Are Domain Generalization Benchmarks with Accuracy on the Line Misspecified?

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Abstract

Spurious correlations are unstable statistical associations that hinder robust decision-making. Conventional wisdom suggests that models relying on such correlations will fail to generalize out-of-distribution (OOD), particularly under strong distribution shifts. However, a growing body of empirical evidence challenges this view, as naive empirical risk minimizers often achieve the best OOD accuracy across popular OOD generalization benchmarks. In light of these counterintuitive results, we propose a different perspective: *many widely used benchmarks for assessing the impact of spurious correlations on OOD generalization are misspecified.* Specifically, they fail to include shifts in spurious correlations that meaningfully degrade OOD generalization, making them unsuitable for evaluating the benefits of removing such correlations. We establish sufficient—and in some cases necessary—conditions under which a distribution shift can reliably assess a model's reliance on spurious correlations. Crucially, under these conditions, we provably should not observe a strong positive correlation between in-distribution and out-of-distribution accuracy—often referred to as accuracy on the line. Yet, when we examine state-of-the-art OOD generalization benchmarks, we find that most exhibit accuracy on the line, suggesting they do not effectively assess robustness to spurious correlations. Our findings expose a limitation in evaluating algorithms for domain generalization, i.e., learning predictors that do not rely on spurious correlations. Our results highlight the need to rethink how we assess robustness to spurious correlations.

1 Introduction

Domain generalization aims to develop predictors that generalize to arbitrarily new and potentially *worst-case* unobserved distributions (Arjovsky et al., 2019; Rosenfeld et al., 2020). Spurious correlations, also referred to as shortcuts, are coincidental statistical associations between features and labels in training data that fail to generalize beyond the training distribution, hindering domain generalization (Nagarajan et al., 2020; Geirhos et al., 2020; Makar et al., 2022). For example, a medical diagnosis model trained on data from one hospital may be less effective in a different hospital with distinct patient demographics, equipment, or clinical practices (Zech et al., 2018; Yang et al., 2024). Similarly, an animal detection model trained on data from one environment and sensor type may be less effective in classifying animals in images captured under a different environment and sensor configuration (Beery et al., 2018; Xiao et al., 2020).

Thus, many algorithms for domain generalization have been focused on learning predictors that ignore these unreliable patterns—i.e., *invariance* or *feature disentanglement* methods (Arjovsky et al., 2019; Krueger et al., 2021; Wang et al., 2019; Parascandolo et al., 2020; Creager et al., 2021; Ahuja et al., 2021; Shi et al., 2021; Zhou et al., 2022; Wang et al., 2022b; Li et al., 2022; Salaudeen & Koyejo, 2024). Zhou et al. (2022); Wang et al. (2022b) provide a more comprehensive survey on domain generalization (Lin et al., 2023; Chen et al., 2022b; 2023b; Zhang et al., 2022). However, empirical evidence suggests that standard empirical risk minimization (ERM), which may leverage spurious correlations (Xiao et al., 2020), often achieves the highest out-of-distribution (OOD) accuracy on widely used benchmarks (Gulrajani & Lopez-Paz, 2020a; Yao et al., 2022; Gagnon-Audet et al., 2022; Yang et al., 2023). Additionally, Taori et al. (2020); Miller et al. (2020; 2021); Wenzel et al. (2022); Baek et al. (2022); Saxena et al. (2024); Nastl & Hardt (2024) demonstrate a strong correlation between in- and out-of-distribution accuracy across several state-of-the-art benchmarks, suggesting

that better in-distribution (ID) performance generally predicts better out-of-distribution performance. Nastl & Hardt (2024) further shows that a model that uses all available features generally results in better OOD performance than a model that uses a select subset of *causal features* for popular distribution shift tabular datasets (Gardner et al., 2023). This observation is seemingly counter to the thesis that there can be statistical patterns in training data that can improve ID performance but worsen OOD performance (Geirhos et al., 2020; Makar et al., 2022). These findings raise a critical question: *Does a better in-distribution classifier imply a better out-of-distribution classifier, challenging the necessity of targeted algorithms for domain generalization?*

We propose and provide evidence for an alternative explanation of ERM's success in domain generalization. Drawing on the concept of underspecification in modern machine learning pipelines (D'Amour et al., 2022), we argue that ERM's apparent superiority may stem from the (mis/under)specification of popular domain generalization benchmarks.

Specifically, we consider the setting where correlations exist in training data and a subset (spurious) shift at test time. We investigate the types of shifts under which predictions from a classifier that relies on these spurious correlations generalize worse out-of-distribution than a classier that only uses the static (domain-general) correlations, i.e., naive ERM is insufficient. We call these types of shifts *well-specified*. We argue that when the best in-distribution classifier is also the best out-of-distribution classifier on a benchmark, this benchmark does not represent the types of settings domain generalization is concerned with. The core of this work studies a benchmark's ability to distinguish between two types of predictors in this setting: one that ignores spurious correlations (domain-general or invariant) and another that leverages all available correlations (including spurious) that maximize in-distribution accuracy (domain-specific).

1.1 Our Contributions

- We establish a negative margin under distribution shift condition (or spurious correlation reversal) that characterizes well-specified domain generalization benchmarks—Theorem 1.
- We show that well-specified benchmarks typically lack a strong correlation between in and out-of-distribution accuracy for a diverse set of classifiers—such a strong correlation has been termed *accuracy on the line* (Miller et al., 2020)—Theorem 3. **When there is positive accuracy on the line, the dataset may be misspecified for evaluating domain generalization.**
- With over 40 total ID/OOD splits across 12 benchmarks, we show that many of the state-of-the-art benchmarks exhibit (*accuracy on the line*) and may be misspecified for domain generalization. Code*

While the theoretical and empirical results we present subsequently provide actionable insights for designing and using domain generalization benchmarks, we will also discuss their implications on benchmarking norms and practices, e.g., model selection, algorithmic fairness, causal representation learning, etc., in the context of both predictive and generative models.

2 Theoretical Analysis: (Mis)specification of Domain Generalization Benchmarks

Following previous work (Wang et al., 2019; Rosenfeld et al., 2020; Ahuja et al., 2021; Salaudeen & Koyejo, 2024) we define domain-general (`dg`) features $\mathcal{Z}_{\text{dg}} \subseteq \mathbb{R}^k$, where the optimal predictor that only uses these domain-general features is desired. Conversely, spurious features (`spu` or domain-specific) $\mathcal{Z}_{\text{spu}} \subseteq \mathbb{R}^l$ contain additional domain-specific information that improves the prediction task in-distribution, but their use can degrade performance out-of-distribution. The observed features $\mathcal{X} \subseteq \mathbb{R}^d$ are a concatenation of $\mathcal{Z}_{\text{dg}}$ and $\mathcal{Z}_{\text{spu}}$, where $d = k + l$. We also define $\mathcal{Y} = \{\pm 1\}$. P's represent probability distributions over X, Y. Let $\mathcal{E}$, where $|\mathcal{E}| > 1$, denote the set of distributions of interest. $P \in \mathcal{E}$ implies that marginals without Z_{spu} are preserved.

We consider classifiers $f \in \mathcal{F} : \mathcal{X} \mapsto \mathcal{Y}$ of the form $f(X) = Z_{\text{dg}}^T \text{w}_{\text{dg}} + Z_{\text{spu}}^T \text{w}_{\text{spu}}$ where $\text{w}_{\text{dg}} \in \mathbb{R}^k$ and $\text{w}_{\text{spu}} \in \mathbb{R}^l$. We note empirical findings support our parameterization of Z_{dg} and Z_{spu} from *non-linear transformation* when a final linear mapping is applied, e.g., kernel regressors or common deep neural networks. Rosenfeld

*https://anonymous.4open.science/status/misspecified_DG_benchmarks-E5FD.

et al. (2022a) demonstrates that improving out-of-distribution performance can be done with new linear classifiers on learned (non-linear) representations. Additional work shows that last-layer retraining improves robustness to spurious correlations (Liu et al., 2021b; LaBonte et al., 2023). These findings suggest that the non-spurious and spurious representations learned by deep models are often linearly separable, suggesting that our assumption is reasonable. Importantly, we allow for a nonlinear shift in spurious correlation.

Within $\mathcal{F}$, $\mathcal{F}_{\text{dg}} \subset \mathcal{F}$ comprises functions f_{dg} that only use domain-general features $f_{\text{dg}}(X) = f_{\text{dg}}([Z_{\text{dg}}; 0])$. Additionally, denote $f_{\text{X}} \in \mathcal{F} \backslash \mathcal{F}_{\text{dg}} := \mathcal{F}_{\text{X}}$, where f_{X} uses both domain-general and domain-specific features. The function $\ell(\cdot, \cdot) \mapsto \mathbb{R}$ denotes a loss function, and $\mathcal{R}^e(f) = \mathbb{E}_{P_e}[\ell(Y, f(X))]$ defines the expected loss for some function $f \in \mathcal{F}$. Additionally, we define the accuracy of $f \in \mathcal{F}$ on distribution P as

$$\text{acc}_P(f) = \mathbb{E}_{(X,Y)\sim P}\left[\mathbf{1}(f(X) \cdot Y > 0)\right] = \Pr\left(f(X) \cdot Y > 0\right) \tag{1}$$

We first formally define spurious features and domain-general features (Definitions 1- 2). The relationship between spurious features and the label we want to predict is allowed to change across domains and can negatively impact out-of-distribution performance, while the relationship between domain-general features and the label is stable across domains. For instance, when predicting medical diagnoses from chest X-rays, the relationship between physiological features and diagnoses is expected to be stable from site to site (domain-general), while the relationship between site-specific markings on X-rays and diagnoses is unstable (spurious); predictors relying on site-specific markings fail out-of-distribution (Zech et al., 2018).

Definition 1 (Domain-General Features Z_{dg})**.** *For all P_i, $P_j \in \mathcal{E}$,*

$$\mathbb{E}_{P_i}[Y \mid Z_{dg}] = \mathbb{E}_{P_j}[Y \mid Z_{dg}]. \tag{2}$$

Definition 2 (Spurious Features Z_{spu})**.** *For all P_i, $P_j \in \mathcal{E}$,*

$$\mathbb{E}_{P_i}[Y \mid Z_{spu}] \neq \mathbb{E}_{P_j}[Y \mid Z_{spu}] \qquad \mathbb{E}_{P_i}[Y \mid Z_{dg}, Z_{spu}] \neq \mathbb{E}_{P_j}[Y \mid Z_{dg}, Z_{spu}]. \tag{3}$$

We assume both types of features are informative about labels (Assumption 1) and are not redundant (Assumption 2). Clearly, if Assumption 1 does not hold, then the learning problem is ill-posed; features are uncorrelated with labels. When Assumption 2 does not hold, spurious features are redundant and have no unique information about the labels. Ahuja et al. (2021) study this setting (*Fully Informative Invariant Features (FIIF)*; we use 'domain-general' instead of 'invariant') and give conditions under which predictors using spurious correlations can achieve equal OOD accuracy as the optimal invariant predictor. Hence, we focus on the partially informative domain-general features setting.

Assumption 1 (Informative Domain-General and Domain-Specific Features)**.** *For all observed training distributions P,*

$$\mathbb{E}_P[Y \mid Z_{dg}] \neq \mathbb{E}_P[Y] \text{ and } \mathbb{E}_P[Y \mid Z_{spu}] \neq \mathbb{E}_P[Y]. \tag{4}$$

Assumption 2 (Non-Redundant Features)**.**

$$Z_{spu} \not\perp\!\!\!\perp Y \mid Z_{dg} \text{ and } Z_{dg} \not\perp\!\!\!\perp Y \mid Z_{spu}. \tag{5}$$

Also referred to as partially informative domain-general features.

Since we consider any feature whose inclusion decreases worst-case performance on the set of distribution of interest $\mathcal{E}$ as spurious, e.g., the set of hospitals one expects a predictor to have to operate in, the definition of spurious is strongly tied to the $\mathcal{E}$ 'worst-case' is with respect to. What is considered spurious for $\mathcal{E}' \neq \mathcal{E}$ may differ, even if $\mathcal{E}' \subset \mathcal{E}$. Notably, this observation implies that domain generalization cannot practically be divorced from domain expertise in defining $\mathcal{E}$. Clearly, too narrow of an $\mathcal{E}$ decreases expected robustness (potentially catastrophically), and too broad of an $\mathcal{E}$ may excessively and unnecessarily decrease overall utility.

We define two predictors: (i) the optimal domain-general predictor, which depends on $\mathcal{E}$ (Definition 3) and (ii) the optimal domain-specific predictor for a given $P \in \mathcal{E}$ (Definition 4).

Definition 3 (Optimal Domain General Predictor $f_{\text{dg}}^{\mathcal{E}}$)**.** *Given a set of distributions of interest* $\mathcal{E} = \{P_i(X, Y) : i = 1, \ldots\}$,

$$f_{dg}^{\mathcal{E}} = \underset{f \in \mathcal{F}_{dg}}{\text{argmax}} \min_{P_i \in \mathcal{E}} acc_{P_i}(f). \tag{6}$$

By construction, $f_{dg}^{\mathcal{E}} \in \mathcal{F}_{dg}$ *does not use spurious features.*

Definition 4 (Optimal Domain Specific Predictor f_{X}^{P})**.** *Given a distribution* P *and*

$$f_X^P = \underset{f \in \mathcal{F}}{\text{argmax}}\, acc_P(f). \tag{7}$$

By construction, $f_X^P \in \mathcal{F}_X$ *uses spurious features (Lemma 1).*

Our first result—Lemma 1—shows that, given informative and non-redundant features (Assumptions 1-2) and for any distribution $P \in \mathcal{E}$, the optimal $\mathcal{E}$-domain-general and P-domain-specific predictor are different, and the optimal domain-specific predictor achieves a lower (convex) loss in-distribution than the optimal domain-general predictor. This also contextualizes the rest of our results in that optimal domain-specific models use all non-redundant features that improve performance.

Lemma 1 (Domain-General and Domain-Specific In-Distribution Error Gap)**.** *Assume non-redundant/non-trivial features, partially informative domain-general features (Assumptions 1-2), and strongly convex loss* ℓ.

$$\min_{f \in \mathcal{F}} \mathcal{R}^e(f) < \min_{f \in \mathcal{F}_{dg}} \mathcal{R}^e(f), \tag{8}$$

where $\mathcal{F} : \mathcal{X} \to \mathbb{R}$ *where* $f(x) = w^\top x = w_{dg}^\top z_{dg} + w_{spu}^\top z_{spu}$, $f \in \mathcal{F}$. *For* $f \in \mathcal{F}_{dg}$, $f(x) = w^\top x = w_{dg}^\top z_{dg}$. *The proof of Lemma 1 is provided in Appendix A.1.*

Given that the in-distribution risk minimizer and the domain-general predictor differ, we now show that given training and test distributions $P_{\text{ID}} \neq P_{\text{OOD}} \in \mathcal{E}$, respectively, the domain-general predictor $f_{\text{dg}}^{\mathcal{E}}$ may also not achieve a higher OOD accuracy than in-distribution risk minimizer $f_{\text{X}}^{P_{\text{ID}}}$ on P_{OOD}. We derive sufficient conditions on P_{OOD} such that $f_{\text{dg}}^{\mathcal{E}}$ achieves a higher OOD accuracy than $f_{\text{X}}^{P_{\text{ID}}}$—Theorem 1. Such $P_{\text{ID}}, P_{\text{OOD}}$ ID/OOD splits make for 'well-specified' benchmarks, as outlined in the following—Definition 5. Our conditions illustrate the need for sufficient misalignment between spurious correlations between P_{ID} and P_{OOD}.

Definition 5 (Well-Specified Domain Generalization Benchmark)**.** *Two ID/OOD splits,* $P_{ID}, P_{OOD} \in \mathcal{E}$, *are 'well-specified' if and only if*

$$acc_{P_{OOD}}\big(f_X^{P_{ID}}\big) < acc_{P_{OOD}}\big(f_X^{\mathcal{E}}\big), \tag{9}$$

where $f_{dg}^{\mathcal{E}}$, $f_X^{P_{ID}}$ *are from Definitions 3 and 4, respectively—note that this definition is w.r.t. to accuracy.*

Analysis Setting. For the rest of this work, we consider sub-Gaussian $Z_{\text{spu}}^{\text{ID}}$ with mean μ_{spu}, covariance Σ_{spu}, and sub-Gaussian parameter κ—importantly, *our results also apply more generally to other classes of random variables (Remark 1).* We define an L_ϕ-Lipschitz function which parametrizes the distribution shift w.r.t. Z_{spu}; $\phi : \mathbb{R}^{l \times l} \to \mathbb{R}^{l \times l}$ such that $Z_{\text{spu}}^{\text{OOD}} = \phi(Z_{\text{spu}}^{\text{ID}})$ and $\mathbb{E}[Z_{\text{spu}}^{\text{OOD}}] = M\mathbb{E}[Z_{\text{spu}}^{\text{ID}}] = M\mu_{\text{spu}}$ where $M \in \mathbb{R}^{l \times l}$ and Σ_ϕ is the covariance of $Z_{\text{spu}}^{\text{OOD}}$.

Our overall goal is to identify shifts where achieving higher transfer accuracy is informative about the reliance of predictions on spurious correlations. Identifying the class of such shifts is necessary as they describe ID/OOD splits for domain-generalization benchmarks where achieving the higher OOD accuracy meaningfully maps to learning domain-general predictors, i.e., the shift is well-specified for the task. Without such well-specified shifts, we would incorrectly assume that a domain-general predictor is ineffective because it achieves a lower accuracy out-of-distribution than a different empirical risk minimizer. Thus, our next result shows that when shifts (ϕ) result in a sufficient misalignment between spurious correlations before and after the shift, the domain-general model achieves a higher out-of-distribution accuracy than a domain-specific predictor.

Remark 1. *Our results consider sub-Gaussian spurious features. However, our results hold for other classes of random variables, e.g., sub-exponential, bounded moments, or other random variables in Orlicz spaces (Krasnoselskii, 1960). In these cases, our proofs remain largely unchanged, with only the constant factors adjusted to account for the different concentration properties of the spurious features.*

Theorem 1 (Sufficient Conditions for Well-Specified Domain Generalization Splits)**.** *Assume Z_{spu}^{ID} is sub-Gaussian with mean μ_{spu}, covariance Σ_{spu}, and parameter κ. Define a nonlinear transformation*

$$\phi : \mathbb{R}^l \to \mathbb{R}^l,$$

that is L_ϕ–Lipschitz, and let

$$Z_{spu}^{OOD} = \phi(Z_{spu}^{ID}).$$

Assume further that

$$\mathbb{E}[Z_{spu}^{OOD}] = \mathbb{E}[\phi(Z_{spu}^{ID})] = M\,\mu_{spu},$$

for some matrix $M \in \mathbb{R}^{l\times l}$. In-distribution, $Z_{spu}^{ID} \sim P_{ID}$ and out-of-distribution, $Z_{spu}^{OOD} \sim P_{OOD}$. Additionally, denote w_{spu} the contribution of Z_{spu}^{ID} to the optimal P_{ID} predictor $f_X^{P_{ID}}$. Then, for any $\delta \in (0,1)$, if

$$w_{spu}^\top (M\,\mu_{spu}) + \sqrt{2\,(L_\phi\,\kappa)^2\,\|w_{spu}\|_2^2 \log(1/\delta)} < 0,$$

(with the understanding that under the Lipschitz assumption, the sub-Gaussian property carries over with parameter $L_\phi\,\kappa$), then with probability at least $1-\delta$ over Z_{spu}^{OOD}, we have

$$acc_{P_{OOD}}(f_X^{P_{ID}}) < acc_{P_{OOD}}(f_{dg}^{\mathcal{E}}),$$

where $f_{dg}^{\mathcal{E}}$ and $f_X^{P_{ID}}$ are the optimal domain–general and domain–specific predictions (Definitions 3–4).

Proof provided in Appendix A.2.

Theorem 2 (Neccesary and Sufficient Conditions for Well-Specified Domain Generalization Splits under Distributional Symmetry)**.** *Assume Z_{spu}^{ID} is a random variable with mean μ_{spu}, covariance Σ_{spu}, and $w_{dg}^\top Z_{dg}^{OOD}$ is symmetric about its mean. Define a nonlinear transformation*

$$\phi : \mathbb{R}^l \to \mathbb{R}^l,$$

that is L_ϕ–Lipschitz, and let

$$Z_{spu}^{OOD} = \phi(Z_{spu}^{ID}).$$

Assume further that

$$\mathbb{E}[Z_{spu}^{OOD}] = \mathbb{E}[\phi(Z_{spu}^{ID})] = M\,\mu_{spu}, \quad \text{and} \quad \mathrm{Var}(Z_{spu}^{OOD}) = \Sigma_\phi$$

for some matrix $M \in \mathbb{R}^{l\times l}$, and that the scalar and $w_{spu}^\top Z_{spu}^{OOD}$ is symmetric about its mean. Then,

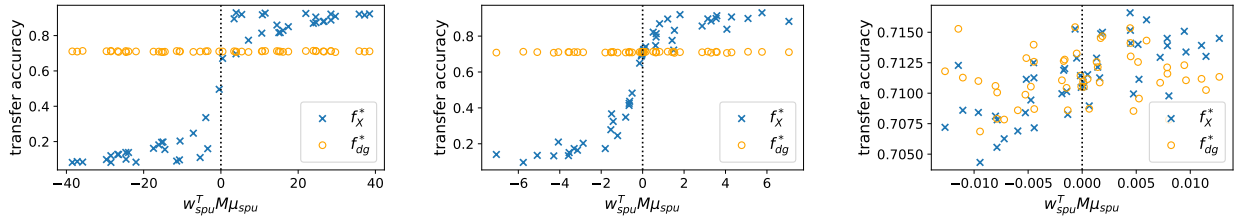
$$acc_{P_{OOD}}(f_X^{P_{ID}}) < acc_{P_{OOD}}(f_{dg}^{\mathcal{E}}) \iff \frac{w_{dg}^\top \mu_{dg} + w_{spu}^\top M \mu_{spu}}{\sqrt{w_{dg}^\top \Sigma_{dg} w_{dg} + w_{spu}^\top \Sigma_\phi w_{spu}}} < \frac{w_{dg}^\top \mu_{dg}}{\sqrt{w_{dg}^\top \Sigma_{dg} w_{dg}}}. \tag{10}$$

Note that the RHS of Equation 10 implies that either:

- Spurious Correlation Reversal: $w_{spu}^\top M \mu_{spu} < 0$ *(related to sufficiency condition without the symmetry assumption in Theorem 1), or*
- *sufficiently large variance for $w_{spu}^\top M \mu_{spu} > 0$, i.e., small SNR.*

Proof provided in Appendix A.3.

Theorem 1 demonstrates that a domain generalization split is well-specified if and only if (i) the OOD spurious correlation is misaligned with the ID spurious correlation and (ii) the variance of spurious features is sufficiently controlled not to undo the effect of misalignment. For symmetrically distributed features, e.g.,



(a) **All random variables are Gaussian.** When Theorem 1 conditions are satisfied OOD (x-axis: $\mathrm{w}_{\mathrm{spu}}^{\top} M \mu_{\mathrm{spu}} + \mathrm{c} < 0$; $c > 0$), the f_{dg}'s outperform f_{X}'s OOD. This result verifies that there needs to be sufficient misalignment between in- and out-of-distribution spurious correlations for the domain-general features to outperform the domain-specific models in OOD accuracy.

(b) $\mathbf{Z}_{\mathrm{spu}}^{\mathrm{ID}}$ **is defined to be a mixture of 4 Gaussian distributions (sub-Gaussian) such that the test distributions are not a mixture of the same 4 Gaussians.** The same conclusions in Figure 1a hold for sub-Gaussian random variables when the test distribution is an extrapolation.

(c) Rosenfeld et al. (2022b) studies domain interpolation, where the target domain is a mixture of the training domain; a variant of ERM is provably worst-shift optimal. $\mathbf{Z}_{\mathrm{spu}}^{\mathrm{ID}}$ **is defined to be a mixture of 4 Gaussian distributions such that the test distributions are a different mixture of the 4 Gaussians.** Overall, there is minimal difference in OOD accuracy between f_{dg} and f_{X} in this setting (domain interpolation).

Figure 1: f_{dg} are trained on (Z_{dg}, Y) pairs and f_{X} are trained on (X, Y) pairs from the same distribution. $X = Z_{\mathrm{dg}} \oplus Z_{\mathrm{spu}}^{\mathrm{ID}}$, where $\oplus$ is concatenation. $Z_{\mathrm{spu}}^{\mathrm{ID}}$ is sub-Gaussian. We evaluate these models on 50 test distributions generated with randomly sampled M such that all other distributions are the same ID and OOD in the test distribution but $Z_{\mathrm{spu}}^{\mathrm{OOD}} = M Z_{\mathrm{spu}}^{\mathrm{ID}}$. Details on the experiments can be found in Appendix B.

Gaussians, these conditions are necessary and sufficient, and hold with certainty (Theorem 1)) illustrates this with Gaussian features..

For example, consider the classic examples of predictors using background, pasture or desert, to predict cows and camels, respectively. This means that the correlation between background and label has to lead to sufficient disagreement (reversed) between training and test distributions such that their use harms OOD (test) predictions.

Importantly, we are concerned with benchmarking predictors designed to not rely on spurious correlations for prediction, even if they are useful and potentially reliable predictors *most of the time*. For example, site-specific markings or orientation of chest X-rays in medical diagnosis (Jabbour et al., 2020), demographic attributes in resource allocation prediction tasks (Chouldechova, 2017; Kumar et al., 2022), etc. Figure 1 verifies these conditions empirically with simulation experiments. Additionally, semi-synthetic examples with variants of the ColoredMNIST dataset is provided in Appendix B.1 (Arjovsky et al., 2019).

Remark 2 (Multisource Domain-Generalization.)**.** *We often have access to a set of distributions for evaluation, and the norm is to evaluate leave-one-domain-out ID and OOD splits (Gulrajani & Lopez-Paz, 2020a). In the context of Definition 5, P_{ID} represents the mixture of ID distributions, and the test split is P_{OOD} (which can also be a mixture). Importantly, a finite mixture of sub-Gaussians is also sub-Gaussian (Appendix A.8 Lemma 6), so our results directly apply without loss of generality. Notably, we focus on ID/OOD split settings. While a set of domains may give many splits, only a subset of the splits may be well-specified.*

So far, we have shown that the learned domain-general and domain-specific predictors on a given training distribution differ, and the domain-general model achieves higher OOD accuracy (well-specified) when spurious correlations ID and OOD are sufficiently misaligned. Next, suppose we observe such misalignment and the split is well-specified; then, we evaluate a set of diverse predictors on held-out ID and OOD test data. We should observe a weak correlation or a strong negative correlation between the predictors' ID and OOD accuracy, i.e., no positive *accuracy on the line. When we observe accuracy on the line, with a high probability, the ID/OOD split is misspecified.*

2.1 Accuracy on the Line

First, we define *accuracy on the line*, the correlation strength between in- and out-of-distribution accuracy—Definition 6.

Definition 6 (Accuracy on the Line; Miller et al. (2021))**.** *Define $a \in \mathbb{R}$, $\epsilon \geq 0$, and Φ^{-1} as the inverse Gaussian cumulative density function. The* correlation property *is defined as*

$$\left|\Phi^{-1}\left(acc_{P_{ID}}(f)\right) - a \cdot \Phi^{-1}\left(acc_{P_{OOD}}(f)\right)\right| \leq \epsilon \, \forall \, f, \tag{11}$$

where f's are distinct predictors.

If there exists an a such that $\epsilon = 0$, then there is a perfect correlation between ID and OOD accuracy. As ϵ grows, the strength of the correlation decreases. If $a > 0$, then the correlation is positive, and if $a < 0$, the correlation is negative. We will call the setting where $a > 0$ positive accuracy on the line and $a < 0$ accuracy on the inverse line. Next, we show that the smaller the ϵ, the smaller the probability of the ID/OOD split being well-specified. The probability of a well-specified ID/OOD split when $\epsilon = 0$ is also 0.

Theorem 3 (Benchmarks with Accuracy on the Line are Misspecified Almost Everywhere.)**.** *Define*

$$\mathcal{W}_\epsilon = \left\{ M \in \mathbb{R}^{l \times l} : \begin{array}{l} w_{spu}^\top (M \, \mu_{spu}) + \sqrt{2 \, (L_\phi \, \kappa)^2 \, \Sigma_{spu} \, \log(1/\delta)} < 0 \quad \textit{(Theorem 1)}, \\ \left|\Phi^{-1}\big(acc_P(f_X)\big) - a \, \Phi^{-1}\big(acc_{P_\phi}(f_X)\big)\right| \leq \epsilon \quad (\epsilon \geq 0) \end{array} \right\} \tag{12}$$

Then:

(i) $\mathcal{W}_0$ has Lebesgue measure zero in $\mathbb{R}^{l \times l}$.

(ii) For any $0 \leq \epsilon_i \leq \epsilon_j$, we have $\mathcal{W}_{\epsilon_i} \subseteq \mathcal{W}_{\epsilon_j}$.

The proof of Theorem 3 and supporting Lemmas provided in Appendix A.6.

The two conditions in Equation 12 are at odds. In particular, as $\epsilon \to 0$ (i.e., perfect accuracy on the line), almost every shift is misspecified, and the Lebesgue measure of the set of well-specified shifts grows monotonically with ϵ, i.e., inversely with accuracy on the line. This means that when we observe accuracy on the line, with a high probability, the ID/OOD split is misspecified. In Appendix A.7, we construct an intuitive example of such (zero-measure) shifts, where both the conditions for well-specified splits and accuracy on the line hold.

Theoretical Results Summary. Overall, Theorem 1 gives conditions for well-specified domain generalization ID/OOD splits, and Theorem 3 demonstrates that there is zero probability that such conditions and *accuracy on the line* for the ID/OOD split simultaneously hold—Figure 2. Moreover, accuracy on the line is at odds with well-specified shifts. Our results suggest that datasets with accuracy on the line may be misspecified for benchmarking domain generalization. In contrast, benchmarks with *accuracy on the inverse line*, or weak correlation between ID and OOD accuracy, are better suited to benchmark domain generalization. This property now gives a test for well-specified benchmarks. Later (Section 4), we apply this test to state-of-the-art domain generalization benchmarks.

3 Related Work

Two influential domain generalization benchmark suites are DomainBed (Gulrajani & Lopez-Paz, 2020a) and WILDS (Koh et al., 2021). **DomainBed** is a collection of object recognition domain generalization benchmarks. For example, PACS (Khosla et al., 2012; Li et al., 2017) includes images of seven classes across four domains: Photos, Art Paintings, Cartoons, and Sketches. Another benchmark is ColoredMNIST (Arjovsky et al., 2019), a semi-synthetic binary classification variation of MNIST (Deng, 2012), which introduces color as a spurious correlation and defines domains by specific color-label associations. Gulrajani & Lopez-Paz (2020a) found that empirical risk minimization achieved the best transfer performance compared to state-of-the-art

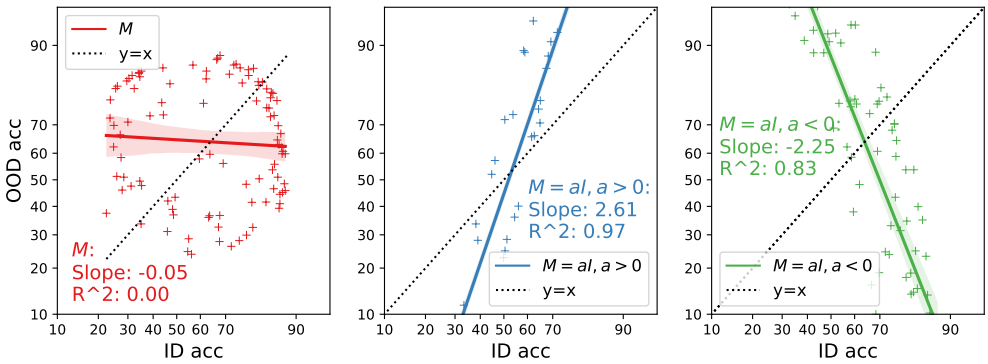


Figure 2: ID vs. OOD accuracy on probit scale. When M satisfies Theorem 1's conditions, the accuracy on the line phenomenon does not occur. For $M_{\text{ID}} = I$ and $M_{\text{OOD}} = aI$, where a is allowed to vary, we observe accuracy on the line. When $a < 0$, we have the spurious correlation reversal condition and have accuracy on the inverse line, where these is a strong but negative correlation between in- and out-of-distribution accuracy. Notably, ID/OOD splits with accuracy on the inverse line are well-specified. More experimental details can be found in Appendix B.

domain generalization algorithms across PACS, ColoredMNIST, and other DomainBed benchmarks. They also observed differences in their findings based on their choice of model selection.

WILDS was designed to better represent real-world shifts across vision and language. For example, Camelyon17 (Zech et al., 2018; AlBadawy et al., 2018) includes images of tissue that may contain tumor tissue (classes) from different hospitals with varying conditions (domains). CivilComments Borkan et al. (2019) includes comments to an online article that may be toxic (class) for different subpopulations defined by demographic identities (domains). These benchmarks illustrate the suite's focus on practical and natural real-world applications. However, even across WILDS benchmarks, no state-of-the-art methods have demonstrated consistent superiority over ERM (Koh et al., 2021).

For **Subpopulation shift**, which we consider a special case of the broader domain generalization task, benchmarks are designed specifically to evaluate distribution shift robustness in scenarios where spurious correlations lead to worse performance on underrepresented subgroups out-of-distribution. For example, the Waterbirds benchmark (Sagawa et al., 2019) introduces a spurious correlation between bird species and backgrounds, such as waterbirds predominantly appearing in water environments. Models that rely on backgrounds rather than bird features when predicting bird type (classes) generalize poorly to new domains where backgrounds are urban areas—birds in urban backgrounds are undersampled in the training data.

Benchmarks addressing **other types of distribution shifts** where domain generalization is desired have also been proposed, e.g., temporal shifts (Yao et al., 2022; Joshi et al., 2023; Zhang et al., 2023).

