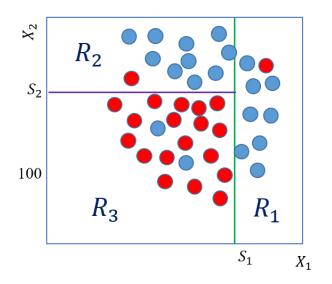
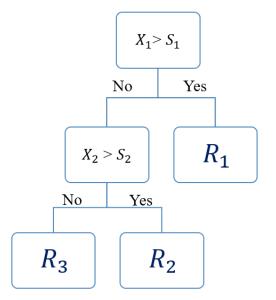


Module 9 Classification and Regression Trees CART



Prof. Pedram Jahangiry









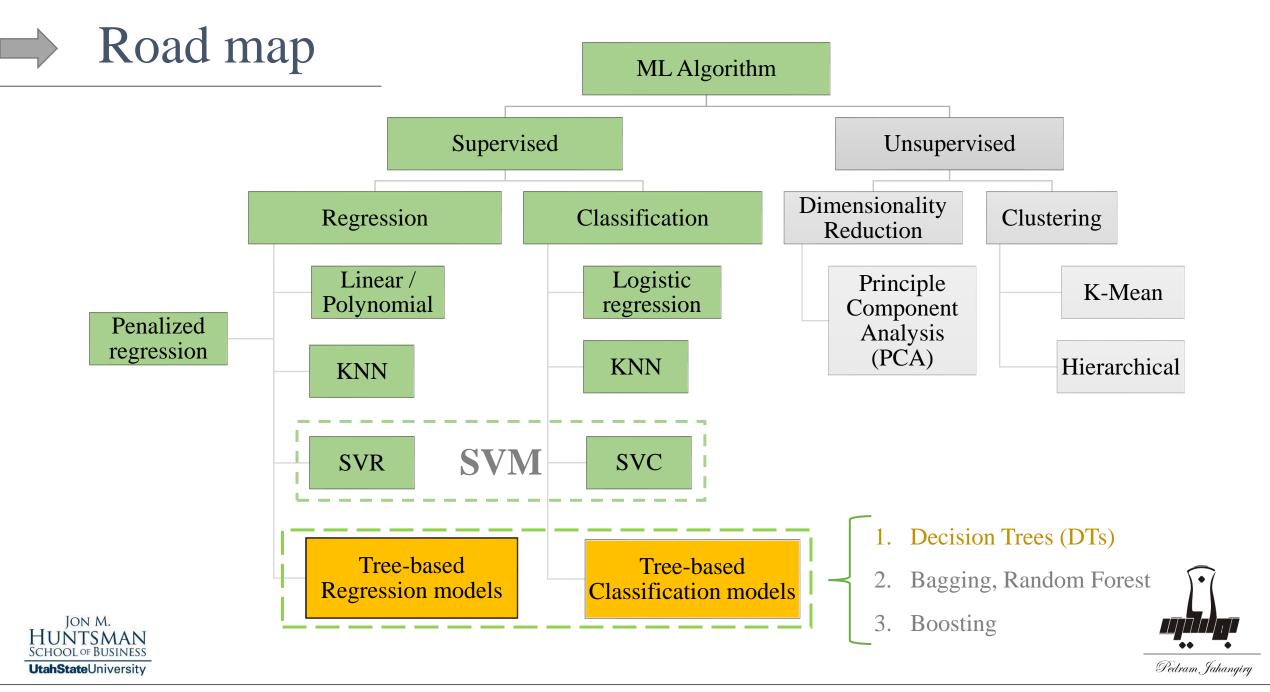


Class Modules

- Module 1- Introduction to Deep Learning
- Module 2- Setting up Machine Learning Environment
- Module 3- Linear Regression (Econometrics approach)
- Module 4- Machine Learning Fundamentals
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Topics

Part I

- 1. Decision Trees Definitions
- 2. Decision Trees criteria
 - MSE
 - Error Rate
 - Gini Index
 - Entropy

