

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 critical domains and applications, such as the healthcare, legal, or finance sectors, where decisions must be transparent and accountable [1].

Interpretability in ML is crucial to enhance trust [34, 35], address potential biases [16], fairness and ethical concerns [21], and ensure compliance with regulations such as the EU’s General Data Protection Regulation (GDPR) [7] and AI Act [8]. To address these challenges, the field of Explainable AI / Interpretable ML has emerged [1]. In the last years, a wide variety of methods has been developed, including model-specific and model-agnostic approaches ranging from local feature attributions to global feature importances and effects¹.

Due to the severity of many applications, it is crucial to utilize these explainability methods correctly to avoid misleading or incorrect conclusions. In general, there are many pitfalls to be aware of [31], 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. Existing works have studied the implications of this choice for many methods, including studies on *Permutation Feature Importance (PFI)* (e.g., in [31]), *Mean Decrease in Impurity (MDI)* of Random Forests [23], or *SHAP* values [24].

However, to the best of our knowledge, there exist no such systematic studies for feature effect methods like *Partial Dependence Plots (PDP)* [11] and *Accumulated Local Effects (ALE)* [2]. Current literature (e.g., [2, 11, 29]) predominantly uses training data without explicit justification, while practitioners also advocate

¹ For an overview of Explainable AI methods, see e.g. [1, 20, 29].

for using unseen validation data (for details, see SECTION 2). Factors that may influence this choice include potential biases arising from overfitting, the dataset size, and computational constraints. Although PDP and ALE are not based on generalization error like PFI, it is not studied if and how they are affected by overfitting. In addition, larger datasets may improve the feature effect estimates, but also increase computation times.

In this paper, we aim to answer the largely unaddressed, fundamental methodological question of whether to use training or validation data to compute feature effects. We perform an empirical simulation study, estimating PDP and ALE on training data, validation data, and in a cross-validated manner for different models and dataset. We decompose the error of PDP and ALE and compare the error components to understand the implications of the data choice. Our main contributions can be summarized 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 datasets.
2. We extend previous work to provide a theoretical framework for decomposing feature effect errors and estimating the corresponding components.
3. We provide an overview of commonly used test functions for simulation studies, including applications and purposes, as guidance for researchers in Interpretable ML.

These contributions have several important implications: Firstly, our empirically grounded recommendations enable practitioners and researchers to make informed decisions about which data to choose for feature effect computation, helping to understand potential implications of their choices. Additionally, our framework for evaluating feature effects and 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 provide an overview of related works on feature effects and test functions for simulation studies. In SECTION 3, introduce the considered feature effect methods PDP and ALE, and demonstrate how to decompose the error of these feature effects, providing corresponding definitions and estimators. We then describe the methodology and set-up of our simulation studies in SECTION 4, present the results in SECTION 5, and conclude our work with a discussion of their implications and limitations in SECTION 6.

2 Related Works

2.1 Feature Effects

Feature Effect Methods. Current literature offers various methods for analyzing feature effects in ML. One of the most popular methods is the *Partial Dependence*

Plot (PDP) [11], which describes the marginal effect of one or two features on the model prediction. The PDP assumes that the features of interest are independent of the remaining features. When this assumption is violated, the method may produce unrealistic data points outside the underlying joint distribution of the data. This extrapolation issue can cause misleading interpretations [29, 31].

M-Plots (Marginal Plots) try to address this issue by considering the conditional distribution of the remaining features given the feature of interest but instead suffer from the omitted variable bias [2, 11].

The *Accumulated Local Effects (ALE)* plot [2] addresses both issues by accumulating local differences in model predictions along the feature of interest, computing conditional expectations within small intervals rather than across the entire feature range.

Alternative approaches include *functional ANOVA (fANOVA)*, which decomposes feature effects into main and interaction effects [19]. Recent extensions have further addressed limitations of these classical methods. For example, *Robust and Heterogeneity-aware ALE (RHALE)* [13] adds heterogeneity quantification to ALE and improving robustness, or *Accumulated Total Derivative Effect (ATDEV)* plots can be decomposed into ALE and *Accumulated Cross Effects (ACE)* [22].

