

Uncertainties in Predictive Inference: Conformal Inference and Cross-Validation

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Regression and Prediction

Data: $(X_i, Y_i)_{i=1}^n$ i.i.d from joint distribution with

$$Y = \mu(X) + \varepsilon$$

where

$$\mathbb{E}(\varepsilon | X) = 0.$$

Goal

1. learn about μ (estimation).
2. predict Y for future observations of X .

Predictive inference

- We would like to quantify the uncertainty of Y for each X observed in the future or in the sample.
 1. Noise uncertainty: even if we knew μ perfectly, we never observe ε .
 2. Sampling uncertainty: empirical distribution as approximation to underlying population.
 3. Modeling uncertainty: popular assumptions, such as Gaussianity of ε , linearity/smoothness of μ , sparsity, etc, may not be exactly correct.

Examples of assumptions

- Classical nonparametric regression
 - μ is smooth (e.g., Hölder class)
 - X has density bounded away from 0
 - $(\varepsilon | X) \sim N(0, \sigma^2)$ or similar
- High dimensional regression
 - $\mu(x) = \beta^T x$ and β is sparse
 - the design matrix is nice (incoherence, RIP, etc)
 - $(\varepsilon | X) \sim N(0, \sigma^2)$ or similar
- Neural network: μ can be written as compositions of (structured) multiple index models.
- Inferences based on these assumptions may not be robust.

Outline

- Conformal inference: reliable prediction band under no structural assumptions (joint work with L. Wasserman, R. J. Tibshirani, M. G'Sell, A. Rinaldo)
- Cross-validation with confidence: make better use of validated loss in sampling-splitting.

A naive prediction band

- Data: $(X_i, Y_i)_{i=1}^n$; Goal: predict Y_{n+1} for a future X_{n+1} .
- Estimate $\hat{\mu}$ (OLS, local polynomial, lasso, NN, etc)
- $R_i = |Y_i - \hat{\mu}(X_i)|$, or any other loss function.
- Prediction band:
$$\hat{\mu}(X_{n+1}) \pm \text{upper } \alpha\text{-quantile of } \{R_i : 1 \leq i \leq n\}.$$
- OK only if $\hat{\mu}$ is very accurate, which requires standard assumptions, as well as good choices of tuning parameters.
- **Overfitting**: this prediction band tends to be too narrow, because the fitted residuals are smaller than the true values.

Conformal Prediction

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- Output $\hat{C}(X_{n+1}) = \{y \in \mathbb{R} : \pi_n(y) \leq 1 - \alpha\}$.

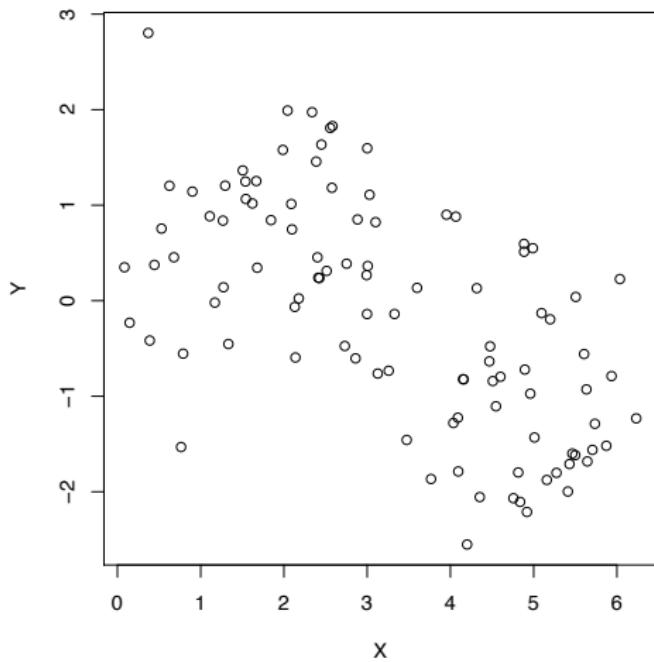
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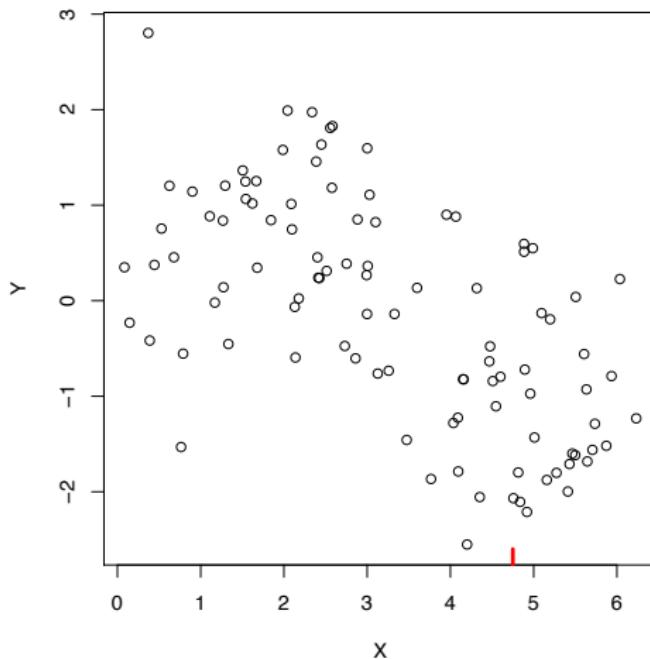
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- **Theorem:** $\mathbb{P}(Y_{n+1} \in \hat{C}(X_{n+1})) \geq 1 - \alpha$, if $(X_i, Y_i)_{i=1}^{n+1}$ is iid.

