 provides an example of PIVOTTREE on the iris dataset, wherein flowers are classified according to their petal characteristics. Starting from a dataset of instances, PIVOTTREE filters down a set of pivots (Fig. 1(a)), i.e., a set of representative flowers. Said *pivots* are then used to learn a case-based model wherein novel instances are represented in terms of their similarity to the induced pivots (Fig. 1(b)). Building on pivot selection, PIVOTTREE then learns a hierarchy of pivots wherein instances are classified. This hierarchy takes the form of a Decision Tree (Fig. 1(c)): novel instances navigate the tree, percolating towards pivots to which they are more similar, ultimately building a chain of similar pivots, and landing in a classification leaf. In this case, given a test instance x: if its similarity to pivot θ is higher than 0.89 (following the left branch), then x is classified as a *Setosa* flower. Otherwise (following the right branch), if x's similarity to pivot 1 is higher than 0.85 (left branch), then x is classified as a pivot 0 flower. If neither condition is met, x is classified as a Virginica flower. In contrast, a traditional Decision Tree (DT) would model the decision boundary with feature-based rules, e.g., "if petal length < 2.4 then Setosa else if petal width < 1.7 then Versicolor else Virginica". However, (i) such an approach can only model axis-parallel splits, and (ii) cannot be employed on data types with features without clear semantics. Hence, improving on traditional DTs, the case-based model learned by PIVOTTREE can provide interpretability even in domains such as images, text, and time series, where by-design interpretable models are both underperforming and lack interpretability. Furthermore, unlike conventional state-of-the-art distance-based predictive models such as KNN [17], our proposal introduces a hierarchical structure to guide similarity-based predictions.

Experiments conducted on 24 datasets of different modalities, i.e., tabular data, time series, images, and text, show that PIVOTTREE yields interpretable predictive models that are as effective as state-of-the-art approaches at a fraction of their complexity expressed as the number of pivots. Qualitative results indicate high effectiveness on different data modalities, while a sensitivity analysis shows stability in the accuracy when varying the number of pivots selected.

After a review of some related works in Sect. 2, in Sect. 3 we illustrate our proposal. Then, Sect. 4 reports the experimental results. Finally, Sect. 5 summarizes our contributions and open research directions.

2 Related Work

Similarity-based methods belong to one of two families: similarity methods, aiming to, given a fixed data representation, learn the proper pivots¹ through similarity on said representation; and *representation* methods, which instead fix a similarity function, and aim to learn a proper instance representation.

Similarity. Underlying similarity methods is the assumption of a fixed data representation. Among them, we can distinguish three subclasses of methods: covering, clustering, and partitioning methods. Covering methods aim to group records around pivots. ε -BALL [6] jointly learns a set of distance-based neighboring *coverages* centered on a set of pivots. Pivots are optimized to be as few as possible, while coverages to be as class-pure as possible. The resulting pivots are thus laid on a "flat" structure where no structure defines the relationship among pivots. Clustering algorithms can provide more nuanced pivot-to-pivot relationships by tackling the lack of inter-pivot relationships. The MINIMAX algorithm [5] builds on agglomerative clustering by identifying cluster representatives, aggregating them in a hierarchical fashion, resulting in a hierarchy of prototypes. PIVOTTREE improves on MINIMAX by greatly improving on its complexity, and by leveraging pivots to perform prediction. *Partitioning* algorithms segment the feature space, assigning a pivot to each segment. PROXIMITYFOR-EST [32] induces a forest of similarity-based Decision Trees routing instances according to two pivots similarities. Notably, pivots are selected randomly, and so is the similarity function, thus yielding highly randomized trees. Unlike covering algorithms, PIVOTTREE constructs hierarchies of pivots, thus improving model interpretability. Like partitioning algorithms, PIVOTTREE partitions the feature space, but unlike PROXIMITYFOREST, it adopts a pivot selection strategy and a fixed similarity function, greatly improving the robustness and variance of its results. Finally, in [45] is presented a related methodology to select the best split for DTs based on the average similarity of instance pairs belonging to each children node. While being comparable to PIVOTTREE as they both determine the best split w.r.t. a similarity function, despite the title, they are inherently different as in [45] are not identified prototypical instances, using traditional feature-based rules for the split.

Representation. Unlike similarity methods, *representation methods* fix a similarity function, and rely on learning a proper representation of the data to find pivots. Unsurprisingly, these methods are often neural models lacking interpretability. [18] and [35] introduce soft Decision Trees, wherein nodes hold pivots, and instances are routed probabilistically towards multiple paths in the

¹ In Sect. 2 we adopt the term *pivot* to refer to the instances selected by different proposals in the state-of-the-art which do not necessarily adopt this term.

tree, thus creating fuzzy chains of pivots. Other approaches improve on the data representation at the cost of the intra-pivot structure. The authors in [2] introduce a neural model that jointly learns the data representation and a set of pivots, which are later used for classification. Similarly, PROTOPNET [10] and HPNET [22] learn a neural network that identifies pivots by learning contrastive representations and employing them for classification. Recently, a set of extensions of PROTOPNET have been proposed. PROTOTEX [12] integrates a similar approach for texts using pre-trained language models. PROTOSENET [33] offers a model where pivots can be refined through user knowledge for general sequence-based data. PROTORYNET [24] improves PROTOSENET by handling longer textual sequences. Finally, by providing even more fine-grained pivots, CNN-TREES [44] learn a neural model that constructs a hierarchy of pivots, each layer more specific than the previous, and each pivot also providing a score indicating its contribution to the final prediction. Unlike neural representation methods, PIVOTTREE learns a crisp and fully interpretable model.

3 Pivot Tree Selection Model

We present here PIVOTTREE, an interpretable hierarchical pivot selection model inspired by Decision Trees [8]. Let us start by formalizing the problem and setting. Without loss of generality, we restrict ourselves to classification tasks and leave other tasks for future work.

Problem Setting. Given a population of instances represented as real-valued m-dimensional feature vectors² in \mathbb{R}^m and a set of class labels $C = \{1, \ldots, c\}$, we assume the existence of an unknown ground-truth function $g : \mathbb{R}^{n \times m} \to C$ mapping each vector in \mathbb{R}^m to one of the c classes in C. In case-based reasoning, the objective is to learn a function $f : \mathbb{R}^m \to C$ approximating g, with f being defined as a function of k exemplary cases named *pivots*. As explained in Sect. 2, similarity-based case-based models define f on a similarity space, often inversely denoted as "distance space", S induced by a similarity function $s : \mathbb{R}^m \times \mathbb{R}^m \to \mathbb{R}$ quantifying the similarity of instances [37].

Given a training set $\langle X, Y \rangle$ with $X = \{x_i\}_{i=1}^n$ of n instances, $Y = \{y_i\}_{i=1}^n$ with $y_i \in C$ the associated class labels, and a similarity function s, our objective is to learn a function $\pi : \mathbb{R}^{n \times m} \to \mathbb{R}^{k \times m}$ that takes as input X and returns a set $P \subseteq X$, i.e., $\pi_s(X) = P$, of k pivots such that the performance of f are maximized. Furthermore, aiming for transparency of the case-based predictive model f, our objective is to employ as an interpretable model f Decision Tree classifiers or k-Nearest Neighbors approaches [20] (kNN).

In practical terms, given a training set $\langle X, Y \rangle$ and a similarity function s, the selection method π selects k pivots P from X. Through the similarity function s and the pivots P, the dataset $X \in \mathbb{R}^m$ is mapped into the similarity space S,

² For the sake of simplicity, we consistently treat data instances as real-valued vectors. Any data transformation employed in the experimental section to maintain coherence with this assumption will be specified when needed.

and thus encoded into a representation $Z \in \mathbb{R}^{n \times k}$ where $Z_{i,j}$ is the similarity between the *i*-th object with the *j*-th pivot in *P*. Hence, the predictive model *f* is trained on $\langle Z, Y \rangle$. Then, given a test instance $x \in \mathbb{R}^m$, *x* is first mapped to a similarity vector $z = \langle s(x, p_1), \ldots, s(x, p_k) \rangle$ yielding its similarity to the set *P* of pivots. Then, *z* is provided to *f*, which performs the prediction.

When f is implemented with a Decision Tree, split conditions will be of the form $s(x, p_i) \ge \beta$, i.e., "if the similarity between instance x and pivot p_i is greater or equal then β , then ...", allowing to easily understand the logic condition by inspecting x and p_i for every condition in the rule.

On the other hand, when f is implemented as a kNN, every decision will be based on the similarity with a few neighbors (typically between one and five) in the similarity space S obtained computing the similarity between each instance with respect to the selected pivots. A human user just needs to inspect x and the similarities with the pivots P and the instances in the neighborhood. When the number of pivots is kept small, the interpretability of both methods increases, limiting the expressiveness. Vice versa, using a selection model π that returns a large number k of pivots can increase the performance at the cost of interpretability. Our proposal aims to balance these two aspects by allowing the selection of a small number of pivots that still guarantee comparable performance to interpretable predictive models.

PivotTree Algorithm. In this paper, with PIVOTTREE, we implement the selection function π . Much like Decision Tree induction algorithms [8], PIVOT-TREE greedily learns a hierarchy of nodes, each node splitting instances towards one of its two children, ultimately reaching terminal leaf nodes, which are associated with a classification label. The splitting is based on *discriminative* pivots and representative pivots. Let X_t be the records constrained by the decision path at iteration t in the tree construction, and Y_t the associated class labels, then a discriminative pivot is an instance of class c, i.e., $p^- \in X_t^{(c)} = \{x_i | x_i \in X_t^{(c)} \}$ $X_t \wedge y_i = c$, such that it maximizes the impurity gain when partitioning X_t w.r.t. the similarity with p^- . Formally, if $X_{t,l} = \{x_i \in X_t | s(x_i, p^-) \geq \beta_t\},\$ $X_{t,r} = \{x_i \in X_t | s(x_i, p^-) < \beta_t\}$ and $Y_{t,l}, Y_{t,r}$ are the associated class labels, respectively, then if $\delta_s(p^-, X_t, Y_t)$, is the Information Gain calculated as in [8] maximizing a task-dependent measure like Entropy or Gini w.r.t. the similarities between p^- and X_t (instead of w.r.t. the features \mathbb{R}^m of X_t), it does not exist another instance \hat{p}^- such that $\delta_s(\hat{p}^-, X_t, Y_t) > \delta_s(p^-, X_t, Y_t)$. Furthermore, besides discriminative pivots, for each iteration v, PIVOTTREE also identifies representative pivots. A representative pivot is an instance of class c, i.e., $p^+ \in \{x_i | x_i \in X_t \land y_i = c\}$ that maximizes the similarity with all the other instances described by the same node and belonging to the same class, i.e., $p^+ = \arg \max_{p' \in X_t^{(c)}} \sum_{x_i \neq p' \in X_t^{(c)}} s(x_i, p').$

In Algorithm 1, we illustrate the pseudo-code for training a PIVOTTREE. Given the dataset and labels $\langle X, Y \rangle$, the similarity function *s*, the maximum tree depth *maxdepth*, it returns the set *P* set of selected pivots, and the trained decision tree *T* (line 5). After initializing the tree and pivots (line 1), PIVOT-TREE induces a similarity matrix *S* between all pairs of instances in *X* (line 2).

Algorithm 1. PIVOTTREE (X, Y)	
Input: $\langle X, Y \rangle$ data and labels, s similarity fun	ction, maxdepth maximum tree depth
Output: P set of pivots, T learned tree	
1: $T \leftarrow \emptyset; P \leftarrow \emptyset;$	Variables initialization
2: $S \leftarrow \langle s(x_i, x_j) \rangle \ \forall x_i, x_j \in X \times X$	Calculate similarity matrix
3: $P, T \leftarrow PTR(X, Y, T, P, S);$	Start of recursive procedure
4: return P, T	
5: function $PTR(X, Y, T, P, S)$	
6: if DEPTH $(T) \leq maxdepth$ then	
7: for $c \in C$ do	
8: $p^- \leftarrow \underset{x' \in X^{(c)}}{\operatorname{argmin}} \delta_s(x', X, Y);$	\triangleright Get discriminative pivot
9: $p^+ \leftarrow \underset{x \in X^{(c)}}{\operatorname{argmax}} \sum_{x \neq x' \in X^{(c)}} s(x, x')$; \triangleright Get representative pivot
10: $P \leftarrow P \cup \{p^-, p^+\};$	$Descript{Add}$ pivots to result set
11: $X_l, X_r, Y_l, Y_r \leftarrow \text{SplitData}(X, Y, P)$; ▷ Split data w.r.t. P
12: $P_l, T_l \leftarrow \operatorname{PTR}(X_l, Y_l, T, P, S)$	▷ Recourse on left child
13: $P_r, T_r \leftarrow \operatorname{PTR}(X_r, Y_r, T, P, S)$	▷ Recourse on right child
14: $T \leftarrow \text{AddSplitToTree}(T, T_l, T_r);$	\triangleright Add split to tree
15: return $P, T;$	Return current pivots and tree
16: else	
17: $p^+ \leftarrow \underset{x \in X^{(c)}}{\operatorname{argmax}} \sum_{x \neq x' \in X^{(c)}} s(x, x');$	\triangleright Get representative pivot
18: $P \leftarrow P \cup \{p^+\};$	$Descript{Add}$ pivots to result set
19: return P , MAKELEAF (T) ;	▷ Return current pivots and leaf

Then, the recursive procedure PTR is started (line 3). If the current depth of the tree DEPTH(T) is lower than the maximum tree depth maxdepth (line 6), then for each class, the most discriminative and most representative pivots are selected and added to the result set P (lines $7-10)^3$. We notice that, since the similarity matrix S is calculated at the beginning, the pairwise similarities to select the most discriminative and representative pivots are available without performing any calculus. The set P of discriminative and representative pivots is then used to select the best split to partition the data with the SPLITDATA function, again maximizing the Information Gain w.r.t. the similarities w.r.t. the pivots in P (line 11). We highlight that, by construction, SPLITDATA selects a discriminative pivot. However, we keep these aspects separated as it is possible to run PIVOTTREE relying only on representative pivots. After that, PIVOTTREE recourses on the left and right subsets X_l, Y_l and X_r, Y_r and composes the tree returned (lines 12–14). On the other hand, if the maximum depth (line 16) or other stopping conditions are met, then the current pivots, augmented with the descriptive pivots of the records in the leaf, and a leaf itself (lines 17–19), are

³ To ease the computational burden, and similarly to other implementations, e.g., scikit-learn, we select only a subset of splits is evaluated.

	Iı	nag	ge					Та	bul	ar							Гext	t		Time Series					
name	cars	cifar	mnist	breast	compas	diva	german	heloc	house	iris	page	sonar	vertebral	wine	imdb	lyrics	news	vicuna	tgpt	devices	ecg	gun	wafer	WOLMS	
tr	· 9k	6k	6k	.4k	5k	9k	.8k	8k	16k	.2k	4k	.2k	.4k	5k	25k	26k	12k	7k	3k	9k	.6k	50	1k	.2k	
ts	9k	1k	1k	.2k	2k	4k	.4k	4k	7k	45	2k	63	93	2k	25k	11k	8k	2k	.8k	8k	5k	.2k	7k	77	
m	768	128	256	30	17	330	61	23	16	4	9	60	6	11	768	768	768	768	768	96	140	150	152	900	
c	196	10	10	2	3	2	2	2	2	3	2	2	2	7	2	2	3	20	2	7	5	2	2	2	

Table 1. Datasets info: tr training and ts test size, m nbr. features, c nbr. classes.

returned. Thus, the complexity of the PIVOTTREE is theoretically bounded by the calculus of the similarity matrix S.

Furthermore, besides being used as a pivot selector method (π) , we underline that PIVOTTREE can be employed as a standalone predictive model by combining the encoding in the similarity space and the tree induction f. In this case, we do not need to train additional interpretable models, as both pivot selection and case-based prediction are already integrated into the model.

Data Agnosticism. By design, PIVOTTREE is a data-agnostic model that leverages the concept of similarity to conduct both selection and prediction tasks simultaneously. While some data types, e.g., relational data, are more amenable than others, e.g., images or text, to similarity computation, with our contribution, we aim to address all data types as one. By decoupling similarity computation and object representation, PIVOTTREE can be applied to any data type supporting a mapping to \mathbb{R}^m , i.e., text through language model, images through vision models, graphs through graph models, etc. In the following experimentation, besides tabular data, we focus on time series, images, and text.

4 Experiments

In this section, we evaluate the performance of PIVOTTREE, which we implemented in Python⁴, on different datasets with different modalities, and against a wide array of competitors. Our objective is to demonstrate that PIVOTTREE is as accurate as state-of-the-art pivot selection methods, while being simpler. With PTS, we indicate PIVOTTREE used as Selector, while with PTC, we refer to PIVOTTREE directly used as Classification model.

Baselines and Competitors. We compare PIVOTTREE with the following baselines and state-of-the-art similarity-based approaches for pivot selection (π) :

- RND: randomly selects instances from the training set to be used as pivots;
- RNC: same as RND, but instances are sampled separately from each class;
- KMS: runs kMeans [40] and adopts the centroids as pivots;
- KMD: runs kMedoids [40] and adopts the medoids as pivots;

 $^{^{4}\} https://github.com/msetzu/pivottree.$

- EBL: selects pivots according to the ε -BALL algorithm⁵ [6].

Regarding model selection, we performed grid searches over the hyper parameter space, selecting the best-performing model on a validation set. On RND, RNC, and KMS, the number of pivots |P| is selected within a grid on $|P| \in [2, 32]$. On EBL, the grid search for ε is performed on an interval between the 2^{nd} and the 40^{th} quantile of the empirical similarity distribution, as suggested in [6]. Regarding the interpretable predictive models (f) to be used on the selected prototypes, we rely on kNN and Decision Tree as implemented by the **sklearn** Python library. For PIVOTTREE, both used as selector or predictor, i.e., PTS or PTC, the best *maxdepth* is searched in an interval [2,4]. Obviously, a deeper PIVOTTREE yields the selection of a larger number of pivots. Finally, to guarantee interpretability for the predictive models, we fix the hyper parameters as follows. Maximum depth equals four for Decision Trees [3], and the maximum number of neighbors for kNN and DT directly trained on the original feature space while preserving hyper parameters.

Evaluation Measures. We evaluated the effectiveness of the selected pivots by measuring the F1-score of the predictive models relying on the different sets of pivots⁶. In line with the literature [7], as proxy of interpretability, we evaluated the *complexity* in terms of k, the number of selected pivots. Note that k can either be user-given, or optimized w.r.t. a given validation set. We experiment in both settings. Finally, to account for differences in datasets, and ease comparison, we turn complexity into *simplicity* as $1 - \frac{k}{|X|}$.

Datasets. In order to show the effectiveness of our proposal for different data types, we experimented with 11 tabular datasets, 5 time series datasets, 3 image datasets, and 5 text datasets. Table 1 reports some dataset details⁷. For tabular datasets, in order to perform a direct distance comparison between instances, we leave unvaried numeric and ordinal features while we one-hot encode categorical ones. We discard instances presenting missing values for one or more features. The datasets are then normalized with a z-score normalization by removing the mean and scaling to unit variance. Time series datasets are left unchanged as they are already preprocessed and normalized. For textual datasets, we first embed the input text with the all-mpnet-base-v2 sentence transformer model⁸, which yields L2-normalized 768-dimensional dense vector with magnitude 1. Finally, for image datasets, we embed each dataset with pretained and fine-tuned vision models. Further details are provided in the project repository. On the basis of these encodings, the similarity s is based on the Euclidean distance. While text embeddings usually rely on cosine similarity, in [29] it is shown that under unit normalization, the two are directly proportional and thus order-preserving.

 $^{^{5}\} https://docs.seldon.io/projects/alibi/en/latest/methods/ProtoSelect.html.$

 $^{^{6}}$ For multi-class datasets we calculate the metric for each label and report the unweighted mean.

⁷ The links to the various repositories and detailed preprocessing steps for the different datasets are available on the project repository.

⁸ https://huggingface.co/sentence-transformers/all-mpnet-base-v2.

Tabular datasets are divided into 70% training and 30% testing, while non tabular data sets come with their own split into training and test set. During model selection, a further split is performed, allocating for each development set 80% of the instances for training and 20% for validation. Thus, for each pivot selection method and classification method of each dataset, we perform a hold-out model-selection procedure, i.e., we find the best-performing hyper parameters configuration on the validation set and use it in the model-assessment phase, training on the whole training set and considering the resulting performances on the test set for final assessment.



Fig. 2. PIVOTTREE prediction and explanation on cifar. Top: selected pivots. Center and bottom: two classification examples. On the left a test instance; in the center, the five nearest neighbors of the selected pivots; and on the right a case-based decision rule.

Qualitative Results. In the following, we illustrate some qualitative examples on different data types to show the usability of PIVOTTREE at prediction and explanation time with DT and kNN with the same set of pivots. PIVOTTREE selects a set of pivots, which are then the training set for either a kNN or a DT. In the latter case, the data is first encoded in a pivot-instance similarity matrix. In Fig. 2, we report two prediction and explanation examples on cifar. The top rows illustrated the pivots selected by PIVOTTREE. The central and bottom rows show two classification examples for the *bird* and *cat* test instances, both on the left of the respective rows. Next to the test instances, we display the five neighbors selected by a kNN on the pivots similarity space. We can notice that for the *bird* example, all the neighbors are indeed birds quite similar in color and shape to the test instance. On the other hand, for the *cat* example, there are also some deers among the neighbors that however are in the same palette as the test instance. Finally, the right column shows the decision rules obtained by training a DT in the similarity space derived by PIVOTTREE. We notice that the bird is recognized thanks to its dissimilarity with the car pivot p_{574} and its

similarity with the bird pivot p_{65} . On the other hand, the cat is classified due to its dissimilarity with p_{351} and p_{781} . Thus, similarly to humans, these kinds of models can also reason by exclusion, suggesting their applicability also in the context of few-shot learning.



Fig. 3. Test time series $(1^{st}$ column), pivots extracted by PIVOTTREE $(2^{nd}$ column), neighbors selected by kNN $(3^{rd}$ column) and decision rule $(4^{tg}$ column) on the gun (top) and ecg (bottom) datasets.

Similarly, Fig. 3 reports two examples from the gun and ecg datasets, classifying tracked hand movements as gun draws and holsterings or not, and heartbeats of five different types, respectively. For both cases, the test instance has a large set of peculiarly similar neighbors, each with minimum variations. For the gun dataset, PIVOTTREE has identified three pivots, two for the not a gun class, both characterized by short starting and ending movements, and interleaved by a long plateau. Here, the movement is sharp, but somewhat smooth, especially when gun is drawn, rather than holstered. The third pivot, associated to the qun class, is instead characterized by minimal motions, interleaved by a sharp draw, and a short plateau. The more pronounced movement closely resembles the test instance, but for a slight shift, and the instance is correctly classified by kNN as gun. The decision rule of the DT instead recognizes the gun class due to the similarity with p_{10} and p_{45} . For ecg, PIVOTTREE identified four pivots, the test instance is very similar to p_{196} in the initial part and to p_{41} in the final part. The kNN classifier correctly retrieves neighbors with this shape and the test is correctly classified as *normal*. The DT instead distinguishes the normal class due to its limited similarity with p_{227} and high similarity with p_{41} .

Quantitative Results. Table 2 and Table 3 report the predictive model performance (F1-score) and complexity (number of pivots), respectively, per data modality and predictive model, i.e., DT and kNN. The bottom rows of the table report the average performance and standard deviations for all methods, and the rank of the pivot selection methods. The best and second-best performers per dataset among the pivot selection methods are in bold and italic, respectively. We can notice that when relying on the original data representation, i.e., when

р	redictor			De	cisio	n T	ree						kNl	V		
s	elector	-	RND	RNC	KMS	KMD	EBL	PTS	PTC	-	RND	RNC	KMS	KMD	EBL	PTS
60	cars	.01	.02	.02	.02	.02	.02	.02	.00	.86	.75	.84	.78	.75	.80	.84
3	cifar	.75	.40	.40	.40	.41	.41	.39	.11	.87	.87	.87	.88	.88	.88	.87
	mnist	.68	.44	.41	.53	.44	.41	.37	.29	.97	.96	.96	.96	.96	.97	.96
	breast	.95	.94	.93	.93	.94	.95	.94	.95	.96	.95	.95	.95	.96	.96	.95
	compas	.50	.47	.48	.46	.48	.48	.52	.49	.46	.47	.48	.48	.48	.46	.47
	german	.58	.54	.53	.48	.48	.61	.50	.48	.67	.59	.59	.59	.59	.65	.60
	heloc	.70	.66	.66	.65	.66	.67	.68	.66	.67	.66	.66	.67	.67	.67	.66
	house	.80	.72	.72	.73	.73	.78	.76	.77	.83	.80	.79	.80	.80	.82	.80
lar	iris	1.0	.94	.91	.96	.96	.90	.92	.92	1.0	.99	1.0	1.0	1.0	1.0	.98
pn	page	.88	.84	.86	.87	.85	.88	.84	.87	.90	.88	.89	.90	.89	.90	.88
Ta	diva	.79	.60	.59	.58	.56	.64	.64	.63	.76	.72	.72	.70	.71	.75	.73
	sonar	.74	.70	.73	.72	.71	.77	.73	.59	.94	.82	.84	.83	.81	.89	.84
	vert.	.72	.68	.69	.66	.69	.65	.71	.68	.73	.69	.69	.73	.76	.74	.78
	wine	.20	.19	.20	.19	.20	.20	.20	.18	.37	.35	.35	.35	.36	.35	.35
	imdb	.70	.73	.72	.75	.74	.79	.78	.78	.78	.78	.77	.79	.80	.79	.82
+	lyrics	.66	.69	.69	.68	.68	.70	.70	.70	.71	.70	.70	.70	.70	.71	.71
ex	news	.12	.16	.16	.19	.18	.16	.24	.01	.69	.55	.50	.62	.60	.66	.65
	tgpt	.84	.80	.80	.80	.81	.84	.79	.84	.92	.88	.88	.90	.90	.89	.89
	vicuna	.63	.57	.55	.55	.59	.64	.63	.59	.68	.69	.69	.67	.71	.73	.72
	devices	.25	.33	.32	.34	.34	.39	.42	.34	.49	.47	.46	.48	.48	.49	.52
ŝ	worms	.54	.54	.53	.57	.56	.56	.56	.56	.60	.58	.61	.58	.58	.61	.70
E.	ecg	.52	.50	.51	.50	.50	.53	.51	.51	.57	.54	.54	.55	.55	.56	.56
Ĥ	gun	.80	.77	.77	.76	.78	.77	.71	.74	.91	.87	.87	.89	.89	.88	.84
	wafer	.90	.93	.93	.94	.94	.95	.92	.93	.99	.97	.98	.98	.98	.98	.98
	avg	.64	.59	.59	.59	.59	.61	.60	.57	.76	.73	.73	.74	.74	.76	.75
	std	.26	.25	.25	.25	.25	.25	.24	.29	.18	.18	.18	.18	.18	.18	.17
	rank		4.8	4.8	4.21	3.9	2.5	3.5	4.3		4.9	4.4	3.6	3.15	2.2	2.9

 Table 2. Predictive model performance as F1-score.

using directly DT or kNN on the training data, we have slightly better performance at the cost of losing the interpretability for non-tabular datasets. Focusing on predictive models relying on pivots, we notice that EBL has, on average, the highest F1-score (Table 2) immediately followed by PTS both for DT and kNN. We observe that the difference in the average of F1-score between EBL and

pr	edictor			$D\epsilon$	ecis	ion	Tr	ee					kN	N		
s	elector	-	RND	RNC	KMS	KMD	EBL	PTS	PTC	_	RND	RNC	KMS	KMD	EBL	PTS
60	cars	-	10	196	6	6	64	778	4	- ,	32	196	32	2 8	64	974
B	cifar	-	32	10	18	28	220	118	12	-	32	20	22	30	18	42
	mnist	-	4	10	4	4	261	2	12	-	32	20	30	30	133	73
	breast	-	32	24	20	28	88	39	6	-	26	28	12	6	99	20
	compas	-	32	32	30	18	70	9	10		18	18	30	32	581	7
	german	-	26	32	24	24	60	22	10	- :	32	32	32	32	72	32
	heloc	-	28	32	24	32	880	9	9	-	32	32	18	22	378	9
	house	-	32	32	16	20	2k	6	13	-	32	26	28	32	1k	30
lar	iris	-	28	32	28	4	69	16	3	-	28	28	32	28	20	10
pn	page	-	32	32	22	24	105	69	10	-	30	32	6	20	112	6
Ja	diva	-	30	32	32	30	528	83	13	-	32	28	30	30	311	13
	sonar	-	32	32	22	22	26	21	4	-	32	20	22	24	21	6
	vert.	-	30	32	28	8	61	53	8	-	32	10	18	8	21	3
	wine	-	28	28	22	22	150	158	14	- ;	24	28	22	32	32	121
	imdb	-	32	32	10	18	531	26	8	-	32	32	8	30	980	26
H	lyrics	-	30	32	30	24	5k	24	2	-	32	32	30	28	156	99
Lex	news	-	30	20	20	22	215	106	13	-	32	20	32	32	215	844
	tgpt	-	26	32	14	26	247	18	12	- ,	32	32	32	28	187	68
	vicuna	-	32	32	32	14	107	40	12	- ,	32	32	32	32	540	30
	devices	-	32	28	24	8	136	408	12	-	32	28	26	30	896	89
e S	worms	-	32	30	12	16	107	25	6	-	8	24	26	14	32	20
<u><u></u></u>	ecg	-	28	28	16	24	43	14	3	-	30	24	30	26	96	37
H	gun	-	28	20	8	30	15	2	2	-	20	24	8	20	5	4
	wafer	-	26	28	12	24	43	13	3	- ;	30	32	32	30	43	45
	avg	-	28	35	20	20	523	86	8	-	29	33	24	26	259	109
	std	-	γ	35	8	8	1k	170	4	-	6	35	8	γ	338	249
	rank		4.8	4.9	3.	3.3	6.6	3.9	1.4	3	3.7	3.1	2.9	2.9	5.1	3.3

 Table 3. Predictive model complexity as number of pivots used.

