

Un-mixing Test-time Adaptation under Heterogeneous Data Streams

Zixian Su*, Jingwei Guo*, Xi Yang†, Qiufeng Wang, Kaizhu Huang†

Abstract—Deploying deep models in real-world scenarios remains challenging due to significant performance drops under distribution shifts between training and deployment environments. Test-Time Adaptation (TTA) has recently emerged as a promising solution, enabling on-the-fly model adaptation without access to source data. However, its effectiveness degrades significantly in the presence of complex, mixed distribution shifts – common in practical settings – where multiple latent domains coexist. Adapting under such intrinsic heterogeneity, especially in unlabeled and online conditions, remains an open and underexplored challenge. In this paper, we study TTA under mixed distribution shifts and move beyond conventional homogeneous adaptation paradigms. By revisiting TTA from a frequency-domain perspective, we observe that distribution heterogeneity often manifests in Fourier space – for instance, high-frequency components tend to carry domain-specific variations. This motivates us to perform domain-aware separation using high-frequency texture cues, making diverse shift patterns more tractable. To this end, we propose FreDA, a novel Frequency-based Decentralized Adaptation framework that decomposes globally heterogeneous data into locally homogeneous components in the frequency domain. It further employs decentralized learning and augmentation strategies to robustly adapt under complex, evolving shifts. Extensive experiments across various environments (corrupted, natural, and medical) demonstrate the superiority of our proposed framework over the state-of-the-arts.

Index Terms—Test-time Adaptation, Transfer Learning

I. INTRODUCTION

DEEP learning models often suffer significant performance degradation when deployed in environments where the data distribution differs from that of the training set – a challenge known as domain shift [1], [2]. Recently, Test-Time Adaptation (TTA) [3]–[10] has emerged as a promising solution by refining model parameters to better align with the encountered data at inference time. It leverages the incoming data stream for real-time adjustments without the need for retraining on a labeled dataset, enabling swift model adaptation to unpredictable data characteristics during deployment.

Despite their success, current TTA models often are limited to ideal testing conditions, typically involving homogeneous test samples with similar types of distribution shifts. In reality, distribution shifts are *mixed, overlapping, and even conflicting* [11]–[15]. For instance, photo management software handles diverse corruptions, such as noise, blur, compression;

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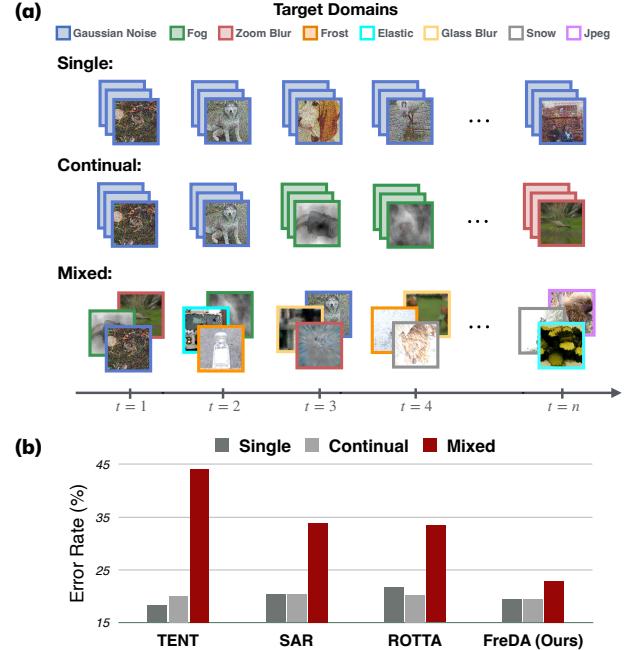


Fig. 1: (a) Illustration of three TTA scenarios with single, continual, and mixed domain shifts. (b) Classification error rate on CIFAR-10-C across different TTA settings: performance of conventional models drops sharply under mixed domain shifts.

medical imaging platforms manage inconsistencies from varied acquisition methods; and autonomous driving systems face fluctuating conditions like lighting, weather, and road types.

Such complexity poses serious challenges for existing methods. While recent efforts have extended TTA to continually changing environments [5], they typically assume a uniform target domain at each time step, as illustrated in the continual setting in Figure 1 (a). Some methods periodically reset the model to its source pre-trained state [7], [9] to mitigate the cumulative effect of sequential distribution shifts on adaptation. Others down-weight outlier target samples [6], [10] to suppress the impact of abrupt or rare shifts. Although these strategies aim to improve robustness under dynamic conditions, they fall short when facing the entangled / mixed real-world distributions (see Figure 1 (b)). Their shortcomings stem from a shared limitation: they lack the capacity to disentangle or localize domain variations. As a result, they often overfit to dominant patterns or suffer from catastrophic forgetting, failing to be deployed in realistic, heterogeneous data streams.

Capturing such intrinsic heterogeneity under unlabeled, online conditions is particularly challenging, and effective adaptation in this setting remains an open problem. To tackle

this, we propose shifting from coarse, homogeneous adaptation to a fine-grained strategy that explicitly disentangles heterogeneous shifts. Our approach leverages the frequency information to disentangle domain variations, which naturally separates data features across frequency bands: high-frequency components capture fine details like edges and textures, while low-frequency components represent global structures such as shape and illumination (Section IV). By decomposing inputs accordingly, we better characterize distributional variations and identify diverse shifts. Moreover, the Fourier transform operates directly on raw pixel-level inputs, which shields the method from performance degradation under large domain gaps, and also enhancing adaptability and deployment efficiency in real-world applications.

Building upon this insight, we introduce a framework, termed Frequency-based Decentralized Adaptation (FreDA). FreDA begins by partitioning incoming data in the Fourier domain, where high-frequency components are used to group samples with similar shift characteristics. This transforms globally heterogeneous inputs into locally homogeneous subsets before any adaptation takes place. Based on these partitions, FreDA deploys multiple local models, each assigned to a specific subset. These models adapt independently to their respective data streams while periodically synchronizing through parameter exchange. The aggregated parameters form a shared base model, which is then used to reinitialize the local models, ensuring continual and coordinated adaptation across domains. This approach not only alleviates clustering errors via collaborative knowledge sharing but also allows local models to capture diverse distribution shifts. To further enhance robustness, we introduce a novel Fourier-based augmentation scheme that improves sample quality and strengthens adaptation to shift-specific characteristics.