Example: conformal prediction interval using smoothing splines

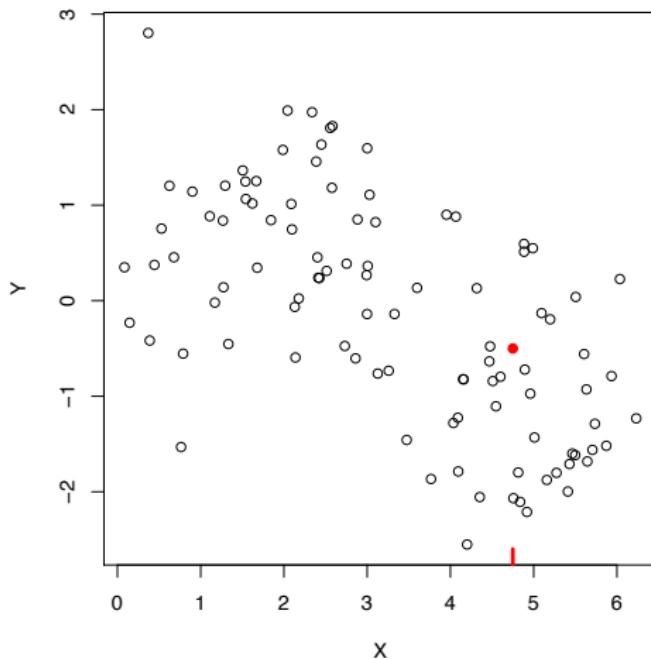


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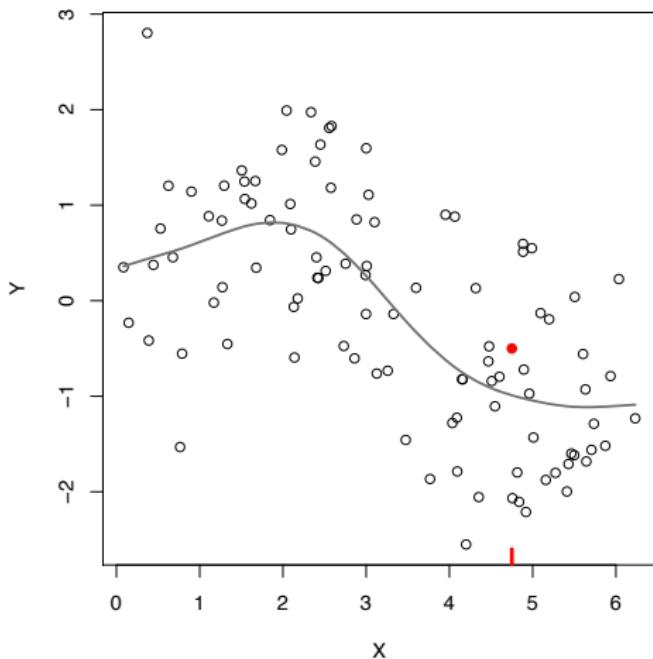
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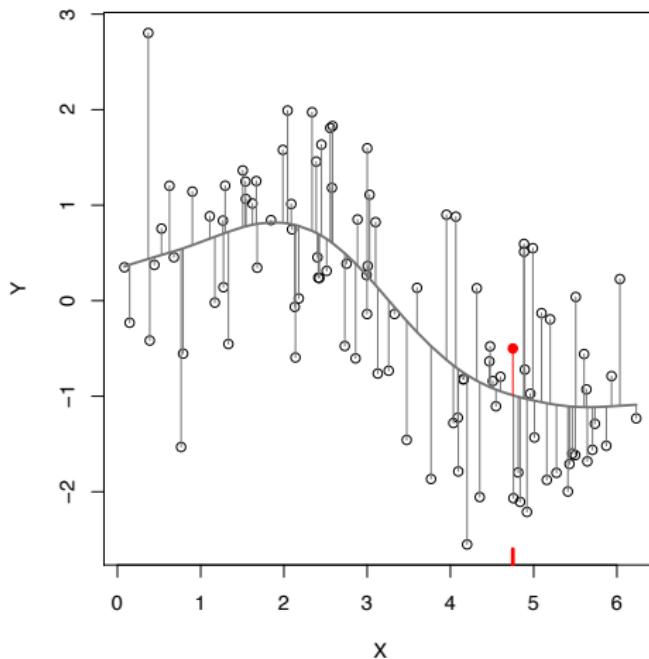
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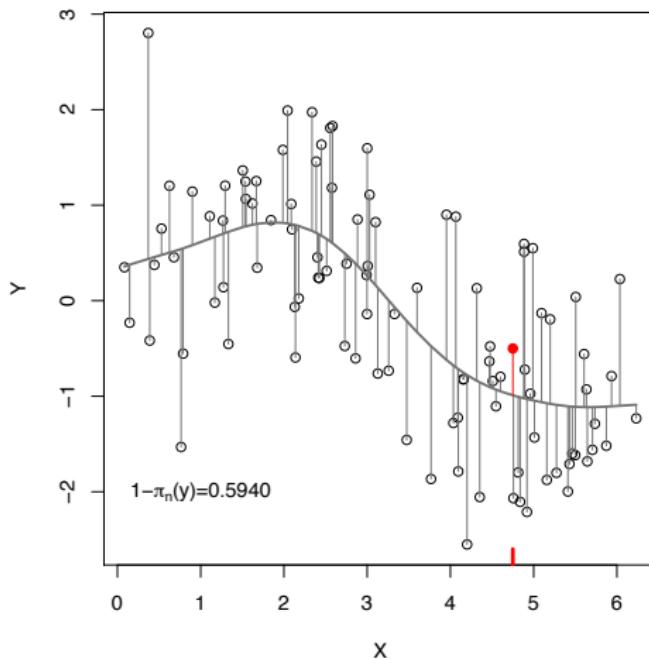
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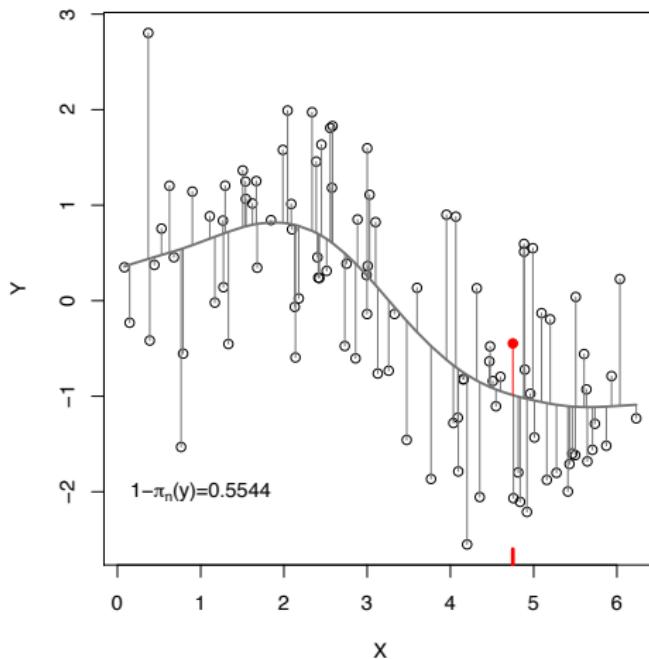
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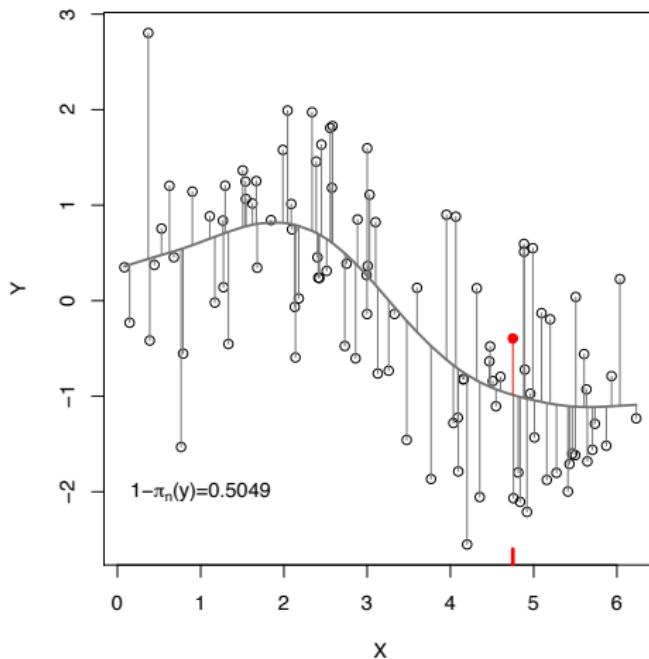
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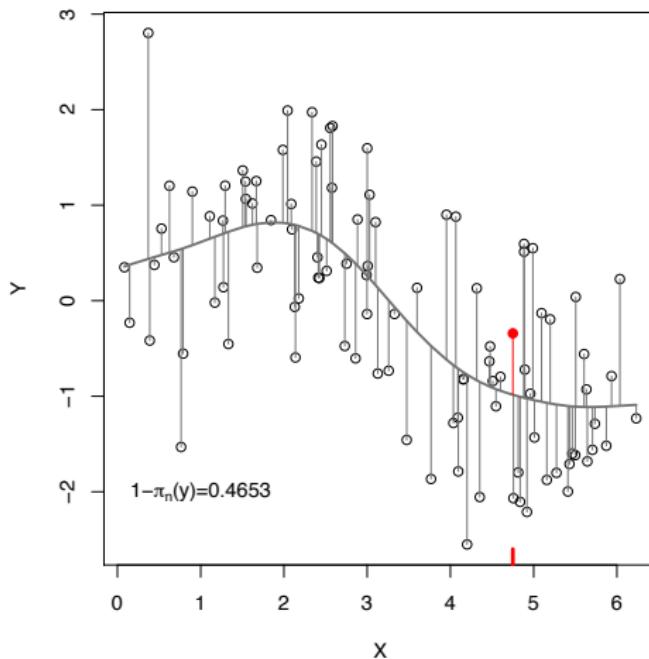
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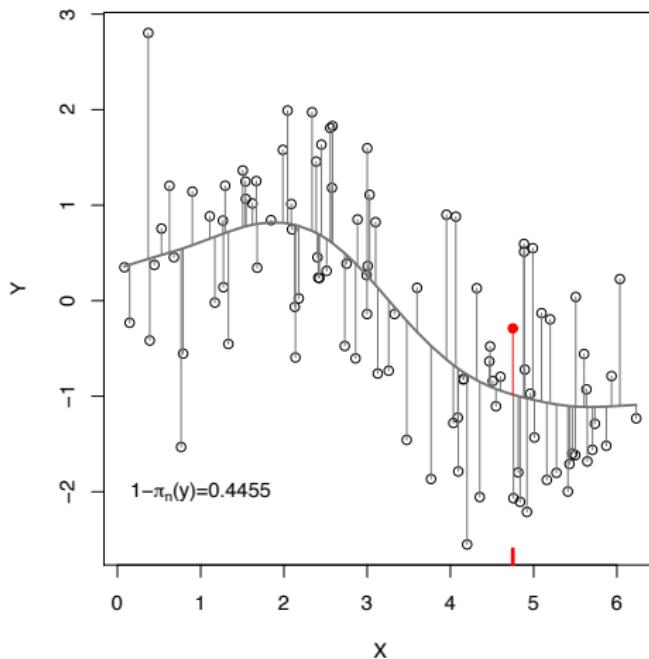
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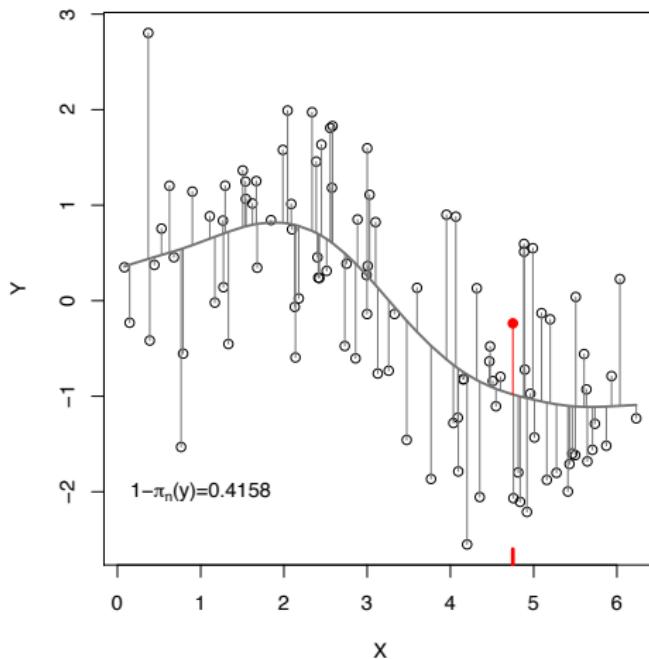
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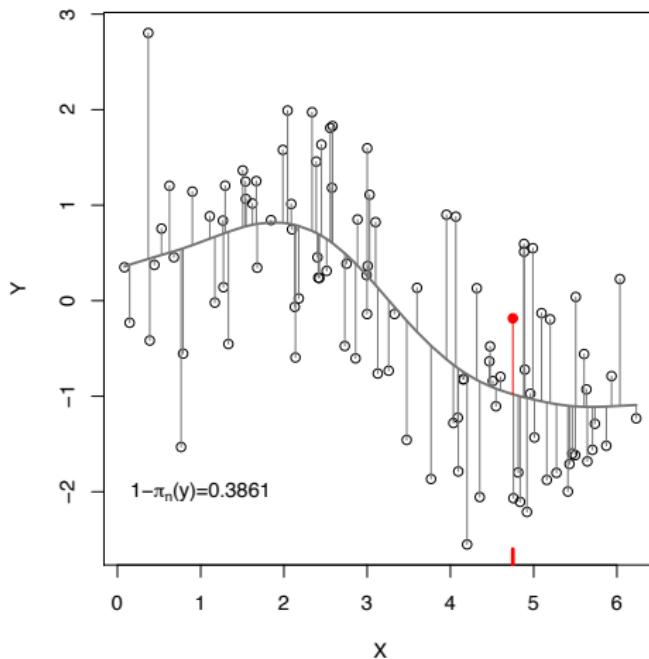
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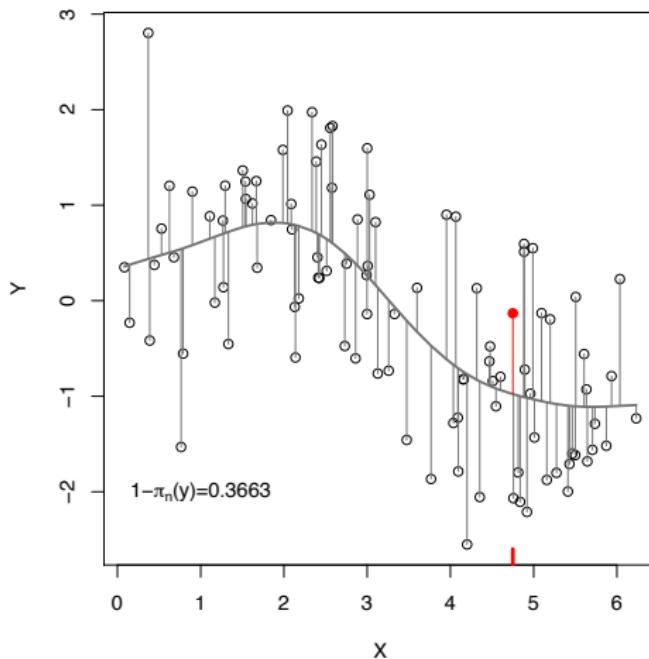
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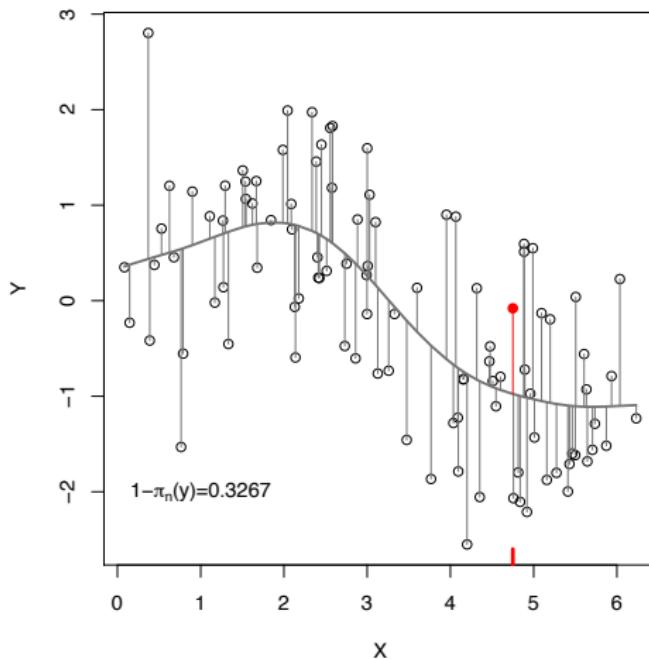