PTS is only 0.1. All the other approaches follow them, with PTC being worse than EBL, thus indicating that PIVOTTREE, in its current implementation, works better as a selector than as a classifier. On the other hand, concerning the complexity (Table 3), even though PTS is not minimizing the number of pivots selected compared to other methods such as KMS, it still requires less than half of the pivots used by EBL to guarantee comparable performance.



Fig. 4. Comparison of model's rank w.r.t. F1-score and complexity against each other with the Nemenyi test. Groups of classifiers that are not significantly different at 95% significance level are connected. Best ranks on the right.

The non-parametric Friedman test compares the average ranks of the various methods over multiple datasets w.r.t. an evaluation measure, in our case, F1-score and complexity. The null hypothesis that all methods are equivalent is rejected (p < 0.001) for all the experiments reported in the various tables. The comparison of the ranks of all methods against each other can be visually represented as shown by the critical difference plots in Fig. 4: lower rank values indicate better models, i.e., best ranks on the right (see [16] for details). In Fig. 4, methods statistically equivalent according to a post-hoc Nemenyi test are connected by black lines. We notice that regardless of the classification model f used, EBL and PTS are tied w.r.t F1-score, while PTS is significantly less complex and untied w.r.t. the number of pivots selected.

In summary, PTS is the best pivot selector, achieving high predictive performance with a smaller number of pivots. Such a result is best appreciated in Fig. 5, where we show the mean and standard deviation of the F1-score and the simplicity of pivot selection methods. Besides, Fig. 5 also highlights the lowest variability of PTS w.r.t EBL in terms of simplicity.

We repeated the experiments in a constrained setting⁹ wherein pivot selection was limited to a maximum of 20 pivots (Table 4 and Fig. 6). While the average

⁹ cars has not been used as it contains 196 classes, and all the methods would have failed.


Fig. 5. Scatter plots for average F1-score and simplicity for pivot selection methods with error bars reporting 10% of the standard deviation.



Fig. 6. Comparison of model's rank w.r.t. F1-score against each other with Nemenyi test. Classifiers that are not significantly different at 95% significance level are connected. Best ranks on the right. Models limited to 20 pivots.



Fig. 7. F1-score varying the number of pivots w.r.t. bins of pivots for datasets with different number of classes.

performance remains more or less unchanged, we notice that PTS is the best performer among the various competitors when DT is used as a classifier. On the contrary, PTC worsens its ranking. In other terms, PIVOTTREE excels in different settings according to the number k of pivots extracted: when k is small, a Decision Tree is best; and when k is large, then kNN is best.

Sensitivity Analysis. Figure 7 reports a sensitivity analysis on PIVOTTREE used as pivot selector (PTS). In particular, we observe the average F1-score

p	redictor		Decision Tree								kNN					
selector		-	RND	RNC	KMS	KMD	EBL	PTS	PTC	-	RND	RNC	KMS	KMD	EBL	PTS
ng	cifar	.75	.39	.40	.40	.41	.41	.42	.11	.87	.86	.87	.88	.88	.88	.80
I	mnist	.68	.44	.41	.53	.44	.42	.37	.29	.97	.96	.96	.96	.96	.96	.95
	breast	.95	.93	.94	.93	.94	.94	.96	.95	.96	.95	.95	.95	.96	.97	.95
	compas	.50	.46	.48	.46	.48	.47	.52	.49	.46	.47	.48	.48	.47	.49	.47
	german	.58	.52	.53	.56	.49	.44	.50	.48	.67	.58	.58	.59	.58	.60	.58
	heloc	.70	.65	.65	.65	.65	.65	.68	.66	.67	.66	.65	.67	.67	.67	.66
	iris	1.0	.94	.92	.98	.96	.95	.95	.92	1.0	.95	1.0	1.0	1.0	1.0	.98
llar	page	.88	.83	.83	.87	.87	.88	.84	.87	.90	.88	.88	.90	.89	.89	.88
q	diva	.79	.60	.60	.58	.55	.59	.63	.63	.76	.72	.71	.69	.70	.71	.73
[Ja	sonar	.74	.71	.70	.71	.71	.74	.72	.59	.94	.82	.84	.82	.83	.89	.84
	vert.	.72	.66	.69	.66	.69	.66	.66	.68	.73	.69	.69	.73	.76	.74	.78
	wine	.20	.19	.19	.19	.19	.18	.19	.18	.37	.35	.35	.35	.35	.35	.36
	imdb	.70	.72	.71	.75	.74	.77	.78	.78	.78	.76	.75	.79	.79	.81	.81
-	lyrics	.66	.68	.68	.68	.68	.70	.70	.70	.71	.69	.69	.70	.68	.71	.68
ex Ex	news	.12	.15	.16	.19	.18	.16	.19	.01	.69	.48	.50	.58	.55	.58	.40
5	tgpt	.84	.79	.79	.80	.82	.84	.79	.84	.92	.87	.86	.88	.88	.90	.88
	vicuna	.63	.55	.55	.52	.59	.57	.64	.59	.68	.67	.67	.66	.69	.72	.71
TimeS	devices	.25	.32	.30	.30	.34	.36	.38	.34	.49	.46	.44	.48	.49	.46	.47
	worms	.54	.52	.52	.57	.56	.61	.56	.56	.60	.58	.61	.60	.58	.51	.70
	ecg	.52	.49	.51	.50	.50	.46	.51	.51	.57	.54	.54	.55	.55	.55	.56
	gun	.80	.77	.77	.76	.77	.77	.71	.74	.91	.87	.86	.89	.89	.88	.84
	wafer	.90	.93	.93	.94	.93	.93	.92	.93	.99	.97	.98	.98	.98	.98	.98
	avg	.66	.60	.60	.62	.61	.61	.62	.58	.76	.72	.72	.73	.73	.74	.73
	std	.23	.22	.22	.22	.22	.23	.22	.27	.18	.19	.19	.18	.19	.19	.19
	rank		5.0	4.6	4.0	3.6	3.7	3.1	3.9		4.8	4.1	3.0	3.1	2.2	3.5

Table 4. Model performance as F1-score with models limited to 20 pivots.

among all datasets with error bars indicating the standard deviations when varying the maximum number of pivots in ranges from 10 to 20, from 20 to 30, etc. Two lines are reported to differentiate the performance between datasets with 2 or 3 classes, i.e., $c \in [2,3]$, versus datasets with 5 to 10 classes, i.e., $c \in [5,10]$. We leave as a future study a sensitivity analysis of datasets with more than 10 classes. The results show that, both for DT and kNN, for datasets with few

classes, the performance is stable independently of the number of pivots selected. Thus, extracting a limited number of highly discriminative and representative pivots can guarantee high performance and high simplicity. On the other hand, for datasets with more than five classes, the results are less stable, and we observe an increase in performance, especially when using kNN, as the DT we relied on is limited by the maximum depth of four, thus practically being limited by its depth and not exploiting all the possible pivots. As a consequence, for datasets with a high number of classes, the tuning of the number of pivots k extracted with PIVOTTREE should be carefully addressed, and it should consider a high number potentially limiting the final interpretability of the predictive model.

Although time complexity is not the primary focus of this paper, here we also report training runtime (in seconds). As example for small datasets, *breast* and *ecg* datasets present fitting runtimes respectively of 4.34 s and 8.29 s. In contrast, *tgpt* and *cifar* show higher training times of 24.91 s and 60.70 s. Larger datasets, both in terms of instances and dimensions, require longer training times, due to the need of finding pivots within a bigger pool. For example, *lyrics* requires 458.81 s for training. In all cases mentioned, prediction times are relatively fast, with all predictions taking under 24.10 s, which is the time needed to perform predictions for the *imdb* test set.

5 Conclusions

We have introduced PIVOTTREE, an interpretable tree-based pivot selection model aimed at facilitating the training of effective interpretable case-based predictive models. In PIVOTTREE, exemplary instances, named *pivots*, guide the construction of a similarity-based case-based model where explanations are a hierarchy of prototypical instances. By design, PIVOTTREE is both a pivot selector and a prediction model, enabling, independently, both the extraction of relevant instances and the construction of an interpretable predictive model. PIVOTTREE is a *data-agnostic* model, which can be seamlessly applied to various data modalities, including tabular data, text, time series, and images. In a wide array of experiments, PIVOTTREE has shown to be on par with state-of-the-art approaches while often retaining lower complexity and higher interpretability.

Given its inherent flexibility, PIVOTTREE lends itself to several future improvements: different data encodings, e.g. TABPFN [23] or ROCKET [15], may further improve instance representation, and thus similarity estimation; joint optimization of pivot selection and case-based reasoning, which is currently decoupled in pivot selection, and tree induction; use of more sophisticated casebased reasoning models; adaptation for other data types such as mobility trajectories [30], and evaluation of the privacy exposure lead by pivots [34]. Another avenue of research lies in integrating prior knowledge or human supervision into prototype learning, as human-machine collaboration could improve the classifier's accuracy and interpretability, as suggested and investigated in [33, 42]. Furthermore, future avenues of research also include assessing PIVOTTREE's interpretability from a human-centric perspective, validating its performance through evaluation schema designed for prototype-based explanations, as described for images in [13,28], time-series in [33], and texts in [12,24]. As such, an extensive comparison of PIVOTTREE's performance and explainability could be conducted against deep learning-based *representations* of the prototypes across different modalities, as well as through feature-based explainability techniques.

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Disentangled Counterfactual Graph Augmentation Framework for Fair Graph Learning with Information Bottleneck

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Abstract. Graph Neural Networks (GNNs) are susceptible to inheriting and even amplifying biases within datasets, subsequently leading to discriminatory decision-making. Our empirical observation reveals that the inconsistent distribution of sensitive attributes conditioned on labels significantly contributes to unfairness. To mitigate this problem, we suggest rectifying this inconsistency of the original dataset through a counterfactual augmentation strategy. Existing methods usually generate counterfactual samples from an entangled representation space, which fail to distinguish the different dependencies on sensitive attributes. Thus, we propose a novel disentangled counterfactual graph augmentation method based on the Information Bottleneck theory, named Fair Disentangled Graph Information Bottleneck (FDGIB). Specifically, FDGIB embeds graphs into two disentangled representation spaces: sensitive-related and sensitive-independent. By satisfying three conditions, FDGIB theoretically guarantees the disentanglement of different sensitive dependencies. We acquire credible counterfactual augmented graphs to facilitate consistency in data distribution and generate fair representations. FDGIB serves as a plug-and-play preprocessing framework that can collaborate with any GNNs. We validate the effectiveness of our model in promoting fairness learning through extensive experiments. Our source code is available at https://github.com/Evanlyf/FDGIB.

Keywords: Graph fairness learning \cdot Information bottleneck \cdot Counterfactual graph \cdot Disentangled representation learning

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L. Zheng and J. Wang—Contribute equally to this work. The appendix is available at the source code link.

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Fig. 1. (a): Empirical investigation conducted on Bail examines the impact of inconsistent distribution of sensitive attributes conditioned on labels. (b): An intuitive example for the impact of disentanglement on counterfactual augmentation.

1 Introduction

Graph-structured data permeates myriad fields of the real world, such as social networks [12,21], recommendation systems [17,38], and traffic forecasting [13,16]. An increasing number of graph mining algorithms [2,32] have been proposed to gain a better understanding of graph data, they strive to map high-dimensional graph nodes into a latent embedding space and demonstrate superior performance on various tasks. Nevertheless, the design of most graph mining algorithms does not contemplate fairness, thus leading to ethical issues. Moreover, the discrimination is further intensified in graph learning since GNNs inherently aggregate features from neighboring nodes, inadvertently aggravate the leakage of sensitive attributes, and amplify bias through network topology [7,11,31].

To further study the possible springhead of discriminations in GNNs, we conduct empirical studies on a vanilla GCN [19] and observe that the inconsistent distribution of sensitive attributes conditioned on labels is a critical cause responsible for the unfairness of GNNs. Specifically, we employ a quantitative metric Δ_r to measure the inconsistency of the sensitive attributes' distribution among different demographic groups with different labels. Given a binary sensitive attribute S and a binary label Y, Δ_r is defined as $\Delta_r = r(Y = 1) - r(Y = 0)$ where $r(Y = y) = \frac{p(S=1|Y=y)}{p(S=0|Y=y)}$ is the probability ratio of S given label Y = y. $\Delta_r = 0$ refers to the sensitive attribute sharing the consistent distribution among groups with different labels. We demonstrate the empirical results in Fig. 1(a)where the y-axis denotes the unfairness metrics (Δ_{DP} and Δ_{EO}) and the xaxis denotes the inconsistency metric Δ_r . We observe that GCN achieves lower unfairness metrics when $|\Delta_r|$ decreases, indicating a positive correlation between inconsistency and unfairness. Inspired by the empirical observation, we suggest generating counterfactual graphs by flipping sensitive attributes to rectify the inconsistent distributions. Theoretical analysis demonstrates that the counterfactual augmentation strategy can ensure distribution consistency.

In addition to rectifying the inconsistent distribution of sensitive attributes, the counterfactual graph augmentation strategy exhibits remarkable flexibility as a versatile preprocessing module [26,27] adaptable to various scenarios. Yet, although existing counterfactual graph augmentation methods demonstrate impressive performance to facilitate fairness, they overlook the disentanglement of different dependencies on sensitive attributes, resulting in generating graphs from an entangled representation space. The entanglement may lead to sub-optimal performance since it may yield undesirable out-of-distribution (OOD) counterfactual samples following distributions that conflict with the real dataset. We provide an intuitive explanation for this problem in Fig. 1(b). For instance, height distribution varies by gender, whereas education level distribution remains consistent across genders in the dataset. The entangled methods fail to distinguish the different dependencies on gender and may lead to counterfactual samples where the males and females share the same height distribution or different education distribution. The OOD counterfactual samples may mislead the graph learning and lead to sub-optimal performance on downstream tasks.

To tackle the above problem, we propose to generate more realistic counterfactual samples by taking into account the different dependencies on the sensitive attributes. Specifically, we propose to explicitly decompose the features of a graph into two disentangled spaces: the *sensitive-related* representation space and the *sensitive-independent* representation space. The sensitive-related space encapsulates the sensitive-related attributes (*e.g.*, the height and weight) which change synergetically when the sensitive attributes are altered. The sensitiveindependent space encapsulates the sensitive-irrelevant attributes (*e.g.*, the education level) and remains invariant when the sensitive attributes change. Within the disentangled spaces, we can align the distributions of sensitiverelated attributes with the sensitive attributes while preserving the distribution of sensitive-independent attributes invariant in the counterfactual samples.

Two challenges hinder the generation of counterfactual graphs in disentangled representation spaces. (i) How to learn the two disentangled representation spaces corresponding to sensitive attributes? Inspired by the Information Bottleneck (IB) principle [3,28], we propose a novel method named Fair Disentangled Graph Information Bottleneck (FDGIB). FDGIB achieves the disentanglement with three conditions: minimal sufficiency, independence, and joint sufficiency. Intuitively, the three conditions squeeze the sensitive-correlated information into the sensitive-related representation space while preserving the sensitive-invariant information into the sensitive-independent space, sharing a similar ideology with IB. (ii) How to optimize the mutual information (MI) involved in FDGIB? The MI is notoriously intractable to optimize for high-dimensional data since it involves integral on the unknown prior distributions. To address this, we derive tractable bounds for the MI terms and experimentally verify the performance.

The main contributions are summarized as follows: (i) We propose a novel disentangled counterfactual graph augmentation strategy to rectify the biased distribution of the original dataset, which serves as a plug-and-play preprocessing module and can be integrated with any GNNs. (ii) We introduce the IB principle into the disentangled counterfactual graph augmentation, which leads to a novel method named Fair Disentangled Graph Information Bottleneck (FDGIB). Further in-depth theoretical analysis demonstrates that FDGIB can achieve

certifiable disentanglement. (iii) Extensive experimental results demonstrate that our method achieves state-of-the-art fairness performance while preserving satisfactory model utility over various benchmark datasets.

2 Preliminaries

2.1 Notation

Given a graph by $\mathcal{G} = (\mathcal{V}, \mathcal{E}, \mathbf{X}, \mathbf{A})$, where $\mathcal{V} = \{v_1, \ldots, v_n\}$ is the set of n nodes, \mathcal{E} is the set of edges, $\mathbf{X} \in \mathbb{R}^{n \times d}$ is the node features matrix. $\mathbf{A} \in \{0, 1\}^{n \times n}$ is the adjacency matrix, where $\mathbf{A}_{ij} = 1$ if the edge between v_i and v_j exists, otherwise $\mathbf{A}_{ij} = 0$. Without loss of generality, we consider an undirected and unweighted graph where each node has a binary-sensitive attribute (e.g. gender, age), and we focus on the counterfactual augmentation at the node level. For simplicity of analysis, we consider the corresponding ego graph \mathcal{G}_i (a k-hop subgraph) for node i since k-layer GNNs only consider the k-hop neighbor nodes. The adjacency matrix and feature matrix of the ego graph are denoted as \mathbf{A}_i and \mathbf{X}_i , separately. The whole graph can be regarded as a set of ego graphs $\mathcal{G} = \{\mathcal{G}_1, \mathcal{G}_2, \cdots, \mathcal{G}_n\}$. We denote the sensitive attribute and the label of node i by $S \in \{0, 1\}$ and $Y \in \{0, 1\}$, separately. Let (R, D) be a pair of disentangled representations of \mathcal{G}_i , where R is the representation of sensitive-related information inherited from \mathcal{G}_i , while D encodes sensitive-independent information. Counterfactual data is annotated with the superscript cf, *i.e.*, \mathcal{G}_i^{cf} , \mathbf{X}_i^{cf} , \mathbf{A}_i^{cf} , \mathbf{R}_i^{cf} and D^{cf} .

2.2 Fairness Metrics

To measure fairness, we adopt two commonly used fairness concepts: demographic parity (Δ_{DP}) and equal opportunity (Δ_{EO}) . Demographic parity assesses whether the algorithm's outcomes \hat{Y} are independent of demographic attributes, while equal opportunity evaluates whether the algorithm provides equal chances of positive outcomes for individuals with similar characteristics.

$$\Delta_{DP} = \left| p\left(\hat{Y} = 1 \, S = 0 \right) - p\left(\hat{Y} = 1 \, S = 1 \right) \right|,$$

$$\Delta_{EO} = \left| p\left(\hat{Y} = 1 | Y = 1, S = 0 \right) - p\left(\hat{Y} = 1 | Y = 1, S = 1 \right) \right|.$$

These fairness metrics provide quantitative measures to evaluate the degree of fairness achieved by the algorithm.

2.3 Information Bottleneck

Built on the notion of mutual information (MI), the Information Bottleneck (IB) compresses the source data to keep task-relevant information and discard task-irrelevant information. Specifically, MI quantifies the relationship between two random variables. The MI between random variables X and Z is formulated as:

$$I(X;Z) = \int_x \int_z p(x,z) \log \frac{p(x,z)}{p(x)p(z)} dxdz.$$

Given the input data X and label Y, the IB method aims to compress X to a "bottleneck" latent representation Z, which keeps the most task-relevant information and discards the task-irrelevant information in X. It can be formally stated in terms of the constrained optimization problem:

$$\min_{Z} \mathcal{L}_{IB} = I(Z;Y) - \beta I(X;Z),$$

where β is the Lagrange multiplier which controls the trade-off and is typically set in [0, 1].

3 Counterfactual Graph Augmentation

To measure the inconsistency of sensitive attributes' distributions conditioned on labels, we adopt the quantitative metric Δ_r stated before, we define the notions of consistent graph and inconsistent graph as follows.

Definition 1. Given a graph consisting of n ego graphs corresponding to n nodes, i.e., $\mathcal{G} = \{\mathcal{G}_1, \mathcal{G}_2, \dots, \mathcal{G}_n\}$, we call \mathcal{G} as a consistent graph if and only if $\Delta_r = 0$. Otherwise, we call \mathcal{G} an inconsistent graph.

Just as we analyze in Fig. 1(a), GNNs learned on the inconsistent graph may suffer more severe unfairness compared to the consistent graph. As a consequence, we suggest rectifying the inconsistent graph to be consistent with the counterfactual graph augmentation strategy. Formally, for ego graph \mathcal{G}_i with label Y = y and sensitive attribute S = s, we generate its corresponding counterfactual sample \mathcal{G}_i^{cf} with the same label Y = y and flipped sensitive attribute S = 1 - s. We theoretically prove this intuition in Theorem 1.

Theorem 1. Suppose a graph consisting of n ego graphs corresponding to n nodes, i.e., $\mathcal{G} = \{\mathcal{G}_1, \mathcal{G}_2, \dots, \mathcal{G}_n\}$. For any inconsistent graph \mathcal{G} with $|\Delta_r| > 0$, we can achieve a consistent graph with augmented counterfactual samples for every involved ego graph, i.e., $\mathcal{G}^c = \{\mathcal{G}_1, \mathcal{G}_2, \dots, \mathcal{G}_n\} \cup \{\mathcal{G}_1^{cf}, \mathcal{G}_2^{cf}, \dots, \mathcal{G}_n^{cf}\}$.

Proof. Since we generate counterfactual samples for every ego graph with flipped sensitive attributes and the same labels, we have p(S = 1|Y = y) = p(S = 0|Y = y) for \mathcal{G}^c . Then we have r(Y = 0) = r(Y = 1), and thus \mathcal{G}^c is consistent.

The above theorem demonstrates the rationality of our counterfactual augmentation strategy. There are some prior works [5,26] that focus on counterfactual graph augmentation. However, they generate counterfactual samples from an entangled representation space, potentially yielding undesirable OOD counterfactuals that follow distributions conflicting with the actual distributions. To tackle the issues, we propose to embed the ego graphs into two disentangled representation spaces: the sensitive-related representation R and the sensitiveindependent representation D. R encapsulates the sensitive-related information involved in the ego graph and changes synergetically when S is flipped, while D remains invariant. To generate the counterfactual samples, we flip S and



Fig. 2. Overall framework of our proposed FDGIB. We extract ego graphs for each node before training and perform a two-stage training strategy.

reconstruct the counterfactual samples from the counterfactual sensitive-related representation space R^{cf} and the invariant sensitive-independent representation space D. The disentangled counterfactual samples can rectify the inconsistency of the original dataset and facilitate the fairness of downstream tasks.

In Fig. 2, we provide an overview of our proposed FDGIB. Specifically, our method works in a two-stage fashion. (i) First stage: disentangled counterfactual augmenter learning. We employ an encoder-decoder architecture to learn a counterfactual graph augmenter. The encoders embed the ego graphs into two disentangled representation spaces to capture different sensitive dependencies. The decoder reconstructs the ego graphs from the disentangled representation space. (ii) Second stage: counterfactual augmentation and fair graph learning. We flip the sensitive attributes of the ego graphs and embed them into the disentangled representation spaces with the encoder. Then we can generate credible counterfactual ego graphs with the disentangled representations. Finally, we learn fair GNNs with the augmented counterfactual graphs.

4 Disentangled Counterfactual Augmenter Learning

In this section, we introduce how we learn the disentangled counterfactual augmenter in the first stage. We introduce how to achieve the disentanglement with theoretical analysis and elucidate the details of our optimization strategy.

4.1 Definition of FDGIB

To learn the disentangled counterfactual augmenter, we adopt the ideology of IB which learns robust representations by preserving the task-related information

while discarding task-irrelevant nuisance. We propose a novel method that can generate counterfactual ego graphs from a disentangled representation space, named Fair Disentangled Graph Information Bottleneck (FDGIB).

