Interpretable Machine Learning

Permutation Feature Importance (PFI)

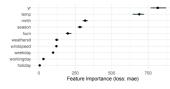


Figure: Bike Sharing Dataset

Learning goals

- Understand how PFI is computed
- Understanding strengths and weaknesses
- Testing Importance



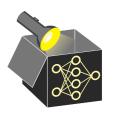
PERMUTATION FEATURE IMPORTANCE (PFI) • Breiman (2001)

Idea: "Destroy" feat. of interest x_i by perturbing it s.t. it becomes uninformative, e.g., randomly permute obs. in x_i (marginal distribution $\mathbb{P}(x_i)$ stays the same). PFI for features x_S using test data \mathcal{D} :





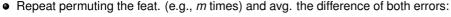
$$\begin{split} \widehat{\mathit{PFI}}_S &= \tfrac{1}{m} \sum_{k=1}^m \mathcal{R}_{\mathsf{emp}}(\hat{t}, \underbrace{\tilde{\mathcal{D}}_{(k)}^S}) - \mathcal{R}_{\mathsf{emp}}(\hat{t}, \mathcal{D}), \\ \text{where } \mathcal{R}_{\mathsf{emp}}(\hat{t}, \mathcal{D}) &= \frac{1}{n} \sum_{(x,y) \in \mathcal{D}} L(\hat{t}(x), y) \end{split}$$



PERMUTATION FEATURE IMPORTANCE (PFI) • Breiman (2001)

Idea: "Destroy" feat. of interest x_i by perturbing it s.t. it becomes uninformative, e.g., randomly permute obs. in x_i (marginal distribution $\mathbb{P}(x_i)$ stays the same). PFI for features x_S using test data \mathcal{D} :



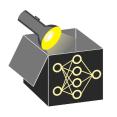


$$\begin{split} \widehat{PFI}_{\mathcal{S}} &= \tfrac{1}{m} \sum_{k=1}^{m} \mathcal{R}_{\mathsf{emp}}(\hat{f}, \tilde{\mathcal{D}}_{(k)}^{\mathcal{S}}) - \mathcal{R}_{\mathsf{emp}}(\hat{f}, \mathcal{D}), \\ \mathsf{where} \ \mathcal{R}_{\mathsf{emp}}(\hat{f}, \mathcal{D}) &= \frac{1}{n} \sum_{(x,y) \in \mathcal{D}} L(\hat{f}(x), y) \end{split}$$

The data \mathcal{D} where x_S is replaced with \tilde{x}^S is denoted as $\tilde{\mathcal{D}}^S$. Example of permuting feature x_S with $S = \{1\}$ and m = 6:

	\mathcal{D}				$ ilde{\mathcal{D}}_{(1}^{S}$)		$ ilde{\mathcal{D}}_{(2}^{S}$)		$ ilde{\mathcal{D}}_{(3)}^{S}$: 3)		$ ilde{\mathcal{D}}_{(4)}^{S}$	-)		$\tilde{\mathcal{D}}_{(5)}^{S}$)		$ ilde{\mathcal{D}}_{(6}^{S}$)
\mathbf{x}_1	X 2	X 3	_ ⇒	$\mathbf{x}_{\mathcal{S}}$	X ₂	X 3	$\mathbf{x}_{\mathcal{S}}$	X ₂	X 3	$\mathbf{x}_{\mathcal{S}}$	X ₂	X 3	$\mathbf{x}_{\mathcal{S}}$	X ₂	X 3	$\mathbf{x}_{\mathcal{S}}$	X ₂	X 3	$\mathbf{x}_{\mathcal{S}}$	X ₂	X 3
1	4	7	7	1	4	7	2	4	7	2	4	7	1	4	7	3	4	7	3	4	7
2	5	8		2	5	8	1	5	8	3	5	8	3	5	8	1	5	8	2	5	8
3	6	9		3	6	9	3	6	9	1	6	9	2	6	9	2	6	9	1	6	9

Note: The S in x_S refers to a **S**ubset of features for which we are interested in their effect on the prediction. Here: We calculate the feature importance for one feature at a time |S| = 1.