Beyond these global feature effect methods, there are regional effect plots such as *REPID* [18], and local methods like *ICE (Individual Conditional Expectation)* curves [14] or SHAP dependence plots [25].

In this paper, we focus on PDP and ALE and refer to them when speaking of feature effects.

Feature Effect Error and Uncertainty. Multiple works have approached measuring uncertainty in feature effects. For probabilistic ML models, Moosbauer et al. [32] derived model-specific confidence bands for PDPs, while applied studies have proposed bootstrap-based approaches [6, 15]. While the former are not model-agnostic, the latter often capture only the variance introduced by Monte Carlo approximation. Nonetheless, the model variance is another source of uncertainty in PDPs and ALEs, and accounting for it requires multiple model fits [2, 31].

Molnar et al. [30] give formalizations of PDP (and PFI) as statistical estimators of ground truth estimands, demonstrate the decomposition of its mean squared error (MSE) into model bias, model variance, and Monte Carlo variance, and provide estimators for both Monte Carlo and overall variance. Our work builds upon and extends this idea.

Data Choice for Feature Effect Estimation. For many interpretability methods, the choice whether to compute explanations in-sample (on training data) or out-of-sample (on validation data) can significantly impact interpretations: For loss-based methods such as *Permutation Feature Importance (PFI)* [5, 9], this choice is crucial and has been extensively studied (e.g., in [31]). Similar concerns have been identified for other explainability methods: both the *Mean Decrease*

in *Impurity (MDI)* of Random Forests and *SHAP* values have been shown to potentially exhibit biases when computed on training data [23, 24].

For feature effect methods, however, the implications of this choice remain largely unexplored. The original works introducing PDP [11] and ALE [2], as well as general introductory literature on Interpretable ML [29], predominantly use training data without explicit justification. In contrast, practitioners often advocate for using holdout data or base their choice on practical constraints such as dataset size². A too large dataset can increase computation times substantially, particularly for PDPs [11]. Moreover, Molnar et al. [30] estimate the PDP on holdout data when aiming to quantify the variance of feature effect estimates. Nonetheless, a systematic study on the implications of the data choice for feature effect estimation is missing so far.

2.2 Test Functions for Simulation Studies

Test functions play a crucial role in research, e.g. for evaluating different methodological approaches, or when conducting simulation studies. In the following, we synthesize 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 rather than giving an exhaustive overview.

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 [33]. 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 [38]. 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 [17]. These classical test function suites are well-established in optimization and may also serve as a basis for Interpretable ML researchers. However, the ability of these artificial test functions to represent complex real-world behavior is often limited [39].

Physics-Inspired Test Functions. Physics-derived functions offer a compelling source of real-world test cases, with the *Feynman Symbolic Regression Database (FSReD)* [37] being a prominent example. FSReD comprises 100 physics equations from the seminal *Feynman’s Lectures on Physics (34–36)*, supplemented

² for examples, see <https://github.com/SauceCat/PDPbox/issues/68> and <https://forums.fast.ai/t/partial-dependence-plot/98465> (both accessed 10/27/2024)

by 20 more challenging equations from other seminal physics texts. These equations span diverse physics domains and involve a varying number of variables and various elementary functions such as arithmetic operations, trigonometric functions, and exponentials. Tabular datasets are generated through random sampling from defined value ranges.

Matsubara et al. [27] addressed several limitations of the original FSReD. They introduce a three-tiered categorization of problems (easy, medium, hard) based on their complexity, incorporate dummy variables to simulate irrelevant features, and implement 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. [14] 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. [22] 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. [36], specifically designed to evaluate the detection of variable interactions. Their functions 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 [4, 10] 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 [4].

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 vari-

ables, 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.

3 Background

3.1 Feature Effects

The *Partial Dependence Plot (PDP)* by Friedman [11] 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 [30]. Molnar et al. [31] recommend using quantile-based over equidistant grids.

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

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

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].

3.2 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. [30] 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. [30]:

Definition 1 (Definition 1 from [30]). *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. [30], 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*³ [28].