Definition 2. (Fair Disentangled Graph Information Bottleneck, FDGIB) Given an ego graph \mathcal{G}_i , let S denotes sensitive attributes, R and D denote the corresponding S-related and S-independent representation. Fair Disentangled Graph Information Bottleneck requires R and D to satisfy the following conditions.

- 1. Minimal sufficiency: R should be minimally sufficient for S, i.e., $I(R; \mathcal{G}_i | S) = 0$ and I(R; S) = H(S);
- 2. Independence: D should be independent of S, i.e., I(D; S) = 0;
- 3. Joint sufficiency: R and D should be sufficient for the ego graph, i.e., $I(R, D; \mathcal{G}_i) = H(\mathcal{G}_i)$.

The main motivation of FDGIB is to isolate the sensitive-related information into R and squeeze the other information into D. Intuitively, the minimal sufficiency of FDGIB requires that R preserves all sensitive-related information (i.e., I(R; S) = H(S)) while discarding sensitive-irrelevant information $(i.e., I(R; G_i|S) = 0)$, which shares a similar idea with IB. The independence condition requires D to be independent of S and remain invariant when S is altered. The joint sufficiency guarantees that we can generate the counterfactual example from the joint space since R and D embed all information of \mathcal{G}_i . Furthermore, we provide an in-depth theoretical analysis to investigate the rationality of FDGIB, which indicates that FDGIB achieves certifiable disentanglement.

Theorem 2. Given an ego graph \mathcal{G}_i , let R an D denote the corresponding sensitive-related representation and sensitive-independent representation which satisfy the conditions of FDGIB. We have(proof is provided in the appendix):

- Mutual disentanglement: R and D are disentangled, i.e., I(R; D) = 0;
- Sensitive disentanglement: Given any attribute F, R can embed its all correlated information with sensitive attributes S, i.e., I(F;S) = I(R;F) while D can encapsulate the remaining information which is irrelevant with the sensitive attributes, i.e., H(F|S) = I(D;F).

The mutual disentanglement demonstrates that FDGIB achieves two disentangled representation spaces. The sensitive disentanglement further proves that for any attributes, R can encapsulate its correlated information with sensitive attributes, while D embeds the remaining information. The theorem provides theoretical support for the rationality of our FDGIB.

4.2 Optimization of FDGIB

In this subsection, we introduce how to generate desirable R and D. Formally, R and D are learned by two GNN encoders, *i.e.*, $q_{\theta_R}(R|\mathcal{G}_i)$ and $q_{\theta_D}(D|\mathcal{G}_i)$. Our goal is to optimize the encoders to learn desirable representations that satisfy the three conditions of FDGIB. The detailed proof is shown in the appendix.

Minimal Sufficiency. We can achieve minimal sufficiency by minimizing $I(R; \mathcal{G}_i|S)$ while maximizing I(R; S), *i.e.*,

$$\min_{\theta_R} \mathcal{L}_{ms} = -I(R;S) + \lambda I(R;\mathcal{G}_i|S) = -I(R;S) + \beta I(R;\mathcal{G}_i), \tag{1}$$

where λ is the Lagrangian multiplier that trade-off two terms, $\beta = \lambda/(1 + \lambda)$ is the reformulation of λ . Then we formulate tractable bounds separately.

To minimize $I(R; \mathcal{G}_i)$, we adopt Contrastive Log-ratio Upper Bound (CLUB) [6] as an estimator of MI upper bound; As for I(R; S), we introduce $q_{\theta_c}(S|R)$ as a variational approximation to p(S|R) and derive a tractable lower bound, *i.e.*,

$$I(R;\mathcal{G}_i) \leq I_{club}(R;\mathcal{G}_i) = \mathbb{E}_{p(R,\mathcal{G}_i)} \log q_{\theta_R}(R|\mathcal{G}_i) - \mathbb{E}_{p(R)p(\mathcal{G}_i)} \log q_{\theta_R}(R|\mathcal{G}_i),$$

$$I(R;S) \geq \mathbb{E}_{p(R,S)} \log q_{\theta_{c1}}(S|R) + H(S),$$

where $q_{\theta_R}(R|\mathcal{G}_i)$ is an approximation to $p(R|\mathcal{G}_i)$ with parameter θ_R , H(S) is a constant that can be ignored. In practice, $q_{\theta_{c1}}(S|R)$ can be viewed as a classifier parameterized with θ_{c1} , which takes R as input to predict S.

Independence. The independence of FDGIB requires that D is independent of S. To achieve this, we propose to optimize D as follows:

$$\min_{\theta_D} \mathcal{L}_{ind} = I(D; S).$$
(2)

Just like I(R; S) analyzed above, I(D; S) can be approximated by a lower bound $\mathbb{E}_{p(D,S)} \log q_{\theta_{c2}}(S|D)$ where $q_{\theta_{c2}}$ is a classifier parameterized by θ_{c2} . Nonetheless, different from the minimal sufficiency in Eq. (1), the independence requires minimization of I(D; S) instead of maximization. As a consequence, we adopt an adversarial training strategy to minimize the MI term:

$$\min_{\theta_D} \max_{\theta_{c2}} \hat{\mathcal{L}}_{ind} = \mathbb{E}_{q_{\theta_D}(D|\mathcal{G}_i)} \mathbb{E}_{p(D,S)} \log q_{\theta_{c2}}(S|D).$$
(3)

The above objective is a min-max optimization problem. Intuitively, the classifier $q_{\theta_{c2}}(S|D)$ attempts to predict the sensitive attributes from representation D, while the encoder $q_{\theta_D}(D|\mathcal{G}_i)$ aims to yield representations embedding no sensitive-related information.

Joint Sufficiency. The joint sufficiency requires that R and D should be sufficient for the ego graph \mathcal{G}_i . To maximize $I(R, D; \mathcal{G}_i)$, we:

$$\min_{\theta_R,\theta_D} \mathcal{L}_{js} = -I(R,D;\mathcal{G}_i).$$
(4)

Similar to I(R; S), we can derive a lower bound for $I(R, D; \mathcal{G}_i)$, where $q_{\theta_{dec}}(\mathcal{G}_i|R, D)$ serves as a decoder parameterized by θ_{dec} to reconstruct \mathcal{G}_i . Intuitively, if we can reconstruct the ego graphs from the representation spaces, we can say that the representations are jointly sufficient for the ego graphs.

In terms of graph features, an MLP decoder is used, taking the concatenated [R; D] as input and producing corresponding reconstructed features. The reconstruction quality is assessed using MSE loss. To reconstruct the adjacency matrix, we first feed the concatenated representation [R; D] into an MLP and then compute the inner product of the output followed by a sigmoid function. The representations are optimized by the cross-entropy loss. Consequently, the joint sufficiency loss consists of two components.

The Whole Objective of FDGIB. With the above optimization strategy, the whole objective of FDGIB can be formulated as:

$$\min_{\Theta} \mathcal{L}_{FDGIB} = \mathcal{L}_{js} + \alpha \mathcal{L}_{ms} + \gamma \mathcal{L}_{ind}, \tag{5}$$

where Θ collects all trainable parameters. The encoder-decoder architecture empowers FDGIB's capability to reconstruct credible counterfactual samples from the disentangled representation space. Based on the augmented counterfactual samples, we can rectify the inconsistent graph and learn fair GNNs.

5 Counterfactual Augmentation and Fair Graph Learning

In this section, we elucidate how we augment the original graph with FDGIB and learn fair GNNs in the second stage. Specifically, we first introduce the details of counterfactual augmentation with FDGIB; Then, we demonstrate how to learn fair GNNs based on the augmented graph. Finally, we summarize the whole pipeline of our method in Algorithm 1, details can be seen in the appendix.

5.1 Counterfactual Augmentation with FDGIB

Given an ego graph \mathcal{G}_i with sensitive attributes S = s and its corresponding disentangled representations R and D, we can generate its counterfactual samples \mathcal{G}_i^{cf} with the help of FDGIB. Formally, we first flip the sensitive attributes as S = 1 - s and then embed the flipped ego graph \mathcal{G}'_i into the sensitive-related representation space $R^{cf} \sim q_{\theta_R}(R|\mathcal{G}'_i)$, where q_{θ_R} is the encoder learned by FDGIB. The counterfactual sensitive-related representation R^{cf} captures the sensitive related information and varies with the flip of the sensitive attributes. Then, the original ego graph is embedded into the sensitive-independent representation space $D \sim q_{\theta_D}(D|\mathcal{G}_i)$. The sensitive-independent representation D encapsulates the invariant information when S is flipped. Based on the R^{cf} and D, we can generate the counterfactual ego graph from the disentangled representation space, *i.e.*, $\mathcal{G}_i^{cf} \sim q_{\theta_{dec}}(\mathcal{G}_i|R^{cf}, D)$. The generation process can be summarized as:

$$\mathcal{G}_{i}^{cf} \sim q_{\theta_{dec}}(\mathcal{G}_{i}|R^{cf}, D)q_{\theta_{D}}(D|\mathcal{G}_{i})q_{\theta_{R}}(R^{cf}|\mathcal{G}_{i}')p(\mathcal{G}_{i}'|S=s, \mathcal{G}_{i})$$

Following the strategy, we can achieve a consistent graph $\mathcal{G}^c = \{\mathcal{G}_1, \mathcal{G}_2, \cdots, \mathcal{G}_n\} \cup \{\mathcal{G}_1^{cf}, \mathcal{G}_2^{cf}, \cdots, \mathcal{G}_n^{cf}\}$ by generating counterfactual samples for every ego graph.

5.2 Fair Graph Learning with FDGIB

FDGIB provides a plug-and-play framework for fair graph learning. For any GNN, we can achieve debiased representations by employing the counterfactual ego graphs, and make fair predictions based on the debiased representations. Specifically, given an ego graph \mathcal{G}_i and its counterfactual ego graph \mathcal{G}_i^{cf} , we can learn representations with the given GNN model, *i.e.*,

$$\mathbf{z}_i = \text{GNN}(\mathcal{G}_i; \theta_f), \quad \mathbf{z}_i^{cf} = \text{GNN}(\mathcal{G}_i^{cf}; \theta_f), \quad \mathcal{L}_{sim} = \frac{1}{n} \sum_{i=1}^n d(\mathbf{z}_i, \mathbf{z}_i^{cf}),$$

where θ_f is the parameters of the GNN. To promote the fairness of the GNN model, we propose to minimize the distance between representations \mathbf{z}_i and \mathbf{z}_i^{cf} because the representations from the original ego graph and counterfactual ego graph should be similar. Furthermore, the representations can be trained in a supervised fashion with the cross-entropy loss:

$$\mathcal{L}_{pred} = \frac{1}{n} \sum_{i=1}^{n} \kappa l \left(f_{\phi} \left(\mathbf{z}_{i} \right), y_{i} \right) + (1 - \kappa) l \left(f_{\phi} \left(\mathbf{z}_{i}^{cf} \right), y_{i} \right),$$

where $l(\cdot)$ is the cross-entropy loss function, f_{ϕ} is a classifier for downstream tasks, y_i is the ground truth of ego graph \mathcal{G}_i and κ is a hyper-parameter.

Finally, the overall loss function for fair graph learning is:

$$\mathcal{L}_{fair} = \mathcal{L}_{pred} + \mu \mathcal{L}_{sim},\tag{6}$$

where μ is a hyperparameter controlling the weight of similarity regularization.

6 Experiments

In this section, we conduct extensive experiments to evaluate the performance of our FDGIB. We initially provide a comprehensive overview of our experiment setting. Subsequently, we engage in a thorough discussion of the experimental results. Specifically, our objective is to answer the following questions:

- (RQ 1) Whether FDGIB benefits the fairness of downstream tasks while maintaining the model utility?
- (RQ 2) Whether FDGIB achieves disentanglement in representation space?
- (RQ 3) Whether the three conditions of FDGIB all benefit the fairness of downstream tasks?

6.1 Experiment Settings

Datasets and Evaluation Metrics. We conduct experiments on four widely used benchmark datasets, namely Bail [18], Income [10], Pokec-z [7], and Pokecn [7]. The details of the datasets can be found in the appendix. We assess the proposed framework in terms of two aspects: model utility and fairness. We are dedicated to achieving fairness while ensuring model utility, aiming to strike a favorable trade-off between them. For model utility, we use Accuracy, F1score, and AUC-ROC to measure node classification performance. For fairness performance, we utilize Δ_{DP} and Δ_{EO} , where smaller values indicate a fairer model.

The second seco	Algorithm	1.	Training	Algorithm	of FDGIB
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Input: $\mathcal{G} = (\mathcal{V}, \mathcal{E}, \mathbf{X}, \mathbf{A}), Y, \alpha, \beta, \gamma, \kappa, n_epoch_1, n_epoch_2;$

Output: Learned encoder q_{θ_R} , q_{θ_D} and predictions \hat{Y} ;

- 1: Extract ego graphs for each node $\mathcal{G} = \{\mathcal{G}_1, \mathcal{G}_2, \cdots, \mathcal{G}_n\}; \#$ Stage1:Disentangled counterfactual augmenter learning
- 2: for epoch $\leftarrow 1$ to n_epoch_1 do
- 3: Compute the representations with two encoders: $R \sim q_{\theta_R}(R|\mathcal{G}_i), D \sim q_{\theta_D}(D|\mathcal{G}_i);$
- 4: Reconstruct the ego graph from the disentangled representation spaces with a decoder: $\mathcal{G}_i \sim q_{\theta_{dec}}(\mathcal{G}_i|R, D);$
- 5: Update the parameters of FDGIB according to the objective in Eq. (5);
- 6: **end for**;

$\# {\tt Stage 2: Counterfactual} \ {\tt augmentation} \ {\tt and} \ {\tt fair \ graph} \ {\tt learning}$

- 7: for epoch $\leftarrow 1$ to n_epoch_2 do
- 8: $\mathcal{G}'_i \leftarrow$ Flip sensitive attributes of \mathcal{G}_i ;
- 9: Compute the disentangled representations from the flipped ego graph and original ego graph, separately: $R^{cf} \sim q_{\theta_R}(R|\mathcal{G}'_i), D \sim q_{\theta_D}(D|\mathcal{G}_i);$
- 10: Reconstruct the counterfactual ego graph from the disentangled representation space: $\mathcal{G}_{i}^{cf} \sim q_{\theta_{dec}}(\mathcal{G}_{i}^{cf}|R^{cf},D);$
- 11: end for
- 12: Learn fair GNNs based on the original ego graphs and the counterfactual graphs, *i.e.*, $\mathcal{G}^c = \{\mathcal{G}_1, \mathcal{G}_2, \cdots, \mathcal{G}_n\} \cup \{\mathcal{G}_1^{cf}, \mathcal{G}_2^{cf}, \cdots, \mathcal{G}_n^{cf}\}$ according to Eq. (6);

Baselines and GNN Backbones. FDGIB is compared with several stateof-the-art fair node representation learning models. We categorize them into two groups: algorithm-based and augmentation-based. (1) Algorithm-based: FairGNN [7] and BIND [10]. (2) Augmentation-based: NIFTY [1], EDITS [9], GEAR [26], FairVGNN [31], and CAF [14]. Additionally, our method serves as a plug-and-play module and is applicable to any downstream GNN classifiers. We conduct a comparative analysis of our method against four backbone GNN models: GraphSAGE [15], GCN [19], GAT [29], and Jumping Knowledge (JK) [35].

Implementation Details. We conducted the training using NVIDIA GeForce RTX 3090 Ti GPUs. We use the default train/valid/test split in [26]. Experimental results are averaged over 3 repeated executions with 3 different seeds to remove any potential initialization bias. The Adam optimizer is adopted to optimize the neural networks with a learning rate of 0.001. We adopt a 2-layer GCN to implement the encoder q_{θ_R} and q_{θ_D} and two MLPs to reconstruct the graph. We perform grid searches to identify the best hyperparameters.

6.2 Prediction Performance and Fairness (RQ 1)

To answer RQ1, we conduct experiments on benchmark datasets with comparison to baselines. We present the experimental results of the average performance with standard deviation across four datasets in Table 1. GraphSAGE is used as the backbone. We can observe:

Dataset	Metrics	FairGNN	BIND	NIFTY	EDITS	GEAR	FairVGNN	CAF	FDGIB
Bail	Accuracy (\uparrow)	86.67 ± 0.52	86.49 ± 0.85	$\underline{87.12\pm0.68}$	84.42 ± 2.87	86.34 ± 0.21	$\textbf{88.41} \pm \textbf{1.29}$	86.68 ± 1.49	86.97 ± 0.23
	F1-score (\uparrow)	81.49 ± 0.35	81.16 ± 0.93	80.66 ± 0.99	77.83 ± 3.79	80.44 ± 0.27	$\textbf{83.58} \pm \textbf{1.88}$	$\underline{82.33 \pm 1.18}$	82.05 ± 0.20
	AUCROC (↑)	91.55 ± 0.16	90.64 ± 0.70	91.31 ± 0.45	89.07 ± 2.26	91.06 ± 0.07	$\underline{91.56 \pm 1.71}$	91.34 ± 1.29	91.74 ± 0.09
	$\Delta_{DP}(\downarrow)$	2.62 ± 0.62	1.15 ± 0.78	6.51 ± 0.11	3.74 ± 3.54	5.02 ± 0.19	$\underline{1.14 \pm 0.67}$	2.19 ± 0.20	$\textbf{0.99} \pm \textbf{1.03}$
	$\Delta_{EO}(\downarrow)$	2.81 ± 0.32	2.13 ± 1.17	5.27 ± 0.78	4.46 ± 3.50	3.42 ± 1.10	1.69 ± 1.13	$\underline{1.65 \pm 1.51}$	1.06 ± 0.52
Income	Accuracy (↑)	74.79 ± 0.20	77.62 ± 1.97	77.00 ± 0.12	73.08 ± 0.03	71.83 ± 0.77	$\textbf{80.76} \pm \textbf{1.87}$	$\underline{78.13 \pm 0.78}$	75.82 ± 2.07
	F1-score (\uparrow)	56.92 ± 0.10	55.24 ± 1.85	53.87 ± 0.05	53.50 ± 0.07	55.16 ± 0.44	54.49 ± 0.86	54.26 ± 1.21	$\underline{56.30\pm0.07}$
	AUCROC (↑)	81.02 ± 0.10	81.44 ± 0.85	80.22 ± 0.12	78.76 ± 0.00	82.75 ± 0.34	80.10 ± 0.44	80.14 ± 0.95	$\underline{82.09\pm0.35}$
	$\Delta_{DP}(\downarrow)$	8.42 ± 0.51	7.93 ± 7.24	13.28 ± 0.14	$\underline{7.48\pm0.10}$	14.95 ± 0.55	7.92 ± 2.07	8.47 ± 3.26	$\textbf{7.10} \pm \textbf{1.02}$
	$\Delta_{EO}(\downarrow)$	$\underline{2.93\pm0.93}$	8.26 ± 2.33	6.09 ± 0.31	5.34 ± 0.61	5.63 ± 1.06	3.41 ± 0.90	4.65 ± 3.06	$\boldsymbol{1.77 \pm 1.01}$
Pokec-z	Accuracy (\uparrow)	68.06 ± 0.05	63.95 ± 0.43	$\underline{68.11 \pm 0.23}$	OOM	67.36 ± 0.71	63.89 ± 2.38	$\textbf{70.17} \pm \textbf{0.22}$	68.10 ± 0.85
	F1-score (\uparrow)	67.71 ± 0.11	63.62 ± 0.57	65.63 ± 0.90	60.62 ± 0.6	65.40 ± 2.81	$\underline{68.83 \pm 1.31}$	67.20 ± 0.51	69.32 ± 0.4
	AUCROC (\uparrow)	73.46 ± 0.01	68.25 ± 0.34	74.11 ± 0.24	66.37 ± 0.7	74.07 ± 0.55	71.35 ± 1.16	$\underline{74.71 \pm 0.25}$	$\textbf{75.37} \pm \textbf{0.59}$
	$\Delta_{DP}(\downarrow)$	5.70 ± 0.10	2.86 ± 1.09	4.95 ± 0.50	2.89 ± 0.4	5.17 ± 1.75	$\underline{2.72 \pm 2.52}$	5.30 ± 0.27	2.69 ± 2.01
	$\Delta_{EO}(\downarrow)$	2.49 ± 0.18	4.29 ± 0.57	2.85 ± 0.84	2.54 ± 0.7	2.83 ± 2.54	$\underline{2.36 \pm 1.88}$	2.58 ± 1.01	1.22 ± 1.20
Pokec-n	Accuracy (\uparrow)	67.26 ± 0.82	60.25 ± 0.65	68.18 ± 0.39	OOM	68.23 ± 1.51	64.75 ± 1.31	$\textbf{70.24} \pm \textbf{0.99}$	$\underline{68.82\pm0.13}$
	F1-score (\uparrow)	$\textbf{67.51} \pm \textbf{1.41}$	57.30 ± 0.57	62.57 ± 1.31	52.53 ± 0.1	65.03 ± 0.76	$\underline{65.65 \pm 1.03}$	63.29 ± 2.10	65.04 ± 1.95
	AUCROC (\uparrow)	74.16 ± 1.48	63.36 ± 0.03	$\underline{74.86\pm0.59}$	62.05 ± 0.6	73.64 ± 1.07	70.79 ± 0.95	73.75 ± 1.90	$\textbf{75.66} \pm \textbf{0.13}$
	$\Delta_{DP}(\downarrow)$	$\underline{1.69 \pm 1.01}$	2.36 ± 1.88	1.73 ± 1.47	2.08 ± 1.2	2.27 ± 1.23	2.41 ± 1.97	1.87 ± 0.31	1.40 ± 1.25
	$\Delta_{EO}(\downarrow)$	$\underline{1.28\pm0.11}$	1.62 ± 1.39	1.38 ± 0.48	1.82 ± 0.9	2.15 ± 0.50	3.05 ± 1.85	1.91 ± 0.55	0.98 ± 0.99
Avg.Rank		3.80	5.45	4.95	6.50	5.60	3.85	3.95	1.75

Table 1. Model utility and fairness of node classification.

* (\uparrow) represents the larger, the better; (\downarrow) represents the opposite. The best result and the runner-up result are shown in **bold** and <u>underlined</u> respectively.

- FDGIB achieves the best fairness across all datasets. The superiority of our method can be ascribed to the disentangled counterfactual augmentation strategy. The disentanglement of FDGIB ensures that the generated counterfactual ego graphs follow a consistent distribution with the real ego graphs of the same sensitive groups. The credible counterfactual ego graphs can significantly facilitate the fairness of downstream GNN classifiers.
- FDGIB strikes an optimal balance between model utility and fairness. We can observe that there is a tradeoff between the model utility and fairness for all methods since none of these methods achieve the best model utility and fairness simultaneously. This observation is consistent with the recent work [37] which theoretically proves that there is an inherent tradeoff between fairness and accuracy in the classification setting. To investigate the overall performance of our method on fairness and model utility, we calculate the average rank of all model utility and fairness evaluation metrics on all datasets. We find that FDGIB ranks 1.75, while the runner-up model ranks 3.80. This demonstrates that our method achieves the best trade-off between fairness and model utility. In other words, our FDGIB can achieve the best fairness while maintaining comparable classification performance. Only on the Accuracy metric of the Income dataset, our method suffers a significant loss. We attribute this observation to the fact that the labels on Income are severely imbalanced (almost 8:2). On the imbalanced dataset, the F1-score is more reliable where our FDGIB achieves the second-best performance.