Other conceptualizations of distribution shift datasets have been extended to benchmark domain generalization. However, not all are designed to evaluate robustness to spurious correlations, nor do they represent multi-source domain generalization settings. For instance, some benchmarks are designed to assess robustness against typical data quality variations encountered in real-world settings, like variations of ImageNet (Deng et al., 2009; Hendrycks & Dietterich, 2019; Hendrycks et al., 2021). Such benchmarks are out of scope for our study as they are not standard benchmarks for the domain generalization tasks we study. Nevertheless, we refer to previous work studying accuracy on the line for these datasets (Taori et al., 2020).

3.1 Accuracy on the Line

The accuracy on the line phenomena has been observed empirically in previous work for many benchmarks in these suites (Recht et al., 2019; Miller et al., 2021; 2020; Taori et al., 2020; Baek et al., 2022; Saxena et al., 2024). However, Liu et al. (2023a) identify real-world tabular datasets with weak or negative linear correlation, and Teney et al. (2023) identify non-tabular real-world datasets where ID and OOD performance

exhibit other patterns between in- and out-of-distribution accuracy beyond strongly positive and linear. Notably, they also found that correlations can be inverted in some datasets. Sanyal et al. (2024) derive noise conditions to achieve shifts with accuracy on the inverse line. Our work uniquely specifies the implications of accuracy on the line (or lack thereof) on the utility of datasets as domain generalization benchmarks.

Other works have studied conditions where the empirical risk minimizer for observed distributions is sufficient for domain generalization. Rosenfeld et al. (2022b) show this to be the case under domain interpolation, i.e., OOD distributions are convex combinations of observed distributions (in an online setting). Additionally, when domain-general features are fully informative, i.e., spurious correlations are redundant, and under some conditions, Ahuja et al. (2021) also show this to be the case. Our results corroborate their findings.

This notion of worst-case stress testing to establish robustness under distribution shift is not new. When evaluating the out-of-distribution generalization of deep learning methods developed on the ImageNet task (Deng et al., 2009) (*ImageNet Large Scale Visual Recognition Challenge (ILSVRC)* (Russakovsky et al., 2015)), Kornblith et al. (2019) assess generalization by applying said methods to contemporary datasets with presumably different distributions than ImageNet, such as CIFAR-10 (Krizhevsky, 2009). However, the datasets they investigated could be considered quite similar in distribution to ImageNet. Alternatively, Salaudeen & Hardt (2024) adversarially constructed a dataset, ImageNot, designed to shift spurious correlations present in the original ImageNet construction strongly. Notably, the findings of Salaudeen & Koyejo (2022) align with those of Kornblith et al. (2019) despite the deliberately adversarial construction of ImageNot. We argue that constructing datasets with such adversarial spurious correlation shifts is essential for rigorously probing a model's use of spurious correlations.

More generally, other works have proposed alternative approaches to developing domain generalization benchmarks. Satisfying our conditions introduces an additional necessary dimension for creating more meaningful evaluations. For example, Zhang et al. (2023) argues that many existing benchmarks are limited by having too few domains and overly simplistic settings, which restrict their ability to simulate the significant distribution shifts observed in real-world scenarios. Similarly, Lynch et al. (2023) contend that benchmarks inadequately capture the complex, many-to-many spurious correlations that can arise in practical applications.

Like previous work, we next investigate the accuracy of the line properties of state-of-the-art domain generalization benchmarks to take an inventory of well-specified datasets.

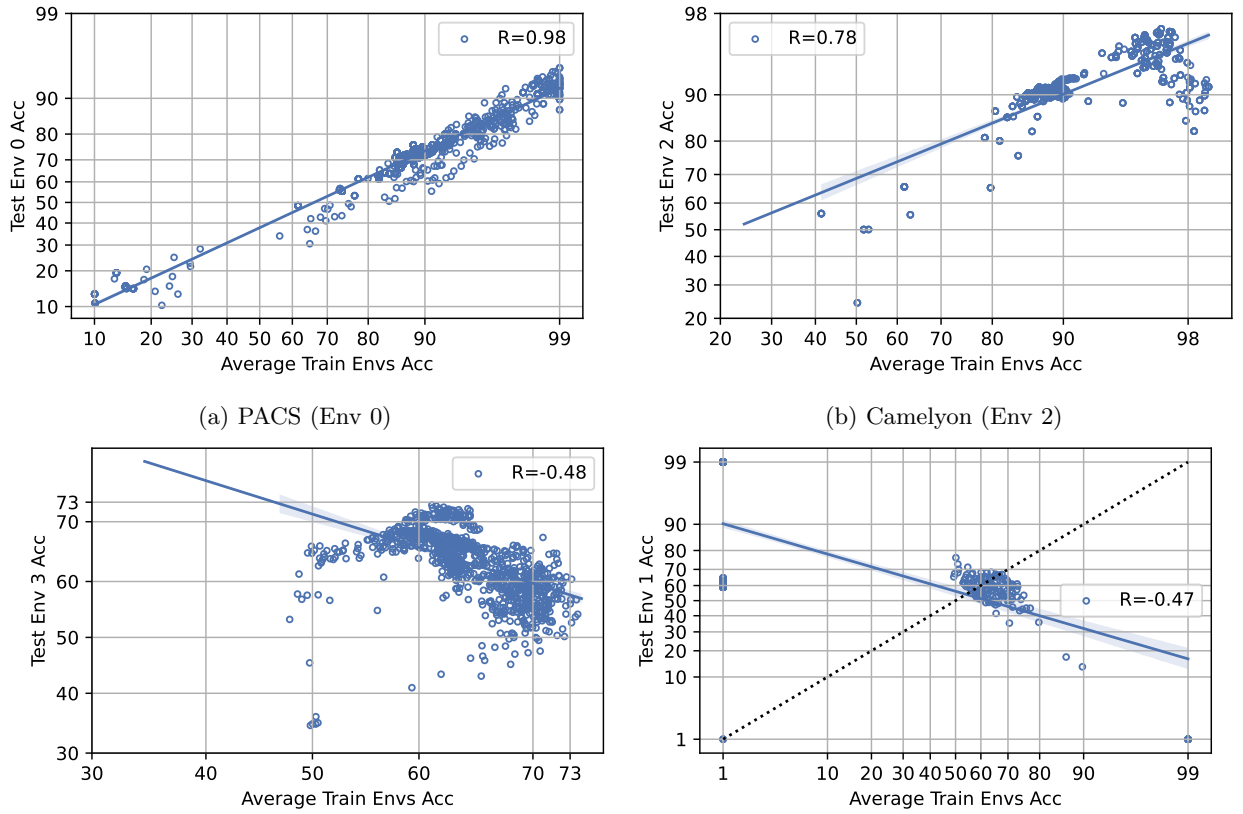
4 Empirical Results

We evaluate the correlation between in-domain (ID) and out-of-domain (OOD) accuracy for benchmarks in the popular DomainBed (Gulrajani & Lopez-Paz, 2020a) and WILDS (Koh et al., 2021) benchmark suites, as well as subpopulation shift benchmarks, e.g., WaterBirds (Sagawa et al., 2019).

Datasets. Specifically, our results include the following datasets: **Camelyon** (Bandi et al., 2018; Koh et al., 2021), **CivilComments** (Borkan et al., 2019; Koh et al., 2021), **ColoredMNIST** (Arjovsky et al., 2019; Gulrajani & Lopez-Paz, 2020a), **Covid-CXR** (Alzate-Grisales et al., 2022; Cohen et al., 2020b; Tabik et al., 2020; Tahir et al., 2021; Suwalska et al., 2023), **FMoW** (Christie et al., 2018; Koh et al., 2021), **PACS** (Li et al., 2017; Gulrajani & Lopez-Paz, 2020a), **Spawrious** (Lynch et al., 2023), **TerraIncognita** (Beery et al., 2018; Gulrajani & Lopez-Paz, 2020a), and **Waterbirds** (Sagawa et al., 2019).

Model Architectures. For vision datasets, we leverage pretrained deep learning architectures such as **ResNet-18/50** (He et al., 2016), **DenseNet-121** (Huang et al., 2017), **Vision Transformers** (Dosovitskiy et al., 2020), and **ConvNeXt-Tiny** (Liu et al., 2022). For language datasets, we utilize pretrained embeddings from **BERT** Jacob et al. (2019) and **DistilBERT** Sanh et al. (2020), and apply lower-capacity machine learning models, such as logistic regression, for downstream classification tasks.

Experimental Setup. The benchmarks we consider include a set of domains—distinct data distributions. As standard in the literature (Gulrajani & Lopez-Paz, 2020a; Koh et al., 2021), the in-distribution (ID) data is defined as a mixture of a subset of the data domains, and the out-of-distribution (OOD) data is not included in the training domains, i.e., we perform a leave-one-domain-out ID/OOD splits for our experiments, where we train on all but one domain and use the left-out domain as OOD. We generate models for our



(a) PACS (Env 0)

(b) Camelyon (Env 2)

(c) Covid-CXR (Env 3)

(d) CivilComments (Env 1).

Figure 3: We show some ID/OOD splits of popular domain-generalization benchmarks with a strong positive, weak, or strong negative correlation between in-distribution and out-of-distribution accuracy. Our results suggest that algorithms that consistently provide models with the best transfer accuracies for these splits are at least partially successful in removing spurious correlations. For Camelyon (b), within some accuracy range, we have accuracy on the inverse line, indicating the importance of a qualitative assessment of these trends.

experiments by varying model and training hyperparameters for each architecture, including the number of training epochs (Appendix C Table 1). Our experiments include training these models end-to-end with varying hyperparameters and data augmentations, as well as pretraining and transfer learning.

Table 3 highlights the prevalence of widely-used domain generalization benchmarks with accuracy on the line, a signature of potential misspecification, while Figure 3 qualitatively illustrates benchmarks with weak or strongly negative correlation between in and out-of-distribution accuracy. We provide a detailed account of our experiments, along with benchmark-specific discussions, in Appendix C.

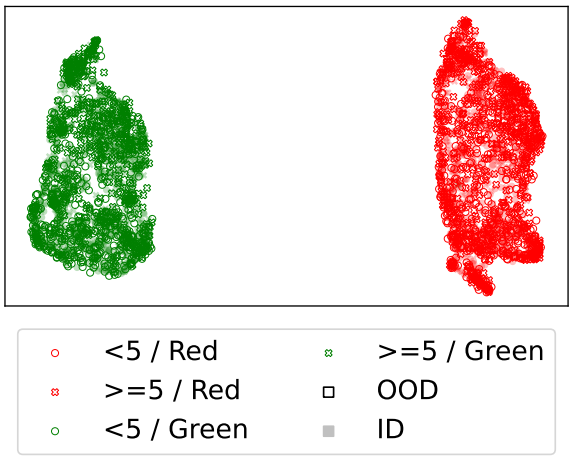
Selecting Number of Models. To ensure a robust estimate of ID-OOD correlation, we follow and extend the approach of prior work by training a diverse set of models that vary in architecture, random seed, data order, and hyperparameters. This diversity mitigates the risk of single-model artifacts influencing our findings. We adopt a simple heuristic: continue adding models until the ID-OOD Pearson correlation changes by less than 1%. As shown in Table 2, this threshold is consistently reached well before exhausting our pool of trained models. In practice, we train 2–10× more models than previous studies, often exceeding thousands per dataset and ensuring our correlation estimates are stable and conservative. For instance, we train over 10,000 models for a single CivilComments split, and our criterion is satisfied with less than 7,000 models. We train over 5,000 models for a PACS environment, and our criterion is satisfied with 910. We trained as many models as computationally feasible to ensure the robustness of our findings.

4.1 Findings

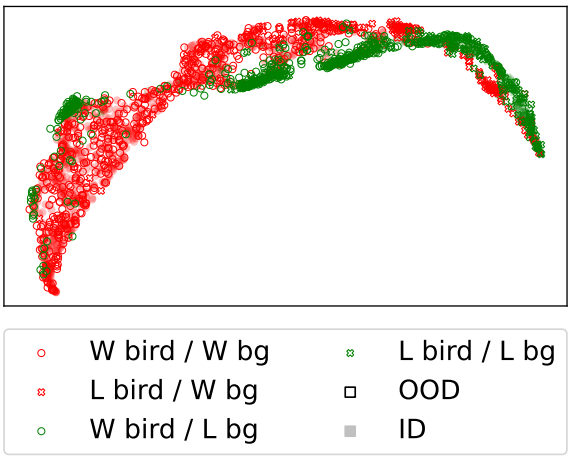
We find that, primarily, semisynthetic datasets satisfy our derived conditions. Particularly, semisynthetic here means real-world datasets that either (i) have artificial spurious correlations introduced (ColoredMNIST, Spawrious, and Waterbirds) and (ii) have some selection process that introduces spurious correlations (CivilComments). The CXR datasets exhibit a relationship that Teney et al. (2024) refers to as 'Not Transfer,' where the OOD accuracy is near constant despite high variance in the ID accuracy. Our results further echo Teney et al. (2024)'s notes on potentially misleading advice from past studies—particularly the focus on improving ID performance to improve OOD robustness (Wenzel et al., 2022), which our work suggests is only a reasonable strategy in settings where distribution shifts are relatively simple and weak.

Our results and findings should not be surprising, given that spurious correlations we aim to mitigate for decision-making tend not to be localized (Teresa-Morales et al., 2022). For instance, correlations between gender and occupation likely persist across naturally collected datasets. Hence, they may not harm performance across different data sources. We find that the subpopulation shift benchmarks we assess often have the desired properties derived in this work. Notably, these benchmarks are (i) constructed such the spurious correlation from training is no longer accurate at testing and (ii) evaluated based on worst-group (worst-case) performance. Our work shows that in less concisely defined domain generalization contexts, such types of shifts are also necessary. We provide detailed results and discussion in Appendix C.

4.2 Discussion



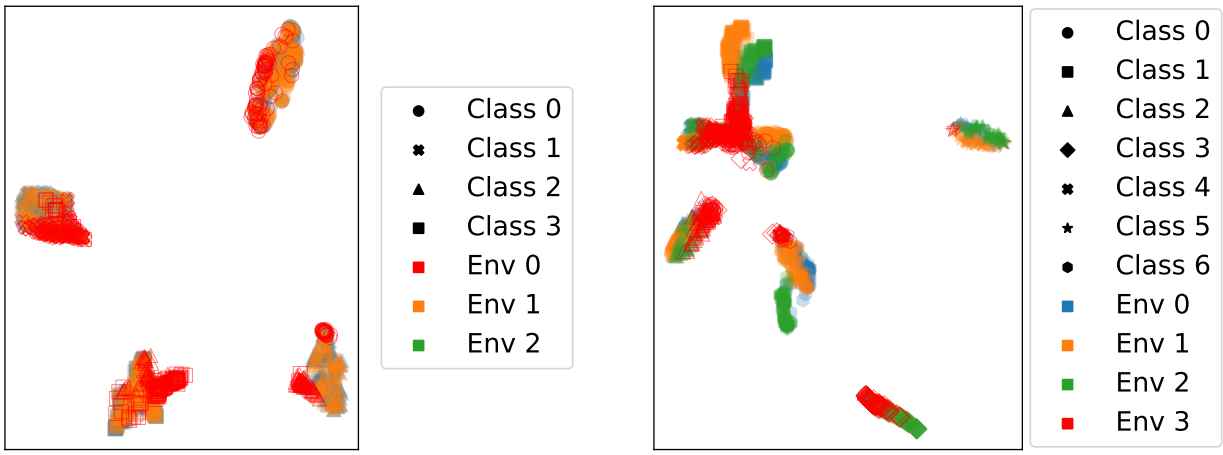
(a) ColoredMNIST. Embeddings of a model trained on Env 1 and 2. The labels are < 5, ≥ 5 and the spurious feature is color (red, green). The UMAP of the embeddings clusters based on color (spurious) rather than digit features (domain-general). This leads to incorrect predictions OOD where the color-label correlation reverses, as observed in our experiments (Section C.2).



(b) Waterbirds. Embeddings trained examples with spurious correlation between bird type and background. The bird type is the label, and the background is the spurious feature. The UMAP of the embeddings clusters based on background type (spurious) more strongly than bird features (domain-general). This leads to incorrect predictions OOD where background-label correlation changes, as observed in our experiments (Section C.9).

Figure 4: We use a UMAP to visualize the embedding clusters for examples. We fit the UMAP to held-out examples of the in-distribution data and also apply it to the out-of-distribution data.

Domain generalization considers worst-case shifts (Arjovsky et al., 2019; Rosenfeld et al., 2022b). While worst-case shifts may have a trade-off with average utility (Salaudeen & Koyejo, 2024; Miller, 2024), it is crucial in fairness-sensitive and high-stakes applications like healthcare (Angwin et al., 2016; Chen et al., 2019; Oakden-Rayner et al., 2020). However, domain generalization benchmarks have not been rigorously assessed for their utility in such scenarios. We address this gap by deriving conditions for well-specified benchmarks, helping practitioners align benchmark choices with their goals. We showed that well-specified



(a) Spawrious O2O Hard. Embeddings of a model trained on Env 1 and 2. The label is animal type ,and the spurious feature is background. Generally, for some classes (e.g., Class 0), some subset of OOD (Env 0) examples cluster very closely to the same classes in in-distribution data. However, for the same class, many OOD examples incorrectly cluster more closely to class 2; there is also smaller variance in these examples. We observe this for most classes, demonstrating that the model is relying on spurious correlations. This matches our observations of OOD performance degradation due to spurious correlations (Section C.3.2)

(b) PACS. Embeddings of a model trained on Env 0-2. The spurious feature is unknown, but it is purported to exist because each environment has a different image style. We find that there is a relatively strong alignment between the in-distribution and out-of-distribution embeddings for each class. This matches our finding that there is not a strong spurious correlation that would cause models to embed examples based on features that have an unstable relationship between labels (Section C.4), like we see in the other examples we analyzed with UMAP.

Figure 5: We use a UMAP to visualize the embedding clusters for examples. We fit the UMAP to held-out examples of the in-distribution data and also apply it to the out-of-distribution data.

benchmarks, i.e., are aligned with worst-case shifts, will not have accuracy on the line. Hence, benchmark users and curators with a goal of addressing such shifts should seek out benchmarks with weak or negative ID and OOD accuracy correlation.

Many benchmark curators define their intended scope carefully. For example, WILDS (Koh et al., 2021) focuses on real-world rather than worst-case shifts. Our findings often validate these intended scopes, yet benchmark users may not always adhere to them. To clarify benchmark suitability, we categorize benchmarks into (i) worst-case and (ii) natural shifts (Koh et al., 2021; Taori et al., 2020), emphasizing that worst-case benchmarks are particularly valuable for auditing uninterpretable predictors, such as detecting demographic biases in model decisions (Ferrer et al., 2021), whereas natural shifts may suffice when prioritizing average OOD performance. Our conditions enable assessing whether spurious correlations (e.g., race in Chest X-Rays (Gichoya et al., 2022)) impact predictions. For robust algorithm development, both worst-case and natural shifts should be considered to ensure broad applicability. Additionally, new spurious correlation benchmarks should undergo *accuracy on the line* evaluations to support their reliability—ideally, there is no positive accuracy on the line. Finally, domain expertise is critical in defining spurious correlations: a narrow potential distribution set may limit robustness, while an overly broad definition of potential distributions can unnecessarily reduce utility (Shen et al., 2024). For instance, Chiou et al. (2024) report that training a model on multiple recording sessions degrades OOD (new session) brain-computer interface (BCI) classification performance compared to training on single-session data.

Constructing Benchmarks Without Positive Accuracy on the Line. Indeed, some of the datasets we study empirically that satisfy our desired conditions are primarily semisynthetic (e.g., ColoredMNIST, Spawrious, Waterbirds). In contrast, datasets such as Covid-CXR and WILDSCamelyon, have in-distribution (ID) and out-of-distribution (OOD) splits that do not exhibit positive accuracy on the line. This suggests

that careful and intentional data collection and curation is likely necessary to obtain naturally occurring datasets without positive accuracy on the line.

One approach to constructing such datasets is to identify settings where a natural experiment or intervention has occurred. For example, the Covid-CXR dataset leverages the natural intervention of the pandemic and regional variations. Without such interventions, there may be little reason to expect that spurious correlations would fail to generalize in the real-world datasets that are most readily available.

Before finalizing a benchmark, we should measure the correlation between ID and OOD performance to ensure that the dataset does not exhibit strong positive accuracy on the line. This validation can help confirm that the benchmark is well-specified and truly tests robustness to spurious correlations.

One approach to constructing such datasets is to identify settings where a natural experiment or intervention has occurred. For example, the Covid-CXR dataset leverages the natural intervention of the pandemic and regional variations. Without such interventions, there may be little reason to expect that spurious correlations would fail to generalize in the datasets that are most readily available.

Qualitatively Assessing Accuracy on the Line. For the Camelyon dataset (Section C.6), we observe a shift in correlation patterns based on model accuracy. Specifically, models with greater than 90% accuracy exhibit a negative correlation between ID and OOD performance, whereas models below this threshold show a positive correlation. Since accuracy on the line (Definition 2) is a global property, this dataset does not meet the criteria for accuracy on the line since there is a strong deviation from the positive correlation in some regimes. This underscores a key limitation: evaluating overall correlation may not sufficiently identify well-specified benchmarks. While this approach is robust against false negatives of misspecification, it may introduce false positives, necessitating qualitative inspection as a complementary assessment tool.

On the Slope vs. the Correlation Coefficient. In this work, we focus on the correlation between ID and OOD accuracy rather than the slope between the two quantities. Recall that for a set of pairs x, y, which have a Pearson R correlation of r, the slope when y is regressed on x is $s = r * (\sigma_y/\sigma_x)$. s depends on variances σ_y and σ_x, which may or may not be relevant to the spurious correlation problem. The slope depends strongly on the definition of $\mathcal{F}$ (Appendix C). This observation is also related to limitations in attributing accuracy drop across distribution only to distributional systematic bias (Salaudeen & Hardt, 2024). While s is informative, its relationship with our derived conditions is not as obvious.

Reconciling Worst-Case and Average-Case Generalization. A strict worst-case approach to generalization can conflict with developing locally (sites) beneficial models, particularly in healthcare (Futoma et al., 2020; Miller, 2024). However, failures even within the same site suggest this tension is unavoidable (Oakden-Rayner et al., 2020), as spurious correlations remain brittle due to other non-local factors like temporal drifts (Ji et al., 2023) or interventions (Birkmeyer et al., 2020). Reliable worst-case robustness benchmarks remain essential, even with a narrower scope. Still, when worst-case focus severely impacts average utility, additional evaluation on alternative benchmarks is warranted. A practical approach is narrowing the model's deployment scope to a smaller set of distributions, which can improve worst-case performance (e.g., when the shifts in the smaller set are weaker) without sacrificing as much average utility. However, practically, maintaining reliable predictions may require scope-dependent model monitoring and updates.

Implications on Key Domain Generalization and Evaluation Practices. ID/OOD splits within the same dataset can exhibit varying Pearson R correlations, meaning some splits provide more reliable benchmarks than others. However, **averaging** over all ID/OOD splits (Gulrajani & Lopez-Paz, 2020a) dilutes this reliability, particularly when only one split is well-specified. The issue worsens when averaging across multiple datasets to compare domain generalization methods (Gulrajani & Lopez-Paz, 2020a). In subpopulation shifts, **worst-group accuracy** is the standard evaluation metric (Koh et al., 2021); adopting a similar norm more generally for domain generalization improves the robustness of evaluation.

A related issue arises in **model selection** via cross-validation, where selecting models based on held-out accuracy—whether IID or a held-out domain—can lead to overfitting to spurious correlations specific to that set. Alternative selection criteria are implied conditional independencies (Salaudeen & Koyejo,

2024), cross-risk minimization (Pezeshki et al., 2023), and confidence-based ensemble aggregation (Chen et al., 2023a). However, model selection under distribution shifts remains a challenge.

Furthermore, our test can act as a pre-check to decide whether algorithmic invariance penalties are worth tuning. However, if there are spurious correlations between a set of observed environments, for any single test environment, it may be the case that models that ignore these spurious correlations perform worse than a model that utilizes them. However, there is a tradeoff between this and other environments where the same spurious correlations lead to failure. This tradeoff is implicit and domain-specific. Analysis of this tradeoff is out of the scope of this work.

Implications on Benchmarking Causal Representation Learning which aims to uncover underlying causal structures. One approach to this is *independent causal mechanisms* (Pearl, 2009; Schölkopf et al., 2021), which has motivated many domain generalization algorithms (Arjovsky et al., 2019; Peters et al., 2016). Since both tasks require distinguishing stable from spurious correlations, our results on evaluating domain generalization also apply to benchmarking causal representation learning. Specifically, when assessing models—including disentangled causal models—based on OOD accuracy, our framework helps determine when domain generalization reliably reflects success in learning causal representations (Salaudeen et al., 2024)—we discuss this further in Appendix D.

Implications on Benchmarking Algorithmic fairness which aims to mitigate biases that cause disparate performance across demographic groups; some definitions of fairness are closely linked to domain generalization (Creager et al., 2021). Group sufficiency particularly aligns with the principle of invariance (Chouldechova, 2017; Liu et al., 2019). We emphasize a straightforward but key insight: when using OOD accuracy to benchmark whether models avoid relying on group information, the benchmark must ensure that group information hinders out-of-distribution performance. In this case, a strong positive correlation between training and worst-group test accuracy suggests that group information generalizes. In contrast, a weak or negative correlation between ID and OOD accuracy is preferable.

Modern **foundation models** are susceptible to spurious correlations (Alabdulmohsin et al., 2024; Zhu et al., 2023; Gerych et al., 2024; Hamidieh et al., 2024). Analyzing the Civil Comments dataset, we find that spurious correlation shifts in language datasets exhibit similar patterns to vision datasets within our framework, showing strong positive, negative, and weak correlations between ID and OOD accuracy. Furthermore, Saxena et al. (2024) report strong positive correlations in large language models for predictive tasks (Q/A) under distribution shift. Our results highlight that the benchmark conditions we establish are also crucial for evaluating spurious correlations in foundation models.

Embeddings Analysis. UMAP (Uniform Manifold Approximation and Projection) is a fast, scalable dimensionality reduction technique that preserves both local and global structure of data (McInnes et al., 2018). Unlike t-SNE (Van der Maaten & Hinton, 2008), UMAP can generalize to new data via a learned embedding function, making it especially useful for visualizing high-dimensional datasets and comparing in-distribution versus out-of-distribution representations. Figures 4-Figures 5 demonstrate the concordance between our findings and the embeddings learned by models. For instance, for ColoredMNIST, Waterbirds, and Spawrious, embeddings of classes are consistent in the training distribution, however, out-of-distribution, the same classes' embeddings are aligned with the wrong class from the in-distribution data, and would likely be labeled incorrectly. This is because the models have learned spurious correlations that change from in-distribution to out-of-distribution. We find that these datasets also have a lower (negative) correlation between in- and out-of-distribution accuracy. For PACS however (Figure 5b), the alignment between in- and out-of-distribution embeddings are consistent, meaning that the same classes in- and out-of-distribution are closely embedded. Our analysis also finds that PACS has a strong correlation between in- and out-of-distribution accuracy. Overall, our qualitative analysis of the embeddings leared by in-distribution optimal models supports our theoretical and empirical findings.

State-of-the-Art Algorithms. Many algorithms have been proposed for domain generalization Gulrajani & Lopez-Paz (2020a); Koh et al. (2021); Yang et al. (2023). On datasets we identify as misspecified—such as PACS, Terra Incognita, WILDSCamelyon, and WILDFMoW—Empirical Risk Minimization (ERM) performs comparably to state-of-the-art domain generalization algorithms (Gulrajani & Lopez-Paz, 2020a; Koh et al.,

2021), particularly when controlling for factors like the use of unconstrained, large-scale pretrained models (e.g., CLIP (Radford et al., 2021), trained on internet-scale data rather than ImageNet). Specific results for WILDS datasets are available at https://wilds.stanford.edu/leaderboard/, maintained by the WILDS team.

In contrast, on datasets we identify as well-specified, several algorithms have consistently outperformed ERM, even under independent validation. For example,(Yang et al., 2023) demonstrates improvements over ERM on Waterbirds and CivilComments.

4.3 Limitations and Future Work

While we have developed probabilistic sufficient conditions for well-specified benchmarks, we derive necessary and sufficient conditions only under the Gaussianity assumption of features. Future work includes exploring whether these conditions are necessary for non-Gaussian features. Demonstrating this may require developing analytical techniques distinct from those applied in this work. Additionally, Yang et al. (2023) demonstrates that while accuracy on the line may hold, other metrics or a combination of metrics may not simultaneously have the same strong positive linear trend. Investigating metrics other than accuracy is left for future work.

Another direction for future work is to develop a more robust automated method for assessing accuracy on the line, beyond simply computing correlation across all training accuracy levels. Currently, qualitative assessment remains necessary to account for cases where models with higher training accuracy exhibit negative accuracy on the line. Empirically, we found that standard change-point detection methods (Killick et al., 2012) are highly sensitive to noise in accuracy measurements. However, incorporating more robust heuristics could improve their reliability, making them a viable approach for automation.

Additionally, while we have evaluated a wide variety of models in this work, continuing to collect data points of ID/OOD accuracies for benchmarks improves the accuracy of the true relationship. Finally, curating additional benchmarks without accuracy on the line, drawn from diverse, real-world scenarios with high-dimensional spurious features, is left for future work. This work, along with others (Recht et al., 2019; Taori et al., 2020; Miller et al., 2021), has characterized this property for a variety of popular benchmarks.

5 Conclusion

Robustness to spurious correlations under worst-case distribution shifts is a critical challenge in machine learning, essential for ensuring the reliability and fairness of models. In this work, we identify significant limitations in current benchmarks designed to address this problem. Specifically, many state-of-the-art benchmarks, which evaluate OOD accuracy by training models on an in-distribution (ID) split and testing on an out-of-distribution (OOD) split, fail to guarantee that models free of spurious correlations will transfer better. We define a benchmark as *well-specified* if such a guarantee exists.

Previous work observed that many benchmarks exhibit the phenomenon of *accuracy on the line*, where improved ID performance directly correlates with improved out-of-distribution performance. Our theoretical findings suggest that this behavior indicates that such benchmarks are misspecified for evaluating domain generalization and emphasize the importance of prioritizing benchmarks that do not exhibit accuracy on the line when addressing worst-case distribution shifts. We aim to provide a clearer path toward developing models robust to spurious correlations by addressing the evaluation ambiguity.

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A Proofs

A.1 Proof of Lemma 1—Domain-Specific Models have Lower In-Domain Error under Partially Informative Domain-General Features

Assume Non-trivial and non-redundant features (Assumption 1-2), and strongly convex ℓ.

$$\min_{f\in\mathcal{F}} \mathbb{E}_{(X,Y)\sim P}\big[\ell(f(X),Y)\big] < \min_{f\in\mathcal{F}_{\mathrm{dg}}} \mathbb{E}_{(X,Y)\sim P}\big[\ell(f(X),Y)\big], \tag{13}$$

where $\mathcal{F}:\mathcal{X}\to\mathbb{R}$ where $f(x)=\mathrm{w}^\top x=\mathrm{w}_{\mathrm{dg}}^\top z_{\mathrm{dg}}+\mathrm{w}_{\mathrm{spu}}^\top z_{\mathrm{spu}}$, $f\in\mathcal{F}$. For $f\in\mathcal{F}_{\mathrm{dg}}$, $f(x)=(\mathrm{w})^\top x=\mathrm{w}_{\mathrm{dg}}^\top z_{\mathrm{dg}}$.

Proof. From Assumption 1-2, non-trivial and non-redundant features, a model that uses both domain-general and spurious features is more expressive than one that does not. Let w be the Bayes optional By the Bayes optimality of w^*, any w achieving the same risk agrees with w^* almost everywhere, i.e.,

$$\mu\left(\left\{x \in \mathcal{X} \mid \mathrm{w}^{*\top} x \neq \mathrm{w}^\top x\right\}\right) = 0,$$

where μ denotes the Lebesgue measure on $\mathcal{X}$.

Let $x = z_{\mathrm{dg}} \oplus z_{\mathrm{spu}}$ and $\mathrm{w} = \mathrm{w}_{\mathrm{dg}} \oplus \mathrm{w}_{\mathrm{spu}}$ such that

$$\mathrm{w}^{*\top} x = \mathrm{w}_{\mathrm{dg}}^\top z_{\mathrm{dg}} + \mathrm{w}_{\mathrm{spu}}^\top z_{\mathrm{spu}}.$$

If we only consider values of x where $\mathrm{w}_{\mathrm{dg}}^\top z_{\mathrm{dg}} \neq 0$ and $\mathrm{w}_{\mathrm{spu}}^\top z_{\mathrm{spu}} \neq 0$, then without loss of generality we have that

$$\mathrm{w}^{*\top} x = \mathrm{w}_{\mathrm{dg}}^\top z_{\mathrm{dg}} + \mathrm{w}_{\mathrm{spu}}^\top z_{\mathrm{spu}} \neq \mathrm{w}_{\mathrm{dg}}^\top z_{\mathrm{dg}}.$$

Given Assumption 1-2,

$$\mu\left(\left\{x \mid \mathrm{w}_{\mathrm{dg}}^\top z_{\mathrm{dg}} \neq 0\right\}\right) > 0,$$

the risk of $\mathrm{w}_{\mathrm{dg}}^\top z_{\mathrm{dg}}$ is strictly greater than that of $\mathrm{w}^{*\top} x$. Equation 13 follows from the strong convexity of the loss. □

A.2 Proof of Theorem 1—Sufficient Conditions for Well-Specified Domain Generalization Benchmark Splits

Assume $Z_{\mathrm{spu}}^{\mathrm{ID}}$ is sub-Gaussian with mean μ_{spu}, covariance Σ_{spu}, and parameter κ. Define a nonlinear transformation

$$\phi : \mathbb{R}^l \to \mathbb{R}^l,$$

that is L_ϕ–Lipschitz, and let

$$Z_{\mathrm{spu}}^{\mathrm{OOD}} = \phi(Z_{\mathrm{spu}}^{\mathrm{ID}}).$$

Assume further that

$$\mathbb{E}[Z_{\mathrm{spu}}^{\mathrm{OOD}}] = \mathbb{E}[\phi(Z_{\mathrm{spu}}^{\mathrm{ID}})] = M\,\mu_{\mathrm{spu}},$$

for some matrix $M \in \mathbb{R}^{l \times l}$. In-distribution, $Z_{\mathrm{spu}}^{\mathrm{ID}} \sim P_{\mathrm{ID}}$ and out-of-distribution, $Z_{\mathrm{spu}}^{\mathrm{OOD}} \sim P_{\mathrm{OOD}}$. Additionally, denote $\mathrm{w}_{\mathrm{spu}}$ the contribution of $Z_{\mathrm{spu}}^{\mathrm{ID}}$ to the optimal P_{ID} predictor $f_{\mathrm{X}}^{P_{\mathrm{ID}}}$. Then, for any $\delta \in (0, 1)$, if

$$\mathrm{w}_{\mathrm{spu}}^\top (M\,\mu_{\mathrm{spu}}) + \sqrt{2\,(L_\phi\,\kappa)^2\,\|\mathrm{w}_{\mathrm{spu}}\|_2^2\,\log(1/\delta)} < 0,$$

(with the understanding that under the Lipschitz assumption the sub-Gaussian property carries over with parameter $L_\phi\,\kappa$), then with probability at least $1 - \delta$ over $Z_{\mathrm{spu}}^{\mathrm{OOD}}$, we have

$$\mathrm{acc}_{P_{\mathrm{OOD}}}(f_{\mathrm{X}}^{P_{\mathrm{ID}}}) < \mathrm{acc}_{P_{\mathrm{OOD}}}(f_{\mathrm{dg}}^{\mathcal{E}}),$$

where $f_{\mathrm{dg}}^{\mathcal{E}}$ and $f_{\mathrm{X}}^{P_{\mathrm{ID}}}$ are the optimal domain–general and domain–specific predictions (Definitions 3–4).