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 f instead of \hat{f} into the estimators in equations (2) and (5) / (6)).

³ for details, see [18]

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. [12]). 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 [30]:

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

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

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:

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

At multiple points, we use the fact that $\mathbb{E}_X[\widehat{PDP}_{\hat{f}}] = PDP_{\hat{f}}$ (cf. [30]).

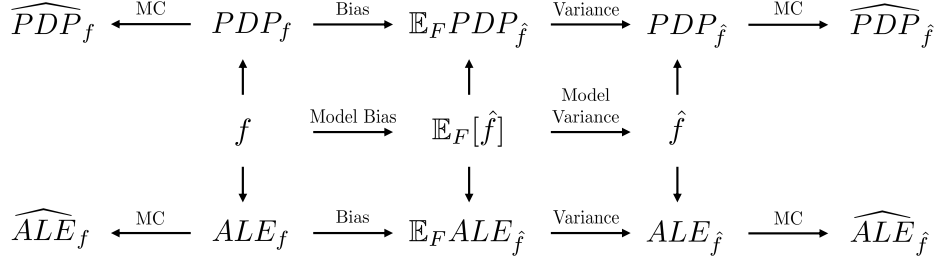


Fig. 1. Error chain in feature effect estimation (modified, original version can be found in [30])

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 $Var_{X_{mc2}}$ 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 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:

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

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

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

Note that these are similar to the approach in [30], 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:

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

and

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

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.

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 2.2, 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 [22] and aims to focus on the impact of correlation and interactions.

- **Friedman1** implements the classical Friedman1 benchmark function [4, 10] 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. [27], two log-uniformly distributed variables on $[0.1, 10]$, two angles uniformly distributed over $[0, 2\pi]$, and two uniformly distributed dummy variables on $[0, 1]$:

$$f_3(\mathbf{x}) = \sqrt{x_1^2 + x_2^2 + 2x_1x_2 \cos(\theta_1 - \theta_2)} \quad (19)$$

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 ϵ is added to each function, scaled by a signal-to-noise ratio of five⁴. 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.

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

⁴ 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.

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

As described earlier, we conduct two ablation studies for more detailed insights into the Monte Carlo variance.

Variance decomposition study. The variance of the feature effects estimated with the procedure described in the main experiment SECTION 4.1 consists of two components: the model variance and the variance due to the Monte Carlo integration. To disentangle these two sources of variance, we aim to additionally estimate the Monte Carlo variance using the estimator in equation (16). This allows us to subtract it from the total variance estimated in the main experiment and obtain the model variance, giving more nuanced insights into the implications of the data choice on feature effect estimation. To obtain this estimate, we maintain the set-up of the main experiment but for each trained model $\hat{f}^{(m)}$ (plus five models from CV $\hat{f}^{(m,1)} - \hat{f}^{(m,5)}$), we draw $K = 30$ new training and validation sets from the DGP and estimate the feature effects on these sets plus in a cross-validated manner with the already fitted models. This allows us to estimate the Monte Carlo variance as in equation (16). Since this study is computationally more expensive, we do this only for XGBoost (OT and OF) with the same datasets and sizes as in the main experiment.

Impact of dataset size on Monte Carlo variance. To investigate the impact of the dataset size on the Monte Carlo variance, we estimate the Monte Carlo variance between the analytical ground truth feature effects and the estimated ground truth feature effects on different dataset sizes. As estimator for the Monte Carlo variance, we use equation (15). With this approach, we entirely omit the

model and thus eliminate other error sources such as model bias and variance (as with a perfect model fit). To estimate the ground truth feature effects, we use 50 different dataset sizes ranging from 10^1 to 10^6 on a logarithmic scale. Note that we only consider the SimpleNormalCorrelated and Friedman1 datasets for this study as an analytical computation of the FeynmanI.29.16 feature effects is infeasible. For other details, we maintain the set-up of the main experiment.

5 Results

6 Conclusions

6.1 Summary & Discussion

6.2 Limitations & Future Work

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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. [30]) 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)* [3] 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.