To further investigate the generalization of our FDGIB to different GNN backbone classifiers, we adopt four different GNN architectures: GraphSAGE, GCN, GAT, and JK. The results are presented in Table 2. We observe that (1)

				r		r		r	
Dataset	Metrics	GraphSAGE		GCN		GAT		JK	
		Vanilla	FDGIB	Vanilla	FDGIB	Vanilla	FDGIB	Vanilla	FDGIB
Bail	Accuracy (\uparrow)	85.99 ± 0.40	$\textbf{86.97} \pm \textbf{0.23}$	84.40 ± 0.32	$\textbf{85.99} \pm \textbf{0.40}$	$\textbf{83.01} \pm \textbf{0.60}$	82.33 ± 0.64	84.51 ± 0.17	$\textbf{86.76} \pm \textbf{0.65}$
	F1-score (\uparrow)	80.05 ± 0.37	$\textbf{82.05} \pm \textbf{0.20}$	79.83 ± 0.28	$\textbf{80.05} \pm \textbf{0.37}$	$\textbf{77.05} \pm \textbf{0.54}$	76.33 ± 0.46	80.00 ± 0.25	81.82 ± 0.85
	AUCROC (\uparrow)	90.44 ± 0.10	91.74 ± 0.09	90.11 ± 0.12	$\textbf{90.44} \pm \textbf{0.10}$	$\textbf{87.43} \pm \textbf{0.26}$	86.72 ± 0.17	$\textbf{90.20} \pm \textbf{0.23}$	89.60 ± 0.81
	$\Delta_{DP}(\downarrow)$	1.35 ± 0.87	$\boldsymbol{0.99 \pm 1.03}$	8.93 ± 0.40	1.35 ± 0.87	4.61 ± 0.63	$\textbf{4.08} \pm \textbf{0.57}$	9.08 ± 0.25	$\textbf{7.27} \pm \textbf{0.83}$
	$\Delta_{EO}(\downarrow)$	2.31 ± 2.39	1.06 ± 0.52	6.07 ± 0.48	$\textbf{2.31} \pm \textbf{2.39}$	2.75 ± 0.81	2.57 ± 0.99	5.40 ± 0.31	$\textbf{4.91} \pm \textbf{1.30}$
Income	Accuracy (↑)	$\textbf{75.98} \pm \textbf{2.34}$	75.82 ± 2.07	$\textbf{76.18} \pm \textbf{1.23}$	74.96 ± 0.73	$\textbf{74.03} \pm \textbf{3.50}$	72.55 ± 0.32	$\textbf{72.23} \pm \textbf{1.21}$	71.44 ± 3.51
	F1-score (\uparrow)	$\textbf{56.85} \pm \textbf{1.59}$	56.30 ± 0.07	$\textbf{48.40} \pm \textbf{1.07}$	45.35 ± 1.23	51.01 ± 1.96	49.57 ± 1.05	43.93 ± 1.79	$\textbf{44.72} \pm \textbf{1.96}$
	AUCROC (\uparrow)	81.90 ± 0.95	82.09 ± 0.35	$\textbf{73.78} \pm \textbf{0.88}$	71.36 ± 0.80	$\textbf{77.41} \pm \textbf{2.03}$	75.95 ± 0.80	$\textbf{71.63} \pm \textbf{0.99}$	70.92 ± 2.57
	$\Delta_{DP}(\downarrow)$	10.09 ± 1.97	$\textbf{7.10} \pm \textbf{1.02}$	10.43 ± 5.34	1.82 ± 0.41	7.31 ± 5.52	5.62 ± 4.50	8.73 ± 3.97	$\textbf{1.67} \pm \textbf{1.16}$
	$\Delta_{EO}(\downarrow)$	2.08 ± 2.10	$\boldsymbol{1.77 \pm 1.01}$	12.93 ± 7.79	$\textbf{2.91} \pm \textbf{1.06}$	8.24 ± 6.03	$\textbf{6.71} \pm \textbf{5.86}$	9.42 ± 4.26	$\textbf{2.68} \pm \textbf{1.45}$
Pokec-z	Accuracy (↑)	68.60 ± 0.20	68.10 ± 0.85	67.64 ± 0.16	67.58 ± 0.53	66.30 ± 0.74	66.39 ± 0.16	66.54 ± 0.60	67.64 ± 0.56
	F1-score (\uparrow)	69.56 ± 0.80	69.32 ± 0.40	69.03 ± 0.20	68.92 ± 0.13	66.27 ± 0.71	$\textbf{67.06} \pm \textbf{0.29}$	69.10 ± 0.52	68.05 ± 1.14
	AUCROC (\uparrow)	75.26 ± 0.07	$\textbf{75.37} \pm \textbf{0.59}$	$\textbf{75.56} \pm \textbf{0.05}$	74.96 ± 0.17	71.84 ± 0.54	$\textbf{72.10} \pm \textbf{0.53}$	74.08 ± 0.53	$\textbf{74.40} \pm \textbf{0.19}$
	$\Delta_{DP}(\downarrow)$	6.74 ± 0.84	$\textbf{2.69} \pm \textbf{2.01}$	2.92 ± 1.94	0.59 ± 0.36	1.94 ± 1.06	$\textbf{0.41} \pm \textbf{0.10}$	4.10 ± 1.67	0.35 ± 0.23
	$\Delta_{EO}(\downarrow)$	4.61 ± 0.57	1.22 ± 1.20	4.32 ± 1.34	0.45 ± 0.36	1.57 ± 0.09	$\bf 1.22 \pm 0.82$	3.68 ± 0.89	0.60 ± 0.67
Pokec-n	Accuracy (↑)	68.86 ± 0.67	68.82 ± 0.13	68.79 ± 0.79	68.97 ± 1.13	$\textbf{68.36} \pm \textbf{1.64}$	67.89 ± 1.20	68.85 ± 0.85	68.56 ± 0.95
	F1-score (\uparrow)	$\textbf{67.19} \pm \textbf{1.29}$	65.04 ± 1.95	68.77 ± 0.42	65.37 ± 1.28	67.25 ± 0.02	66.46 ± 0.63	68.58 ± 0.37	64.62 ± 1.68
	AUCROC (\uparrow)	$\textbf{75.85} \pm \textbf{0.30}$	75.66 ± 0.13	$\textbf{76.94} \pm \textbf{0.26}$	75.39 ± 1.38	$\textbf{74.45} \pm \textbf{0.93}$	73.76 ± 0.92	$\textbf{76.38} \pm \textbf{0.31}$	74.95 ± 0.24
	$\Delta_{DP}(\downarrow)$	6.94 ± 1.12	1.40 ± 1.25	8.74 ± 2.20	$\textbf{2.02} \pm \textbf{1.38}$	8.05 ± 1.43	$\textbf{4.48} \pm \textbf{2.68}$	9.36 ± 0.35	$\textbf{3.31} \pm \textbf{0.70}$
	$\Delta_{EO}(\downarrow)$	4.13 ± 0.47	$\textbf{0.98} \pm \textbf{0.99}$	6.08 ± 3.20	$\textbf{2.74} \pm \textbf{1.16}$	7.67 ± 1.20	$\textbf{4.93} \pm \textbf{1.75}$	5.81 ± 0.12	$\textbf{3.12} \pm \textbf{0.03}$

 Table 2. Comparison of model utility and fairness between vanilla GNNs and GNNs with our FDGIB.

Compared with vanilla GNNs, our FDGIB significantly improves fairness while maintaining competitive model utility. We observe that FDGIB either outperforms vanilla GNNs on the model utility or slightly sacrifices model utility to achieve superior fairness. This further indicates that our model strikes a balance between model utility and fairness. (2) Among four backbone classifiers, Graph-SAGE achieves the best model utility. In terms of fairness performance, Graph-SAGE outperforms other models on the Bail and Pokec-n datasets. However, on the Income and Pokec-z dataset, the JK encoder achieves the best results, albeit with a potential trade-off on the model utility.

6.3 Disentanglement Analysis (RQ 2)

To answer RQ2, we conduct experiments to investigate whether our model successfully achieves disentanglement for sensitive-related and irrelevant information. Just as Theorem 2 demonstrates, the disentanglement of FDGIB can be analyzed in two levels: the mutual disentanglement and the sensitive disentanglement. We first investigate the mutual disentanglement. Specifically, we measure the MI between the sensitive-related representation R and sensitive-independent representation D in Fig. 3(a). We adopt the MINE [4], a mutual information estimator to quantitatively measure the MI. FDGIB-w/o-ind indicates the variant of FDGIB without the minimal sufficiency and independence terms, *i.e.*, an entangled method that only considers the join sufficiency. From the figure, we can observe that our FDGIB can significantly facilitate mutual disentanglement between R and D and achieve lower MI.

Then we investigate the sensitive disentanglement. Specifically, we employ two classifiers to predict the sensitive attributes based on the disentangled representations R and D. As illustrated in Fig. 3(b), we observe a significant difference in the

prediction performance of sensitive attribute S between R and D. This observation indicates that R encapsulates sensitive-related information while D embeds sensitive-independent information. The empirical disentanglement analyses further verify the validity of our theoretical analysis in Theorem 2.



Fig. 3. The performance of the disentangled learning of FDGIB. (a) The MI between representations R and D; (b) Sensitive attributes prediction with R and D respectively.



Fig. 4. Hyper-parameter study on Bail dataset.

6.4 Ablation Study(RQ 3)

To answer RQ3, we conduct ablation studies to investigate whether the three conditions of our FDGIB facilitate the performance. Specifically, we consider three variants: FDGIB-w/o-js, FDGIB-w/o-ms, and FDGIB-w/o-ind that respectively discard the joint sufficiency, the minimal sufficiency, and the independence. GraphSAGE is adopted as the backbone classifier. As shown in Table 3, we can observe that most of the variants exhibit inferior fairness performance compared to FDGIB, except for FDGIB-w/o-is on Income. Nonetheless, the improvement of FDGIB-w/o-js in fairness might be achieved at the cost of prediction utility, since it achieves a much lower F1 score than others on the imbalanced Income dataset. This indicates FDGIB-w/o-js tends to make random predictions regardless of the sensitive attributes and thus achieve better fairness. Moreover, FDGIB-w/o-ms achieves competitive prediction performance compared to all variants, which can be attributed to the fact that the minimal sufficiency condition may inevitably sacrifice a certain degree of model utility. This further empirically verifies that there is a tradeoff between model utility and fairness in our FDGIB. The ablation results demonstrate that the three conditions contribute to FDGIB.

Dataset	Method	Accuracy (\uparrow)	F1-score (\uparrow)	AUCROC (\uparrow)	$\Delta_{DP} (\downarrow)$	$\Delta_{EO} (\downarrow)$
Bail	FDGIB-w/o-js	83.71 ± 0.75	73.30 ± 1.64	90.82 ± 0.18	1.86 ± 0.69	2.59 ± 1.53
	FDGIB-w/o-ms	$\textbf{87.33} \pm \textbf{0.42}$	82.36 ± 0.26	92.03 ± 0.13	4.13 ± 0.84	2.75 ± 0.55
	${\rm FDGIB}\text{-}{\rm w/o\text{-}ind}$	87.32 ± 0.86	82.27 ± 1.10	91.66 ± 0.37	2.72 ± 0.49	1.15 ± 0.89
	FDGIB	86.97 ± 0.23	82.05 ± 0.20	91.74 ± 0.09	$\textbf{0.99} \pm \textbf{1.03}$	1.06 ± 0.52
Income	FDGIB-w/o-js	81.05 ± 0.27	49.45 ± 1.35	81.64 ± 1.66	$\textbf{6.74} \pm \textbf{1.21}$	2.12 ± 0.61
	FDGIB-w/o-ms	76.61 ± 0.53	56.32 ± 1.24	81.67 ± 0.71	8.12 ± 3.94	4.12 ± 2.56
	${\rm FDGIB}\text{-}{\rm w/o\text{-}ind}$	74.29 ± 0.28	55.93 ± 0.11	81.77 ± 0.24	9.83 ± 0.67	2.75 ± 0.69
	FDGIB	75.82 ± 2.07	56.30 ± 0.07	82.09 ± 0.35	7.10 ± 1.02	$\textbf{1.77} \pm \textbf{1.01}$
Pokec-z	FDGIB-w/o-js	67.13 ± 0.20	68.44 ± 1.97	73.30 ± 1.42	4.06 ± 1.36	2.41 ± 0.99
	FDGIB-w/o-ms	68.38 ± 0.77	68.51 ± 1.58	$\textbf{76.04} \pm \textbf{0.29}$	7.09 ± 2.12	4.58 ± 2.01
	${\rm FDGIB}\text{-}{\rm w/o\text{-}ind}$	67.32 ± 1.21	68.04 ± 2.09	74.86 ± 0.57	5.49 ± 1.84	4.45 ± 0.81
	FDGIB	68.10 ± 0.85	69.32 ± 0.4	75.37 ± 0.59	2.69 ± 2.01	1.22 ± 1.20
Pokec-n	FDGIB-w/o-js	67.71 ± 1.12	61.91 ± 2.85	73.75 ± 1.67	4.84 ± 2.10	3.83 ± 0.72
	FDGIB-w/o-ms	69.42 ± 0.37	68.37 ± 0.47	75.92 ± 0.51	3.16 ± 1.96	2.04 ± 1.34
	FDGIB-w/o-ind	68.91 ± 1.40	65.81 ± 1.83	75.58 ± 0.58	3.32 ± 0.87	1.90 ± 0.61
	FDGIB	68.82 ± 0.13	65.04 ± 1.95	75.66 ± 0.13	$\textbf{1.40} \pm \textbf{1.25}$	$\textbf{0.98} \pm \textbf{0.99}$

 Table 3. Ablation results on three conditions.

6.5 Hyper-Parameter Sensitivity Analysis

Besides the above experimental results, we also conduct parameter studies to investigate the influence of hyper-parameters, i.e., α , β , and γ in Eq. (5), which control the weight of minimal sufficiency term, the IB term and the independence term, separately. The results over different settings of parameters in Bail are shown in Fig. 4. We can observe that the model utility metrics remain robust across various hyper-parameter values while the fairness metrics prefer larger hyper-parameters. This indicates that larger hyper-parameters, which enhance the constraints on disentanglement, may benefit the learning of fair GNNs.

7 Related Work

Fairness Learning on Graphs. According to the strategies employed to achieve fairness, recent works can be roughly divided into algorithm-based and augmentation-based methods. Specifically, algorithm-based methods [8,23,34] endeavor to design appropriate objective functions to learn fair embeddings by incorporating regularization terms or modifying weights. For example, FairGO [34] utilizes a composition of filters to transform the original embeddings into a fair embedding space. However, the necessity of specific constraints suffers generalization problems. Different from that, augmentation-based methods [20,24] address the data biases by generating augmented samples through modifying graphs. For example, FairAug [20] reduces the impact of sensitive attributes on the model by feature masking, edge perturbation, and node sampling. Nonetheless, existing fairness methods focus on augmenting the original graph from the

entangled representation space, leading to suboptimal performance on downstream tasks. To the best of our knowledge, only CAF [14] considers the learning of fair disentangled graph representations. However, CAF finds counterfactual samples directly from the original graph based on the l_2 norm of the disentangled representations, which may fail when no counterfactual nodes exist in the original data for the given target node.

Disentangled Representation Learning (DRL). DRL [22, 30, 36] seeks to learn a model that can discern the factors embedded within the observable data by representing them in a disentangled manner. However, the high dimensionality and structural complexity of graph data pose challenges for DRL on graphs. To address this, DisenGCN [25] proposes a neighborhood routing mechanism, which is capable of dynamically identifying the latent factor that may have caused the edge between nodes. Wu et al. [33] design a disentangled graph attention network by leveraging relation-aware aggregation and mutual information to determine weights for disentangled components. Nonetheless, the performance of DRL has not been investigated on the counterfactual graph augmentation task.

8 Conclusion

In this paper, we investigate the counterfactual graph augmentation problem to rectify the inconsistent distribution of sensitive attributes in the real dataset. Existing augmentation-based methods fail to disentangle different dependencies on sensitive attributes, leading to sub-optimal performance on downstream tasks. To tackle this problem, we introduce the IB theory into the counterfactual graph augmentation task and propose a novel disentangled counterfactual augmentation method named Fair Disentangled Graph Information Bottleneck (FDGIB). FDGIB consists of three conditions, namely Minimal Sufficiency, Independence, and Joint Sufficiency. With these conditions, FDGIB can embed the information of the original graph into two disentangled spaces: sensitive-related and sensitive-independent. We derive tractable bounds for optimizing the IB-based objective and offer theoretical analysis to substantiate the effectiveness of our proposed approach. Extensive experimentation on various datasets demonstrates the superior performance of our FDGIB in improving fairness.

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A New Framework for Evaluating the Validity and the Performance of Binary Decisions on Manifold-Valued Data

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Abstract. In this paper, we introduce a new framework that can be used for evaluating the validity and the performance of machine learning models on manifold-valued data. More particularly, two methods are detailed with theoretical properties for spherical and functional data. In a general setting, we develop a new set of procedures for nonparametric hypothesis testing on manifolds within a desired error level. These tests encompass probability distributions constrained to specific domains, which can pose significant challenges for commonly used techniques. The resulting statistical concepts are primarily characterized by computational simplicity and are grounded in relevant contexts, making them extendable to a wide range of applications. The algorithms and the theoretical analysis of the proposed methods are substantiated by many and varied experimental results on simulated and real data.

Keywords: Machine Learning \cdot Hypothesis testing \cdot Manifold-valued data \cdot Spherical Data \cdot Functional Data \cdot Probability Density Functions

1 Introduction

In machine learning and data science, binary decision has taken an important part of applications ranging from science to industry [8,17,18]. For example, in medical applications, a binary decision is required to determine whether a patient has disease or not using different features. This has been solved by different strategies among which the most popular remains binary decision based on optimization and hypothesis testing based on inference. If the first one was well generalized for manifold-valued data, the second one has been little or not investigated. It is reminiscent of its usefulness for evaluating the validity and the performances of machine learning models and its important role in reducing subjectivity, by making choices more objective and more transparent for interpretation [3, 4]. In this paper, we generalize hypothesis testing for manifoldvalued data and use it as two-sample testing for binary decision. Furthermore, we make it possible to use it as a metric for evaluating machine learning models. Nonlinear data have been widely used and analyzed in meteorology, astronomy, forest sciences, and biology with particular machine learning models [1, 10,29,31]. For example, directional data require non-standard methods [11,30], while redefining fundamental concepts [13]. The exploration of directional data traces back to fundamental analysis, focusing on the unit circle S^1 and the unit sphere S^2 [6]. In hypothesis tests for spherical data, circular cases have been studied in [24] and then extended to the unit sphere in [25]. Further developments explored tests for dispersion on a sphere by [26], building on the distribution proposed in [7], and discussing properties of the spherical median and equivalents for the sign test. More recently, [19] introduced a method for multivariate functional data which was extended in [27] via a large-scale Monte Carlo simulation spherical data.

In this paper, we introduce a novel method for analyzing directional data on the sphere [14]. Our approach involves employing random projections along fixed directions to estimate single and double spherical means. We emphasize the importance of projection onto the tangent space of the sphere, leveraging its advantages in addressing various problems. Tests are designed to assess alignment with a population's average direction and shared population polarization. We provide a thorough exploration of background theory before deriving these inference tests.

Recently, many widely used models involve high-dimensional probability distributions constrained within specific domains. These models often pose significant computational challenges,: Transforming the constrained domain to the entire Euclidean space for convenience can be computationally intractable. To address this issue, many works have explored connections between Fisher geometry and the geometry of the infinite Hilbert sphere [9,23]. Much of this research focuses on probability density functions and shape analysis. Particularly, geometrical tools and optimal transport techniques in order to quantify the divergence between distributions.

In this study, we study a subclass of functional data with a focus on Probability Density Functions (PDFs) using an adaptive structure derived from the Fisher-Rao metric [12]. This simplifies the analysis of data (nonlinear) in infinitedimensional spaces, making them interpretable through histograms or probability density representations. Among the main contributions of this framework, extending hypothesis testing from finite-dimensional spherical data to PDFs equipped with the Fisher-Rao metric is a considerable novelty. Hence, we are able to make binary decisions from functional data by measuring divergence between their PDFs [15,23].

The rest of the paper is organized as follows. Section 2 provides background and fundamental concepts. Section 3 details methodology for extending the multivariate central limit theorem. Section 4 presents statistical tools for analyzing spherical data. Section 5 establishes the connection between finite-dimensional and Hilbert spheres. Section 6 discusses experimental results, and Sect. 7 offers concluding remarks. For simple and fluid reading, technical proofs are given in the Appendix.

2 Background

Before we present our main results we first recall some notions to make the rest easier to understand [28] and then give some required tools for its extended version to the finite-dimensional sphere. Let $\mathbf{v} = (v^1, \ldots, v^n)^T \in \mathbb{R}^n$ be a random vector of a mean $\mathbb{E}(\mathbf{v}) = \mathbf{m} \in \mathbb{R}^n$ and a covariance matrix $\operatorname{Var}(\mathbf{v}) = \mathbb{E}[(\mathbf{v} - \mathbf{m})(\mathbf{v} - \mathbf{m})^T] = \boldsymbol{\Sigma} \in \mathbb{R}^{n \times n}$.

Theorem 1. Let \mathbf{v}_i (i = 1, ..., N) be a sequence of independent random variables, but not necessary identically distributed, each with finite expected value \mathbf{m}_i and covariance $\boldsymbol{\Sigma}_i$. Define the resultant sum of \mathbf{v}_i by $\mathbf{S}_N = \sum_{i=1}^N \mathbf{v}_i$. Then, under the Lyapunov condition: "If the $(2 + \varepsilon)^{\text{th}}$ moment with $\varepsilon > 0$ exists for a statistical distribution of independent random variates \mathbf{v}_i , the means \mathbf{m}_i and variances $\boldsymbol{\Sigma}_i$ are finite", i.e.,

$$\lim_{N \to \infty} \left(\sum_{i=1}^{N} \boldsymbol{\Sigma}_{i}\right)^{-(2+\varepsilon)/2} \sum_{i=1}^{N} \mathbb{E}\left(|\mathbf{v}_{i} - \mathbf{m}_{i}|^{2+\varepsilon}\right) = 0$$

we have

$$\left(\sum_{i=1}^{N} \boldsymbol{\Sigma}_{i}\right)^{-1/2} (\mathbf{S}_{N} - \sum_{i=1}^{N} \mathbf{m}_{i}) \xrightarrow{D} \mathcal{N}_{n}(0, \boldsymbol{I}) \quad as \ N \to \infty.$$
(1)

Here, $\mathcal{N}_n(\mathbf{m}, \boldsymbol{\Sigma})$ refers to the multivariate Gaussian distribution of dimension n with mean \mathbf{m} and covariance $\boldsymbol{\Sigma}$.

Let $\mathbf{v}_1 = (v_1^1, \dots, v_1^n)^T$ and $\mathbf{v}_2 = (v_2^1, \dots, v_2^n)^T$ be two vectors in \mathbb{R}^n . We remind that the Euclidean inner product between them is $\langle \mathbf{v}_1, \mathbf{v}_2 \rangle_2 = \mathbf{v}_1^T \mathbf{v}_2 = \sum_{j=1}^n v_j^1 v_2^j$ implying that the Euclidean norm of $\mathbf{v} = (v^1, \dots, v^n)^T \in \mathbb{R}^n$ is $||\mathbf{v}||_2 = \langle \mathbf{v}, \mathbf{v} \rangle_2^{1/2} = (\sum_{j=1}^n v_j^{1/2})^{\frac{1}{2}}$. The finite-dimensional unit sphere in \mathbb{R}^n is defined by

$$\mathcal{S}^{n-1} := \Big\{ \mathbf{z} \in \mathbb{R}^n \mid ||\mathbf{z}||_2 = 1 \Big\}.$$
(2)

Endowed with the Euclidean inner product, the tangent space of S^{n-1} locally at \mathbf{z} is

$$T_{\mathbf{z}}(\mathcal{S}^{n-1}) := \Big\{ \mathbf{v} \in \mathbb{R}^n \mid \big\langle \mathbf{z}, \mathbf{v} \big\rangle_2 = 0 \Big\}.$$
(3)

The distance between two vectors on the sphere $\mathbf{z}_1, \mathbf{z}_2 \in S^{n-1}$, called the geodesic distance, is given by the shortest arc connecting \mathbf{z}_1 to \mathbf{z}_2 and satisfying

$$d_{\mathcal{S}^{n-1}}(\mathbf{z}_1, \mathbf{z}_2) = \arccos\left(\left\langle \mathbf{z}_1, \mathbf{z}_2 \right\rangle_2\right). \tag{4}$$

Given N spherical vectors $\mathbf{z}_1, \ldots, \mathbf{z}_N \in S^{n-1}$, their natural mean on S^{n-1} is the Fréchet mean, denoted $\hat{\boldsymbol{\mu}}$, that minimizes the Fréchet variance. We note

$$\hat{\boldsymbol{\mu}} = \underset{\mathbf{z}\in\mathcal{S}^{n-1}}{\operatorname{argmin}} \sum_{i=1}^{N} d_{\mathcal{S}^{n-1}}^{2} (\mathbf{z}, \mathbf{z}_{i}).$$
(5)

The Fréchet mean is unique on any part of the sphere with an injectivity radius less than π . It can be found using a gradient approach detailed in [23]. In this paper, we adopt a strategy of using linear methods for spherical data, achieved by projecting vectors onto a tangent space to form a linear plane instead of a nonlinear manifold (sphere). Additionally, the Euclidean distance on the tangent plane can be used instead of the geodesic distance.

3 Formulation on the Tangent Space of the Sphere



Fig. 1. An illustration of the measure of concentration around the mean.

Let $\mathbf{z} = (z^1, \ldots, z^n)^T$, $\boldsymbol{\mu} = (\boldsymbol{\mu}^1, \ldots, \boldsymbol{\mu}^n)^T \in S^{n-1}$ such that \mathbf{z} belongs to the cut locus of $\boldsymbol{\mu}$, which means that the inner product $\langle \mathbf{z}, \boldsymbol{\mu} \rangle_2 > 0$ almost surely and the geodesic distance $d_{S^{n-1}}(\mathbf{z}, \boldsymbol{\mu}) < \pi/2$ almost surely. In other words, there exists a radius $r \ge 0$ such that the distance can be bounded as follows: $d_{S^{n-1}}(\boldsymbol{\mu}, \mathbf{z}) \le$ $r < \pi/2$. We will be using these two vectors to indicate direction only. Let \mathbf{N} be a unit normal vector on S^{n-1} and $\mathbf{T} \perp \mathbf{N}$ be the corresponding unit tangent vector. Therefore, $\{\mathbf{N}, \mathbf{T}\}$ constitutes together an orthonormal basis in the ambient space \mathbb{R}^n . Let $\mathbf{x} \in \mathbb{R}^n$ be the intersection between the limit of \mathbf{z} and the tangent space spanned by \mathbf{T} and orthogonal to \mathbf{N} as illustrated in Fig. 1. By adjusting the basis such that $\mathbf{N} = \boldsymbol{\mu}$ and writing $\mathbf{x} = c.\mathbf{z}$ where $c = ||\mathbf{x}||_2 \ge 1$, we get

$$\mathbf{x} = 1.\boldsymbol{\mu} + (c^2 - 1)^{1/2}.\mathbf{T}.$$
 (6)

So, with c as a scalar random variable, we have **x** is a random vector in \mathbb{R}^n . Applying the expectation we get

$$\mathbb{E}(\mathbf{x}) = 1.\boldsymbol{\mu} + \mathbb{E}((c^2 - 1)^{1/2}).\mathbf{T}.$$
(7)

The quantity $\mathbb{E}(c)$ is a measure of the concentration of the value of \mathbf{z} around the direction $\boldsymbol{\mu}$. If the radius r is close to zero then $\mathbb{E}(c)$ will be close to 1. Conversely, if the radius r is close to $\pi/2$ then $\mathbb{E}(c)$ takes large but finite values. In all cases, we have $1 \leq \mathbb{E}(c) < \infty$.

Let \mathbf{x}_i (i = 1, ..., N) be a sequence of independent random variables in \mathbb{R}^n . From 7 it can be expressed as: $\mathbf{x}_i = 1.\boldsymbol{\mu} + \sqrt{c_i^2 - 1}.\mathbf{T} = c_i.\mathbf{z}_i$ where $\mathbf{z}_i \in \mathcal{S}^{n-1}$. **Theorem 2.** Define the resultant sum of \mathbf{x}_i by $\mathbf{S}_N = \sum_{i=1}^N \mathbf{x}_i$. Then, we have

$$\boldsymbol{\Sigma}^{-1/2}(\mathbf{S}_N - N\mathbb{E}(c)\bar{\mathbf{z}}) \xrightarrow{D} \mathcal{N}_n(0, \boldsymbol{I}) \quad as \ N \to \infty,$$
(8)

and

$$\boldsymbol{\sigma}^{-1}(\bar{\mathbf{z}}^T \mathbf{S}_N - N\mathbb{E}(c) ||\bar{\mathbf{z}}||_2^2) \xrightarrow{D} \mathcal{N}(0, 1) \quad as \ N \to \infty,$$
(9)

where $\bar{\mathbf{z}}$ refers to the empirical mean of $\mathbf{z}_1, \ldots, \mathbf{z}_N$, $\boldsymbol{\Sigma} = Var(c) \sum_{i=1}^N \mathbf{z}_i \mathbf{z}_i^T$ and $\boldsymbol{\sigma}^2 = \bar{\mathbf{z}}^T \boldsymbol{\Sigma} \bar{\mathbf{z}} = Var(c) \bar{\mathbf{z}}^T \sum_{i=1}^N \mathbf{z}_i \mathbf{z}_i^T \bar{\mathbf{z}}$. Here, $\mathcal{N}(.,.)$ refers to the univariate Gaussian distribution.