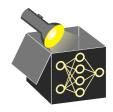




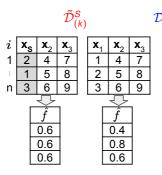
 \mathcal{I}

i	xs	\mathbf{x}_2	\mathbf{x}_3
1	2	4	7
:	1	5	8
n	3	6	9

X ₁	X ₂	x ₃
1	4	7
2	5	8
3	6	9

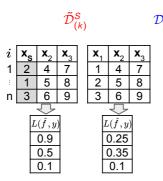


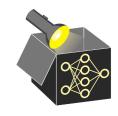
- **1. Perturbation:** Sample feature values from the distribution of $x_S(P(X_S))$.
 - \Rightarrow Randomly permute feature x_S
 - \Rightarrow Replace original feature with permuted feature \tilde{x}_S and create data $\tilde{\mathcal{D}}^S$ containing \tilde{x}_S





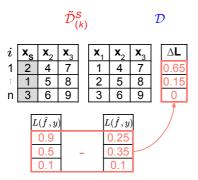
- **1. Perturbation:** Sample feature values from the distribution of x_S ($P(X_S)$).
 - \Rightarrow Randomly permute feature x_S
 - \Rightarrow Replace original feature with permuted feature \tilde{x}_S and create data $\tilde{\mathcal{D}}^S$ containing \tilde{x}_S
- **2. Prediction:** Make predictions for both data, i.e., \mathcal{D} and $\tilde{\mathcal{D}}^{\mathcal{S}}$

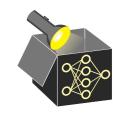




3. Aggregation:

Compute the loss for each observation in both data sets





3. Aggregation:

- Compute the loss for each observation in both data sets
- Take the difference of both losses ΔL for each observation

$$\mathcal{R}_{\mathsf{emp}}(\hat{f}, ilde{\mathcal{D}}^{\mathcal{S}}_{(k)}) - \mathcal{R}_{\mathsf{emp}}(\hat{f}, \mathcal{D})$$

i	xs	\mathbf{x}_2	\mathbf{x}_3
1	2	4	7
:	1	5	8
n	3	6	9

X ₁	X ₂	\mathbf{x}_3
1	4	7
2	5	8
3	6	9

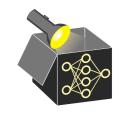


3. Aggregation:

- Compute the loss for each observation in both data sets
- Take the difference of both losses ΔL for each observation
- \bullet Average this change in loss across all observations Note: This is equivalent to computing \mathcal{R}_{emp} on both data sets and taking the difference

0.35

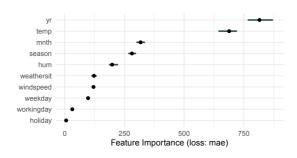
= 0.4



3. Aggregation:

- Compute the loss for each observation in both data sets
- Take the difference of both losses ΔL for each observation
- Average this change in loss across all observations
- Repeat perturbation and average over multiple repetitions

EXAMPLE: BIKE SHARING DATASET





Interpretation:

- Year (yr) and Temperature (temp) are most important features
- Destroying information about yr by permuting it increases mean absolute error of model by 816
- 5% and 95% quantile of repetitions due multiple permutations are shown as error bars

 Interpretation: PFI is the increase of model error when feature's information is destroyed



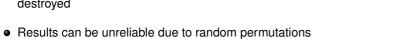
- Interpretation: PFI is the increase of model error when feature's information is destroyed
- Results can be unreliable due to random permutations
 - \Rightarrow Solution: Average results over multiple repetitions



- Interpretation: PFI is the increase of model error when feature's information is destroyed
- Results can be unreliable due to random permutations
 Solution: Average results over multiple repetitions
- Permuting features despite correlation with other features can lead to unrealistic combinations of feature values (since under dependence $\mathbb{P}(x_i, x_{-i}) \neq \mathbb{P}(x_i)\mathbb{P}(x_{-i})) \rightsquigarrow \text{Extrapolation issue}$



 Interpretation: PFI is the increase of model error when feature's information is destroyed



⇒ Solution: Average results over multiple repetitions



- Permuting features despite correlation with other features can lead to unrealistic combinations of feature values (since under dependence $\mathbb{P}(x_i, x_{-i}) \neq \mathbb{P}(x_i)\mathbb{P}(x_{-i}) \rightarrow \text{Extrapolation}$ issue
- PFI automatically includes importance of interaction effects with other features
 - \Rightarrow Permutation also destroys information of interactions where permuted feature is involved
 - \Rightarrow Importance of all interactions with the permuted feature are contained in PFI score

