

# Understanding Implications of Dataset Choice for Feature Effect Estimation: A Simulation-Based Investigation through Error Decomposition

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**Abstract.** The abstract should briefly summarize the contents of the paper in 150–250 words.

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## 1 Introduction

Most Machine Learning (ML) models can be considered black boxes — opaque systems that intrinsically do not allow insight into their internal reasoning, making it often impossible to explain their decisions. However, this can be problematic in many domains and applications, such as the healthcare, legal, or finance sectors, where decisions must be transparent and accountable [1].

Interpretability is crucial to enhance trust [29, 30], address potential biases [13], fairness and ethical concerns [18], and ensure compliance with regulations such as the EU’s General Data Protection Regulation (GDPR) [5] and AI Act [6]. To address these challenges, the field of Explainable AI / Interpretable ML has emerged [1]. Although there are many different methods<sup>1</sup>, we will focus on feature effect methods like *Partial Dependence Plots (PDP)* [9] and *Accumulated Local Effects (ALE)* [2].

Due to the severity of many applications, it is crucial to utilize these explainability methods correctly. In general, there are many pitfalls to be aware of [27], including whether to compute explanations in-sample, i.e. on training data, or out-of-sample, which we refer to as validation data in the following. In loss-based methods such as *Permutation Feature Importance (PFI)* [4, 7], this choice is crucial and has already been studied (e.g., in [27]). Nevertheless, other explainability methods, such as the *Mean Decrease in Impurity (MDI)* (or *Gini Importance*) of Random Forests, or *SHAP* values [20, 21], have also been found to exhibit biases when computed on training data.

However, to the best of our knowledge, there exist no similar studies for feature effect methods like PDP and ALE. Existing works, including the original papers of PDP and ALE, rely on training data without further justification (e.g., [2, 9, 25]). In contrast, practitioners often advocate for using unseen

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<sup>1</sup> For an overview of Explainable AI methods, see e.g. [1, 17, 25].

test/validation data<sup>2</sup>, or base their choice on practical constraints like dataset size<sup>3</sup>. While the training set is usually larger and might thus lead to less variance in the feature effect estimates, a too large dataset can increase computation times substantially, particularly for the PDP [9]. On the other hand, although feature effects are not based on generalization error like PFI, it is not clear how much they are affected by overfitting or distribution shifts between training and validation data.

In this paper, we aim to answer this largely unaddressed, fundamental methodological question of whether to use training or validation data to compute feature effects. We perform an empirical simulation study, comparing feature effect error and uncertainty for PDP and ALE across training data, validation data, and cross-validation scenarios, considering various data scenarios and model types. Our main contributions in this paper are as follows.

1. We shed light on the question of whether to compute feature effects on training data, validation data, or in a cross-validated manner, grounded through our comprehensive simulation study, considering feature effect error, bias, and variance across different models and data scenarios.
2. We define methods to quantify feature effect error and uncertainty in a theoretical framework, building upon previous work in this area.
3. We provide an overview of commonly used test functions for simulation studies in Interpretable ML, including applications and purposes.

These contributions have several important implications: Our empirically grounded recommendations enable practitioners and researchers to make informed decisions about which dataset to select for feature effect computation, helping to understand potential implications of their choices. Additionally, our framework for evaluating feature effects and our systematic collection of test functions provide a foundation for future research in Interpretable ML, with the latter specifically facilitating test function choice for simulation studies.

The remainder of this paper is structured as follows. In SECTION 2, we introduce the considered feature effect methods PDP and ALE, as well as related works on feature effect error and uncertainty, and give an overview of common test functions for simulation studies. In SECTION ??, we extend existing works and give our definitions of feature effect errors and uncertainty. We then describe the methodology and set-up of our simulation studies in SECTION 4, present the results in SECTION 5, and discuss their implications and limitations in SECTION ??. In SECTION 6, we briefly conclude our work.

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<sup>2</sup> see, e.g. <https://github.com/SauceCat/PDPbox/issues/68> (10/27/2024)

<sup>3</sup> see, e.g. <https://forums.fast.ai/t/partial-dependence-plot/98465> (10/27/2024)

## 2 Background & Related Work

### 2.1 Feature Effects

The *Partial Dependence Plot (PDP)* by Friedman [9] describes the marginal effect of one or two features on the prediction of a model  $\hat{f}$ . For a feature set  $X_S$  (with  $S \subseteq \{1, \dots, p\}$ ,  $|S| = 1$  or  $|S| = 2$ ), the PDP is defined as

$$PDP_{\hat{f},S} = \mathbb{E}_{X_C}[\hat{f}(x_S, X_C)] = \int f(x_S, x_C) d\mathbb{P}(x_C), \quad (1)$$

where  $X_C$  is the complement feature subset.  $PDP_{\hat{f},S}$  is a function of  $x_S$  and can be estimated by Monte Carlo integration:

$$\widehat{PDP}_{\hat{f},S}(x_S) = \frac{1}{n} \sum_{i=1}^n \hat{f}(x_S, x_C^{(i)}). \quad (2)$$

Here,  $x_C^{(i)}$  are the actual complement feature values from the dataset of  $n$  instances. To plot this function, a grid of  $G$  grid points  $\{(x_S^{(g)}, \widehat{PDP}_{\hat{f},S}(x_S^{(g)}))\}_{g=1}^G$  can be used [26]. Molnar et al. [27] recommend using quantile-based over equidistant grids.