Based on the Law of Large Numbers we get the following results of convergence in probability.

Lemma 1.

1. $\frac{\mathbf{S}_{N}}{N} \xrightarrow{P} \mathbb{E}(c)\bar{\mathbf{z}} \quad as \ N \to \infty.$ 2. $\frac{||\mathbf{S}_{N}||_{2}}{N} \xrightarrow{P} \mathbb{E}(c)||\bar{\mathbf{z}}||_{2} \quad as \ N \to \infty.$ 3. $\frac{\mathbf{S}_{N}}{||\mathbf{S}_{N}||_{2}} \xrightarrow{P} \frac{\bar{\mathbf{z}}}{||\bar{\mathbf{z}}||_{2}} \quad as \ N \to \infty.$

Combining Theorem 2 and Lemma 1 we obtain the following proposition.

Proposition 1. We have

$$N\boldsymbol{\sigma}^{-1}(\frac{||\bar{\mathbf{z}}||_2||\mathbf{S}_N||_2}{N} - \mathbb{E}(c)||\bar{\mathbf{z}}||_2^2) \xrightarrow{D} \mathcal{N}(0,1) \quad as \ N \to \infty.$$
(10)

We can now use Proposition 1 to derive hypothesis tests for spherical data.

4 The Finite-Dimensional Unit Sphere

In this section, we develop tests of single and double spherical means.

4.1 Hypothesis Tests of a Single Spherical Mean

We assess a test to determine if the spherical directions $\mathbf{z}_1, \ldots, \mathbf{z}_N$ come from a population characterized by a mean direction $\boldsymbol{\mu}^0$. We use Proposition 2 to test the null hypothesis of a specific mean direction $\boldsymbol{\mu}^0$.

Proposition 2. Define the resultant sum of \mathbf{x}_i depending on $\boldsymbol{\mu}^0$ by $\mathbf{S}_N^{\boldsymbol{\mu}^0} = \sum_{i=1}^N c_i^{\boldsymbol{\mu}^0} \mathbf{z}_i$ where $c_i^{\boldsymbol{\mu}^0} = \frac{||\mathbf{x}_i||_2}{\langle \mathbf{x}_i, \boldsymbol{\mu}^0 \rangle_2}$. Under the hypothesis $H_0: \boldsymbol{\mu} = \boldsymbol{\mu}^0$, the statistic $N\boldsymbol{\sigma}^{-1}(\frac{||\mathbf{\bar{z}}||_2||\mathbf{S}_N^{\boldsymbol{\mu}^0}||_2}{N} - \mathbb{E}(c)||\mathbf{\bar{z}}||_2^2)$ is approximately distributed as $\mathcal{N}(0,1)$ when $N \to \infty$.

In the above term we can estimate $\mathbb{E}(c)$ from Lemma 1 by

$$\widehat{\mathbb{E}}(c) = \frac{1}{N} \frac{||\mathbf{S}_N||_2}{||\mathbf{\bar{z}}||_2}.$$
(11)

Likewise, we can estimate Var(c) in order to evaluate σ^2 by

$$\widehat{\operatorname{Var}}(c) = \widehat{\mathbb{E}}(c^2) - \widehat{\mathbb{E}}(c)^2,$$

$$= \frac{1}{N} \sum_{i=1}^N \frac{||\mathbf{x}_i||_2^2}{\langle \mathbf{x}_i^2, \hat{\boldsymbol{\mu}} \rangle_2^2} - \left(\frac{1}{N} \frac{||\mathbf{S}_N||_2}{||\bar{\mathbf{z}}||_2}\right)^2,$$
(12)

where $\hat{\mu}$ is the Fréchet mean estimate.

4.2 Hypothesis Tests for Equality of Two Spherical Means

Consider two independent resultant vectors \mathbf{S}_{N_1} and \mathbf{S}_{N_2} obtained from samples of sizes N_1 and N_2 respectively, i.e., $\mathbf{S}_{N_j} = \sum_{i=1}^{N_j} \mathbf{x}_i^j = \sum_{i=1}^{N_j} c_i^j \cdot \mathbf{z}_i^j$; j = 1, 2. Utilizing Proposition 1, we deduce the subsequent result.

Proposition 3. Under the hypothesis $H_0: \mathbb{E}(c^1)||\bar{\mathbf{z}}^1||_2^2 = \mathbb{E}(c^2)||\bar{\mathbf{z}}^2||_2^2$ where $\bar{\mathbf{z}}^1$ and $\bar{\mathbf{z}}^2$ refers to the empirical mean of spherical samples: \mathbf{z}_i^1 and \mathbf{z}_i^2 of sizes N_1 and N_2 , the statistic $\left(\frac{\sigma_1^2}{N_1^2} + \frac{\sigma_2^2}{N_2^2}\right)^{-1/2} \left(\frac{||\bar{\mathbf{z}}^1||_2||\mathbf{S}_{N_1}||_2}{N} - \frac{||\bar{\mathbf{z}}^2||_2||\mathbf{S}_{N_2}||_2}{N}\right)$ is approximately distributed as $\mathcal{N}(0,1)$ when $N_1, N_2 \to \infty$.

In order to evaluate σ_1^2 and σ_2^2 , the variances of c^1 and c^2 can be estimated by

$$\widehat{\operatorname{Var}}(c^{j}) = \frac{1}{N_{j}} \sum_{i=1}^{N_{j}} \frac{||\mathbf{x}_{i}^{j}||_{2}^{2}}{\left\langle \mathbf{x}_{i}^{j}, \hat{\boldsymbol{\mu}}^{j} \right\rangle_{2}^{2}} - \left(\frac{1}{N_{j}} \frac{||\mathbf{S}_{N_{j}}||_{2}}{||\bar{\mathbf{z}}^{j}||_{2}}\right)^{2}; \quad j = 1, 2,$$
(13)

where $\hat{\mu}^{j}$ is the Fréchet mean of *j*-th spherical sample of size N_{j} and \mathbf{x}_{i}^{j} denotes the *i*-th observation from *j*-th sample.

5 The Manifold of PDFs

Before we give details of the hypothesis test on the set of PDFs we introduce some tools about the geometry of Riemannian representations and their structures. The metric is fixed to be the Fisher-Rao the only metric invariant to reparametrizations [23].

5.1 From Spherical Data to Distribution Functions

Let p be a PDF of a real-valued random variable. The set of all PDFs defined on I = [0, 1] is a simplex satisfying

$$\mathcal{P} := \Big\{ p : I \to \mathbb{R} \mid p \text{ is nonnegative and } \int_{I} p(t)dt = 1 \Big\}.$$
(14)

The Fisher-Rao metric on \mathcal{P} is defined by

$$\left\langle g_1, g_2 \right\rangle_p := \int_I \frac{g_1(t)g_2(t)}{p(t)} dt, \tag{15}$$

where $g_1, g_2 \in T_p(\mathcal{P})$ are two tangent vectors belonging to the tangent space of \mathcal{P} locally at p and satisfying

$$T_p(\mathcal{P}) := \Big\{ g : I \to \mathbb{R} \ \big| \ \int_I g(t) dt = 0 \Big\}.$$
(16)

As a second representation we introduce the set of Square-root Density Functions (SRDFs) with

$$\mathcal{H} := \left\{ \psi : I \to \mathbb{R} \mid \psi \text{ is nonnegative, and } ||\psi||_{\mathbb{L}^2} := \left(\int_I \psi(t)^2 dt \right)^{1/2} = 1 \right\} (17)$$

 \mathcal{H} results to be the Hilbert upper-hemisphere (nonnegative part) endowed with the \mathbb{L}^2 metric. Note that the tangent space of \mathcal{H} locally at ψ is

$$T_{\psi}(\mathcal{H}) := \left\{ f: I \to \mathbb{R} \mid \left\langle \psi, f \right\rangle_{\mathbb{L}^2} = \int_I \psi(t) f(t) dt = 0 \right\}.$$
(18)

Associated with each $p \in \mathcal{P}$ is a unique $\psi \in \mathcal{H}$ (isometrically) expressed as

$$\psi(t) = \sqrt{p(t)}; \quad t \in I.$$
(19)

The advantage of the representation $\psi \in \mathcal{H}$ is that it greatly simplifies the underlying geometry of \mathcal{P} with some nice tools on the Hilbert sphere. For instance, the geodesic distance between ψ_1 and ψ_2 in \mathcal{H} is given by

$$d_{\mathcal{H}}(\psi_1, \psi_2) = \arccos\left(\left\langle \psi_1, \psi_2 \right\rangle_{\mathbb{L}^2}\right). \tag{20}$$

Given $\psi_1, \ldots, \psi_N \in \mathcal{H}$ their natural mean function is the Fréchet mean denoted \hat{h} belonging to \mathcal{H} and minimizing the Fréchet variance

$$\hat{h} = \underset{\psi \in \mathcal{H}}{\operatorname{argmin}} \sum_{i=1}^{N} d_{\mathcal{H}}^{2} (\psi, \psi_{i}).$$
(21)

Let $p \in \mathcal{P}$ be a PDF and $\psi \equiv \sqrt{p} \in \mathcal{H}$ be its corresponding SRDF. Since ψ is an element of $\mathbb{L}^2(I, \mathbb{R})$ then it can be represented as a convergent orthogonal series expansion satisfying

$$\psi(t) = \sum_{l=1}^{\infty} a_l \phi_l(t), \qquad (22)$$

where $(\phi_l)_l$ is a complete orthonormal basis in $\mathbb{L}^2(I,\mathbb{R})$. In practice, it seems natural to consider a truncated version of ψ at order *m* expressed as

$$\psi^m(t) = \sum_{l=1}^m a_l \phi_l(t),$$
(23)

and the rest of the sum is considered as an approximation error: $e^m(t) = \sum_{l=m+1}^{\infty} a_l \phi_l(t)$. Note that $\psi^m(t)$ can be re-written as

$$\psi^m(t) = \boldsymbol{\Phi}(t)^T \mathbf{A},\tag{24}$$

for $\mathbf{A} = (a_1, \ldots, a_m)^T$ and $\boldsymbol{\varPhi}(t) = (\phi_1(t), \ldots, \phi_m(t))^T$. From Eq. 17 we state that the truncated SRDF ψ^m belongs to \mathcal{H} if and only if ψ^m is nonnegative and $||\psi^m||_{\mathbb{L}^2} = 1$. The condition that ψ^m is nonnegative does not impose any additional calculation and we can maintain it in practice. Consequently, $\psi^m(t)$ is a SRDF if and only if in addition to non-negativity the Euclidean norm of \mathbf{A} takes one from the following identity

$$||\psi^{m}||_{\mathbb{L}^{2}}^{2} = \int_{I} \psi^{m}(t)^{2} dt = \sum_{l=1}^{m} a_{l}^{2} \int_{I} \phi_{l}(t)^{2} dt = \sum_{l=1}^{m} a_{l}^{2} = ||A||_{2}^{2}.$$
 (25)

5.2 Hypothesis Tests

Let $p \in \mathcal{P}$ be a PDF and $\psi^m \in \mathcal{H}$ be the truncated version of its corresponding SRDF depending on **A** as detailed in Eq. 24. We also assume that h is a spherical mean function in \mathcal{H} truncated by $h^m(t) = \mathbf{\Phi}(t)^T \mathbf{W}$ such that **W** is the corresponding spherical mean direction in \mathcal{S}^{m-1} . By analogy to the finite-dimensional case detailed in Sect. 3 and from Fig. 1 the limit of $\psi^m(t)$'s intersection with the orthonormal tangent vector at $h^m(t)$ is

$$f^m(t) = d.\psi^m(t),\tag{26}$$

where d is a scalar random variable giving the concentration of $\psi^m(t)$ around $h^m(t)$ and satisfying: $d = \frac{||f^m||_{\mathbb{L}^2}}{\langle f^m, h^m \rangle_{\mathbb{L}^2}} = \frac{1}{\langle \psi^m, h^m \rangle_{\mathbb{L}^2}}$. Now, we will show how Eq. 26 is equivalent to

$$\mathbf{B} = d.\mathbf{A},\tag{27}$$

where **B** is the intersection of the limit of **A** with the orthonormal tangent vector at **W**. From the isometry result in Eq. 25 we state that $d = \frac{1}{\langle \mathbf{A}, \mathbf{W} \rangle_2}$. Multiplying terms in Eq. 26 by $\boldsymbol{\Phi}(t)$ we get

$$\boldsymbol{\Phi}(t)f^{m}(t) = \frac{1}{\left\langle \mathbf{A}, \mathbf{W} \right\rangle_{2}} \boldsymbol{\Phi}(t) \boldsymbol{\Phi}(t)^{T} \mathbf{A}.$$
(28)

Integrating the above expression we obtain

$$\int_{I} \boldsymbol{\Phi}(t) f^{m}(t) dt = \frac{1}{\left\langle \mathbf{A}, \mathbf{W} \right\rangle_{2}} \Big(\int_{I} \boldsymbol{\Phi}(t) \boldsymbol{\Phi}(t)^{T} dt \Big) \mathbf{A}.$$
(29)

Using the fact that $\int_{I} \boldsymbol{\Phi}(t) \boldsymbol{\Phi}(t)^{T} dt = \mathcal{I}$ where \mathcal{I} refers to the $m \times m$ identity matrix we get

$$\int_{I} \boldsymbol{\Phi}(t) f^{m}(t) dt = \frac{1}{\left\langle \mathbf{A}, \mathbf{W} \right\rangle_{2}} \mathbf{A} = d.\mathbf{A}, \tag{30}$$

which matches Eq. 27 for $\mathbf{B} = \int_{I} \boldsymbol{\Phi}(t) f^{m}(t) dt$. Therefore, Proposition 2 and Proposition 3 can be updated for simple and double hypothesis tests on PDFs replacing \mathbf{x}_{i} by \mathbf{B}_{i} , \mathbf{z}_{i} by \mathbf{A}_{i} and $\boldsymbol{\mu}$ by \mathbf{W} .

6 Experimental Results

We empirically evaluate the proposed method's effectiveness through separate assessments on simulation studies and real-world datasets. A Python demo code of the method is provided at: Nonparametric-Statistical-Testing-for-Spherical-Data.

6.1 Results on Simulated Data

To begin, we validate the performance of the proposed framework using simulated datasets on both the finite-dimensional sphere and the space of PDFs.

Case 1: The Finite-Dimensional Sphere. We conduct experiments involving two hypothesis tests on the unit sphere, specifically on S^1 (n = 2) and S^2 (n = 3). For these experiments, we generate spherical data using the von Mises-Fisher (vMF) distribution [2], a fundamental, unimodal, and isotropic distribution defined on S^{n-1} . The probability density function of this distribution is represented as:

$$p_{\mu,\kappa}(t) = (\frac{\kappa}{2})^{n/2 - 1} \frac{1}{\Gamma(n/2)I_{n/2 - 1}} \exp(\kappa \mu^T t).$$

In this expression, Γ denotes the gamma function, and I_{ν} represents the modified Bessel function of the first kind at order ν . The parameter $\mu \in S^{n-1}$ specifies the unique mode, while $\kappa \geq 0$ indicates the degree of concentration.

i) Hypothesis tests of a simple spherical mean. From Proposition 2, when considering a significance level $\alpha \in [0, 1]$, the critical region for rejecting the null hypothesis $H_0: \mu = \mu^0$ is defined as follows:

$$CR_{\alpha} = \left\{ \frac{||\bar{\mathbf{z}}||_{2}||\mathbf{S}_{N}^{\mu^{0}}||_{2}}{N} < \mathbb{E}(c)|\bar{\mathbf{z}}||_{2}^{2} - q_{1-\alpha/2}\frac{\sigma}{N} \right\} \cup \left\{ \frac{|\bar{\mathbf{z}}||_{2}||\mathbf{S}_{N}^{\mu^{0}}||_{2}}{N} > \mathbb{E}(c)|\bar{\mathbf{z}}||_{2}^{2} + q_{1-\alpha/2}\frac{\sigma}{N} \right\}$$

Here, q_{α} represents the α -quantile of the standard Gaussian distribution. The mean and the variance of the concentration parameter c will be empirically estimated as described in Eq. 11 and Eq. 12.

In these experiments, we generate N = 1000 spherical data points \mathbf{z}_i from the von Mises-Fisher distribution $p_{\mu^0,\kappa}$ with $\kappa = 5$. For n = 2, we use $\mu^0 = (0,1)$, and for n = 3, we use $\mu^0 = (0, 0, 1)$. A few examples of \mathbf{z}_i are visualized in Fig. 2 (top) (a) on the circle and in Fig. 2 (bottom) (a) on the sphere. The Fréchet mean, denoted as $\hat{\boldsymbol{\mu}}$ and obtained from Eq. 5, is marked as a green dots. We also illustrate the tangent line and plane at μ^0 that contain the corresponding



Fig. 2. (a) Data simulated from the vMF distribution (blue) projected onto the tangent plane of the sphere at μ^0 (brown) and their Fréchet mean (green).(b) The distribution of $||\bar{\mathbf{z}}||_2 ||\mathbf{S}_N^{\mu^0}||_2$ under H_0 . (c) The acceptance region $\operatorname{CR}_{\alpha}^c$ (blue) with test statistic value (red) and *p*-value (gray)(Color figure online).

projected data, shown in brown. In Fig. 5 (top) (b) and Fig. 2 (bottom) (b), we provide visualizations of the Gaussian distribution of $||\bar{\mathbf{z}}||_2||\mathbf{S}_N^{\mu^0}||_2$ under the null hypothesis $H_0: \boldsymbol{\mu} = \boldsymbol{\mu}^0$, exhibiting a mean of $N\mathbb{E}(c)||\bar{\mathbf{z}}||_2^2$ and a variance of σ^2 for the circle and the sphere, respectively. Further details of the hypothesis testing are illustrated in Fig. 2 (top) (c) and Fig. 2 (bottom) (c). The horizontal red line represents the test statistic value $\frac{||\bar{\mathbf{z}}||_2||\mathbf{S}_N^{\mu^0}||_2}{N}$. The blue-shaded area indicates the region CR_{α}^c as the complement of CR_{α} where the null hypothesis H_0 is accepted. This region varies for different values of $\alpha \in [0, 1]$. The vertical gray line represents the test statistic value, assuming the null hypothesis is true. In this case, the *p*-value is equal to 0.8 for both datasets. Therefore, we can only accept the alternative hypothesis H_1 for high levels of significance, exceeding 80%.

ii) Hypothesis tests for equality of two spherical means. Proposition 3 defines the critical region for rejecting the null hypothesis H_0 : $\mathbb{E}(c^1)||\bar{\mathbf{z}}^1||_2^2 = \mathbb{E}(c^2)||\bar{\mathbf{z}}^2||_2^2$ as follows:

$$CR_{\alpha} = \left\{ \frac{||\mathbf{\bar{z}}^1||_2||\mathbf{S}_{N_1}||_2}{N_1} - \frac{||\mathbf{\bar{z}}^2||_2||\mathbf{S}_{N_2}||_2}{N_2} < -q_{1-\alpha/2}\sqrt{\frac{\sigma_1^2}{N_1^2} + \frac{\sigma_2^2}{N_2^2}} \right\} \cup \left\{ \frac{||\mathbf{\bar{z}}^1||_2||\mathbf{S}_{N_1}||_2}{N_1} - \frac{||\mathbf{\bar{z}}^2||_2||\mathbf{S}_{N_2}||_2}{N_2} > q_{1-\alpha/2}\sqrt{\frac{\sigma_1^2}{N_1^2} + \frac{\sigma_2^2}{N_2^2}} \right\}.$$



Circular data with $\mu^1 = (\frac{1}{\sqrt{2}}, \frac{1}{\sqrt{2}})$ and $\mu^2 = (-\frac{1}{\sqrt{2}}, \frac{1}{\sqrt{2}})$.



Spherical data with $\mu^1 = (\frac{1}{\sqrt{2}}, 0, \frac{1}{\sqrt{2}})$ and $\mu^2 = (-\frac{1}{\sqrt{2}}, 0, \frac{1}{\sqrt{2}})$.

Fig. 3. (a) Data simulated from the vMF distribution with μ^1 (blue) and μ^2 (red) projected onto the tangent space of the sphere (brown and pink, respectively) at μ^1 and μ^2 and their Fréchet means $\hat{\mu}^1$ and $\hat{\mu}^2$ (green). (b) The acceptance region CR^c_{α} (blue) with test statistic value (red) and *p*-value (gray)(Color figure online).

Here, the variance of the concentration parameter c^{j} is empirically estimated as described in Eq. 13 for j = 1, 2.

In this experiment, we simulate datasets with $N_1 = N_2 = 1000$ circular data points from $p_{\mu^1,\kappa}$ and $p_{\mu^2,\kappa}$. Specifically, we use $\mu^1 = (\frac{1}{\sqrt{2}}, \frac{1}{\sqrt{2}})$ and $\mu^2 = (-\frac{1}{\sqrt{2}}, \frac{1}{\sqrt{2}})$. In Fig. 3 (top) (a), we provide examples of \mathbf{z}_i with a mean of μ^1 represented by blue dots and a mean of μ^2 represented by red dots. The Fréchet means, denoted as $\hat{\mu}^1$ and $\hat{\mu}^2$, are shown as green dots. The tangent lines at μ^1 and μ^2 are depicted in black, along with the corresponding projected data in brown and pink, respectively. In Fig. 3 (top) (b), we have tested the null hypothesis $H_0 : \mathbb{E}(c^1) ||\bar{\mathbf{z}}^1||_2^2 = \mathbb{E}(c^2) ||\bar{\mathbf{z}}^2||_2^2$. The horizontal red line represents the test statistic value $\frac{||\bar{\mathbf{z}}^1||_2||\mathbf{S}_{N_1}||_2}{N_1} - \frac{||\bar{\mathbf{z}}^2||_2||\mathbf{S}_{N_2}||_2}{N_2}$. The blue area illustrates the region CR^c_{α} for different values of $\alpha \in [0, 1]$. The vertical gray line represents the *p*-value, which is zero in this case. Consequently, we confidently accept the alternative hypothesis $H_1 : \mathbb{E}(c^1) ||\bar{\mathbf{z}}^1||_2^2 \neq \mathbb{E}(c^2) ||\bar{\mathbf{z}}^2||_2^2$ for any level of significance α . The same process is repeated for testing spherical data in Fig. 3 (bottom) using $\mu^1 = (\frac{1}{\sqrt{2}}, 0, \frac{1}{\sqrt{2}})$ and $\mu^2 = (-\frac{1}{\sqrt{2}}, 0, \frac{1}{\sqrt{2}})$. The *p*-value remains zero, indicating no overlap between the red line and the blue area, and consequently leading to the acceptance of the alternative hypothesis H_1 .
Case 2: The Manifold of PDFs. To assess the computational complexity of the proposed procedure, we conducted a hypothesis test on populations of PDFs. We set the truncation order of the convergent orthogonal series expansion to m = 30, following the approach in [9]. As an orthonormal basis, we selected $\phi_l(t) = \sqrt{2} \sin(l\pi t)$ for $l = 1, \ldots, 30$.

We considered two datasets of simulated PDFs: beta and inverse gamma distributions. This experiment involved $N_1 = N_2 = 1000$ pairs of PDFs, with slight differences between the two populations in each dataset. Each observation p_i represented a PDF, with random uniform noise added to the initial parameters. For the beta class, we used $p_i^1 = \mathcal{B}(2 + \epsilon_i, 2)$ for the first population and $p_i^2 = \mathcal{B}(1.8 + \epsilon_i, 2)$ for the second, where $\epsilon_i \sim \mathcal{U}([-0.2, 0.2])$ is a realization of a uniform distribution. For the inverse gamma class, we employed $p_i^1 = \mathcal{IG}(3 + \epsilon_i, 0.1)$ for the first population and $p_i^2 = \mathcal{IG}(2.8 + \epsilon_i, 0.1)$ for the second, again with ϵ_i drawn from a uniform distribution. Examples of p_i^j are displayed in Fig. 4 (a) and Fig. 5 (a), with different colors (blue and red) representing the two populations.

We also illustrated the corresponding SRDFs $\psi_i^j \equiv \sqrt{p_i^j}$ in Fig. 4 (b) and Fig. 5 (b) for beta and inverse gamma distributions, respectively. In Fig. 4 (c) and Fig. 5 (c), we display the truncated SRDFs $\psi_i^{m,j}$. Furthermore, we computed the Fréchet mean of \mathbf{W}_i^j (i = 1, ..., N; j = 1, 2), denoted as $\hat{\mathbf{W}}^j$, using Eq. 5. This allowed us to illustrate the Fréchet mean of $\psi_i^{m,j}$, denoted $\hat{h}^{m,j}$ in green, with $\hat{h}^{m,j}(t) = \boldsymbol{\Phi}(t)^T \hat{\mathbf{W}}^j$. In Fig. 4 (d) and Fig. 5 (d), we presented the results of the double hypothesis test of population means. In both cases, we found that the *p*-value is zero. Therefore, we confidently reject the null hypothesis H_0 for any significance level α .

6.2 Real Data

In this section, we evaluate the proposed methods on two datasets of PDFs. The first dataset encompasses clinical growth charts for children aged 1 to 12 years, consisting of 100 girls and 100 boys [21]. This dataset provides a typical example of biological dynamics observed over a span of 132 months and has been widely used as a motivating case for the analysis of functional data [20]. Each growth chart represents the size increment (in centimeters) of a child over these 132 months. It's worth noting that standard techniques encounter difficulties when dealing with high-dimensional or functional data inputs. However, in this context, we've opted to use the corresponding PDFs instead of the original data. Consequently, all the growth charts are now represented by PDFs of child sizes, registered within the interval I = [0, 1]. In Fig. 6 (a), we provide visual representations of some examples using a nonparametric kernel method with bandwidth selection carried out via the method described in [5]. Our objective for this dataset is to assess whether the mean population size of girls is statistically equivalent to that of boys or not. The results, as depicted in Fig. 6 (d), indicate a p-value of 13%.



Fig. 4. (a) The observed PDFs of beta distribution and (b) their associated SRDFs with two different colors: The first population (blue) and the second population (red). (c) The truncated SRDFs with estimated Fréchet means (green). (d) The acceptance region CR^{c}_{α} (blue) with test statistic value (red) and *p*-value (gray).



Fig. 5. (a) The observed PDFs of inverse gamma distribution and (b) their associated SRDFs with two different colors: The first population (blue) and the second population (red). (c) The truncated SRDFs with estimated Fréchet means (green). (d) The acceptance region CR^{α}_{α} (blue) with test statistic value (red) and *p*-value (gray).



Fig. 6. (a) The observed PDFs from growth charts and (b) their associated SRDFs with two different colors: Boy (blue) and girl (red). (c) The truncated SRDFs with estimated Fréchet means (green). (d) The acceptance region CR^{c}_{α} (blue) with test statistic value (red) and *p*-value (gray).