Proof. Define

$$Z_{\mathrm{spu}}^{\mathrm{OOD}} = \phi(Z_{\mathrm{spu}}^{\mathrm{ID}}).$$

From Equation 1 and the law of total probability, the out-of-distribution (OOD) accuracy of $f_{\mathrm{X}}^{P_{\mathrm{ID}}}$ is equivalently

$$\mathrm{acc}_{\mathrm{OOD}}(f_{\mathrm{X}}^{P_{\mathrm{ID}}}) = \Pr\left(\mathrm{w}_{\mathrm{dg}}^\top Z_{\mathrm{dg}} + \mathrm{w}_{\mathrm{spu}}^\top Z_{\mathrm{spu}} > 0\right),$$

and

$$\mathrm{acc}_{\mathrm{OOD}}(f_{\mathrm{dg}}^{\mathcal{E}}) = \Pr\Big(\mathrm{w}_{\mathrm{dg}}^\top Z_{\mathrm{dg}} > 0\Big).$$

It suffices to show that

$$\Pr\Big(\mathrm{w}_{\mathrm{dg}}^{\top} Z_{\mathrm{dg}} > -\mathrm{w}_{\mathrm{spu}}^{\top} Z_{\mathrm{spu}}^{\mathrm{OOD}}\Big) < \Pr\Big(\mathrm{w}_{\mathrm{dg}}^{\top} Z_{\mathrm{dg}} > 0\Big) \tag{14}$$

with high probability.

Since $Z_{\mathrm{spu}}^{\mathrm{ID}}$ is sub-Gaussian with parameter κ, by the Lipschitz property of ϕ the random variable $\mathrm{w}_{\mathrm{spu}}^{\top} Z_{\mathrm{spu}}^{\mathrm{OOD}}$ is sub-Gaussian with mean

$$\mathbb{E}\Big[\mathrm{w}_{\mathrm{spu}}^{\top} Z_{\mathrm{spu}}^{\mathrm{OOD}}\Big] = \mathrm{w}_{\mathrm{spu}}^{\top} \mathbb{E}\Big[Z_{\mathrm{spu}}^{\mathrm{OOD}}\Big] = \mathrm{w}_{\mathrm{spu}}^{\top}(M\,\mu_{\mathrm{spu}})$$

and sub-Gaussian parameter at most $L_\phi\,\kappa$ (i.e., with variance proxy bounded by $(L_\phi\,\kappa)^2\,\|\mathrm{w}_{\mathrm{spu}}\|_2^2$). Thus, for any $t > 0$,

$$\Pr\Big(\mathrm{w}_{\mathrm{spu}}^{\top} Z_{\mathrm{spu}}^{\mathrm{OOD}} > \mathrm{w}_{\mathrm{spu}}^{\top}(M\,\mu_{\mathrm{spu}}) + t\Big) \le \exp\Big(-\frac{t^2}{2(L_\phi\,\kappa)^2\,\|\mathrm{w}_{\mathrm{spu}}\|_2^2}\Big).$$

Choose

$$t = \sqrt{2(L_\phi\,\kappa)^2\,\|\mathrm{w}_{\mathrm{spu}}\|_2^2\,\log(1/\delta)}.$$

Then,

$$\Pr\Big(\mathrm{w}_{\mathrm{spu}}^{\top} Z_{\mathrm{spu}}^{\mathrm{OOD}} > \mathrm{w}_{\mathrm{spu}}^{\top}(M\,\mu_{\mathrm{spu}}) + t\Big) \le \delta.$$

Therefore, with probability at least $1-\delta$,

$$\mathrm{w}_{\mathrm{spu}}^{\top} Z_{\mathrm{spu}}^{\mathrm{OOD}} < \mathrm{w}_{\mathrm{spu}}^{\top}(M\,\mu_{\mathrm{spu}}) + \sqrt{2(L_\phi\,\kappa)^2\,\|\mathrm{w}_{\mathrm{spu}}\|_2^2\,\log(1/\delta)}.$$

Assume that

$$\mathrm{w}_{\mathrm{spu}}^{\top}(M\,\mu_{\mathrm{spu}}) + \sqrt{2(L_\phi\,\kappa)^2\,\|\mathrm{w}_{\mathrm{spu}}\|_2^2\,\log(1/\delta)} < 0.$$

Then, with probability at least $1-\delta$, we have

$$\mathrm{w}_{\mathrm{spu}}^{\top} Z_{\mathrm{spu}}^{\mathrm{OOD}} < 0.$$

In this case,

$$\Big\{\mathrm{w}_{\mathrm{dg}}^{\top} Z_{\mathrm{dg}} > -\mathrm{w}_{\mathrm{spu}}^{\top} Z_{\mathrm{spu}}^{\mathrm{OOD}}\Big\} \subset \Big\{\mathrm{w}_{\mathrm{dg}}^{\top} Z_{\mathrm{dg}} > 0\Big\},$$

which implies

$$\Pr\Big(\mathrm{w}_{\mathrm{dg}}^{\top} Z_{\mathrm{dg}} > -\mathrm{w}_{\mathrm{spu}}^{\top} Z_{\mathrm{spu}}^{\mathrm{OOD}}\Big) < \Pr\Big(\mathrm{w}_{\mathrm{dg}}^{\top} Z_{\mathrm{dg}} > 0\Big).$$

Equivalently,

$$\begin{aligned}
\mathrm{acc}_{\mathrm{OOD}}(f_{\mathrm{X}}^{P_{\mathrm{ID}}}) &= \Pr\Big(\mathrm{w}_{\mathrm{dg}}^{\top} Z_{\mathrm{dg}} + \mathrm{w}_{\mathrm{spu}}^{\top} Z_{\mathrm{spu}}^{\mathrm{OOD}} > 0\Big) \\
&= \Pr\Big(\mathrm{w}_{\mathrm{dg}}^{\top} Z_{\mathrm{dg}} > -\mathrm{w}_{\mathrm{spu}}^{\top} Z_{\mathrm{spu}}^{\mathrm{OOD}}\Big) \\
&< \Pr\Big(\mathrm{w}_{\mathrm{dg}}^{\top} Z_{\mathrm{dg}} > 0\Big) \\
&= \mathrm{acc}_{\mathrm{OOD}}(f_{\mathrm{dg}}^{\mathcal{E}}).
\end{aligned}$$

Therefore, with probability at least $1-\delta$,

$$\mathrm{acc}_{\mathrm{OOD}}(f_{\mathrm{X}}^{P_{\mathrm{ID}}}) < \mathrm{acc}_{\mathrm{OOD}}(f_{\mathrm{dg}}^{\mathcal{E}}).$$

□

Lemma 2 (Monotonicity of 0–1 Accuracy under Symmetry)**.** *Let W_0 be a* continuous*, nondegenerate random variable satisfying*

$$W_0 \stackrel{d}{=} -W_0,$$

i.e. centrally symmetric about zero, and assume its CDF F_{W_0} is strictly increasing on $\mathbb{R}$ with $F_{W_0}(0) = \frac{1}{2}$. Fix $p = \Pr(Y = 1) \in (0, 1)$, and define the conditional scores

$$U \mid (Y = 0) = \sigma W_0, \qquad U \mid (Y = 1) = \mu_1 + \sigma W_0,$$

with $\sigma > 0$ and $\mu_1 \in \mathbb{R}$. Let

$$r = \frac{\mu_1 - 0}{\sigma} = \frac{\mu_1}{\sigma}.$$

Then the 0–1 accuracy of the threshold-at-zero classifier $\widehat{Y} = \mathbf{1}\{U > 0\}$ can be written as

$$acc = p\, F_{W_0}(r) + (1 - p)\, \Pr(W_0 \le 0) = p\, F_{W_0}(r) + (1 - p)\, \tfrac{1}{2},$$

and this accuracy is a strictly increasing *function of the signal-to-noise ratio r.*

Proof. We first define the accuracy decomposition. Writing $\widehat{Y} = 1 \iff U > 0$, we have

$$\begin{aligned} \mathrm{acc} &= \Pr(\widehat{Y} = Y) = \Pr(\widehat{Y} = 1, Y = 1) + \Pr(\widehat{Y} = 0, Y = 0) \\ &= p\, \Pr(U > 0 \mid Y = 1) + (1 - p)\, \Pr(U \le 0 \mid Y = 0). \end{aligned}$$

Then we use a change of variables using symmetry. Since $U \mid Y = 1 = \mu_1 + \sigma W_0$,

$$\Pr(U > 0 \mid Y = 1) = \Pr\big(W_0 > -\tfrac{\mu_1}{\sigma}\big) = 1 - F_{W_0}(-r) = F_{W_0}(r),$$

and since $U \mid Y = 0 = \sigma W_0$ with $\Pr(W_0 \le 0) = F_{W_0}(0) = \frac{1}{2}$,

$$\Pr(U \le 0 \mid Y = 0) = \tfrac{1}{2}.$$

Substituting into the decomposition gives

$$\mathrm{acc} = p\, F_{W_0}(r) + (1 - p)\, \tfrac{1}{2} = G(r).$$

Because F_{W_0} is strictly increasing and $p > 0$, G is strictly increasing in r. Hence accuracy increases in the signal-to-noise ratio $r = \mu_1/\sigma$. □

A.3 Theorem 2—Necessary Conditions for Well-Specified Domain Generalization Benchmark Splits

Assume $Z^{\mathrm{ID}}_{\mathrm{spu}}$ is a random variable with mean μ_{spu}, covariance Σ_{spu}, and $\mathrm{w}_{\mathrm{dg}}^\top Z^{\mathrm{OOD}}_{\mathrm{dg}}$ is symmetric about its mean. Define a nonlinear transformation

$$\phi : \mathbb{R}^l \to \mathbb{R}^l,$$

that is L_ϕ–Lipschitz, and let

$$Z^{\mathrm{OOD}}_{\mathrm{spu}} = \phi(Z^{\mathrm{ID}}_{\mathrm{spu}}).$$

Assume further that

$$\mathbb{E}[Z^{\mathrm{OOD}}_{\mathrm{spu}}] = \mathbb{E}[\phi(Z^{\mathrm{ID}}_{\mathrm{spu}})] = M\, \mu_{\mathrm{spu}}, \quad \text{and} \quad \mathrm{Var}(Z^{\mathrm{OOD}}_{\mathrm{spu}}) = \Sigma_\phi$$

for some matrix $M \in \mathbb{R}^{l \times l}$, and that the scalar and $\mathrm{w}_{\mathrm{spu}}^\top Z^{\mathrm{OOD}}_{\mathrm{spu}}$ is symmetric about its mean. Then,

$$\mathrm{acc}_{P_{\mathrm{OOD}}}(f_{\mathrm{X}}^{P_{\mathrm{ID}}}) < \mathrm{acc}_{P_{\mathrm{OOD}}}(f_{\mathrm{dg}}^{\mathcal{E}}) \iff \frac{\mathrm{w}_{\mathrm{dg}}^\top \mu_{\mathrm{dg}} + \mathrm{w}_{\mathrm{spu}}^\top M \mu_{\mathrm{spu}}}{\sqrt{\mathrm{w}_{\mathrm{dg}}^\top \Sigma_{\mathrm{dg}} \mathrm{w}_{\mathrm{dg}} + \mathrm{w}_{\mathrm{spu}}^\top \Sigma_\phi \mathrm{w}_{\mathrm{spu}}}} < \frac{\mathrm{w}_{\mathrm{dg}}^\top \mu_{\mathrm{dg}}}{\sqrt{\mathrm{w}_{\mathrm{dg}}^\top \Sigma_{\mathrm{dg}} \mathrm{w}_{\mathrm{dg}}}}. \tag{15}$$

Note that the RHS of Equation 15 implies that either:

- *Spurious Correlation Reversal:* $\mathrm{w}_{\mathrm{spu}}^\top M \mu_{\mathrm{spu}} < 0$ (related to sufficiency condition without the symmetry assumption in Theorem 1), or
- sufficiently large variance when $\mathrm{w}_{\mathrm{spu}}^\top M \mu_{\mathrm{spu}} > 0$, i.e., small SNR.

Proof. Directly applying Lemma 2,

$$\mathrm{acc}_{P_{\mathrm{OOD}}}(f_{\mathrm{X}}^{P_{\mathrm{ID}}}) < \mathrm{acc}_{P_{\mathrm{OOD}}}(f_{\mathrm{dg}}^{\mathcal{E}}) \iff \frac{\mathrm{w}_{\mathrm{dg}}^\top \mu_{\mathrm{dg}} + \mathrm{w}_{\mathrm{spu}}^\top M \mu_{\mathrm{spu}}}{\sqrt{\mathrm{w}_{\mathrm{dg}}^\top \Sigma_{\mathrm{dg}} \mathrm{w}_{\mathrm{dg}} + \mathrm{w}_{\mathrm{spu}}^\top \Sigma_\phi \mathrm{w}_{\mathrm{spu}}}} < \frac{\mathrm{w}_{\mathrm{dg}}^\top \mu_{\mathrm{dg}}}{\sqrt{\mathrm{w}_{\mathrm{dg}}^\top \Sigma_{\mathrm{dg}} \mathrm{w}_{\mathrm{dg}}}}.$$

□

A.4 Lemma 4—Accuracy on the Line

Lemma 3. *Assume Z_{spu}^{ID} is sub-Gaussian with mean μ_{spu}, covariance Σ_{spu}, and parameter κ. Define a nonlinear mapping*

$$\phi : \mathbb{R}^l \to \mathbb{R}^l,$$

that is L_ϕ–Lipschitz, and let

$$Z_{spu}^{OOD} = \phi(Z_{spu}^{ID}).$$

Assume further that

$$\mathbb{E}[Z_{spu}^{OOD}] = \mathbb{E}[\phi(Z_{spu}^{ID})] = M\,\mu_{spu}, \text{ and } \Sigma_\phi = Z_{spu}^{OOD}(Z_{spu}^{OOD})^\top$$

and that

$$\|M\,\mu_{spu} - \mu_{spu}\| \le \epsilon_1, \tag{16}$$

$$\left\| w_{spu}^\top \Sigma_\phi^\top\, w_{spu} - w_{spu}^\top \Sigma_{spu}\, w_{spu} \right\| \le \epsilon_2. \tag{17}$$

Moreover, assume there exists a constant $B > 0$ such that for sufficiently small t (a Tsybakov-type condition),

$$\Pr\Big(|f_X(X)| \le t\Big) \le Bt.$$

Then, for any $\delta > 0$, with probability at least $1 - \delta$ over Z_{spu}^{ID}, the following holds for any classifier $f_X \in \mathcal{F}$:

$$\Big| acc_P(f_X) - acc_{P_\phi}(f_X) \Big| \le B\,\epsilon,$$

where

$$\epsilon = \|w_{spu}\|\epsilon_1 + C\sqrt{\log(1/\delta)} + \sqrt{\epsilon_2},$$

and for some small constant $c > 0$,

$$C = c\kappa \cdot \max\Big\{ \|w_{spu}\|,\ \|M\| \cdot \|w_{spu}\| \Big\}.$$

Proof. Define

$$\Delta(X) = f(X) - f(X'),$$

where $X \sim P$ and $X' \sim P_\phi$, respectively, (i.e. X' is obtained by replacing $Z_{\mathrm{spu}}^{\mathrm{ID}}$ with $Z_{\mathrm{spu}}^{\mathrm{OOD}}$). Since

$$f(X) = \mathrm{w}_{\mathrm{dg}}^\top Z_{\mathrm{dg}} + \mathrm{w}_{\mathrm{spu}}^\top Z_{\mathrm{spu}}^{\mathrm{ID}} \quad \text{and} \quad f(X') = \mathrm{w}_{\mathrm{dg}}^\top Z_{\mathrm{dg}} + \mathrm{w}_{\mathrm{spu}}^\top Z_{\mathrm{spu}}^{\mathrm{OOD}},$$

we have

$$\Delta(X) = \mathrm{w}_{\mathrm{spu}}^\top Z_{\mathrm{spu}}^{\mathrm{ID}} - \mathrm{w}_{\mathrm{spu}}^\top Z_{\mathrm{spu}}^{\mathrm{OOD}} = \mathrm{w}_{\mathrm{spu}}^\top \Big(Z_{\mathrm{spu}}^{\mathrm{ID}} - Z_{\mathrm{spu}}^{\mathrm{OOD}} \Big).$$

We now decompose $\Delta(X)$ into a deterministic part $g(X)$ and a stochastic part $h(X)$:

$$\Delta(X) = \underbrace{\Big[\mathrm{w}_{\mathrm{spu}}^{\top}\mathbb{E}[Z_{\mathrm{spu}}^{\mathrm{ID}}] - \mathrm{w}_{\mathrm{spu}}^{\top}\mathbb{E}[Z_{\mathrm{spu}}^{\mathrm{OOD}}]\Big]}_{g(X)} + \tag{18}$$

$$\underbrace{\Big((\mathrm{w}_{\mathrm{spu}}^{\top}Z_{\mathrm{spu}}^{\mathrm{ID}} - \mathbb{E}[\mathrm{w}_{\mathrm{spu}}^{\top}Z_{\mathrm{spu}}^{\mathrm{ID}}]) - (\mathrm{w}_{\mathrm{spu}}^{\top}Z_{\mathrm{spu}}^{\mathrm{OOD}} - \mathbb{E}[\mathrm{w}_{\mathrm{spu}}^{\top}Z_{\mathrm{spu}}^{\mathrm{OOD}}])\Big)}_{h(X)}. \tag{19}$$

Since

$$\mathbb{E}[Z_{\mathrm{spu}}^{\mathrm{ID}}] = \mu_{\mathrm{spu}} \quad \text{and} \quad \mathbb{E}[Z_{\mathrm{spu}}^{\mathrm{OOD}}] = M\,\mu_{\mathrm{spu}},$$

we have

$$\begin{aligned}
|g(X)| &= \big|\mathrm{w}_{\mathrm{spu}}^{\top}\mu_{\mathrm{spu}} - \mathrm{w}_{\mathrm{spu}}^{\top}(M\,\mu_{\mathrm{spu}})\big| && (20)\\
&= \big|\mathrm{w}_{\mathrm{spu}}^{\top}(\mu_{\mathrm{spu}} - M\,\mu_{\mathrm{spu}})\big| && (21)\\
&\le \|\mathrm{w}_{\mathrm{spu}}\|\,\|\mu_{\mathrm{spu}} - M\,\mu_{\mathrm{spu}}\| && (22)\\
&\le \|\mathrm{w}_{\mathrm{spu}}\|\,\epsilon_1. && (23)
\end{aligned}$$

Next, consider the stochastic term $h(X)$. Since $Z_{\mathrm{spu}}^{\mathrm{ID}}$ is sub-Gaussian with parameter κ, both $\mathrm{w}_{\mathrm{spu}}^{\top}Z_{\mathrm{spu}}$ and $\mathrm{w}_{\mathrm{spu}}^{\top}Z_{\mathrm{spu}}^{\mathrm{OOD}}$ are sub-Gaussian with parameters $\kappa\|\mathrm{w}_{\mathrm{spu}}\|$ and $L_{\phi}\kappa\|\mathrm{w}_{\mathrm{spu}}\|$ respectively.

Therefore, for any $t > 0$,

$$\Pr\Big(|\mathrm{w}_{\mathrm{spu}}^{\top}Z_{\mathrm{spu}}^{\mathrm{ID}} - \mathbb{E}[\mathrm{w}_{\mathrm{spu}}^{\top}Z_{\mathrm{spu}}]| > t\Big) \le 2\exp\Big(-\frac{t^2}{2(\kappa\|\mathrm{w}_{\mathrm{spu}}\|)^2}\Big),$$

and

$$\Pr\Big(|\mathrm{w}_{\mathrm{spu}}^{\top}Z_{\mathrm{spu}}^{\mathrm{OOD}} - \mathbb{E}[\mathrm{w}_{\mathrm{spu}}^{\top}Z_{\mathrm{spu}}^{\mathrm{OOD}}]| > t\Big) \le 2\exp\Big(-\frac{t^2}{2(L_{\phi}\kappa\|\mathrm{w}_{\mathrm{spu}}\|)^2}\Big).$$

Applying the union bound, with probability at least $1-\delta$ we have

$$\Big|\mathrm{w}_{\mathrm{spu}}^{\top}Z_{\mathrm{spu}} - \mathbb{E}[\mathrm{w}_{\mathrm{spu}}^{\top}Z_{\mathrm{spu}}]\Big| + \Big|\mathrm{w}_{\mathrm{spu}}^{\top}Z_{\mathrm{spu}}^{\mathrm{OOD}} - \mathbb{E}[\mathrm{w}_{\mathrm{spu}}^{\top}Z_{\mathrm{spu}}^{\mathrm{OOD}}]\Big| \le C\sqrt{\log(1/\delta)},$$

where with a small constant factor $c > 0$,

$$C = c\kappa \cdot \max\Big\{\|\mathrm{w}_{\mathrm{spu}}\|,\ L_{\phi}\cdot\|\mathrm{w}_{\mathrm{spu}}\|\Big\}.$$

Additionally, by assumption,

$$\Big|\mathrm{w}_{\mathrm{spu}}^{\top}(M\,\mu_{\mathrm{spu}}) - \mathrm{w}_{\mathrm{spu}}^{\top}\mu_{\mathrm{spu}}\Big| \le \|\mathrm{w}_{\mathrm{spu}}\|\,\epsilon_1,$$

and

$$\Big|\mathrm{w}_{\mathrm{spu}}^{\top}\left(\Sigma_{\phi}\mathrm{w}_{\mathrm{spu}}\right) - \mathrm{w}_{\mathrm{spu}}^{\top}\left(\Sigma_{\mathrm{spu}}\mathrm{w}_{\mathrm{spu}}\right)\Big| \le \epsilon_2.$$

Combining these bounds, with probability at least $1-\delta$ we obtain

$$\Delta(X) \le \|\mathrm{w}_{\mathrm{spu}}\|\epsilon_1 + C\sqrt{\log(1/\delta)} + \sqrt{\epsilon_2} = \epsilon.$$

Finally, the classifier's accuracy difference is determined by the probability that $f(X)$ and $f(X')$ disagree in sign. Given a Tsybakov-type condition, $\Pr(|f_{\mathrm{X}}(X)| \le t) \le Bt$, this probability is controlled by $B\epsilon$. It follow that,

$$\Big|\mathrm{acc}_P(f) - \mathrm{acc}_{P_{\phi}}(f)\Big| \le B\epsilon.$$

This completes the proof. □

Lemma 4. *Assume Z_{spu}^{ID} is sub-Gaussian with mean μ_{spu}, covariance Σ_{spu}, and parameter κ. Define a nonlinear mapping*

$$\phi : \mathbb{R}^l \to \mathbb{R}^l,$$

which is L_ϕ–Lipschitz, and let

$$Z_{spu}^{OOD} = \phi(Z_{spu}^{ID}).$$

Assume further that

$$\mathbb{E}[Z_{spu}^{OOD}] = \mathbb{E}[\phi(Z_{spu}^{ID})] = M\,\mu_{spu}, \text{ and } \Sigma_\phi = Z_{spu}^{OOD}(Z_{spu}^{OOD})^\top$$

and that

$$\|M\,\mu_{spu} - \mu_{spu}\| \le \epsilon_1, \tag{24}$$

$$\left\| w_{spu}^\top \Sigma_\phi w_{spu} - w_{spu}^\top \Sigma_{spu}\, w_{spu} \right\| \le \epsilon_2, \tag{25}$$

where the second inequality is understood to control the difference in the covariance (or concentration) of the spurious features after transformation. Moreover, assume there exists a constant $B > 0$ such that for sufficiently small t (a Tsybakov-type condition),

$$\Pr\Big(|f_X(X)| \le t\Big) \le Bt,$$

and there exists $\alpha > 0$ such that

$$acc_P(f_X) \in [\alpha, 1-\alpha] \quad \text{and} \quad a\, acc_{P_\phi}(f_X) \in [\alpha, 1-\alpha].$$

Then for any $\delta \in (0,1)$, with probability at least $1-\delta$,

$$\Big|\Phi^{-1}\big(acc_P(f_X)\big) - a\,\Phi^{-1}\big(acc_{P_\phi}(f_X)\big)\Big| \le \widetilde{\epsilon}, \tag{26}$$

for any classifier $f_X \in \mathcal{F}$, where

$$\widetilde{\epsilon} = LB\Big(\|w_{spu}\|\epsilon_1 + C\sqrt{\log(1/\delta)} + \sqrt{\epsilon_2}\Big) + \zeta,$$

with—for a small constant factor $c > 0$—

$$C = c\kappa \cdot \max\Big\{\|w_{spu}\|,\ \|M\| \cdot \|w_{spu}\|\Big\},$$

and

$$\zeta = a|1-a| \max_{x \in [\alpha, 1-\alpha]} \Big|\Phi^{-1}(x)\Big|,$$

where Φ is the Gaussian cumulative distribution function, and L is its Lipschitz constant on $[\alpha, 1-\alpha]$, i.e., for $p, q \in [\alpha, 1-\alpha]$,

$$\Big|\Phi^{-1}(p) - \Phi^{-1}(q)\Big| \le L|p-q|.$$

Proof. By Lemma 3, with probability at least $1-\delta$ we have

$$\Big|\mathrm{acc}_P(f_\mathrm{X}) - a\,\mathrm{acc}_{P_\phi}(f_\mathrm{X})\Big| \le B\Big(\|\mathrm{w_{spu}}\|\epsilon_1 + C\sqrt{\log(1/\delta)} + \sqrt{\epsilon_2}\Big).$$

Since $\mathrm{acc}_P(f_\mathrm{X})$ and $a\,\mathrm{acc}_{P_\phi}(f_\mathrm{X})$ lie in $[\alpha, 1-\alpha]$, the function Φ^{-1} is Lipschitz on this interval with constant L. Therefore, if we set $p = \mathrm{acc}_P(f_\mathrm{X})$ and $q = \mathrm{acc}_{P_\phi}(f_\mathrm{X})$, then

$$\Big|\Phi^{-1}(p) - \Phi^{-1}(q)\Big| \le L|p-q|.$$

Taking into account the scaling factor a yields

$$\Big|\Phi^{-1}(p) - a\,\Phi^{-1}(q)\Big| \le L|p-q| + |1-a|\Big|\Phi^{-1}(q)\Big|.$$

Since $|p-q|$ is bounded by the result of Lemma 3, we obtain

$$\left|\Phi^{-1}\big(\mathrm{acc}_P(f_{\mathrm{X}})\big) - a\,\Phi^{-1}\big(\mathrm{acc}_{P_\phi}(f_{\mathrm{X}})\big)\right| \le LB\Big(\|\mathrm{w}_{\mathrm{spu}}\|\epsilon_1 \tag{27}$$

$$+ C\sqrt{2\log(4/\delta)} + \sqrt{\epsilon_2}\Big) + |1-a| \max_{x\in[\alpha,1-\alpha]}\left|\Phi^{-1}(x)\right|. \tag{28}$$

Defining the right-hand side as $\widetilde{\epsilon}$ completes the proof. □

A.5 Lemma 5—Tradeoff Between Accuracy on The Line and Well-Specification

Lemma 5. *Assume Z^{ID}_{spu} is sub-Gaussian with mean μ_{spu}, covariance Σ_{spu}, and parameter κ. Define a nonlinear mapping*

$$\phi : \mathbb{R}^l \to \mathbb{R}^l,$$

which is L_ϕ–Lipschitz, and let

$$Z^{OOD}_{spu} = \phi(Z^{ID}_{spu}).$$

Assume further that

$$\mathbb{E}[Z^{OOD}_{spu}] = \mathbb{E}[\phi(Z^{ID}_{spu})] = M\,\mu_{spu}, \text{ and } \Sigma_\phi = Z^{OOD}_{spu}(Z^{OOD}_{spu})^\top$$

and that

$$\|M\,\mu_{spu} - \mu_{spu}\| \le \epsilon_1, \tag{29}$$

$$\left\|w^\top_{spu}\Sigma_\phi w_{spu} - w^\top_{spu}\Sigma_{spu}\,w_{spu}\right\| \le \epsilon_2. \tag{30}$$

Fix $w_{spu} \in \mathbb{R}^l$ so that $w^\top_{spu}\mu_{spu} > 0$. Suppose that M satisfies the spurious correlation reversal condition

$$w^\top_{spu}(M\,\mu_{spu}) + \sqrt{2(L_\phi\kappa)^2\|w_{spu}\|_2^2\log(1/\delta)} \le -\gamma < 0,$$

for some margin $\gamma > 0$. Moreover, assume there exists a constant $B > 0$ such that for sufficiently small t (a Tsybakov-type condition),

$$\Pr\Big(|f_X(X)| \le t\Big) \le Bt,$$

and that there exists some $\alpha > 0$ such that

$$acc_P(f_X),\ a\,acc_{P_\phi}(f_X) \in [\alpha, 1-\alpha].$$

Then, with probability at least $1-\delta$,

$$\left|\Phi^{-1}\big(acc_P(f_X)\big) - a\,\Phi^{-1}\big(acc_{P_\phi}(f_X)\big)\right| \ge C\,\|w_{spu}\|\sqrt{\log(1/\delta)}\,\|M\,\mu_{spu} - \mu_{spu}\| - \zeta, \tag{31}$$

where

$$\zeta = |1-a| \max_{x\in[\alpha,1-\alpha]}\left|\Phi^{-1}(x)\right|,$$

and C is a positive constant (depending on α and the local slope of Φ^{-1}, Lipschitzness of ϕ, and concentration of Z^{OOD}_{spu}). Moreover,

$$\|M\,\mu_{spu} - \mu_{spu}\| \ge \|w_{spu}\|^{-1}\Big(\gamma + w^\top_{spu}\mu_{spu}\Big),$$

so that the right-hand side of (31) *is strictly positive whenever $\gamma + w^\top_{spu}\mu_{spu} > 0$.*

Proof. Since the spurious correlation reversal condition yields

$$\mathrm{w}^\top_{\mathrm{spu}}(M\,\mu_{\mathrm{spu}}) \le -\gamma < 0,$$

and since $\mathrm{w}^\top_{\mathrm{spu}}\mu_{\mathrm{spu}} > 0$, we have

$$\mathrm{w}^\top_{\mathrm{spu}}(M\,\mu_{\mathrm{spu}} - \mu_{\mathrm{spu}}) = \mathrm{w}^\top_{\mathrm{spu}}(M\,\mu_{\mathrm{spu}}) - \mathrm{w}^\top_{\mathrm{spu}}\mu_{\mathrm{spu}} \le -\gamma - \mathrm{w}^\top_{\mathrm{spu}}\mu_{\mathrm{spu}}.$$

By the Cauchy–Schwarz inequality,

$$\|\mathrm{w}_{\mathrm{spu}}\|\,\|M\,\mu_{\mathrm{spu}}-\mu_{\mathrm{spu}}\| \geq \left|\mathrm{w}_{\mathrm{spu}}^{\top}(M\,\mu_{\mathrm{spu}}-\mu_{\mathrm{spu}})\right| \geq \gamma + \mathrm{w}_{\mathrm{spu}}^{\top}\mu_{\mathrm{spu}},$$

so that

$$\|M\,\mu_{\mathrm{spu}}-\mu_{\mathrm{spu}}\| \geq \frac{\gamma + \mathrm{w}_{\mathrm{spu}}^{\top}\mu_{\mathrm{spu}}}{\|\mathrm{w}_{\mathrm{spu}}\|}.$$

Next, note that

$$\mathrm{acc}_P(f_{\mathrm{X}}) = \Pr\Big(\mathrm{w}_{\mathrm{dg}}^{\top}Z_{\mathrm{dg}} > -\mathrm{w}_{\mathrm{spu}}^{\top}Z_{\mathrm{spu}}^{\mathrm{ID}}\Big) \quad \text{and} \quad \mathrm{acc}_{P_\phi}(f_{\mathrm{X}}) = \Pr\Big(\mathrm{w}_{\mathrm{dg}}^{\top}Z_{\mathrm{dg}} > -\mathrm{w}_{\mathrm{spu}}^{\top}Z_{\mathrm{spu}}^{\mathrm{OOD}}\Big).$$

Since $\mathrm{w}_{\mathrm{spu}}^{\top}(M\,\mu_{\mathrm{spu}})$ is very negative relative to the random fluctuations of $\mathrm{w}_{\mathrm{spu}}^{\top}Z_{\mathrm{spu}}^{\mathrm{OOD}}$ (by the sub-Gaussian concentration inequality with parameter κ) and since $\mathrm{w}_{\mathrm{spu}}^{\top}\mu_{\mathrm{spu}} > 0$, one can apply standard concentration arguments to show that with probability at least $1-\delta$

$$\Big|\mathrm{acc}_P(f_{\mathrm{X}}) - a\,\mathrm{acc}_{P_\phi}(f_{\mathrm{X}})\Big| \geq C_0\kappa\,\|\mathrm{w}_{\mathrm{spu}}\|\sqrt{\log(1/\delta)}\,\|M\,\mu_{\mathrm{spu}}-\mu_{\mathrm{spu}}\|$$

for some constant $C_0 > 0$. Since by assumption $\mathrm{acc}_P(f_{\mathrm{X}})$ and $a\,\mathrm{acc}_{P_\phi}(f_{\mathrm{X}})$ lie in $[\alpha, 1-\alpha]$, the inverse Gaussian CDF Φ^{-1} is L-Lipschitz on this interval. Thus, we have

$$\begin{aligned}
\Big|\Phi^{-1}\big(\mathrm{acc}_P(f_{\mathrm{X}})\big) - a\,\Phi^{-1}\big(\mathrm{acc}_{P_\phi}(f_{\mathrm{X}})\big)\Big| &\geq L\Big|\mathrm{acc}_P(f_{\mathrm{X}}) - a\,\mathrm{acc}_{P_\phi}(f_{\mathrm{X}})\Big| - |1-a|\max_{x\in[\alpha,1-\alpha]}\Big|\Phi^{-1}(x)\Big| \\
&\geq L\Big(C_0\kappa\,\|\mathrm{w}_{\mathrm{spu}}\|\sqrt{\log(1/\delta)}\,\|M\,\mu_{\mathrm{spu}}-\mu_{\mathrm{spu}}\|\Big) \\
&\quad - |1-a|\max_{x\in[\alpha,1-\alpha]}\Big|\Phi^{-1}(x)\Big|.
\end{aligned}$$

Defining $C = C_0 L\kappa$ and $\zeta = |1-a|\max_{x\in[\alpha,1-\alpha]}|\Phi^{-1}(x)|$ completes the proof:

$$\Big|\Phi^{-1}\big(\mathrm{acc}_P(f_{\mathrm{X}})\big) - a\,\Phi^{-1}\big(\mathrm{acc}_{P_\phi}(f_{\mathrm{X}})\big)\Big| \geq C\,\|\mathrm{w}_{\mathrm{spu}}\|\sqrt{\log(1/\delta)}\,\|M\,\mu_{\mathrm{spu}}-\mu_{\mathrm{spu}}\| - \zeta.$$

Since $\|M\,\mu_{\mathrm{spu}}-\mu_{\mathrm{spu}}\| \geq \frac{\gamma+\mathrm{w}_{\mathrm{spu}}^{\top}\mu_{\mathrm{spu}}}{\|\mathrm{w}_{\mathrm{spu}}\|}$, the right-hand side is strictly positive whenever $\gamma + \mathrm{w}_{\mathrm{spu}}^{\top}\mu_{\mathrm{spu}} > 0$. □

A.6 Proof of Theorem 3—Benchmarks with Accuracy on the Line are Misspecified Almost Everywhere.

