Frontier Topics in Empirical Economics: Week 3 Machine Learning and Model Selection

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- We now have many tools in our box beyond linear regression
 - Series regression, partial linear regression

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- Which method should we choose?

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 Data driven method (Machine learning)

 Price causal structure (DAG)
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- We select models only using data
- We do not put our economic knowledge into the process
- Let's first introduce a major statistical concept: Bias-variance tradeoff

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A traditional linear mode

$$y = x\beta + \epsilon \tag{1}$$

A model with quadratic term

$$y = x\beta + x^2\alpha + \epsilon \tag{2}$$

A non-parametric mode

$$y = g(x) + \epsilon \tag{3}$$

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A model with quadratic term

$$-x^{2}+x^{2}$$

A non-parametric model

$$v = g(x) + \epsilon \tag{3}$$

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Model A

$$y = x_1'\beta + \epsilon \tag{4}$$

Model B

$$v = x_1' \beta_1 + x_2' \beta_2 + \epsilon \tag{5}$$

- Why not always the second one?
- Always better to have a more complicated model?

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$$Y = f(X) + \epsilon$$

- $\hat{f}(x)$ is a model trained by some data
- **IIII** It will be changed when sample is changed: $f^1(x)$, $f^2(x)$...
- \blacksquare Expectation E[f(x)] is taken over different sampless
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= irreducible error + Bias² + Variance

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- Overfit current data ⇒ Poor out-of-sample prediction

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Consider a data generating process

$$Y = 1 + 1.5X + \epsilon$$
$$\epsilon \sim N(0, 100)$$

- Simulate 30 observations from this process
- Let's start to fit it with different polynomials
- Green line is the true DGP
- Red line is the fitting function

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Figure: First Order (Linear) Fitting

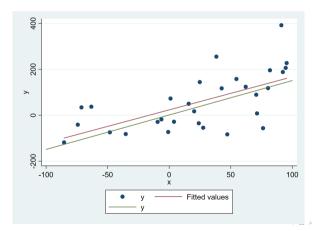


Figure: Second Order (Quadratic) Fitting

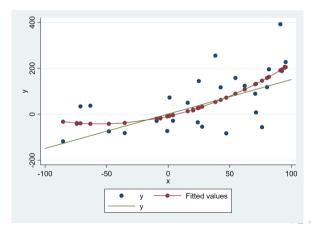


Figure: Third Order (Cubic) Fitting

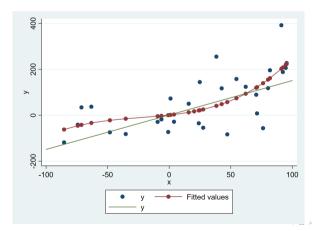


Figure: Fourth Order Fitting

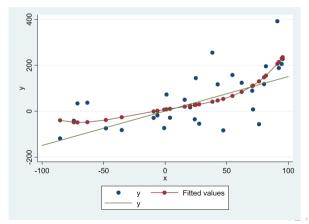


Figure: Fifth Order Fitting

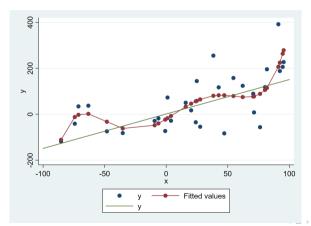


Figure: Sixth Order Fitting

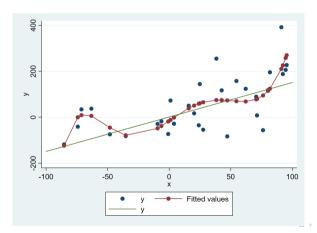
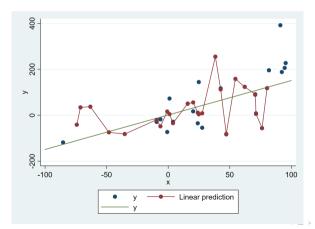


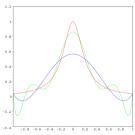
Figure: Twentieth Order Fitting





High order polynomials: Picking up noises, not signals!!! Bad out-of-sample prediction!!!

