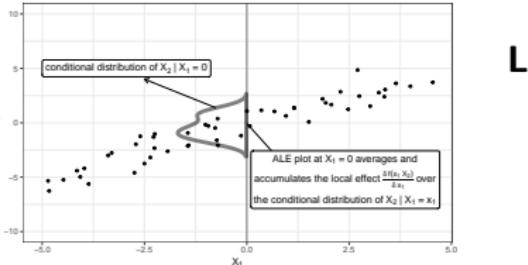


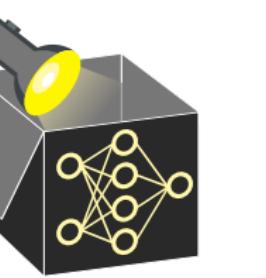
Interpretable Machine Learning

Regional Effects



Learning goals

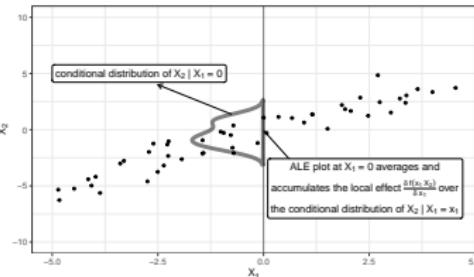
- Difference between feature effects and feature interactions
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Interpretable Machine Learning

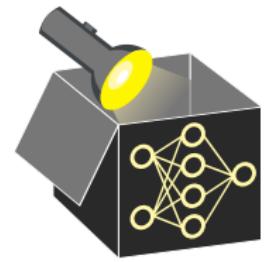
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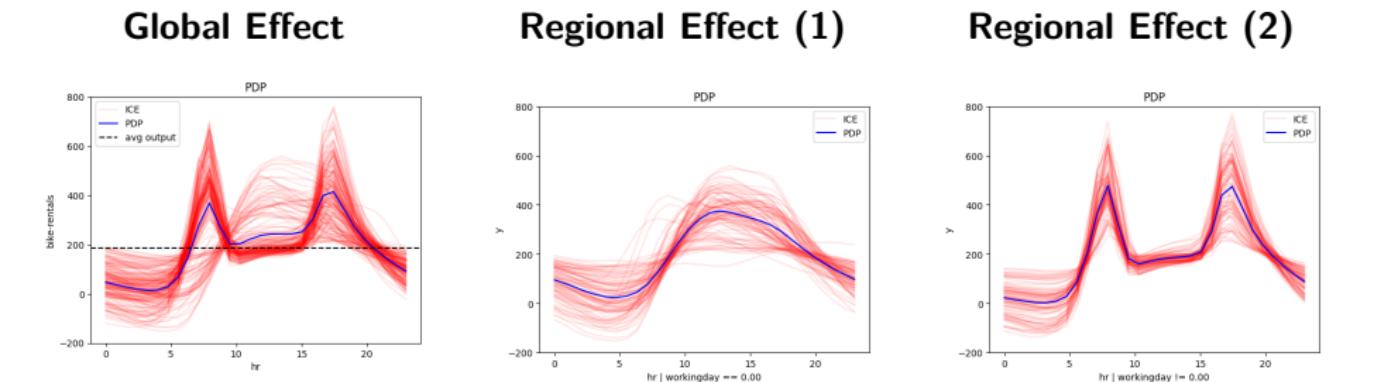
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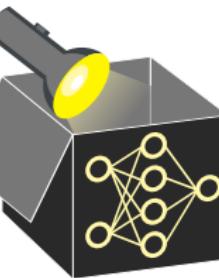
WHY REGIONAL EXPLANATIONS?

Problem: PD & ICE plots can be confounded by feature interactions.

Solution: Group homogeneous ICE curves in such a way that reduces the presence of individual interaction effects within a group \rightsquigarrow Regional effect plots (REPs).



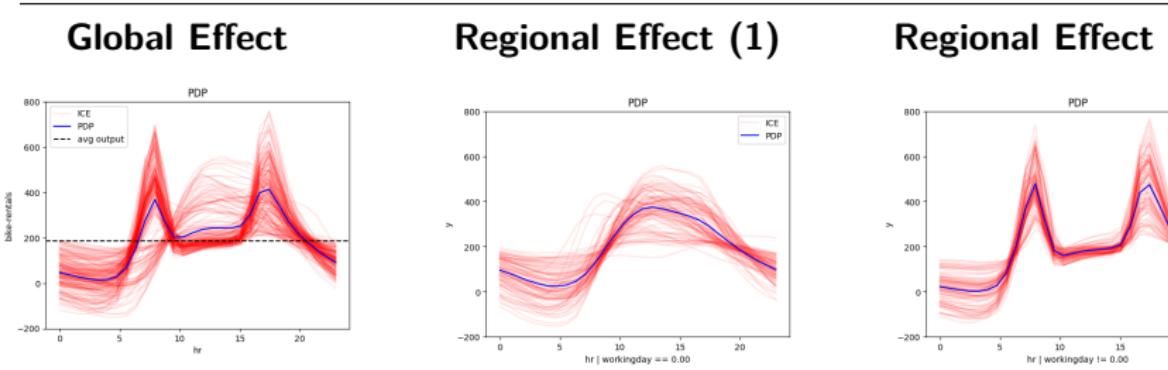
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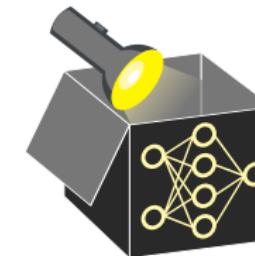
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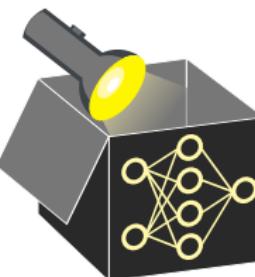


ICE CURVE: LOCAL FEATURE EFFECTS

Question: How do feature changes affect the prediction for **one observation**?

Idea: Split $\mathbf{x} = (x_j, \mathbf{x}_{-j})$ into x_j (feature of interest) and \mathbf{x}_{-j} (remaining features)

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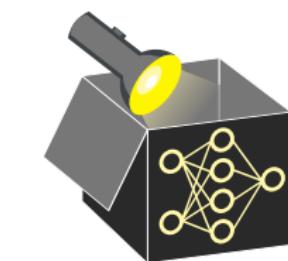