To summarize, the main contributions of this work are three-fold:

- We identify a key limitation of most existing TTA methods – their neglect of real-world data heterogeneity – which leads to suboptimal performance when confronted with mixed and diverse distribution shifts.
- We propose FreDA, a frequency-based decentralized adaptation framework that leverages spectral decomposition and localized adaptation to effectively address heterogeneous distribution shifts at test-time.
- We validate FreDA through extensive experiments on corrupted, natural, and medical benchmarks, demonstrating consistent improvements over state-of-the-art methods across various TTA scenarios.

II. CONNECTIONS TO PREVIOUS STUDIES

A. Transfer Learning, Domain Adaptation, Test-time Training, and Test-time Adaptation

Deep neural networks often experience performance degradation when deployed in environments different from their training settings, due to distribution shifts between source (training) and target (testing) domains. This issue is extensively studied under the umbrella of transfer learning [16], [17], which aims to transfer knowledge across domains. A key

TABLE I: Comparison of different transfer learning settings.

Topic	Source Data	Target Labels	Online Adaptation	Model Agnostic
Supervised Domain Adaptation	✓	✓	✗	–
Unsupervised Domain Adaptation	✓	✗	✗	–
Source-free Domain Adaptation	✗	✗	✗	–
Test-time Training	✗	✗	✓	✗
Test-time Adaptation	✗	✗	✓	✓

Note: TTA is the most practical setting for real-world deep model deployment.

branch of transfer learning is domain adaptation (DA) [18]–[21], where models trained on a labeled source domain are adapted to a target domain. Depending on the availability of source and target data, DA can be classified into: Supervised DA using labeled data in both source and target domains during training; Unsupervised DA, which relies on labeled source data and unlabeled target data; and Source-free DA, that adapts to the target domain without access to source data, typically due to privacy or transmission constraints.

While traditional DA methods assume access to the whole target distribution during adaptation, such access is not always possible in real-world deployment. To address this, two prominent test-time adaptation paradigms have been proposed: Test-time Training (TTT) [22]–[24] introduces an auxiliary task – often self-supervised – during pre-training. At test time, the model optimizes this auxiliary objective to adapt to the target distribution accordingly; Test-time Adaptation (TTA) [3], [7], [8] poses a more demanding and practical scenario, where the model must adapt on-the-fly to the test stream without any prior modification during training. This setting emphasizes the need for rapid, real-time model updates to effectively capture the continuously incoming data. A brief comparison of these adaptation settings is summarized in TABLE I. Our work falls under the TTA setting, where we propose a generalizable and robust adaptation framework that maintains strong performance across diverse distribution shifts and corruptions – without relying on source data or altering the pre-training process.

B. Non-i.i.d. Test-time Adaptation

While conventional TTA methods primarily focus on single domain shifts under the idealized independent and identically distributed (i.i.d.) assumption, real-world deployment often violates these conditions due to inherent data heterogeneity. This discrepancy has motivated recent efforts to extend TTA to more realistic settings, with two primary challenges:

1) *Mixed Domains*: Modern TTA methods designed for continual domain shifts [5], [6], [9], [10], [25] have pushed beyond the single adaptation setting. However, these approaches assume that the target domain is uniform at each time point – a special case of mixed domains where the change occurs across distinct time periods. In reality, data streams often involve genuinely mixed domains with heterogeneous distributions present simultaneously. Existing approaches primarily focus on stabilizing model updates through periodic parameter

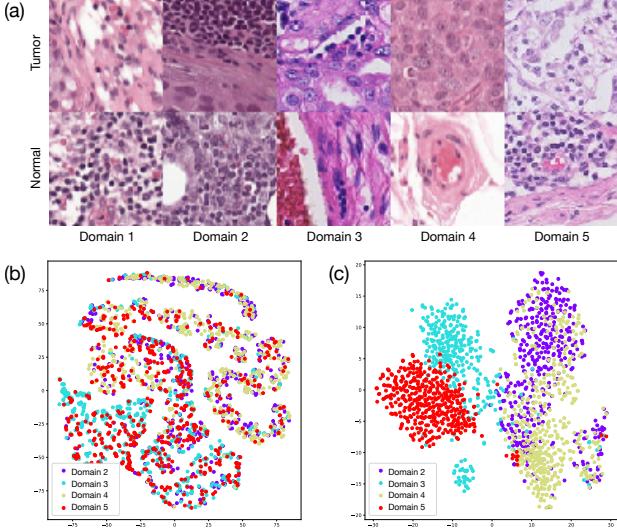


Fig. 2: (a) Visual heterogeneity across five healthcare centers in Camelyon17 dataset: Example patches showcase diverse domain shifts that characterize mixed target domains. (b) Conventional sample latent features from pretrained models fail to separate different target subdomains, showing significant overlap. (c) High-frequency information enable distinct separation of target subdomains, contrasting with the conventional ones.

resets [7], [9] or importance weighting of target samples [6], [9], [10], but these techniques are ill-equipped to disentangle and manage the complex interleaving of domain shifts, leading to degraded performance. In contrast, our work adopts a data-centric perspective by explicitly addressing mixed domains via frequency-space decomposition, which facilitates domain separation and proactive distribution alignment prior to model adaptation. While recent efforts like [7] acknowledge mixed distribution challenges under the broader “Dynamic Wild World” paradigm, their unified treatment of diverse real-world factors does not systematically address data heterogeneity. Conversely, our study directly targets the core issue of mixed domains in TTA, introducing tailored strategies for disentangling and adapting to co-occurring heterogeneous distributions.

2) *Dependent Sampling*: The second challenge stems from class-level dependencies introduced by temporal data sampling. This topic has garnered significant research attention, with a growing body of work [25]–[30] actively addressing it through strategies such as pseudo-label-based rebalancing, or extended observation windows. These approaches aim to correct the temporal imbalance caused by skewed class distributions over time. However, unlike these studies that focus on class-level imbalance, our work diverges by targeting sample-level heterogeneity, which remains underexplored. Specifically, we address style diversity and distributional variation at test-time – a challenge that persists even under class-balanced conditions – to improve model robustness under mixed-domain scenarios.

