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# Responsible Machine Learning

with Interpretable Models, Post-hoc Explanation, and Disparate Impact Testing

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**Abstract:** This text outlines a viable approach for training and evaluating machine learning (ML) systems for high-stakes, human-centered, or regulated applications using common Python programming tools. The accuracy and intrinsic interpretability of two types of constrained models, monotonic gradient boosting machines (MGBM) and explainable neural networks (XNN), a deep learning architecture well-suited for structured data, are assessed on simulated data with known feature importance and sociological bias characteristics and on realistic, publicly available lending data. For maximum transparency and the potential generation of personalized adverse action notices, the constrained models are analyzed using post-hoc explanation techniques including plots of partial dependence (PD) and individual conditional expectation (ICE) and global and local gradient-based or Shapley feature importance. The constrained model predictions are also tested for disparate impact (DI) and other types of sociological bias using straightforward group fairness measures. By combining innovations in interpretable models, post-hoc explanation, and bias testing with accessible software tools, this text aims to provide a template workflow for important ML applications that require high accuracy and interpretability and low disparate impact.

**Keywords:** Machine Learning; Neural Network; Gradient Boosting Machine; Interpretable; Explanation; Fairness; Disparate Impact; Python

## 0. Introduction

ML models can be inaccurate and unappealable black-boxes, even with the application of newer post-hoc explanation techniques [1].<sup>1</sup> ML models can perpetuate and exacerbate social biases [2], [3], [4], [5]. ML models can be hacked, resulting in manipulated model outcomes or the exposure of proprietary intellectual property or sensitive training data [6], [7], [8]. The authors make no claim that the interdependent issues of opaqueness, unwanted sociological bias, or security vulnerabilities in ML have been solved, even as singular entities, much less as complex intersectional phenomena, e.g. the fairwashing or scaffolding of biased models with ML explanations or the privacy harms of ML explanations [9], [10], [11]. However, this text does present an attempt to address the technological aspects of these vexing problems with interpretable models, post-hoc explanation, and DI testing implemented in widely available, free, and open source Python tools. Section 1 describes methods and materials herein, including simulated and collected training datasets, interpretable and constrained architectures used to train highly transparent models, post-hoc explanations used to create an *appealable*

<sup>1</sup> See: "When a Computer Program Keeps You in Jail".

decision-making framework, tests for DI and other unwanted sociological bias, and public and open source software resources. In Section 2, interpretable and constrained modeling results are compared to less interpretable and unconstrained models and post-hoc explanation and DI testing results are also presented. Section 3 then discusses some nuances of the presented modeling, explanation, and DI testing methods and results. Finally, Section 4 closes this text with a brief outline of proposed additional steps to increase human trust and understanding in ML and also touches on the authors' future work.

## 1. Materials and Methods

Section 1 presents descriptions of notation, training data, ML models, post-hoc explanation techniques, DI testing methods, and software resources as follows:

- §1.1 Notation: for spaces, datasets, and models.
- §1.2 Training data: simulated data with known feature importance and social bias characteristics and collected home mortgage data.
- §1.3 ML models: interpretable XNN and constrained MGBM models.
- §1.4 Post-hoc explanation techniques: PD, ICE, and Shapley values.
- §1.5 DI testing methods: .
- §1.6 Software resources: GitHub repository for the presented results; utilized and potentially useful Python packages.

### 1.1. Notation

To facilitate descriptions of data and modeling, explanatory, and DI testing techniques, notation for input and output spaces, datasets, and models is defined here.

#### 1.1.1. Spaces

- Input features come from the set  $\mathcal{X}$  contained in a  $P$ -dimensional input space,  $\mathcal{X} \subset \mathbb{R}^P$ . An arbitrary, potentially unobserved, or future instance of  $\mathcal{X}$  is denoted  $\mathbf{x}$ ,  $\mathbf{x} \in \mathcal{X}$ .
- Labels corresponding to instances of  $\mathcal{X}$  come from the set  $\mathcal{Y}$ .
- Learned output responses come from the set  $\hat{\mathcal{Y}}$ .

#### 1.1.2. Datasets

- The input dataset  $\mathbf{X}$  is composed of observed instances of the set  $\mathcal{X}$  with a corresponding dataset of labels  $\mathbf{Y}$ , observed instances of the set  $\mathcal{Y}$ .
- Each  $i$ -th observation of  $\mathbf{X}$  is denoted as  $\mathbf{x}^{(i)} = [x_0^{(i)}, x_1^{(i)}, \dots, x_{P-1}^{(i)}]$ , with corresponding  $i$ -th labels in  $\mathbf{Y}$ ,  $\mathbf{y}^{(i)}$ , and corresponding predictions in  $\hat{\mathbf{Y}}$ ,  $\hat{\mathbf{y}}^{(i)}$ .
- $\mathbf{X}$  and  $\mathbf{Y}$  consist of  $N$  tuples of observations:  $[(\mathbf{x}^{(0)}, \mathbf{y}^{(0)}), (\mathbf{x}^{(1)}, \mathbf{y}^{(1)}), \dots, (\mathbf{x}^{(N-1)}, \mathbf{y}^{(N-1)})]$ .
- Each  $j$ -th input column vector of  $\mathbf{X}$  is denoted as  $X_j = [x_j^{(0)}, x_j^{(1)}, \dots, x_j^{(N-1)}]^T$ .