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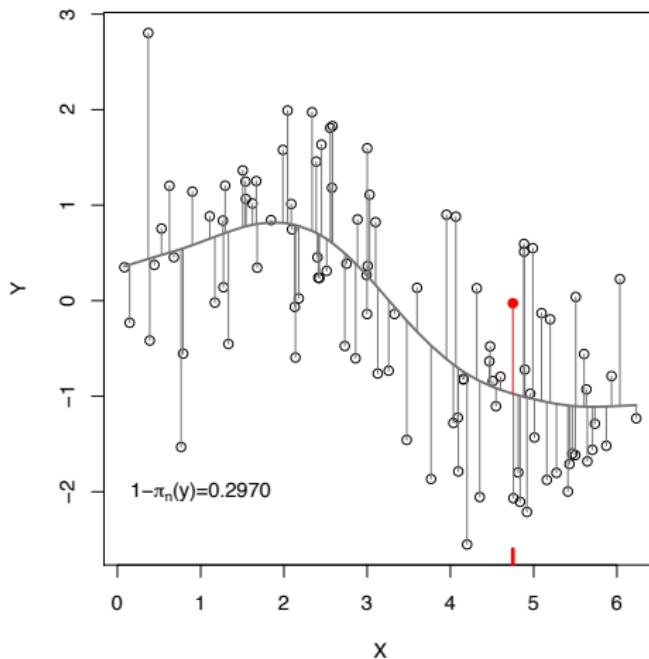
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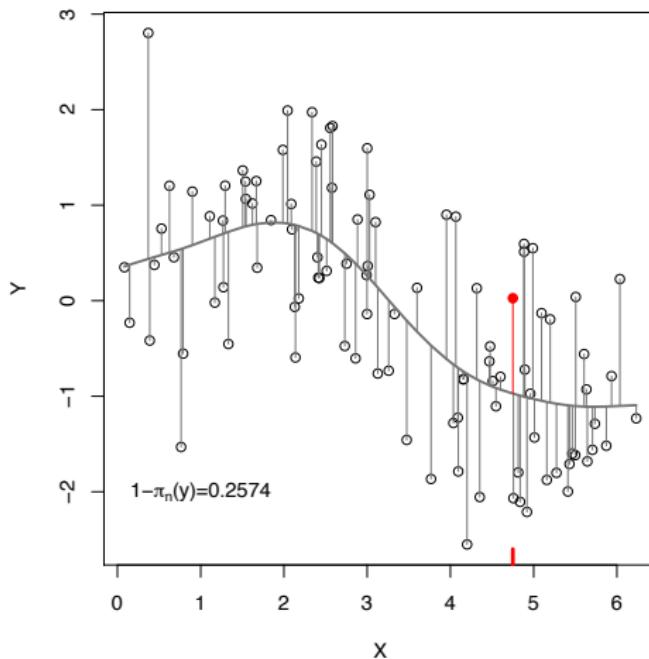
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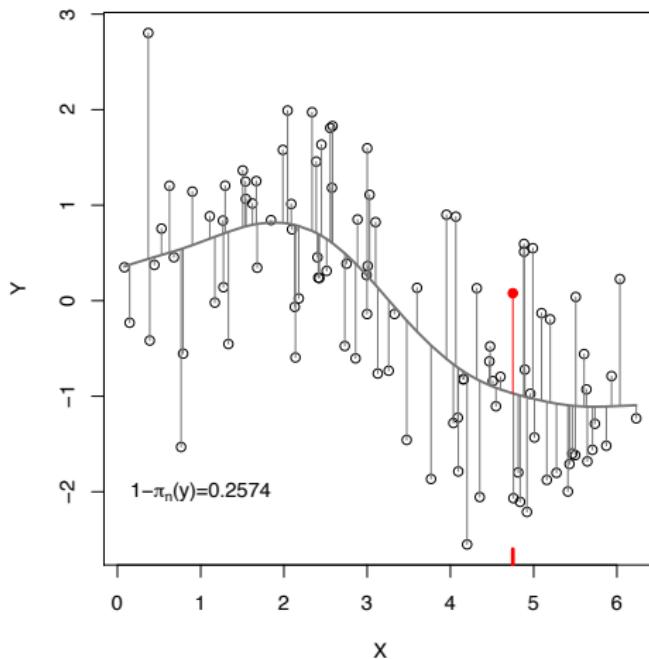
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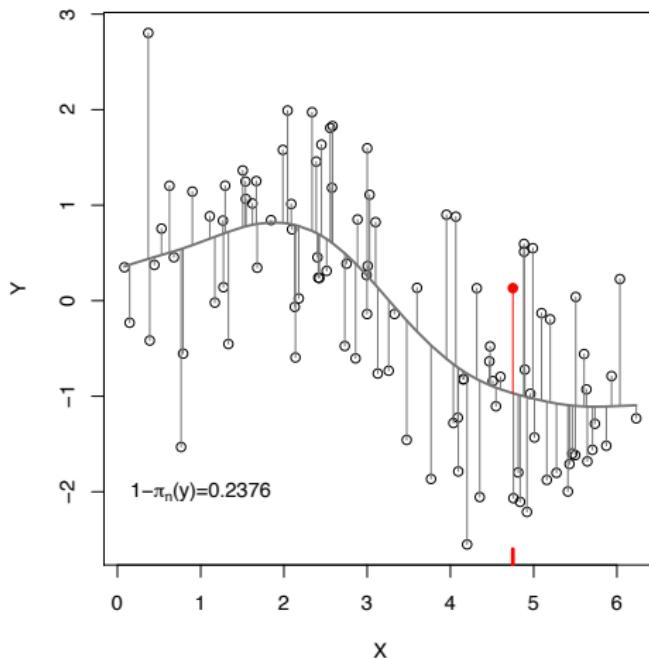
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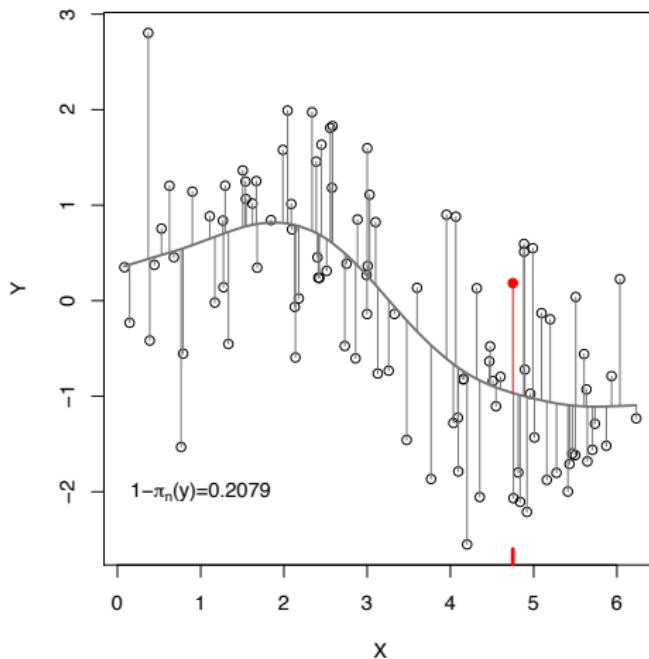
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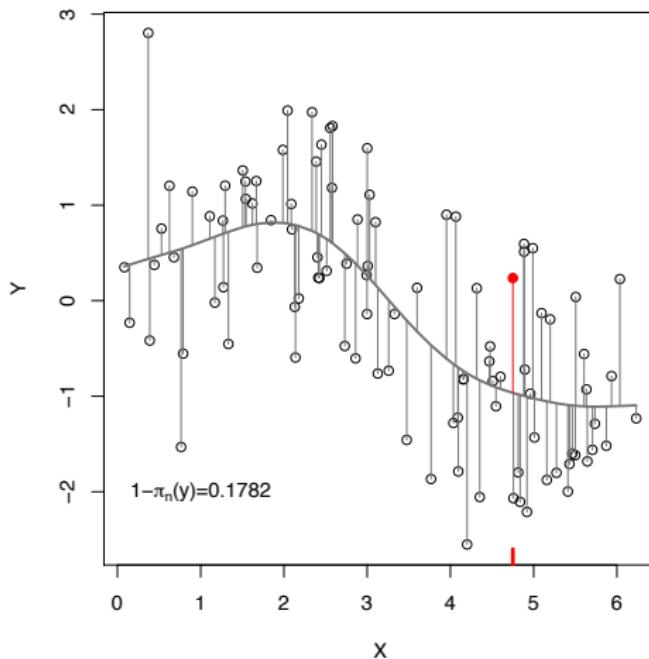
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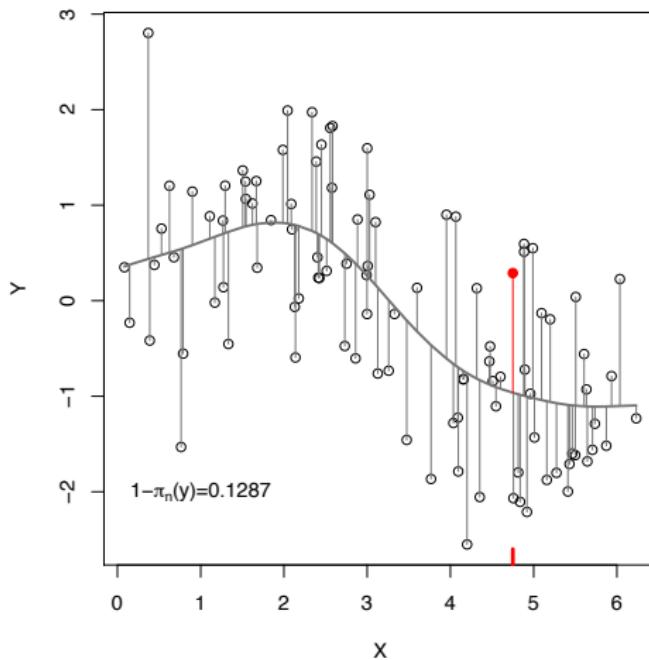
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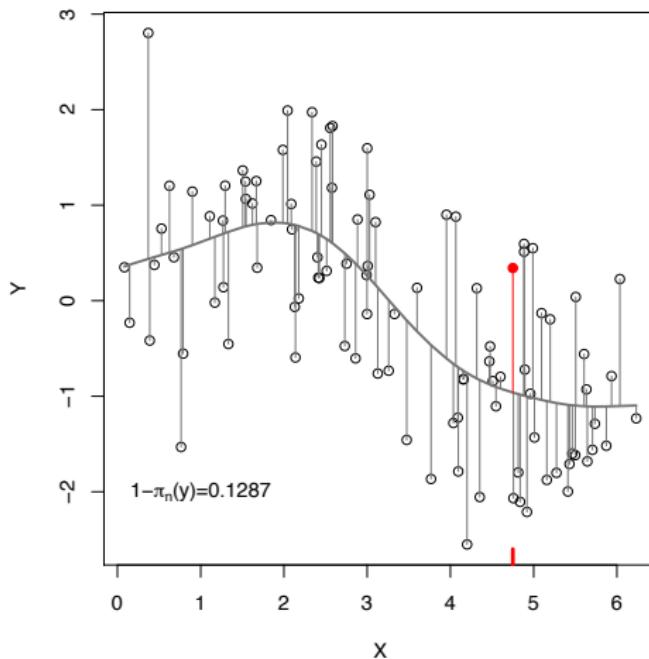
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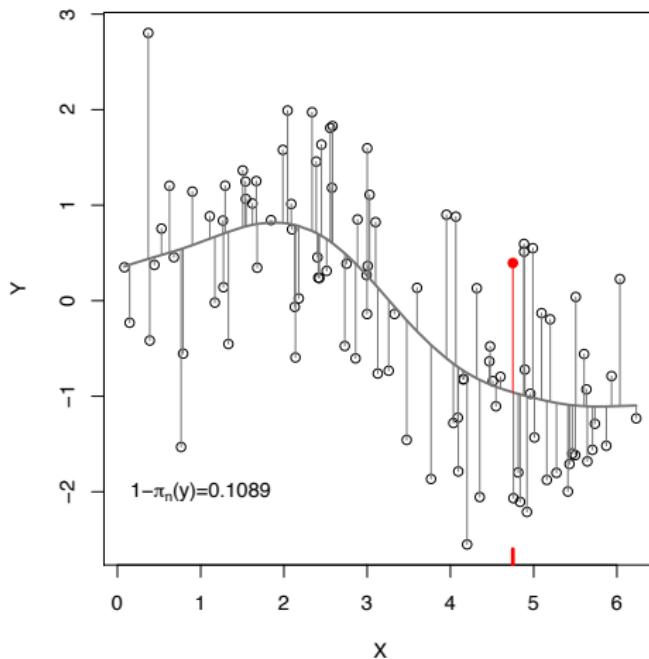
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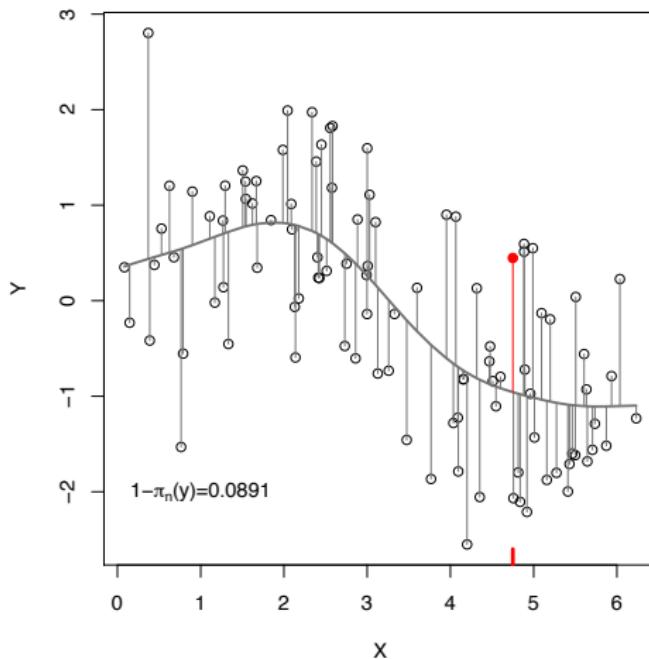
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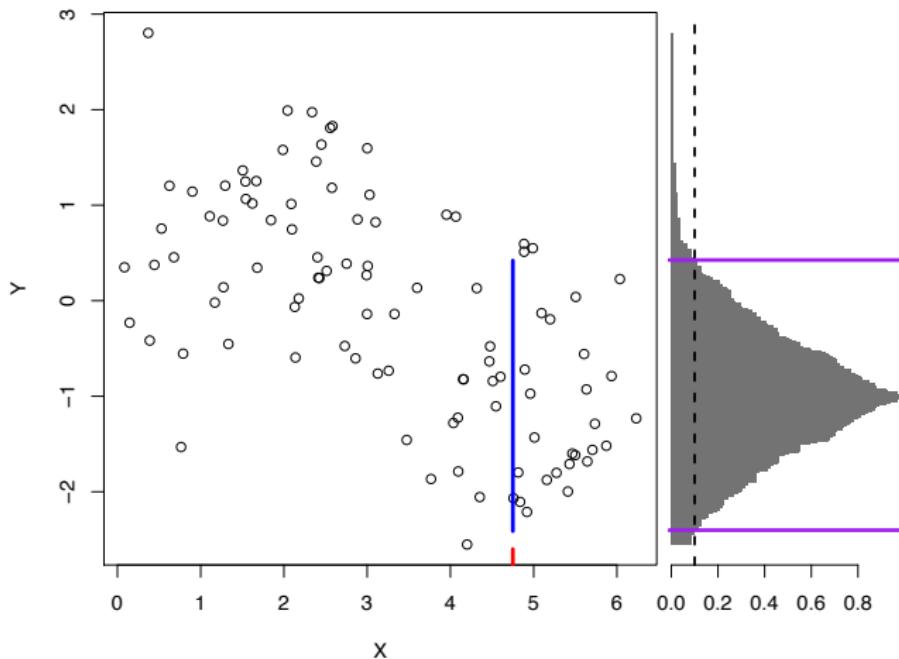
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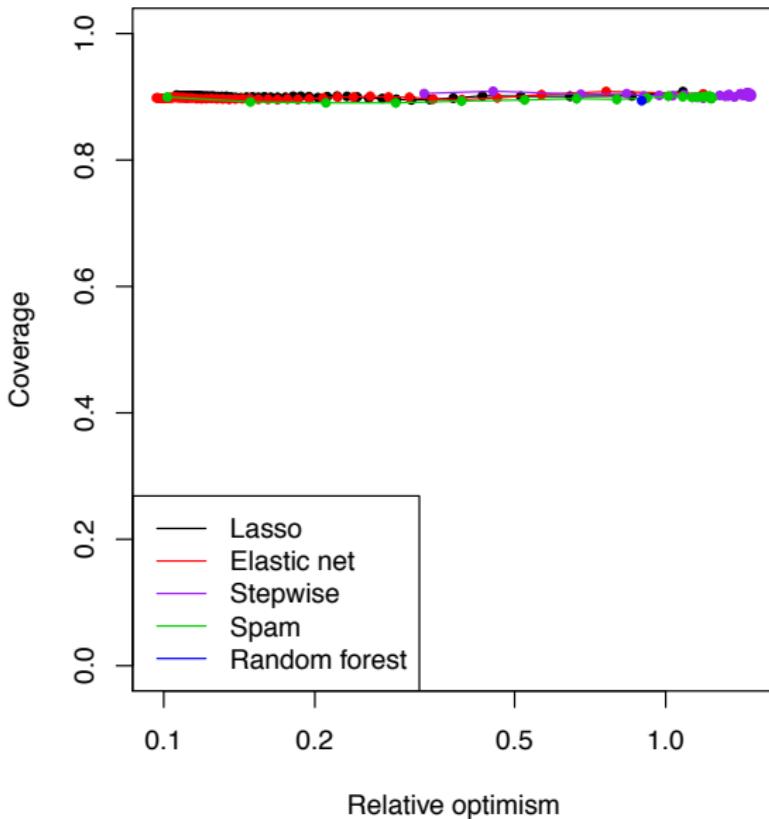