Methods	C10	C100	IN
TBN (Centralized)	33.8	45.8	82.5
TBN (Decentralized)	28.5	43.2	77.6

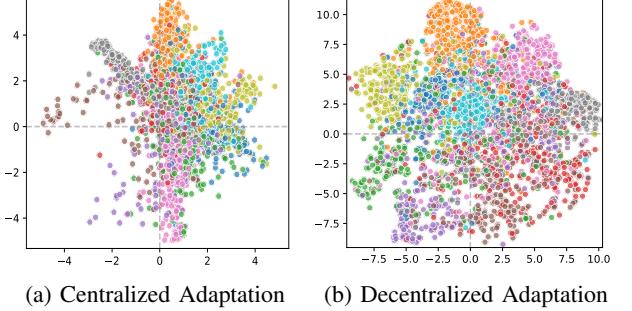


Fig. 3: Experimental Result (error rate) and t-SNE feature visualization using TBN [31] as the pseudo-labeling method under mixed distribution shifts, comparing centralized and decentralized adaptation. Different colors represent different classes. (a) Centralized Adaptation: Global BN parameters are applied to the entire batch. (b) Decentralized Adaptation: Localized BN parameters are tailored for clusters separated based on high-frequency features. C10, C100, IN respectively denotes CIFAR-10-C, CIFAR-100-C and ImageNet-C datasets.

III. PROBLEM DEFINITION OF MIXED DOMAIN SHIFTS

Test-time adaptation (TTA) aims to adjust a model $q_\theta(y|x)$, initially trained on a source dataset $\mathcal{D}_s = \{(x, y) \sim p_s(x, y)\}$, to a target domain $\mathcal{D}_t = \{(x, y) \sim p_t(x, y)\}$ on a data stream without accessing source data or target labels. TTA handles covariate shift by assuming $p_s(y|x) = p_t(y|x)$ while $p_s(x) \neq p_t(x)$. This challenge intensifies when \mathcal{D}_t contains multiple non-i.i.d sub-distributions $p_{t_i}(x)$, such as

$$p_t(x) \leftarrow \{p_{t_1}(x), p_{t_2}(x), \dots, p_{t_N}(x)\}.$$

Specifically, we have $\mathcal{D}_t = \mathcal{D}_{t_1} \cup \mathcal{D}_{t_2} \cup \dots \cup \mathcal{D}_{t_N}$ where $x \in \mathcal{D}_{t_1}$ satisfying $x \sim p_{t_1}(x)$. This scenario requires the model $q_\theta(y|x)$ to effectively handle the heterogeneous and evolving target distribution to maintain robust performance. TTA strategies must therefore refine the model to optimize its predictive accuracy across these diverse sub-domains, ensuring consistent and reliable performance amidst significant distributional variability.

IV. TTA UNDER MIXED DISTRIBUTION SHIFTS: A FOURIER PERSPECTIVE

While Test-Time Adaptation (TTA) methods excel under single-type distribution shifts, their performance degrades catastrophically when facing mixed distribution shifts – a ubiquitous challenge in real-world deployment. As demonstrated on CIFAR-10-C (Figure 1 (b)), leading methods like TENT [3], SAR [7] and RoTTA [25] suffer 16% average accuracy drop compared to single-shift scenarios. Conventional TTA paradigms attempt to mitigate this through passive robustness measures, such as entropy regularization [3], gradient masking [7], or model resetting [9]. These methods treat mixed

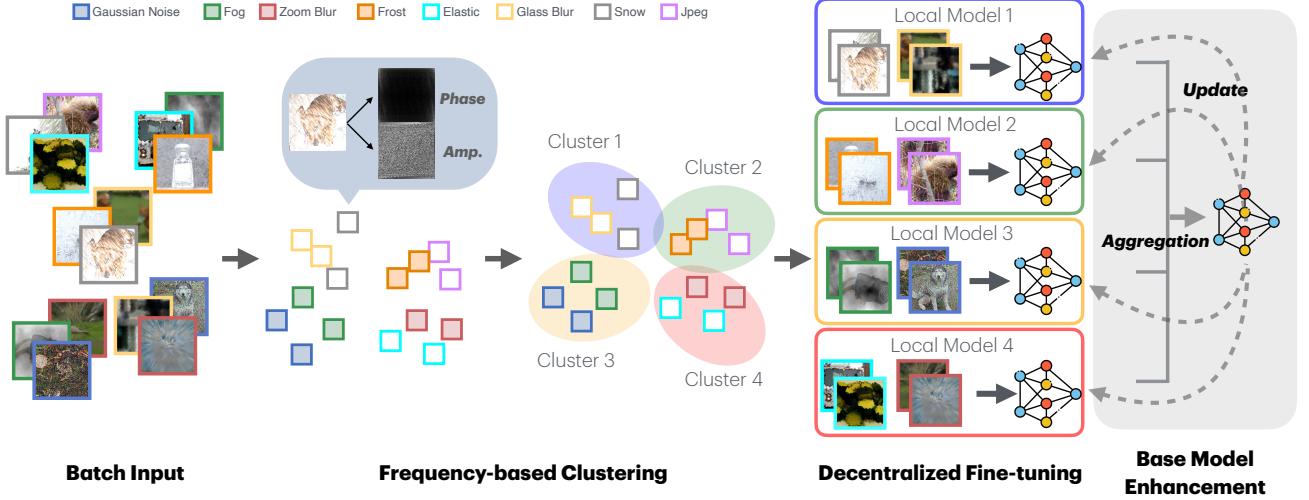


Fig. 4: Illustration of Frequency-based Decentralized Learning.

distribution shifts as irreducible noise, deferring corrective actions until model failures occur – a reactive paradigm that inevitably compromises adaptation efficacy.