Assume that $Z_{\mathrm{spu}}^{\mathrm{ID}}$ is sub-Gaussian with mean μ_{spu}, covariance Σ_{spu}, and parameter κ. Define a nonlinear mapping

$$\phi : \mathbb{R}^l \to \mathbb{R}^l,$$

which is L_ϕ–Lipschitz, and let

$$Z_{\mathrm{spu}}^{\mathrm{OOD}} = \phi(Z_{\mathrm{spu}}^{\mathrm{ID}}).$$

Assume further that

$$\mathbb{E}[Z_{\mathrm{spu}}^{\mathrm{OOD}}] = \mathbb{E}[\phi(Z_{\mathrm{spu}}^{\mathrm{ID}})] = M\,\mu_{\mathrm{spu}},$$

and that

$$\|M\,\mu_{\mathrm{spu}}-\mu_{\mathrm{spu}}\| \leq \epsilon_1, \tag{32}$$

$$\Big\|\mathrm{w}_{\mathrm{spu}}^{\top}\Sigma_\phi\,\mathrm{w}_{\mathrm{spu}} - \mathrm{w}_{\mathrm{spu}}^{\top}\Sigma_{\mathrm{spu}}\,\mathrm{w}_{\mathrm{spu}}\Big\| \leq \epsilon_2. \tag{33}$$

Suppose that $M \in \mathbb{R}^{l\times l}$ satisfies the *spurious correlation reversal condition*

$$\mathrm{w}_{\mathrm{spu}}^{\top}(M\,\mu_{\mathrm{spu}}) + \sqrt{2\,(L_\phi\,\kappa)^2\,\|\mathrm{w}_{\mathrm{spu}}\|_2^2\,\log(1/\delta)} < 0,$$

and assume that there exists a constant $B > 0$ such that for sufficiently small t (a Tsybakov-type condition),

$$\Pr\Big(|f_{\mathrm{X}}(X)| \le t\Big) \le Bt,$$

and there exists some $\alpha > 0$ such that

$$\mathrm{acc}_P(f_{\mathrm{X}}),\ a\,\mathrm{acc}_{P_\phi}(f_{\mathrm{X}}) \in [\alpha, 1-\alpha],$$

where $\mathrm{acc}_{P_\phi}(f_{\mathrm{X}})$ is the accuracy when the out-of-distribution features are given by $\phi(Z^{\mathrm{ID}}_{\mathrm{spu}})$, and Φ denotes the Gaussian cumulative distribution function.

Define

$$\mathcal{W}_\epsilon = \left\{ M \in \mathbb{R}^{l\times l} : \begin{array}{l} \mathrm{w}_{\mathrm{spu}}^\top (M\,\mu_{\mathrm{spu}}) + \sqrt{2\,(L_\phi\,\kappa)^2\,\|\mathrm{w}_{\mathrm{spu}}\|_2^2\,\log(1/\delta)} < 0, \\ \Big|\Phi^{-1}\big(\mathrm{acc}_P(f_{\mathrm{X}})\big) - a\,\Phi^{-1}\big(\mathrm{acc}_{P_\phi}(f_{\mathrm{X}})\big)\Big| \le \epsilon \end{array} \right\}. \tag{34}$$

Then:

(i) $\mathcal{W}_0$ has Lebesgue measure zero in $\mathbb{R}^{l\times l}$.

(ii) For any $0 \le \epsilon_i \le \epsilon_j$, we have $\mathcal{W}_{\epsilon_i} \subseteq \mathcal{W}_{\epsilon_j}$.

In particular, as $\epsilon \to 0$ (i.e., perfect accuracy on the line), almost every shift is misspecified, and the Lebesgue measure of the set of well–specified shifts grows monotonically with ϵ.

Proof. From Lemma 4 have the inequality

$$\Big|\Phi^{-1}\big(\mathrm{acc}_P(f_{\mathrm{X}})\big) - a\,\Phi^{-1}\big(\mathrm{acc}_{P_\phi}(f_{\mathrm{X}})\big)\Big| \ge C\Big(\|\mathrm{w}_{\mathrm{spu}}\|\sqrt{\log(1/\delta)}\,\|M\mu_{\mathrm{spu}} - \mu_{\mathrm{spu}}\|\Big) - |1-a| \max_{x\in[\alpha,1-\alpha]}\Big|\Phi^{-1}(x)\Big|,$$

where C is a positive constant (depending on the concentration bounds), and L is the Lipschitz constant of Φ^{-1} on $[\alpha, 1-\alpha]$. Suppose

$$\Big|\Phi^{-1}\big(\mathrm{acc}_P(f_{\mathrm{X}})\big) - a\,\Phi^{-1}\big(\mathrm{acc}_{P_\phi}(f_{\mathrm{X}})\big)\Big| \le \epsilon,$$

then for ϵ sufficiently small, it must be that

$$\|M\mu_{\mathrm{spu}} - \mu_{\mathrm{spu}}\| \le \frac{\epsilon + |1-a|\max_{x\in[\alpha,1-\alpha]}\Big|\Phi^{-1}(x)\Big|}{C\,\|\mathrm{w}_{\mathrm{spu}}\|\sqrt{\log(1/\delta)}}. \tag{35}$$

Thus, as $\epsilon \to 0$ we must have

$$\|M\mu_{\mathrm{spu}} - \mu_{\mathrm{spu}}\| = 0,$$

i.e.,

$$M\mu_{\mathrm{spu}} = \mu_{\mathrm{spu}} \implies \mathrm{w}_{\mathrm{spu}}^\top (M\mu_{\mathrm{spu}}) = \mathrm{w}_{\mathrm{spu}}^\top \mu_{\mathrm{spu}} \ge 0.$$

The second equality follows from $\mathrm{w}_{\mathrm{spu}}$ being the optimal contribution of $Z^{\mathrm{ID}}_{\mathrm{spu}}$ to $f_{\mathrm{X}}^{P_{\mathrm{ID}}}$.

Let

$$S = \{M \in \mathbb{R}^{l\times l} : M\mu_{\mathrm{spu}} = \mu_{\mathrm{spu}}\}.$$

Since $\mu_{\mathrm{spu}} \neq 0$, S is an affine subspace of $\mathbb{R}^{l\times l}$ with dimension strictly less than l^2 and hence has Lebesgue measure zero. Since

$$\mathcal{W}_0 \subset S,$$

it follows that $\mathcal{W}_0$ has Lebesgue measure zero.

The monotonicity claim follows immediately from Equations 34-35: if $0 \le \epsilon_i \le \epsilon_j$, then by definition

$$\mathcal{W}_{\epsilon_i} \subseteq \mathcal{W}_{\epsilon_j}.$$

This completes the proof. □

A.7 Example of Shifts with Accuracy on the Line that are Well-Specified

Let $Z_{\text{spu}} \in \mathbb{R}^k$ be Gaussian with mean $\mu_{\text{spu}} = \mathbb{E}[Z_{\text{spu}}] \neq 0$, and covariance Σ_{spu}. Fix $\mathrm{w}_{\text{spu}} \in \mathbb{R}^k$ with $\mathrm{w}_{\text{spu}}^\top \mu_{\text{spu}} \neq 0$ and let $a > 0$. Assume the mapping is linear, i.e. $\phi(u) = M\,u$, so that

$$Z_{\text{spu}}^{\text{OOD}} = \phi(Z_{\text{spu}}^{\text{ID}}) = M\,Z_{\text{spu}}^{\text{ID}}.$$

Then for any $\epsilon > 0$ and $\delta \in (0,1)$, there exists a matrix $M \in \mathbb{R}^{k\times k}$ such that:

$$\mathrm{w}_{\text{spu}}^\top M\mu_{\text{spu}} + \sqrt{2\,\mathrm{w}_{\text{spu}}^\top M\Sigma_{\text{spu}}M^\top \mathrm{w}_{\text{spu}}\,\log(1/\delta)} < 0, \tag{36}$$

$$\left|\Phi^{-1}\big(\mathrm{acc}_P(f_{\mathrm{X}})\big) - a\,\Phi^{-1}\big(\mathrm{acc}_{P_M}(f_{\mathrm{X}})\big)\right| \leq \epsilon, \tag{37}$$

with probability at least $1-\delta$, where $\mathrm{acc}_P(f_{\mathrm{X}})$ and $\mathrm{acc}_{P_M}(f_{\mathrm{X}})$ are defined as in Lemma 4.

For the construction, let

$$v = \frac{\mathrm{w}_{\text{spu}}}{\|\mathrm{w}_{\text{spu}}\|}$$

be the unit vector in the direction of w_{spu}, and define its reflection matrix

$$R = I - 2vv^\top.$$

Note that $Rv = -v$ and $R^2 = I$. Choose a scalar $\alpha > 0$ and define

$$M = \alpha R.$$

Then, we compute:

$$\mathrm{w}_{\text{spu}}^\top(M\mu_{\text{spu}}) = \mathrm{w}_{\text{spu}}^\top(\alpha R\mu_{\text{spu}}) = \alpha\,(\mathrm{w}_{\text{spu}}^\top R\mu_{\text{spu}}) = -\alpha\,\mathrm{w}_{\text{spu}}^\top\mu_{\text{spu}},$$
$$\mathrm{w}_{\text{spu}}^\top(M\Sigma_{\text{spu}}M^\top)\mathrm{w}_{\text{spu}} = \alpha^2\,\mathrm{w}_{\text{spu}}^\top(R\Sigma_{\text{spu}}R^\top)\mathrm{w}_{\text{spu}} = \alpha^2\,\mathrm{w}_{\text{spu}}^\top\Sigma_{\text{spu}}\mathrm{w}_{\text{spu}},$$

since $R\mathrm{w}_{\text{spu}} = -\mathrm{w}_{\text{spu}}$ and R is orthogonal.

The spurious correlation reversal condition (36) becomes

$$-\alpha\,\mathrm{w}_{\text{spu}}^\top\mu_{\text{spu}} + \sqrt{2\,\alpha^2\,\mathrm{w}_{\text{spu}}^\top\Sigma_{\text{spu}}\mathrm{w}_{\text{spu}}\,\log(1/\delta)} < 0.$$

This can be written as

$$\alpha\left(-\mathrm{w}_{\text{spu}}^\top\mu_{\text{spu}} + \alpha\sqrt{2\,\mathrm{w}_{\text{spu}}^\top\Sigma_{\text{spu}}\mathrm{w}_{\text{spu}}\,\log(1/\delta)}\right) < 0.$$

In particular, since $\mathrm{w}_{\text{spu}}^\top\mu_{\text{spu}} > 0$, it suffices to choose α such that

$$\alpha > \frac{\sqrt{2\,\mathrm{w}_{\text{spu}}^\top\Sigma_{\text{spu}}\mathrm{w}_{\text{spu}}\,\log(1/\delta)}}{\mathrm{w}_{\text{spu}}^\top\mu_{\text{spu}}}.$$

At the same time, we want the errors induced by M to be small. Define the following error terms:

$$\epsilon_1 = \|M\mu_{\text{spu}} - \mu_{\text{spu}}\| = \|\alpha R\mu_{\text{spu}} - \mu_{\text{spu}}\|,$$

and

$$\epsilon_2 = |\alpha^2 - 1| \cdot \left|\mathrm{w}_{\text{spu}}^\top\Sigma_{\text{spu}}\mathrm{w}_{\text{spu}}\right|.$$

We want to choose α close to 1 (so that ϵ_1 and ϵ_2 are small) while also satisfying the above inequality. Hence, we set

$$\alpha = \max\left\{1+\eta,\ \frac{\sqrt{2\,\mathrm{w}_{\text{spu}}^\top\Sigma_{\text{spu}}\mathrm{w}_{\text{spu}}\,\log(1/\delta)}}{\mathrm{w}_{\text{spu}}^\top\mu_{\text{spu}}}\right\},$$

for some small $\eta > 0$ chosen so that $|\alpha - 1|$ is below the desired threshold. By choosing α accordingly, we ensure that:

1. The spurious correlation reversal condition (36) holds.
2. The induced errors ϵ_1 and ϵ_2 are small enough so that, by Lemma 4, we have

$$\left|\Phi^{-1}(\mathrm{acc}_P(f_{\mathrm{X}})) - a\,\Phi^{-1}(\mathrm{acc}_{P_M}(f_{\mathrm{X}}))\right| \le B\Big(\|\mathrm{w}_{\mathrm{spu}}\|\epsilon_1 + C\sqrt{2\log(4/\delta)} + \sqrt{\epsilon_2}\Big) \le \epsilon.$$

Thus, $M = \alpha R$ satisfies both conditions (36) and (37) with probability at least $1-\delta$. Note, however, that the set of such M has Lebesgue measure zero in $\mathbb{R}^{l\times l}$.

A.8 Lemma 6—Finite Mixtures of sub-Gaussians are sub-Gaussian

Lemma 6. *Let X be a finite mixture of sub-Gaussian random variables $X_1, X_2, \ldots, X_k$ with parameters $c_1, c_2, \ldots, c_k$ respectively. That is, $\forall t \in \mathbb{R}$ and each $i \in \{1, \ldots, k\}$,*

$$\mathbb{E}\big[e^{t(X_i - \mathbb{E}[X_i])}\big] \le e^{c_i t^2}.$$

Assume the mixture probabilities $p_1, p_2, \ldots, p_k$ satisfy $\sum_{i=1}^k p_i = 1$ and $p_i \ge 0$. Then X is also sub-Gaussian. Specifically, there exists a constant $c > 0$ such that $\forall t \in \mathbb{R}$,

$$\mathbb{E}\big[e^{t(X - \mathbb{E}[X])}\big] \le e^{ct^2}.$$

Proof. Since X is a mixture, we have

$$\mathbb{E}\big[e^{t(X-\mathbb{E}[X])}\big] = \sum_{i=1}^k p_i\,\mathbb{E}\big[e^{t(X_i-\mathbb{E}[X])}\big].$$

For each i, write

$$X_i - \mathbb{E}[X] = \big(X_i - \mathbb{E}[X_i]\big) + \big(\mathbb{E}[X_i] - \mathbb{E}[X]\big).$$

Thus,

$$\mathbb{E}\big[e^{t(X_i-\mathbb{E}[X])}\big] = e^{t(\mathbb{E}[X_i]-\mathbb{E}[X])}\,\mathbb{E}\big[e^{t(X_i-\mathbb{E}[X_i])}\big] \le e^{t(\mathbb{E}[X_i]-\mathbb{E}[X])}\,e^{c_i\,t^2}.$$

Let

$$\Delta = \max_{1\le i\le k}\big|\mathbb{E}[X_i] - \mathbb{E}[X]\big| \quad \text{and} \quad C = \max_{1\le i\le k} c_i.$$

Then, since $e^{t(\mathbb{E}[X_i]-\mathbb{E}[X])} \le e^{|t|\Delta}$ for each i, it follows that

$$\mathbb{E}\big[e^{t(X-\mathbb{E}[X])}\big] \le \sum_{i=1}^k p_i\,e^{|t|\Delta}\,e^{C\,t^2} = e^{|t|\Delta}\,e^{C\,t^2}.$$

Then we have $\forall t \in \mathbb{R}$

$$e^{|t|\Delta} \le e^{\frac{1}{2}\Delta^2} e^{\frac{1}{2}t^2}$$

and

$$\mathbb{E}\big[e^{t(X-\mathbb{E}[X])}\big] \le e^{\frac{1}{2}\Delta^2}\,e^{\left(C+\frac{1}{2}\right)t^2}.$$

Defining

$$c = C + \frac{1}{2} + \frac{\frac{1}{2}\Delta^2}{t^2},$$

note that the factor $e^{\frac{1}{2}\Delta^2}$ is independent of t and can be absorbed into a constant. In particular, there exists a constant $c' > 0$ (which may depend on Δ and C) such that

$$\mathbb{E}\big[e^{t(X-\mathbb{E}[X])}\big] \le e^{c'\,t^2} \quad \forall t \in \mathbb{R}.$$

Thus, X is sub-Gaussian. □

B Simulation Experiment Setup

Simulation Experiments. We evaluate our results so far empirically. We define an initial distribution with $Z_{\text{dg}} \in \mathbb{R}^2$ as a Gaussian with mean $Y \cdot \mu_{\text{dg}}$, where $\mu_{\text{dg}} = [1; 1]$ and unit variance, and $Z_{\text{spu}}^{\text{ID}} \in \mathbb{R}^2$ as a Gaussian with mean $Y \cdot \mu_{\text{spu}}$, where $\mu_{\text{spu}} = [1; 1]$ and unit variance. The input $X \in \mathbb{R}^4$ and label $y \in \{0, 1\}$. We define a domain by M where $Z_{\text{spu}}^{\text{OOD}} = MZ_{\text{spu}}^{0}$ and all other random variables' distribution is preserved. We consider settings where the training domain is (i) Gaussian and (ii) Sub-Gaussian (mixture of Gausians). We define a set of 50 Gaussian test domains defined by randomly sampled M.

We train two types of models: *domain general*, which are logistic regression models trained and evaluated with only Z_{dg} features but still trained only on the training distribution, and *domain specific*, which are logistic regression models trained an evaluated with X features but still trained only on the training distribution. Details on the experiments can be found in Appendix B.

Figure 1a demonstrates the setting where the training domain is defined by $M = I_{[2]}$, i.e., $Z_{\text{spu}}^{\text{ID}}$ is a multivariate Gaussian. In this setting, we observe the expected behavior derived in Theorem 1. That is, when the spurious correlation reversal and controlled spurious feature variance hold out-of-distribution, the domain-general models outperform the domain-specific models.

Figure 1 demonstrates the setting where the training domain is a mixture of M's, i.e., a mixture of Gaussians making a Sub-Gaussian distribution. Figure 1c demonstrates the setting where M is unconstrained. Here, the test domains can be written as an interpolation of the training domains, i.e., there are positive and negative definite M's mixed to create the training domain. In this setting, Rosenfeld et al. Rosenfeld et al. (2022b) show that the training domain empirical risk minimizer solves the worst-case domain generalization problem. Indeed, figure 1c's results show that there is not generally a difference in OOD performance between the domain-general and domain-specific models.

Figure 1c demonstrates the setting where the testing domains are not a convex combination of the training domain – test domains can be outside the bounds of the training M's. Here, there is a clear difference in the OOD performance between the domain-general and domain-specific models. Furthermore, the expected conditions derived in Theorem 1 are observed. That is, when the spurious correlation reversal and controlled spurious feature variance hold out-of-distribution, the domain-general models outperform the domain-specific models.

Clearly, in natural datasets, it is often impractical to conduct such experiments to determine when a domain-general model achieves the best transfer accuracy on a benchmark. Typically, the domain-general features are unknown, and we lack the ability to manipulate natural datasets. However, we demonstrate below that the absence of a strong positive correlation between in- and out-of-distribution accuracy for arbitrary predictors—referred to as accuracy on the line (Miller et al., 2021), Definition 6—can identify well-specified benchmarks that reliably evaluate domain generalization via transfer accuracy. *We will show that well-specified domain generalization benchmarks exhibit either weak in- and out-of-distribution accuracy correlation or a strong inverse correlation.*

Parameters. We use the following parameters across our experiments: $\mu = [1, 1]$, $\Sigma_{\text{dg}} = \text{diag}([1, 1])$, $\mu_{\text{spu}} = [1, 1]$. We expect our results to hold independent of these parameters. We chose these parameters for the ease of intuition of the results on the simulated dataset. We use a sample size of 1000 for each domain.

$$P_M = \begin{cases} Y & = \text{Bern}(0.5) \\ Z_{\text{dg}} & = \mathcal{N}\big(Y \cdot \mu_{\text{dg}}, \Sigma_{\text{dg}}\big) \\ Z_{\text{spu}} & = \mathcal{N}\big(Y \cdot M\mu_{\text{dg}}, M\Sigma_{\text{spu}}M^{\top}\big) \end{cases} \tag{38}$$

In Figure 1a, We pick a P_I as our train domain and then randomly sample M's to construct P_M's. We then train a domain-specific logistic regression model for $f_{\text{X}} : \mathcal{X} \mapsto \mathcal{Y}$ and a domain-general logistic regression model for $f_{\text{dg}} : \mathcal{Z}_{\text{dg}} \mapsto \mathcal{Y}$ for P_I. We retrain each model

Algorithm 1: Generative Mechanism for ColoredMNIST

Input : MNIST dataset with grayscale images z_{dg} and binary labels $y \in \{0, 1\}$
Output : ColoredMNIST dataset with colorized images x and labels y
Define color mapping probability $P(z_{\text{spu}}|y)$ based on a chosen spurious correlation
Sample $y \sim P(y)$ from the original MNIST dataset
Sample grayscale image z_{dg} corresponding to y
`// Introduce spurious correlation`
With probability p, assign color z_{spu} based on $P(z_{\text{spu}}|y)$
With probability $1 - p$, assign color z_{spu} randomly (breaking correlation)
Apply color transformation $T(z_{\text{dg}}, z_{\text{spu}})$ to obtain x
return (x, y)

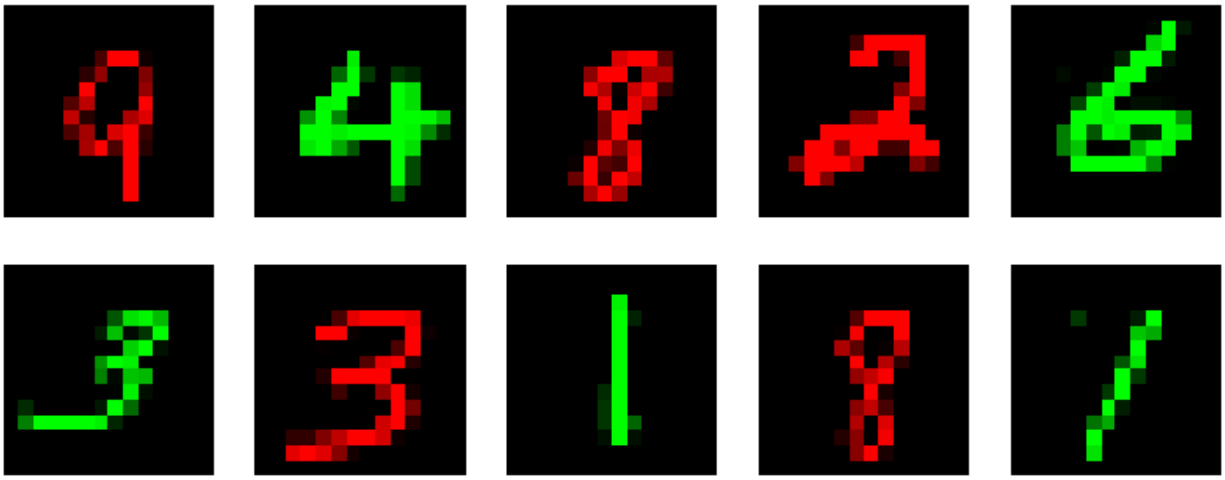


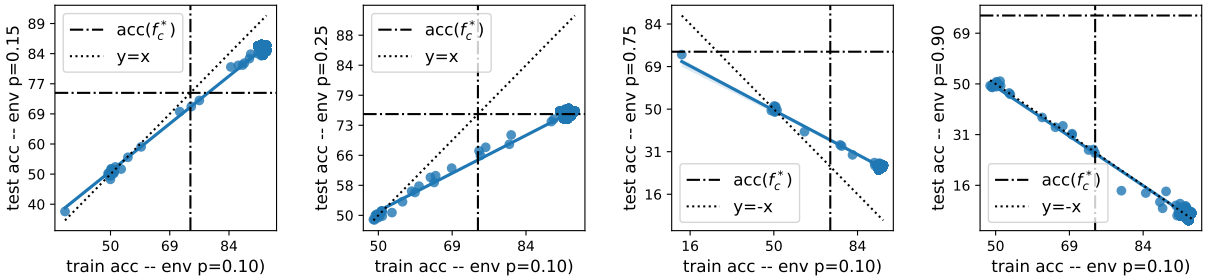
Figure 6: Colored MNIST image examples

B.1 ColoredMNIST Case Study

The ColoredMNIST dataset (Arjovsky et al., 2019) intuitively illustrates the complexity of benchmarking domain generalization. ColoredMNIST modifies the gray-scale MNIST (Deng, 2012) dataset by adding color as a spurious correlation. The digits labels are binary with $+1$ when 'digit ≥ 5' and -1 otherwise. The observed (training) labels, however, contain 25% label noise, i.e., a predictor that uses digit information can achieve 75% accuracy at most, in/out-of-distribution. Additionally, the digit images are colored. The color of the digit matches the noisy observed labels with probability p_e, inducing a *spurious correlation* or *shortcut* of strength p_e. p_e defines a distinct distribution.

Since the observed labels are noisy versions of the true digit labels, the color potentially correlates more with the observed labels than the digit itself. For example, consider a training domain where $p_e^i = P_i(Y = +1 \mid \text{color} = +1) = 0.9$. A color-based predictor would achieve 90% accuracy in-domain, while a domain-general predictor that ignores color would achieve 75% accuracy at a maximum. Furthermore, under a shift where $p_e^j = P_j(Y = 1 \mid \text{color} = +1) > 0.75$, a color-based model trained on p_e^i model will still outperform the domain-general model in OOD accuracy. However, when $p_e^j = P_j(Y = 1 \mid \text{color} = +1) < 0.75$, the same color-based model will transfer worse than the domain-general model. This simple example underscores that for a domain generalization benchmark to be well-specified, w.r.t. to OOD accuracy, spurious correlations from training to test domains must change enough for the domain-general model to achieve the highest possible OOD accuracy.

Figure 7 demonstrates that domain-general models need not transfer the best OOD. To demonstrate this, we test a set of models on a ColoredMNIST training domain where $P_{\text{tr}}(Y = 1 \mid \text{color} = \text{green}) = 0.1$ and



(a) $m = 0.79$
Pearson $R = 0.97$

(b) $m = 0.50$
Pearson $R = 0.97$

(c) m = -0.52
Pearson R = -0.96

(d) m = -0.98
Pearson R = -0.98

Figure 7: Correlations between model performance In-Distristribution vs. Out-of-Distribution on ColoredMNIST variations. m is the slope of the line, and R is the Pearson correlation coefficient. The axis-parallel dashed lines denote the maximum within-domain accuracy of 75%, and $y = x$ represents invariant performance across training and test (target) domains. Models achieving above 75% accuracy use color as a predictor. Figures 7a and 7b represent shifts where color-based predictors achieve the highest OOD accuracy—above 75% accuracy. Without domain knowledge, one might conclude that the best ERM solution is the most domain-general. However, Figures 7c and 7d show that these models are not domain-general; some features that improve ID accuracy hurt OOD performance.

across various test domains with $P_{\text{te}}(Y = 1 \mid \text{color} = \text{green}) = p_e^j$. The observation of the variance in the domain-general and color-based model transfer gap in Figure 7 underscores this work's key question on which ID-OOD shifts allow for reliable domain-generalization evaluation.

We leverage a ConvNet architecture for the ColoredMNIST dataset (Table 4); we vary hyperparameters enumerated in (Gulrajani & Lopez-Paz, 2020a). We vary the hyperparameters in Table 1 and whether or not we use data augmentation.

C Additional Results and Discussion

C.1 Model Training

Data Augmentation. When data augmentation is applied, the transformation consists of a series of preprocessing steps applied to images before they are used for training. First, the image undergoes a **random resized crop** to a size of 224×224 pixels, with a scaling factor ranging from 70% to 100% of the original size. Next, a **random horizontal flip** is applied to introduce variability in orientation. The transformation also includes **color jittering**, which adjusts brightness, contrast, saturation, and hue with a factor of 0.3 each, followed by **random grayscale conversion**, which randomly turns images into grayscale with a certain probability.

Experimental Setup. We follow the following general experimental procedure. When experiments deviate from this, it is specified in their respective sections.

Each dataset consists of E domains, each corresponding to a unique data distribution. Our experiments involve ID/OOD splits using a leave-one-domain-out approach. Specifically, for each domain indexed as $i \in [1..E]$, we train on the subset $\mathcal{E}^i_{\text{train}} = \{e_1, \ldots, e_{i-1}, e_{i+1}, \ldots, e_E\}$ and test on the held-out domain $\mathcal{E}^i_{\text{test}} = \{e_i\}$.

For each i, we train the following models on $P^{\mathcal{E}^i_{\text{train}}}$: ResNet18, ResNet50, DenseNet121, and ConvNeXt_Tiny. For each model, we consider ImageNet pretrained variants: (i) Fine-tuned – end-to-end training on $P^{\mathcal{E}^i_{\text{train}}}$ and

Table 1: Models Generation

Hyperparameter	**Range**
Learning Rate (lr)	10^{-5} to $10^{-3.5}$
Weight Decay	10^{-6} to 10^{-2}
Batch Size	2^3 (8) to $2^{5.5}$ (≈ 45)
Data Augmentation	{True, False}
Transfer Learning	{True, False}
Model Architecture	{ResNet18, ResNet50, DenseNet121, ViT-B-16, and ConvNeXt_Tiny}
Dropout	$\{0.0, 0.1, 0.5\}$
Epoch	–

(ii) Transfer learning – retraining only the last layer on $P^{\mathcal{E}^i_{\text{train}}}$. Models are generated with hyperparameters in Table 1; we also take models at different checkpoints during training.

Table 2: Dataset environments with minimum samples such that change in ID and OOD accuracies correlation changes less than 1% with new models, and total samples included. We do this for each dataset and ID/OOD split. Our analysis accounts for sampling error and ensures this threshold is met with 95% confidence, determined by bootstrapping with 1000 resamplings.