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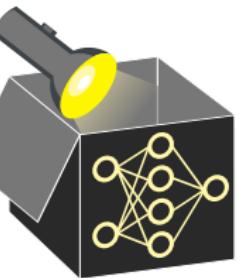
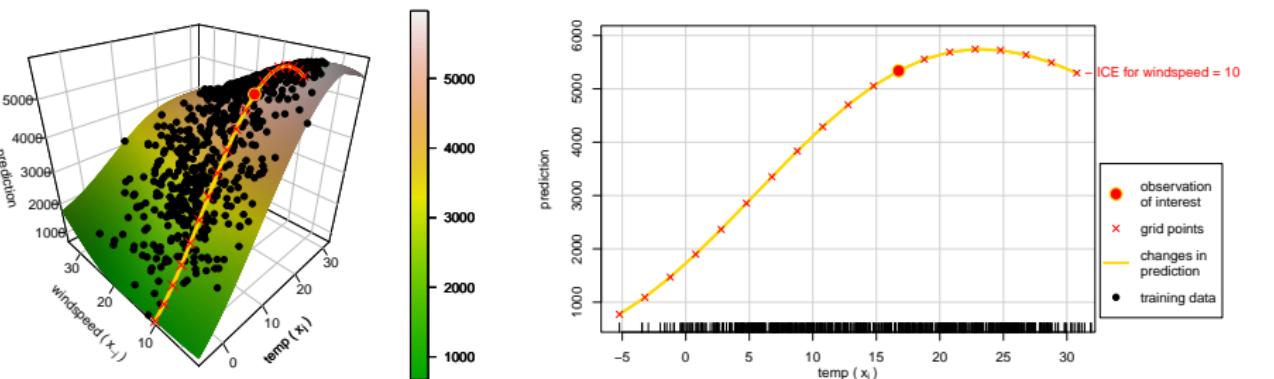
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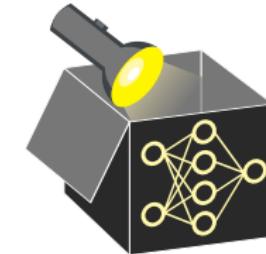
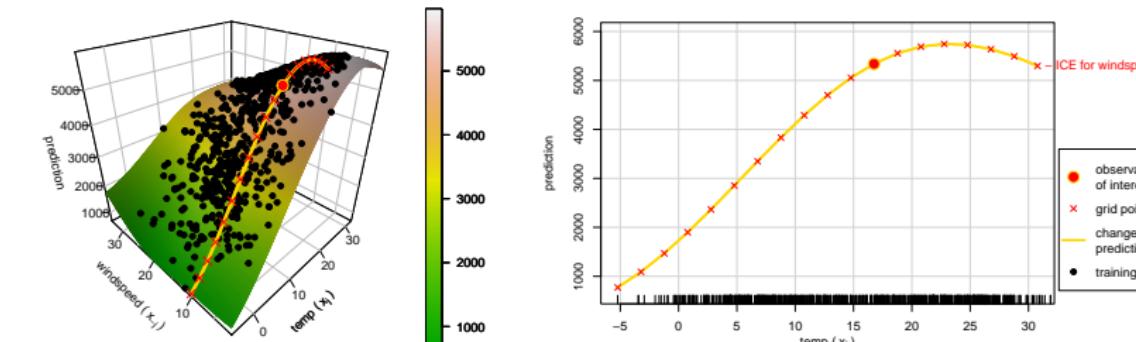
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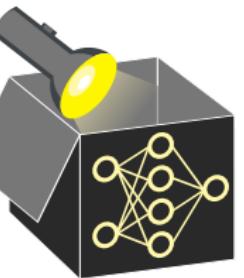
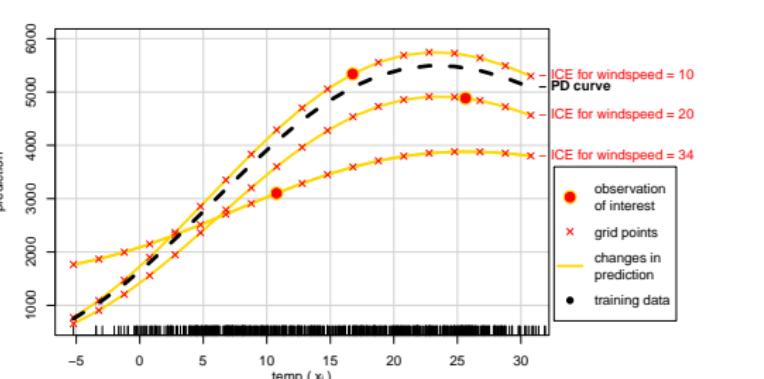
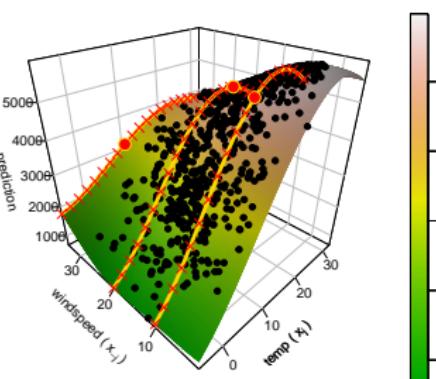
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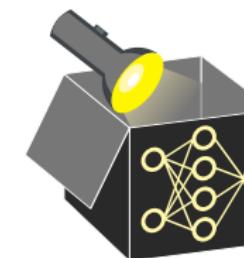
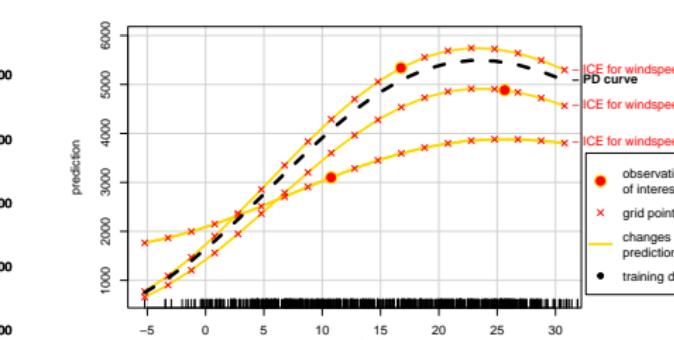
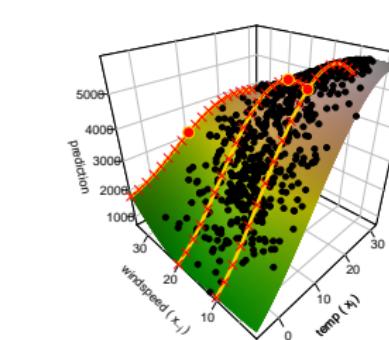
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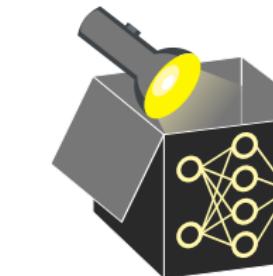
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FEATURE INTERACTIONS

Hooker (2004, 2007): Functional ANOVA decomposition of a function

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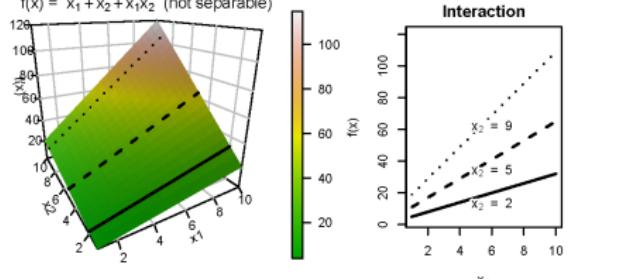


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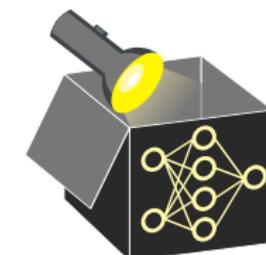
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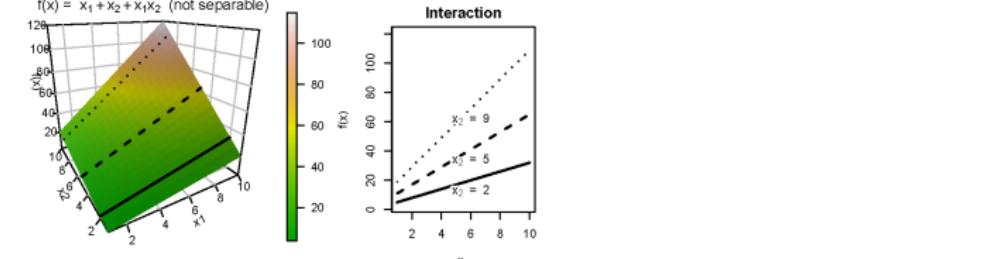


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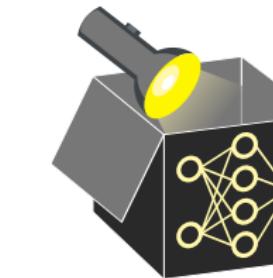
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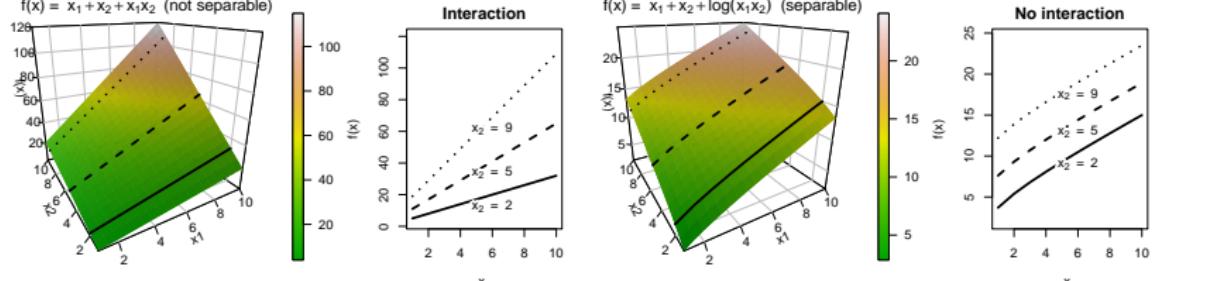