#### 1.1.3. Models

- A type of ML model  $g$ , selected from a hypothesis set  $\mathcal{H}$ , is trained to represent an unknown signal-generating function  $f$  observed as  $\mathbf{X}$  with labels  $\mathbf{Y}$  using a training algorithm  $\mathcal{A}$ :  $\mathbf{X}, \mathbf{Y} \xrightarrow{\mathcal{A}} g$ , such that  $g \approx f$ .
- $g$  generates learned output responses on the input dataset  $g(\mathbf{X}) = \hat{\mathbf{Y}}$ , and on the general input space  $g(\mathcal{X}) = \hat{\mathcal{Y}}$ .
- The model to be explained and tested for unwanted social bias is denoted as  $g$ .

## 1.2. Data Description

Results are presented for simulated data with known feature importance and social bias characteristics and for more realistic lending data sourced from the Home Mortgage Disclosure Act (HMDA) database.<sup>2</sup>

### 1.2.1. Simulated Data

### 1.2.2. Lending Data

Lending data is sampled from the public HMDA database to create a real-world dataset on which to assess interpretable models, post-hoc explanations, and DI tests. The sampled lending data contains recent and anonymized consumer loans issued by many major United States (US) financial institutions. Consumers and loans in the dataset are characterized by Bayesian improved surname geocoding (BISG) demographic features, consumer credit scores and various consumer financial features, and loan purpose, amount, and term features.<sup>3</sup> The lending data is divided randomly into training and test partitions. The training data contains 33 total features and 144,000 rows, each representing a unique loan, and a fold identifier to ensure consistent 5-fold cross-validation accuracy and error measurements across different types of models. Consumer financials and loan descriptors are used for training. Demographic features are not used in model training. The lending test data contains 36,000 loans.

## 1.3. Model Description

### 1.3.1. Explainable Neural Network

XNNs are an alternative formulation of additive index models in which the ridge functions are neural networks [12]. XNNs also bare a strong resemblance to generalized additive models (GAMs) and so-called explainable boosting machines (EBMs or GA<sup>2</sup>M), i.e. GAMs which consider main effects and a small number of 2-way interactions and incorporate boosting in their training [13], [14]. Hence, XNNs enable users to tailor interpretable neural network architectures to a given prediction problem and to visualize model behavior by plotting ridge functions. XNNs are composed of a global bias term,  $\mu_0$ ,  $K$  individually specified neural networks,  $n_k$  with scale parameters  $\gamma_k$ , and the inputs to each  $n_k$  are themselves a linear combination of modeling inputs,  $\sum_{j=0}^J \beta_{k,j} x_j$ .

$$g^{\text{XNN}}(\mathbf{x}) = \mu_0 + \sum_{k=1}^K \gamma_k n_k \left( \sum_{j=1}^J \beta_{k,j} x_j \right) \quad (1)$$

$g^{\text{XNN}}$  is comprised of 3 meta-layers:

1. The first and deepest meta-layer, composed of  $K$  linear  $\sum_j \beta_{k,j} x_j$  hidden units, is known as the *projection layer* and is fully connected to each input feature,  $X_j$ .
2. The second meta-layer contains  $K$  hidden and separate  $n_k$  ridge functions, or *subnetworks*. Each  $n_k$  is a neural network, which can be parameterized to suite a given modeling task. To facilitate direct visualization, the input to each subnetwork is the 1-dimensional output of its associated projection layer hidden unit,  $\sum_j \beta_{k,j} x_j$ .
3. The output meta-layer, called the *combination layer*, is another linear unit comprised of a global bias term,  $\mu_0$ , and the  $K$  weighted 1-dimensional outputs of each subnetwork,  $\gamma_k n_k(\sum_j \beta_{k,j} x_j)$ . Again, subnetwork output is restricted to 1-dimension for visualization purposes.

<sup>2</sup> See: Mortgage data (HMDA).

<sup>3</sup> See: Using publicly available information to proxy for unidentified race and ethnicity.