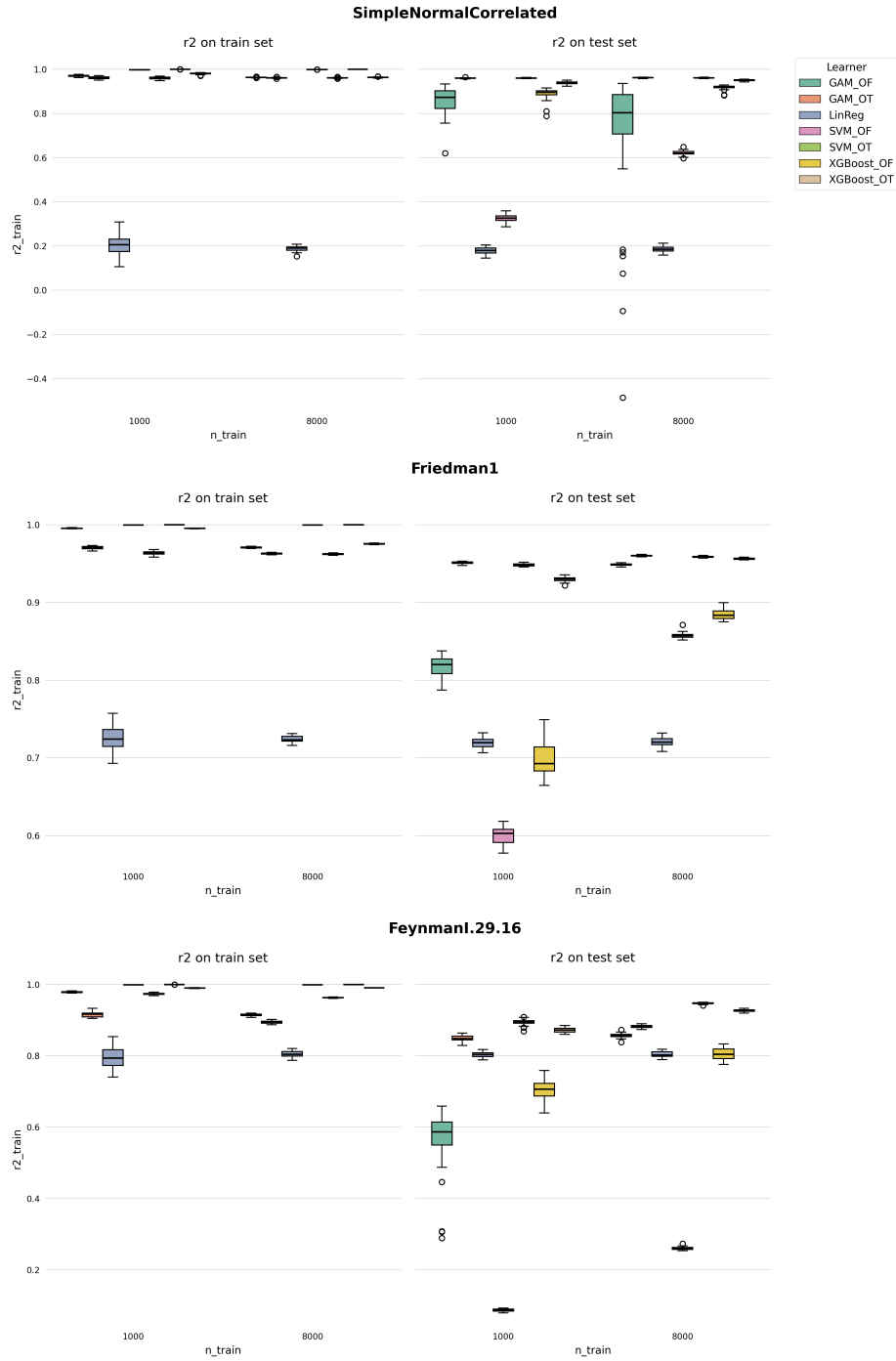


Fig. 2. R2-score of the models on training (left) and holdout test data (right), each boxplot aggregates 30 repetitions