Part II

- 1. Regression Trees
- 2. Classification Trees



Part III

- 1. Pruning a tree
- 2. Hyperparameters

Part IV

- 1. Pros and Cons
- 2. Applications in Finance



Part I Decision Trees definitions and criteria



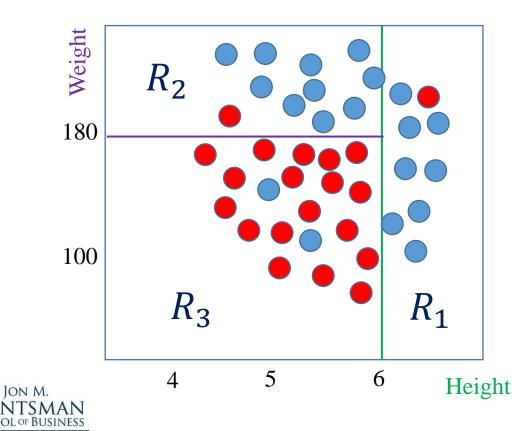


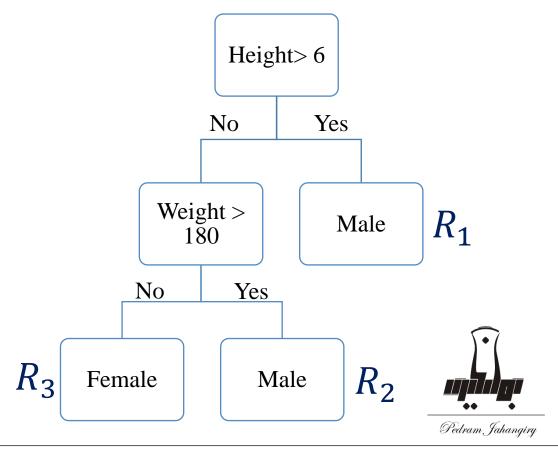


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Decision Trees Definitions

- DTs are ML algorithms that progressively divide data sets into smaller data groups based on a descriptive feature, until they reach sets that are small enough to be described by some label.
- DTs apply a top-down approach to data, trying to group and label observations that are similar.



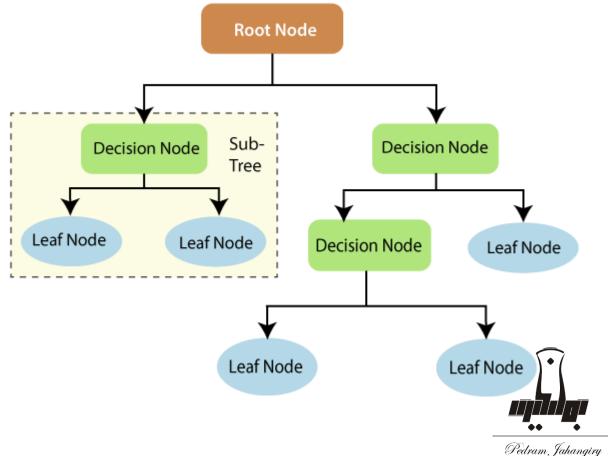




Decision Trees Definitions

- When the target variable consists of real numbers: regression trees
- When the target variable is categorical: classification trees
- Terminology:
 - ✓ Root node
 - ✓ Splitting
 - ✓ Branch
 - ✓ Decision node (internal node)
 - ✓ Leaf node (terminal node)
 - ✓ Sub-tree
 - ✓ Depth (level)







Decision Trees Criteria

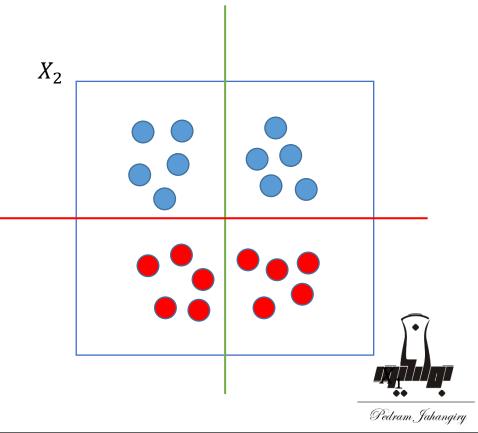
• Which split adds the most information gain (minimum impurity)?