 Interpretation: PFI is the increase of model error when feature's information is destroyed

- Results can be unreliable due to random permutations
 - ⇒ Solution: Average results over multiple repetitions
- Permuting features despite correlation with other features can lead to unrealistic combinations of feature values (since under dependence $\mathbb{P}(x_i, x_{-i}) \neq \mathbb{P}(x_i)\mathbb{P}(x_{-i})) \rightsquigarrow \text{Extrapolation}$ issue
- PFI automatically includes importance of interaction effects with other features
 - \Rightarrow Permutation also destroys information of interactions where permuted feature is involved
 - \Rightarrow Importance of all interactions with the permuted feature are contained in PFI score
- Interpretation of PFI depends on whether training or test data is used

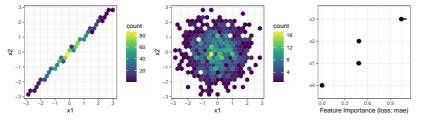
COMMENTS ON PFI - EXTRAPOLATION

Example: Let $y=x_3+\epsilon_y$ with $\epsilon_y\sim N(0,0.1)$ where $x_1:=\epsilon_1, x_2:=x_1+\epsilon_2$ are highly correlated $(\epsilon_1\sim N(0,1),\epsilon_2\sim N(0,0.01))$ and $x_3:=\epsilon_3, x_4:=\epsilon_4$, with $\epsilon_3,\epsilon_4\sim N(0,1)$. All noise terms are independent. Fitting a LM yields $\hat{f}(\mathbf{x})\approx 0.3x_1-0.3x_2+x_3$.



COMMENTS ON PFI - EXTRAPOLATION

Example: Let $y=x_3+\epsilon_y$ with $\epsilon_y\sim N(0,0.1)$ where $x_1:=\epsilon_1, x_2:=x_1+\epsilon_2$ are highly correlated $(\epsilon_1\sim N(0,1),\epsilon_2\sim N(0,0.01))$ and $x_3:=\epsilon_3, x_4:=\epsilon_4$, with $\epsilon_3,\epsilon_4\sim N(0,1)$. All noise terms are independent. Fitting a LM yields $\hat{f}(\mathbf{x})\approx 0.3x_1-0.3x_2+x_3$.

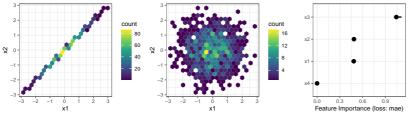


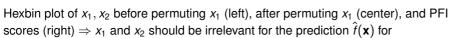
Hexbin plot of x_1 , x_2 before permuting x_1 (left), after permuting x_1 (center), and PFI scores (right)



COMMENTS ON PFI - EXTRAPOLATION

Example: Let $y=x_3+\epsilon_y$ with $\epsilon_y\sim N(0,0.1)$ where $x_1:=\epsilon_1, x_2:=x_1+\epsilon_2$ are highly correlated $(\epsilon_1\sim N(0,1),\epsilon_2\sim N(0,0.01))$ and $x_3:=\epsilon_3, x_4:=\epsilon_4$, with $\epsilon_3,\epsilon_4\sim N(0,1)$. All noise terms are independent. Fitting a LM yields $\hat{f}(\mathbf{x})\approx 0.3x_1-0.3x_2+x_3$.





 $\{\mathbf{x}: \mathbb{P}(\mathbf{x})>0\}$ as $0.3x_1-0.3x_2\approx 0$ \Rightarrow PFI evaluates model on unrealistic obs. outside $\mathbb{P}(\mathbf{x})\rightsquigarrow x_1, x_2$ are considered relevant (PFI >0)

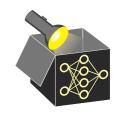


COMMENTS ON PFI - INTERACTIONS

Example: Let x_1, \ldots, x_4 be independently and uniformly sampled from $\{-1, 1\}$ and

$$y := x_1x_2 + x_3 + \epsilon_Y \text{ with } \epsilon_Y \sim N(0,1)$$

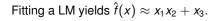
Fitting a LM yields $\hat{f}(x) \approx x_1 x_2 + x_3$.