The PDP assumes that the features in  $S$  are independent of the features in  $C$ . If this is violated, the perturbations may produce unrealistic data points outside the underlying joint distribution of the data. This extrapolation issue can cause misleading interpretations [25, 27].

The *Accumulated Local Effects (ALE)* plot is an alternative to the PDP that solves the extrapolation issue [2]. Using the notation above, for  $|S| = 1$ , the ALE plot is defined as

$$ALE_{\hat{f},S}(x_S) = \int_{x_{\min,S}}^{x_S} \mathbb{E}_{X_C|X_S} [\hat{f}^S(X_S, X_C) | X_S = z_S] dz_S - \text{constant} \quad (3)$$

$$= \int_{x_{\min,S}}^{x_S} \int_{x_C} \hat{f}^S(z_S, x_C) \mathbb{P}(x_C | z_S) dx_C dz_S - \text{constant}, \quad (4)$$

where  $\hat{f}^S(x_S, x_C) = \frac{\partial \hat{f}(x_S, x_C)}{\partial x_S}$ . The constant is chosen so that  $\widehat{ALE}_{\hat{f},S}(X_S)$  is centered with a mean of 0 w.r.t. the marginal distribution of  $X_S$ . The uncentered ALE can be estimated by

$$\widehat{\widehat{ALE}}_{\hat{f},S}(x) = \sum_{k=1}^{k_S(x)} \frac{1}{n_S(k)} \sum_{i: x_S^{(i)} \in N_S(k)} [\hat{f}(z_{k,S}, x_C^{(i)}) - \hat{f}(z_{k-1,S}, x_C^{(i)})]. \quad (5)$$

Here,  $\{N_S(k) = (z_{k-1,S}, z_{k,S}]\}_{k=1}^K$  partitions the samples  $\{x_S^{(i)}\}_{i=1}^n$  into  $K$  intervals or neighborhoods  $N_S(k)$ .  $n_S(k)$  denotes the number of observations in

the  $k$ th interval  $N_S(k)$ ,  $k_S(x)$  represents the index of the interval to which a particular value  $x$  of feature  $x_S$  belongs. The uncentered ALE is centered by

$$\widehat{ALE}_{f,S}(x) = \widehat{ALE}_{f,S}(x) - \frac{1}{n} \sum_{i=1}^n \widehat{ALE}_{f,S}(x_S^{(i)}) \quad (6)$$

to have a mean effect of 0. For the grid that defines the intervals, the quantiles of the empirical distribution of  $\{x_S^{(i)}\}_{i=1}^n$  can be used [2].

Other feature effect methods include the *M-Plot (Marginal Plot)*, which, however, suffers from the omitted variable bias [2, 9], or *functional ANOVA (fANOVA)*, which decomposes feature effects into main and interaction effects [16].

Furthermore, methods exist that extend previous ones, such as *Robust and Heterogeneity-aware ALE (RHALE)* [11], or *Accumulated Total Derivative Effect (ATDEV)* plots, which can be decomposed into ALE and *Accumulated Cross Effects (ACE)* [19].

In addition to these global feature effect methods, there are regional effect plots such as *REPID* [15], as well as local methods, including *ICE (Individual Conditional Expectation)* curves [12] or SHAP dependence plots (e.g., [25]) based on SHAP values [22].

For the remainder of this paper, we will focus on PDP and ALE as the most popular global feature effect methods and refer to them when speaking of feature effects.

## 2.2 Feature Effect Error Decomposition

To quantify the error of a computed feature effect, a “ground truth” needs to be defined first. We follow the approach of Molnar et al. [26] and define ground truth versions of PDP and ALE directly on the data generating process (DGP) by applying PDP and ALE to the underlying ground truth function  $f$  (instead of model  $\hat{f}$ ).

For PDP, we can directly use the definition of Molnar et al. [26]:

**Definition 1 (Definition 1 from [26]).** *The PDP ground truth is the PDP applied to function  $f : \mathcal{X} \rightarrow \mathcal{Y}$  of the data generating process.*

$$PDP_{f,S}(x_S) = \mathbb{E}_{X_C}[f(x_S, X_C)] \quad (7)$$

As stated by Molnar et al. [26], their results also apply to conditional variants of the PDP such as ALE. We now make this definition explicit:

**Definition 2.** *The ALE ground truth is the ALE applied to function  $f : \mathcal{X} \rightarrow \mathcal{Y}$  of the data generating process.*

$$ALE_{f,S}(x_S) = \int_{x_{min,S}}^{x_S} \mathbb{E}_{X_C|X_S}[f^S(X_S, X_C)|X_S = z_S] dz_S - \text{constant} \quad (8)$$

where  $f^S(x_S, x_C) = \frac{\partial f(x_S, x_C)}{\partial x_S}$  and constant chosen such that the effect has a mean of 0 w.r.t. the marginal distribution of  $X_S$ .

Note that different ground truth effects may also be defined, and our choices come with certain implications and limitations, such as omitting the *aggregation bias*<sup>4</sup> [24].