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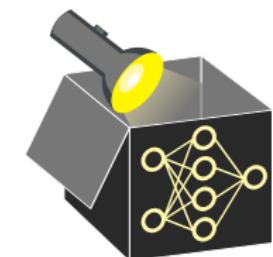
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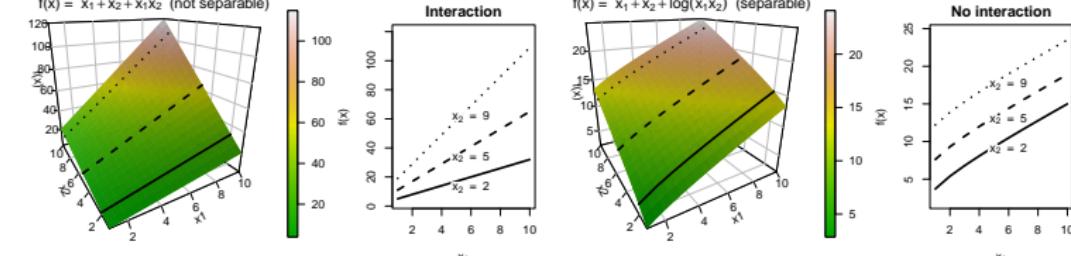


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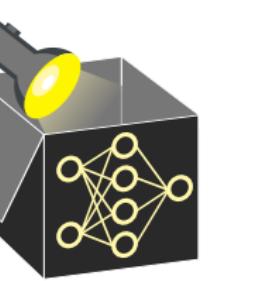
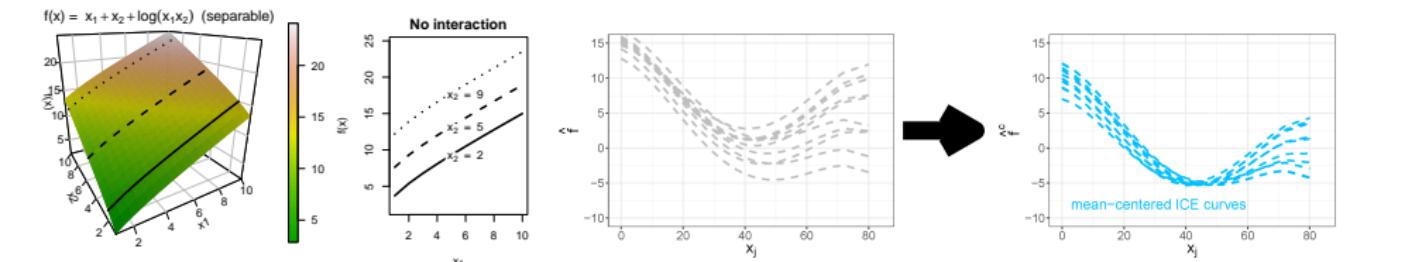
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Recall: Different shapes of ICE curves indicate interactions (ignore vertical shifts)
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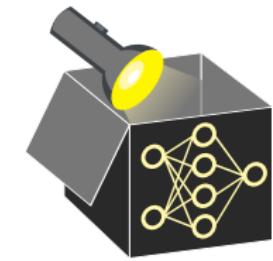
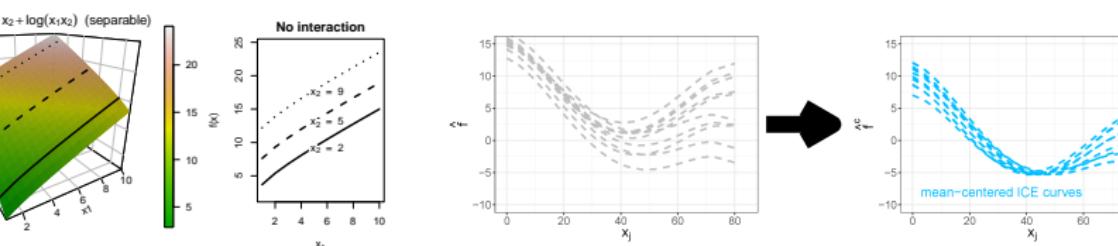
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REGIONAL EFFECTS - SYNTHETIC EXAMPLE

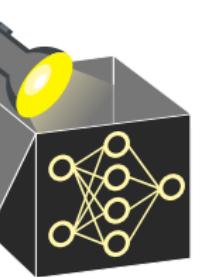
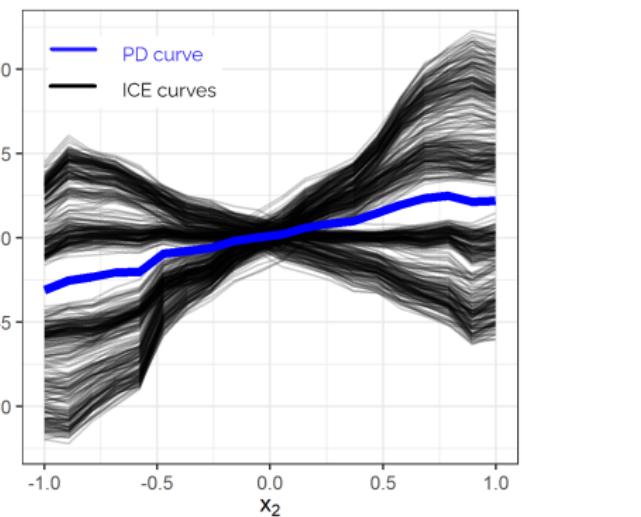
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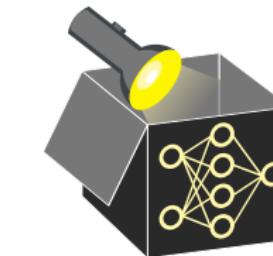
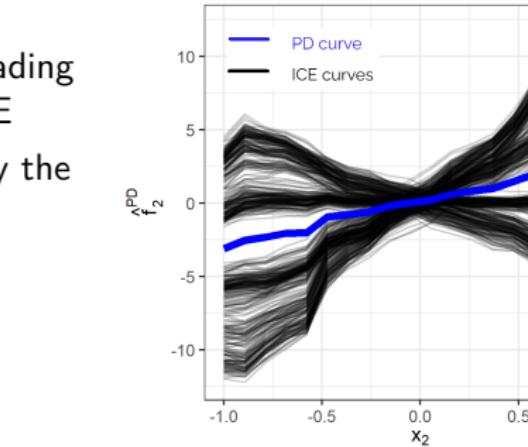
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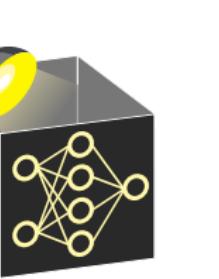
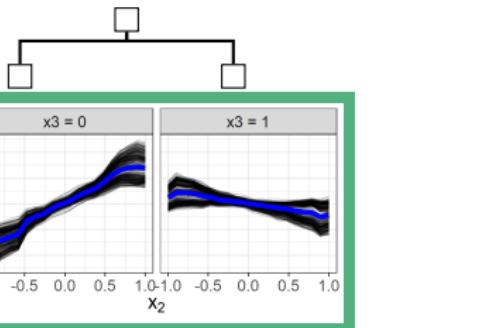
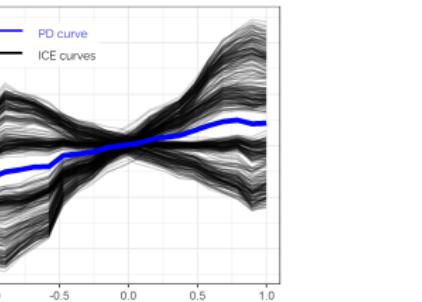
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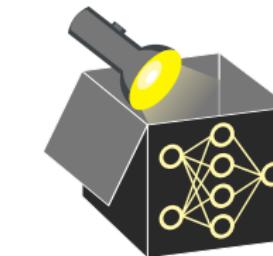
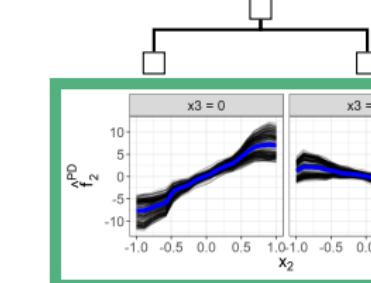
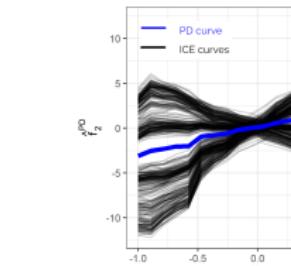
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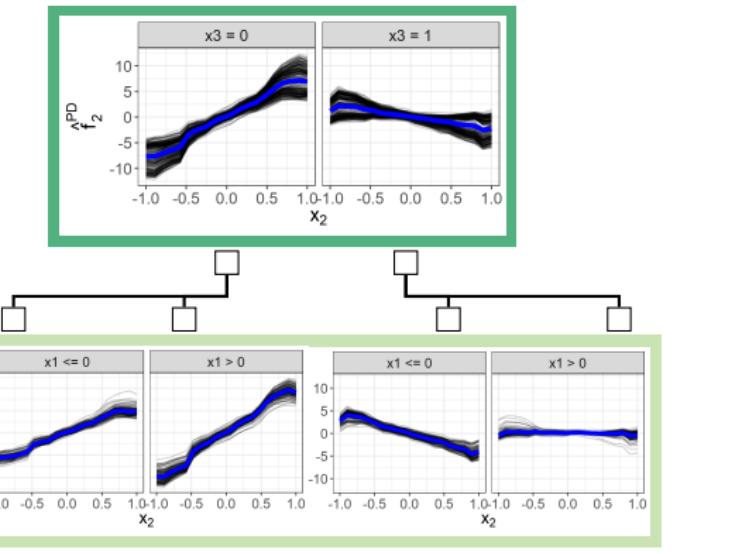
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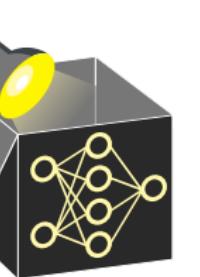
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Regional effect (blue curves) $\hat{\approx}$
Estimate PD curve in each region