• Regression trees: MSE

• Classification trees:

- 1. Error rate
- 2. Entropy
- 3. Gini Index

Control how a Decision Tree decides to split the data





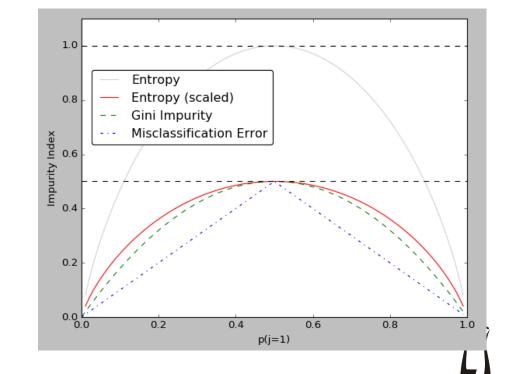


Decision Trees Criteria

- Entropy: Measures the impurity or randomness (uncertainty) in the data points
- Gini Index: Measure how often a randomly chosen element would be incorrectly labeled
- For both Entropy and Gini, 0 expresses all the elements belong to a specified class (pure)
- Different decision tree algorithms utilize different impurity metrics

$$entropy = -\sum_{j} p_{j} \log_{2}(p_{j})$$

$$Gini = 1 - \sum_{i} p_j^2$$



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Part II Regression / Classification Trees! How does a decision tree work?



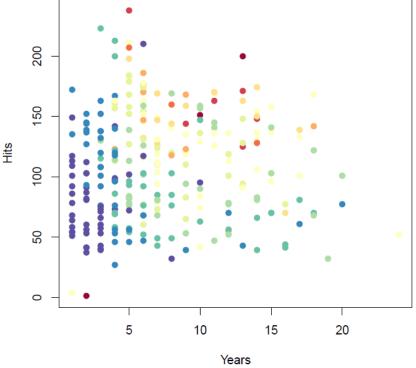




Regression Trees

Baseball Salary is color-coded from low (blue, green) to high (yellow, red)

- DTs apply a top-down approach to data, trying to group and label observations that are similar.
- The main questions in every decision-making process:
- 1. Which feature to start with?
- 2. Where to put the split (cut off)?

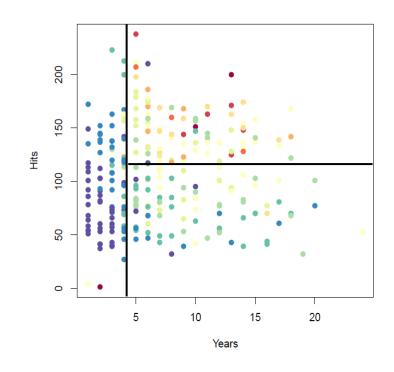






Interpreting the results

- Based on color-coded salary, it seems that years is the most important factor in determining salary.
- For less experienced players, the number of hits seems irrelevant.
- Among more experienced players thought, players with more hits tend to have higher salaries.
- As one can see, the model is very easy to display, interpret and explain.





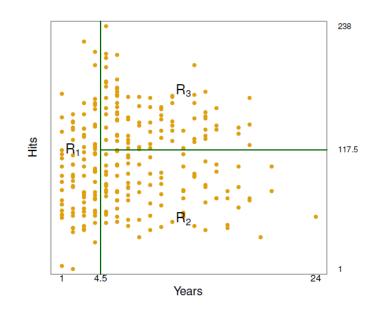




Tree building process

- Divide the feature space into J distinct and non-overlapping regions.
- For every observation that falls into the region R_j , we make the same prediction, which is simply the mean of the target values for the training observations in R_j .
- The goal is to find rectangles $R_1, R_2, ..., R_j$ that minimize the RSS:

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



• Where \hat{y}_{R_i} is the mean target for the training observations within the j^{th} rectangle.







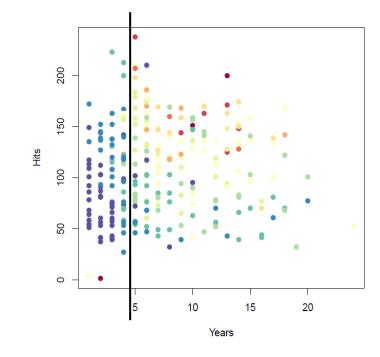
Tree building process: Recursive Binary Splitting

- How does the algorithm select X_i and the split s?
- X_j and s are selected such that splitting the feature space into the regions $\{X|X_j < s\}$ and $\{X|X_j \ge s\}$ leads to the largest possible reduction in RSS.

$$R_1(j,s) = \{X | X_j < s\} \text{ and } R_2(j,s) = \{X | X_j \ge s\}$$

• Seeking for the value of *j* and *s* that minimized the following equation:

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



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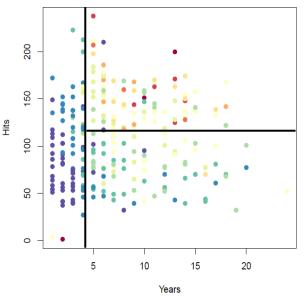
• The best split is made at that particular step, rather than looking ahead and picking a split that will lead to a better tree in some future step.