With more complex ground truth functions  $f$ , it may become increasingly difficult to derive the ground truth feature effects analytically, especially for ALE. In these cases, we therefore propose to also estimate those effects by Monte Carlo integration, yielding  $\widehat{PDP}_{f,S}(x_S)$  and  $\widehat{ALE}_{f,S}(x_S)$  (obtained by plugging in  $\hat{f}$  instead of  $\hat{f}$  into the estimators in equations (2) and (5) / (6)).

Summarizing, we have now defined four quantities per feature effect:  $PDP_{f,S}$ ,  $\widehat{PDP}_{f,S}$ ,  $PDP_{\hat{f},S}$ , and  $\widehat{PDP}_{\hat{f},S}$  (analogue for ALE). We can now define different errors between each of these quantities. In this paper, we focus on the MSE, as it can be decomposed into bias and variance (see e.g. [10]). Taking, for example,  $PDP_{f,S}$  as ground truth, we can define the MSE of  $PDP_{\hat{f},S}$  at a point  $x_S$  as follows [26]:

$$\text{MSE}(x_S; PDP_{f,S}, PDP_{\hat{f},S}) = \mathbb{E}_F[(PDP_{f,S}(x_S) - \widehat{PDP}_{\hat{f},S}(x_S))^2] \quad (9)$$

$$= \underbrace{(PDP_{f,S}(x_S) - \mathbb{E}_F[PDP_{\hat{f},S}(x_S)])^2}_{\text{Bias}^2} + \underbrace{\text{Var}_F[PDP_{\hat{f},S}(x_S)]}_{\text{Variance}} \quad (10)$$

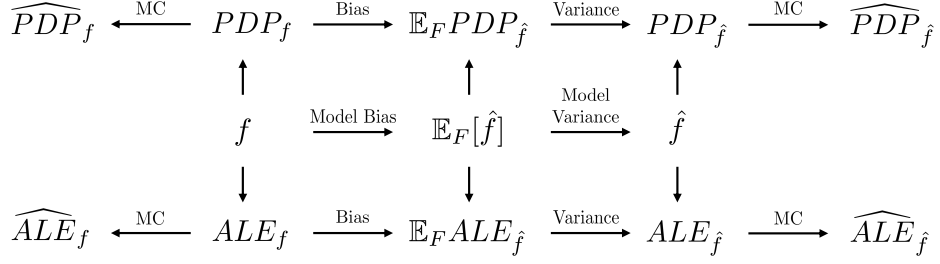
Here,  $F$  denotes the distribution of trained models. The bias is linked to the bias of the model, the variance comes from the variance in the model fits (randomness in training data, randomness in model training procedure).

Since we usually cannot determine  $PDP_{\hat{f},S}$ , we need to estimate it by Monte Carlo integration, yielding  $\widehat{PDP}_{\hat{f},S}(x_S)$ . This, however, introduces an additional variance term (we use the random variable  $X_{mc}$  for the Monte Carlo samples (e.g., training or validation data)):

$$\begin{aligned} \text{MSE}(x_S; PDP_{f,S}, \widehat{PDP}_{\hat{f},S}) &= \underbrace{(PDP_{f,S}(x_S) - \mathbb{E}_F[PDP_{\hat{f},S}(x_S)])^2}_{\text{Bias}^2} \\ &\quad + \underbrace{\text{Var}_F[PDP_{\hat{f},S}(x_S)]}_{\text{Variance}} + \underbrace{\mathbb{E}_F \text{Var}_{X_{mc}}[\widehat{PDP}_{\hat{f},S}(x_S)]}_{\text{MC-Variance}} \end{aligned} \quad (11)$$

*Proof.* For better readability, we will omit the subscript  $S$  as well as the point  $x_S$  and use  $X = X_{mc}$  in this proof:

<sup>4</sup> for details, see [15]



**Fig. 1.** Error chain in feature effect estimation (modified, original version can be found in [26])

$$\begin{aligned}
\text{MSE}(PDP_f, \widehat{PDP}_{\hat{f}}) &= \mathbb{E}_F \mathbb{E}_X [(PDP_f - \widehat{PDP}_{\hat{f}})^2] \\
&= \mathbb{E}_F \mathbb{E}_X [PDP_f^2 - 2PDP_f \widehat{PDP}_{\hat{f}} + \widehat{PDP}_{\hat{f}}^2] \\
&= PDP_f^2 - 2PDP_f \mathbb{E}_F [PDP_{\hat{f}}] + \mathbb{E}_F \mathbb{E}_X [\widehat{PDP}_{\hat{f}}^2] \\
&= PDP_f^2 - 2PDP_f \mathbb{E}_F [PDP_{\hat{f}}] + \mathbb{E}_F \text{Var}_X [\widehat{PDP}_{\hat{f}}] + \mathbb{E}_F [\mathbb{E}_X [\widehat{PDP}_{\hat{f}}]^2] \\
&= PDP_f^2 - 2PDP_f \mathbb{E}_F [PDP_{\hat{f}}] + \mathbb{E}_F \text{Var}_X [\widehat{PDP}_{\hat{f}}] + \text{Var}_F (\mathbb{E}_X [\widehat{PDP}_{\hat{f}}]) \\
&\quad + \mathbb{E}_F (\mathbb{E}_X [\widehat{PDP}_{\hat{f}}])^2 \\
&= PDP_f^2 - 2PDP_f \mathbb{E}_F [PDP_{\hat{f}}] + \text{Var}_F [PDP_{\hat{f}}] + \mathbb{E}_F [PDP_{\hat{f}}]^2 + \mathbb{E}_F \text{Var}_X [\widehat{PDP}_{\hat{f}}] \\
&= (PDP_f - \mathbb{E}_F [PDP_{\hat{f}}])^2 + \text{Var}_F [PDP_{\hat{f}}] + \mathbb{E}_F [\text{Var}_X (\widehat{PDP}_{\hat{f}})]
\end{aligned}$$