Dataset	**OOD**	**Minimum Samples**	**Total Samples**
CivilComments	Env 0	6,710	10,350
CivilComments	Env 1	7,210	10,350
CivilComments	Env 2	7,610	10,350
CivilComments	Env 3	10,310	10,350
CivilComments	Env 4	6,210	10,350
CivilComments	Env 5	9,810	10,350
CivilComments	Env 6	9,810	10,350
CivilComments	Env 7	10,310	10,350
CivilComments	Env 8	10,310	10,350
CivilComments	Env 9	4,910	10,350
CivilComments	Env 10	5,910	10,350
CivilComments	Env 11	5,510	10,350
CivilComments	Env 12	4,810	10,350
CivilComments	Env 13	5,610	10,350
CivilComments	Env 14	2,510	10,350
CivilComments	Env 15	5,910	10,350
ColoredMNIST	Env 0	14,110	19,758
ColoredMNIST	Env 1	3,110	20,130
ColoredMNIST	Env 2	1,810	20,020
Covid-CXR	Env 0	3,410	7,140
Covid-CXR	Env 1	1,410	7,140
Covid-CXR	Env 2	3,310	7,140
Covid-CXR	Env 3	4,110	7,140
Covid-CXR	Env 4	4,610	7,140
PACS	Env 0	2,310	4,407
PACS	Env 1	3,110	4,761
PACS	Env 2	910	5,043
PACS	Env 3	1,210	4,806
SpawriousM2M Easy	Env 0	1,310	6,630
SpawriousM2M Easy	Env 1	3,010	6,630

Continued on next page

Table 2 continued from previous page

Dataset	OOD	Minimum Samples	Total Samples
SpawriousM2M Easy	Env 2	1,610	6,732
SpawriousM2M Hard	Env 0	5,610	5,916
SpawriousM2M Hard	Env 1	2,910	5,916
SpawriousM2M Hard	Env 2	2,910	5,916
SpawriousO2O Easy	Env 0	4,810	6,630
SpawriousO2O Easy	Env 1	5,610	6,630
SpawriousO2O Easy	Env 2	5,610	6,630
SpawriousO2O Hard	Env 0	4,810	5,916
SpawriousO2O Hard	Env 1	4,310	5,916
SpawriousO2O Hard	Env 2	5,910	5,916
Terra Incognita	Env 0	910	3,186
Terra Incognita	Env 1	1,410	3,186
Terra Incognita	Env 2	1,010	3,135
Terra Incognita	Env 3	1,710	3,165
WILDS Camelyon	Env 0	1,610	8,400
WILDS Camelyon	Env 1	2,010	9,228
WILDS Camelyon	Env 2	3,110	7,408
WILDS Camelyon	Env 3	2,810	8,548
WILDS Camelyon	Env 4	4,710	7,480
WILDS FMoW	Env 0	810	27,340
WILDS FMoW	Env 1	2,310	27,245
WILDS FMoW	Env 2	710	28,725
WILDS FMoW	Env 3	4,110	25,550
WILDS FMoW	Env 4	4,310	23,930
WILDS FMoW	Env 5	7,010	26,540
WILDS Waterbirds	Env 0	2,010	8,823
WILDS Waterbirds	Env 1	510	9,027

Table 3: We train on a set of ID distributions and test on a left-out OOD distribution. We present the Pearson R of ID and OOD probit-transformed accuracies and the slope and intercept of OOD accuracy regressed on ID accuracy. (*) OOD for waterbirds refers to the group where $y = 0$ and $a = 0$. The ID dataset is the mixture of groups at train time. Additional datasets and analysis are provided in Appendix C, which also includes complete tables with all splits for each dataset. With a threshold of $R < 0.5$, only a subset of datasets satisfy our derived conditions.

Dataset	**OOD**	$R <$ **0.5**	**slope**	**offset**	**R**	**p-value**	**std error**
ColoredMNIST	Env 2 acc	✓	-1.56	0.47	-0.74	0.00	0.01
CXR	Env 1 acc	✓	-0.60	0.56	-0.48	0.00	0.03
SpawriousO2O hard	Env 0 acc	✓	0.32	-0.21	0.50	0.00	0.05
SpawriousM2M hard	Env 0 acc	✓	0.76	-0.26	0.94	0.00	0.01
SpawriousO2O easy	Env 0 acc	✗	0.48	-0.29	0.74	0.00	0.04
SpawriousM2M easy	Env 0 acc	✗	0.34	0.26	0.60	0.00	0.00
PACS	Env 1 acc	✗	0.68	-0.68	0.84	0.00	0.01
TerraIncognita	Env 1 acc	✗	0.83	-1.41	0.74	0.00	0.02
Camelyon	Env 2 acc	✗	0.62	0.49	0.78	0.00	0.01
FMoW	Env 5 acc	✗	0.76	-0.61	0.87	0.00	0.01
CivilComments	Env 1 acc	✓	-0.49	0.16	-0.47	0.00	0.03
WaterBirds	Env 0 (*) acc	✓	-0.13	1.58	-0.13	0.00	0.03

C.2 ColoredMNIST

ColoredMNIST (Arjovsky et al., 2019). A variant of the MNIST handwritten digit classification dataset (LeCun, 1998). Domain $d \in \{0.1, 0.2, 0.9\}$ contains a disjoint set of digits colored either red or blue. The label is a noisy function of the digit and color, such that color bears a correlation of d with the label and the digit bears a correlation of 0.75 with the label. This dataset contains 70,000 examples of dimension (2, 28, 28) and 2 classes.

Experimental Details. We leverage a ConvNet architecture for the ColoredMNIST dataset (Table 4).

Table 4: MNIST ConvNet architecture.

#	**Layer**
1	Conv2D (in=d, out=64)
2	ReLU
3	GroupNorm (groups=8)
4	Conv2D (in=64, out=128, stride=2)
5	ReLU
6	GroupNorm (groups=8)
7	Conv2D (in=128, out=128)
8	ReLU
9	GroupNorm (groups=8)
10	Conv2D (in=128, out=128)
11	ReLU
12	GroupNorm (8 groups)
13	Global average-pooling

Discussion. Despite colored MNIST's apparent simplicity, the spurious correlation between color and the label is quite strong – particularly generalization to text environment 2, going from domains with spurious correlation probability of $0.1, 0.2 \rightarrow 0.9$. in Gulrajani & Lopez-Paz (2020a)'s evaluation of standard domain generalization methods at the time, they found that no model could mitigate the effect of this spurious

correlation. We note that this ID/OOD split has a strong accuracy on the inverse line. In test environment 1, we observe that the training distributions are such that the spurious correlations cancel out (0.1 vs. 0.9), and the domain-general model is also the best ID empirical risk minimizer.

Knowledge of the spurious correlation mechanism in each domain makes it relatively easy to identify the type of features a model uses due to the predictability of expected accuracy between models that use color and those that don't. Due to the potential ambiguity of benchmarking results when spurious correlation mechanisms are unknown, semisynthetic benchmarks are vital in the evaluation process.

Table 5: ColoredMNIST ID vs. OOD properties.

OOD	slope	intercept	Pearson R	p-value	standard error
Env 0 acc	1.90	-0.58	0.82	0.00	0.01
Env 1 acc	0.96	0.01	0.94	0.00	0.00
Env 2 acc	-1.56	0.47	-0.74	0.00	0.01

Table 6: ColoredMNIST ID vs. OOD properties.

OOD	ID	slope	intercept	Pearson R	p-value	standard error
Env 0 acc	Env 1 acc	1.23	-0.12	0.99	0.00	0.00
Env 0 acc	Env 2 acc	-0.73	0.87	-0.36	0.00	0.02
Env 1 acc	Env 0 acc	0.91	0.04	0.98	0.00	0.00
Env 1 acc	Env 2 acc	0.67	0.16	0.72	0.00	0.01
Env 2 acc	Env 0 acc	-1.34	0.53	-0.82	0.00	0.01
Env 2 acc	Env 1 acc	-1.65	0.29	-0.64	0.00	0.02

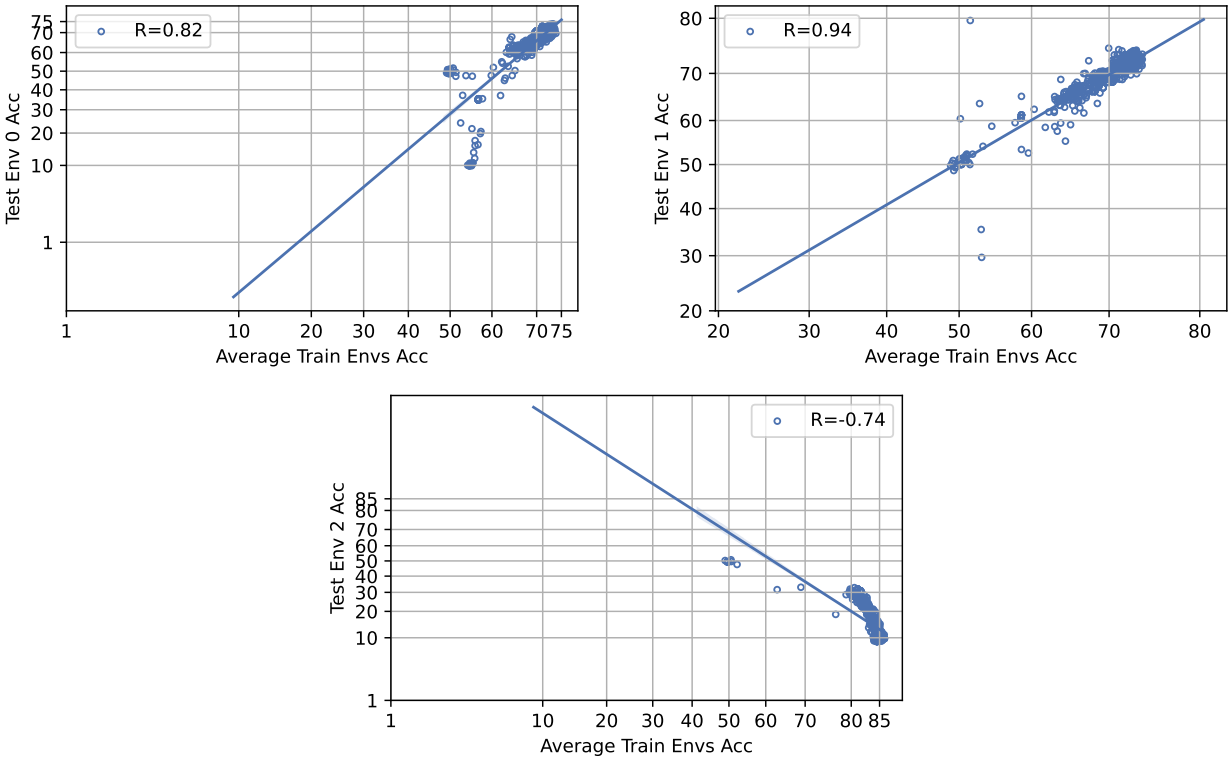


Figure 8: ColoredMNIST: Average train Env Accuracy vs. Test Env Accuracy.

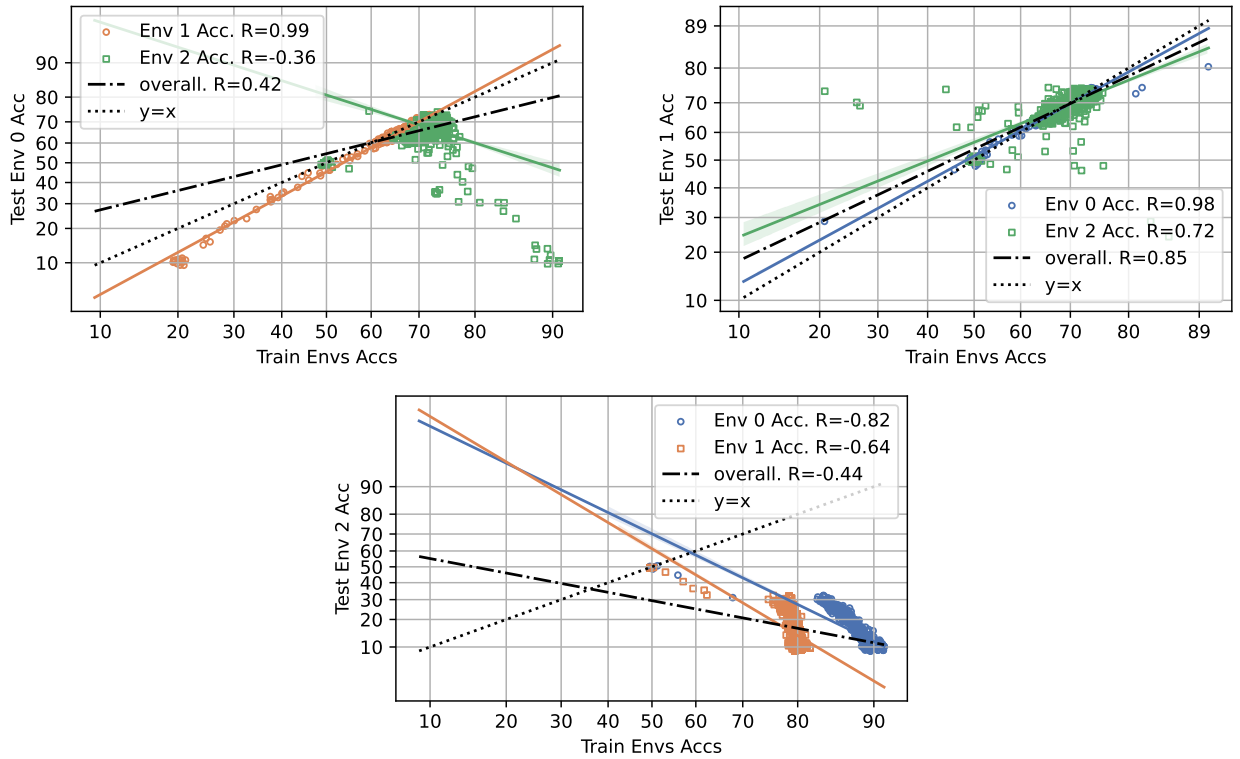


Figure 9: ColoredMNIST: Train Env Accuracy vs. Test Env Accuracy.

C.3 Spawrious

Spawrious (Lynch et al., 2023). The Spawrious image classification benchmark suite consists of six different datasets, including one-to-one (O2O) spurious correlations, where a single spurious attribute correlates with a binary label, and many-to-many (M2M) spurious correlations across multiple classes and spurious attributes. Each benchmark task is proposed with three difficulty levels: Easy, Medium, and Hard. The dataset contains images of four dog breeds $c \in \{bulldog, dachshund, labrador, corgi\}$ found in six backgrounds $b \in \{beach, desert, dirt, jungle, mountain, sand\}$. Images are generated using text-to-image models and filtered using an image-to-text model for quality control. This benchmark suite consists of 152,064 images of dimensions (3, 224, 224).

For the O2O task, the class (dog breed) and background combinations are sampled such that $\mu\%$ of the images per class contain a spurious background b^{sp} and $(100-\mu)\%$ contain a generic background b^{ge}. While the generic background is held constant for each class, each spurious background is observed in only one class ($p_{train}(b_i^{sp} \mid c_j) = 1$ if $i = j$ and 0 if $i \neq j$). Two separate training domains are defined by varying the value of μ. These induced spurious correlations are reverted to yield a test domain with an unseen class-background pair for each class ($p_{test}(b_i \mid c_i) = 1$).

For the M2M task, disjoint class and background groups are constructed $\mathcal{B}_1, \mathcal{B}_2, \mathcal{C}_1, \mathcal{C}_2$, each with two elements. To introduce the training domains, class-background combinations (c, b) are selected with $c \in \mathcal{C}_i$ and $b \in \mathcal{B}_i$. Each training domain consists of a single background per class such that $p^e_{train}(b_k \mid c_k) = e$, with domain index $e \in \{0, 1\}$, $b_k \in \mathcal{B}_i$, $c_k \in \mathcal{C}_i$. In contrast, the test domain is generated by selecting combinations from $c \in \mathcal{C}_i$ and $b \in \mathcal{B}_j$ with $i \neq j$ and sampling backgrounds such that $p_{test}(b_1 \mid c_k) = p_{test}(b_2 \mid c_k) = 0.5$ for $c_k \in \mathcal{C}_i$, $\{b_1, b_2\} = \mathcal{B}_j$.

The difficulty level (Easy, Medium, Hard) differs due to the splits in the available class-background combinations. These splits were empirically determined, and the full details of the final data combinations are found in Table 2 of Lynch et al. (2023).

Discussion. We observed that test environments 1 and 2 have a strong correlation between ID and OOD accuracy and have a slope of 1, making them misspecified for benchmarking domain generalization. Appropriately, Lynch et al. (2023) propose transferring to test environment '0' as the spurious correlation task. The correlation for test environment 0 is much weaker than the others, indicating that ID improvement does not as strongly imply OOD improvement. While there is still a positive linear correlation, the interpretation of these benchmarking results is informative because of the knowledge of the spurious correlation mechanism. Lynch et al. (2023) give examples of informative analysis of benchmarking results on this dataset. Notably, the O2O_easy setting has a weaker correlation by design, and the accuracy on the line strength increases. We see similar behavior for the M2M_ setting. However, this task is much harder than the O2O task, which is reflected in weaker accuracy on the line.

C.3.1 SpawriousO2O Easy

Table 7: SpawriousO2O_easy ID vs. OOD properties.

OOD	slope	intercept	Pearson R	p-value	standard error
Env 0 acc	0.48	-0.29	0.74	0.00	0.04
Env 1 acc	1.05	-0.13	0.98	0.00	0.02
Env 2 acc	0.95	-0.11	0.97	0.00	0.02

Table 8: SpawriousO2O_easy ID vs. OOD properties.

OOD	ID	slope	intercept	Pearson R	p-value	standard error
Env 0 acc	Env 1 acc	0.50	-0.33	0.75	0.00	0.04
Env 0 acc	Env 2 acc	0.47	-0.23	0.72	0.00	0.04
Env 1 acc	Env 0 acc	1.09	-0.33	0.93	0.00	0.04
Env 1 acc	Env 2 acc	1.01	0.11	0.98	0.00	0.02
Env 2 acc	Env 0 acc	0.93	-0.12	0.94	0.00	0.03
Env 2 acc	Env 1 acc	0.94	-0.02	0.98	0.00	0.01

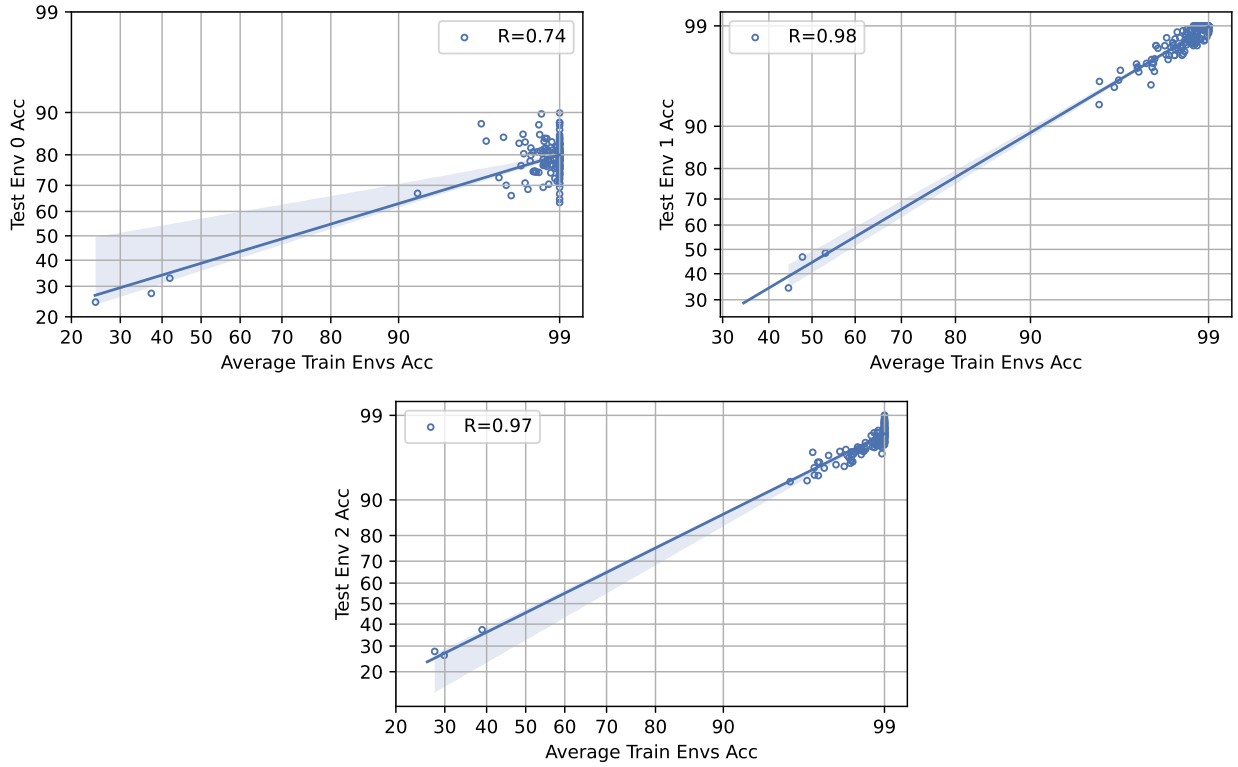


Figure 10: SpawriousO2O easy: Average train Env Accuracy vs. Test Env Accuracy.

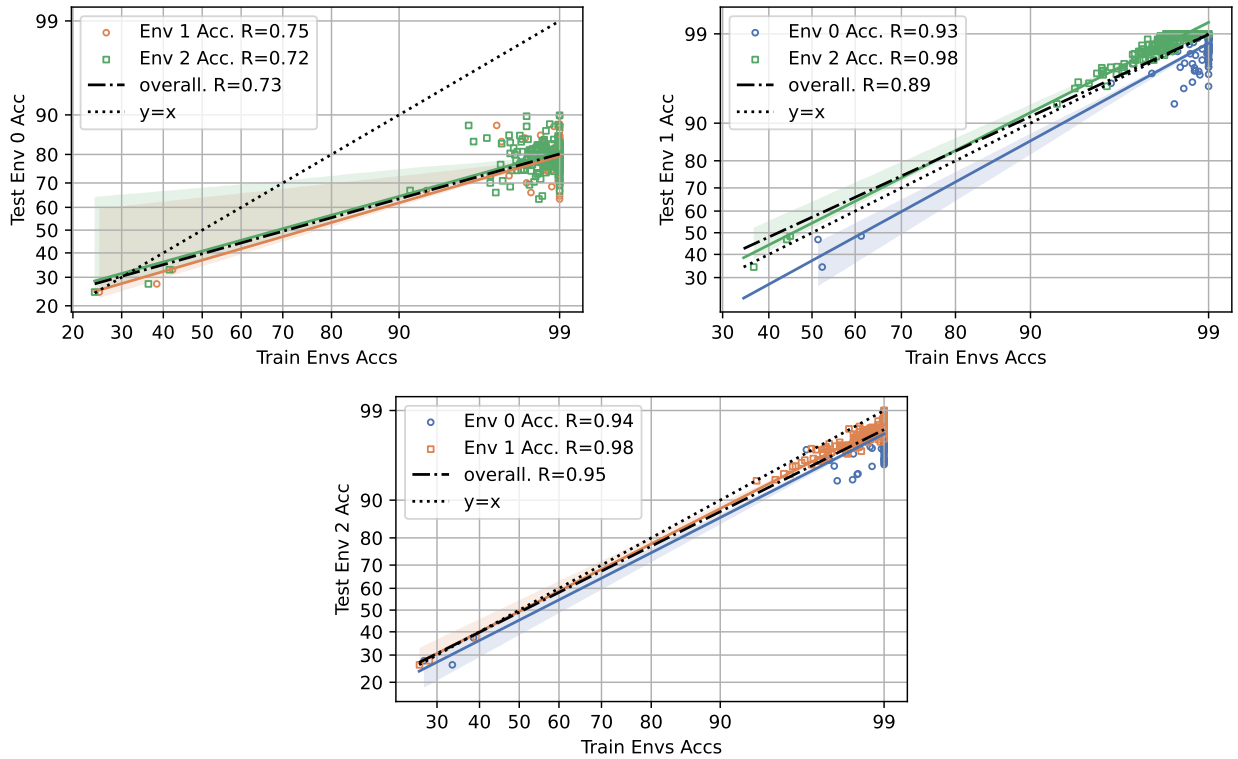


Figure 11: SpawriousO2O easy: Train Env Accuracy vs. Test Env Accuracy.

C.3.2 SpawriousO2O Hard

Table 9: SpawriousO2O_hard ID vs. OOD properties.

OOD	slope	intercept	Pearson R	p-value	standard error
Env 0 acc	0.32	-0.21	0.50	0.00	0.05
Env 1 acc	0.98	0.06	0.96	0.00	0.02
Env 2 acc	0.94	-0.07	0.96	0.00	0.02

Table 10: SpawriousO2O_hard ID vs. OOD properties.

OOD	ID	slope	intercept	Pearson R	p-value	standard error
Env 0 acc	Env 1 acc	0.32	-0.23	0.49	0.00	0.05
Env 0 acc	Env 2 acc	0.32	-0.19	0.50	0.00	0.04
Env 1 acc	Env 0 acc	0.90	0.14	0.89	0.00	0.04
Env 1 acc	Env 2 acc	1.01	0.12	0.97	0.00	0.02
Env 2 acc	Env 0 acc	0.92	-0.05	0.93	0.00	0.03
Env 2 acc	Env 1 acc	0.92	0.01	0.97	0.00	0.02

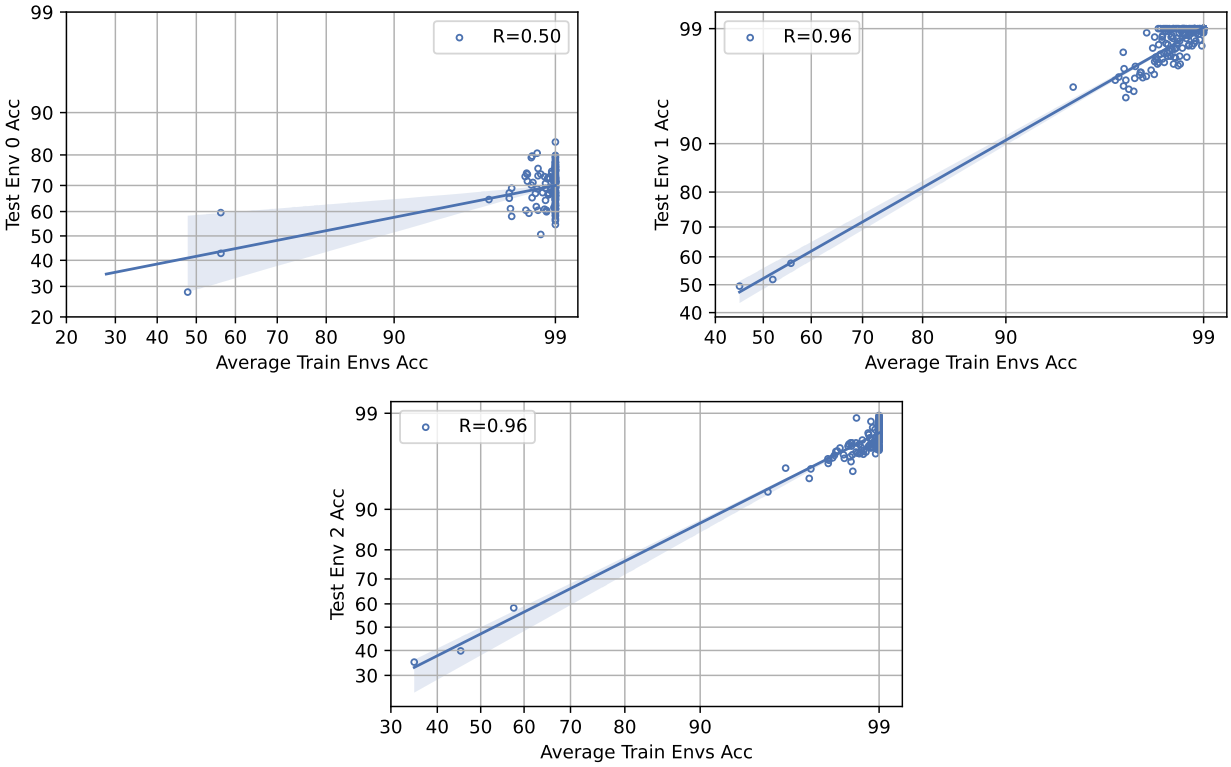


Figure 12: SpawriousO2O hard: Average train Env Accuracy vs. Test Env Accuracy.

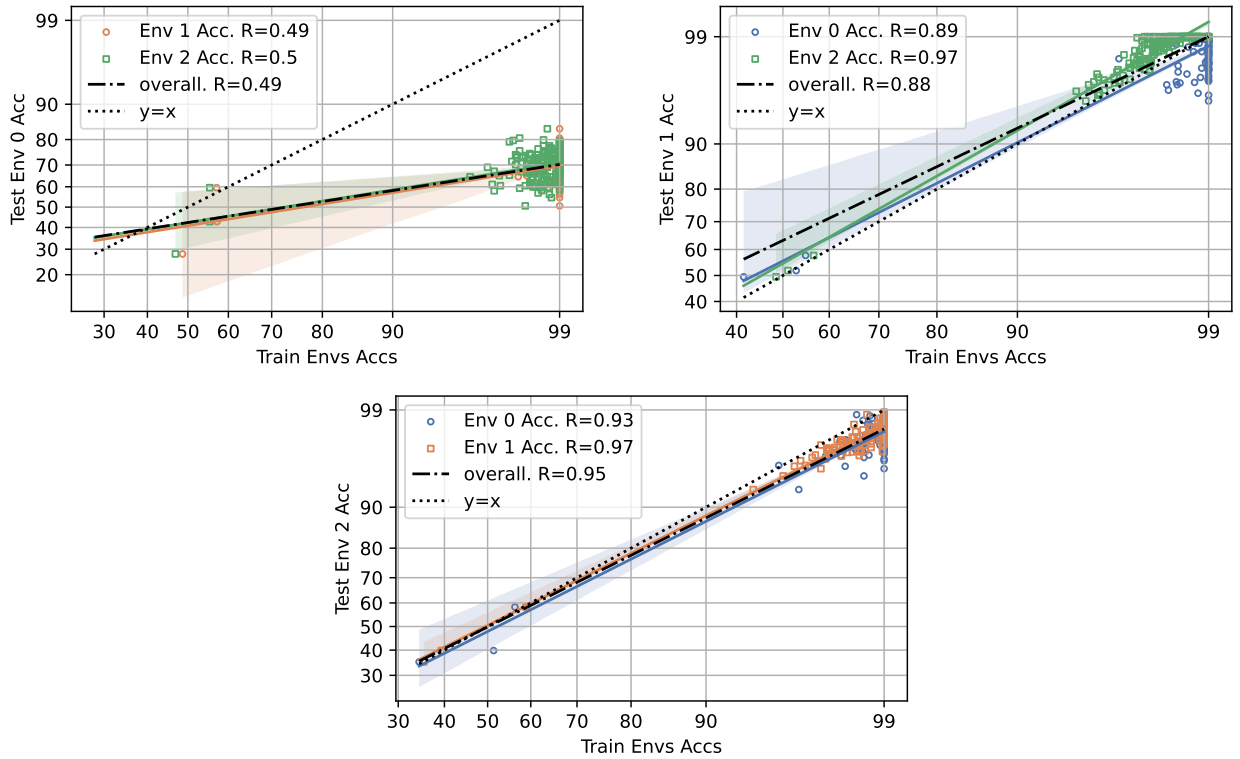


Figure 13: SpawriousO2O hard: Train Env Accuracy vs. Test Env Accuracy.

C.3.3 SpawriousM2M Easy

Table 11: SpawriousM2M_easy ID vs. OOD properties.

OOD	slope	intercept	Pearson R	p-value	standard error
Env 0 acc	0.34	0.26	0.60	0.00	0.01
Env 1 acc	0.65	-0.08	0.95	0.00	0.00
Env 2 acc	0.65	0.02	0.93	0.00	0.00

Table 12: SpawriousM2M_easy ID vs. OOD properties.

OOD	ID	slope	intercept	Pearson R	p-value	standard error
Env 0 acc	Env 1 acc	0.35	0.23	0.61	0.00	0.01
Env 0 acc	Env 2 acc	0.32	0.29	0.58	0.00	0.01
Env 1 acc	Env 0 acc	0.64	-0.09	0.95	0.00	0.00
Env 1 acc	Env 2 acc	0.63	-0.05	0.94	0.00	0.00
Env 2 acc	Env 0 acc	0.67	-0.02	0.94	0.00	0.00
Env 2 acc	Env 1 acc	0.61	0.08	0.90	0.00	0.01

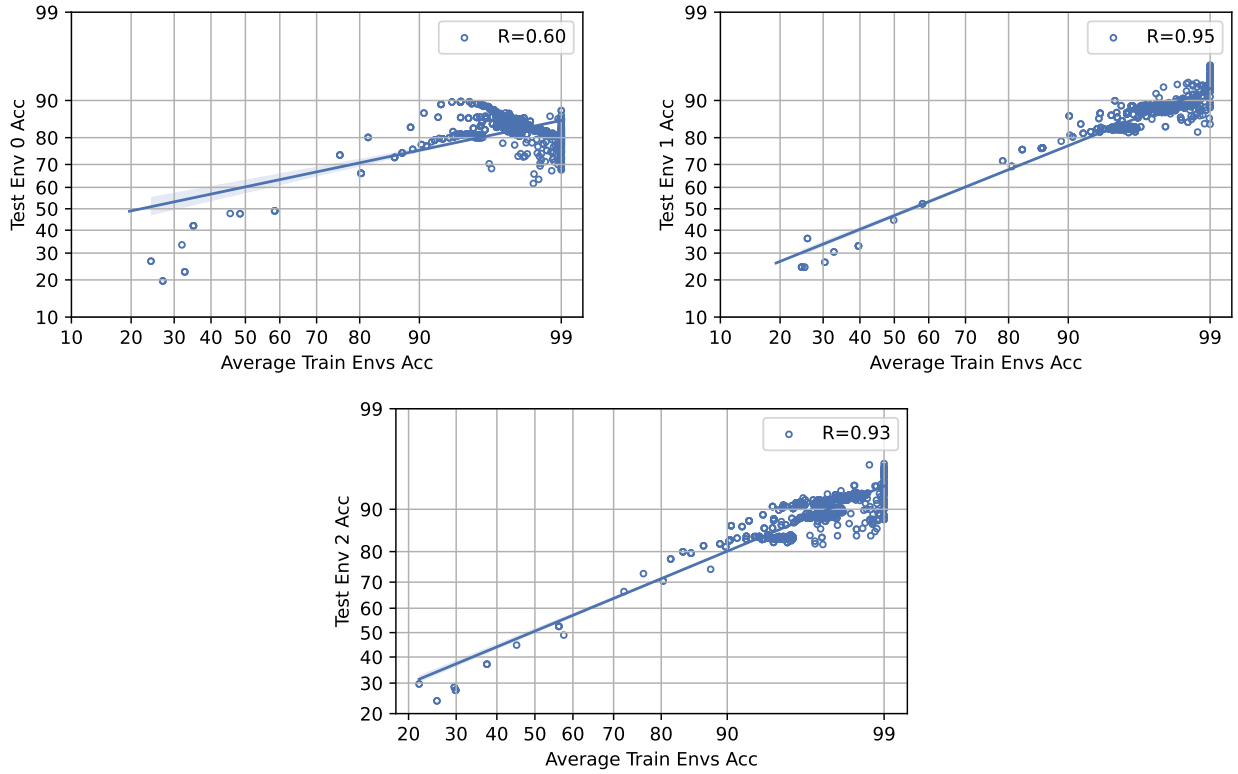


Figure 14: SpawriousO2O easy: Average train Env Accuracy vs. Test Env Accuracy.