The second dataset comprises hand force signals obtained from a population consisting of 60 healthy individuals and 90 patients with arthritis [22]. The medical protocol involved the continuous measurement of hand force over a period of time, with the objective of studying endurance during a test. Consequently, it is expected that members of the healthy group would exhibit greater hand strength compared to patients experiencing pain. In Fig. 7 (a), we present some PDFs generated by registering and normalizing arthritis curves. Our objective



Fig. 7. (a) The observed PDFs from arthritis curves and (b) their associated SRDFs with two different colors: Patient (blue) and healthy (red). (c) The truncated SRDFs with estimated Fréchet means (green). (d) The acceptance region CR^{c}_{α} (blue) with test statistic value (red) and *p*-value (gray).

with this dataset is to ascertain whether the mean functions of the healthy and patient samples are statistically equivalent or not. The findings, as depicted in Fig. 7 (d), indicate that we can only accept the equality between means at a significance level lower than 11%.

6.3 Comparison

In this section, we use real data to the proposed methods against some stateof-the-art methods, providing a robust means to assess the disparities in means between two populations within each dataset. When working with functional data, a simple approach involves running individual tests on each time point to identify potential differences. However, this strategy may lead to situations where individual tests indicate no significant difference, even though a collective examination of all variables may reveal significant distinctions. To better suit the characteristics of functional data, we opt for the "interval testing procedure" designed for two populations [16] as a more appropriate testing method. For the sake of comparison, we consider three different functional bases: B-spline, Fourier, and phase-amplitude Fourier. Specifically, we employ the fdatest R-package, which necessitates that functional data be evaluated on a uniform grid.

Method	Growth charts	Arthritis curves
Dataset		
B-spline	18%	15%
Fourier	15%	25%
Phase-amplitude Fourier	12%	64%
Proposed	13%	11%

 Table 1. p-values of various methods.

Table 1 summarizes the *p*-values from various methods used in our analysis. Our proposed test outperforms B-spline, Fourier, and Phase-amplitude Fourier tests for arthritis data but is slightly less effective for growth chart data compared to Phase-amplitude Fourier. Specifically, our test yields *p*-values of approximately 13% for growth data and 11% for arthritis data, smaller than comparison tests. This indicates a significant difference in means between populations (girls and boys) for growth data, while other methods may not detect it. Our approach, utilizing PDFs with the Fisher-Rao metric, excels in detecting skewness, leading to a highly efficient hypothesis test.

7 Conclusion

We have introduced a new framework for evaluating the validity and the performance of binary decision models of manifold-valued data with applications on spherical data and probability density functions. The proposed methods are flexible, computationally efficient, and extendable to other manifolds. The experimental results demonstrate demonstrate their robustness with simulation studies and their consistency with real data. Additionally, we have provided comparisons and have shown, very often, better performance compared to other methods.

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The Future is Different: Predicting Reddits Popularity with Variational Dynamic Language Models

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Abstract. Large pre-trained language models (LPLM) have shown spectacular success when fine-tuned on downstream supervised tasks. It is known, however, that their performance can drastically drop when there is a *distribution shift* between the data used during training and that used at inference time. In this paper we focus on data distributions that naturally change over time and introduce four REDDIT datasets, namely the WALLSTREETBETS, ASKSCIENCE, THE DONALD, and POLI-TICS sub-reddits. First, we empirically demonstrate that LPLM can display average performance drops of about 79% in the best cases, when predicting the popularity of future posts. We then introduce a methodology that leverages neural variational dynamic topic models and attention mechanisms to infer temporal language model representations for regression tasks Our models display performance drops of only about 33% in the best cases when predicting the popularity of future posts, while using only about 7% of the total number of parameters of LPLM and providing interpretable representations that offer insight into real-world events, like the GameStop short squeeze of 2021. Source code to reproduce our experiments is available online.

Keywords: Temporal Large Language Models \cdot Prediction \cdot Topic Models

1 Introduction

The modern natural language processing (NLP) paradigm leverages massive datasets, data-scalable (deep) attention mechanisms and minimal inductive biases [36] in the form of large pre-trained language models (LPLM) which are subsequently fine-tuned on new learning tasks and datasets [6,31,32].¹ This approach has proven a tremendous success in supervised learning, where applications

¹ https://github.com/cvejoski/Supervised-Dynamic-LLM

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include question answering, sentiment analysis, named entity recognition and textual entailment, just to name a few (see e.g. [13,21,24,29,30] for a review). Yet, many recent works have also reported the strong sensitivity of LPLM to *distribution shifts* between the data used during training and that used at inference time [7,16,22,27,40]. In other words, fine-tuned LPLM are known to suffer significantly at zero-shot when applied to different data domains.



Fig. 1. Dynamic features of WALLSTREETBETS. Top row: 30 most frequent words for all documents collected within the first, middle and last 8-week windows of the WALL-STREETBETS datasets. Middle row: time evolution of topic proportions from three randomly chosen topics, inferred by a neural Dynamic Topic Model. Bottom row: top 30 words associated with each of the topics in the middle row. The figure illustrates that topic representations, as opposite to simple Bag-of-Word representations, capture (some of) the dynamic components of the dataset.

In this work we focus on a particular type of natural distribution shift which arises in documents collected over long periods of time. Indeed, document collections such as magazines, academic journals, news articles and social media content not only feature trends and themes that change with time, but also employ their language differently as time evolves [11]. LPLM fine-tuned on documents collected up to some given time (i.e. on a given observation time window) might therefore perform poorly when evaluated in future documents, if the latter *differ enough* from the previously observed ones (in either content or language usage). That is, if the dataset of interest evolves in a non-stationary fashion over time. The question is then how to characterize such non-stationary features.

Recent studies have shown that the performance of LPLM on downstream tasks, like classification or named entity recognition, does indeed degrade when the target and train distribution are temporarily misaligned. We complement these works by investigating the performance of LPLM when *predicting* the popularity (i.e. number of comments or reactions) of REDDIT post—the news aggregator, content rating and discussion website, given the history (and content) of past ones.

In what follows we introduce and study four datasets that we extract from REDDIT. We are principally concerned with the two questions, namely (i) do these datasets exhibit strong enough distribution shift across time to affect the performance of LPLM, fine-tuned on the history of past posts? and, if so, (ii) how can we deal with such natural domain change problems? Each of the RED-DIT datasets consists, as usual, of a single sequence of document collections (i.e. each time point in the sequence consists of an aggregate of documents with a given timestamp). This practical aspect entails, in particular, that the inference of representations capturing their relevant dynamic components (i.e. the non-stationary features from above)—if at all present—should be done via low capacity models. Bayesian generative models for sequences, such as Kalman filters or Gaussian processes, are good examples well suited for such a task, and Fig. 1 illustrates this point. In the first row of the figure we report the 30 most frequent words from all documents collected within the first, middle and last 8-week windows of one of our datasets, which spans about one year in total. There is no discernible change between these three time points—in this representation of the data—and one might jump to the conclusion that the dataset shows no dynamics (or that the data distribution is stationary). In the second row, however, we report how the proportions of three randomly chosen topics, inferred via a neural variational variant of Dynamic Topic Models (DTM) [5], changes as time evolves. Some of the dynamic features of the dataset are now evident in this representation. Note that the last row in the figure shows the top 30 words associated with each of the topics in the second row.

Our first contribution is to (empirically) show that LPLM, the likes of BERT [13] and RoBERTa [24], fine-tuned on histories of past REDDIT posts, display average performance drops of about 79% (in the best case!) when predicting the popularity of future posts. In sharp contrast, we shall observe that LPLM perform very well on test sets extracted from the history of past posts. This result thus responds affirmatively to our first question above.

Our second contribution consists of an novel methodology that explicitly models the dynamic components of the data, to deal with the kind of temporal distribution shift we observe in REDDIT. Indeed, we strive to retain the expressiveness of neural language models (NLM) for treating the low-level word statistics composing the posts, while deploying DTM for encoding the kind of high-level document sequence dynamics shown in Fig. 1. If one interprets the inferred topics as representing the *domains* of the dataset, their inferred dynamics can, at least in principle, account for (some aspects of) the temporal domain changes present in the dataset. Note that taking the view of topics as domains to deal with distribution shift problems has also been taken in the past, albeit in static settings [15, 18, 28]. A bit more in detail, our approach consists of mainly two components. First, we use neural variational DTM to infer the time- and document-dependent proportions of a set of latent topics that best describe the data collections. We represent this set of topics via learnable topic embeddings. Second, we deploy NLM to encode the word sequences composing the posts into sequences of contextualised word representations. Given these word representations, we then modify a recently proposed attention mechanism [37] to construct temporal post representations sensitive to the temporal domain changes. These depend on both the NLM word representations of the post in question and the history of the dataset, as represented by the latent topics and their time-dependent proportions. The resulting representations can be used to predict the popularity of future posts.

Below we show our approach significantly outperforms LPLM. Indeed, our models display performance drops of only about 33% (in the best cases) when predicting the popularity of future posts, while using only about 7% of the total number of parameters of LPLM, and providing interpretable representations that offer insight into real-world events, like the GameStop short squeeze of 2021.

2 Related Work

On Natural Distribution Shifts in NLP. WILDS, the benchmark introduced by Koh et al. [19], is a very recent dataset collection which explores different types of real-world distribution shifts. Section 8 of Koh et al. [19] focus on nontemporal distribution shifts in NLP and we refer the reader to it for details and references. Additional to these are the aforementioned works which use topic models in static settings, to tackle domain change problems [15, 18, 28], as well as those works which report performance drops of LPLM, again under nontemporal distribution shifts [7, 16, 27, 40].

Koh et al. [19] includes in Appendix F a section about *temporal* distributions shift for review data. They report only modest performance drops, a result we do not find surprising. Review data typically deals with items (e.g. products, restaurants, etc.) whose basic features change moderately with time (see also Luu et al. [26] for similar observations). In contrast, several recent works report significant performance drops when the target and train distribution are temporarily misaligned [1–3, 14, 17, 22, 23, 25, 26, 33, 35]. For example, Lazaridou et al. [22] systematically shows that Transformer-XL language models perform worse when used for predicting future utterances not included in the training period.

Some of these works propose to deal with the temporal distribution shift by incorporating temporal information directly into the model, either as input [14] or into the self-attention mechanism itself [34]. Yet none of them model time *explicitly*.

On Dynamic Language Models for Supervised Downstream Tasks. Incorporating temporal information into (neural) models of text is key to capture the constant state of flux typical of streaming text datasets, the likes of news article collections or social media content. Very early in the game, Yogatama et al. [39] considered

using temporal, non-linguistic data to condition n-gram language models and predict economics-related content at a given time. More recently Delasalles et al. [12] learned, via recurrent neural models, hidden variables encoding time information, which are then used both to condition neural language models and in classification tasks. Similarly, Cvejoski et al. [10] leveraged recurrent neural point process models to infer dynamic representations that help model both content and arrival times of Yelp reviews. Other recent work have also used both temporal and text information, but to predicting review ratings [9,38] instead. Different from all these works, we use DTM to infer and explicitly model the dynamic components of our corpora, and attention mechanisms to connect them with neural language models for regression tasks.

3 Model



Fig. 2. Left: D-TAM-GRU model. The input to the model is the document's word sequence $\mathbf{X}_{t,d}$ and BoW representations $\mathbf{w}_{t,d}$ and \mathbf{W}_t , the output is a continuous label r_d . The model consists of four components: (i) the DTM, (ii) the NLM encoder, (iii) the attention module and (iv) a regressor module, which takes the output from the attention module and predicts the label r_d . Right: Detailed view of the Attention Module. This module takes as input the document's word representations $\mathbf{u}_{1:N}^d$ from the NLM and the topic proportion $\boldsymbol{\theta}_{t,d}$ from the DTM. See equations in the main text for the definition of all these variables.

In this section we propose a novel methodology to deal with a class of distribution shift that naturally arises in corpora collected over long periods of time. Suppose we are given an ordered collection of corpora $\mathcal{D} = \{D_1, D_2, \ldots, D_T\}$, so that the *t*th corpus D_t is composed of N_t documents (REDDIT posts), all received within the *t*th time window. Let the *d*th document in D_t be defined by the tuple $(\mathbf{w}_{t,d}, \mathbf{X}_{t,d}, r_{t,d})$, where $\mathbf{w}_{t,d}$ denotes the Bag-of-Words (BoW) representation of the document, $\mathbf{X}_{t,d} = (\mathbf{x}_1^{t,d}, \ldots, \mathbf{x}_M^{t,d})$ denotes the sequence of words comprising the document and $r_{t,d}$ labels the document's rating. Similarly, let \mathbf{W}_t denote the BoW representation for the entire document set within D_t . Given a new document d' at time T + 1—that is, given $\mathbf{X}_{T+1,d'}, \mathbf{w}_{T+1,d'}$ —the task is to predict the rating $r_{t+1,d'}$.

Our approach takes the perspective that latent topics, understood as word aggregates grouped together by means of word co-occurrence information within the corpora, can be understood as representing the *domains of the dataset*. To allow the distribution of topics within the documents to change with time allows, at least in principle, to model domain changes within the collection as time evolves. We shall use such temporal information to weight the importance of the words composing the new document $\mathbf{X}_{T+1,d'}$, with respect to the topic (domain) proportions in that document, at time T + 1, by means of an attention mechanism.

The model thus requires the introduction of two components, namely (i) a neural variational DTM and (ii) an attention module that leverages the representations obtained by both the DTM and a low-level NLM, to create a temporal REDDIT post representation. The resulting representation can then be used by a neural regressor model to predict $r_{t+1,d'}$.

In the following we introduce all the different components of the model in detail.

Neural Variational Dynamic Topic Model. Let us suppose the corpora collection \mathcal{D} is described by a set of K unknown topics (domains). We then assume there is a sequence of global hidden variables $\eta_1, \ldots, \eta_T \in \mathbb{R}^{\dim(\eta)}$ which encodes how the topic proportions change among the corpora as time evolves (i.e. as one moves from D_t to D_{t+1}). We also assume there is a *local* hidden variable $\zeta_{t,d} \in \mathbb{R}^{\dim(\zeta)}$, conditioned on η_t , which encodes the content of the dth document in D_t , in terms of the K topics.

Generative Model. Let us denote with ψ the set of parameters of our generative model. We generate the *d*th document in D_t by first sampling the topic proportions $\boldsymbol{\theta}_{t,d} \in [0,1]^K$ as follows

$$\boldsymbol{\eta}_t \sim \mathcal{N}\left(\boldsymbol{\mu}_{\psi}^{\eta}(\boldsymbol{\eta}_{t-1}), \delta \mathbf{I}\right),$$
(1)

$$\boldsymbol{\zeta}_{t,d} \sim \mathcal{N} \left(\mathbf{W}_{\psi}^{\zeta} \, \boldsymbol{\eta}_t + \mathbf{c}_{\psi}^{\zeta}, \, 1 \right), \tag{2}$$

$$\boldsymbol{\theta}_{t,d} = \operatorname{softmax}(\mathbf{f}_{\psi}^{\zeta}(\boldsymbol{\zeta}_{t,d})), \tag{3}$$

where $\mathbf{f}_{\psi}^{\zeta} : \mathbb{R}^{\dim(\zeta)} \to \mathbb{R}^{K}$ is neural network with parameters in ψ and $\mathbf{W}_{\zeta} \in \mathbb{R}^{\dim(\zeta) \times \dim(\eta)}, \mathbf{c}_{\xi} \in \mathbb{R}^{\dim(\zeta)} \subset \psi$ are trainable parameters. Furthermore, and just as in Deep Kalman Filters [20], η_t is Markovian and evolves under a Gaussian noise with mean $\boldsymbol{\mu}_{\psi}^{\eta} : \mathbb{R}^{\dim(\eta)} \to \mathbb{R}^{\dim(\eta)}$, defined via a neural network with parameters ψ , and variance δ . The latter being a hyperparameter of the model. Finally, we choose the prior $\eta_1 \sim \mathcal{N}(0, 1)$.

Once we have $\boldsymbol{\theta}_{t,d}$ we generate the corpora sequence by sampling

$$z_{t,d,n} \sim \text{Categorical}(\boldsymbol{\theta}_{t,d}), \qquad \qquad w_{t,d,n} \sim \text{Categorical}(\boldsymbol{\beta}_{z_{t,d,n}}), \qquad (4)$$

where $z_{t,d,n}$ is the time-dependent topic assignment for $w_{t,d,n}$, which labels the *n*th word in document $d \in D_t$, and $\beta \in \mathbb{R}^{K \times V}$ is a learnable topic distribution over words. We define the latter as

$$\boldsymbol{\beta} = \operatorname{softmax}(\boldsymbol{\alpha} \otimes \boldsymbol{\rho}), \tag{5}$$

with $\boldsymbol{\alpha} \in \mathbb{R}^{K \times E}$, $\boldsymbol{\rho} \in \mathbb{R}^{V \times E}$ learnable topic and word embeddings, respectively, for some embedding dimension E, and \otimes denoting tensor product.

Neural Topic Attention Model. Given a document d, composed of the word sequence $\mathbf{x}_1, \ldots, \mathbf{x}_M$, the task is to predict its rating r_d . One straightforward approach to this problem is to consider a (deep) neural language model to infer contextualized word representations of the form $\mathbf{u}_1, \mathbf{u}_2, \ldots, \mathbf{u}_M = s_{\rho}(\mathbf{x}_1, \mathbf{x}_2, \ldots, \mathbf{x}_M)$, where s_{ρ} is any neural sequence processing model (as e.g. a GRU [8] or BERT model) with parameters ρ .

With the representations $\mathbf{u}_1, \mathbf{u}_2, \ldots, \mathbf{u}_M$ at hand one can define a summary representation \mathbf{s}_d (by e.g. averaging over the **u**'s or using the CLS token of BERT-like models) and use it as input to a neural regressor \mathbf{f}_{ρ}^r (also with parameters ρ) to predict r_d . Yet, if the document d is received at time T + 1, i.e. out of the observation window on which the parameter ρ was optimised, and if the dataset in question displays temporal domain changes, we might expect this simple model to underperform.

Here we use the DTM of Sect. 3, which is assumed to model the domain changes in the corpora, to define a temporal summary representation sensible to the natural distribution shifts of the dataset. Indeed, we follow Wang et al. [37] and use an attention mechanism to construct \mathbf{s}_d as follows

$$\mathbf{s}_{t,d} = \sum_{j}^{M} \sum_{i}^{K} (\theta_{t,d}^{i} - \delta) \times \operatorname{softmax}(\operatorname{MLP}(\mathbf{u}_{j})^{T} \cdot \boldsymbol{\alpha}_{i}) \odot \mathbf{u}_{j},$$
(6)

where the softmax function is taken with respect to the word sequence, the α_i label the set of K global topic embeddings and the $\theta_{t,d}$ denote the time- and document-dependent topic proportions, as defined in Eq. 3. To make the obtained attention weight more diverse among topics we substract from the mixture of topics constant δ , where $0 < \delta < 1$.

It follows that the proposed document representation is nothing but the sum of the projections of each of the document's word representations onto topic space, weighted by the *time-dependent* topic proportions of each dimension. It can also be understood as an attentive representation in which each word is queried by the weighted topics.

In what follows we let \mathbf{s}_{ρ} be modeled by a GRU network [8] and name the model thus defined D-TAM-GRU. Our proposed framework is shown in Fig. 2.

Inference Model. We approximate the true posterior distribution of the hidden variables with a variational (and structured) posterior of the form

$$q_{\varphi}(\boldsymbol{\eta}_{t}, \boldsymbol{\zeta}_{t,d} | \mathbf{w}_{t,d}, \mathbf{W}_{1:T}) = \prod_{t}^{T} q_{\varphi}(\boldsymbol{\eta}_{t} | \boldsymbol{\eta}_{1:t-1}, \mathbf{W}_{1:T}) \times \prod_{d}^{N_{t}} q_{\varphi}(\boldsymbol{\zeta}_{t,d} | \mathbf{w}_{t,d}, \boldsymbol{\eta}_{t})$$
(7)

where $\mathbf{W}_{1:T} = (\mathbf{W}_1, \dots, \mathbf{W}_T)$ is the ordered sequence of BoW representations for the corpus collection and φ labels the variational parameters. The posterior distribution over the local variables is chosen to be Gaussian

$$q_{\varphi}(\boldsymbol{\zeta}_{t,d}|\mathbf{w}_{t,d},\boldsymbol{\eta}_{t}) = \mathcal{N}(\boldsymbol{\mu}_{\varphi}^{\zeta}[\mathbf{w}_{t,d},\boldsymbol{\eta}_{t}], \boldsymbol{\sigma}_{\varphi}^{\zeta}[\mathbf{w}_{t,d},\boldsymbol{\eta}_{t}]),$$
(8)

with μ_{φ}^{ζ} and σ_{φ}^{ζ} neural networks with parameter φ . Likewise, the posterior distribution over the global variables is also Gaussian, but now depends not only on the latent variables at time t - 1, but also on the entire sequence of BoW representations $\mathbf{W}_{1:T}$. Explicitly we write

$$q_{\varphi}(\boldsymbol{\eta}_t | \boldsymbol{\eta}_{t-1}, \mathbf{W}_{1:T}) = \mathcal{N}(\boldsymbol{\mu}_{\varphi}^{\eta}[\boldsymbol{\eta}_{t-1}, \mathbf{h}_T^{\eta}], \boldsymbol{\sigma}_{\varphi}^{\eta}[\boldsymbol{\eta}_{t-1}, \mathbf{h}_T^{\eta}]),$$
(9)

where $\mu_{\varphi}^{\eta}, \sigma_{\varphi}^{\eta}$ are too given by neural networks and \mathbf{h}_{t}^{η} is a recurrent representation encoding the sequence $\mathbf{W}_{1:T}$. Indeed,

$$\mathbf{h}_t^{\eta} = g_{\varphi}^{\eta}(\mathbf{W}_t, \mathbf{h}_{t-1}^{\eta}), \tag{10}$$

with g_{φ} a neural model for sequence processing (like e.g. a GRU [8]). Figure 2 illustrates the architecture of the complete model.

Training Objective DTM. To optimize the DTM parameters $\{\psi, \varphi\}$ we minimize the variational lower bound on the logarithm of the marginal likelihood $p_{\psi}(w_{t,d,n}|\boldsymbol{\beta})$. Following standard methods [4], the latter can readily be shown to be

$$\mathcal{L}[\boldsymbol{\beta}, \boldsymbol{\psi}, \boldsymbol{\varphi}] = \sum_{t=1}^{T} \sum_{d=1}^{N_t} \left(\mathbb{E}_{\{\boldsymbol{\zeta}_{t,d}, \boldsymbol{\eta}_t\}} \Big\{ \log p_{\boldsymbol{\psi}}(\mathbf{w}_{t,d} | \boldsymbol{\beta}, \boldsymbol{\zeta}_{t,d}, \boldsymbol{\eta}_t) - \mathrm{KL}\left[q_{\boldsymbol{\varphi}}(\boldsymbol{\zeta}_{t,d} | \mathbf{w}_{t,d}, \boldsymbol{\eta}_t); p_{\boldsymbol{\psi}}(\boldsymbol{\zeta}_{t,d} | \boldsymbol{\eta}_t)\right] \Big\} \right)$$
(11)
$$-\mathrm{KL}\left[q_{\boldsymbol{\varphi}}(\boldsymbol{\eta}_1 | \mathbf{W}_{1:T}); p(\boldsymbol{\eta}_1)\right] - \sum_{t=2}^{T} \mathrm{KL}\left[q_{\boldsymbol{\varphi}}(\boldsymbol{\eta}_t | \boldsymbol{\eta}_{1:t-1}, \mathbf{W}_{1:T}); p_{\boldsymbol{\psi}}(\boldsymbol{\eta}_t | \boldsymbol{\eta}_{t-1})\right],$$

where KL labels the Kullback-Leibler divergence and β is given in (5).

Training Objective. To optimize the complete set of model parameters $\{\psi, \varphi \rangle$ we minimize the objective

$$\mathcal{L} = \mathcal{L}_{TM}[\boldsymbol{\beta}, \boldsymbol{\psi}, \boldsymbol{\varphi}] + \alpha_y \mathcal{L}_r[\boldsymbol{\rho}], \qquad (12)$$

where \mathcal{L}_{TM} denotes the DTM loss (defined in Eq. 11) and \mathcal{L}_r denotes the regression loss. The latter is defined as the root-mean-squared error between the target rating r_d and the prediction of a regressor model, that is

$$\mathcal{L}_{r}[\boldsymbol{\rho}] = \text{RMSE}(r_{d}, \mathbf{f}_{\rho}^{r}(\mathbf{s}_{d})), \qquad (13)$$

with $\mathbf{f}^r : \mathbb{R}^{\dim(s)} \to \mathbb{R}$ a neural network and \mathbf{s}_d defined by Eq. 6.

Prediction. In order to predict the rating $r_{T+N,d}$ of a new document, N steps into the future, given its word sequence $\mathbf{X}_{T+N,d}$, we rely on the generative process of our model albeit conditioned on the past. Essentially one must generate Monte Carlo samples from the posterior distribution and propagate the global latent representations into the future with the help of the prior transition function Eq. 1. This procedure is depicted on the conditional predictive distribution (for a single step) of our model

$$p(r_{T+1,d}|\mathcal{D}, \mathbf{X}_{T+1,d}) = \int p(r_{T+1,d}|\boldsymbol{\eta}_{T+1}, \boldsymbol{\zeta}_{T+1,d}, \mathbf{X}_{T+1,d}) \\ \times p(\boldsymbol{\zeta}_{T+1,d}|\boldsymbol{\eta}_{T+1}) p(\boldsymbol{\eta}_{T+1}|\boldsymbol{\eta}_{T}) \\ \times p(\boldsymbol{\eta}_{1:T}|\mathcal{D}) d\boldsymbol{\eta}_{1:T+1} d\boldsymbol{\zeta}_{T+1,d},$$
(14)

where $p(\eta_{1:t}|\mathcal{D})$ is the exact posterior over the dynamical global variables, which we approximate with our variational expression Eq. 9, and where

$$p(r_{T+1,d}|\boldsymbol{\eta}_{T+1}, \boldsymbol{\zeta}_{T+1,d}, \mathbf{X}_{T+1,d}) = \delta(r_{T+1,d} - \mathbf{f}_{\rho}^{r}(\mathbf{s}_{T+1,d}[\boldsymbol{\eta}_{T+1}, \boldsymbol{\zeta}_{T+1,d}, \mathbf{X}_{T+1,d}])), \quad (15)$$

with \mathbf{f}_{ρ}^{r} the neural regresor, $\mathbf{s}_{T+1,d}$ defined in Eq. (6) and δ the Dirac delta function.