At multiple points, we use the fact that  $\mathbb{E}_X [\widehat{PDP}_{\hat{f}}] = PDP_{\hat{f}}$  (cf. [26]).

We see that the variance due to MC integration also depends on the model distribution  $F$ . Similarly, one could use the estimate  $\widehat{PDP}_{f,S}$  as groundtruth, introducing an additional variance term  $\text{Var}_{X_{mc2}}$  when estimated on different MC sample (proof in APPENDIX A). An overview of all error terms in this chain can be found in **Fig. 1**.

$\mathbb{E}_F$  and  $\text{Var}_F$  could be estimated by averaging over multiple models of the same inducer fitted to  $M$  different training data sets sampled independently from the DGP. We propose the following estimators:

$$\widehat{\text{MSE}}(x_S; PDP_{f,S}, \widehat{PDP}_{\hat{f},S}) = \frac{1}{M} \sum_{m=1}^M (PDP_{f,S}(x_S) - \widehat{PDP}_{\hat{f}(m),S}(x_S))^2 \quad (12)$$

$$\widehat{\text{Bias}}(x_S; PDP_{f,S}, \widehat{PDP}_{\hat{f},S}) = (PDP_{f,S}(x_S) - \frac{1}{M} \sum_{m=1}^M \widehat{PDP}_{\hat{f}(m),S}(x_S)) \quad (13)$$

$$\widehat{\text{Variance}}(x_S; \widehat{PDP}_{\hat{f},S}) = \frac{1}{M-1} \sum_{m=1}^M \left( \widehat{PDP}_{\hat{f}^{(m)},S}(x_S) - \frac{1}{M} \sum_{m=1}^M \widehat{PDP}_{\hat{f}^{(m)},S}(x_S) \right)^2 \quad (14)$$

Note that these are similar to the approach in [26], but we do not specify which data points to use for MC integration. The variance captures both the variance in the model fits and the variance due to MC integration. To estimate the MC variance, we propose the following estimators:

$$\widehat{\text{Variance}}_{MC}(x_S; PDP_f, \widehat{PDP}_f) = \frac{1}{K} \sum_{k=1}^K (PDP_f(x_S) - \widehat{PDP}_f^{(k)}(x_S))^2 \quad (15)$$

and

$$\widehat{\text{Variance}}_{MC}(x_S; \widehat{PDP}_{\hat{f}}) = \frac{1}{M(K-1)} \sum_{m=1}^M \sum_{k=1}^K \left( \widehat{PDP}_{\hat{f}^{(m)},S}^{(k)}(x_S) - \frac{1}{K} \sum_{k=1}^K \widehat{PDP}_{\hat{f}^{(m)},S}^{(k)}(x_S) \right)^2 \quad (16)$$

For more convenient analysis of the errors, one could also aggregate them over the marginal distribution of  $X_S$  (e.g., estimated by averaging over the grid points if chosen appropriately) to obtain a single error measure per feature effect.

While our definitions are based on the PDP, they can be directly applied to the ALE as well.

### 3 Test Functions for Simulation Studies

Test functions play a crucial role in research, e.g. for evaluating and comparing different methodological approaches, or when conducting simulation studies. This section synthesizes commonly used test functions across various domains, providing structured guidance for researchers — particularly those in Interpretable ML — in selecting appropriate test functions for their simulation studies. By examining test functions from different fields and their purposes of application, we aim to facilitate experimental design decisions. While this overview is not exhaustive, it highlights key test functions and their characteristics to help researchers make informed choices for their specific research needs.

*Test Functions in Optimization.* The field of optimization, where test functions are essential to enable the assessment and comparison of optimization algorithms, has established a rich foundation of test functions. A fundamental approach involves using simple mathematical expressions like the sphere function [28]. These basic functions are often combined with more complex ones like Branin or Rosenbrock to create comprehensive test suites that incorporate important properties such as nonlinearity, non-separability, and scalability [33]. A notable framework in this domain is the Comparing Continuous Optimizer (COCO) platform with its Black Box Optimization Benchmark (BBOB), offering

a structured approach to testing continuous optimization algorithms through artificial test functions [14]. These classical test function suites are well-established in optimization and may also serve as a basis for Interpretable ML researchers. However, it is important to note that the ability of these artificial test functions to represent complex real-world behavior is limited [34].