To address this, we propose a paradigm shift to *proactive distribution management* – instead of demanding models to handle arbitrary mixtures, we first resolve the heterogeneity via frequency-space decomposition. Our key insight is that mixed distributions exhibit spectral signatures that can be disentangled before model adaptation begins. As evidenced in Figure 2 (c), high-frequency components naturally separate distinct shifts without relying on error-prone model features (see Figure 2 (b)). This theoretically grounded decomposition converts the ill-posed mixed-shift adaptation into multiple well-conditioned homogeneous-shift problems, effectively create an ideal adaptation environment from chaotic real-world streams. Crucially, this decomposition enables a *decentralized adaptation* where we can deploy domain-specific models tailored to high-frequency-separated clusters. Take TBN [31] as an example: while its centralized variant recalculates batch normalization (BN) statistics globally (treating heterogeneous data as a single distribution), our decentralized adaptation computes localized BN parameters within each spectrally disentangled cluster. As shown in Figu 3, this context-aware estimation preserves semantic discriminability: samples exhibit tighter within-class cohesion and sharper between-class boundaries compared to the blurred separability of *centralized adaptation*. Building on these insights, the following section details how leveraging the frequency domain enhances TTA methods for realistic scenarios with mixed distribution shifts.

V. FREQUENCY-BASED DECENTRALIZED ADAPTATION

The previous discussion highlights how heterogeneity within target distributions can hinder effective model adaptation. This naturally raises the question: *How can we manage such distributional heterogeneity to enable better adaptation?* As outlined earlier, distinguishing samples associated with different distribution shifts offers a promising pathway. Building on this insight, we address the TTA problem by

leveraging high-frequency components in the data and propose a novel framework termed Frequency-based Decentralized Adaptation (FreDA). It partitions target samples into multiple homogeneous subdomains in the Fourier space, facilitating more accurate model adaptation. This is further enhanced by a frequency-based augmentation strategy that enriches each subdomain with diverse samples, strengthening model robustness.

A. Frequency-based Decentralized Learning

Fourier transform offers an effective method to extract different frequency components from images, with high-frequency information particularly useful for capturing fine-grained details such as texture and noise. These details often highlight subtle variations among different distribution shifts. Based on this insight, we propose a module termed Frequency-based Decentralized Learning. It leverages frequency information extracted from the pixel space to systematically partition data into multiple homogeneous subsets, enabling multiple local models to specialize in capturing each distribution shift individually. Concurrently, our method enables collaborative learning by allowing periodic parameter sharing among these local models via an enhanced base model, thereby boosting overall adaptability to diverse distribution shifts.

1) *Frequency Feature Extraction:* We start by extracting frequency domain features from the input images. Let $\mathbf{X} \in \mathbb{R}^{n \times c \times h \times w}$ denote a batch of input images, where n is the batch size, c is the number of channels, h and w are the height and width. We first apply a Fourier transform \mathcal{F} to each image \mathbf{x}_i . This transform converts the image from the spatial domain to the frequency domain, producing a complex-valued representation $\mathcal{F}(\mathbf{x}_i) \in \mathbb{C}^{h \times w \times c}$ that contains both real $R(\mathbf{x}_i)$ and imaginary part $I(\mathbf{x}_i)$. Next, we compute the amplitude spectrum $A(\mathbf{x}_i)(u, v)$ using $A(x)(u, v) = \sqrt{R^2(x)(u, v) + I^2(x)(u, v)}$. It reveals the intensity of the frequency content, e.g., high-frequency amplitudes highlight edges and fine details while low-frequency amplitudes emphasize the overall structure and gradual changes. Then, we

filter out low-frequency elements using mask $M(u, v) = \mathbb{1}((u < \frac{h}{4} \vee u > \frac{3h}{4}) \vee (v < \frac{w}{4} \vee v > \frac{3w}{4}))$ to emphasize the high-frequency components $G(x)(u, v)$ that are more likely to indicate shifts in distribution:

$$G(x)(u, v) = A(x)(u, v) \cdot M(u, v). \quad (1)$$

2) *Frequency-Based Clustering*: We then employ a clustering algorithm (e.g., K-means) to partition the frequency features into K clusters, each corresponding to a different type of distribution shift. The process is formalized as:

$$\min_{\mathbf{C}, \mathbf{Z}} \sum_{i=1}^n \|\mathbf{A}_{hf,i} - \mathbf{C}_{\mathbf{Z}_i}\|_2^2, \quad (2)$$

where $\mathbf{A}_{hf,i} = \text{vec}(G(x_i))$, $\mathbf{C} \in \mathbb{C}^{K \times d}$, $\mathbf{Z} \in \{1, \dots, K\}^n$ denotes the 1D high-frequency component of the amplitude spectrum, the centroids of the clusters and the cluster assignments for each image. hf refers to high-frequency components, and $d = h \times w \times c$ is flattened dimension.

3) *Decentralized Fine-tuning*: Test-time fine-tuning is then decentralized across these clusters, allowing for specialized adaptation within each subgroup: For each cluster k , we adapt a specialized model $q_{\theta_k}(y|x)$ that is fine-tuned using only the data within that cluster:

$$\theta_k^* = \arg \min_{\theta_k} \mathbb{E}_{x \sim p_{t,k}} [\mathcal{L}(q_{\theta_k}(x))], \quad (3)$$

where $p_{t,k}$ represents the data distribution within cluster k , and \mathcal{L} is the loss function. The predictions of each iteration are collected and sorted after the local fine-tuning.

4) *Base Model Enhancement*: To integrate knowledge from all subnetworks and prevent degradation on specific subdomains, we periodically aggregation their parameters at intervals of time T :

$$\theta_{\text{base}} = \sum_{k=1}^K \left(\frac{|\mathcal{D}_k|}{\sum_{j=1}^K |\mathcal{D}_j|} \theta_k \right), \quad (4)$$

where $|\mathcal{D}_k|$ denotes sample number in cluster k . This aggregation step combines the parameter updates from each subnetwork proportionally to its cluster size. The updated parameters θ_{base} are then distributed back to each subnetwork, initializing them for the next batch of training: $\theta_k \leftarrow \theta_{\text{base}}$.