COMMENTS ON PFI - INTERACTIONS

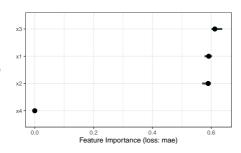
Example: Let x_1, \ldots, x_4 be independently and uniformly sampled from $\{-1, 1\}$ and

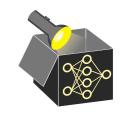
$$y := x_1 x_2 + x_3 + \epsilon_Y \text{ with } \epsilon_Y \sim N(0,1)$$



Although x_3 alone contributes as much to the prediction as x_1 and x_2 jointly, all three are considered equally relevant.

 \Rightarrow PFI does not fairly attribute the performance to the individual features.

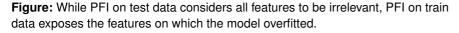




COMMENTS ON PFI - TEST VS. TRAINING DATA

Example: x_1, \ldots, x_{20}, y are independently sampled from $\mathcal{U}(-10, 10)$. An xgboost model with default hyperparameters is fit on a small training set of 50 observations. The model overfits heavily.







COMMENTS ON PFI - TEST VS. TRAINING DATA

Example: x_1, \ldots, x_{20}, y are independently sampled from $\mathcal{U}(-10, 10)$. An xgboost model with default hyperparameters is fit on a small training set of 50 observations. The model overfits heavily.

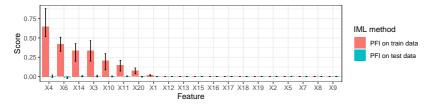




Figure: While PFI on test data considers all features to be irrelevant, PFI on train data exposes the features on which the model overfitted.

Why? PFI can only be nonzero if the permutation breaks a dependence in the data. Spurious correlations help the model perform well on train data but are not present in the test data.

⇒ If you are interested in which features help the model to generalize, apply PFI on test data.

IMPLICATIONS OF PFI

Can we get insight into whether the \dots

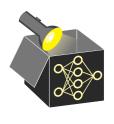
- feature x_i is causal for the prediction?
 - $PFI_j \neq 0 \Rightarrow$ model relies on x_j
 - As the training vs. test data example demonstrates, the converse does not hold



IMPLICATIONS OF PFI

Can we get insight into whether the ...

- feature x_i is causal for the prediction?
 - $PFI_j \neq 0 \Rightarrow$ model relies on x_j
 - As the training vs. test data example demonstrates, the converse does not hold
- 2 feature x_j contains prediction-relevant information?
 - $PFI_j \neq 0 \Rightarrow x_j$ is dependent of y or it's covariates x_{-j} or both (due to extrapolation)
 - x_j is not exploited by model (regardless of whether it is useful for y or not)
 ⇒ PFI_j = 0



IMPLICATIONS OF PFI

Can we get insight into whether the ...

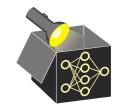
- feature x_i is causal for the prediction?
 - $PFI_j \neq 0 \Rightarrow$ model relies on x_j
 - As the training vs. test data example demonstrates, the converse does not hold



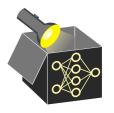
- $PFI_j \neq 0 \Rightarrow x_j$ is dependent of y or it's covariates x_{-j} or both (due to extrapolation)
- x_j is not exploited by model (regardless of whether it is useful for y or not)
 ⇒ PFI_j = 0
- \odot model requires access to x_i to achieve it's prediction performance?
 - As the extrapolation example demonstrates, such insight is not possible



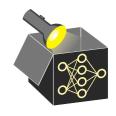
• PIMP was originally introduced for random forest's built-in permutation feature importance



- PIMP was originally introduced for random forest's built-in permutation feature importance
- PIMP investigates whether the PFI score significantly differs from 0
 - ⇒ Useful because PFI can be non-zero due to stochasticity



- PIMP was originally introduced for random forest's built-in permutation feature importance
- PIMP investigates whether the PFI score significantly differs from 0
 - ⇒ Useful because PFI can be non-zero due to stochasticity
- PIMP tests the H_0 -hypothesis: Feature is independent of the target y (unimportant)



- PIMP was originally introduced for random forest's built-in permutation feature importance
- PIMP investigates whether the PFI score **significantly** differs from 0
 - ⇒ Useful because PFI can be non-zero due to stochasticity
- PIMP tests the H_0 -hypothesis: Feature is independent of the target y (unimportant)
- Sampling under H_0 : Permute target y, retrain model, compute PFI scores (repeat)
 - ⇒ Permuting y breaks relationship to all features
 - \Rightarrow By computing PFI scores again, we obtain distribution of PFI scores under H_0