*Physics-Inspired Test Functions.* Physics-derived functions offer a compelling source of real-world test cases, with the Feynman Symbolic Regression Database (FSReD) being a prominent example. FSReD comprises 100 physics equations from the seminal *Feynman’s Lectures on Physics (34–36)*, supplemented by 20 more challenging equations from other seminal physics texts [32]. These equations span diverse physics domains including classical mechanics, electromagnetism, and quantum mechanics. They involve between one and nine independent variables and incorporate various elementary functions such as arithmetic operations, trigonometric functions, and exponentials. From that, tabular datasets were generated through random sampling from defined value ranges.

Building upon this foundation, Matsubara et al. [23] addressed several limitations of the original FSReD. Their enhanced framework introduces a three-tiered categorization of problems (easy, medium, hard) based on their complexity, incorporates dummy variables to simulate irrelevant features, and implements more realistic sampling ranges and strategies. Detailed specifications for all formulas, including their sampling parameters, are available in their work.

While initially developed for symbolic regression tasks, many Feynman equations may serve as suitable test functions for simulation studies in Interpretable ML. Their basis in physical principles provides real-world relevance, though researchers should carefully select equations that align with their specific analytical objectives and complexity requirements.

*Test Functions for Interpretable Machine Learning.* The Interpretable ML field itself has developed several specialized test functions designed to evaluate specific aspects of interpretability methods.

Goldstein et al. [12] used several simple test functions to demonstrate the behavior of Individual Conditional Expectation (ICE) curves. These include a simple additive function to demonstrate the absence of interactions, simple interactions to reveal heterogeneity that might be obscured by averaging procedures such as PDPs, and a specially designed function with an empty quadrant for assessing extrapolation behavior.

Similarly, Liu et al. [19] focus on simple functions before progressing to more complex ones. They begin with basic two-variable scenarios — using additive functions, interaction functions, and combinations thereof — and examine these under both independent and correlated feature conditions to compare various feature effect methods. The advantage of these simple test functions is that solutions (e.g., feature effects) can also be computed analytically, and that they allow for deeper and more fine-grained analysis of individual aspects.

A more complex test function suite was proposed by Tsang et al. [31], specifically designed to evaluate the detection of variable interactions. Their func-



tions incorporate various types of interactions with different orders, strengths, non-linearities, and overlaps. While this makes them particularly valuable for interaction detection, they are also useful for evaluating other interpretability methods in scenarios with complex interactions.

The Friedman functions [3, 8] serve as classical benchmarks applicable across various interpretability tasks. These three functions combine linear and non-linear effects with interactions, incorporating dummy variables and random noise terms to reflect realistic complexity. For detailed specifications, see [3].

When choosing test functions for simulation studies in Interpretable ML, researchers should consider several criteria, including the specific aspects of interpretability being evaluated, the desired complexity level and number of variables, the presence of specific challenges such as correlation between features or interactions, the need for analytical solutions for validation, as well as the relevance to real-world applications in the domain of interest.

## 4 Methodology & Experimental Set-Up

To address the research question of which data to choose to estimate feature effects, we conduct a comprehensive simulation study. For different models, datasets, and dataset sizes, we estimate the feature effects PDP and ALE on training data, validation data, and in a cross-validated manner. We then compute the feature effect error as MSE with respect to the ground truth feature effects and decompose it into bias and variance components. By comparing these error terms across the different estimation strategies (training, validation, cross-validation), we aim to provide nuanced empirical insights into the implications of the data choice on feature effect estimation. For deeper analysis, we conduct two ablation studies: one decomposing the feature effect variance into model variance and Monte Carlo variance, and another examining the impact of the dataset size on the Monte Carlo variance. In the following, we describe our experimental set-up for the main simulation study (SECTION 4.1) and the ablation studies (SECTION 4.2).

### 4.1 Main Experiment

*Datasets.* Building upon SECTION 3, we employ three distinct datasets of varying complexity for our simulation study:

- **SimpleNormalCorrelated** consists of four standard-normally distributed features, where the first two features exhibit strong correlation ( $\rho = 0.9$ ) while the others are independent dummy variables. The target variable is given by this simple formula:

$$f_1(\mathbf{x}) = x_1 + \frac{x_2^2}{2} + x_1x_2 \quad (17)$$

This test function is inspired by [19] and aims to focus on the impact of correlation and interactions.

- **Friedman1** implements the classical Friedman1 benchmark function [3, 8] with seven uniformly distributed features between 0 and 1, all mutually independent:

$$f_2(\mathbf{x}) = 10 \sin(\pi x_1 x_2) + 20(x_3 - \frac{1}{2})^2 + 10x_4 + 5x_5 \quad (18)$$

This dataset includes a mix of linear and different non-linear effects with interactions.

- **Feynman I.29.16** is based on the Feynman equation I.29.16 describing wave interference, comprising six independent features. Using the refined sampling strategies from Matsubara et al. [23], two log-uniformly distributed variables on  $[0.1, 10]$ , two angles uniformly distributed over  $[0, 2\pi]$ , and two uniformly distributed dummy variables on  $[0, 1]$ :

$$f_3(\mathbf{x}) = \sqrt{x_1^2 + x_2^2 + 2x_1 x_2 \cos(\theta_1 - \theta_2)} \quad (19)$$

This dataset is designed to reflect a physics-based relationship for more real-world relevance.