B. Frequency-based Augmentation

While decentralized learning effectively mitigates batch-level heterogeneity, it may falls short in accurately characterizing individual distribution shifts – largely due to limited batch size and noise introduced by coarse clustering. To improve target data quality, TTA methods commonly adopt data augmentation as a practical strategy to enhance model generalization. However, conventional augmentation techniques – typically borrowed from standard computer vision tasks such as rotation, cropping, and mixup – are primarily tailored for single-shift scenarios. They often fail to provide the targeted, distribution shifts-aware augmentation as required under mixed domain shifts. To this end, we propose a frequency-based augmentation strategy tailored for TTA in mixed-shift

Algorithm 1 Framework of Frequency-based Decentralized Learning and Augmentation

Require: Step t , Input batch $\mathbf{X} = \{x_1, x_2, \dots, x_n\} \in \mathbb{R}^{n \times h \times w \times c}$, Pretrained source model q_θ , Initialize Feature Repository and Local Sample Pool $\mathcal{R}, \mathcal{S}_k \leftarrow \emptyset$, CLUSTER_NUM K , KMEANS_SIZE N , COMM_INTERVAL f ;

Step 1: Extract Frequency Features

- 1: **for** $i = 1$ to n **do**
- 2: $\mathbf{A}_{hf,i} \leftarrow \text{vec}(G(\mathbf{x}_i))$ ▷ Extract high-freq components
- 3: **end for**

Step 2: Dynamic Clustering

- 4: $\mathcal{R} \leftarrow \mathcal{R} \cup \{\mathbf{A}_{hf,i}\}_{i=1}^n$ ▷ Frequency Information Repository
- 5: $\mathcal{R} \leftarrow \mathcal{R}[(|\mathcal{R}| - N + 1) :]$ ▷ Keep the last N entries for kmeans clustering
- 6: $(\mathbf{C}_t, \mathbf{Z}) \leftarrow \text{K-means}(\mathcal{R}, K, \mathbf{C}_{t-1})$ ▷ Obtain Cluster Labels
- 7: $\mathbf{Z} = \{Z_i\}_{i=1}^n$ (Eq. 2)

Step 3: Local Model Training

- 7: **for** cluster $k \in \{1, \dots, K\}$ **do**
- 8: $\mathcal{S}_k \leftarrow \mathcal{S}_k \cup \{x_i \mid Z_i = k\}$ ▷ Gather samples for cluster k
- 9: $\mathcal{S}_k \leftarrow \mathcal{S}_k[(|\mathcal{S}_k| - n + 1) :]$ ▷ Keep the last batch_size = n entries
- 10: $\mathcal{S}'_k \leftarrow \text{select_samples}(\mathcal{S}_k)$ ▷ Select samples (Eq. 5)
- 11: **for** each $x_i \in \mathcal{S}'_k$ **do**
- 12: $\tilde{x}_i \leftarrow \text{augment}(x_i)$ ▷ Augment data (Eq. 7)
- 13: Train($q_{\theta_k}, x_i, \tilde{x}_i$) ▷ Train local model (Eq. 3)
- 14: **end for**
- 15: **end for**

Step 4: Compile Predictions

- 16: $\mathbf{Y} \leftarrow \text{collect_sort}(\{q_{\theta_k}(\mathbf{X})\})$ ▷ Collect and sort predictions

Step 5: Base Model Enhancement

- 17: If $t \% f == 0$: ▷ Model Communication with interval f (Eq.4)
- 18: $\theta_{\text{global}} \leftarrow \sum_{k=1}^K w_k \theta_k$
- 19: $\theta_k \leftarrow \theta_{\text{global}}$

environments. Specifically, our method perturbs the amplitude components of target samples in the Fourier space, enabling targeted augmentation with respect to each sample’s underlying distribution. By manipulating the frequency-domain characteristics, our approach generates diverse yet distribution-consistent variations, thereby enriching subdomain representations and facilitating robust model adaptation

1) *Sample Selection Mechanism*: We first select the reliable samples in each local model leveraging a criterion derived from the weighted entropy framework used in ETA [6] based on two primary conditions:

$$\text{Cri} = \mathbb{1}[(H(\mathbf{y}_t) < H_0) \wedge (|\cos(\mathbf{y}_t, \bar{\mathbf{y}}_{t-1})| < \epsilon)]. \quad (5)$$

The entropy $H(\mathbf{y}_t)$ measures the uncertainty in the current predictions. The cosine similarity $\cos(\mathbf{y}_t, \bar{\mathbf{y}}_{t-1})$ denotes the deviation between the current sample’s class probabilities \mathbf{y}_t and the aggregated class probabilities $\bar{\mathbf{y}}_{t-1}$. ϵ is the threshold for cosine similarity, and H_0 is the fixed entropy threshold. This ensures that selected samples exhibit significant deviations from previous predictions in class distribution and lower prediction uncertainty.

2) *Frequency-Based Augmentation*: The augmentation process involves perturbing the amplitude spectrum. Let $A(x_i)$ represent the amplitude spectrum of a selected sample x_i . To generate a perturbed amplitude spectrum $\tilde{A}(x_i)$, we apply a random Gaussian perturbation:

$$\tilde{A}(x_i) = (1 + \alpha \cdot \Delta) \cdot A(x_i), \quad (6)$$

where $\Delta \sim \mathcal{N}(0, \sigma^2)$ is a perturbation matrix sampled from a Gaussian distribution, and α is a scaling factor. Then, the augmented sample \tilde{x}_i is reconstructed via the inverse Fourier

transform \mathcal{F}^{-1} to the perturbed amplitude spectrum, combined with the original phase spectrum $P(x_i)$:

$$\tilde{\mathbf{X}}_i = \mathcal{F}^{-1} \left(\tilde{A}(x_i), P(x_i) \right). \quad (7)$$

3) Loss Function: The training objective combines the entropy loss of the selected samples with a consistency loss from the augmented samples. The total loss is defined as:

$$\mathcal{L}_{\text{total}} = \frac{1}{n} \sum_{i=1}^n H(\mathbf{y}_i) + \lambda \cdot \frac{1}{n} \sum_{i=1}^n \mathcal{L}_{\text{con}}(\hat{\mathbf{y}}_i, \tilde{\mathbf{y}}_i), \quad (8)$$

where the entropy loss $H(\mathbf{y}_i)$ for the original sample x_i is given by $H(\mathbf{y}_i) = -\sum_{j=1}^C \mathbf{y}_{i,j} \log \mathbf{y}_{i,j}$ with \mathbf{y}_i being the predicted probability over the C classes, and the consistency loss $\mathcal{L}_{\text{con}}(\hat{\mathbf{y}}_i, \tilde{\mathbf{y}}_i) = -\sum_{j=1}^C \hat{\mathbf{y}}_{i,j} \log \tilde{\mathbf{y}}_{i,j}$ is defined as the cross-entropy between the prediction $\hat{\mathbf{y}}_i$ of the augmented sample \tilde{x}_i and the pseudo-label $\tilde{\mathbf{y}}_i$ from the original sample.