\Rightarrow Additive decomposition of global feature effect



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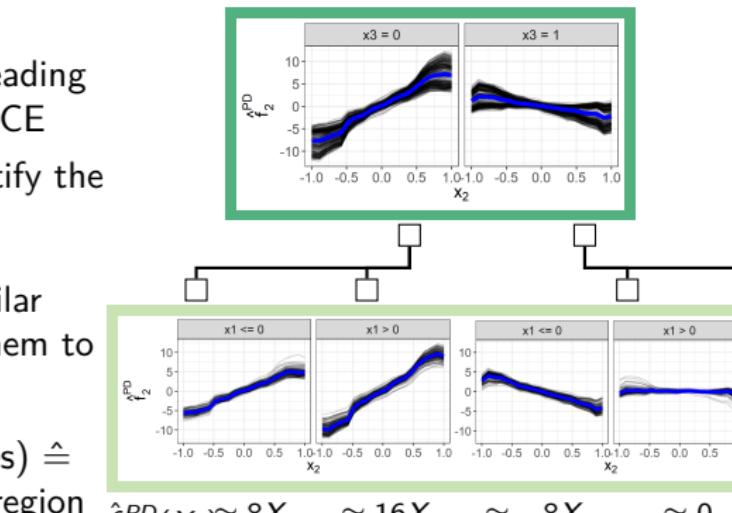
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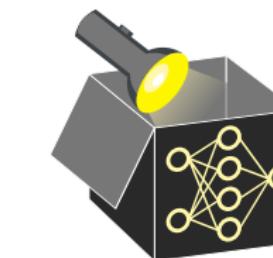
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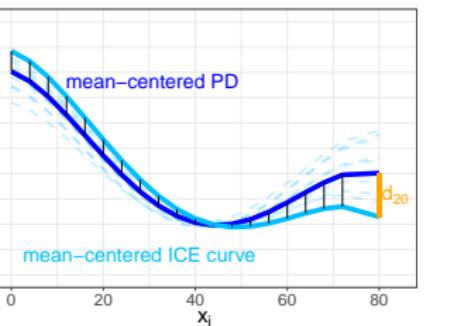


REGIONAL EFFECTS - DETAILS

Question: How to split the curves into regions?

Define risk as L2 loss of mean-centered ICE curves:

$$\mathcal{R}_j(\mathcal{N}) = \sum_{x \in \mathcal{N}} \sum_{k=1}^m \underbrace{(\hat{f}^c(\tilde{x}_j^{(k)}, x_{-j}) - \hat{f}_{j|\mathcal{N}}^{PD,c}(\tilde{x}_j^{(k)}))^2}_{d_k}$$



with the average feature effect in region $\mathcal{N} \subseteq \mathcal{X}$:

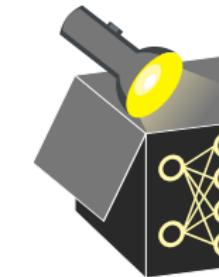
$$\hat{f}_{j|\mathcal{N}}^{PD,c}(\tilde{x}_j) = \frac{1}{|\mathcal{N}|} \sum_{x \in \mathcal{N}} \hat{f}^c(\tilde{x}_j, x_{-j})$$

- ⇒ Measures interaction-related heterogeneity (variance) of ICE curves in \mathcal{N}
- ⇒ Recursive partitioning (CART): Find best feature-split combination that solves

$$\arg \min_{z,t} \mathcal{R}_j(\mathcal{N}_{left}) + \mathcal{R}_j(\mathcal{N}_{right})$$

- $\mathcal{N}_{left} = \{x \in \mathcal{N} | x_z \leq t\}$
- $\mathcal{N}_{right} = \{x \in \mathcal{N} | x_z > t\}$
- Split point t for feature $x_z, z \in -j$

Intuition: Is another feature x_z responsible for the heterogeneity (measured by \mathcal{R}_j)?

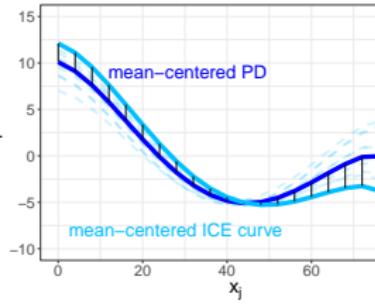


REGIONAL EFFECTS - DETAILS

Question: How to split curves into regions?

Define risk as L2 loss of mean-centered ICE curves:

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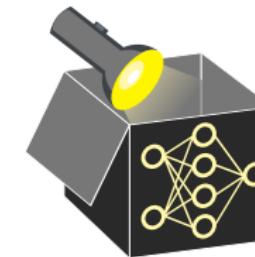
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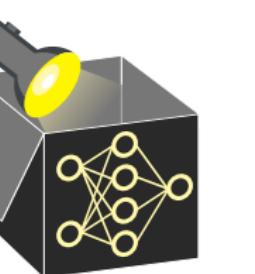
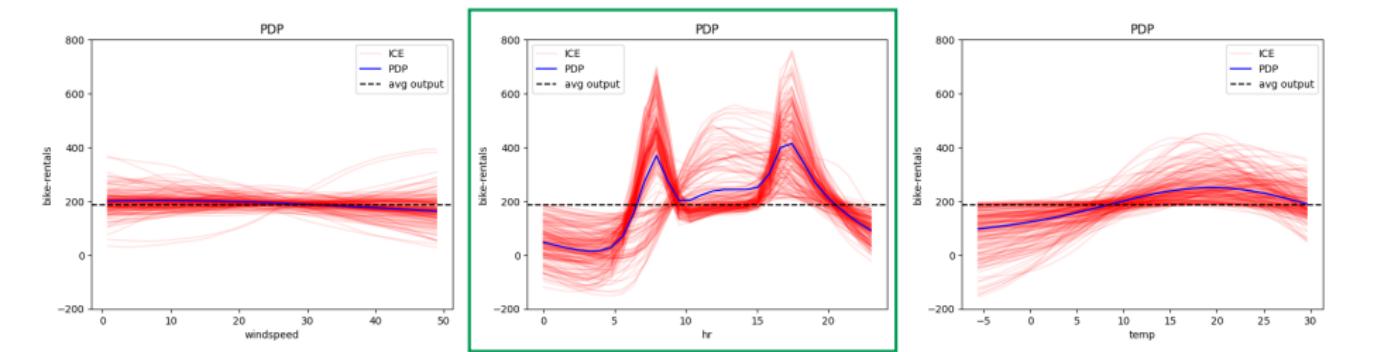
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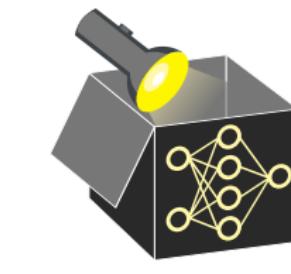
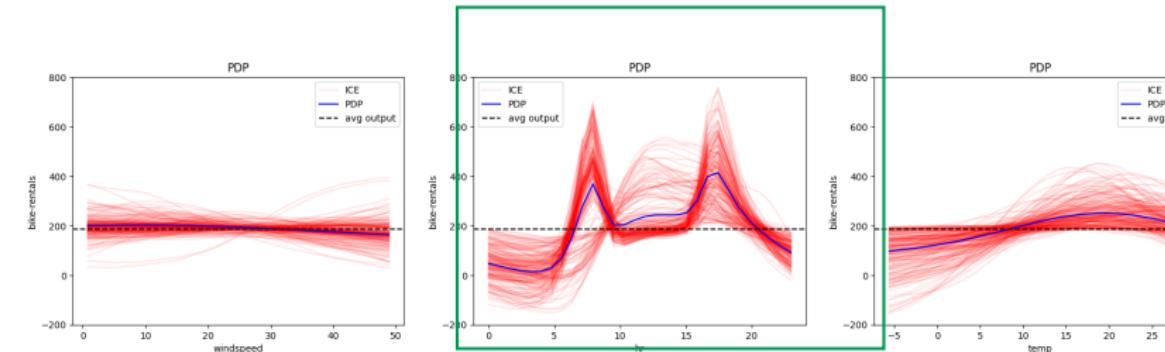
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REGIONAL EFFECT PLOTS - REAL EXAMPLE