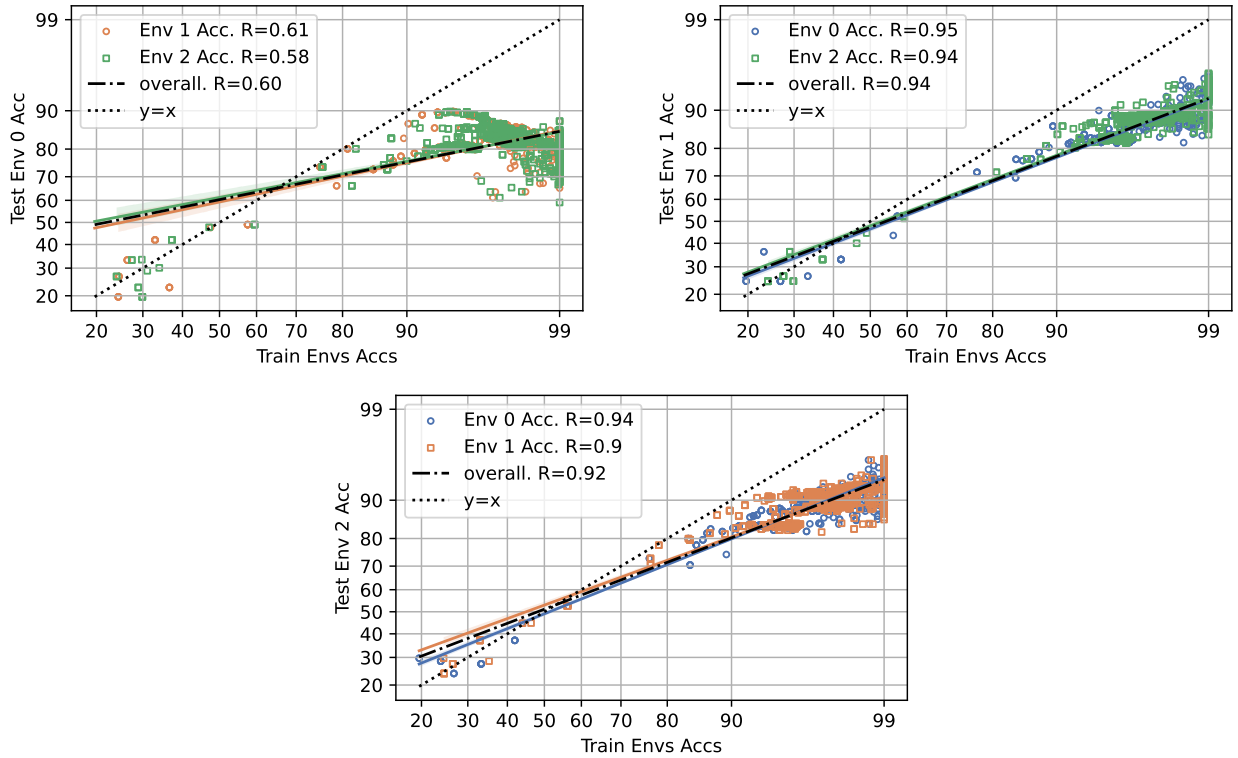


Figure 15: SpawriousO2O easy: Train Env Accuracy vs. Test Env Accuracy.

C.3.4 SpawriousM2M Hard

Table 13: SpawriousM2M_hard ID vs. OOD properties.

OOD	slope	intercept	Pearson R	p-value	standard error
Env 0 acc	0.16	-0.04	0.29	0.00	0.01
Env 1 acc	0.76	-0.26	0.94	0.00	0.01
Env 2 acc	0.66	-0.10	0.91	0.00	0.01

Table 14: SpawriousM2M_hard ID vs. OOD properties.

OOD	ID	slope	intercept	Pearson R	p-value	standard error
Env 0 acc	Env 1 acc	0.17	-0.07	0.33	0.00	0.01
Env 0 acc	Env 2 acc	0.14	-0.02	0.27	0.00	0.01
Env 1 acc	Env 0 acc	0.78	-0.27	0.95	0.00	0.00
Env 1 acc	Env 2 acc	0.73	-0.23	0.92	0.00	0.01
Env 2 acc	Env 0 acc	0.68	-0.11	0.92	0.00	0.01
Env 2 acc	Env 1 acc	0.62	-0.08	0.89	0.00	0.01

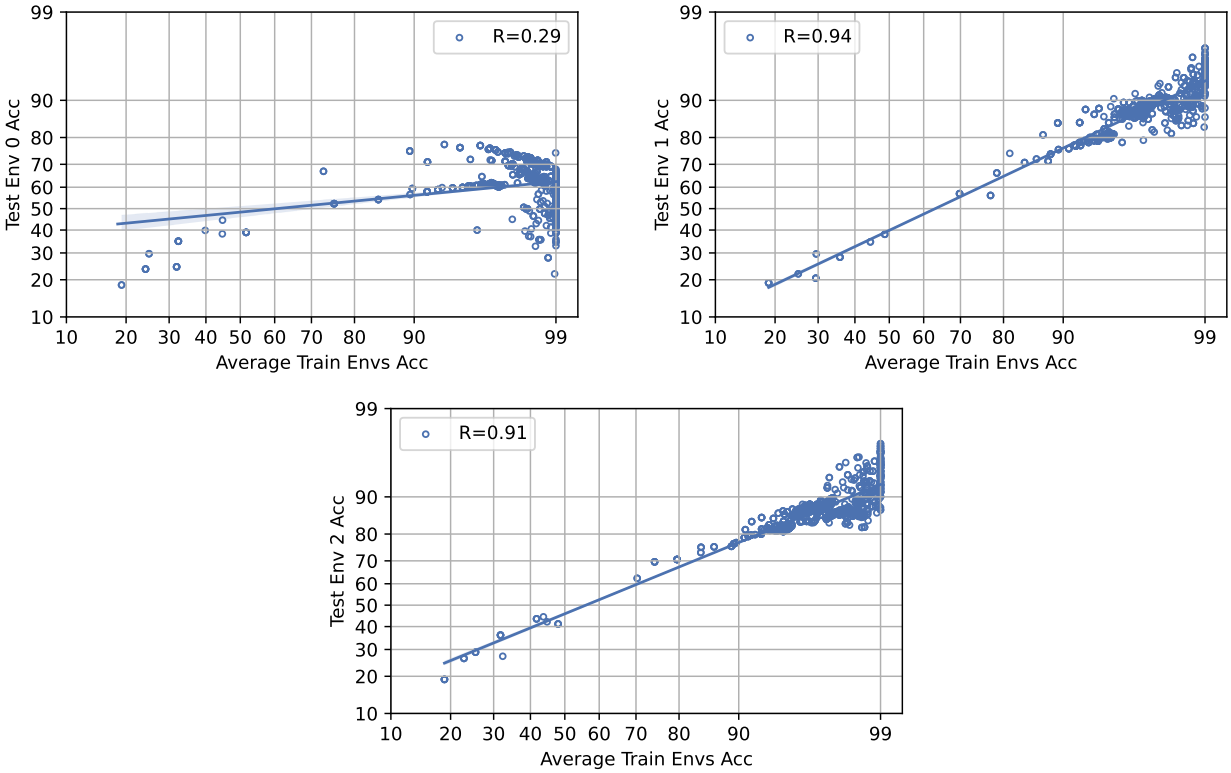


Figure 16: SpawriousM2M Hard: Average train Env Accuracy vs. Test Env Accuracy.

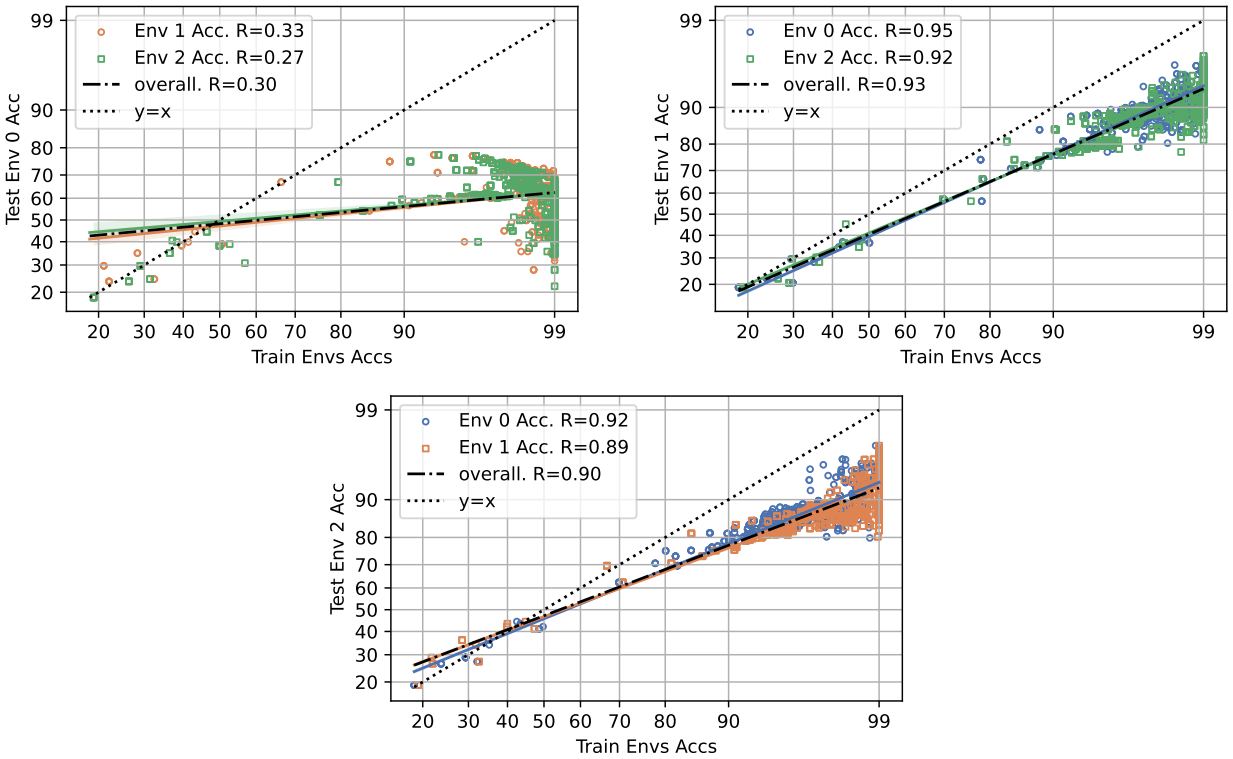


Figure 17: SpawriousM2M Hard: Train Env Accuracy vs. Test Env Accuracy.

C.4 PACS

PACS (Li et al., 2017). A dataset comprised of four domains $d \in \{art, cartoons, photos, sketches\}$. This dataset contains 9,991 examples of dimension (3, 224, 224) and 7 classes.

Discussion. In general, we find that PACS does not strongly represent worst-case shifts for any split. Our results suggest that this benchmark may not accurately benchmark an algorithm's ability to give models free of spurious correlations.

Table 15: PACS ID vs. OOD properties.

OOD	slope	intercept	Pearson R	p-value	standard error
Env 0 acc	0.74	-0.31	0.98	0.00	0.00
Env 1 acc	0.68	-0.68	0.84	0.00	0.01
Env 2 acc	1.00	0.32	0.86	0.00	0.01
Env 3 acc	0.76	-0.87	0.86	0.00	0.01

Table 16: PACS ID vs. OOD properties.

OOD	ID	slope	intercept	Pearson R	p-value	standard error
Env 0 acc	Env 1 acc	0.71	-0.10	0.96	0.00	0.01
Env 0 acc	Env 2 acc	0.64	-0.47	0.91	0.00	0.01
Env 0 acc	Env 3 acc	0.64	0.14	0.90	0.00	0.01
Env 1 acc	Env 0 acc	0.75	-0.59	0.89	0.00	0.01
Env 1 acc	Env 2 acc	0.51	-0.67	0.71	0.00	0.01
Env 1 acc	Env 3 acc	0.71	-0.29	0.98	0.00	0.00
Env 2 acc	Env 0 acc	1.04	0.23	0.87	0.00	0.01
Env 2 acc	Env 1 acc	0.90	0.45	0.82	0.00	0.02
Env 2 acc	Env 3 acc	0.78	0.76	0.75	0.00	0.02
Env 3 acc	Env 0 acc	0.76	-0.79	0.83	0.00	0.01
Env 3 acc	Env 1 acc	0.80	-0.75	0.92	0.00	0.01
Env 3 acc	Env 2 acc	0.50	-0.83	0.64	0.00	0.02

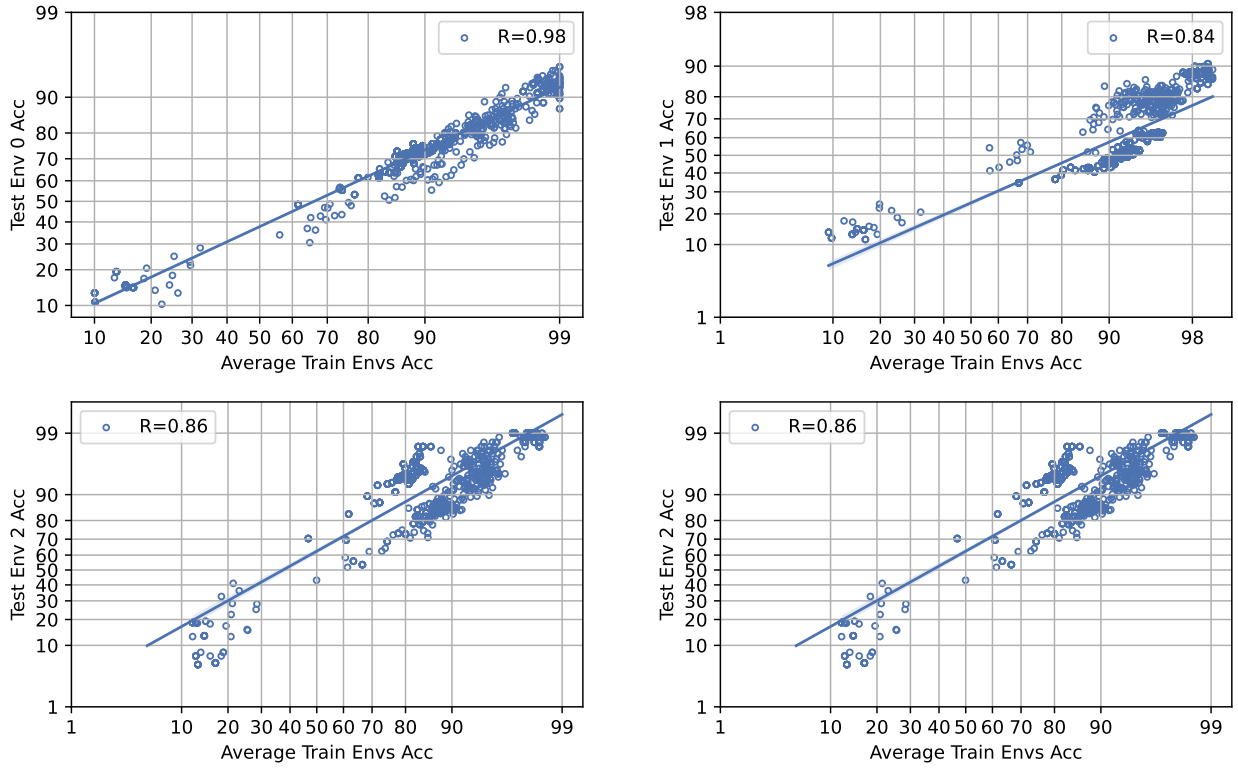


Figure 18: PACS: Average train Env Accuracy vs. Test Env Accuracy.

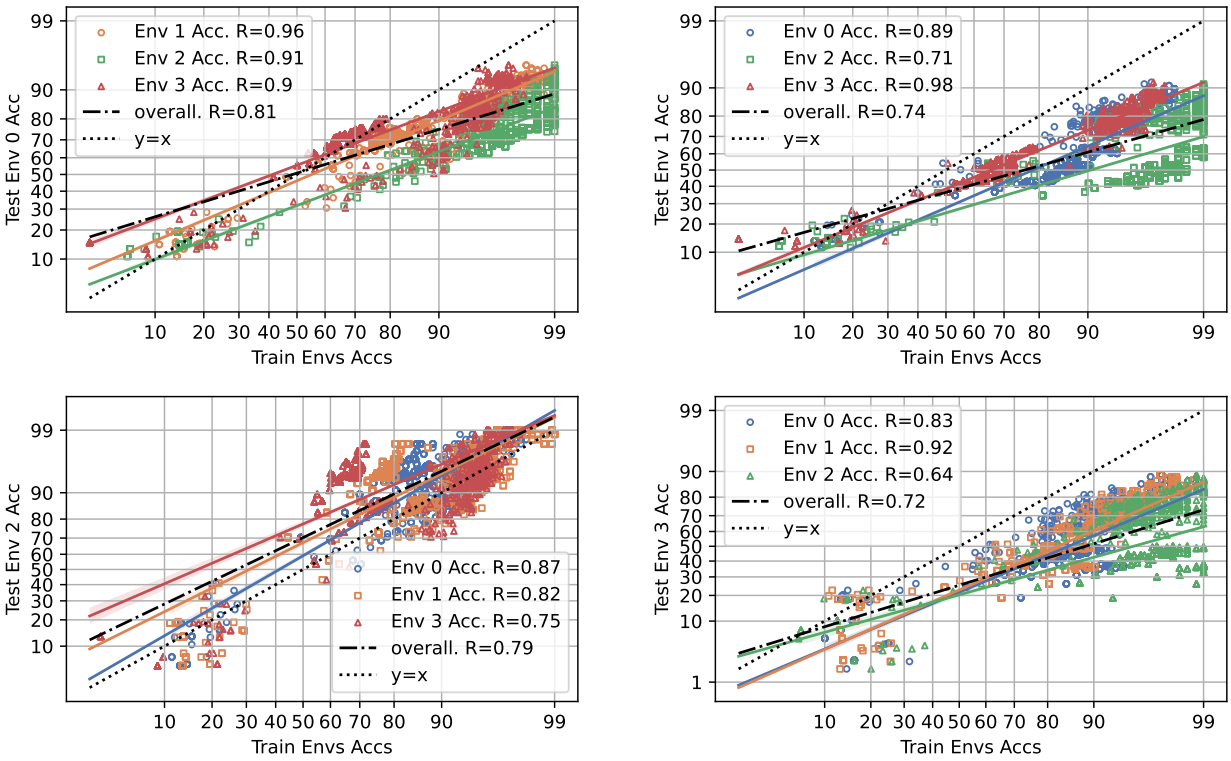


Figure 19: PACS: Train Env Accuracy vs. Test Env Accuracy.

C.5 Terra Incognita

Terra Incognita (Beery et al., 2018). A dataset that contains photographs of wild animals taken by camera traps at locations $d \in \{L100, L38, L43, L46\}$. This dataset contains 24,788 examples of dimensions (3, 224, 224) and 10 classes: Bird, Bobcat, Cat, Coyote, Dog, Empty, Fox, Horse, Mouse, Opossum, Rabbit, Raccoon, Rat, Skunk, Squirrel, Weasel.

Discussion. In general, we find that Terra Incognita does not strongly represent worst-case shifts for any split. Ahuja et al. (2021) consider Terra Incognita domain-general features to be fully informative, i.e., labels did not need to rely on spurious features such as the background to generate labels. Our results suggest that this benchmark may not accurately benchmark an algorithm's ability to give models free of spurious correlations. For Env 1, we observe that the slope of the line varies quite a bit. Particularly, there is a near-zero slope for models greater than 80% accuracy.

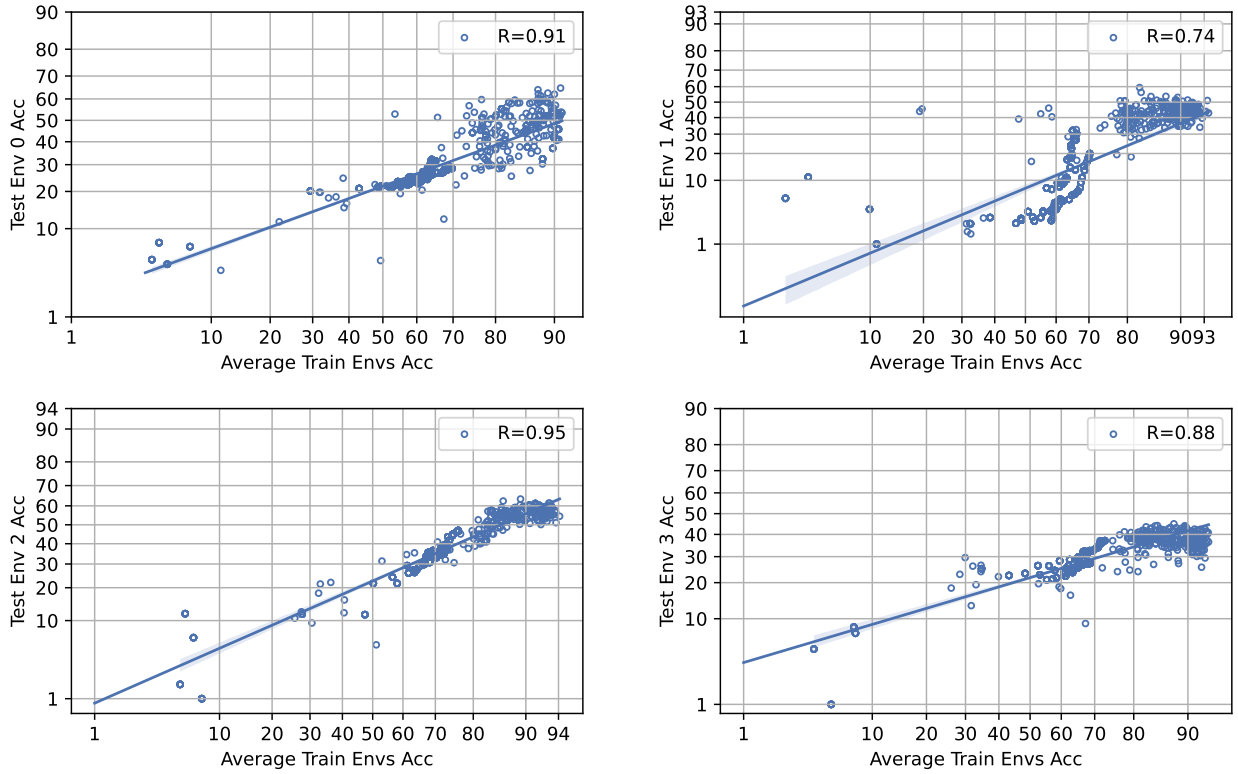


Figure 20: Terra Incognita: Average train Env Accuracy vs. Test Env Accuracy.

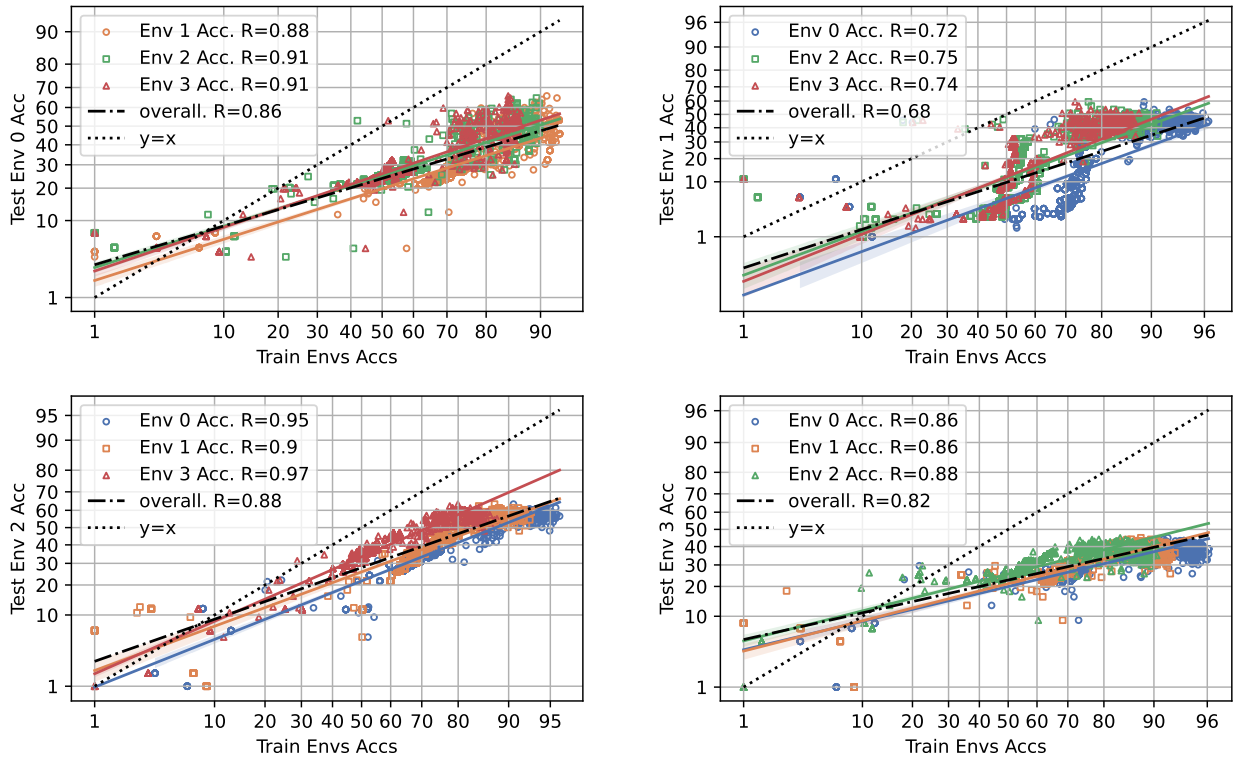


Figure 21: Terra Incognita: Train Env Accuracy vs. Test Env Accuracy.

C.6 Camelyon

Camelyon (Bandi et al., 2018; Koh et al., 2021). A dataset that contains histopathological images of lymph node tissue, collected from two hospitals, denoted as Hospital A, Hospital B. This dataset contains 327,680 examples of dimension (3, 96, 96) and 2 classes (tumor, non-tumor).

Discussion. We find that overall, there is a strong correlation between ID and OOD accuracy. However, we observe that for some ID/OOD splits, a regime of training accuracy has a negative correlation (environments 0 and 2), suggesting that within a certain accuracy range, these splits may be well-specified for benchmarking spurious correlations for models in the regime with negative correlation. This highlights the importance of qualitative evaluation as opposed to quantitative evaluation.

Table 17: WILDSCamelyon ID vs. OOD properties.

OOD	slope	intercept	Pearson R	p-value	standard error
Env 0 acc	0.78	0.33	0.90	0.00	0.01
Env 1 acc	0.71	-0.00	0.88	0.00	0.01
Env 2 acc	0.62	0.49	0.78	0.00	0.01
Env 3 acc	0.63	0.49	0.88	0.00	0.01
Env 4 acc	0.63	0.40	0.78	0.00	0.01

Table 18: WILDSCamelyon ID vs. OOD properties.

OOD	ID	slope	intercept	Pearson R	p-value	standard error
Env 0 acc	Env 1 acc	0.73	0.44	0.88	0.00	0.01
Env 0 acc	Env 2 acc	0.79	0.25	0.90	0.00	0.01
Env 0 acc	Env 3 acc	0.83	0.17	0.90	0.00	0.01
Env 0 acc	Env 4 acc	0.74	0.28	0.91	0.00	0.01
Env 1 acc	Env 0 acc	0.71	-0.00	0.89	0.00	0.01
Env 1 acc	Env 2 acc	0.69	0.02	0.85	0.00	0.01
Env 1 acc	Env 3 acc	0.74	-0.05	0.89	0.00	0.01
Env 1 acc	Env 4 acc	0.69	-0.03	0.89	0.00	0.01
Env 2 acc	Env 0 acc	0.64	0.41	0.81	0.00	0.01
Env 2 acc	Env 1 acc	0.59	0.58	0.74	0.00	0.01
Env 2 acc	Env 3 acc	0.67	0.37	0.79	0.00	0.01
Env 2 acc	Env 4 acc	0.61	0.44	0.81	0.00	0.01
Env 3 acc	Env 0 acc	0.66	0.41	0.90	0.00	0.01
Env 3 acc	Env 1 acc	0.57	0.64	0.84	0.00	0.01
Env 3 acc	Env 2 acc	0.63	0.47	0.87	0.00	0.01
Env 3 acc	Env 4 acc	0.61	0.45	0.88	0.00	0.01
Env 4 acc	Env 0 acc	0.63	0.35	0.79	0.00	0.01
Env 4 acc	Env 1 acc	0.60	0.49	0.74	0.00	0.01
Env 4 acc	Env 2 acc	0.62	0.39	0.79	0.00	0.01
Env 4 acc	Env 3 acc	0.67	0.29	0.78	0.00	0.01

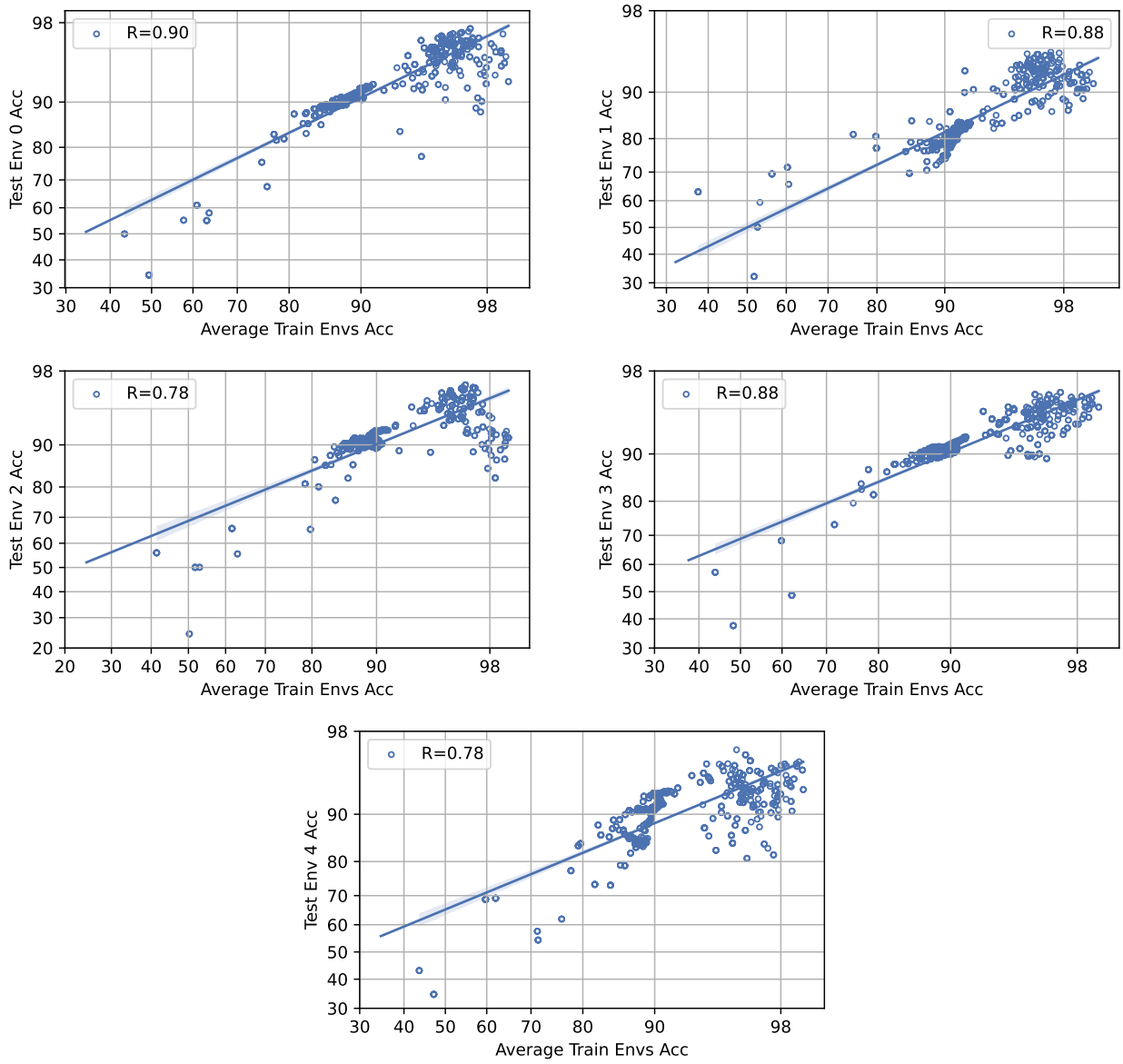


Figure 22: Camelyon: Average train Env Accuracy vs. Test Env Accuracy.