Invert p-values to get conformal interval

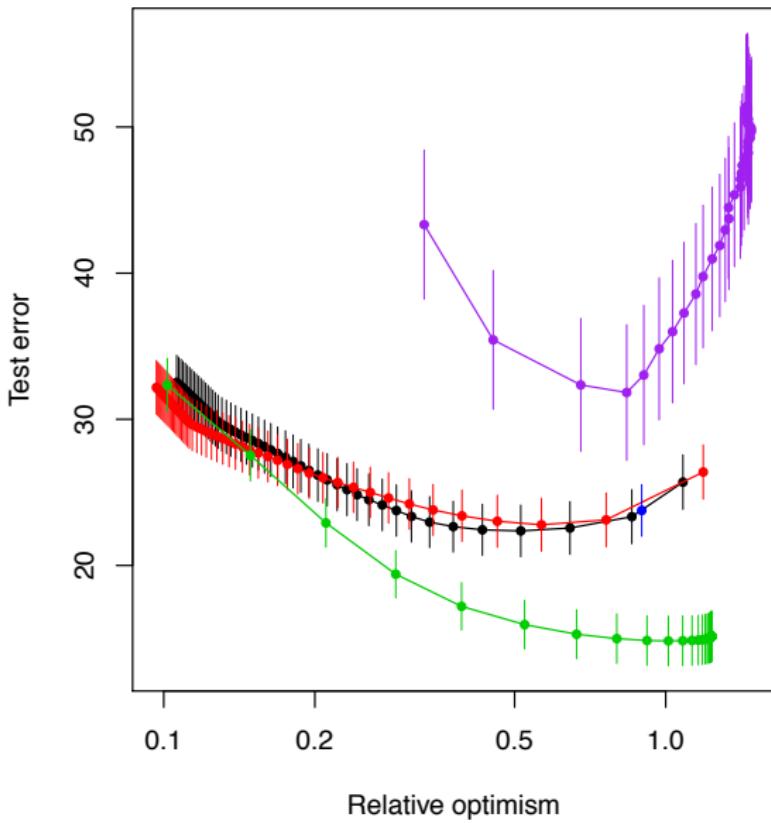
A high-dimensional example

- $n = 200, p = 2000$
- $\mathbb{E}(Y|X)$ is mixed additive B-splines on 5 variables.
- $X \sim N(0, I_{2000})$.
- $(\varepsilon | X = x) \sim t_2$