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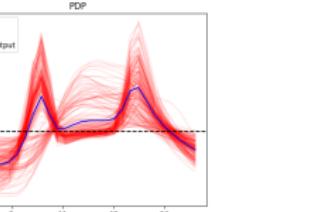


- Identify feature with highly heterogeneous local effects
 - ~~ **hour: Most important and highly heterogeneous feature (highest variance)**
- Find regions in feature space where this heterogeneity is minimal
 - ~~ Partition feature space using CART to minimize variance of mean-centered ICE curves within each region

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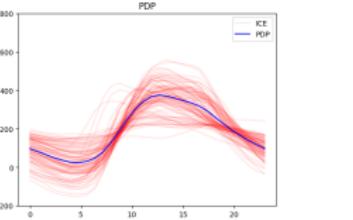
REGIONAL EFFECT PLOTS - REAL EXAMPLE

Regional effects of hour



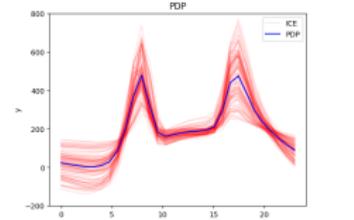
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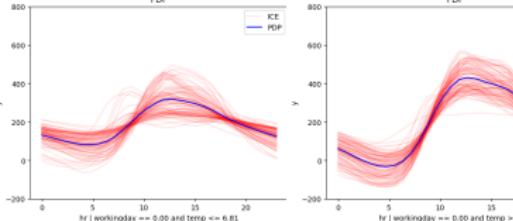
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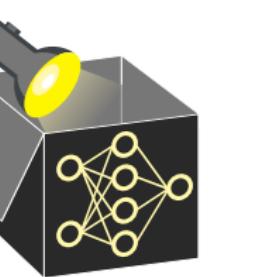
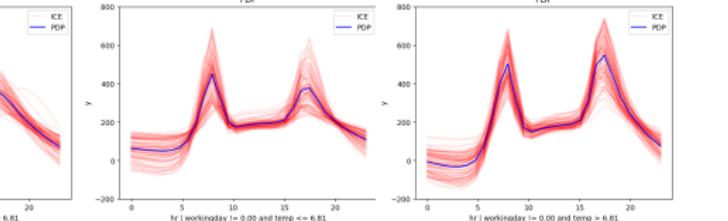
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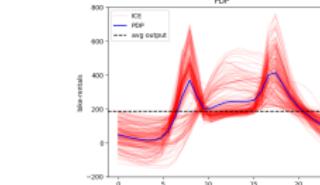
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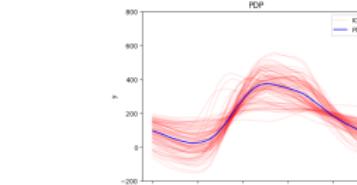
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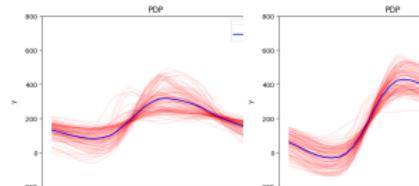
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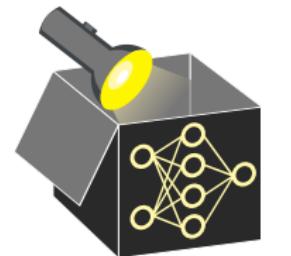
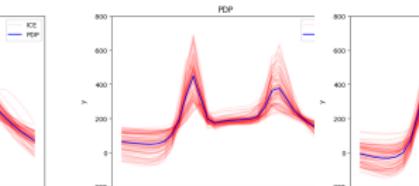
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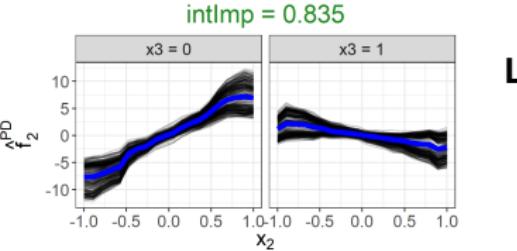
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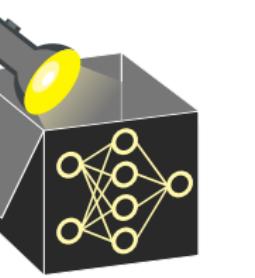
Interpretable Machine Learning

Regional Effects



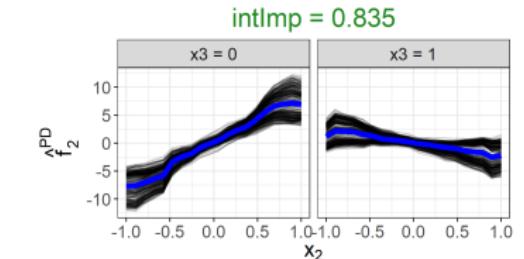
Learning goals

- Interaction quantification
- REPID interaction importance



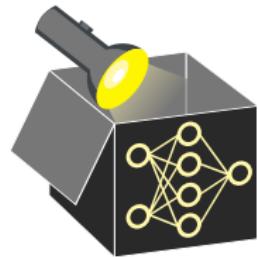
Interpretable Machine Learning

Regional Effects Interaction importance



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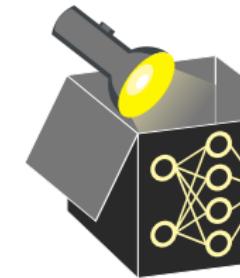


INTERACTION QUANTIFICATION

It's helpful to know not just how another feature changes the marginal effect of x_S but how strong that interaction is and want to rank features by it.

Approaches:

- **H-Statistics:** Variance of the deviation between the joint PDP and the sum of marginal PDPs (larger variance \Rightarrow stronger interaction).
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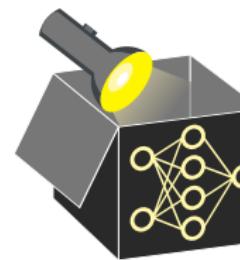


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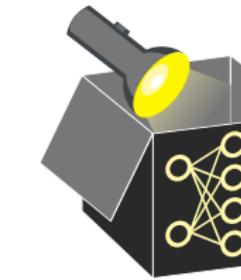
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Pitfalls:

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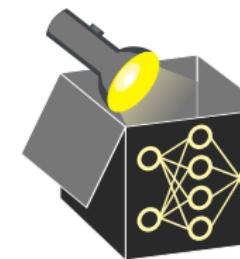
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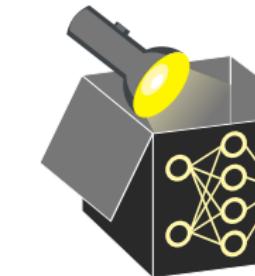
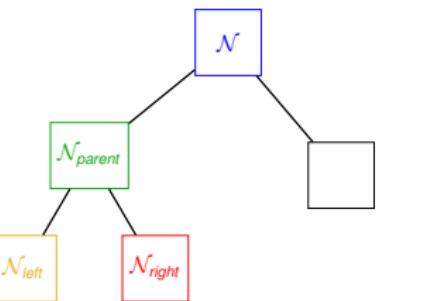


REPID INTERACTION IMPORTANCE

On parent node level (for \mathcal{N}_{parent}):

$$intImp(\mathcal{N}_{parent}) = \frac{\mathcal{R}(\mathcal{N}_{parent}) - (\mathcal{R}(\mathcal{N}_{left}) + \mathcal{R}(\mathcal{N}_{right}))}{\mathcal{R}(\mathcal{N})}$$

Interpretation: Reduction of ICE curve variance after one split of \mathcal{N}_{parent} into \mathcal{N}_{left} and \mathcal{N}_{right} relative to the ICE curve variance in the root node \mathcal{N} .