$g^{\text{XNN}}$  is typically trained by mini-batch stochastic gradient descent (SGD).  $L_1$  regularization is often applied to both the projection and combination layers to induce a sparse and interpretable model, where each  $n_k$  subnetwork and corresponding combination layer  $\gamma_k$  are ideally associated with an important  $X_j$  or combination thereof.

### 1.3.2. Monotonically Constrained Gradient Boosting Machine

MGBMs constrain typical GBM training to consider only tree splits that obey user-defined positive and negative monotonicity constraints. The MGBM remains an additive combination of  $B$  trees trained by gradient boosting,  $T_b$ , but each tree learns a set of splitting rules that respect monotonicity constraints,  $\Theta_b^{\text{mono}}$ .

$$g^{\text{mono}}(\mathbf{x}) = \sum_{b=1}^B T_b(\mathbf{x}; \Theta_b^{\text{mono}}) \quad (2)$$

As in unconstrained GBM,  $\Theta_b^{\text{mono}}$  is selected in a greedy, additive fashion by minimizing a regularized loss function that considers known target labels,  $\mathbf{y}$ , the predictions of all subsequently trained trees in the MGBM,  $g_{b-1}^{\text{mono}}(\mathbf{X})$ , and a regularization term that penalizes complexity in the current tree,  $\Omega(T_b)$ . For the  $b$ -th iteration, the loss function,  $\mathcal{L}_b$ , can generally be defined as:

$$\mathcal{L}_b = \sum_{i=0}^{N-1} l(y^{(i)}, g_{b-1}^{\text{mono}}(\mathbf{x}^{(i)}), T_b(\mathbf{x}^{(i)}; \Theta_b^{\text{mono}})) + \Omega(T_b) \quad (3)$$

In addition to  $\mathcal{L}_b$ ,  $g^{\text{mono}}$  training is characterized by additional splitting rules and constraints on tree node weights. Each binary splitting rule,  $\theta_{b,j,k} \in \Theta_b$ , is associated with a feature,  $X_j$ , is the  $k$ -th split associated with  $X_j$  in  $T_b$ , and results in left and right child nodes with a numeric weights,  $\{w_{b,j,k,L}, w_{b,j,k,R}\}$ . For terminal nodes,  $\{w_{b,j,k,L}, w_{b,j,k,R}\}$  can be direct numeric components of some  $g^{\text{mono}}$  prediction. For two values of some feature  $X_j$ ,  $x_j^\alpha \leq x_j^\beta$ , where the prediction for each value results in  $T_b(x_j^\alpha; \Theta_b) = w_\alpha$  and  $T_b(x_j^\beta; \Theta_b) = w_\beta$ ,  $\Theta_b$  is restricted to be positive monotonic w.r.t.  $X_j$  by the following rules and constraints.

1. For the first and highest split in  $T_b$  involving  $X_j$ , any  $\theta_{b,j,0}$  resulting in the left child weight being greater than the right child weight,  $T(x_j; \theta_{b,j,0}) = \{w_{b,j,0,L}, w_{b,j,0,R}\}$  where  $w_{b,j,0,L} > w_{b,j,0,R}$ , is not considered.
2. For any subsequent left child node involving  $X_j$ , any  $\theta_{b,j,k \geq 1}$  resulting in  $T(x_j; \theta_{b,j,k \geq 1}) = \{w_{b,j,k \geq 1,L}, w_{b,j,k \geq 1,R}\}$  where  $w_{b,j,k \geq 1,L} > w_{b,j,k \geq 1,R}$ , is not considered.
3. Moreover, for any subsequent left child node involving  $X_j$ ,  $T(x_j; \theta_{b,j,k \geq 1}) = \{w_{b,j,k \geq 1,L}, w_{b,j,k \geq 1,R}\}$ ,  $\{w_{b,j,k \geq 1,L}, w_{b,j,k \geq 1,R}\}$  are bound by the parent set of node weights,  $\{w_{b,j,k-1,L}, w_{b,j,k-1,R}\}$ , such that  $\{w_{b,j,k \geq 1,L}, w_{b,j,k \geq 1,R}\} \leq \frac{w_{b,j,k-1,L} + w_{b,j,k-1,R}}{2}$ .
4. (1) and (2) are also applied to all right child nodes, except that for right child nodes  $\{w_{b,j,k \geq 1,L}, w_{b,j,k \geq 1,R}\} \geq \frac{w_{b,j,k-1,L} + w_{b,j,k-1,R}}{2}$ .