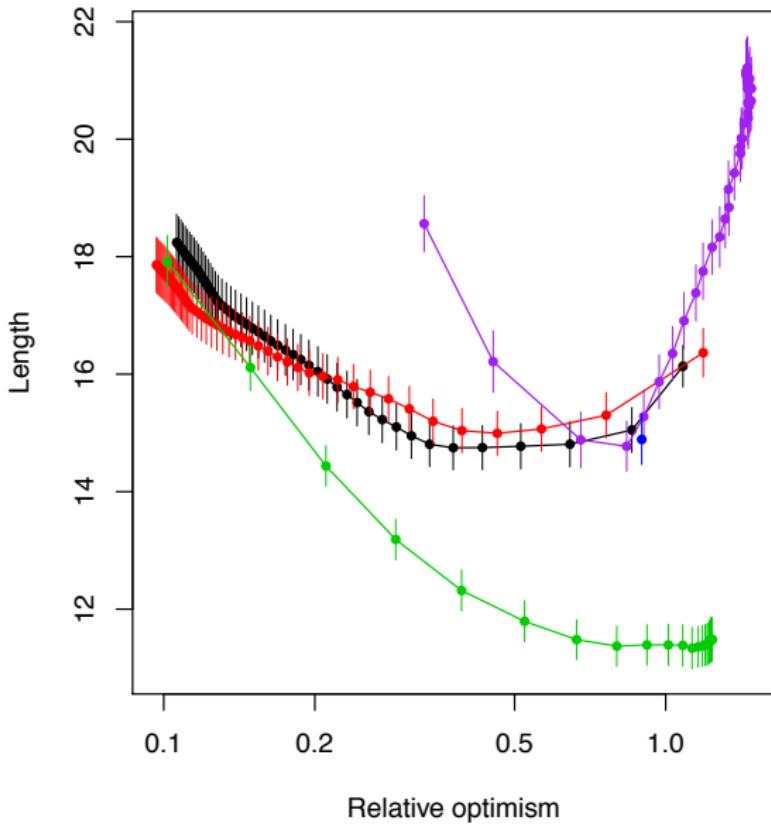
Coverage, Setting B



Test Error, Setting B



Length, Setting B



Remarks

- The coverage is always $1 - \alpha$ (anti-conservative) regardless of fitting method and value of tuning parameter.
- Good $\hat{\mu}$ gives short prediction intervals.
- The coverage guarantee is **marginal**, over the $(n + 1)$ -tuple $(X_i, Y_i)_{i=1}^{n+1}$.
- Can be combined with almost any point estimator $\hat{\mu}$.

A brief history of conformal prediction

- Developed, since 1996, by V. Vovk and collaborators as a generic tool for online sequential prediction.
- Lei, Robins, & Wasserman (2013): tolerance region.
- Lei & Wasserman (2014): nonparametric regression.
- Lei (2014): binary classification.
- Lei, Rinaldo, & Wasserman (2015): functional clustering.
- Sadinle, Lei, & Wasserman (2015): multi-class classification.
- Lei, G'Sell, Rinaldo, Tibshirani, Wasserman (2016): high dimensional regression, variable importance, further insights, R package “conformalInference”.
- Lei (2017): Fast computation for the Lasso.
- Chernozhukov et al (2018): time series.

Variable importance

- Assume $X \in \mathbb{R}^d$, where d can be large; $\hat{\mu}$ is a fitting algorithm.
- For $j = 1, \dots, d$, let $\hat{\mu}_{-j}$ be fitted without the j th coordinate of X .
- The j th variable is important if $|Y - \hat{\mu}_{-j}(X)|$ is larger than $|Y - \hat{\mu}(X)|$.
- Need to watch out for overfitting when using $|Y_i - \hat{\mu}_{-j}(X_i)| - |Y_i - \hat{\mu}(X_i)|$.

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- Need to watch out for overfitting when using $|Y_i - \hat{\mu}_{-j}(X_i)| - |Y_i - \hat{\mu}(X_i)|$.
- Idea: make a conformal prediction interval for

$$D_{ij} = |Y'_i - \hat{\mu}_{-j}(X_i)| - |Y'_i - \hat{\mu}(X_i)|$$

where Y'_i is a fresh draw from $(Y|X = X_i)$.

Variable importance

- Let $\tilde{C}(X_i)$ be a valid prediction interval for Y'_i and define

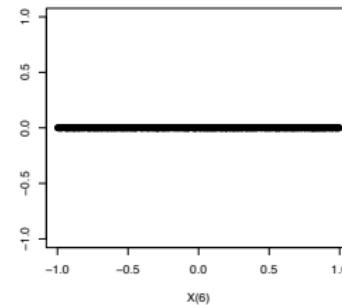
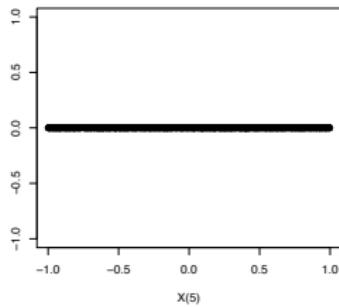
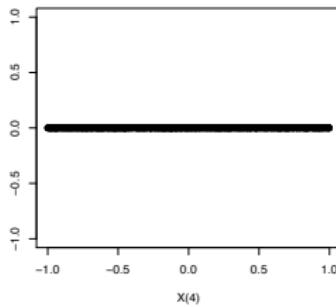
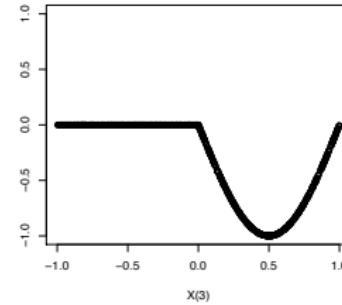
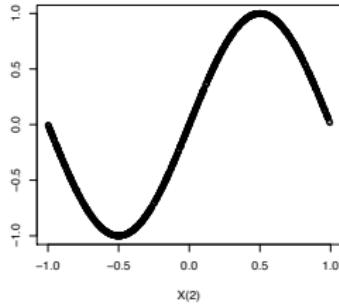
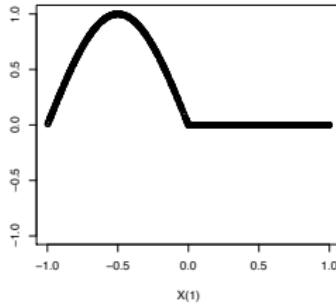
$$V_{ij} = \{|y - \hat{\mu}_{-j}(X_i)| - |y - \hat{\mu}(X_i)| : y \in \tilde{C}(X_i)\}$$

- Fact:** $Y'_i \in \tilde{C}(X_i) \Rightarrow D_{ij} \in V_{ij}$, and $\mathbb{P}(D_{ij} \in V_{ij}, \forall j) \geq 1 - \alpha$.
- Can construct conformal prediction band $\tilde{C}(X)$ such that

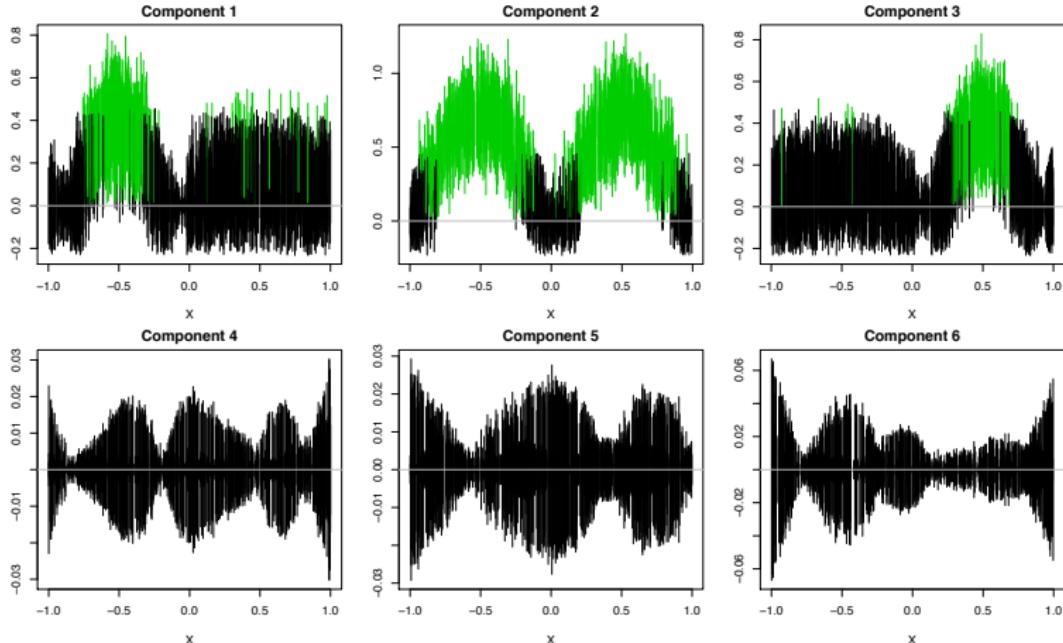
$$\mathbb{P}\left[n^{-1} \sum_{i=1}^n \mathbf{1}(D_{ij} \in V_{ij}, \forall j) \geq 1 - \alpha - \varepsilon\right] \geq 1 - 2e^{-cn\varepsilon^2}$$

Example: Additive Model

$$Y = \sum_{j=1}^6 f_j(X(j)) + N(0, 1)$$



How do V_{ij} 's look like?