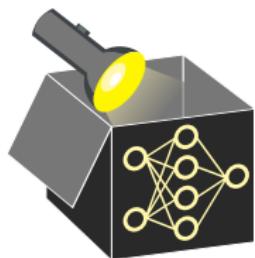
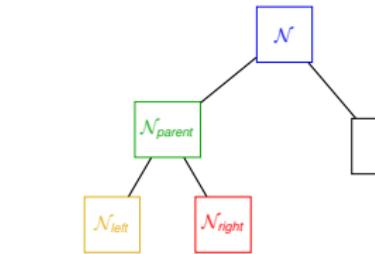


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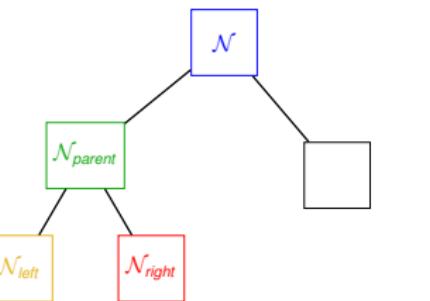


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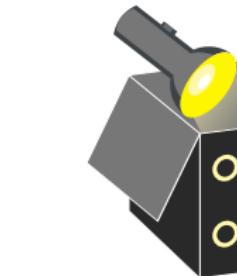
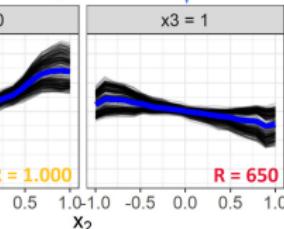
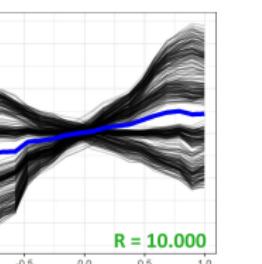
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Split reduces 83.5% of variance.

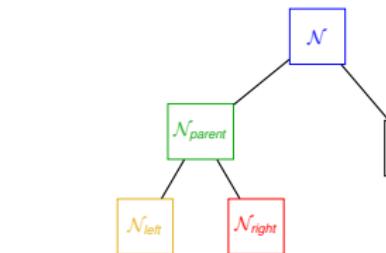


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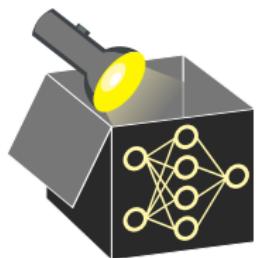
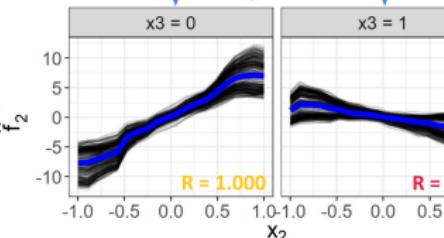
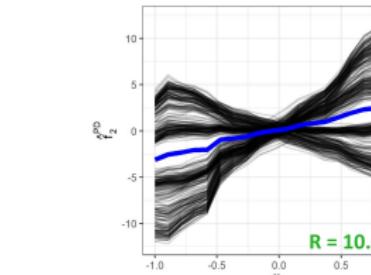
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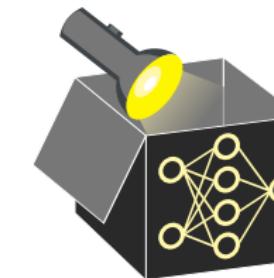
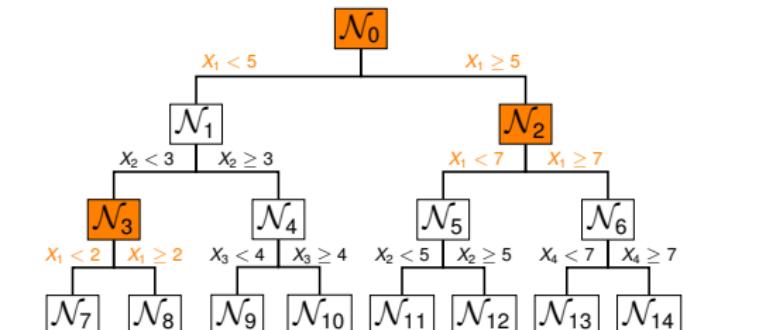
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$$intImp_j = \sum_{i \in \mathcal{B}_j} intImp(\mathcal{N}_i)$$

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Interpretation: Overall reduction of ICE curve variance due to splits by X_j (in %).

Example: For $X_1 \Rightarrow \mathcal{B}_1 = \{0, 2, 3\}$



REPID INTERACTION IMPORTANCE

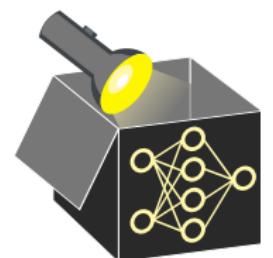
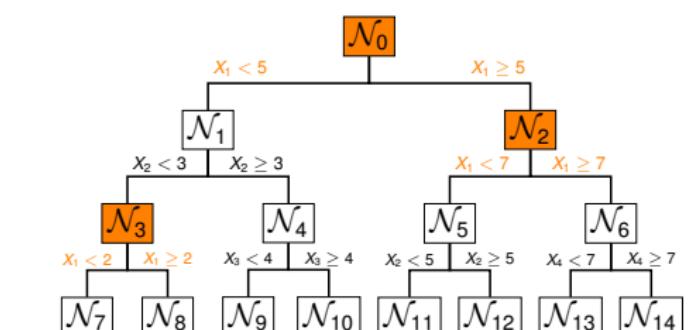
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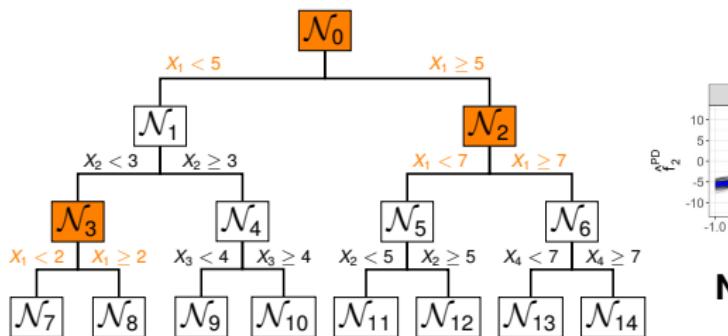
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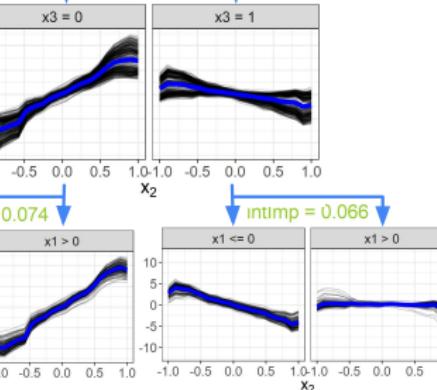
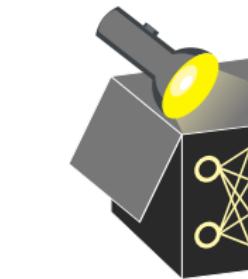
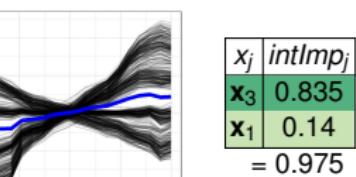
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Note: $intImp$ can also be used as a stopping criterion.