C. Overall Framework

We provide the overall pipeline algorithm of FreDA in Algorithm 1. During implementation, we leverage a memory bank strategy [4], [25], [32], [33] that is updated in real time. This design serves two purposes: **1)** to ensure accurate clustering—since an overly small batch could impede the effective separation of data – and **2)** to maintain the number of samples processed by the local model consistent with the original batch size, thereby preventing performance degradation due to a drastic reduction in batch size (e.g., reducing by a factor of the cluster number).

VI. THEORETICAL INSIGHTS

To facilitate theoretical understanding, we leverage the expansion-based analysis framework proposed in [34]. We first revisit the core concepts and then highlight the theoretical advantages of our method under mixed domain shifts.

A. Expansion Theory

Definition 1 ((a, c)-expansion). A class-conditional distribution P_i satisfies (a, c) -expansion if $\forall S \subseteq \mathcal{X}$ with $P_i(S) \leq a$:

$$\mathcal{P}_i(\mathcal{N}(S)) \geq \min(c\mathcal{P}_i(S), 1)$$

where $\mathcal{N}(S)$ refers to the neighborhood of S under data augmentations.

Definition 2 (Separation). P is (μ, r) -separated if:

$$\mathbb{E}_{x \sim \mathcal{P}} \left[\max_{x' \in B_r(x)} \mathbf{1}(G(x) \neq G(x')) \right] \leq \mu$$

where $B_r(x)$ is an ℓ_2 -ball of radius r .

The core theorem from [34] states:

Theorem 1 (Pseudo-label Denoising). Under (a, c) -expansion and (μ, r) -separation, any classifier G of:

$$\min_G \frac{c+1}{c-1} \mathcal{L}_{\text{pl}}(G) + \frac{2c}{c-1} \mathcal{R}_B(G)$$

achieves error:

$$\text{Err}(G) \leq \frac{2}{c-1} \text{Err}(G_{\text{pl}}) + \frac{2c}{c-1} \mu$$

where \mathcal{L}_{pl} is pseudo-label loss and \mathcal{R}_B is consistency regularizer.

B. Analysis of Our Method

Our method reduces both terms in Theorem 1 to achieve a tighter bound through two mechanisms:

1) Frequency-Coherent Partitioning.: Let $\{P_{t_k}\}_{k=1}^K$ be the K sub-domains identified by frequency-based partitioning. For measurable sets $\{S_k\}_{k=1}^K$ where $S_k \subseteq X_k$ (the support of P_{t_k}), define:

$$\text{Err}(G_{\text{pl}}^k) \leq \text{Err}(G_{\text{pl}}) - \Delta_k$$

where $\Delta_k = \mathcal{P}_{t_k}(E_c)$ and $E_c = \{x \in X_k \mid G_{\text{pl}}(x) \neq y(x)\}$. This contributes to the reduction of the first term in Theorem 1, which is further supported by experimental results in Figure 3.

2) Augmentation-Induced Expansion: Our frequency augmentation expands neighborhoods:

$$\widehat{\mathcal{N}}(S) = \mathcal{N}_{\text{base}}(S) \cup \left\{ x' : \inf_{x \in S} |A(\mathcal{F}(x')) - A(\mathcal{F}(x))|_2 \leq \epsilon \right\}$$

where $\mathcal{N}_{\text{base}}(S)$ is the base augmentation neighborhood. It is worth noting that, unlike most methods, we do not apply base augmentation. The term $\mathcal{N}_{\text{base}}(S)$ is retained in the formulation for theoretical completeness. Our choice of this representation is intended to highlight the expansion gain brought by incorporating frequency augmentation.

The expansion gain becomes:

$$\begin{aligned} \widehat{c} &= c_{\text{base}} + \gamma, \\ \gamma &= \inf_{S: P(S) \leq a} \frac{P(\widehat{\mathcal{N}}(S) \setminus \mathcal{N}_{\text{base}}(S))}{P(S)} \end{aligned}$$

Substituting into Theorem 1, the error becomes:

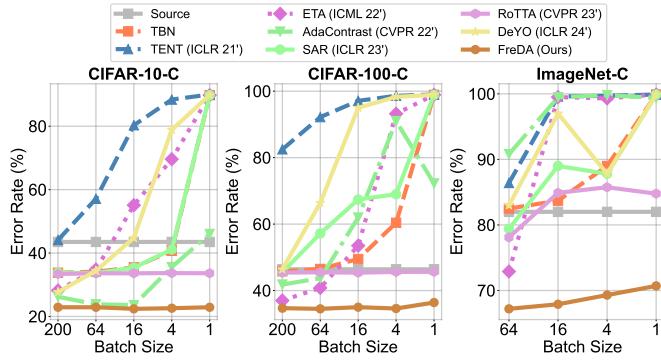
$$\begin{aligned} \text{Err}(G) &\leq \underbrace{\frac{2}{(c_{\text{base}} + \gamma) - 1} \left(\text{Err}(G_{\text{pl}}) - \sum_{k=1}^K \Delta_k \right)}_{\text{Reduced by both terms}} \\ &\quad + \underbrace{\frac{2(c_{\text{base}} + \gamma)}{(c_{\text{base}} + \gamma) - 1} \mu}_{\text{Reduced by } O(\gamma/c^2)} \end{aligned}$$

Notation Summary

- c : Expansion factor (Def. 1).
Class-wise connectivity metric: Larger c implies stronger neighborhood propagation of local consistency to global predictions.
- μ : Inter-class separation probability (Def. 2).
Robustness measure: Probability of different classes having overlapping neighborhoods under perturbations, lower μ indicates clearer decision boundaries.
- \mathcal{R}_B : Consistency regularizer (Thm. 1).
Stability term: Penalizes prediction inconsistency between original inputs and their augmented variants.
- Δ_k : Error reduction in k -th sub-domain.
Specialization gain: Reduced pseudo-label noise in sub-domain k due to frequency-based partitioning.
- γ : Expansion gain from frequency augmentation.
Neighborhood enhancement: Additional expansion capability measured as the relative increase of augmented neighborhoods.

