Quant III

Lab 12: Tree-based Method

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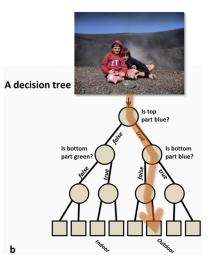
December 11, 2020

Announcement

- Final Exam:
 - I will distribute it Friday, December 18th, 9:45 am (EST)
 - Due: Sunday, December 20th, 11:59 pm (EST)
- Replication: due Friday, December 18th, 11:59 pm (EST)
- Final homework:
 - Due: Friday, December 18th, 11:59 pm (EST)
 - You should submit a pdf and a r object,
 - which includes a object for user: The input being a design matrix, and output being predictions.
- I add an extra office hour for next week, check here

A Binary Decision Tree

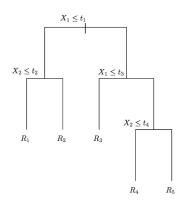
• binary tree: each node has either 2 children or 0 children

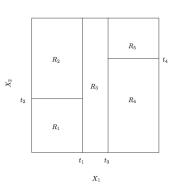


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Binary Decision Tree on R^2

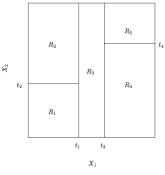
• Consider a binary tree on $\{(X_1, X_2) \mid X_1, X_2 \in R\}$

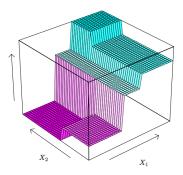




Binary Regression Tree on R^{2}

• Consider a binary tree on $\{(X_1, X_2) \mid X_1, X_2 \in R\}$





Recursive binary splitting,

- Split the predictor space on one dimension (on predictor) into two sets.
- Split each of the resulting sets into two sets again.
- Repeat until you have only a small number of observations left in each of the terminal sets.

Recursive binary splitting

• For a given j and s, split the space of each predictor x_j , j = 1, ..., p, into two regions:

$$R_{-}(j,s) = \{ \mathbf{x} | x_j < s \}$$

 $R_{+}(j,s) = \{ \mathbf{x} | x_j \ge s \}$

 $R_{-}(j,s)$ is the region of predictor space where $x_i < s$.

• The splitting is done by finding j and s that minimizes

$$\sum_{i:x_i \in R_-(j,s)} (y_i - \hat{y}_{R_-})^2 + \sum_{i:x_i \in R_+(j,s)} (y_i - \hat{y}_{R_+})^2$$

where $\hat{y_R} = E(y|x_i \in R)$ - Each resulting region is then split further until each region contains no more than N^* observations (where N^* is a small number).

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Classification tree

- Suppose that the outcome variable is nominal $y_i \in \{c_1, ..., c_K\}$ so that y_i can be one of K categories.
- Prediction for observation $i: \hat{p_{ik}}$
- Prediction in that region R_i : $\hat{p_{jk}}$
- The idea is exactly the same except that when calculating \hat{y}_{iR_m} we use the most frequent outcome category for observations in the set R_m .
- The only difference is that RSS is not used to find splits or to calculate the optimal complexity parameter α .

Impurity Measures

Classification error:

$$1 - \max_k(\hat{p}_{jk})$$

where \hat{p}_{jk} is the proportion of observations in the region R_j that are from the class k. If a region R_j contains observations from many classes, then it does not classify them well. - Gini index

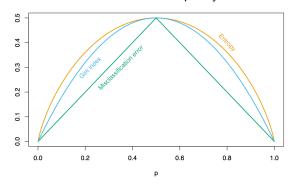
$$\sum_{k=1}^K \hat{p}_{jk} (1 - \hat{p}_{jk})$$

Cross-entropy:

$$-\sum_{k=1}^K \hat{p}_{jk} \ln \hat{p}_{jk}$$

Two-Class Node Impurity Measures

- Consider binary classification
- Let p be the relative frequency of class 1.
- Here are three node impurity measures as a function of p



Bagging

- Combine predictions from multiple regression trees to improve their individual predictive capacity:
 - **1** Take a bootstrap sample b = 1, ..., B from your data (with replacement).
 - ② Estimate a deep regression tree for sample b (without prunning).
 - **3** Calculate prediction $\hat{g}^b(x)$ given that tree.
 - Calculate the ensemble prediction

$$\hat{g}(\mathbf{x}) = \sum_{b=1}^{B} \hat{g}^b(\mathbf{x}) / B$$

 Averaging over a large number of high-variance low-bias trees results in lower-variance bag of trees.

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Random Forest

- Two disadvantages of bagging:
 - Computationally complex
 - Each tree are too similar
- Random forest solve these issues:
 - **1** Take a bootstrap sample b = 1, ..., B from your data (with replacement).
 - Build regression tree by randomly selecting at each step $\mathbf{m} < \mathbf{p}$ predictors (usually $m \approx \sqrt{p}$).
 - 3 Calculate prediction $\hat{g}^b(x)$ from that tree.
 - Calculate the ensemble prediction

$$\hat{g}(\mathbf{x}) = \sum_{b=1}^{B} \hat{g}^b(\mathbf{x}) / B$$

• Mathematically, RF should recover true g(x)

Different Forest

- You may see different forests, e.g.
 - Generalized Random Forest (Athey, et al.)
- Most the difference lies in the spliting criterion.

Optional: Boosting

- Ensemble methods combine multiple models
- Parallel ensembles:
 - · each model is built independently
 - e.g. bagging and random forests
 - Main Idea: Combine many (high complexity, low bias) models to reduce variance
- Sequential ensembles:
 - Models are generated sequentially
 - Try to add new models that do well where previous models lack

Intuition

- Suppose we want to solve a classification problem (spam or not)
- A weak learner is a classifier that does slightly better than random.
 - Weak learners are like "rules of thumb":
 - If an email has "Viagra" in it, more likely than not it's spam.
 - Email from a friend is probably not spam.
 - A linear decision boundary.
- Can we combine a set of weak classifiers to form single classifier that makes accurate predictions
- Yes! Boosting solves this problem. [Rob Schapire (1990).]
- In general, you can use different classifier. (e.g., Grimmer et al, 2017)

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Basic Idea

• Linear combination of basis functions:

$$f(x) = \sum_{m=1}^{M} \nu_m g_m(x)$$

• We'll consider learning by empirical risk minimization:

$$\hat{f} = \underset{f \in \mathbb{F}}{\operatorname{argmin}} \frac{1}{n} \sum_{i=1}^{n} \ell(y_i, f(x_i)),$$

for some **loss function** $\ell(y, \hat{y})$.

• Write ERM objective function as

$$J(v_1, \ldots, v_M, h_1, \ldots, h_M) = \frac{1}{n} \sum_{i=1}^n \ell\left(y_i, \sum_{m=1}^M v_m h_m(x)\right).$$

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• How to optimize *J*? i.e. how to learn?

Gradient-Based Methods

• **Suppose** our base hypothesis space is parameterized by $\Theta = R^b$:

$$J(v_1,\ldots,v_M,\theta_1,\ldots,\theta_M) = \frac{1}{n}\sum_{i=1}^n \ell\left(y_i,\sum_{m=1}^M v_m h(x;\theta_m)\right).$$

- Can we can differentiate J w.r.t. v_m 's and θ_m 's? Optimize with SGD?
- For some hypothesis spaces and typical loss functions, yes!
- Neural networks fall into this category! (h_1, \ldots, h_M) are neurons of last hidden layer.)
- We have gradient boost machine, Adaboost, etc.
- For trees, we have gradient tree boosting