- We have actually learned two kinds of overfitting
- Runge phenomenon and Gibbs phenomenon



- Adjusted R-squared: the proportion of explained variations in y Still remember why we need to adjust for the number of regressors?
- AIC: Akaike Information Criterion AIC = 2k + nln(RSS/n), k is the number of regressors
- BIC: Bayesian Information Criterion
 This is motivated by the Bayesian approach to model selection

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Another important measure is Cross-Validation (CV)

- The basic idea is to separate all samples into training sample and validation sample
- Training sample is used to train (estimate) the model
- Validation sample is then used to check the "out-of-sample" prediction
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- First, we separate all samples into K parts
- Each time, we choose K-1 parts to train (estimate) the model
- We then use the remaining one part k to calculate the mean squared predicted error MSE_k
- we rotate the samples K times so that each part is used as the validation sample once, and have K pieces of MSE_k
- We take the average of them to have: $CV = \frac{1}{k} \sum_{k=1}^{K} MSE_{k}$
- This is called "K-fold Cross-Validation"



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Here is the process of CV



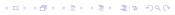
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- You have some data that is not used in the estimation and use it to check your estimation validity
- It helps you to determine which model fits better to the data, in terms of out-of-sample prediction
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- Main target: How complicated the model should be should be
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Machine Learning and Model Selection: Penalized Regressions

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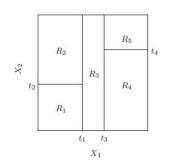
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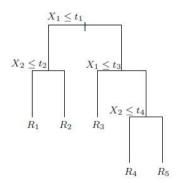
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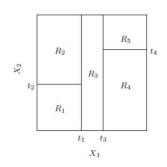
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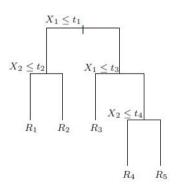
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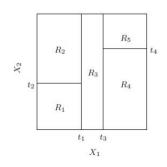


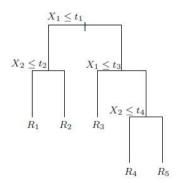
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- Two choices: continue partitioning or stop + where to partition
- Greedy algorithm
- For each region R_m (leaf), we define:

Size (# of obs):
$$N_m = \{x_i \in R_m\}$$

Fitted value (mean as fit): $\hat{c}_m = \frac{1}{N_m} \sum_{x \in R_m} y_i$
SSE (error in leaf): $Q_m(T) = \frac{1}{N_m} \sum_{x \in R} (y_i - \hat{c}_m)^2$

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- For j th predictor, cut position s
- Define half plane $R_1(j,s) = \{X | X_j \le s\}, R_2(j,s) = \{X | X_j > s\}$
- How to find (j,s) in each branch? Minimize SSE (Easy)

$$\min_{j,s} [\min_{c_1} \sum_{x_i \in R_1(j,s)} (y_i - c_1)^2 + \min_{c_2} \sum_{x_i \in R_2(j,s)} (y_i - c_2)^2]$$

 \blacksquare Here c_1 and c_2 are conditional means (in leaf 1 and 2)

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lpha as the tuning parameter; $|\mathcal{T}|$ as number of terminal nodes

- Total SSE (bias) + Size penalty
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- Using sub-sampling or bagging to reduce variance of a single tree
- Draw a lot of different samples (1,2,...B) with sub-sampling (n < N) (Jackknife) or bagging (n = N) (Bootstrap)
- \blacksquare De-correlation: In each split, randomly select m variables to do the partition

$$\hat{f}^{B}(x) = \frac{1}{B} \sum_{b=1}^{B} T_{b}(x)$$
$$V(\hat{f}) \approx \rho \sigma^{2} + \frac{1 - \rho}{B} \sigma^{1}$$

- Random Forests = Tree Method + Sampling average (Many De-correlated Trees)
- To reduce V(f): $B \uparrow$ (more sampling), $\rho \downarrow$ (smaller correlation)

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Algorithm 15.1 Random Forest for Regression or Classification.

- 1. For b = 1 to B:
 - (a) Draw a bootstrap sample \mathbf{Z}^* of size N from the training data.
 - (b) Grow a random-forest tree T_b to the bootstrapped data, by recursively repeating the following steps for each terminal node of the tree, until the minimum node size n_{min} is reached.
 - i. Select m variables at random from the p variables.
 - ii. Pick the best variable/split-point among the m.
 - iii. Split the node into two daughter nodes.
- 2. Output the ensemble of trees $\{T_b\}_1^B$.

To make a prediction at a new point x:

Regression:
$$\hat{f}_{rf}^B(x) = \frac{1}{B} \sum_{b=1}^B T_b(x)$$
.

Classification: Let $\hat{C}_b(x)$ be the class prediction of the bth random-forest tree. Then $\hat{C}_{rf}^B(x) = majority\ vote\ \{\hat{C}_b(x)\}_1^B$.

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 Making predictions using weighted averages of "nearby" observation
- Difference: Weighting scheme
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Machine Learning and Model Selection: Application of Causal Forests

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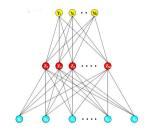


FIGURE 11.2. Schematic of a single hidden layer, feed-forward neural network

X - Input; Z- Hidden layer/unit; Y - Output

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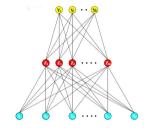


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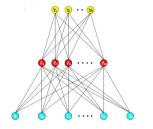


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- Because it was first developed as models for the human brain
- Each unit represents a neuron
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