To generate the datasets, a standard normally distributed noise term  $\epsilon$  is added to each function, scaled by a signal-to-noise ratio of five<sup>5</sup>. For each test function, we consider two dataset sizes:  $(n_{train} = 1000, n_{val} = 250)$  and  $(n_{train} = 8000, n_{val} = 2000)$ .

*Models.* We consider a set of seven complementary learners, spanning different modeling paradigms:

- **LinReg**: a simple linear regression model as a baseline.
- **GAM\_OT**: a Generalized Additive Model (GAM) with spline terms and tensor splines for first-order interactions, number of splines and penalization optimally tuned.
- **GAM\_OF**: as above, but with hyperparameters chosen to overfit.
- **SVM\_OT**: a Support Vector Machine (SVM) with optimally tuned hyperparameters.
- **SVM\_OF**: a Support Vector Machine (SVM) with hyperparameters chosen to overfit.
- **XGBoost\_OT**: an XGBoost model with optimally tuned hyperparameters.
- **XGBoost\_OF**: an XGBoost model with hyperparameters chosen to overfit.

Hyperparameters are pre-selected, i.e. carefully hand-picked for overfitting scenarios and tuned on a different dataset for optimal tuning. Details on the hyperparameters and performances of the models can be found in APPENDIX B.

<sup>5</sup> To determine the factor by which the noise is multiplied, the standard deviation of the signal is computed over 100'000 randomly drawn samples of  $y$  and divided by the signal-to-noise ratio of five.

*Feature Effect Estimation.* For the trained models, we estimate the feature effects  $\widehat{PDP}_{f,S}$  and  $\widehat{ALE}_{f,S}$  per feature using equations (2) and (5). For Monte Carlo integration, we use for each the full training and validation set as well as a cross-validation strategy. In the latter case, we use a 5-fold cross-validation on the combined samples of training and validation set, in each fold fit the model on four folds, and estimate the feature effects on the remaining fold. We then average the feature effects over all folds. We also estimate the ground truth feature effects  $\widehat{PDP}_{f,S}$  and  $\widehat{ALE}_{f,S}$  since the complexity of the Feynman equation makes analytical computation infeasible. For the Monte Carlo integration, we use 10000 samples additionally sampled from the DGP, which introduces only a negligible additional variance term as we show in (...). We use 100 grid points, defined by the quantiles of the theoretical distribution of  $X_S$  to enable a comparison across feature effects. Additionally, we center the curves after estimation to have a mean effect of 0 w.r.t. the grid points, and we omit the first and the last grid point after estimation to avoid boundary effects especially occurring in ALE plots.

*Feature Effect Errors.* To quantify the error of the estimated feature effects, we compute the MSE of the estimated model feature effects with respect to the estimated ground truth features effects, as well as their bias and variance. For that, we use the estimators defined in equations (12) to (14), using  $\widehat{PDP}_{f,S}$  instead of  $PDP_{f,S}$  as ground truth (analogously for ALE). We repeat each dataset-size-model combination  $M = 30$  times to estimate the error terms, where each repetition involves drawing a new training and validation set from the DGP, refitting the models, and estimating the feature effects. To aggregate the errors into a single measure per feature effect, we average the errors over the marginal distribution of  $X_S$  by averaging over the grid points.

## 4.2 Ablation Experiments

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## 5 Results

## 6 Conclusions

### 6.1 Summary & Discussion

### 6.2 Limitations & Future Work

**Acknowledgments.** A bold run-in heading in small font size at the end of the paper is used for general acknowledgments, for example: This study was funded by X (grant number Y).

**Disclosure of Interests.** The authors have no competing interests to declare that are relevant to the content of this article.

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## A Proof: Monte Carlo Variance for Groundtruth

*Proof.* For better readability, we will omit the subscript  $S$  as well as the point  $x_S$  and use  $X = X_{mc}$  in this proof:

$$\begin{aligned}
\mathbb{E}_F \mathbb{E}_X [(PDP_f - \widehat{PDP}_f)^2] &= \mathbb{E}_F \mathbb{E}_X [PDP_f^2 - 2PDP_f \widehat{PDP}_f + \widehat{PDP}_f^2] \\
&= PDP_f^2 - 2PDP_f \mathbb{E}_F \mathbb{E}_X [\widehat{PDP}_f] + \mathbb{E}_F \mathbb{E}_X [\widehat{PDP}_f^2] \\
&= PDP_f^2 - 2PDP_f^2 + \mathbb{E}_F \text{Var}_X [\widehat{PDP}_f] + \mathbb{E}_F [\mathbb{E}_X [\widehat{PDP}_f]^2] \\
&= PDP_f^2 - 2PDP_f^2 + \mathbb{E}_F \text{Var}_X [\widehat{PDP}_f] + PDP_f^2 \\
&= \mathbb{E}_F \text{Var}_X [\widehat{PDP}_f] \\
&= \text{Var}_X [\widehat{PDP}_f]
\end{aligned}$$

Again we use  $\mathbb{E}_X [\widehat{PDP}_f] = PDP_f$  (cf. [26]) as well as the fact that all quantities based on  $f$  do not depend on the model distribution  $F$ .

## B Model Hyperparameters and Performance

In this appendix, we give details on the models used in the simulation study. Specifically, we provide details on the hyperparameter selection and the finally used hyperparameters, as well as the performance of the models.