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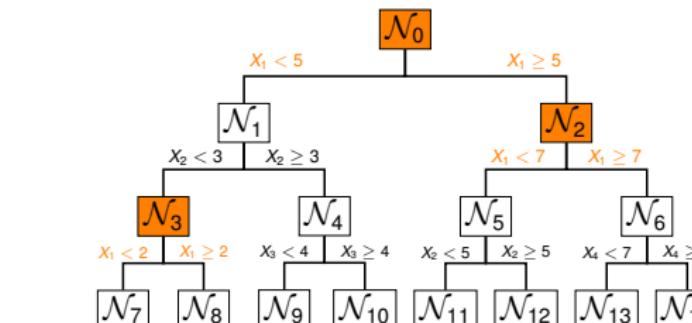
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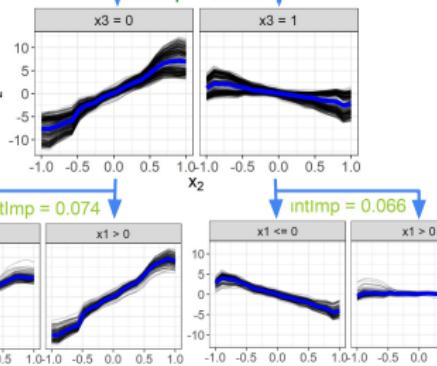
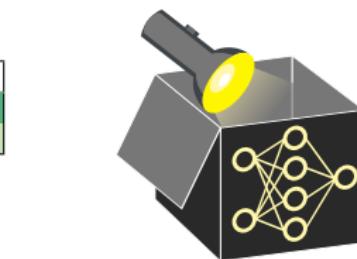
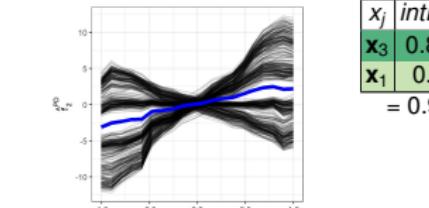
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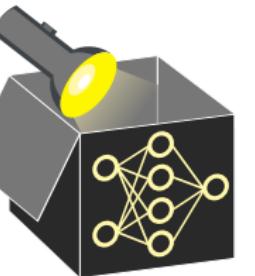


OUTPERFORMING SOTA

Simulation setting

- Draw 1000 i.i.d. samples from $X_1, \dots, X_4 \sim \mathcal{U}(-1, 1)$
- True underlying function: $f(\mathbf{x}) = \sum_{j=1}^4 \mathbf{x}_j + \mathbf{x}_1\mathbf{x}_2 + \mathbf{x}_2\mathbf{x}_3 + \mathbf{x}_1\mathbf{x}_3 + \mathbf{x}_1\mathbf{x}_2\mathbf{x}_3 + \epsilon$
- Fit a correctly specified linear model (interactions with \mathbf{x}_4 are excluded)
- 30 repetitions, measure interaction strength between \mathbf{x}_2 and all other 3 features

Which methods are sensitive to changes in main effect sizes or feature correlations?



Pitfall	REPID	H-Statistic	Greenwell	SHAP
sensitive to changes of main effect	No	Yes	Yes	No
sensitive to changes of correlation between \mathbf{x}_j and other features	No	Yes	No	Yes

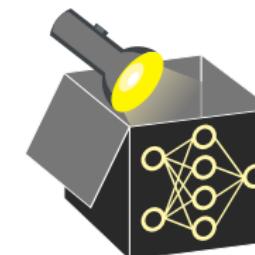
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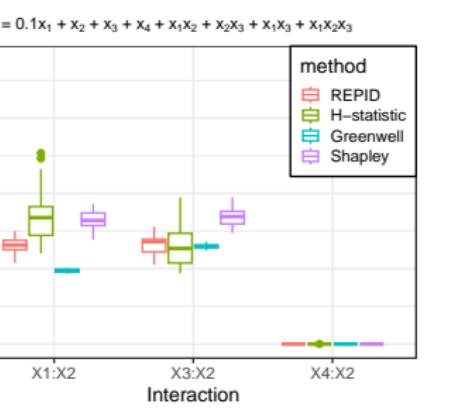
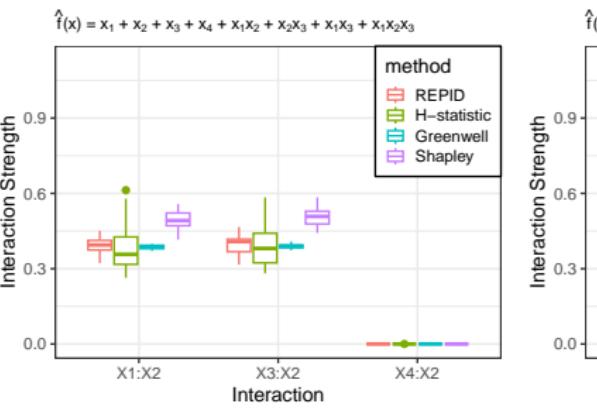
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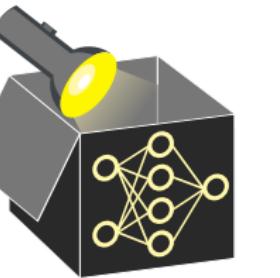
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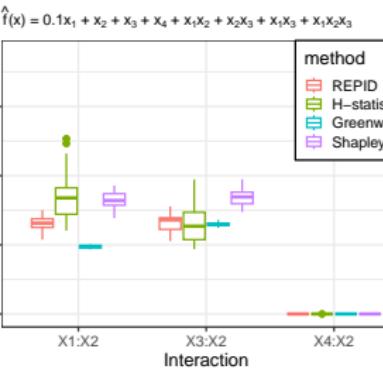
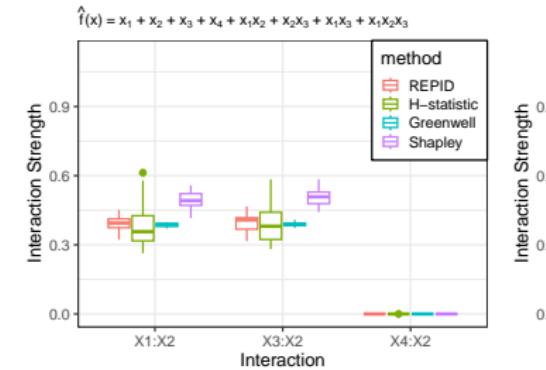
OUTPERFORMING SOTA



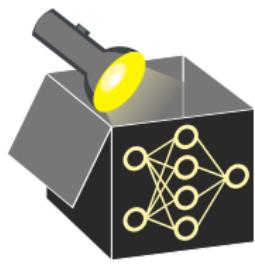
- **Left (initial setting):** Interaction strength of $x_1:x_2$ and $x_3:x_2$ similar; $x_4:x_2$ no interaction
- **Right:** Set main effect $\beta_1 = 0.1$
 - **Expectation:** Interaction strengths should not change
 - **Fail:** H-statistic ($x_1:x_2 > x_3:x_2$) and Greenwell ($x_1:x_2 < x_3:x_2$)



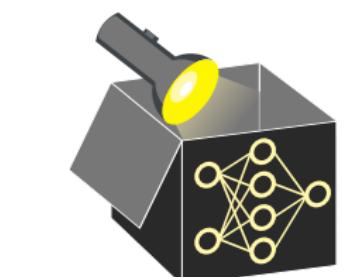
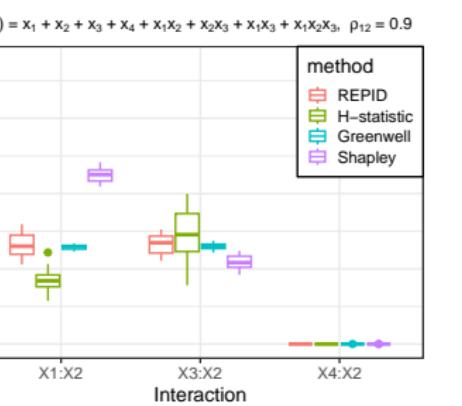
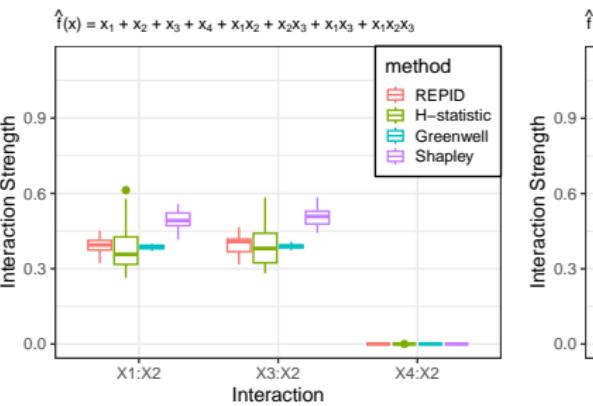
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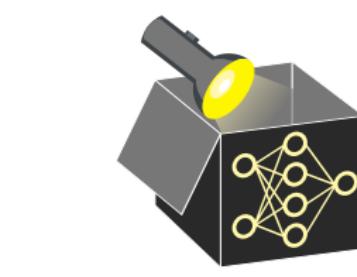
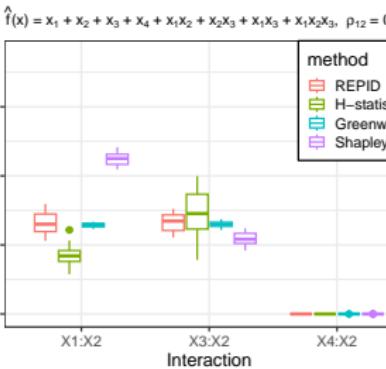
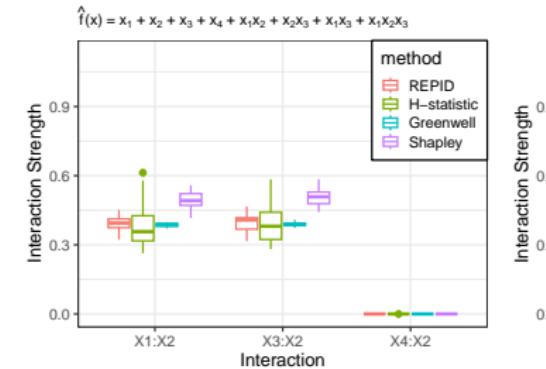


