Ch 8.1: Decision Trees

Lecture 16 - CMSE 381

Michigan State University

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Dept of Computational Mathematics, Science & Engineering

Wed, March 20, 2024

Announcements

Last time:

Cubic Splines

This lecture:

• 8.1 Decision Trees

Announcements:

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• Mid-term exam 2: next week

Section 1

Decision Trees

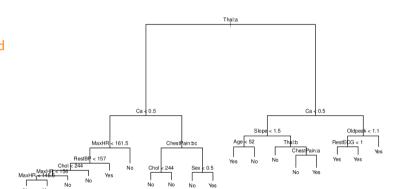
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Big idea

 Previously, we've always had some form of function for prediction:

$$f(X_1,\cdots,X_p)=\hat{Y}$$

 Now, we make a sequence of decisions to make a prediction, either regression or classification version



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Subset of Hitters data

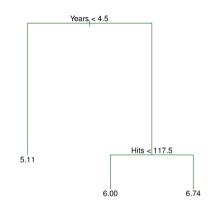
	Hits	Years	Salary	LogSalary
1	81	14	475.0	6.163315
2	130	3	480.0	6.173786
3	141	11	500.0	6.214608
4	87	2	91.5	4.516339
5	169	11	750.0	6.620073
317	127	5	700.0	6.551080
318	136	12	875.0	6.774224
319	126	6	385.0	5.953243
320	144	8	960.0	6.866933
321	170	11	1000.0	6.907755

- Remove observations missing salary values
- log transform salary for something closer to bell shape
- Goal: predict log salary (can reverse by returning exp(x) if model returns x)

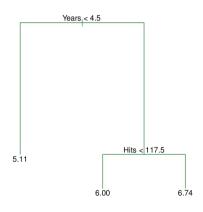
A simpler decision tree example

	Hits	Years	LogSalary
1	81	14	6.163315
2	130	3	6.173786
3	141	11	6.214608
4	87	2	4.516339
5	169	11	6.620073
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- Top split assigns observations with *Years* < 4.5 to left branch
- Return mean response for players with that.
- Predictions:
 - ▶ mean log salary is 5.107, so returns exp(5.107) = \$165.174 thousand dollars
 - ► 5.999 ⇒ \$402,834
 - ► 6.740 ⇒ \$845, 346



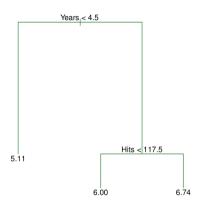
Interpretation of example



- Years most important factor for determining salary
- Players with less experience earn lower salaries than more experienced
- For the less experienced players, number of hits plays little role in salary
- For more experienced players, number of hits affects it
- Likely an oversimplification of real relationship, but easier to interpret and has nice graphical representation

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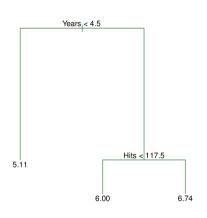
Regions defined by the tree

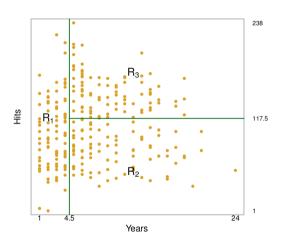


- Point out tree is "upside down"
- Leaves of the tree:
 - $R_1 = \{X \mid Years < 4.5\}$
 - ► R₂ = {X | Years >= 4.5, Hits < 117.5}
 - $R_3 = \{X \mid Years >= 4.5, Hits >= 117.5\}$

- Other splits are called Internal Nodes
- Segments that connect nodes called Branches or Edges

Viewing Regions Defined by Tree

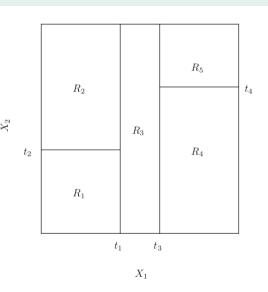




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Viewing Regions Defined by Tree

- We divide the predictor space that is, the set of possible values for X₁, X₂, ··· , X_p — into J distinct and non-overlapping regions, R₁, R₂, ··· , R_J.
- ② For every observation that falls into the region R_j , we make the same prediction = the mean of the response values for the training observations in R_j .