4 Task, Dataset and Experimental Setup

In this sections we first introduce our popularity prediction task. Next, we present detailed information about our proposed REDDIT dataset for temporal distribution shifts analysis. Finally, we describe the experimental setup that we use to train and evaluate the models and introduce the baselines we compare against.

Task. We define the popularity of a given REDDIT post as the number of comments the submission receives. This quantity measures the impact of the post's content on the REDDIT community at a given moment of time. Table 1 in the Appendix contains the histograms of the number of comments per submission. We infer such distributions with a continuous variable (r_d in our notation), and frame the prediction task as regression. We thus quantify the performance of our models with the *coefficient of determination* (R^2), the root-mean-square error (RMSE) and the mean-absolute error (MAE). We report all results wrt. the last two metrics in the Appendix due to lack of space. Finally, we also evaluate the performance of the topic models, whenever these are used. See also Appendix. Table 1. Regression results (wrt R2 metric): We report our results for three different topic models, each with 25, 50 and 100 hidden topics. MLP, BERT and RoBERTa have no topic model associated to them. Gray columns show the PREDICTION (out-of-distribution) results for all models. White columns show the results for the UP-TO-DATE dataset. For each sub-column, we underline the best model. We highlight with boldface the best model overall, separately for PREDICTION and UP-TO-DATE datasets.

Model	$\mathbf{R2}^{\uparrow}$ (Prediction)			R2 \uparrow (up-to-date)		
	25	50	100	25	50	100
GG						
MLP	0.011 ± 0.012			0.0007		
Y						
BERT	-1.029 ± 0.756			0.0712		
RoBERTa	0.052 ± 0.158			0.0484		
TAM-GRU	0.075 ± 0.096	0.150 ± 0.118	0.164 ± 0.141	0.0053	0.0555	0.0449
TAM-BERT	0.054 ± 0.103	0.134 ± 0.171	0.154 ± 0.170	0.0103	0.0967	0.0867
TAM-RoBERTa	-0.019 ± 0.008	-0.019 ± 0.008	0.092 ± 0.099	-0.0175	-0.0169	0.0295
D-ST	0.008 ± 0.030	-0.010 ± 0.010 -	-0.006 ± 0.008	-0.0175	-0.0138	-0.0175
D-TAM-GRU	0.152 ± 0.105	$\underline{0.196\pm0.175}$	0.173 ± 0.161	0.0387	0.0386	0.0397
DOLITIC	-5.27 ± 13.577			0.6278		
BERT	-8.89 ± 21.951			0.6820		
RoBERTa	-0.028 ± 0.239			0.7306		
TAM-GRU	-0.448 ± 1.101	-0.050 ± 0.053 -	-0.078 ± 0.042	0.6993	0.7106	0.7251
TAM-BERT	-0.051 ± 0.024	-0.041 ± 0.121 -	-0.022 ± 0.010	0.6577	0.6608	0.5576
TAM-RoBERTa	-0.091 ± 0.047	0.014 ± 0.227	0.008 ± 0.408	-0.0191	0.6329	0.6305
D-ST	-0.072 ± 0.216	-0.191 ± 0.127	-0.178 ± 0.291	0.3370	0.3631	0.2372
D-TAM-GRU	0.133 ± 0.163	-0.078 ± 0.042 -	-0.038 ± 0.012	0.7118	0.7515	0.7028
THE DONALL	-0.006 ± 0.004			0.4162		
BERT	-0.032 ± 0.018			0.6674		
RoBERTa	0.110 ± 0.258			0.5290		
TAM-GRU	0.176 ± 0.334	0.153 ± 0.224	0.133 ± 0.225	0.5821	0.5561	0.5813
TAM-BERT	-0.009 ± 0.005	0.169 ± 0.345	0.026 ± 0.164	0.0288	0.4523	0.4157
TAM-RoBERTa	-0.116 ± 0.072	-0.036 ± 0.024 -	-0.032 ± 0.203	0.5992	0.4878	0.5906
D-ST	-0.011 ± 0.011	-0.012 ± 0.011 -	-0.011 ± 0.010	-0.0014	-0.0015	-0.0012
D-TAM-GRU	$\underline{0.203 \pm 0.278}$	$\underline{0.179 \pm 0.314}$	0.228 ± 0.369	0.6553	$\underline{0.7322}$	0.7071
WALLSTREETBETS	0.324 ± 0.662			0.5862		
BERT	-4.632 ± 12.553			0.6850		
RoBERTa	-0.003 ± 0.012			0.5484		
TAM-GRU	0.218 ± 0.530	0.274 ± 0.577	0.293 ± 0.565	0.6345	0.5645	0.5865
TAM-BERT	0.353 ± 0.188	-0.015 ± 0.017	0.153 ± 0.705	0.5652	0.5717	<u>0.8117</u>
TAM-RoBERTa	-0.014 ± 0.025	-0.008 ± 0.005 -	-0.280 ± 1.139	-0.0084	0.5994	0.0218
D-ST	0.540 ± 0.296	-0.008 ± 0.006	$\underline{0.376 \pm 0.165}$	-0.0038	-0.0022	0.1823
D-TAM-GRU	0.640 ± 0.249	0.101 ± 0.417	0.160 ± 0.103	0.6577	0.8085	0.6047

Dataset. In this work we propose a new REDDIT² dataset for temporal distribution shifts analysis. REDDIT is a news aggregator, content rating and discussion website. Users can post content on the site, like images, text links and videos, which are rated and commented by other users. The posts which are called *submissions* are organized by subject in groups or *subreddits*. We crawled the posts for the ASKSCIENCE, POLITICS, THE DONALD and the WALLSTREETBETS subreddits. The ASKSCIENCE is a subreddit in which science questions are posted and answered. POLITICS is a subreddit where news and politics in the U.S. are discussed. THE DONALD was a subreddit where supporters of former U.S. president Donald Trump were initiating discussions. Lastly, the WALLSTREETBETS is a subreddit where stock trading is discussed. This subreddit played a major role in the GameStop short squeeze that caused losses³ for some U.S. firms in early 2021.

Time Window and Out-of-Distribution Selection. After pre-processing, we first split each of the datasets into the UP-TO-DATE and PREDICTION (out-of-distribution) datasets. We create such a distinction to study temporal distribution shifts in the dataset. Specifically, we take the last 20 time points as prediction (out-of-distribution data) and the rest for the up-to-date posts (in-distribution data). In this way we ensure that we do not train on documents that come from the future, which is what we actually want to model. That is, we do not violate causality. Likewise we ensure there is a clear distinction between past and future, which will allow us to uncover temporal distribution shifts, if present.

Next, we split the UP-TO-DATE submissions randomly into train, validation and test sets (80%, 10%, 10%, respectively). Additionally, to evaluate the DTM on the document completion (i.e. generalization) task, we split the documents of the test set into two halves. The first half is used as input to the topic model; the second half is used to measure the document completion perplexity.

Baseline Models. The baseline models are introduced in order of increasing complexity. These models are generally composed of two modules, namely (i) an *encoder* module, which takes as input either the word sequence $\mathbf{X}_{t,d}$ or the BoW $\mathbf{w}_{t,d}$ of the document and outputs a summary representation $\mathbf{s}_{d,t}$; (ii) a *regressor* module, which takes as input the representation $\mathbf{s}_{d,t}$ and predicts the rating $r_{t,d}$ of the document.

The simplest baseline model we consider defines both encoder and regressor as MLPs, and takes as input the BoW representation $\mathbf{w}_{t,d}$ of the document. We name it MLP. Next we introduce baselines with attention-based models [36] as encoder and MLPs as regressor. We use two attention-based encoder architectures: BERT [13] and RoBERTa [24]. The input to these models is the word sequence $\mathbf{X}_{t,d}$ and we use their CLS embedding as input to the regressor module. The third baseline is TAM, the neural topic attention model for supervised learning

² https://www.reddit.com.

³ https://www.bloomberg.com/news/articles/2021-01-25/gamestop-short-sellers-reloadbearish-bets-after-6-billion-loss.

proposed by Wang et al. [37]. TAM combines topic models and NLM to produce the representation $\mathbf{s}_{d,t}$, just as in Eq. 6 above, but with *static* topic proportions. The original TAM version uses a GRU as NLM. We call this version TAM-GRU. We also extend this model by replacing the GRU with either BERT or RoBERTa. Accordingly we name these baselines TAM-BERT and TAM-RoBERTa, respectively. Finally, we use our DTM from Sect. 3 as encoder module, and input the inferred local hidden variable $\zeta_{t,d}$ to the regressor, which here too is defined by an MLP. We name this last baseline D-ST.



Fig. 3. Cumulative average R^2 score for each time step into the future obtained by using the predictions of the best transformer, the best *static* TAM and the best *dynamic* TAM models. Our models perform in all the datasets better than the static ones, and comparably well in the THE DONALD dataset which exhibits more stationary behavior, and in this case our model is not suitable.

Training and Evaluation Metrics. We use grid search during training to find the best hyper-parameters of each model type. All models are trained on the training subset of the UP-TO-DATE submissions, and the validation subset is used for choosing the best hyper-parameters. For all models that rely on a DTM module we use 25, 50 and 100 topics. Details regarding the values of other hyper-parameters can be found in the Appendix (section Model Training Setup).

5 Results and Discussion

In this section we discuss our results on the task of predicting the popularity of future submissions (posts) on the REDDIT platform, by predicting the number of comments the submissions will receive. As explained above, this task is defined by training all models on the UP-TO-DATE submission set and evaluating them on the PREDICTION set, which consists of submissions received in the future.

One of the key takeaways of the present work is to highlight that LPLM, finetuned on the UP-TO-DATE submissions, fails at predicting the popularity of future posts. To see this let us first examine the UP-TO-DATE results in the second column of Table 1, which show the performance of all models, evaluated on the test subset. Note how LPLM provides the best results in two out of four subreddits, yielding R^2 scores higher than the dynamic models, including our D-TAM-GRU. In contrast, Table 1 first column shows the PREDICTION (out-of-distribution) results. Comparing the performance of LPLM on the in-distribution set against



Fig. 4. Time evolution of topics' proportion. The time series are obtained by taking the mean and two standard deviations of the activity θ_k of topic k in all the documents for a given time step t. We present three randomly picked topics for each dataset. One could immediately notice that there is almost no dynamics for the THE DONALD dataset.

their performance on the out-of-distribution set, we observe performance drops of about 79% (in the best cases!). LPLM thus fail at predicting the popularity of future posts, and we understand these findings as being consequence of temporal distributions shifts between the UP-TO-DATE and PREDICTION sets.

The second important observation we can make from Table 1 is that, D-TAM-GRU not only outperforms all LPLM on the PREDICTION dataset, but also displays performance drops of only about 33% (in the best cases) when predicting the popularity of future posts. Thus, our methodology does in fact help dealing with the temporal distributions shifts of the REDDIT dataset.

Our ablation study also shows that (static) topic attention models expand the capabilities of LPLM. See Table 1 and Fig. 3. These findings comply with results in the literature, which indicate that topic models largely improve model performance under distribution shifts [15, 18, 22, 28]. In fact, TAM-GRU (the best static model for prediction) displays performance drops of 40% in the best cases. Comparing this value with the performance drop of its dynamic counterpart, we can conclude that explicitly modelling the dynamic components of the data makes language models more robust against temporal distribution shifts.

Finally, Fig. 4 displays the time series for the topic proportions of three randomly selected topics from each dataset. Note how the topics exhibit a different range of dynamic behavior, accounting for seasonality, trendiness and bursty as well as simply random behavior. As a whole, the ability of D-TAM-GRU to leverage such dynamic information in the prediction task strongly depends on the nature of these dynamical patterns, as well as the overall weight obtained by the topic attention mechanism. Indeed, one requires enough *relevant* topics with dynamical information. In what can be thought of as a kind of distributed signal-to-noise ratio, we speculate that in order for the prediction capabilities of NLMs to be improved by our approach, the dataset at hand must be such that there are enough topics with non-stationary behavior, i.e. topics that exhibit a distribution change over time, and that such topics are important for the prediction task, above other topics with stationary dynamical behavior (i.e. no change in time). THE DONALD dataset shows qualitative behavior that is overall stationary, as the topic proportions present strong noisy behavior, which explains why D-TAM-GRU performs comparably to Roberta in Fig. 3.



Fig. 5. Evolution of topic proportion in the WALLSTREETBETS dataset. Topics showing community culture (profane jargon) and the behavior of the population during to the "GameStop" short squeeze event.

Interpretability and WALLSTREETBETS. An added advantage of our model is the interpretable character of the representations inferred by our DTM, as we have seen in Fig. 4. Let us now take a look at Fig. 5. The WALLSTREETBETS subreddit has become a success story when it comes to the power the Web has to impact society, as retail investors organized themselves in the platform to create major shifts in the stock market, thereby playing a major role in short squeeze of the "GameStop" stock (an American video game retailer). This event can be directly observed in Fig. 5-right, where the model inferred a rapid increase in the importance of a topic about "gme" (ticker value of the "GameStop" stock), previous to the sudden increase of the stock price by January 28, 2021. Now, due to the rapid increase of the stock price, some brokerages such as "Robinhood" halted trading. The reaction of the community to this decision can also be observed in the rise of the topic shown in Fig. 5-middle, in which"Robinhood" is paired with derogatory jargon. New insights into the behavior of the population are uncovered by our model too. Figure 5-left shows how profane-related topic decays in importance with time, which means that the language used by the WALLSTREETBETS subreddit community is shifting. Beyond these qualitative results, the ability of our model to predict the popularity of posts, allows us to quantify the impact of several topics in the platform, as well as to predict popularity shifts within the user population. As WALLSTREETBETS continues to gain ground with retail investors, our methodology opens a window to quantitatively study possible future rises in the popularity of futures stocks in the REDDIT platform.

6 Conclusion

We studied the prediction capabilities of LPLM and showed that, for a newly introduced dataset with rich dynamic behavior, temporal distribution shifts cause a sharp drop in their performance. We introduced a neural variational DTM with attention that outperforms LPLM and overcomes (some of) the difficulties created by the temporal distribution shifts. Remarkably, our models use only about 7% of the total number of parameters of LPLM and provide interpretable representations that offer insight into real-world events.

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CIRCUITVQA: A Visual Question Answering Dataset for Electrical Circuit Images

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Abstract. A visual question answering (VQA) system for electrical circuit images could be useful as a quiz generator, design and verification assistant or an electrical diagnosis tool. Although there exists a vast literature on VQA, to the best of our knowledge, there is no existing work on VQA for electrical circuit images. To this end, we curate a new dataset, CIRCUITVQA, of 115K+ questions on 5725 electrical images with ~70 circuit symbols. The dataset contains schematic as well as hand-drawn images. The questions span various categories like counting, value, junction and position based questions. To be effective, models must demonstrate skills like object detection, text recognition, spatial understanding, question intent understanding and answer generation. We experiment with multiple foundational visio-linguistic models for this task and find that a finetuned BLIP model with component descriptions as additional input provides best results. We make the code and dataset publicly available (https://github.com/rahcode7/Circuit-VQA).

Keywords: Visual Question Answering \cdot Vision and Language \cdot Multimodal Large Language Models \cdot VQA for circuits \cdot electricalVQA \cdot circuitVQA

1 Introduction

A visual question answering (VQA) system for electrical circuit images could be useful in several scenarios. It can be used as a teaching tool or a quiz generator for students who are learning about electrical circuits. It can also provide feedback and hints to help students solve circuit problems. It can be used as a design assistant or a verification tool for engineers who are creating or modifying electrical circuits. It can also suggest improvements or optimizations for the circuit design. It can be used as a debugging or a diagnosis tool for technicians who are repairing or testing electrical circuits. It can also identify faults or errors in the circuit functionality or performance. Such a system can also be used as an



Fig. 1. Sample circuit image from CIRCUITVQA, and question-answer pairs per question type

accessibility tool for visually impaired users who want to interact with or learn about electrical circuits. Finally, it can serve as an analysis tool for researchers who want to study or compare electrical circuits.

VQA aims at answering a text question in the context of an image [3]. Several VQA datasets have been proposed in the literature [3,12,21,37]. Most methods for VQA use basic multimodal fusion of language and image embeddings [20], attention-based multimodal fusion [42] or neural module networks [2,14]. More recently, newer problem settings have been proposed as extensions of the basic VQA framework like Text VQA [36], Visual Dialog [8], Video QA [44], retrieval-based VQA [32] and knowledge-based VQA for videos [11,13]. Extending on this rich literature, we propose a novel problem setting: VQA in the context of electrical circuit images.

With the advent of Transformers, multiple vision language models (VLMs) [22,23,40], have been proposed and they have showcased remarkable capabilities for VQA datasets. However, such models do not directly generalize to out-of-domain samples especially when there is large variety. Electrical circuit images can be very complex and diverse, with different layouts, components, symbols, labels, and connections. Hence, we first curate a novel dataset, CIRCUITVQA, to finetune these models. CIRCUITVQA has 115K+ questions across 5725 circuit images with ~71 popular circuit symbols. Questions in CIRCUITVQA have been designed such that, to be able to handle different types of questions in the dataset, a model should possess skills like object recognition, attribute classification, counting, spatial reasoning, common sense reasoning, and circuit analysis. CIRCUITVQA comprises a mix of hand-drawn as well as schematic images. Figure 1 shows a few examples of such questions for a sample image.

We systematically evaluate performance of popular vision-language models for various types of questions: counting, value-based, junction-based and position-based. We also investigate the effectiveness of various kinds of input image representations to be passed as input to these models: just the raw image itself, optical character recognition (OCR) and object detection outputs. We also explore if augmenting the text question with description of relevant electrical components helps improve the performance. Overall, we make the following contributions in this paper. (1) We construct a novel and diverse circuit-based VQA dataset, CIRCUITVQA, with 115K+ questions. The dataset contains carefully designed questions across 5 types which test multiple visio-linguistic skills of multimodal models. (2) We conduct a holistic evaluation of state-of-the-art vision language models: (a) fine-tuning based evaluation of BLIP [23], GIT [40] and Pix2Struct [22] on train part of CIRCUITVQA, and (b) zero-shot evaluation of instruction-tuned models like LLaVA [26], InstructBLIP [7] and GPT4V [31]. (3) We conduct extensive experimentation by combining external modules like Optical Character Recognition (OCR), Object detection and supplying detailed description of electrical components to improve and understand the capabilities of these vision-language models for the CIRCUITVQA task. (4) We propose a novel hallucination score based metric that can be widely applied to any VQA task.

2 Related Work

VQA for Science: Unlike general VQA [3] which focuses on natural images, VQA for Science is a subfield of VQA that focuses on answering questions about scientific images, such as diagrams, graphs, charts, and illustrations. Popular datasets include ScienceQA [27] (on science lectures), AI2D [19] (on diagrams), ChartQA [28] (on chart summaries), FigureQA [18] (on scientific-style figures from five classes: line plots, dot-line plots, vertical and horizontal bar graphs, and pie charts), DVQA [17] (on bar-charts), PlotQA [29] (on plots), LeafQA [5] (on figures/charts), and BizGraphQA [4] (on graph-structured diagrams from business domains). Although these datasets contain diverse range of diagrams, they do not particularly contain any questions related to electrical circuits.

ML for Electrical Circuits: The increasing complexity of electronic design automation (EDA) tasks has aroused large interest in incorporating ML to solve EDA tasks [15] and electronic circuit design. Tasks include recognition of hand-drawn electrical and electronic circuit components, and fault diagnosis of analog circuits. In this work, we extend this line of work by introducing the task of VQA for circuit images.

Hallucinations for VLMs: For VLMs, hallucination [16] refers to contradictions between the visual input (taken as 'fact') and the text output of a VLM [24,35]. CHAIR [35] evaluates object hallucinations in image captioning by quantifying differences of objects between model generation and ground-truth captions. POPE [24] formulates a binary - yes or no questions about the object presence in the images such as "Is there a person in the image?". Since no specific hallucination evaluation metrics have been proposed for the VQA task specifically, we fill that gap by proposing a new metric, HVQA, in this paper.

	Type	# Images	Description	Frequent Object Classes
D1	Schematic	1284	Electrical circuits with 7141 annotations for object detection across 7 classes.	resistor, current-source, inductor, capacitor, voltage-ac, voltage-dc, arrow
D2	Hand-drawn	2304	Hand-drawn electrical circuit diagram images as well as 212K bounding box annotations across 59 object classes, and segmentation ground-truth files. Also has junction, cross-over and text annotations.	resistor, terminal input, diode, transistor, GND, LED, voltage, thyristor, switch, inductor, VSS, speaker, AND, NOT, varistor
D3	Hand-drawn	487	Electrical circuits with 8353 annotations for object detection across 14 classes.	junction, text, resistor, current-source, inductor, capacitor-unpolarized, voltage-dc, voltage-dc_ac, multi-cell-battery, gnd, diode, terminal, single-cell-battery, crossover
D4	Schematic	1273	Digital circuit images with 2398 annotations for object detection across 7 classes.	and, nand, not, or, xor, nor, xnor
D5	Hand-drawn	1679	Electrical circuits with 58K annotations for object detection across 45 classes.	junction, text, resistor, terminal, diode, capacitor-unpolarized, crossover, transistor, gnd, inductor, voltage-dc, thyristor, switch

Table 1. Details of source datasets for CIRCUITVO

3 CIRCUITVQA Dataset Curation and Analysis

In this section, we discuss two aspects of the CIRCUITVQA dataset construction: (a) collecting circuit images from various sources, (b) generating question answer pairs using either human annotations or automatically using available metadata.

3.1 Collection of Circuit Images

We gather the images in CIRCUITVQA from five datasets available on public platforms like Roboflow and Kaggle. The original source of many of these datasets can be traced back to the Handwritten Circuit Diagram Images (CGHD) [38]. These images are of two types: schematic and hand-drawn. Besides the images, the dataset contains metadata like human annotated bounding boxes and the corresponding component classes like resistor, ammeter etc. Table 1 shows details of the five source datasets: Roboflow Circuit recognition $(D1)^1$, Kaggle CGHD $(D2)^2$, Roboflow CGHD-Supplement $(D3)^3$, Roboflow Circuit Recognition Electronics $(D4)^4$ and Roboflow CGHD-Full Supplement $(D5)^5$. D1

¹ https://universe.roboflow.com/rp-project/circuit-recognition.

² https://www.kaggle.com/datasets/johannesbayer/cghd1152.

 $^{^{3}\} https://universe.roboflow.com/development-tohnm/cghd-supplement-g34fl.$

⁴ https://universe.roboflow.com/rp-project/circuit-recognition-electronics/.

 $^{^{5}}$ https://universe.roboflow.com/development-tohnm/cghd-full-supplemented.

and D4 are schematic while others are hand-drawn. The datasets also differ in terms of the kind of electrical components. While D1 has just 7 object classes, D2 has 59. We aggregate data across these five datasets leading to a collection of 7027 images. Next, we identify potential duplicate images using perceptual hashing [43]. We then keep only one copy of these images by deleting similar ones with a Hamming distance >3. This leads to our final unified dataset of 5725 images of which 3175 are hand-drawn and 2550 are schematic. We make the dataset publicly available¹.

3.2 Generation of Question Answer Pairs

We generate five categories of questions: Simple Counting, Spatial counting, Position based, Value Based and Junction based. Figure 1 shows example questionanswer pairs for each question type for a sample circuit image. To generate these questions, we utilize the metadata associated with the images like the associated components and their bounding boxes. For each type, we obtain question templates using ChatGPT [30] and then instantiate questions using these templates. A full list of generated question templates is mentioned in Table 2. In the following we discuss the question-answer generation process for each question type. Table 3 summarizes the answer type for every type of question.

1. Simple Counting Questions. Given an image, in a simple counting question, we ask for the count of each component type in the image. We prompt Chat-GPT with this prompt: "Paraphrase the following text in 20 ways - How many X does the circuit have?" This leads to 20 different paraphrases which are used as question templates to generate simple counting questions in CIRCUITVQA. For every image, we randomly sample a question template and replace the placeholder X with the actual component name to get an instantiated question. This can be done because each image has the component names and their counts as associated metadata. The metadata is also used to obtain the actual answer. Answering such questions requires a model to possess object recognition and counting skills.

2. Spatial Counting Questions. Given an image, in a spatial counting question, we ask how many components of a certain type are connected directly to the left, right, top or bottom of the given component. Thus, for datasets D1, D2, D3 and D5, we use this question template "How many Y are connected directly to the $\langle \text{direction} \rangle$ of X?" where direction can be any of left, right, top or bottom. For dataset D4 which is based on digital gates, we use the following question templates: "How many gates are providing an input to X?", "How many gates are connected to the right of X?", "How many Y gates are connected to the right of X?", and "How many Y gates are connected to the left of X?" For every image in these datasets, we randomly sample a question template and replace the placeholders X and Y (from the set of components mentioned in metadata) with the actual component name to get an instantiated question. Since there is no automated way of generating an answer using associated metadata, we perform human annotation to annotate answers. The first author performed

Question Type	Question Templates
Simple Counting	How many Xs are there in the specified circuit? What number of X are included in the given circuit? What is the total count of Xs in the circuit? Can you determine the number of Xs in the circuit? How numerous are the Xs in the circuit? What is the quantity of Xs present in the circuit? Are there multiple Xs in the circuit? What is the total X count within the circuit? Could you provide the number of Xs in the circuit? How many components are there in the circuit that function as Xs? What is the X tally in the circuit? Could you indicate the quantity of Xs present in the circuit? How many X devices are there in the circuit? What is the total X count in the given circuit? Do you know how many Xs are present in the circuit? Could you specify the quantity of Xs in the circuit? Could you specify the quantity of Xs in the circuit? Could you provide the count of Xs included in the circuit? What is the tally of components offering X in the circuit?
Spatial Counting	How many Y are connected directly to the left of X? How many Y are connected directly to the right of X? How many Y are connected directly to the top of X? How many Y are connected directly to the bottom of X? How many gates are providing an input to X? How many gates are connected to the right of X? How many Y gates are connected to the right of X? How many Y gates are connected to the left of X?
Value Based	What are the current reading displayed by the XX? Please provide the values displayed on the XX. What does the XX show in terms of reading? What numerical value is being shown on the XX? What reading does the XX display? What are the value depicted on the XX? Can you provide the current measurement given by the XX? What are the current value indicated on the XX? What does the XX read at the moment? What are the present reading on the XX? Could you share the current reading that the XX shows?