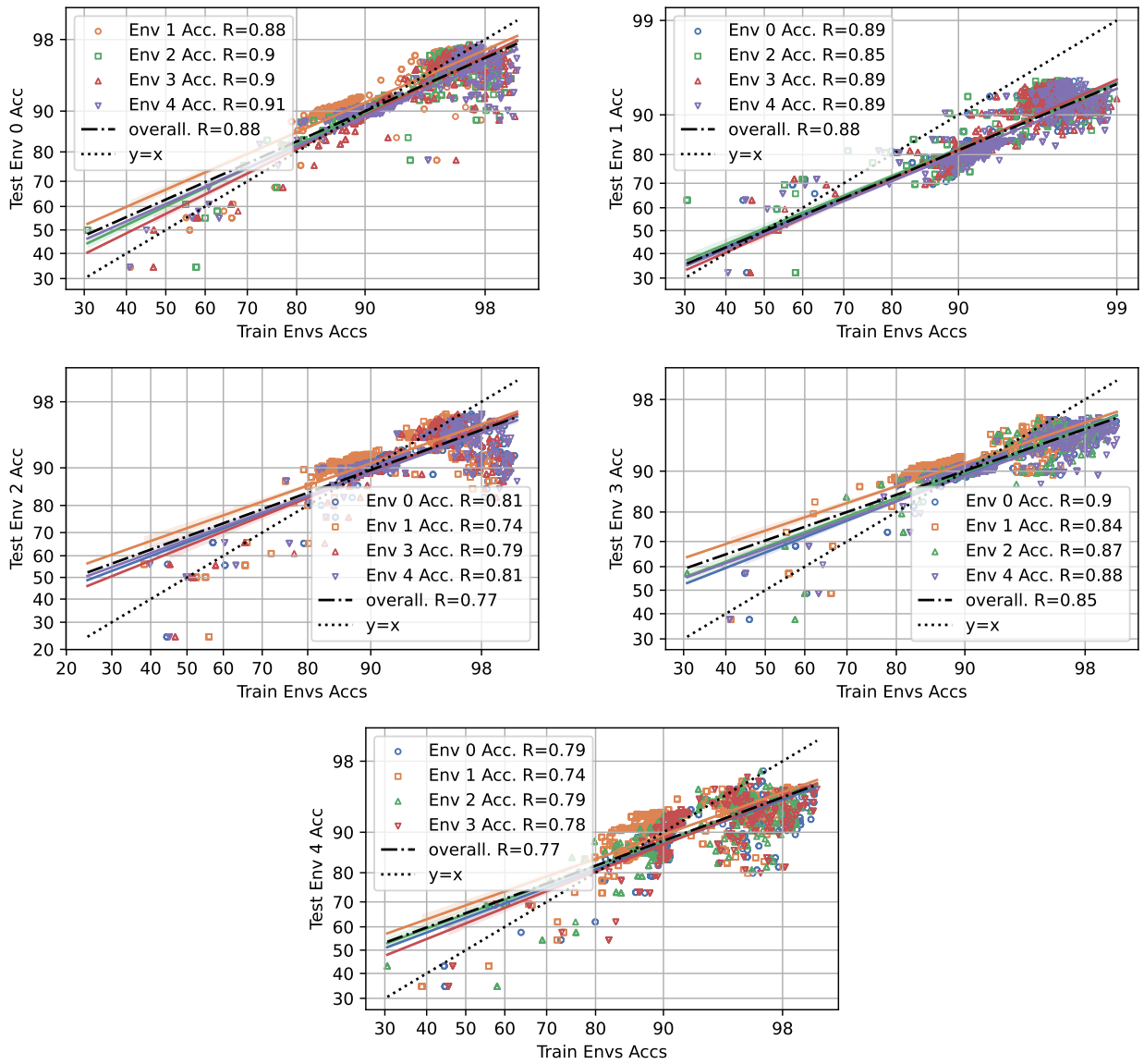


Figure 23: Camelyon: Train Env Accuracy vs. Test Env Accuracy.

C.7 Covid-CXR

Covid-CXR (Alzate-Grisales et al., 2022; Cohen et al., 2020b; Tabik et al., 2020; Tahir et al., 2021; Suwalska et al., 2023). A dataset that aggregates five different benchmark Covid-19 datasets, represented as the domain $d \in$ {Cov-Caldas (Columbia), Covid-Chest-X-Ray (Global), Covid-GR (Spain), Covid-Qu-Ex (Global), and PolCovid (Poland)}. These open-source Covid-19 datasets are collected from around the globe, accessible online. We are provided with *chest X-ray image (x)* and *binary diagnosis (y)*. The objective is to maintain consistent performance across different datasets d, which may incorporate data from singular or multiple sources.

Dataset	# Train	# Test	% Pos Train	% Neg Train	% Pos Test	% Neg Test
Cov-Caldas	3247	688	0.566	0.434	0.565	0.435
Covid-Chest-X-Ray	625	157	0.376	0.624	0.376	0.624
Covid-GR	681	171	0.501	0.499	0.497	0.503
Covid-Qu-Ex	21715	6788	0.353	0.647	0.353	0.647
PolCovid	4343	450	0.249	0.751	0.333	0.667

Table 19: Composition for Covid-19 Datasets.

- Cov-Caldas (Columbia, Alzate-Grisales et al. (2022)): This dataset was sourced from a single institution, S.E.S. Hospital Universitario de Caldas, located in the State of Caldas, Colombia. Labels were assigned based on positive results from conventional laboratory tests, such as PCR.
- Covid-Chest-X-Ray (Global, Cohen et al. (2020b)): This dataset, compiled from web sources, publications, and volunteer contributions, includes data on five types of pneumonia and Covid-19, along with metadata such as sex, age, and symptoms. The images are sourced from medical websites and are part of an open-source public project, where contributors can submit pull requests to add new images. The data is compiled from 138 unique locations.
- Covid-GR (Spain, Tabik et al. (2020)): In collaboration with four expert radiologists from Hospital Universitario Clínico San Cecilio in Granada, Spain, the authors developed a protocol for selecting and annotating chest X-ray (CXR) images for the dataset. A CXR image is labeled as Covid-19 positive if both the RT-PCR test and the radiologist's assessment confirm the diagnosis within 24 hours.
- Covid-Qu-Ex (Global, Tahir et al. (2021)): This dataset aggregates chest X-rays from six subdatasets, including the Covid-19 CXR dataset, RSNA CXR dataset (non-Covid infections and normal CXRs), Chest-Xray-Pneumonia dataset, PadChest dataset, Montgomery and Shenzhen CXR lung mask datasets, and QaTa-Cov19 CXR infection mask dataset. Designed to serve as a benchmark, it combines multiple publicly available datasets and repositories, which were originally dispersed and formatted differently. The authors performed quality control to ensure consistency. This dataset, specifically portions of their cited Covid-19 CXR dataset, overlaps with Covid-Chest-X-Ray (Env 1), thus deviating slightly from our standard experimental procedure of leave-one-domain-out.
- PolCovid (Poland, Suwalska et al. (2023)): Chest X-rays were collected from 15 Polish hospitals using a variety of devices and parameters due to differences in equipment between medical centers. The dataset includes patients with Covid-19, pneumonia, and healthy controls.

Experimental Details. Using the DomainBed suite by Gulrajani & Lopez-Paz (2020b), we employ the ResNet-50 architecture (with and without AugMix data augmentation), along with ResNet-18, DenseNet-121, and ConvNeXt-Tiny, on the Covid-CXR dataset. Each of the five datasets is treated as a distinct domain, d. For each model, we apply two pretrained variants using ImageNet: (i) Fine-tuned and (ii) Transfer learning.

Discussion. Covid-CXR is a real-world medical dataset containing chest X-ray images of Covid-19 patients, where spurious correlations emerge naturally due to the complex and multifaceted nature of medical data. These correlations are not explicitly observed or intentionally introduced but arise from underlying factors such as imaging artifacts, patient demographics, or co-occurring medical conditions that are often unmeasured or unaccounted for in the dataset. As observed, there exist strong, weak, and inverse correlations between ID and OOD performance. While environments 0, 2, and 4 are primarily positively correlated in OOD performance, environments 1 and 3 demonstrate the inverse accuracy-on-the-line phenomenon. Additionally, several observations show weak correlations, with slopes close to zero. Existing literature suggests that a horizontal line (i.e. slope of 0) is indicative of a severe distribution shift, where it prevents any meaningful transfer learning between the training data and the OOD data Teney et al. (2024). This could be attributed to the significantly more severe distribution shifts present in this dataset. These shifts, as discussed by Cohen et al. (2020a), may arise due to errors in labeling, discrepancies between institutions and radiologists, biases in clinical practices, and interobserver variability.

Moreover, Covid-Qu-Ex (Env 3), the largest and most diverse dataset of those explored in this work, demonstrates low OOD transfer accuracy. For OOD performance for ID Env 3, we observe slopes closest to 0, indicating a weak or near-zero correlation. The high ID accuracy (up to $\approx 99\%$) suggests that the model may be learning misleading features in OOD domains. However, for environments evaluated on this dataset, a strongly negative relationship is present, suggesting that improved ID performance may be associated with reduced OOD accuracy. In this dataset, the authors scale by concatenating existing chest X-Ray datasets. But, as discussed in Cohen et al. (2020a), Shen et al. (2024), and Teney et al. (2024), simply increasing the amount of data may not address the core issue of distribution shift, as more data could exacerbate overfitting to domain-specific artifacts or noise, rather than improving generalization. Shen et al. (2024) coins this as the *Data Addition Dilemma*, where adding data can both improve and worsen performance.

Table 20: CXR ID vs. OOD properties.

OOD	slope	intercept	Pearson R	p-value	standard error
Env 0 acc	0.54	-0.27	0.55	0.00	0.02
Env 1 acc	-0.38	0.13	-0.50	0.00	0.02
Env 2 acc	0.44	0.05	0.54	0.00	0.02
Env 3 acc	-0.60	0.56	-0.48	0.00	0.03
Env 4 acc	0.53	-0.04	0.31	0.00	0.04

Table 21: CXR ID vs. OOD properties.

OOD	ID	slope	intercept	Pearson R	p-value	standard error
Env 0 acc	Env 1 acc	0.56	-0.23	0.57	0.00	0.02
Env 0 acc	Env 2 acc	0.31	-0.21	0.41	0.00	0.02
Env 0 acc	Env 3 acc	0.16	-0.26	0.84	0.00	0.00
Env 0 acc	Env 4 acc	0.43	-0.39	0.68	0.00	0.01
Env 1 acc	Env 0 acc	-0.43	0.09	-0.61	0.00	0.01
Env 1 acc	Env 2 acc	-0.16	0.06	-0.27	0.00	0.01
Env 1 acc	Env 3 acc	-0.07	0.06	-0.51	0.00	0.00
Env 1 acc	Env 4 acc	-0.21	0.12	-0.46	0.00	0.01
Env 2 acc	Env 0 acc	0.39	0.07	0.54	0.00	0.01
Env 2 acc	Env 1 acc	0.36	0.06	0.46	0.00	0.02
Env 2 acc	Env 3 acc	0.09	0.06	0.61	0.00	0.00
Env 2 acc	Env 4 acc	0.28	-0.04	0.51	0.00	0.01
Env 3 acc	Env 0 acc	-0.68	0.54	-0.49	0.00	0.03
Env 3 acc	Env 1 acc	-0.47	0.51	-0.46	0.00	0.02
Env 3 acc	Env 2 acc	-0.38	0.51	-0.37	0.00	0.02
Env 3 acc	Env 4 acc	-0.26	0.54	-0.29	0.00	0.02
Env 4 acc	Env 0 acc	-0.03	0.19	-0.02	0.48	0.04
Env 4 acc	Env 1 acc	0.56	-0.01	0.40	0.00	0.03
Env 4 acc	Env 2 acc	0.47	-0.07	0.37	0.00	0.03
Env 4 acc	Env 3 acc	0.10	0.04	0.36	0.00	0.01

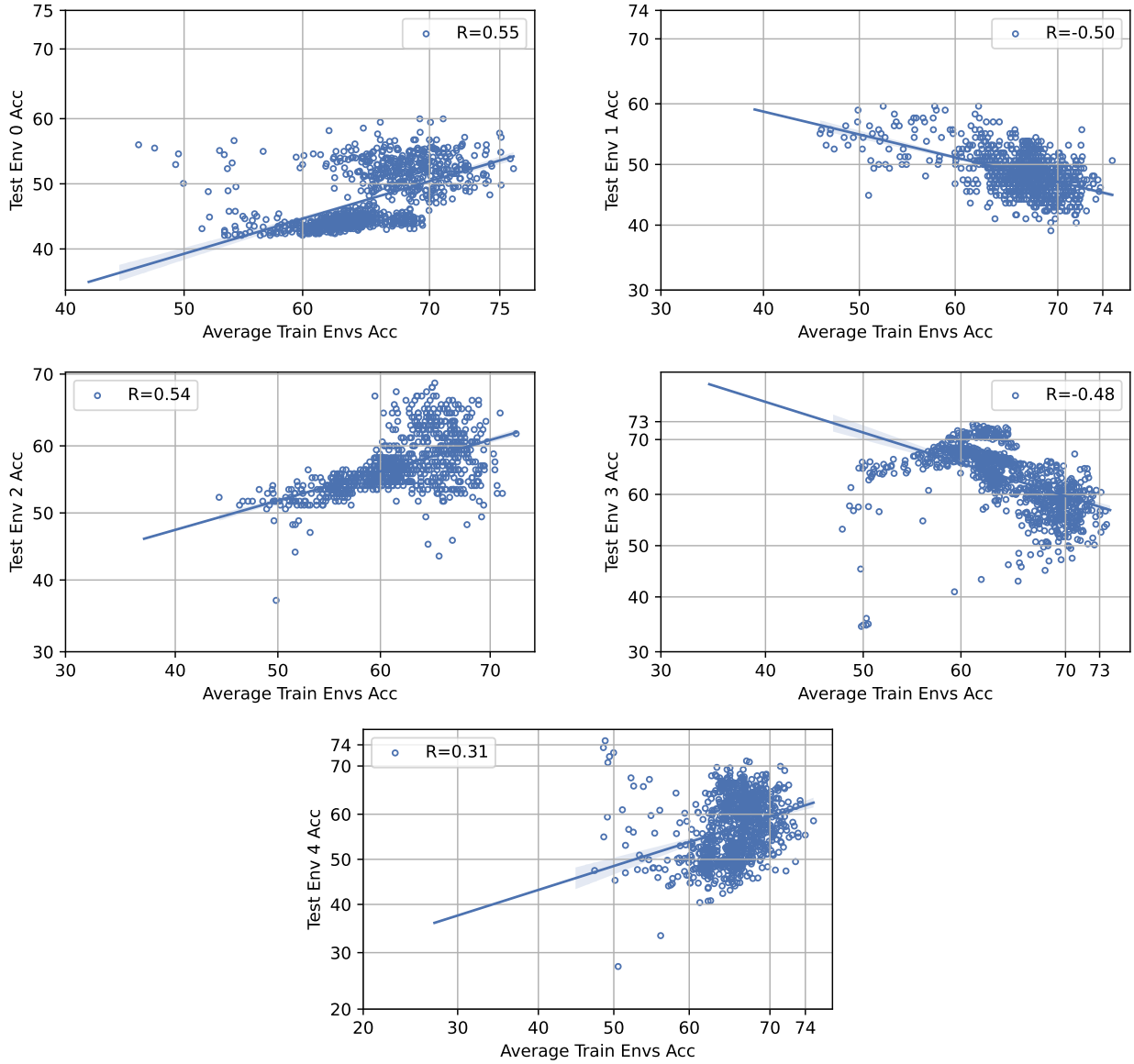


Figure 24: Covid-CXR: Average train Env Accuracy vs. Test Env Accuracy.

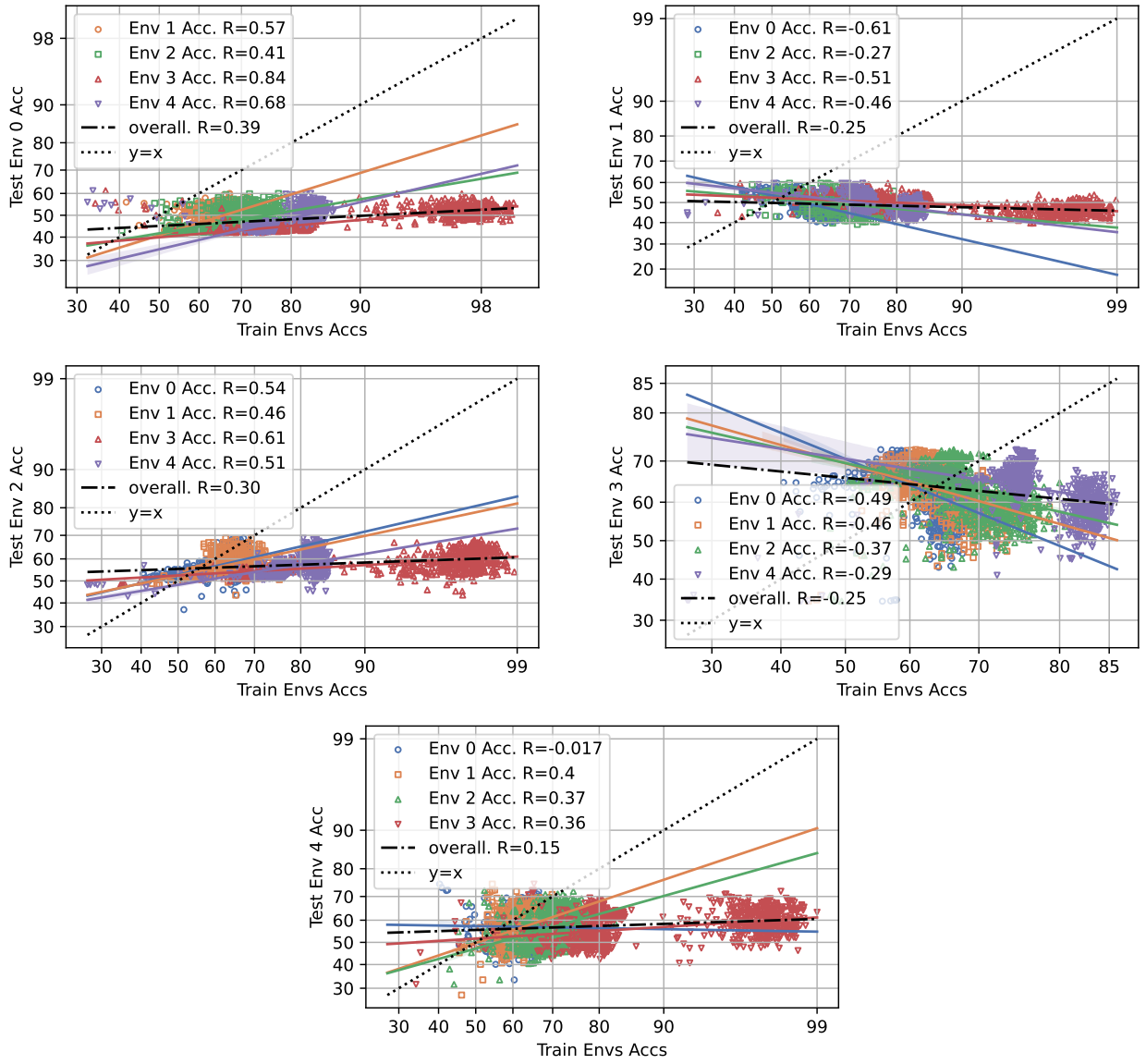


Figure 25: Covid-CXR: Train Env Accuracy vs. Test Env Accuracy.

C.8 FMoW

FMoW (Bandi et al., 2018; Koh et al., 2021). A dataset consisting of 141,696 RGB satellite images from 2022 - 2017 (resized to 224 x 224 pixels), where the label is one of 62 building or land use categories. This dataset simultaneously considers a domain generalization task, where two domains are defined by the year of image acquisition $t \in \{before2016, after2016\}$, and a subpopulation shift task, where the domains are denoted by the geographic region of the image $r \in \{Africa, Americas, Oceana, Asia, Europe\}$.

Discussion. We find that WILDFMoW has accuracy on the line for all splits, suggesting that this benchmark may be misspecified for benchmarking spurious correlations.

Table 22: WILDSFMoW ID vs. OOD properties.

OOD	slope	intercept	Pearson R	p-value	standard error
Env 0 acc	0.75	-0.62	0.98	0.00	0.00
Env 1 acc	0.78	-0.34	0.96	0.00	0.00
Env 2 acc	0.65	-0.48	0.94	0.00	0.00
Env 3 acc	0.83	-0.34	0.99	0.00	0.00
Env 4 acc	0.96	-0.18	0.99	0.00	0.00
Env 5 acc	0.76	-0.61	0.87	0.00	0.01

Table 23: WILDSFMoW ID vs. OOD properties.

OOD	ID	slope	intercept	Pearson R	p-value	standard error
Env 0 acc	Env 1 acc	0.78	-0.52	0.98	0.00	0.00
Env 0 acc	Env 2 acc	0.69	-0.72	0.97	0.00	0.00
Env 0 acc	Env 3 acc	0.71	-0.52	0.97	0.00	0.00
Env 0 acc	Env 4 acc	0.48	-0.89	0.96	0.00	0.00
Env 0 acc	Env 5 acc	0.43	-1.11	0.94	0.00	0.00
Env 1 acc	Env 0 acc	0.75	-0.26	0.97	0.00	0.00
Env 1 acc	Env 2 acc	0.78	-0.43	0.96	0.00	0.00
Env 1 acc	Env 3 acc	0.80	-0.20	0.98	0.00	0.00
Env 1 acc	Env 4 acc	0.53	-0.62	0.95	0.00	0.00
Env 1 acc	Env 5 acc	0.49	-0.88	0.94	0.00	0.00
Env 2 acc	Env 0 acc	0.58	-0.50	0.94	0.00	0.00
Env 2 acc	Env 1 acc	0.70	-0.47	0.93	0.00	0.00
Env 2 acc	Env 3 acc	0.64	-0.49	0.92	0.00	0.00
Env 2 acc	Env 4 acc	0.43	-0.80	0.91	0.00	0.00
Env 2 acc	Env 5 acc	0.39	-1.01	0.92	0.00	0.00
Env 3 acc	Env 0 acc	0.74	-0.30	0.98	0.00	0.00
Env 3 acc	Env 1 acc	0.91	-0.25	0.99	0.00	0.00
Env 3 acc	Env 2 acc	0.79	-0.49	0.97	0.00	0.00
Env 3 acc	Env 4 acc	0.53	-0.67	0.96	0.00	0.00
Env 3 acc	Env 5 acc	0.49	-0.96	0.92	0.00	0.00
Env 4 acc	Env 0 acc	0.89	-0.13	0.98	0.00	0.00
Env 4 acc	Env 1 acc	1.02	-0.08	0.99	0.00	0.00
Env 4 acc	Env 2 acc	0.92	-0.33	0.98	0.00	0.00
Env 4 acc	Env 3 acc	0.95	-0.08	0.99	0.00	0.00
Env 4 acc	Env 5 acc	0.54	-0.87	0.90	0.00	0.00
Env 5 acc	Env 0 acc	0.74	-0.56	0.89	0.00	0.01
Env 5 acc	Env 1 acc	0.82	-0.54	0.85	0.00	0.01
Env 5 acc	Env 2 acc	0.70	-0.74	0.84	0.00	0.01
Env 5 acc	Env 3 acc	0.74	-0.55	0.85	0.00	0.01
Env 5 acc	Env 4 acc	0.52	-0.90	0.86	0.00	0.00

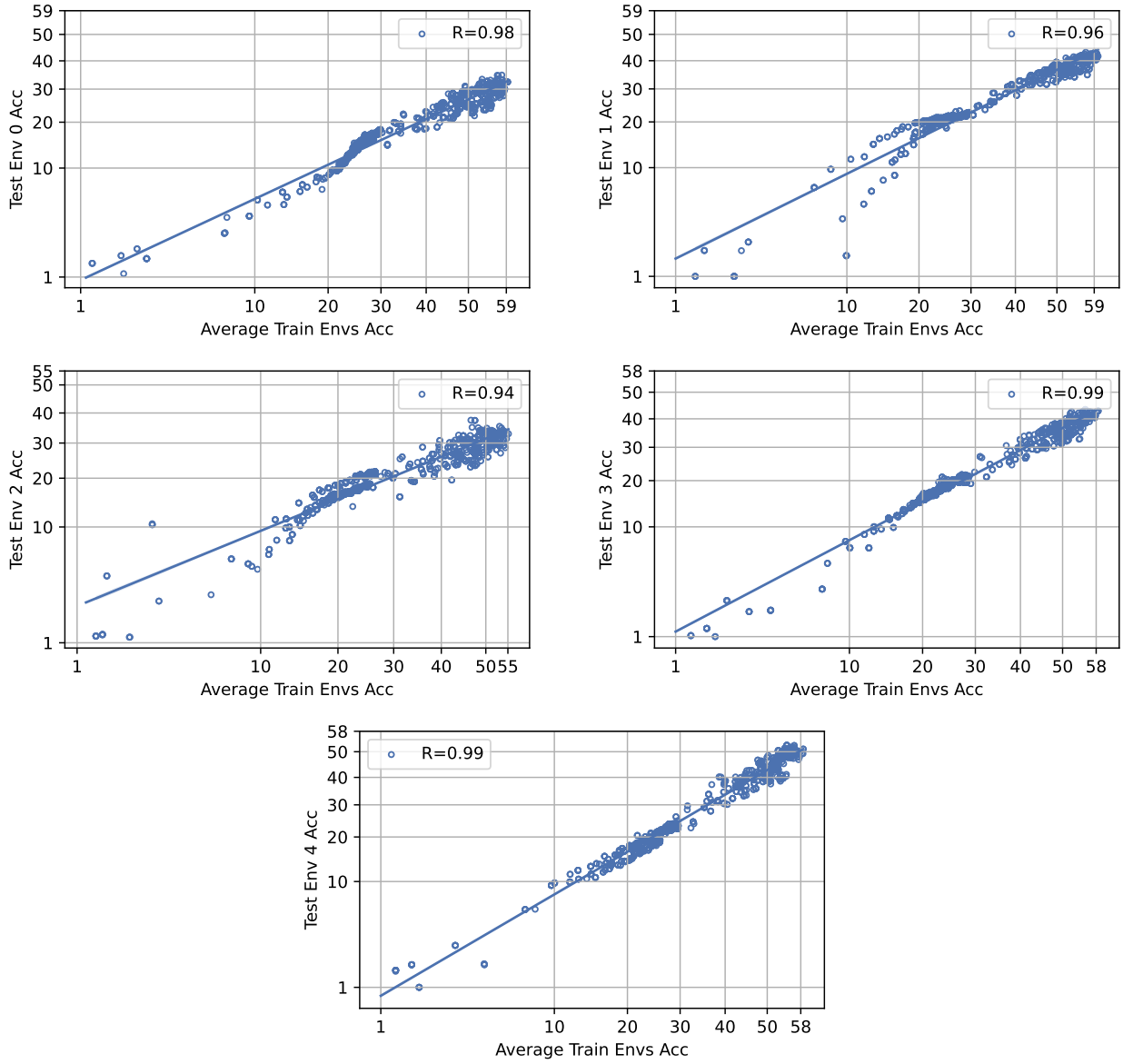


Figure 26: FMoW: Average train Env Accuracy vs. Test Env Accuracy.

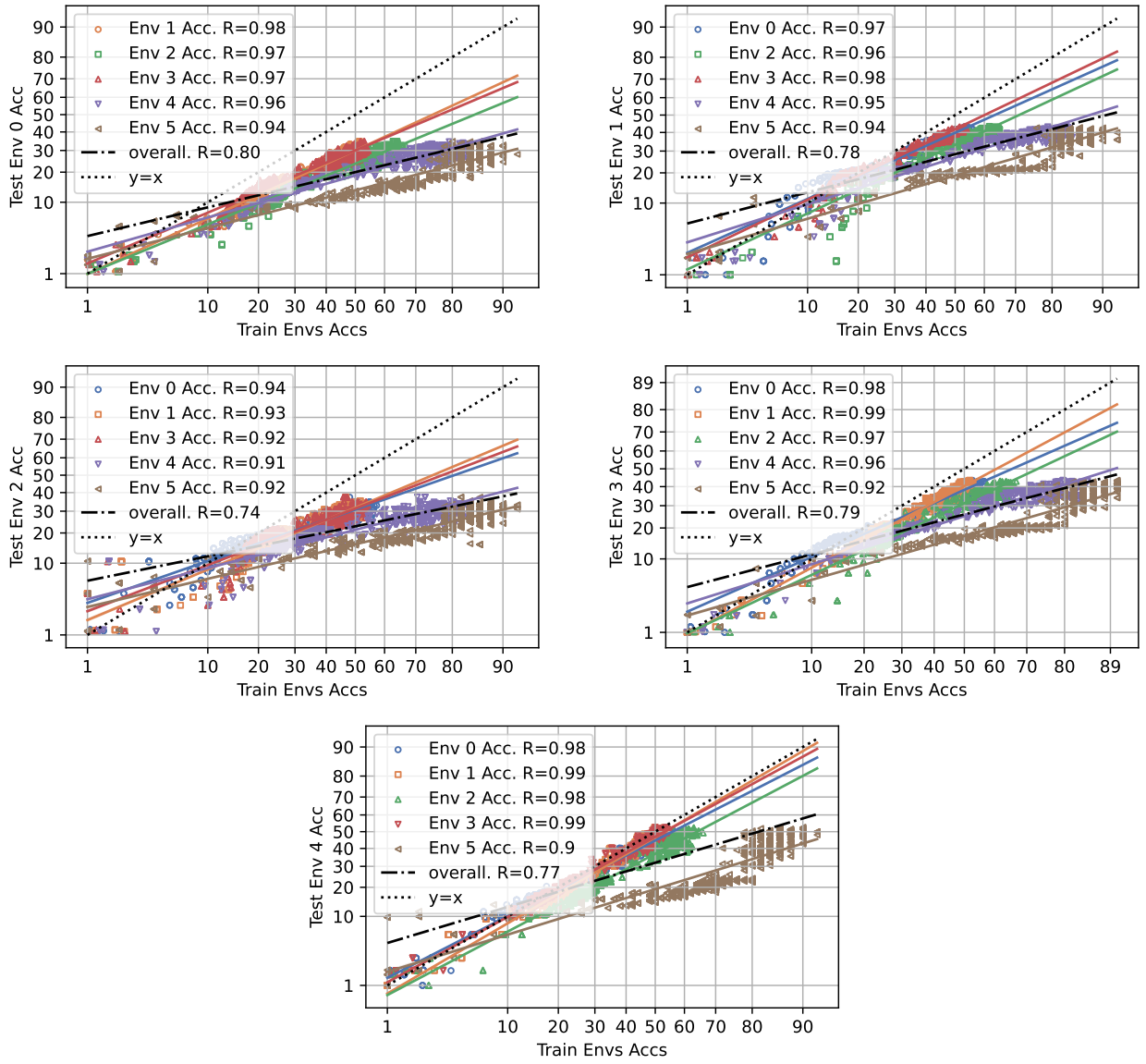


Figure 27: FMoW: Train Env Accuracy vs. Test Env Accuracy.

C.9 Waterbirds

Waterbirds (Sagawa et al., 2019; Koh et al., 2021). The Waterbirds dataset is a modification of the CUB dataset (Welinder et al., 2010) constructed to induce a subpopulation shift in the association between bird type $b \in \{waterbird, landbird\}$ in the foreground and the image background $c \in \{water, land\}$. Waterbird labels are assigned to seabirds and waterfowl; all other bird types are labeled as landbirds. Image backgrounds are obtained from the Places dataset (Zhou et al., 2016) and subset to include water backgrounds (categories: ocean or natural lake) and land backgrounds (categories: bamboo forest or broadleaf forest). The training set consists of 95% of all waterbirds with a water background and the remaining 5% with a land background. Similarly, 95% of all landbirds are displayed on a land background with the remaining 5% on a water background. The validation and test sets include an equal distribution of waterbirds and landbirds on each background. This dataset consists of 11,788 examples of size (3 x 224 x 224).

Experimental Details. Environment 0 consists of the full training split from the Waterbirds dataset and Environment 1 is a concatenation of the validation and test splits from the Waterbirds dataset (Sagawa et al., 2019). In addition to plotting the average ID vs. OOD accuracies for each test environment, we also include average ID vs. group-specific accuracies and pairwise combinations of group-specific ID vs. group-specific OOD accuracy.

Discussion. In our evaluation of average ID and OOD performance, we find that the Waterbirds dataset does not strongly represent worst-case shifts. However, plotting group-specific OOD accuracies, we find no linear correlation for Environment 1 group $(y = 0, a = 1)$, representing a degradation in worst-group accuracy (WGA) with improvements to Environment 0 ID average accuracy. We examine the pairwise group-specific accuracies for each environment and generally find that OOD group accuracy positively correlates with ID accuracy for groups sharing the same label. Since Waterbirds groups are defined by label (bird), attribute (background) pairs, this result is consistent with the studies that suggest worst-class accuracy (WCA) is a good proxy for WGA when group membership is unknown (Yang et al., 2023). Similarly, OOD group accuracy negatively correlates with ID accuracy for groups of the opposite label, demonstrating the trade-off between majority and minority group/class performance under subpopulation shifts.

Table 24: WILDSWaterbirds average ID vs. OOD properties.

ID	OOD	slope	intercept	Pearson R	p-value	standard error
Env 0 acc	Env 1 acc	0.83	0.35	0.92	0.00	0.01
Env 1 acc	Env 0 acc	0.47	0.16	0.69	0.00	0.02

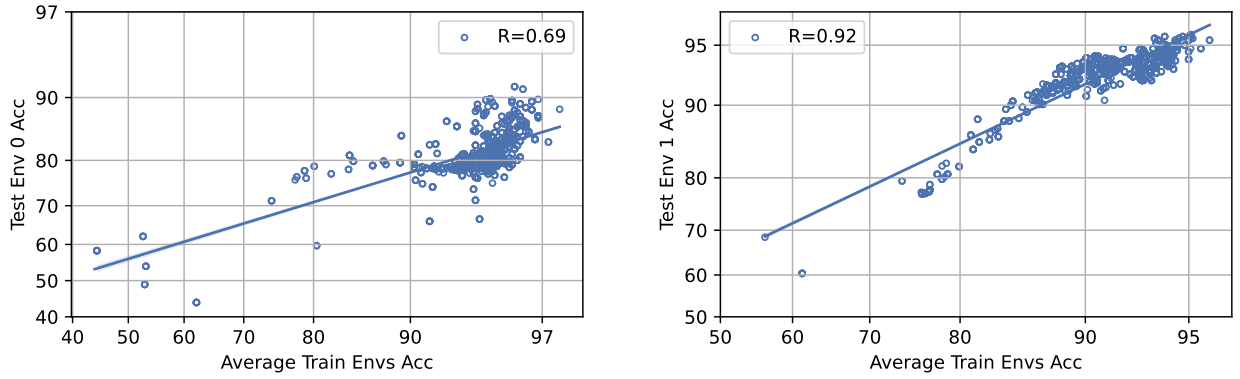


Figure 28: Waterbirds average ID vs. average OOD accuracies

Table 25: Waterbirds average ID vs. group-specific OOD properties.