Note that for any one  $X_j$  and  $T_b \in g^{\text{mono}}$  left subtrees will always produce lower predictions than right subtrees, and that any  $g^{\text{mono}}(\mathbf{x})$  is an addition of each  $T_b$  output, with the application of a monotonic logit or softmax link function for classification problems. Moreover, each tree's root node corresponds to some constant node weight that by definition obeys monotonicity constraints,  $T(x_j^\alpha; \theta_{b,0}) = T(x_j^\beta; \theta_{b,0}) = w_{b,0}$ . Together these additional splitting rules and node weight constraints ensure that  $g^{\text{mono}}(x_j^\alpha) \leq g^{\text{mono}}(x_j^\beta) \forall x_j^\alpha \leq x_j^\beta \in X_j$ . For a negative monotonic constraint, i.e.  $g^{\text{mono}}(x_j^\alpha) \geq g^{\text{mono}}(x_j^\beta) \forall x_j^\alpha \leq x_j^\beta \in X_j$ , left and right splitting rules and node weight constraints are switched.

## 1.4. Explanatory Method Description

### 1.4.1. Partial Dependence and Individual Conditional Expectation

PD plots are a widely-used method for describing the average predictions of a complex model  $g$  across some partition of data  $\mathbf{X}$  for some interesting input feature  $X_j$  [13]. ICE plots are a newer method that describes the local behavior of  $g$  for a single instance  $\mathbf{x} \in \mathcal{X}$ . PD and ICE can be combined in the same plot to compensate for known weaknesses of PD, to identify interactions modeled by  $g$ , and to create a holistic portrait of the predictions of a complex model for some  $X_j$  [15].

Following Friedman *et al.* [13] a single feature  $X_j \in \mathbf{X}$  and its complement set  $\mathbf{X}_{(-j)} \in \mathbf{X}$  (where  $X_j \cup \mathbf{X}_{(-j)} = \mathbf{X}$ ) is considered.  $\text{PD}(X_j, g)$  for a given feature  $X_j$  is estimated as the average output of the learned function  $g(\mathbf{X})$  when all the observations of  $X_j$  are set to a constant  $x \in \mathcal{X}$  and  $\mathbf{X}_{(-j)}$  is left unchanged.  $\text{ICE}(x_j, \mathbf{x}, g)$  for a given instance  $\mathbf{x}$  and feature  $x_j$  is estimated as the output of  $g(\mathbf{x})$  when  $x_j$  is set to a constant  $x \in \mathcal{X}$  and all other features  $\mathbf{x} \in \mathbf{X}_{(-j)}$  are left untouched. PD and ICE curves are usually plotted over some set of constants  $x \in \mathcal{X}$ .

### 1.4.2. Shapley Values

Shapley explanations, including Tree SHAP (SHapley Additive exPlanations), are a class of additive, locally accurate feature contribution measures with long-standing theoretical support [16]. Shapley explanations are the only possible locally accurate and globally consistent feature contribution values, meaning that Shapley explanation values for input features always sum to  $g(\mathbf{x})$  and that Shapley explanation values can never decrease for some  $x_j$  when  $g$  is changed such that  $x_j$  truly makes a stronger contribution to  $g(\mathbf{x})$  [16].

For some observation  $\mathbf{x} \in \mathcal{X}$ , Shapley explanations take the form:

$$g(\mathbf{x}) = \phi_0 + \sum_{j=0}^{j=\mathcal{P}-1} \phi_j \mathbf{z}_j \quad (4)$$

In Equation 4,  $\mathbf{z} \in \{0, 1\}^{\mathcal{P}}$  is a binary representation of  $\mathbf{x}$  where 0 indicates missingness. Each  $\phi_j$  is the local feature contribution value associated with  $x_j$  and  $\phi_0$  is the average of  $g(\mathbf{X})$ .

Shapley values can be estimated in different ways. Tree SHAP is a specific implementation of Shapley explanations that relies on traversing internal tree structures to estimate the impact of each  $x_j$  for some  $g(\mathbf{x})$  of interest [17].

$$\phi_j = \sum_{S \subseteq \mathcal{P} \setminus \{j\}} \frac{|S|!(\mathcal{P} - |S| - 1)!}{\mathcal{P}!} [g_{\mathbf{x}}(S \cup \{j\}) - g_{\mathbf{x}}(S)] \quad (5)$$

## 1.5. Social Bias Test Description

### 1.6. Software Resources

Python code to reproduce the results presented in this text are available at: <https://github.com/h2oai/article-information-2019>. The authors primarily make use of the [datatable](#), [h2o](#), [matplotlib](#), [pandas](#), [scikit-learn](#), [seaborn](#), and [shap](#) packages for data manipulation, modeling, and reporting results.

## 2. Results

### 2.1. Simulated Data Results

### 2.2. Loan Data Results

## 3. Discussion

## 4. Conclusion

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**Conflicts of Interest:**

## Abbreviations

The following abbreviations are used in this manuscript: ML – machine learning, MGBM – monotonic gradient boosting machine, XNN – explainable neural network, PD – partial dependence, ICE – individual conditional expectation, DI – disparate impact, SGD – stochastic gradient descent.

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