 Table 2. Question Templates for various question types

(continued)

 Table 2. (continued)

Question Type	Question Templates
Junction based	Does a X exist between junction Y and junction Z? Is there a X present from junction Y to junction Z? Does a X occupy the space between junction Y and junction Z? Is there a X connecting junction Y to junction Z? Can a X be found between junction Y and junction Z? Does junction Y have a X leading to junction Z? Is there a X in the path from junction Y to junction Z? Can we observe a X between junction Y and junction Z? Does the circuit between junction Y and junction Z? Does the circuit between junction Y and junction Z? Does there impedance in the connection between junction Y and junction Z? Can you confirm the presence of a X between junction Y and junction Z? Does the circuit at junction Y and junction Z? Is there any resistance between junction Y and junction Z? Is a X located along the path from junction Y to junction Z? Can you verify if there is a X between junction Y and junction Z? Is there a X linking junction Y and junction Z? Is there a X linking junction Y to junction Z? Is there a X linking junction Y to junction Z? Is there a X bridging the gap between junction Y and junction Z? Does junction Y to junction Z through a X? Is there any resistance encountered from junction Y to junction Z? Is a X placed in the line connecting junction Y and junction Z?
Position based	Which circuit symbol is on the far X? Identify the circuit symbol that is at the extreme X. What is the circuit symbol located on the Xmost side? Tell me the circuit symbol positioned at the Xmost end. Point out the circuit symbol that is furthest to the X. Which circuit symbol is on the very X-hand side? Please indicate the circuit symbol situated all the way to the X. What is the name of the circuit symbol at the Xmost position? Which circuit symbol is on the extreme X? Find the circuit symbol on the Xmost side. Locate the circuit symbol positioned at the very X. Can you tell me which circuit symbol is at the Xmost position? Which circuit symbol positioned at the extreme X end? Point me to the circuit symbol on the Xmost side. What is the circuit symbol is placed at the extreme X end? Point me to the circuit symbol on the Xmost side. What is the circuit symbol is placed at the extreme X end? Point me to the circuit symbol on the Xmost side. What is the circuit symbol is placed at the extreme X end? Point me to the circuit symbol on the Xmost side. What is the circuit symbol's name that appears on the Xmost? Show me the circuit symbol positioned to the far X. Among the circuit symbols which one is at the Xmost position?

manual annotations for this objective and well-defined labeling task. Answering these questions requires the model to have an understanding of the way components are connected to each other spatially, i.e., object detection and localization skills.

3. Value Based Questions. Given an image, in a value based question, we ask what is the value associated with a particular electrical component. We prompt ChatGPT with this prompt: "Paraphrase the following sentence in 20 ways. What is the reading on X?" This leads to 20 different paraphrases⁶ which are used as question templates to generate value based questions in CIRCUITVQA. Again, we instantiate these templates to generate questions. If there are multiple components of type X in the image, the system is expected to provide a list of all of their values as the answer. Answering such questions requires a model to possess the optical character recognition skills, object recognition skills, and also the capability to link text labels with components.

Image metadata does not contain values associated with components. But the values are mentioned in the image. To generate answers automatically we used Google Vision APIs to perform OCR. The value text label is then linked with the closest bounding box (from associated metadata), and hence to a relevant component. However, on manual inspection, we found that this led to poor results because (i) OCR quality is bad especially for hand-drawn images, and (ii) closest bounding box heuristic often fails. Hence, finally we resorted to manual answer labeling done by the first author.

4. Junction Based Questions. Given an image, in a junction based question, we would like to know whether a component exists between two junctions. Thus, these are binary questions. Datasets D2, D3 and D5 also have labeled bounding boxes for junctions. We prompt ChatGPT with this prompt: "Paraphrase the following text in 20 ways - Does a X exist between junction Y and junction Z?" The generated paraphrases are used as question templates to generate junction based questions. To instantiate these templates for a positive answer (i.e., answer="yes"), we need valid triples of component X, junction Y and junction Z. First, we randomly choose a junction Y. Next, based on its Euclidean distance with other junctions (computed using centers of their bounding boxes), we choose a junction Z which is closest to Y. Lastly for every component in the image, we find its distance to every junction, and choose a component X such that its sum of distances to junctions Y and Z is minimum compared to any other pair of junctions. Such a $\langle X, Y, Z \rangle$ triple helps generate a question with answer="yes". Next, we randomly sample a component X' from the image metadata, of a different type from X. Such a $\langle X', Y, Z \rangle$ triple helps generate a question with answer=""". Answering junction-based questions requires a model to possess object detection and localization, as well as spatial reasoning skills.

5. Position Based Questions. Given an image, in a position based question, we want to know the component at the left-most, right-most, top-most or bottom-most of the image. We prompt ChatGPT with this prompt: "Paraphrase

 $^{^{6}}$ On manual inspection, we removed a few templates which did not make sense.

the following in 20 ways - Which is the Xmost circuit symbol?" The resultant paraphrases are used as question templates to generate position based questions. For every image, we randomly sample a question template and replace the placeholder X with one of left, right, top or bottom to get an instantiated question. To get the answer, we utilize the bounding boxes of the components present in the image and find their minimum and maximum X and Y coordinates to decide the left-most, right-most, top-most or bottom-most components in the image. If there is no unique answer, we eliminate those questions. Answering such questions requires the model to possess object detection and localization skills.



Fig. 2. Frequency distribution of value-based questions across component names.



Fig. 3. Frequency distribution of count-based questions across number of components of a particular type in images in CIRCUITVQA. Left: Simple Counting, Right: Spatial Counting.

3.3 CIRCUITVQA Dataset Analysis

We split the images into 70%, 20% and 10% split for training, testing and validation sets. Table 4 provides the count of questions by question type for train, test and validation splits. Figure 2 shows the frequency distribution of value-based questions across component names in CIRCUITVQA. Components like

Question Type	Answer Type
Simple Counting	Count (number)
Spatial Counting	Count (number)
Junction based	Binary
Position based	Component Name
Value based	List of values with units

 Table 3. Answer types for every question type

Table 4. # question-answer pairs per question type

Question Type	Training	Test	Val	Total
Simple Counting	16249	4776	2332	23357
Spatial Counting	624	170	236	1030
Junction based	45948	13998	6640	66586
Position based	14904	4232	2151	21287
Value based	2823	137	362	3322
Total	80548	23313	11721	115582

"resistor", "gnd", "and gate", "nand gate", and "inductor" are the most frequent in value-based questions. Figure 3 shows the frequency distribution of count-based questions across number of components of a particular type in images in CIR-CUITVQA. The left plot is for simple counting questions while the right plot is for spatial counting. For simple counting questions, although several questions have count as $1, \sim 52\%$ questions have the answer count greater than 1. similarly, there is good variety in answers for spatial counting questions.

4 Methods for CIRCUITVQA

To solve the CIRCUITVQA problem, we leverage two kinds of multimodal large language models as discussed in the following and detailed in Table 5.

4.1 Generative Models

BLIP [23]. BLIP (Bootstrapping language-image pre-training) is a multimodal mixture of encoder-decoder which operates with unimodal encoders for image and text. The model comprises of an image-grounded text encoder, image-grounded text decoder based on BERT [9] and image encoder based on vision transformers (ViT) [10]. In Visual Question Answering setting, we follow the same methodology described in the paper for finetuning on train part of CIR-CUITVQA. Specifically, we provide a circuit image-question pair to the image and text encoders separately, then compute the multimodal embeddings and provide it to the final text decoder (along with shifted outputs). The VQA model

is fine tuned with Language modelling loss which utilizes ground-truth answer as the target labels.

GIT [40]. GIT (Generative Image-to-text Transformer) is a decoder-only transformer that leverages CLIP [33] as a vision backbone. We fine-tune GIT on our task. Specifically, we concatenate the question and ground-truth answer as a special caption and apply the language modelling loss to the answer and [EOS] token.

Pix2Struct [22]. Pix2Struct is a generative model for visual understanding that converts image to text. It has an image encoder and a text decoder. We provide the images and the questions to the input image encoder. The model renders the questions on top of the image. It scales images up or down to extract maximal patches that fit within the sequence length parameter. For fair comparison with other models, we evaluate its performance using input images that have been resized to 384×384 dimensions.

 Table 5. Details of generative and instruction-tuned models that we experiment with for the CIRCUITVQA task.

	Architecture Initialization			Pretraining	Size
Model	Text Encoder	Image Encoder	Text Decoder	Objective	(Parameters)
BLIP-Base	BERT-base	ViT-B/16	BERT-base	Image captioning, image-text contrastive (ITC), image-text matching (ITM)	129M
GIT-Base	No text encoder	ViT-B/16	BERT	Image captioning	129M
Pix2Struct-Base	No text encoder	ViT	BERT	Screenshot parsing	282M
LLaVA	No text encoder	ViT-L/14	LLaMA	Auto-regressive loss for Conversation, detailed description, complex reasoning	6.76B
InstructBLIP	No text encoder	ViT + QFormer	Vicuna-7B	Language modeling on 26 datasets	7.91B

4.2 Instruction Tuned Models

LLaVA [26]. LLaVA (Large Language and Vision Assistant) is an end-to-end trained large multimodal model trained to follow human intent to complete visual tasks. It connects a vision encoder (ViT) with massive LLM based on LLaMA [39] or Vicuna [6]. At finetune time, the visual encoder weights are frozen but both the pre-trained weights of the projection layer and LLM are updated.

InstructBLIP [7]. InstructBLIP is the instruction fine-tuned version of BLIP2. Just like BLIP2, it is pretrained in two stages. Instruction-aware Q-former module takes in the instruction text tokens as additional input. While performing instruction tuning, the image encoder and the LLM are frozen. Tuning is done using 26 publicly available diverse datasets.

GPT4V [31]. GPT-4 with vision (GPT-4V) enables users to instruct GPT-4 to analyze image inputs provided by the user. GPT-4V shows unprecedented ability

in understanding and processing an arbitrary mix of input images, sub-images, texts, scene texts, and visual pointers. Its capabilities include open-world visual understanding, visual description, multimodal knowledge, commonsense, scene text understanding, document reasoning, coding, temporal reasoning, abstract reasoning, emotion understanding, and many more [41].

4.3 Language Modeling Loss

We used cross entropy loss for finetuning. The loss is computed only over answer tokens. We also experiment with a class-weighted version of cross entropy loss. Here each class is represented as a group of tokens. Specifically, we calculate weight of a class at the token level as a inverse count of that token in the dataset.

4.4 Input Representations

In the base variant of our experiments, we pass the original image and text as input to various models discussed in the previous subsection. Further, we also experiment with passing other forms of input representations as input. These include OCR text, bounding box information from object detection, and visual description of components. Table 6 shows how such information is included as part of the input prompt to instruction-tuned models.

	Variant	Prompt
LLaVA	Base	Given the image, answer the following question: Q
	Desc	The question is about the circuit component (Component-Name). Its definition is as follows: (ChatGPT-description). Now, given the image, answer the following question: Q
	OCR	Here is the OCR information (OCR). You can use it to answer the following question. Now, given the image, answer the following question: Q
	BBox	Here are the bounding box coordinates of each component in the given image in the format of a pair of component name and coordinates. (Bounding-box-coordinates). Now, given the image, answer the following question: Q
	BBox +Segments	Here are the bounding box coordinates and segment of each component in the given image in the format of a triple of component name, coordinates, and segment name. (Bounding-box-segments). Now, given the image, answer the following question: Q
Instruct BLIP	Base	Q
	Desc	The question is about the circuit component (Component-Name). Its definition is as follows: (ChatGPT-description). Q
	OCR	Here is the OCR information (OCR) . Q
	BBox	Here are the bounding box coordinates of each component in the given image in the format of a pair of component name and coordinates. (Bounding-box-coordinates). Q
	BBox +Segments	Here are the bounding box coordinates and segment of each component in the given image in the format of a triple of component name, coordinates, and segment name. (Bounding-box-segments). Q
GPT4V	Base	Q
	Desc	Use the following description of the electrical component to answer the question: (ChatGPT-description). Now, respond to this question: Q
	OCR	Use the following OCR output to answer the question: $\langle OCR \rangle$. Now, respond to this question: Q
	BBox	Use the following bounding box output comprising of the components and their coordinates in the image: (Bounding-box-coordinates). Now, respond to this question: Q
	BBox +Segments	Use the following bounding box output comprising of the components and their corresponding positions in the image : (Bounding-box-segments). Now, respond to this question: Q

 Table 6. Input Prompt Templates for Instruction-based Models

OCR Text: Since some questions relate to actual text labels in the image, the models may benefit from outputs of an external OCR module. Therefore, we

conduct an experiment to provide the OCR extracted tokens as an input to the vision-language models. We utilize Google Vision API⁷ to collect the OCR outputs from the circuit image. Then, we append the OCR output as a prefix to the question separated by an [OCR] token for fine-tuning the generative models. We also experiment with passing filtered OCR text as input by keeping only the numbers and units typically expected by electrical measurements. In this setting, we retain any OCR output tokens that contain any of the symbols in [' Ω ', 'H', 'A', 'F', 'V', 'W', 'k', 'K', '.', ' κ ', 'M'] or a combination of these symbols with a digit or only digits.

Bounding Box Information: Bounding boxes identified using object detection methods help in attending to the relevant local parts of the image [25], and their usage has been shown to improve the performance in transformers [1]. Therefore to increase the spatial awareness of the components in images, we utilize an object detection module.

Metadata in CIRCUITVQA contains human annotated bounding boxes for various components in electrical circuit images. We use this dataset to (a) finetune the YOLOv8 [34] object detection model, and (b) use them in our fine tuning experiments of vision-language models.

We fine-tune the pretrained YOLOv8 model for 300 epochs on image size of 384. The batch size was kept at 16 and patience (early stopping criterion) was set at 50 epochs. The learning rate was determined automatically and set at 0.01 and SGD optimizer was used with momentum 0.9. On validation set, YOLOv8 finetuned Objection detection leads to precision of 78.1, recall of 63.9, mAP50 of 69.8 and mAP(50-95) of 51.3. Figure 4 shows a few object detection examples. The figure shows that our fine-tuned model is able to identify electrical components from circuit images effectively. We fine-tuned YOLOv8 for these classes: __background__, acv, ammeter, and, antenna, arr, block, capacitor, capacitor-unpolarized, capacitor.adjustable, capacitor-polarized, crossover, crystal, current-source, diac, diode, diode.light emitting, diode.thyrector, fuse, gnd, diode.zener, inductor, inductor.coupled, inductor.ferrite, inductor2, integrated circuit, integrated circuit.ne555, integrated circuit.voltage regulator, junction, lamp, magnetic, mechanical, microphone, motor, multi-cell-battery, nand, nor, not, operational amplifier, operational amplifier.schmitt trigger, optical, optocoupler, or, probe, probe.current, relay, resistor, probe.voltage, resistor.adjustable, resistor.photo, single-cell-battery, socket, speaker, switch, terminal, text, thyristor, transformer, transistor, transistor-photo, transistor.bjt, transistor.fet, triac, unknown, varistor, voltage-ac, voltage-dc, voltage-dc ac, voltage.battery, voltmeter, vss, xnor, and xor.

We experiment with two ways of providing the bounding box information as input to our vision-language models. In the BBox method, for each detected component, along with the component name, we pass bounding boxes in the $\langle x, y, w, h \rangle$ format where x, y are box center, w and h indicate width and height. The model may not be able to process the numerical information; hence we

⁷ https://cloud.google.com/vision.


Fig. 4. Examples of Object detection results using our finetuned YOLOv8.

abstract out this information by assigning each bounding box to one of the 9 segments depending on its position in the image "upper left", "upper middle", "upper right", "left", "middle", "right", "lower left", "lower middle", "lower right". Based on this segment assignment, in the BBox+Segment method, for each detected component, along with the component name, we pass bounding boxes in the $\langle x, y, w, h \rangle$ format as well as the segment name.

Visual Description of Components: For every electrical component in our CIRCUITVQA dataset, we first obtain a short description using ChatGPT [30] with the following prompt "Describe the electrical component $\langle \text{component} \rangle$ in 50 words". In the Desc method, we pass the component description of relevant circuit component as a prefix to the question with a special token [DESC] separator. For example, description for capacitor is "Capacitor: Symbolized by two parallel lines with a gap, it stores and releases electrical energy, acting as a temporary energy reservoir in a circuit."

5 Experiments and Results

5.1 Experimental Setup

Generative Models: For GIT, the learning rates are set to 1e-5 and 2e-5 for the image encoder and the text decoder respectively. Rest of the hyperparameters are set to default values. For BLIP and Pix2Struct, learning rate for the text decoder is set to 2e-5. For all models, we use cosine learning rate scheduler. We use AdamW optimizer with a weight decay of 0.05. For Pix2Struct, we use default patch size of 16×16 and sequence length of 4096. For the text decoder of all models, we used hidden size of 768.

All models are trained for 10 epochs. The batch size is set to 4 for all the experiments. For fine-tuning and inference, we used a machine with 8 NVIDIA 32GB V100s. The computation time was 20-40 h for various models. All models are trained to optimize for cross-entropy loss (with label smoothing of 0.1) except for Pix2Struct where we found weighted cross-entropy loss to perform better. Also, we perform all experiments using an input image size of 384×384 .

Instruction-Based Models: To utilize InstructBLIP in zero shot settings, we set the number of beams to 5 and min length of the sequence to be generated to

1 and max length to 256. We keep the probability value for top p sampling to 0.9. Also, we set the temperature to 1 after trying out few different temperature settings. For LLaVA, we set the number of beams for beam search to 1. And provide the max length of the tokens to be generated to 512. Finally, we set the temperature to 0. For GPT4V, we set temperature to 0.15, max tokens as 350, top-p as 0.8, frequency penalty as 1 and presence penalty as 1.

5.2 Metrics

For every model, we measure exact-match accuracy and hallucination score as the two metrics. A good model should not just generate accurate answers but also not hallucinate. Hallucinations for visual question answering deserve specific definitions. Hence, we discuss these definitions and propose a new metric HVQA in the following.

Hallucination in VQA systems could be in terms of predictions of non-existing in-domain objects, over-counting of existing objects, or predictions with out-of-domain objects. Accordingly, we define Hallucination Score for Visual Question Answering (HVQA) as average of three scores: (a) $HVQA_{count}$ (captures over-counting of existing objects), (b) $HVQA_{in-domain}$ (captures predictions of non-existing in-domain objects), and (c) $HVQA_{out-domain}$ (captures predictions with out-of-domain objects). Each of these are fractions with total number of predicted objects as the denominator. Since we perform object detection on the input image as part of generating the answer, we can directly use the object detection outputs to compute the above scores. $HVQA_{count}$ is applicable for simple counting, spatial counting and value based questions. $HVQA_{in-domain}$ and $HVQA_{out-domain}$ are both applicable for position-based questions. HVQA is a general metric applicable to any VQA task.

5.3 Results

Main Results: Table 7 shows our main results where we compare various methods under different input representations on the CIRCUITVQA test set with respect to Accuracy (Acc) and hallucination score (HVQA).

BLIP provides the best accuracy while LLaVa and GPT4V provide the lowest hallucination scores. Our best model is a fine-tuned BLIP model with an accuracy of 91.7, when it is paired with prompts of visual description of the component (we call it as BLIP-Desc). It also maintains one of the lowest HVQA scores among all models. When an external OCR output is provided to these models, we observe a drop in their respective performances. This could be due to a lot of noise in the output of the OCR module. However, after postprocessing of the OCR output, there was a significant improvement in GIT (accuracy 71.2 vs 68.4) and InstructBLIP (accuracy 13.2 vs 12.5) when compared to the OCR output used directly. Also, when bounding boxes with their coordinates for each component were provided, we observe a drop in performance of the finetuned smaller models. However, the larger LLAVA zero-shot model can utilize that information and shows significant accuracy gains (42.9 vs 35.6 for the base model). Notably, the accuracy further increases to 44.6 with BBox+Seg.

	Model	Base		OCR		OCR-Post		Desc		BBox		BBox+Seg	
		Acc	HVQA	Acc	HVQA	Acc	HVQA	Acc	HVQA	Acc	HVQA	Acc	HVQA
Fine- Tuned	BLIP	84.4	5.9	81.8	5.8	80.8	5.9	91.7	5.5	75.6	6.4	74.0	6.2
	GIT	72.5	6.3	68.4	6.2	71.2	5.9	55.3	6.7	40.2	7.6	48.7	6.1
	Pix2Struct	71.2	6.3	69.1	6.2	41.9	6.7	70.3	6.1	44.2	4.1	36.6	4.5
Zero-Shot	LLaVA	35.6	3.8	35.4	5.2	35.4	5.4	35.6	3.8	42.9	2.8	44.6	3.8
	InstructBLIP	6.8	19.2	12.5	14.3	13.2	13.7	35.0	5.5	6.8	19.2	6.8	19.2
	GPT4V	34.5	4.8	41.2	2.2	34.1	4.0	33.7	5.9	32.1	3.0	32.3	3.7

Table 7. Main Results on CIRCUITVQA test set. Acc (\uparrow) , HVQA (\downarrow) .

Table 8. Accuracy results per question type for the Desc variants of the models onCIRCUITVQA test set.

Model	Simple Counting	Spatial Counting	Junction based	Position based	Value based
BLIP	83.5	57.6	97.9	84.1	18.2
GIT	46.5	34.7	66.9	29.4	0.7
Pix2Struct	48.2	44.7	90.1	32.6	11.7
LLaVA	18.8	0.6	7.8	50.6	0.7
InstructBLIP	35.8	14.1	0.0	0.9	0.0
GPT4V	12.5	10.0	50.6	6.4	0.7

Table 9. Hallucination scores. A = count, B = in-domain, C = out-domain.

Model	А	В	С
BLIP	0.9	15.6	0
GIT	0.6	19.6	0
Pix2Struct	0.3	18.2	0
LLaVA	5.8	0	5.5
InstructBLIP	16.6	0	0
GPT4V	0.07	12.3	5.4

Results per Question Type: For our best model (model that uses description), we analyze the accuracy results per question type in Table 8. Table 9 shows hallucination scores across various models on the CIRCUITVQA test set, where $A = HVQA_{count}$, $B = HVQA_{in-domain}$, $C = HVQA_{out-domain}$. We observe that BLIP-Desc outperforms all other models for each question type. It also hallucinates less on in-domain objects compared to its fine-tuned counterparts GIT



Fig. 5. Examples of images from CIRCUITVQA dataset

and Pix2Struct. Also, fine tuning broadly ensures that the models (BLIP, GIT and Pix2Struct) do not hallucinate out-of-domain objects. On the other hand, instruction-tuned models like LLaVA and GPT4V have a significantly higher HVQA_{out-domain}. LLaVA predicts out-of-domain objects like 'circle', 'square', 'A', 'B', 'D', 'F', 'triangle', 'carlin', 'nano', 'peizo-keeper', 'trigger', 'Snake Snake Detector'. InstructBLIP is very cautious and has neither out-of-domain nor indomain hallucinations, possibly because of its failure to understand positionbased or value-based questions.

For counting of objects, Pix2Struct hallucinates the least (HVQA_{count} of 0.3), while our best model BLIP-Desc hallucinates a little more, but is twice accurate compared to Pix2Struct. Among all visual description based models, InstructBLIP hallucinates the most on counting (HVQA_{count} of 16.6).

Case Studies: Table 10 show examples of questions and predicted answers associated with a few circuit images from the test set as shown in Fig. 5. For value based question, we can see that the model is able to accurately extract various values associated with the respective component. For junction question types, the model can correctly answer the respective question about two junctions even when there are more than 40 junctions in the image. We also observe that the model can correctly answer spatial counting questions by understanding the id associated with each component and then reasoning over the image to answer the question. Similarly the model can easily count values between 1 to 5, as shown in the examples.

Image	Question Type	Question	Prediction
(A)	Simple Counting	Could you indicate the quantity of resistors present in the circuit?	5
	Position-based	What is the circuit symbol's name that appears on the rightmost?	resistor
	Value-based	What is the value depicted on the ammeter?	['3a']
	Position-based	Which circuit symbol is on the extreme bottom?	resistor
	Value-based	What do the resistors read at the moment?	['10hm', '10hm', '10hm', '10hm', '30hm']
	Simple Counting	How many components are there in the circuit that function as ammeters?	1
(B)	Spatial Counting	How many gates are providing an input to Gate4	2
	Spatial Counting	How many or gates are connected to the right of Gate3 ?	1
	Simple Counting	Could you indicate the quantity of ors present in the circuit?	2
	Spatial Counting	How many and gates are connected to the left of Gate1 ?	1
	Position-based	Can you tell me which circuit symbol is at the leftmost position?	not
	Position-based	Which circuit symbol is placed at the extreme bottom end?	or
(C)	Junction-based	Is a integrated_circuit placed in the line connecting junction 4 and junction 12 ?	yes
	Simple Counting	How numerous are the capacitor-unpolarized in the circuit?	1
	Simple Counting	What number of transformer are included in the given circuit	1
	Simple Counting	How many components are there in the circuit that function as integrated _circuits?	2
	Junction-based	Is there a transformer interposed between junction 37 and junction 36 ?	yes

Table 10. Examples of Predictions from our best model for questions related to images shown in Fig. 5.

Table 11. Examples of error cases from our best model for questions related to images shown in Fig. 5.

Image	Question Type	Question	Answer	Prediction	Error Category
(C)	Value-based	What does the resistor.adjustable read at the moment?	['220kohm']	['100kohm']	Wrong values
	Junction-based	Is there a capacitor between junction 18 and junction 16?	no	yes	-
(D)	Spatial Counting	How many voltmeter are connected directly to the right of C4?	0	1	Over-counting
	Position-based	Which circuit symbol is placed at the extreme left end?	voltage.battery	resistor	Near miss
(E)	Simple Counting	Could you provide the count of resistors included in the circuit?	4	2	Under-counting

Error Analysis: We manually analyzed 100 test cases where our system leads to an error, 20 for each question type. Among the 20 value-based questions, 4 errors can be attributed to incorrect units, 5 were a result of both units and values being wrong, and the majority (11 errors), were due to incorrect values. For 20 junctionbased questions, 12 errors were for images with \geq 40 junctions and 8 for images with <40 junctions. Broadly, we observe that accuracy drops with increase in number of junctions in input image. For position-based questions, for 9 samples, the predicted component was physically the second closest to the correct answer component; remaining 11 predictions were far from the actual answer. In simple counting questions, we identified 11 over-counting errors, all within a range of 1 to 5, while there were 9 instances of under-counting. Spatial counting questions had 4 cases of over-counting and 16 examples of under-counting. Table 11 shows a few error examples from our best model for questions related to a few circuit images from the test set as shown in Fig. 5.

6 Conclusion

In this paper, we proposed the problem of visual question answering for electrical circuit images. We curated a dataset, CIRCUITVQA, for the task with five question types. We hope that this dataset will help the VQA community to focus on the critical problem of VQA for circuit images. We performed extensive evaluation of several state-of-the-art vision language models. We also experimented with different forms of input representation including OCR text, bounding boxes based on object detection and detailed description of relevant circuit components. Our experiments reveal that the BLIP model with description of components provide the highest VQA accuracy across most question types, and the lowest hallucination score.

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