TABLE III: Classification error rate (\downarrow) on DomainNet126 and Camelyon17 under **Mixed Distribution Shifts**.

Methods	DomainNet126					Camelyon17					
	Real	Painting	Clipart	Sketch	Avg.	C1	C2	C3	C4	C5	Avg.
Source	45.2	41.6	49.5	45.3	45.4	21.6	43.6	52.5	47.4	47.6	42.5
TBN	45.5	39.9	45.9	37.5	42.2	26.5	38.5	31.7	39.4	32.8	33.8
TENT (ICLR 21')	42.2	37.8	44.7	37.5	40.6	44.7	50.5	49.9	49.1	48.6	48.6
ETA (ICML 22')	41.1	37.3	43.4	36.4	39.5	47.4	52.5	47.9	49.9	39.2	47.4
SAR (ICLR 23')	43.2	38.5	44.8	37.0	40.9	26.5	38.5	31.7	39.4	32.8	33.8
DeYO (ICLR 24')	40.9	36.4	43.6	36.9	39.4	50.4	50.3	48.8	51.7	50.5	50.4
FreDA (ours)	40.2	36.1	40.0	33.6	37.5	18.6	24.7	24.8	40.5	30.8	27.9

Fig. 5: Averaged classification error rate (\downarrow) on CIFAR-10-C/100-C and ImageNet-C with various batch size under **Mixed Domains**.

epochs with other training specifications outlined in the Wilds benchmark [44].

4) *Hyperparameter Configuration*: The batch size is set to 200, 64, 128 and 32 for CIFAR-10/100-C, ImageNet-C, DomainNet126 and Camelyon17 following the previous methods. The SGD optimizer is used with learning rates adjusted to 0.01, 0.0001, 0.001 and 0.00005, respectively. The learning rate is proportionally decreased in the experiment studying the effect of batch size. The Kmeans Size is 512, Clutser Number is 4, Communication Interval is 10 across all the tasks. The perturbation magnitude α is fixed to 0.1 and the coefficient λ in loss function is fixed to 0.5. Two threshold in Eq. 5 is set to the same value for corruption datasets and DomainNet126 following ETA [6]. While for Camelyon17, the class diversity related threshold is adjusted to 0.9 empirically.

B. Main Results

1) *FreDA improves across diverse distribution shifts*: Our method consistently attains the lowest error rates across all evaluated datasets (see TABLE V and III). Notably, on the Camelyon17 dataset, FreDA reduced the error rate to 27.9%, outperforming the next best method by 5.9%. This significant improvement is particularly noTABLE where other approaches falter if compared with no training (TBN), meaning they struggle to adapt to the complex medical imaging data. By effectively handling high variability and intricate patterns in the data, FreDA maintains superior accuracy and adaptability.

2) *FreDA remains stable under various batch size*: To simulate deployment with constrained batch sizes, we evaluate models under both varying batch sizes and mixed distributions.

TABLE IV: Ablation study of FreDA.

DT	SS	SA	C10	C100	IN
			44.1	82.5	86.4
✓			24.8	54.2	81.2
	✓		29.6	37.5	71.0
✓	✓	✓	39.4	71.7	92.9
✓	✓	✓	24.3	36.3	69.4
✓	✓	✓	27.7	36.2	65.9
✓	✓	✓	24.4	50.2	77.7
✓	✓	✓	22.9	34.7	67.2

In Figure 5, we present the results on CIFAR-10-C, CIFAR-100-C, and ImageNet-C using batch sizes ranging from 200 (64) down to 1. Unlike other methods that significantly degrade as batch size decreases – for example the error rate of DeYO increases from 27.7% to 89.8% when batch size drops from 200 to 1 on CIFAR-10-C – FreDA consistently maintains strong performance. This stability demonstrates FreDA’s robustness, making it highly suitable for real-world applications where large batches is not always feasible.

3) *FreDA enhances adaptation via synergistic designs*: This section validates our designs by ablating three key modules – Decentralized Training (DT), Sample Selection (SS), and Sample Augmentation (SA). The baseline here leverages only the entropy loss. From TABLE VI, we have the following observations: 1) Implementing decentralized training and sample selection alone results in substantial improvements, reducing error rates dramatically across all datasets. 2) Sample augmentation alone has the possibility to increase error rates, suggesting that although this approach introduces useful variability, it may introduce unexpected noise under the absence of proper selection or decentralized training. 3) The combined approach delivers the best performance across all datasets, showing the synergistic effect of our different designs.

C. More analysis and Discussion

1) *Performance with Transformer Backbone*: In addition to evaluating our model on commonly compared CNN backbones, we further assess its performance under transformer-based architecture, specifically using the ViT-Base backbone. Here, we report results on the ImageNet-C benchmark using ViTBase-LN [45] (see TABLE V), where the pretrained weights are obtained from `torchvision`. Importantly, all experimental configurations are kept consistent with those used in the CNN-based experiments. As shown, our method continues to deliver strong performance, demonstrating its robustness and adaptability across different backbone architectures.

integrating Fourier-based augmentation, we expand the range of confident samples for each distribution shift, further boosting model performance. The experimental results underscore the efficacy of FreDA, highlighting its potential to influence the field and guide future research in adapting to dynamic and diverse data shifts. While FreDA addresses a critical challenge in handling heterogeneous data streams, providing a solid pipeline for this issue, there are still avenues for further enhancement. Our current aggregation approach, which averages models based on cluster counts, has been effective in solving the problem at hand. However, exploring alternative strategies – such as weighting models by the divergence between clusters – might lead to incremental improvements. Additionally, refining the sample selection process from a original sample-level focus to a more granular patch-level could extend FreDA’s applicability to tasks such as segmentation, further enhancing its versatility in real-world scenarios.