The j th variable is likely to be important if some of $\{D_{ij} : 1 \leq i \leq n\}$ are above 0.

A higher dimensional example

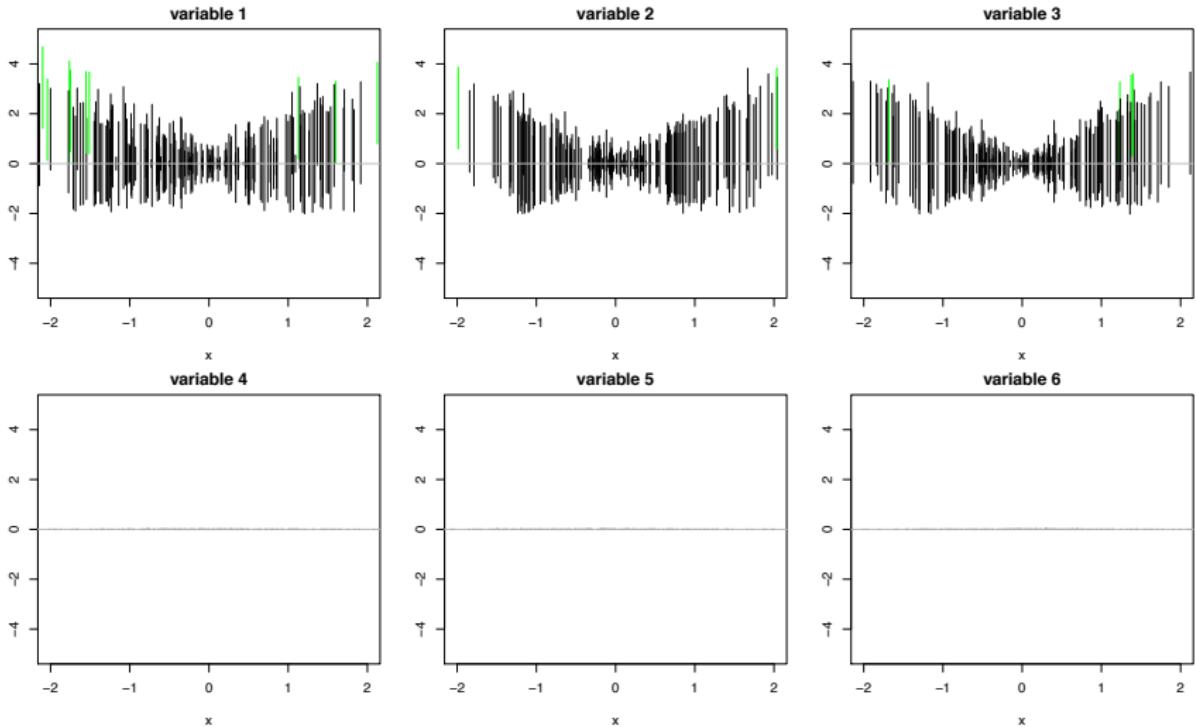
- $n = 200, p = 100$
- $Y = X^T \beta + \varepsilon$
- $\varepsilon \sim N(0, 1)$, independent of X
- $\beta = (2, 2, 2, 0, \dots, 0)^T$
- Design matrix

Case 1: $\mathbb{E}(XX^T) = I$ (all standard assumptions hold)

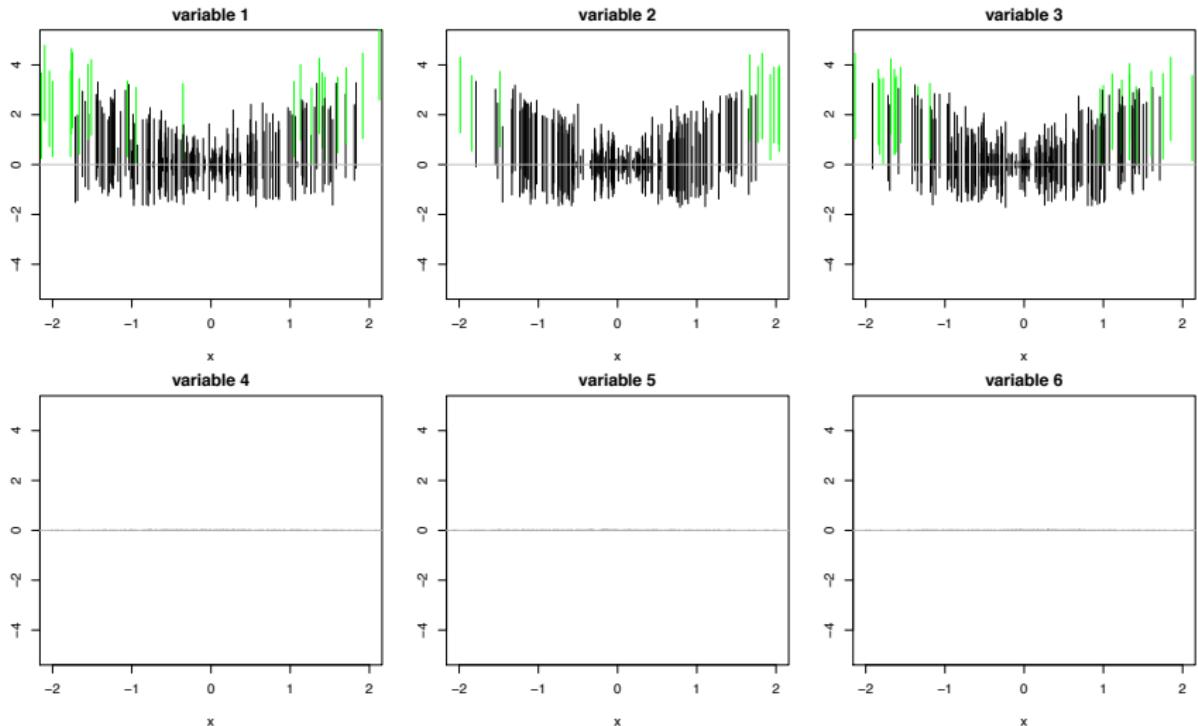
Case 2: $\text{corr}(X(j), X(j')) = 0.7$ if $j \neq j'$ (strong correlation)

- Fitting methods
 - (a) Lasso with $\lambda = 0.3$
 - (b) Forward Stepwise with 3 steps