OUTPERFORMING SOTA



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- **REPID is the only method which always leads to correct rankings for these settings**

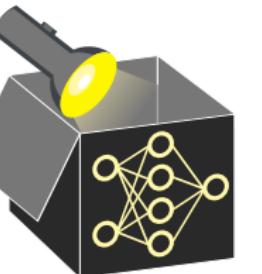
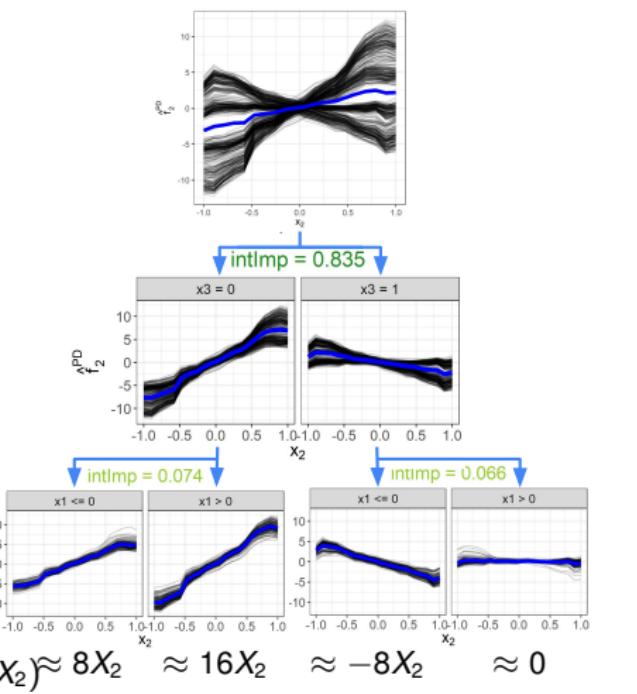
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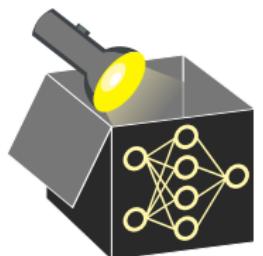
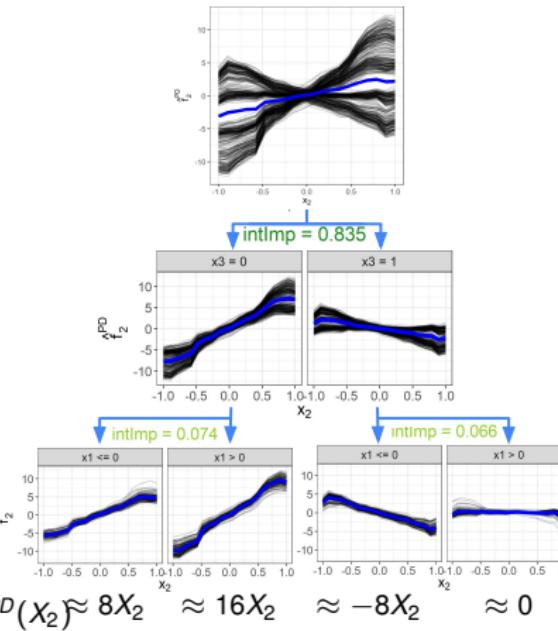
LIMITATIONS OF REPID

- 1) Restricted to one feature of interest
⇒ Different regions for different features



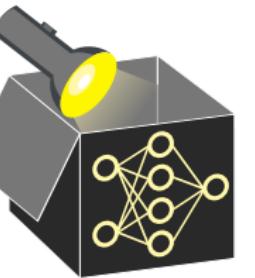
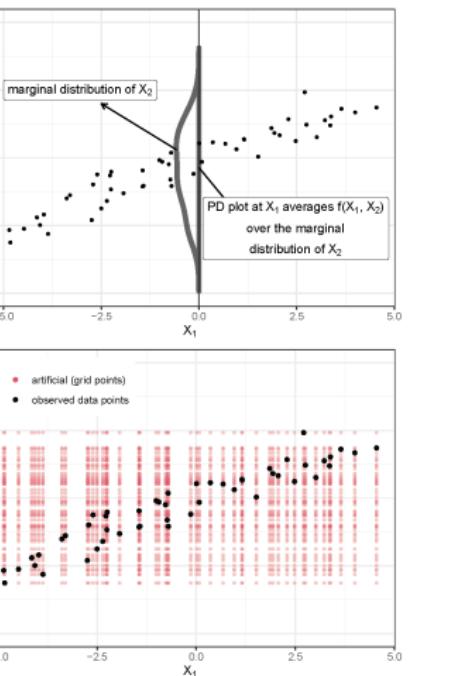
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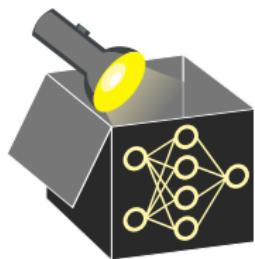
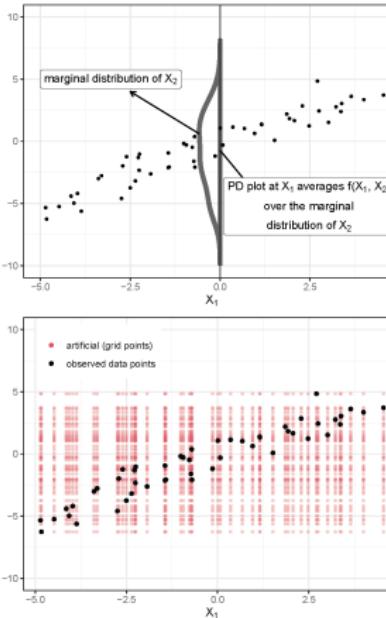
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CONCLUSION

Summary of Contributions (REPID):

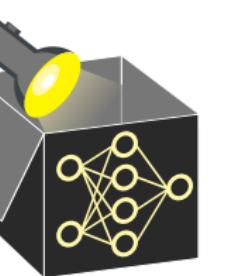
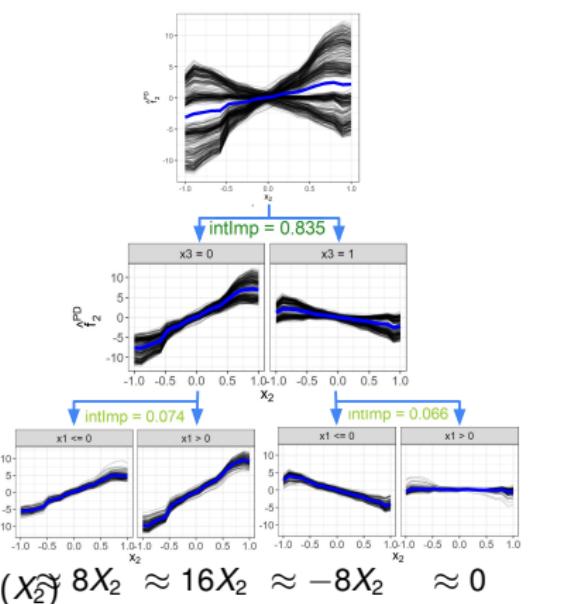
- Regional effects in interpretable regions
- Additive decomposition of feature effect
- Quantify feature interactions
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Further Directions:

Pruning, GADGET as a predictor, comparing regions across models, efficient implementation, more efficient testing and splitting approach, ...



CONCLUSION

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