### B.1 Hyperparameters

An overview of all hyperparameters used for the different models can be found in **Table 1**. Note that the linear regression is excluded as we do not have any hyperparameters to tune.

Hyperparameters for the overfitting models (OF) were carefully hand-picked to achieve strong performance on the training data while performing relatively poorly on the validation data.

The optimal hyperparameters (OT) were chosen by tuning the models on a separate data sample. Training and validation data were sampled independently from the correspondings DGPs, with a training size of  $n_{train}$  and a validation size of 10000 to get a reliable performance estimate. While this scenario is unrealistic in practice, it allows as to consider hyperparameters as “pre-selected” and avoid costly nested resampling strategies for the simulation study. Each model was tuned for 200 trials using a *Tree-structured Parzen Estimator (TPE)* [?] using validation MSE as objective to minimize (equivalent to maximizing R2-score).

### B.2 Model Performance Evaluation

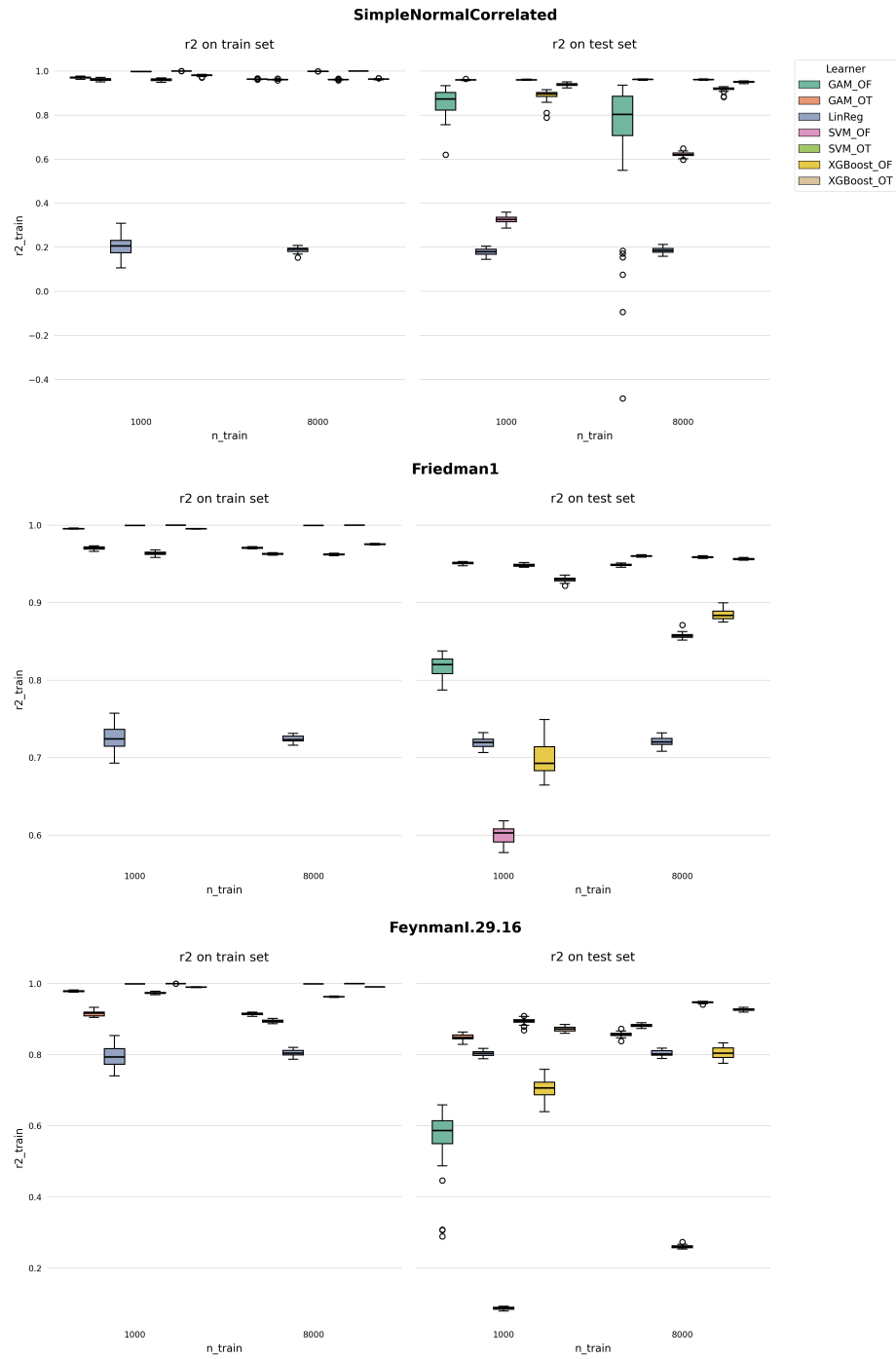
To ensure that the models perform as expected, we evaluate their performance both on the training data and a holdout test data with the latter consisting of