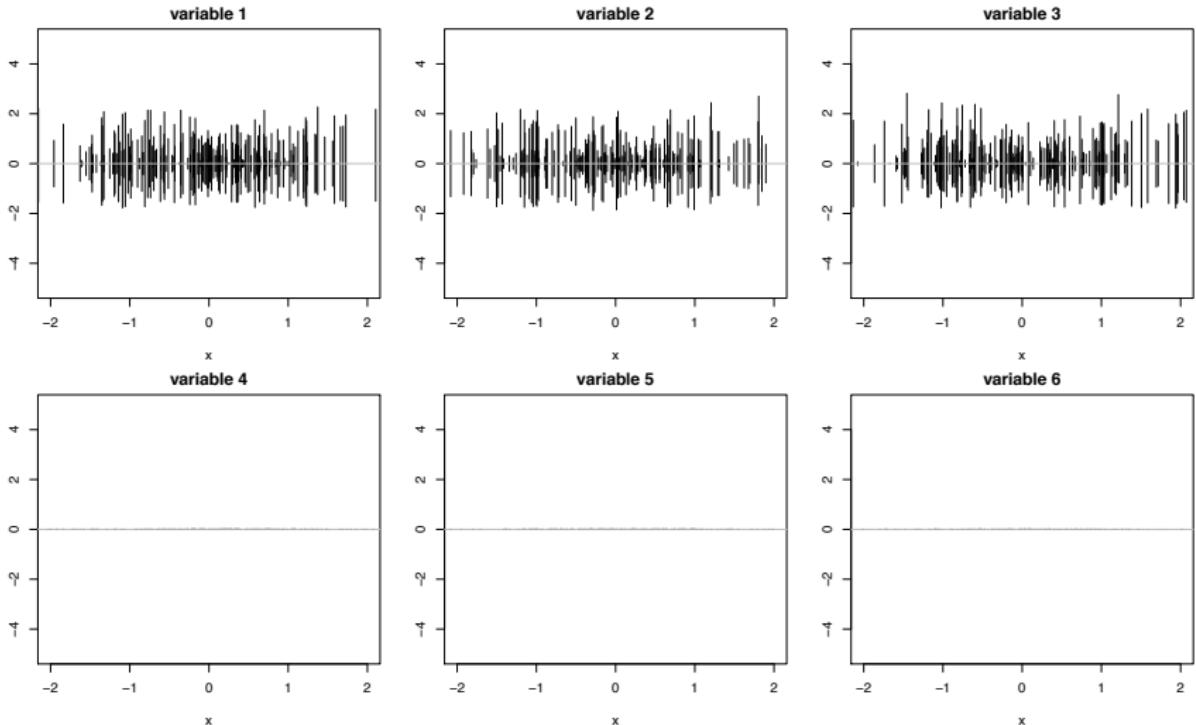
Uncorrelated case, Lasso



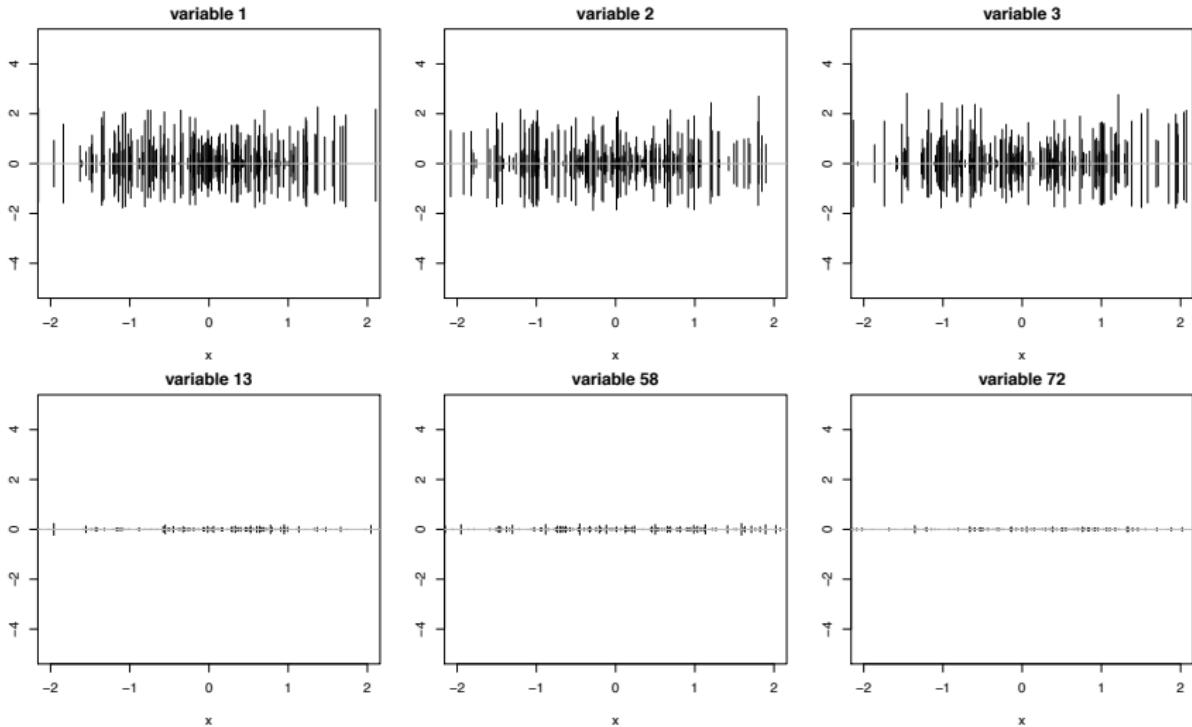
Uncorrelated case, Forward Stepwise



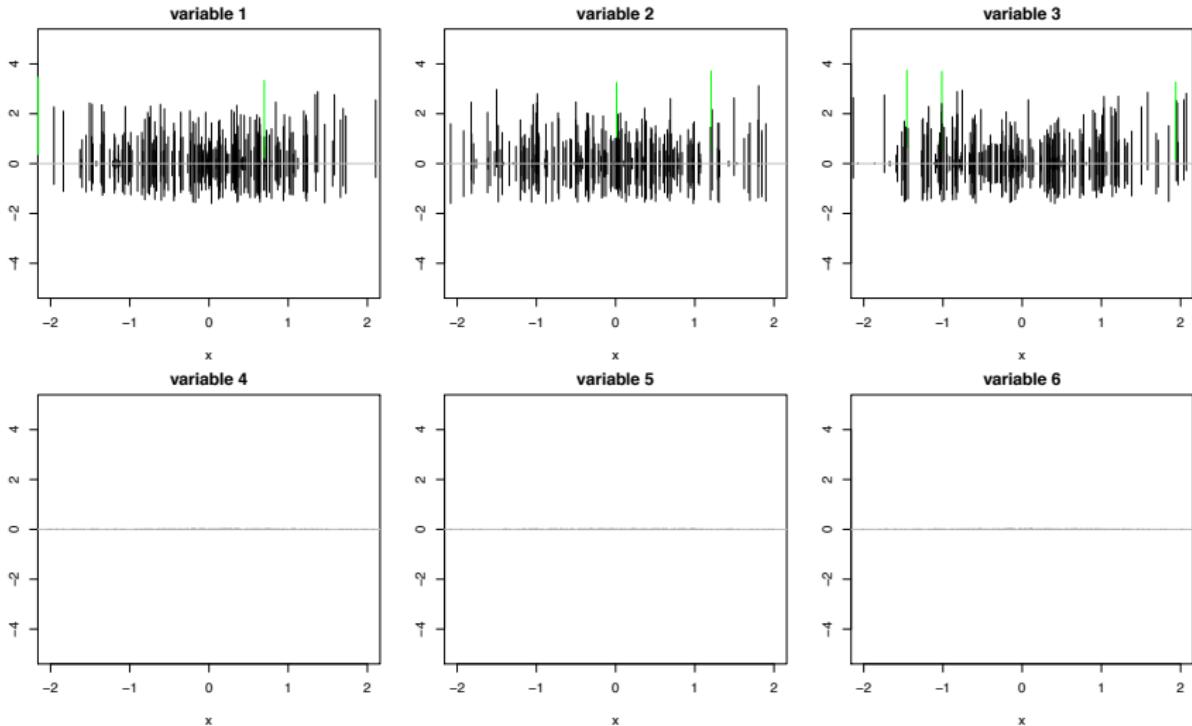
Correlated case, Lasso



Correlated case, Lasso



Correlated case, Forward Stepwise



Construction of $\tilde{C}(X)$

In-sample split conformal:

1. Split data into \mathcal{D}_1 and \mathcal{D}_2

2. For $k = 1, 2$

2.1 Let $\hat{\mu}_k$ be fitted using \mathcal{D}_k , $k = 1, 2$.

2.2 Let \hat{F}_k be the empirical CDF of $\{|Y_i - \hat{\mu}_{3-k}(X_i)| : (X_i, Y_i) \in \mathcal{D}_k\}$.

2.3 For each $X_i \in \mathcal{D}_k$,

$$\tilde{C}(X_i) = [\hat{\mu}_{3-k}(X_i) \pm \hat{F}_k^{-1}(1 - \alpha)]$$

Requires only two fits and two order statistics of cross-validated residuals.

Other topics

- Fast computation: avoid re-fitting $\hat{\mu}$ with extra data point (X_{n+1}, y) for all values of X_{n+1} and all y .
- Higher order correction: conformal prediction band with adaptive width.
- Theory: when $\hat{\mu}$ is a good estimator, then the conformal band is nearly optimal (requires standard assumptions, mainly relies on stability of $\hat{\mu}$).

From conformalization to cross-validation

- The construction of $\tilde{C}(X)$ reminds us of cross-validation, with just one difference:

CV looks at the empirical mean of the validated loss, while $\tilde{C}(X)$ looks at the empirical quantiles.
- Idea: there is more information in the validated loss than just the empirical mean.

Cross-validation with confidence

	Parameter est.	Model selection
Point est.	MLE, M-est., ...	Cross-validation
Interval est.	Confidence interval	CVC

In the regression setting

- Data: $D = \{(X_i, Y_i) : 1 \leq i \leq n\}$, i.i.d from joint distribution P on $\mathbb{R}^p \times \mathbb{R}^1$
- $Y = \mu(X) + \varepsilon$, with $E(\varepsilon | X) = 0$
- Loss function: $\ell(\cdot, \cdot) : \mathbb{R}^2 \mapsto \mathbb{R}$
- Goal: find $\hat{\mu} \approx \mu$ so that

$$Q(\hat{\mu}) \equiv \mathbb{E}[\ell(\hat{\mu}(X), Y) | \hat{\mu}]$$

is small.

Model selection

- Candidate set: $\mathcal{M} = \{1, \dots, M\}$. Each $m \in \mathcal{M}$ corresponds to a candidate model.
- Given m and data D , there is an estimate $\hat{\mu}(D, m)$ of μ .
- Model selection: find the best m such that it minimizes $Q(\hat{\mu})$ over all $m \in \mathcal{M}$ with high probability.

Cross-validation

- Sample split: Let I_{tr} and I_{te} be a partition of $\{1, \dots, n\}$.
- Fitting: $\hat{\mu}_m = \hat{\mu}(D_{\text{tr}}, m)$, where $D_{\text{tr}} = \{(X_i, Y_i) : i \in I_{\text{tr}}\}$.
- Validation: $\hat{Q}(\hat{\mu}_m) = n_{\text{te}}^{-1} \sum_{i \in I_{\text{te}}} \ell(\hat{\mu}_m(X_i), Y_i)$.
- CV model selection: $\hat{m}_{\text{cv}} = \arg \min_{m \in \mathcal{M}} \hat{Q}(\hat{\mu}_m)$.
- V-fold cross-validation:
 1. For $V \geq 2$, split the data into V folds.
 2. Rotate over each fold as I_{tr} to obtain $\hat{Q}^{(v)}(\hat{\mu}_m^{(v)})$
 3. $\hat{m} = \arg \min V^{-1} \sum_{v=1}^V \hat{Q}^{(v)}(\hat{\mu}_m^{(v)})$
 4. Popular choices of V : 10 and 5.
 5. $V = n$: leave-one-out cross-validation

A simple negative example

- Model: $Y = \mu + \varepsilon$, where $\varepsilon \sim N(0, 1)$.
- $\mathcal{M} = \{1, 2\}$. $m = 1$: $\mu = 0$; $m = 2$: $\mu \in \mathbb{R}$.
- Truth: $\mu = 0$
- Consider a single split: $\hat{\mu}_1 \equiv 0$, $\hat{\mu}_2 = \bar{\varepsilon}_{\text{tr}}$.
- $\hat{m}_{\text{cv}} = 1 \Leftrightarrow 0 < \hat{Q}(\hat{\mu}_2) - \hat{Q}(\hat{\mu}_1) = \bar{\varepsilon}_{\text{tr}}^2 - 2\bar{\varepsilon}_{\text{tr}}\bar{\varepsilon}_{\text{te}}$.
- If $n_{\text{tr}}/n_{\text{te}} \asymp 1$, then $\sqrt{n}\bar{\varepsilon}_{\text{tr}}$ and $\sqrt{n}\bar{\varepsilon}_{\text{te}}$ are independent normal random variables with constant variances. So $\mathbb{P}(\hat{m}_{\text{cv}} = 1)$ is bounded away from 1.

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- (Shao 93, Zhang 93, Yang 07) \hat{m}_{cv} is inconsistent unless $n_{\text{tr}} = o(n)$.
- V -fold does not help!

Cross-Validation with Confidence

- Now suppose we have a set of candidate models $\mathcal{M} = \{1, \dots, M\}$.
- Split the data into D_{tr} and D_{te} , and use D_{tr} to obtain $\hat{\mu}_m$ for each m .
- Recall that the model quality is $Q(\hat{\mu}) = \mathbb{E} [\ell(\hat{\mu}(X), Y) \mid \hat{\mu}]$.
- For each m , test hypothesis (conditioning on $\hat{\mu}_1, \dots, \hat{\mu}_M$)

$$H_{0,m} : \min_{j \neq m} Q(\hat{\mu}_j) \geq Q(\hat{\mu}_m).$$

- Let \hat{p}_m be a valid p -value.
- $\mathcal{A}_{\text{cvc}} = \{m : \hat{p}_m > \alpha\}$ is our confidence set for the best fitted model: $\mathbb{P}(m^* \in \mathcal{A}_{\text{cvc}}) \geq 1 - \alpha$, where $m^* = \arg \min_m Q(\hat{\mu}_m)$.

