How to Build a Tree?

Three elements:

- 1. Where to split?
- 2. When to stop?
- 3. How to predict at each leaf node?

Prediction at Leaf Nodes

Each leaf node (corresponds to region R_m) contains some samples.

Assign the prediction for a leaf node to be the average (of the response variable Y).

$$\hat{f}(X) = \sum_{m} c_m I\{X \in R_m\}.$$

$$\min_{c_m} \sum_{i=1, x_i \in R_m}^{n} (y_i - c_m)^2,$$

 $\implies c_m = \text{ average of } y_i \text{'s whose } x_i \in R_m$

Where to Split?

- ullet A split is denoted by (j,s): split the data into two parts based on whether "var $j < {\sf value}\ s$ ".
- For each split, define a split criterion $\Phi(j,s)$
 - deduction of RSS for regression
- Trees are built in a top-down greedy fashion. Start with the root: try all possible variables j=1:p and all possible split values^a, and pick the best split, i.e., the split having the best Φ value. Now, data are divided into the left node and right node. Repeat this procedure in each node.

^aFor each variable j, sort the n values (from n samples), and choose s to be a middle point of two adjacent values. So at most (n-1) possible values for s.

Goodness of Split $\Phi(j,s)$

For Regression tree, we look at the deduction of RSS if we split samples at node t into t_R and t_L :

$$\Phi(j,s) = \mathsf{RSS}(t) - \Big[\mathsf{RSS}(t_R) + \mathsf{RSS}(t_L)\Big],$$

where

$$RSS(t) = \sum_{x_i \in t} (y_i - c_t)^2,$$

$$c_t = AVE\{y_i : x_i \in t\}.$$

Note that $\Phi(j,s)$ is always positive if we split the data into two groups (even randomly), unless the mean of the left node and the one of right node are the same.

Issues: Split Categorical Predictors

- For a categorical predictor with m levels, there are $2^{m-1}-1$ possible partitions of the m labels into two groups.
- However, for regression with square error, the computation simplifies: order the m levels by their mean values of Y, and then split the categorical variable as if it were an ordered predictor there are only (m-1) potential splits.

Issues: Missing Predictor Values

- Discard any observation with missing values serious depletion of the training set.
- Splitting criteria are evaluated on non-missing observations.
- Once a split (j, s) is determined, what to do with observations missing X_j ?

- Find surrogate variables that can predict the binary outcome $"X_j < s"$ and $"X_j \ge s"$ using a one-split tree.
- Rank those surrogate variables along with the blind rule "go with majority".
- Any observation that is missing X_j is then classified with the first surrogate variable, or if missing that, the second surrogate variable (or the blind rule) is used, and etc.

When to Stop?

- A simple one : stop splitting at a node if the gain from any split is less than some pre-specified threshold.
- BUT, this is short-sighted.
- Another strategy: grow a large tree and then prune it (i.e., cut some branches).