ID	OOD	slope	intercept	Pearson R	p-value	standard error
Env 0 avg acc	Env 1 y=0,a=0 acc	-0.13	1.58	-0.13	0.00	0.03
Env 0 avg acc	Env 1 y=0,a=1 acc	0.00	1.32	0.00	0.95	0.03
Env 0 avg acc	Env 1 y=1,a=0 acc	0.22	1.17	0.84	0.00	0.00
Env 0 avg acc	Env 1 y=1,a=1 acc	0.32	1.13	0.81	0.00	0.01
Env 1 avg acc	Env 0 y=0,a=0 acc	0.70	-0.02	0.65	0.00	0.03
Env 1 avg acc	Env 0 y=0,a=1 acc	-0.07	1.61	-0.11	0.00	0.02
Env 1 avg acc	Env 0 y=1,a=0 acc	0.27	1.63	0.56	0.00	0.01
Env 1 avg acc	Env 0 y=1,a=1 acc	0.30	1.21	0.60	0.00	0.01

Table 26: Waterbirds pairwise group-specific ID vs. OOD properties.

ID	OOD	slope	intercept	Pearson R	p-value	standard error
Env 0 y=0,a=0 acc	Env 1 y=0,a=0 acc	0.76	0.58	0.86	0.00	0.01
Env 0 y=0,a=1 acc	Env 1 y=0,a=0 acc	0.72	0.73	0.73	0.00	0.02
Env 0 y=1,a=0 acc	Env 1 y=0,a=0 acc	-0.15	2.19	-0.43	0.00	0.01
Env 0 y=1,a=1 acc	Env 1 y=0,a=0 acc	-0.13	2.17	-0.36	0.00	0.01
Env 0 y=0,a=0 acc	Env 1 y=0,a=1 acc	0.45	1.05	0.42	0.00	0.03
Env 0 y=0,a=1 acc	Env 1 y=0,a=1 acc	0.65	0.72	0.54	0.00	0.03
Env 0 y=1,a=0 acc	Env 1 y=0,a=1 acc	-0.03	1.96	-0.06	0.08	0.01
Env 0 y=1,a=1 acc	Env 1 y=0,a=1 acc	-0.05	1.99	-0.10	0.00	0.01
Env 0 y=0,a=0 acc	Env 1 y=1,a=0 acc	-1.16	3.05	-0.33	0.00	0.10
Env 0 y=0,a=1 acc	Env 1 y=1,a=0 acc	-0.96	2.58	-0.25	0.00	0.12
Env 0 y=1,a=0 acc	Env 1 y=1,a=0 acc	1.24	0.13	0.93	0.00	0.02
Env 0 y=1,a=1 acc	Env 1 y=1,a=0 acc	1.34	0.12	0.92	0.00	0.02
Env 0 y=0,a=0 acc	Env 1 y=1,a=1 acc	-0.70	2.02	-0.32	0.00	0.07
Env 0 y=0,a=1 acc	Env 1 y=1,a=1 acc	-0.87	2.24	-0.34	0.00	0.08
Env 0 y=1,a=0 acc	Env 1 y=1,a=1 acc	0.84	0.19	0.95	0.00	0.01
Env 0 y=1,a=1 acc	Env 1 y=1,a=1 acc	0.92	0.20	0.97	0.00	0.01
Env 1 y=0,a=0 acc	Env 0 y=0,a=0 acc	0.77	0.55	0.96	0.00	0.01
Env 1 y=0,a=1 acc	Env 0 y=0,a=0 acc	0.16	2.18	0.35	0.00	0.01
Env 1 y=1,a=0 acc	Env 0 y=0,a=0 acc	-0.07	2.23	-0.18	0.00	0.01
Env 1 y=1,a=1 acc	Env 0 y=0,a=0 acc	0.02	2.26	0.05	0.10	0.01
Env 1 y=0,a=0 acc	Env 0 y=0,a=1 acc	0.54	-0.49	0.39	0.00	0.04
Env 1 y=0,a=1 acc	Env 0 y=0,a=1 acc	0.73	0.23	0.88	0.00	0.01
Env 1 y=1,a=0 acc	Env 0 y=0,a=1 acc	-0.13	0.61	-0.23	0.00	0.02
Env 1 y=1,a=1 acc	Env 0 y=0,a=1 acc	-0.50	1.25	-0.69	0.00	0.02
Env 1 y=0,a=0 acc	Env 0 y=1,a=0 acc	-0.33	0.50	-0.18	0.00	0.05
Env 1 y=0,a=1 acc	Env 0 y=1,a=0 acc	-0.40	-0.01	-0.32	0.00	0.04
Env 1 y=1,a=0 acc	Env 0 y=1,a=0 acc	0.61	0.17	0.84	0.00	0.01
Env 1 y=1,a=1 acc	Env 0 y=1,a=0 acc	0.88	-1.21	0.83	0.00	0.02
Env 1 y=0,a=0 acc	Env 0 y=1,a=1 acc	0.05	1.08	0.03	0.38	0.05
Env 1 y=0,a=1 acc	Env 0 y=1,a=1 acc	-0.50	1.50	-0.47	0.00	0.03
Env 1 y=1,a=0 acc	Env 0 y=1,a=1 acc	0.44	1.44	0.60	0.00	0.02
Env 1 y=1,a=1 acc	Env 0 y=1,a=1 acc	0.94	0.13	0.96	0.00	0.01

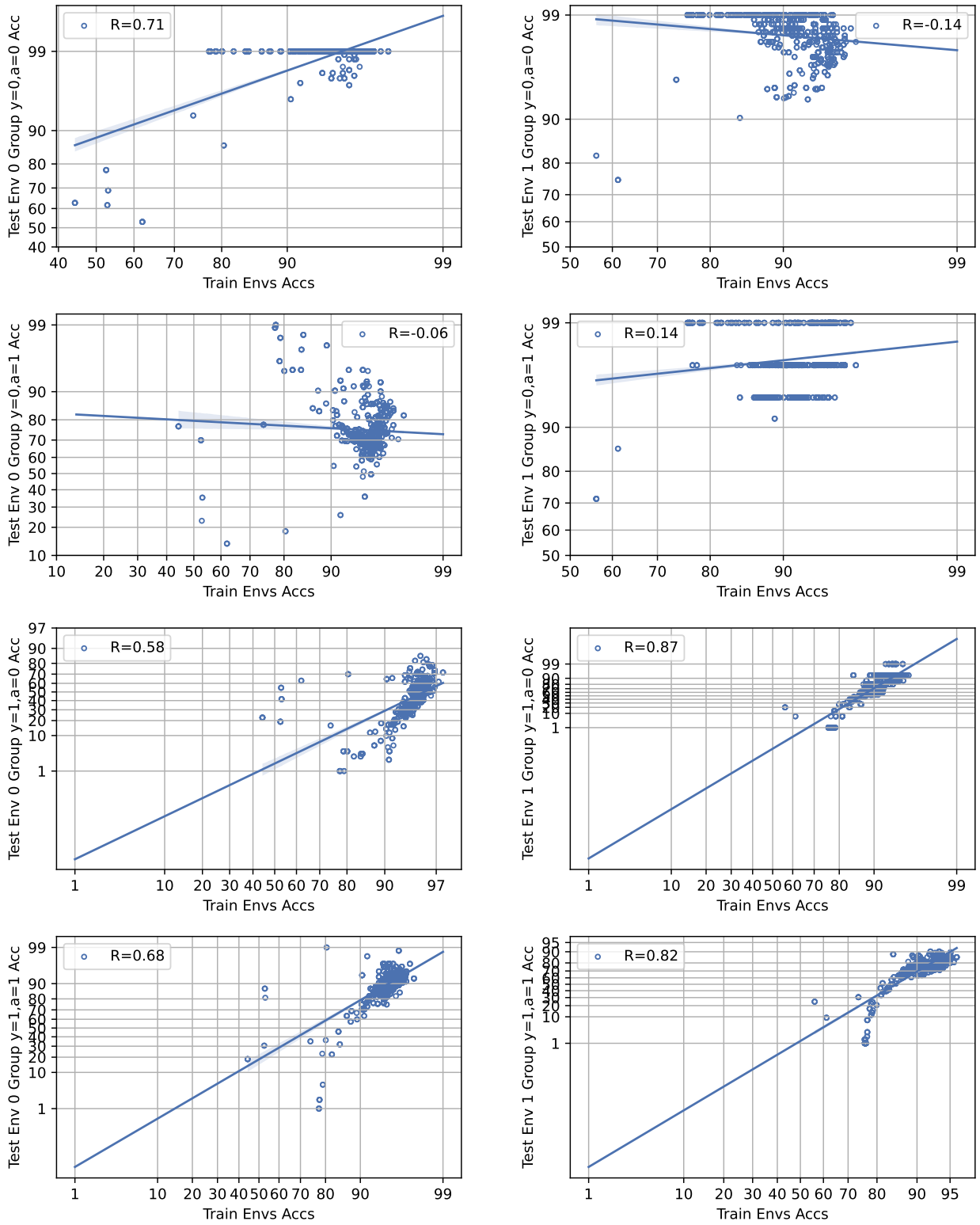


Figure 29: Waterbirds average ID vs. group-specific OOD accuracies

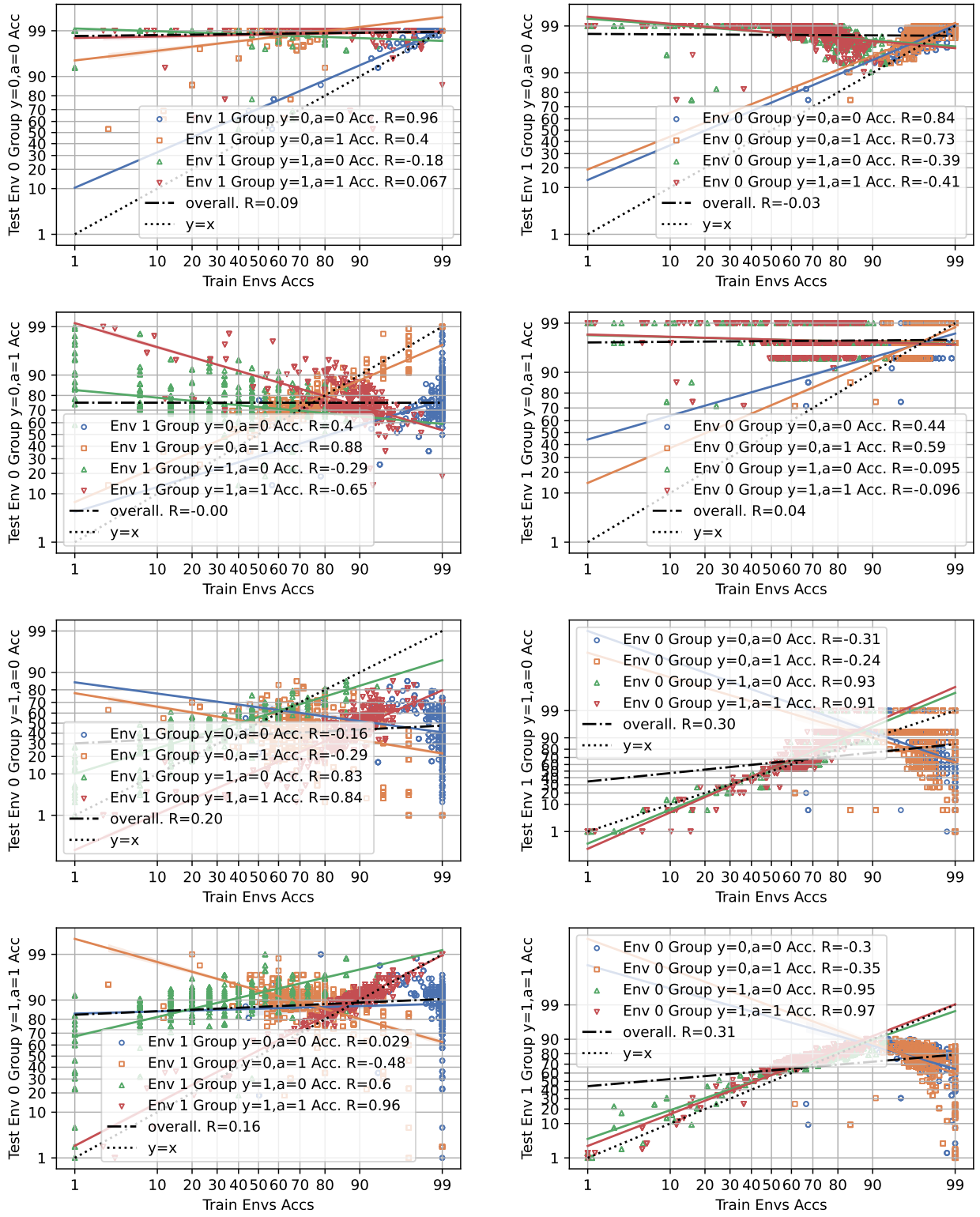


Figure 30: Waterbirds pairwise group-specific accuracies

C.10 CivilComments

CivilComments (Koh et al., 2021) A dataset comprised of multiple subpopulations, which correspond to different demographic identities. The domain d is a multi-dimensional binary vector with 8 dimensions, each corresponding to whether a given comment mentions one of the 8 demographic identities: *male, female, LGBTQ, Christian, Muslim, other religions, Black*, and *White*. For each of the 8 identities, we form 2 groups corresponding to the toxicity label, generating a total of 16 groups (e.g., male_toxic, male_non-toxic, etc.). By construction, since each comment may belong to more than one group, this experimental procedure differs slightly from the standard subpopulation shift framework discussed in this work. Regardless, the experimental results with and without group overlaps display similar accuracy on the line and accuracy on the inverse line patterns.

Experimental Details. We leverage both BERT and DistilBERT architectures for the CivilComments dataset.

Discussion. The CivilComments dataset exhibits both an accuracy on a line and an accuracy on the inverse line phenomenon across all test environments in the Leave-One-Domain-Out procedure. The spurious correlation, which involves the presence or absence of 8 different demographic identities (e.g., male, white, Christian, Muslim, etc.), is rather strongly correlated with the toxicity label. Empirically, the correlation coefficient (R) values range from as low as -0.47 to as high as 0.43. There appear to be many ID/OOD splits that can be derived from this dataset for benchmarking domain generalization.

Table 27: CivilComments ID vs. OOD properties.

OOD	slope	intercept	Pearson R	p-value	standard error
Env 0 acc	0.36	-0.29	0.28	0.00	0.04
Env 1 acc	-0.49	0.16	-0.47	0.00	0.03
Env 2 acc	0.39	-0.17	0.28	0.00	0.04
Env 3 acc	-0.49	0.12	-0.42	0.00	0.03
Env 4 acc	0.58	0.20	0.43	0.00	0.04
Env 5 acc	-0.54	0.00	-0.43	0.00	0.04
Env 6 acc	0.46	-0.04	0.30	0.00	0.05
Env 7 acc	-0.50	0.16	-0.43	0.00	0.03
Env 8 acc	0.49	0.03	0.36	0.00	0.04
Env 9 acc	-0.49	0.06	-0.41	0.00	0.03
Env 10 acc	0.53	0.17	0.34	0.00	0.05
Env 11 acc	-0.57	-0.09	-0.46	0.00	0.03
Env 12 acc	0.46	0.05	0.30	0.00	0.05
Env 13 acc	-0.54	-0.04	-0.45	0.00	0.03
Env 14 acc	0.41	-0.10	0.29	0.00	0.04
Env 15 acc	-0.52	0.06	-0.43	0.00	0.04

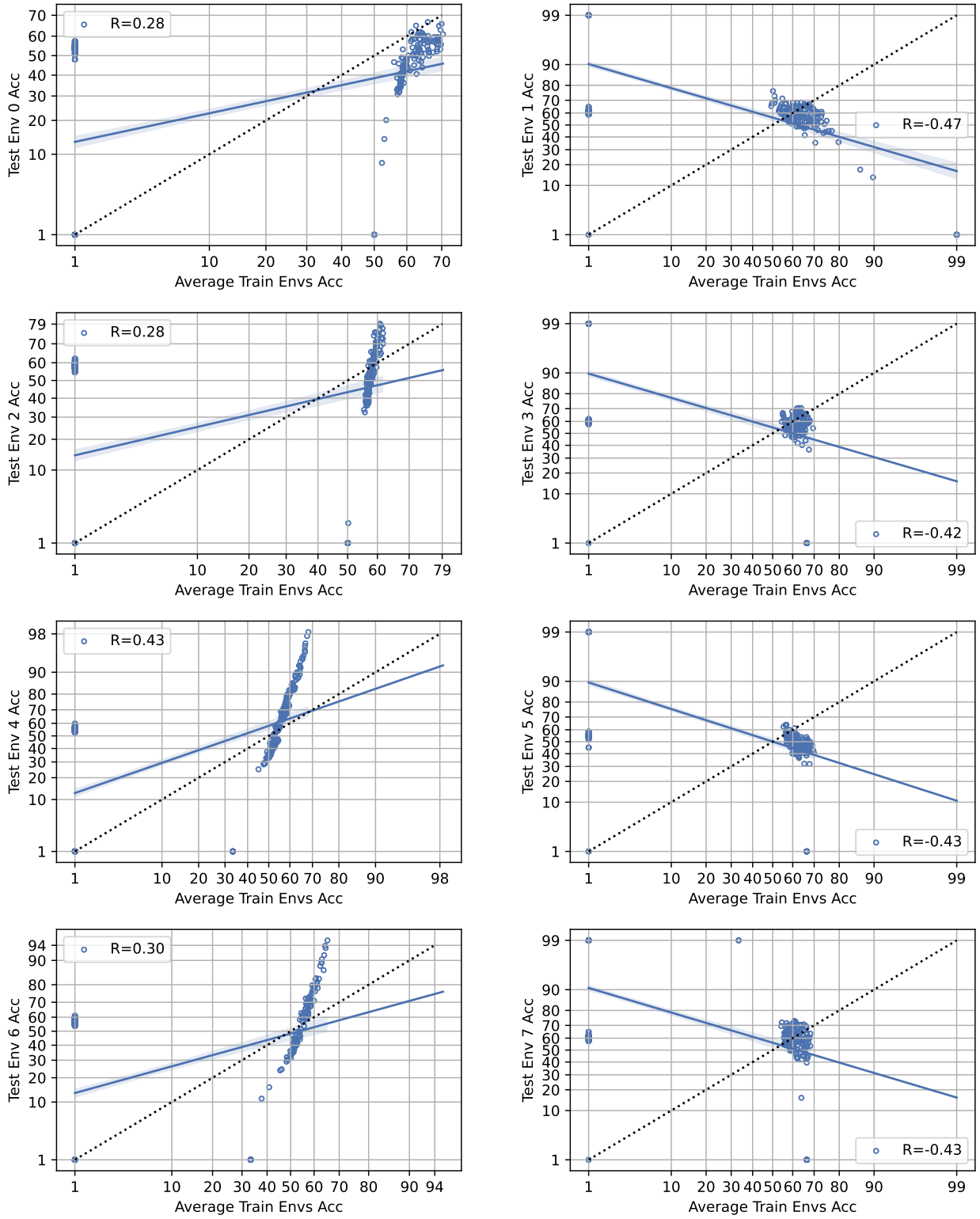


Figure 31: CivilComments: Average train Env Accuracy vs. Test Env Accuracy.

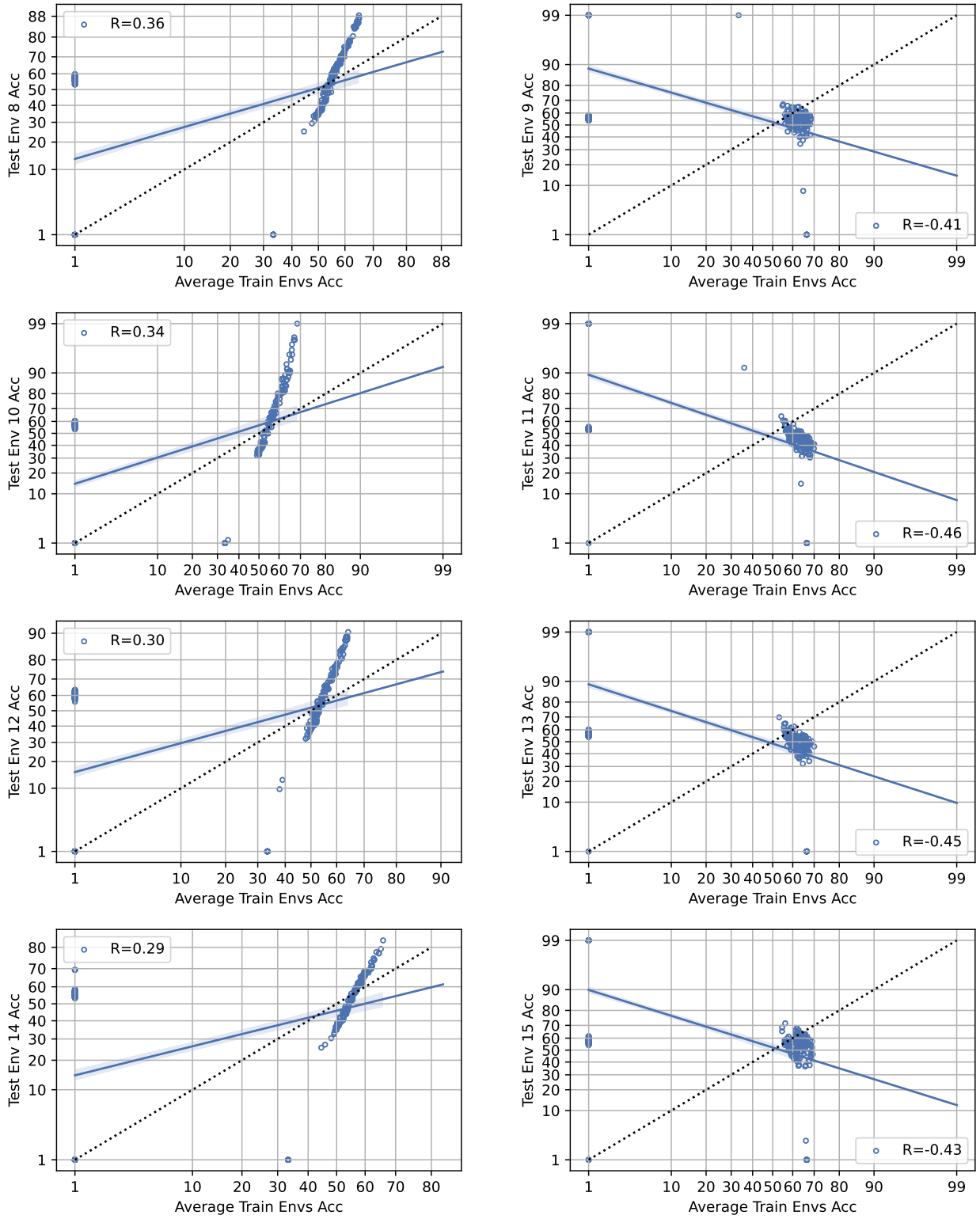


Figure 32: CivilComments: Average train Env Accuracy vs. Test Env Accuracy.

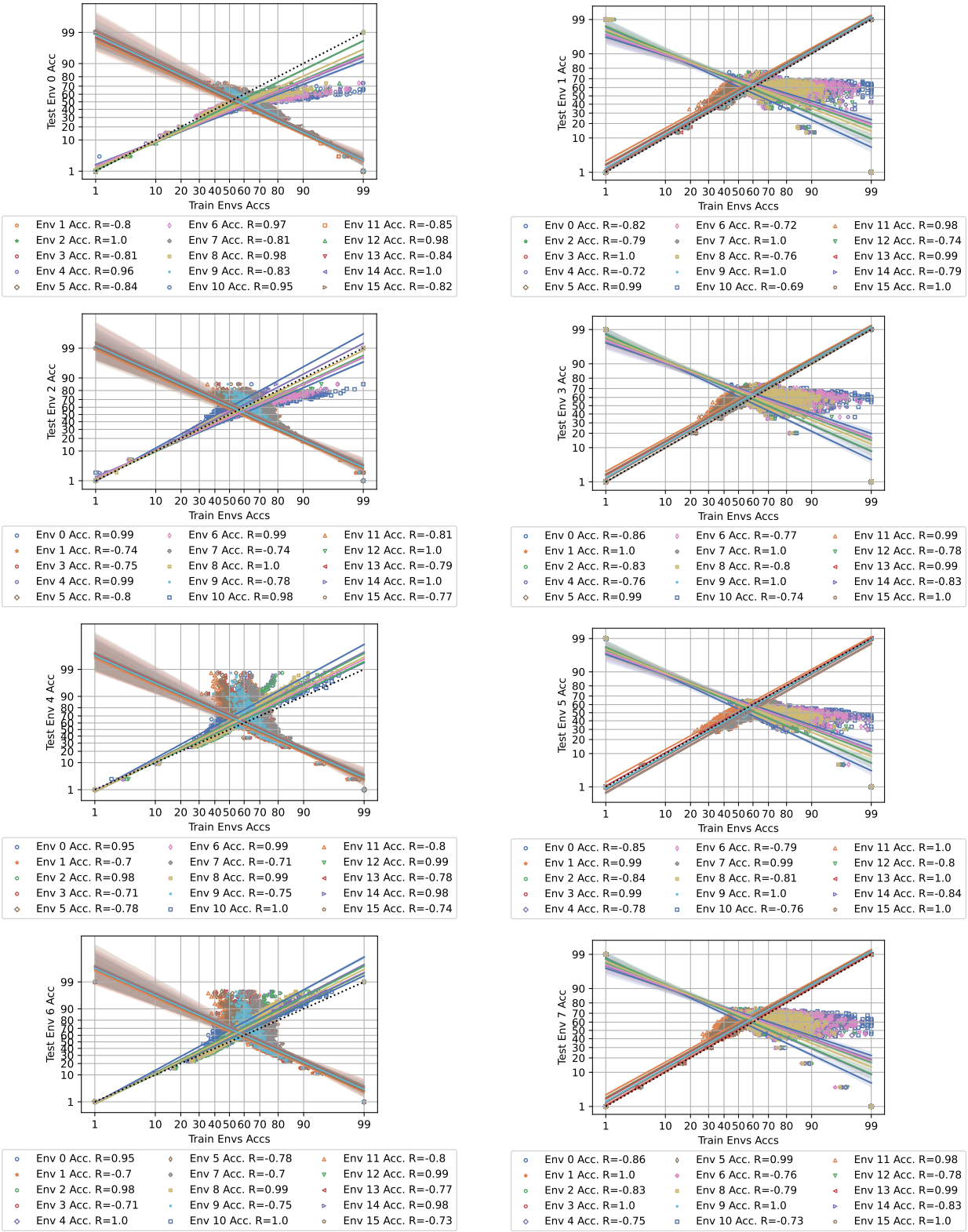


Figure 33: CivilComments: Train Env Accuracy vs. Test Env Accuracy.

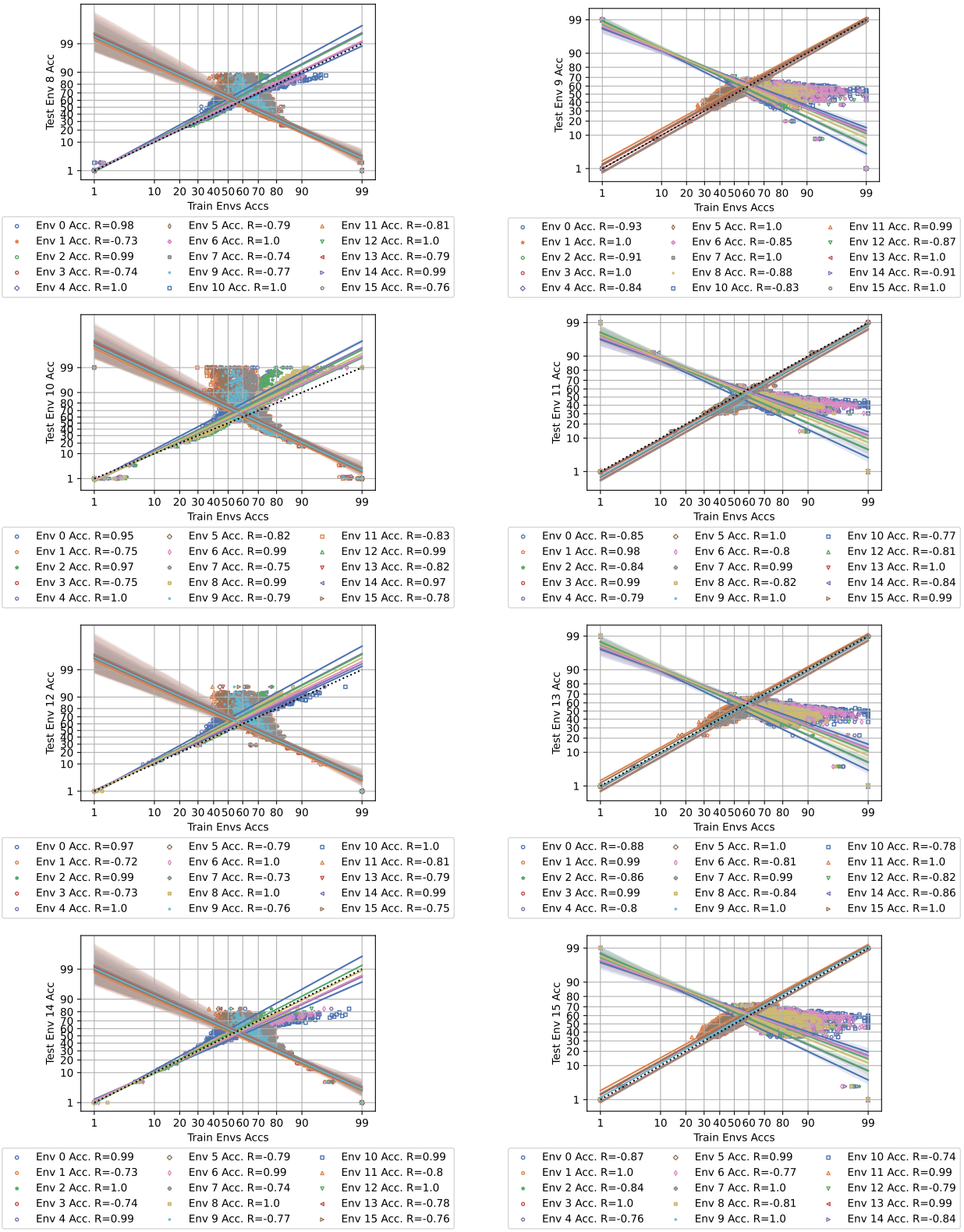


Figure 34: CivilComments: Train Env Accuracy vs. Test Env Accuracy.

D Benchmarking Causal Representation Learning

Models that incorporate or learn structural knowledge of the domains they are applied to have been shown to be more efficient and generalize better across different settings (Parascandolo et al., 2020; Sanchez-Gonzalez et al., 2020; Gondal et al., 2019; Goyal et al., 2019; Battaglia et al., 2016; Bapst et al., 2019; Makar et al., 2022; Zheng & Makar, 2022; Wang et al., 2022a). An example of such a structure is the principle of *independent causal mechanisms* (Haavelmo, 1944; Aldrich, 1989; Hoover, 1990; Pearl, 2009; Schölkopf et al., 2012; Janzing et al., 2012; Peters et al., 2016; Schölkopf et al., 2021), which posits that the generative process of a system's variables consists of autonomous components, or mechanisms, that operate independently and do not inform one another. This implies that the conditional distribution of each variable, given its causes (mechanisms), is independent of the other variables and mechanisms (Peters et al., 2017). Learning causal representations is an active area of research (Schölkopf et al., 2021). Datasets for causal representation learning are primarily (semi)parametric where (some) causal variables are known and potentially intervenable (Von Kügelgen et al., 2021; Lippe et al., 2022a;b; Ahmed et al., 2020; Liu et al., 2023b). Then, success is assessed by how well learned disentangled representations (mechanisms) explain outcome variance, using R^2 or MCC (Mathew's Correlation Coefficient) (Lopez et al., 2023). However, the task of causal representation learning with complex datasets with limited knowledge or control over generative mechanisms remains a challenge, especially without requiring most (or at least some) relevant causal variables to be directly observed Lopez-Paz et al. (2017)—we identify that benchmarking causal representation learning in this setting is also challenging.

Causal representation learning is closely tied to domain generalization, which aims to learn representations from multiple observed domains that give predictors whose performance is invariant to new domains (new data distributions). Many works in domain generalization (Arjovsky et al., 2019; Salaudeen & Koyejo, 2022; 2024; Mahajan et al., 2021; Liu et al., 2021a; Lv et al., 2022; Chen et al., 2022a; Eastwood et al., 2022). have been motivated by the principle of independent causal predictors, which aims to identify causal predictors from observational data by searching for feature sets that maintain stable (invariant) predictive accuracy across interventional distributions Peters et al. (2016); Heinze-Deml et al. (2018); Arjovsky et al. (2019). Additionally, more recent work motivates learning causal representations from multiple datasets arising from unknown interventions (von Kügelgen et al., 2024).

Thus, one may naively consider domain generalization as a proxy task to benchmark causal representation learning in more complex settings. This work studies when performance on a domain generalization task is informative of the causal representation learning task. Specifically, when benchmarking a set of models, including a disentangled causal model, based on transfer accuracy, our results on evaluating domain generalization apply, where non-causal correlations are spurious (Salaudeen et al., 2024).