Calculating \hat{p}_m

- Recall $H_{0,m} : \min_{j \neq m} Q(\hat{\mu}_j) \geq Q(\hat{\mu}_m)$.
- Consider $n_{\text{te}} \times (M - 1)$ matrix (I_{te} is the index set of D_{te})

$$\left[\xi_{m,j}^{(i)} \right]_{i \in I_{\text{te}}, j \neq m}, \text{ where } \xi_{m,j}^{(i)} = \ell(\hat{\mu}_m(X_i), Y_i) - \ell(\hat{\mu}_j(X_i), Y_i)$$

- Multivariate mean testing. $H_{0,m} : \mathbb{E}(\xi_{m,j}) \leq 0, \forall j \neq m$.

Calculating \hat{p}_m

- $H_{0,m} : \mathbb{E}(\xi_{m,j}) \leq 0, \forall j \neq m.$
- Let $\hat{\mu}_{m,j}$ and $\hat{\sigma}_{m,j}$ be the sample mean and standard deviation of $(\xi_{m,j}^{(i)} : i \in I_{\text{te}})$.
- Naturally, one would reject $H_{0,m}$ for large values of

$$\max_{j \neq m} \frac{\hat{\mu}_{m,j}}{\hat{\sigma}_{m,j}}.$$

- Approximate the null distribution using high dimensional Gaussian comparison [Chernozhukov et al '12].

Studentized Gaussian Multiplier Bootstrap

1. $T_m = \max_{j \neq m} \sqrt{n_{\text{te}}} \frac{\hat{\mu}_{m,j}}{\hat{\sigma}_{m,j}}$
2. Let B be the bootstrap sample size. For $b = 1, \dots, B$,
 - 2.1 Generate iid standard Gaussian $\zeta_i, i \in I_{\text{te}}$.
 - 2.2 $T_b^* = \max_{j \neq m} \frac{1}{\sqrt{n_{\text{te}}}} \sum_{i \in I_{\text{te}}} \frac{\xi_{m,j}^{(i)} - \hat{\mu}_{m,j}}{\hat{\sigma}_{m,j}} \zeta_i$
3. $\hat{p}_m = B^{-1} \sum_{b=1}^B \mathbf{1}(T_b^* > T_m)$. correlation.

Properties of CVC

- $\mathcal{A}_{\text{cvc}} = \{m : \hat{p}_m > \alpha\}.$
- Let $\hat{m}_{\text{cv}} = \arg \min_m \hat{Q}(\hat{\mu}_m).$

Proposition

If $\alpha < 0.5$, then $\mathbb{P}(\hat{m}_{\text{cv}} \in \mathcal{A}_{\text{cvc}}) \rightarrow 1$ as $B \rightarrow \infty$.

- Can view \hat{m}_{cv} as the “center” of the confidence set.

Coverage of \mathcal{A}_{cvc}

- Recall $\xi_{m,j} = \ell(\hat{\mu}_m(X), Y) - \ell(\hat{\mu}_j(X), Y)$.
- Let $\mu_{m,j} = \mathbb{E}[\xi_{m,j} | \hat{\mu}_m, \hat{\mu}_j]$, $\sigma_{m,j}^2 = \text{Var}[\xi_{m,j} | \hat{\mu}_m, \hat{\mu}_j]$.

Theorem

Assume that $(\xi_{m,j} - \mu_{m,j})/(A_n \sigma_{m,j})$ has sub-exponential tail for all $m \neq j$ and some $A_n \geq 1$ such that for some $c > 0$

$$A_n^6 \log^7(M \vee n) = O(n^{1-c}).$$

1. If $\max_{j \neq m} \left(\frac{\mu_{m,j}}{\sigma_{m,j}} \right)_+ = o\left(\sqrt{\frac{1}{n \log(M \vee n)}}\right)$, then $\mathbb{P}(m \in \mathcal{A}_{\text{cvc}}) \geq 1 - \alpha + o(1)$.
2. If $\max_{j \neq m} \left(\frac{\mu_{m,j}}{\sigma_{m,j}} \right)_+ \geq CA_n \sqrt{\frac{\log(M \vee n)}{n}}$ for some constant C , and $\alpha \geq n^{-1}$, then $\mathbb{P}(m \in \mathcal{A}_{\text{cvc}}) = o(1)$.

Proof of coverage

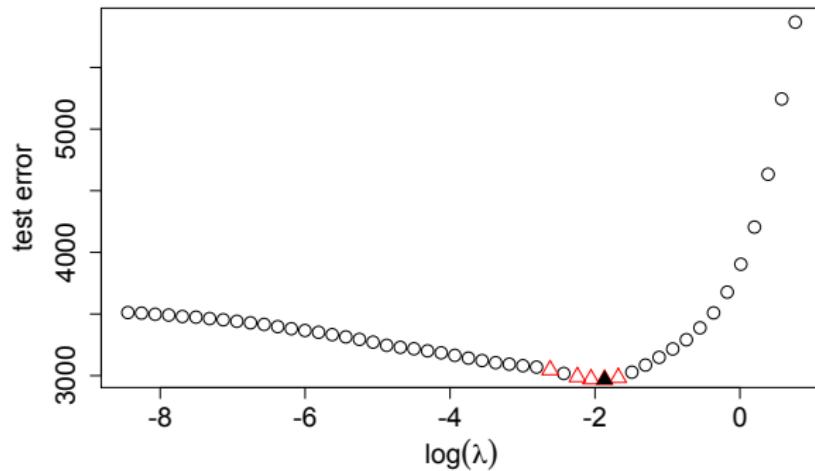
- Let $Z(\Sigma) = \max N(0, \Sigma)$, and $z(1 - \alpha, \Sigma)$ its $1 - \alpha$ quantile.
- Let $\hat{\Gamma}$ and Γ be sample and population correlation matrices of $(\xi_{m,j}^{(i)})_{i \in I_{\text{te}}, j \neq m}$. When $B \rightarrow \infty$,

$$\mathbb{P}(\hat{p}_m \leq \alpha) = \mathbb{P}\left[\max_j \sqrt{n_{\text{te}}} \frac{\hat{\mu}_{m,j}}{\hat{\sigma}_{m,j}} \geq z(1 - \alpha, \hat{\Gamma})\right]$$

- Tools (2, 3 are due to Chernozhukov et al.)
 1. Concentration: $\sqrt{n_{\text{te}}} \frac{\hat{\mu}_{m,j}}{\hat{\sigma}_{m,j}} \leq \sqrt{n_{\text{te}}} \frac{\hat{\mu}_{m,j} - \mu_{m,j}}{\sigma_{m,j}} + o(1/\sqrt{\log M})$
 2. Gaussian comparison: $\max_j \sqrt{n_{\text{te}}} \frac{\hat{\mu}_{m,j} - \mu_{m,j}}{\sigma_{m,j}} \stackrel{d}{\approx} Z(\Gamma) \stackrel{d}{\approx} Z(\hat{\Gamma})$
 3. Anti-concentration: $Z(\hat{\Gamma})$ and $Z(\Gamma)$ have densities $\lesssim \sqrt{\log M}$

Example: the diabetes data (Efron et al 04)

- $n = 442$, with 10 covariates: age, sex, bmi, blood pressure, etc.
- Response is diabetes progression after one year.
- Including all quadratic terms, $p = 64$.
- 5-fold CVC with $\alpha = 0.05$, using Lasso with 50 values of λ .



Triangle: models in \mathcal{A}_{cv} , solid triangle: \hat{m}_{cv} .

The most parsimonious model in \mathcal{A}_{cvc}

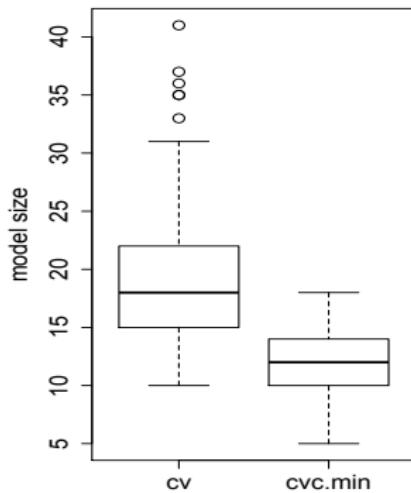
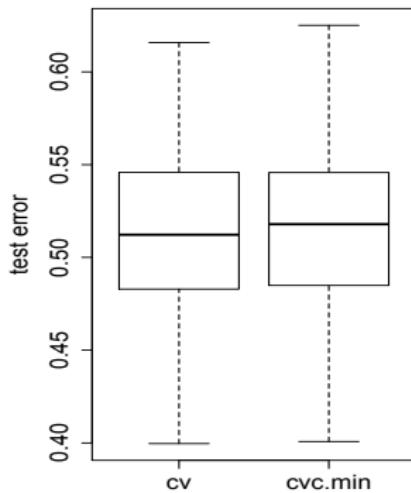
- Let J_m be the subset of variables selected using model m

$$\hat{m}_{\text{cvc},\min} = \arg \min_{m \in \mathcal{A}_{\text{cvc}}} |J_m|.$$

- $\hat{m}_{\text{cvc},\min}$ is the simplest model that gives a similar predictive risk as \hat{m}_{cv} .
- Consistent in low-dimensional linear models with conventional V-fold implement.

The diabetes data revisited

- Split $n = 442$ into 300 (estimation) and 142 (risk approximation).
- 5-fold CVC applied on the 300 sample points, with a final re-fit.
- The final estimate is evaluated using the 142 hold-out sample.
- Repeat 100 times, using Lasso with 50 values of λ .



Summary

- Conformal prediction uses symmetry and out-of-sample fitting to add protection against model misspecification.
- CVC uses hypothesis tests to produce confidence sets for model selection
- Both methods are applicable to many learning algorithms, even black-box type algorithms.

Thanks!

Questions?

“Distribution Free Predictive Inference for Regression”

arXiv:1604.04173 with Wasserman, Tibshirani, G’Sell, Rinaldo

“Cross-Validation with Confidence”, arxiv.org/1703.07904

<http://www.stat.cmu.edu/~jinglei/talk.shtml>