- PIMP was originally introduced for random forest's built-in permutation feature importance
- PIMP investigates whether the PFI score **significantly** differs from 0
 - ⇒ Useful because PFI can be non-zero due to stochasticity
- PIMP tests the H_0 -hypothesis: Feature is independent of the target y (unimportant)
- Sampling under H_0 : Permute target y, retrain model, compute PFI scores (repeat)
 - \Rightarrow Permuting v breaks relationship to all features
 - \Rightarrow By computing PFI scores again, we obtain distribution of PFI scores under H_0
- Compute p-value the tail probability under H_0 and use it as a new importance measure



TESTING IMPORTANCE (PIMP)

PIMP algorithm:

- For $m \in \{1, \ldots, n_{repetitions}\}$:
 - Permute response vector y
 - Retrain model with data X and permuted y
 - Compute feature importance PFI_i^m for each feature j (under H_0)



TESTING IMPORTANCE (PIMP)

PIMP algorithm:

- For $m \in \{1, \ldots, n_{repetitions}\}$:
 - Permute response vector y
 - Retrain model with data X and permuted y
 - Compute feature importance PFI_i^m for each feature j (under H_0)
- Train model with X and unpermuted y



TESTING IMPORTANCE (PIMP)

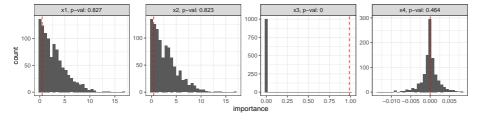
PIMP algorithm:

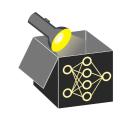
- For $m \in \{1, \ldots, n_{repetitions}\}$:
 - Permute response vector y
 - Retrain model with data X and permuted y
 - Compute feature importance PFI_i^m for each feature j (under H_0)
- Train model with X and unpermuted y
- **3** For each feature $j \in \{1, ..., p\}$:
 - Fit probability distribution of the feature importance values PFl_j^m , $m \in \{1, \dots, n_{repetitions}\}$ (choice between Gaussian, lognormal, gamma or non-parametric)
 - Compute feature importance PFI_j for the model without permutation of y (under H₁)
 - Retrieve the p-value of *PFI_i* based on the fitted distribution



PIMP FOR EXTRAPOLATION EXAMPLE

Recall: $y = x_3 + \epsilon_y$ with $\epsilon_y \sim N(0, 0.1)$, x_1 , x_2 highly correlated but independent of y, x_4 is independent of y and all other variables. Fitting a LM yields $\hat{f}(\mathbf{x}) \approx 0.3x_1 - 0.3x_2 + x_3$.

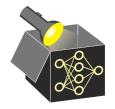




- Histograms: H_0 distribution of PFI scores after permuting y (1000 repetitions)
- Red: PFI score estimated on unpermuted y (under H₁) → compare against H₀ distribution
- Results: Although PFI for x_1 and x_2 is nonzero (red), PIMP considers them not significantly relevant (p-value > 0.05)

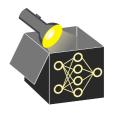
► Romano et al. (2010)

• When should we reject the H_0 -hypothesis for a feature?



➤ Romano et al. (2010)

- When should we reject the *H*₀-hypothesis for a feature?



▶ Romano et al. (2010)

- When should we reject the *H*₀-hypothesis for a feature?
- Accounting for multiplicity of individual tests can be achieved by controlling an appropriate error rate, e.g., the family-wise error rate (FWE: probability of at least one type-I error)



► Romano et al. (2010)

- When should we reject the H_0 -hypothesis for a feature?
- The larger the number of features, the more tests need to be performed by PIMP → Multiple testing problem: If multiplicity of tests is not taken into account, the probability that some of the true H₀-hypothesis is rejected (type-I error) by chance may be large
- Accounting for multiplicity of individual tests can be achieved by controlling an appropriate error rate, e.g., the family-wise error rate (FWE: probability of at least one type-I error)
- One classical method to control the FWE is the **Bonferroni correction** which rejects a null hypothesis if its p-value is smaller than α/m with m as the number of performed parallel tests