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How to build the tree?

Step 1, grow the tree iteratively: in each step, we decide which region R_j to add. Step 2, prune the tree: cut the unnecessary branches

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Step 1: How do we decide on R_j s?

Training error: For any fixed partition, R_1, \dots, R_J , the training error is defined as

$$RSS = \sum_{j=1}^{J} \sum_{i \in R_j} (y_i - \hat{y}_{R_j})^2$$

 \hat{y}_{R_j} = mean response for training observations in jth box **Goal**:

Find optimal boxes R_1, \dots, R_J that minimize

$$\sum_{j=1}^J \sum_{i \in R_j} (y_i - \hat{y}_{R_j})^2$$

- Can't actually check every possible partition
- Instead, go for top-down greedy approach called Recursive binary splitting
- Begins at top of tree with all data points

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Uses best split at every step

Recursive Binary Splitting

In the first iteration,

• Pick X_j and s, so that splitting into $\{X \mid X_j < s\}$ and $\{X \mid X_j \ge s\}$ results in largest possible reduction in RSS

$$R_1(j,s) = \{X \mid X_j < s\}$$

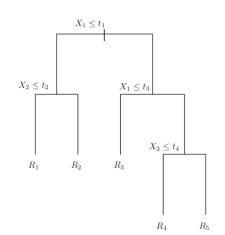
$$R_2(j,s) = \{X \mid X_j \ge s\}$$

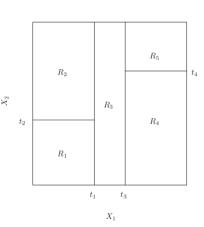
$$\sum_{i|x_i \in R_1(j,s)} (y_i - \hat{y}_{R_1})^2 + \sum_{i|x_i \in R_2(j,s)} (y_i - \hat{y}_{R_2})^2$$

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Repeat the process

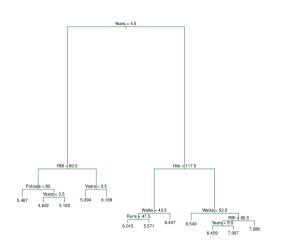
- Do this repeatedly
- Each time can split one of the previously identified regions
- Keep going until some stopping criterion is reached. E.g. until each region has at most 5
 observations





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Step 2: Pruning



- Big trees leave you open to potential overfitting
- Could just stop building earlier, but that's short sighted
- Instead, grow a big tree and prune it back
- Find subtree with best test error rate

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Too many subtrees to test them all

Weakest Link Pruning

Also called Cost complexity pruning

For every α , there is a subtree T that minimizes:

$$\sum_{m=1}^{|T|} \sum_{i|x_i \in R_m} (y_i - \hat{y}_{R_m})^2 + \alpha |T|$$

- |T| = number of terminal nodes of T
- R_m is rectangle for mth terminal node
- \hat{y}_{R_m} is mean of training observations in R_m

- $\alpha = 0$ gets entire tree
- ullet Increasing lpha penalizes size of tree
- Branches pruned from tree in nested and predictable fashion
- \bullet Easy to get trees for all vales of α