APPENDIX

A. Adaptation Scenarios

1) *Mixed Domains*: In this scenario, the model processes a long sequence of test samples where each sample $x_i \sim p_{e_i}(x)$ is independently drawn from a randomly selected target domain $\mathcal{D}_{e_i} \in \{\mathcal{D}_{t_1}, \mathcal{D}_{t_2}, \dots, \mathcal{D}_{t_N}\}$ and a randomly selected class c_i among classes $\{1, 2, \dots, C\}$. The sequence is represented as:

$$\left\{ x_1^{\mathcal{D}_{e_1}, c_1}, x_2^{\mathcal{D}_{e_2}, c_2}, \dots, x_k^{\mathcal{D}_{e_k}, c_k} \right\},$$

where each target domain index $e_i \in \{1, 2, \dots, N\}$ and class number $c_i \in \{1, 2, \dots, C\}$ are independently and randomly selected for each sample x_i .

2) *Continual Domain Adaptation*: In this setting, the model adapts online to a sequentially presented series of test domains, where each domain shift occurs only once and does not reappear. The sequence progresses through distinct target domains in a fixed order $D_1 \rightarrow D_2 \rightarrow \dots \rightarrow D_N$, with samples from each domain appearing contiguously. The class labels within each domain are independently and randomly selected. The sequence is structured as:

$$\begin{aligned} & \underbrace{\{(x_1^{D_1, c_1}), (x_2^{D_1, c_2}), \dots, (x_{l_1}^{D_1, c_{l_1}})\}}_{\text{Samples from domain } D_1} \\ & \rightarrow \underbrace{\{(x_{l_1+1}^{D_2, c_{l_1+1}}), \dots, (x_{l_1+l_2}^{D_2, c_{l_1+l_2}})\}}_{\text{Samples from domain } D_2} \\ & \rightarrow \dots \rightarrow \underbrace{\{(x_{l_1+\dots+l_{N-1}+1}^{D_N, c_{l_1+\dots+l_{N-1}+1}}), \dots, (x_k^{D_N, c_k})\}}_{\text{Samples from domain } D_N} \end{aligned}$$

where:

- D_i denotes the i -th target domain in the fixed sequence, with $i \in \{1, 2, \dots, N\}$.
- $c_m \in \{1, 2, \dots, C\}$ is the randomly selected class label for the m -th sample, independent of domain transitions.

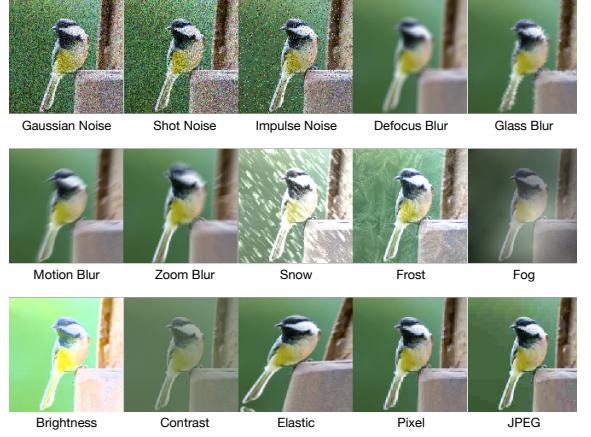


Fig. 6: Examples from ImageNet-C under common image corruptions. The images showcase a range of corruption types (e.g., noise, blur, and weather distortions) at varying severity levels.



Fig. 7: Samples from DomainNet126 across four subdomains (Real, Sketch, Painting, Clipart). These visualizations reflect the stylistic and perceptual variations inherent in each domain.

B. Dataset Visualization

To further illustrate the characteristics of the datasets used in our evaluation, we present visualizations of the data distribution across different corruption types (Fig. 6), natural domain shifts (Fig. 7), and medical centers (Fig. 2). These figures highlight the diverse challenges that our models face in each evaluation scenario, providing insight into the complexity of the test conditions.

C. Relation to Frequency Domain Learning

Recent advances highlight frequency-based techniques as powerful tools for domain transfer. In domain generalization, frequency analysis has revealed critical insights into model robustness and learning dynamics [46]–[53]. For domain adaptation, interpolating image amplitude spectra across styles has proven effective in reducing domain gaps and preventing overfitting to low-level statistics [54]–[57]. Motivated by these advancements, we discover that frequency information inherently captures domain characteristics, making it a valuable medium for decoupling mixed target domains – an aspect largely unexplored in prior work. On this basis, we propose

FreDA, the first framework addressing TTA under mixed domain shifts in Fourier space, with a decentralized adaptation and perturbation mechanism.

D. Relation to Multi-Target Domain Adaptation

TTA under mixed distribution shifts shares similarities with multi-target unsupervised domain adaptation (MT-UDA) [11], [12], [58], [59], yet diverges critically in complexity: while MT-UDA assumes static, predefined target domains and leverages labeled source data for explicit domain alignment, TTA operates with no access to source data and must adapt to dynamic, unpredictable target streams in an online manner. This eliminates direct source-target discrepancy computation and demands robust incremental adaptation rather than offline multi-domain optimization. These differences highlight the need for novel methodologies tailored specifically to TTA, going beyond the solutions developed for MT-UDA.

E. Relation to Decentralized, Federated, Distributed Learning

This work also intersects with decentralized, federated, and distributed learning due to splitting data batches into disjoint subsets and applying decentralized model adaptation. First, while decentralized learning focuses on non-i.i.d. data that is naturally distributed across multiple nodes [60], our approach starts with a centralized batch of target samples. By proactively splitting this data into disjoint subsets, we expose latent non-i.i.d. characteristics, enabling the effective use of decentralized learning techniques. Second, federated learning considers data privacy and model collaborations within decentralized learning [61]. In our case, as target samples are mixed in a batch, data privacy is not a concern. However, similar to federated learning, our approach also involves weight aggregation over subnetworks to enhance their base models before next batch adaptation. Third, distributed learning aims to improve training efficiency on large-scale datasets by partitioning data for synchronized training [62]. In contrast, our method operates in a real-time fine-tuning context with limited data at one time, hence scalability is less of a concern.

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