Dataset	n_train	Model	Hyperparameters
Simple Normal Correlated	1000	GAM_OF	n_bases: 50; lam: 0.0005;
		GAM_OT	n_bases: 20; lam: 15.6807;
		SVM_OF	C: 800; gamma: 10;
		SVM_OT	C: 917.9061; gamma: 0.0030;
		XGBoost_OF	n_estimators: 1200; max_depth: 16; learning_rate: 0.35; subsample: 1.0; min_child_weight: 1; colsample_bytree: 1.0; colsample_bylevel: 1.0; lambda: 0; alpha: 0;
		XGBoost_OT	n_estimators: 1640; max_depth: 5; learning_rate: 0.0062; subsample: 0.5601; min_child_weight: 1.6999; colsample_bytree: 0.7632; colsample_bylevel: 0.6944; lambda: 0.0156; alpha: 0.0660;
	8000	GAM_OF	n_bases: 64; lam: 1e-05;
		GAM_OT	n_bases: 5; lam: 0.0010;
		SVM_OF	C: 1000; gamma: 10;
		SVM_OT	C: 864.4724; gamma: 0.0085;
		XGBoost_OF	n_estimators: 1500; max_depth: 18; learning_rate: 0.4000; subsample: 1.0; min_child_weight: 1; colsample_bytree: 1.0; colsample_bylevel: 1.0; lambda: 0; alpha: 0;
		XGBoost_OT	n_estimators: 2586; max_depth: 5; learning_rate: 0.0044; subsample: 0.9484; min_child_weight: 1.4257; colsample_bytree: 0.8471; colsample_bylevel: 0.8672; lambda: 5.1002; alpha: 0.0026;
Friedman1	1000	GAM_OF	n_bases: 50; lam: 0.0001;
		GAM_OT	n_bases: 21; lam: 0.0402;
		SVM_OF	C: 1000; gamma: 15;
		SVM_OT	C: 917.1949; gamma: 0.2102;
		XGBoost_OF	n_estimators: 1000; max_depth: 14; learning_rate: 0.3; subsample: 1.0; min_child_weight: 1; colsample_bytree: 1.0; colsample_bylevel: 1.0; lambda: 0; alpha: 0;
		XGBoost_OT	n_estimators: 2621; max_depth: 8; learning_rate: 0.0335; subsample: 0.6192; min_child_weight: 5.4066; colsample_bytree: 0.7651; colsample_bylevel: 0.5224; lambda: 11.6021; alpha: 4.5342;
	8000	GAM_OF	n_bases: 80; lam: 1e-08;
		GAM_OT	n_bases: 22; lam: 0.0657;
		SVM_OF	C: 1000; gamma: 18;
		SVM_OT	C: 901.1903; gamma: 0.2602;
		XGBoost_OF	n_estimators: 1200; max_depth: 14; learning_rate: 0.3; subsample: 1.0; min_child_weight: 1; colsample_bytree: 1.0; colsample_bylevel: 1.0; lambda: 0; alpha: 0;
		XGBoost_OT	n_estimators: 3691; max_depth: 5; learning_rate: 0.0070; subsample: 0.6643; min_child_weight: 1.4075; colsample_bytree: 0.8403; colsample_bylevel: 0.8186; lambda: 0.0399; alpha: 5.0734;
Feynman I.29.16	1000	GAM_OF	n_bases: 50; lam: 0.0001;
		GAM_OT	n_bases: 31; lam: 0.3260;
		SVM_OF	C: 200; gamma: 8;
		SVM_OT	C: 11.4317; gamma: 0.1394;
		XGBoost_OF	n_estimators: 1000; max_depth: 14; learning_rate: 0.3; subsample: 1.0; min_child_weight: 1; colsample_bytree: 1.0; colsample_bylevel: 1.0; lambda: 0; alpha: 0;
		XGBoost_OT	n_estimators: 4246; max_depth: 9; learning_rate: 0.0327; subsample: 0.8004; min_child_weight: 6.3792; colsample_bytree: 0.8420; colsample_bylevel: 0.8357; lambda: 14.1131; alpha: 6.0556;
	8000	GAM_OF	n_bases: 64; lam: 5e-07;
		GAM_OT	n_bases: 32; lam: 0.4500;
		SVM_OF	C: 400; gamma: 10;
		SVM_OT	C: 22.5167; gamma: 0.1114;
		XGBoost_OF	n_estimators: 1000; max_depth: 14; learning_rate: 0.3; subsample: 1.0; min_child_weight: 1; colsample_bytree: 1.0; colsample_bylevel: 1.0; lambda: 0; alpha: 0;
		XGBoost_OT	n_estimators: 3962; max_depth: 7; learning_rate: 0.0372; subsample: 0.9351; min_child_weight: 4.0962; colsample_bytree: 0.8640; colsample_bylevel: 0.7728; lambda: 29.2036; alpha: 5.1631;

Table 1. Hyperparameters for the models used in the simulation study



10000 samples to get a reliable performance estimate. The performances evaluated over the 30 repetitions are aggregated in **Fig. 2** showing the R2-scores. As intended, the overfitting models perform better on the training data than the optimally tuned models, while being outperformed on the holdout test data and also exhibiting higher variance in their generalization performance. Note that the linear regression model serves only as a baseline. The overfitted SVM (SVM\_OF) show substantially worse performances on test data compared to the other models. Therefore, we excluded it from further analysis.



**Fig. 2.** R2-score of the models on training (left) and holdout test data (right), each boxplot aggregates 30 repetitions