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• Pick α via CV

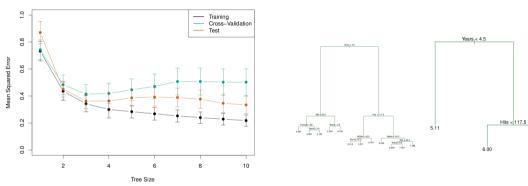
Algorithm version

Algorithm 8.1 Building a Regression Tree

- Use recursive binary splitting to grow a large tree on the training data, stopping only when each terminal node has fewer than some minimum number of observations.
- 2. Apply cost complexity pruning to the large tree in order to obtain a sequence of best subtrees, as a function of α .
- 3. Use K-fold cross-validation to choose α . That is, divide the training observations into K folds. For each $k = 1, \ldots, K$:
 - (a) Repeat Steps 1 and 2 on all but the kth fold of the training data.
 - (b) Evaluate the mean squared prediction error on the data in the left-out kth fold, as a function of α .
 - Average the results for each value of α , and pick α to minimize the average error.
- 4. Return the subtree from Step 2 that corresponds to the chosen value of α .

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Messing with α



Result of pruning is the three leaf tree on the right

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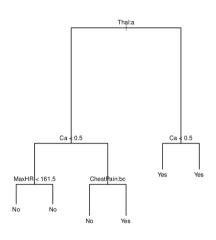
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Section 2

Classification Decision Tree

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Basic idea



- Example from heart data
- Can't use RSS, need error rate
- Could decide splits by classification error rate
- Gives too much emphasis on large classes, so use something else
- Use two other options
- \hat{p}_{mk} = proportion of training observations in R_m from the kth class
- $E = 1 \max_{k}(\hat{p}_{mk})$ Fraction of training observations not in the most common class

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Gini index

$$G = \sum_{k=1}^K \hat{
ho}_{mk} (1-\hat{
ho}_{mk})$$

- Measure of total variance across K classes
- small value if all \hat{p}_{mk} 's close to zero or 1
- Small value means node contains mostly observations from one class

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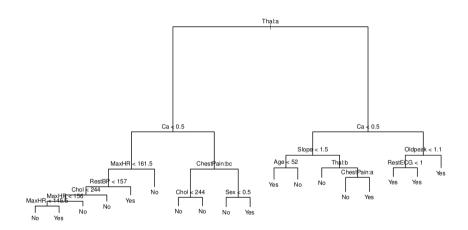
Entropy

$$D = -\sum_{k=1}^{K} \hat{p}_{mk} \log \hat{p}_{mk}$$

- Positive because $0 \le \hat{p}_{mk} \le 1$
- ullet Near zero if $\hat{
 ho}_{mk}$ all near 0 or near 1
- Small value if *m*th node has majority one class

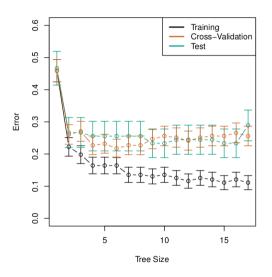
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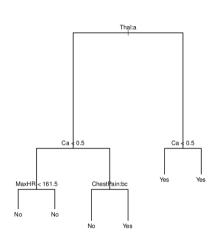
Example



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Pruning the example



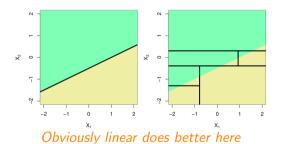


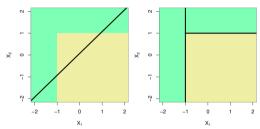
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Some coding!

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Linear models vs trees





Not going to beat this case though

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Pros/Cons

Pros:

- Trees are very easy to explain to people. Often easier to explain than linear regression!
- Trees can be displayed graphically, and are easily interpreted even by a non-expert (especially if they are small).
- Trees can easily handle qualitative predictors without the need to create dummy variables.

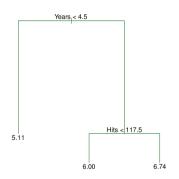
Cons:

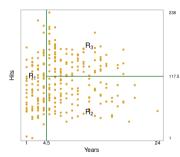
- Not as accurate as other methods of classification and regression
- Not robust: small change in data can cause large change in estimated tree
- Fix.... aggregate many decision trees

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Summarize

- Split into regions by greedily decreasing RSS
- Prune tree by using cost complexity
- Not robust Next time, figure out how to aggregate trees





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