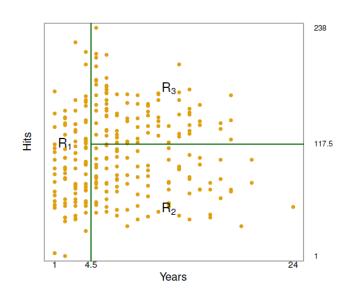


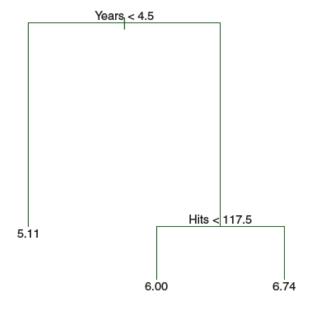


Tree building process: Recursive Binary Splitting

- Next, the algorithm repeats the process, looking for the best feature and best split in order to split the data further to minimize the RSS within each of the resulting regions.
- The process continues until a stopping criterion is reached; for instance, continues until no region contains more than a fixed number of observations.



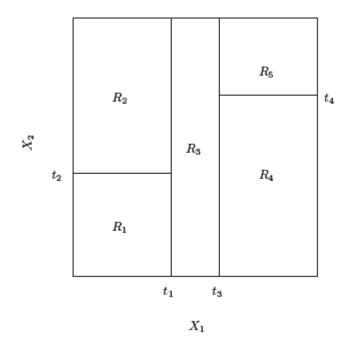


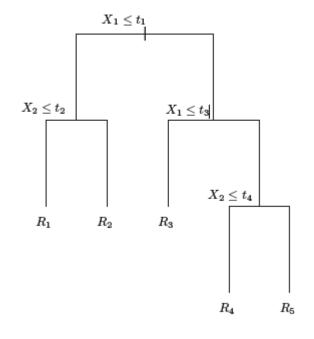


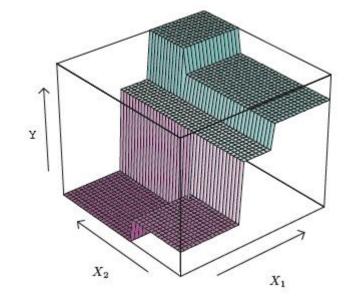




A Five-Region Example of Recursive Binary Splitting







The output of recursive binary splitting on a two-dimensional example

A tree corresponding to the partition in the left panel.

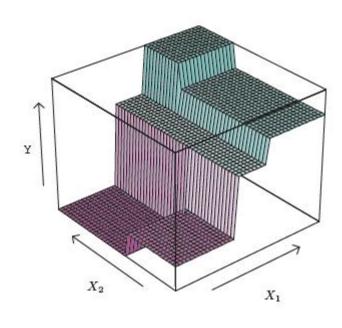
A perspective plot of the prediction surface corresponding to that tree.







Overfitting?



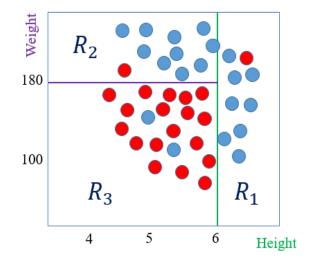


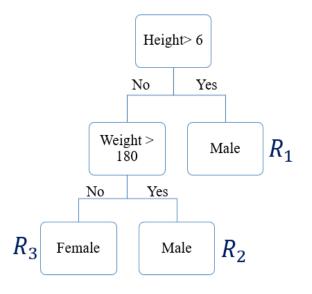




Classification Trees

- Classification trees are very similar to regression trees, except that it is used to predict a qualitative response rather than aquantitative one.
- The prediction of the algorithm at each terminal node will be the category with the majority of data points i.e., the most commonly occurring class.





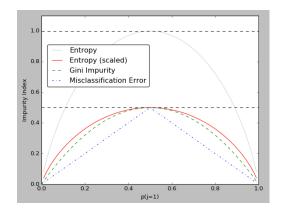






Classification Trees (details)

- Just as in the regression setting, the recursive binary splitting is used to grow a classification tree. However, instead of RSS we will be using one of the following impurity criteria:
 - 1. Classification error rate: \longrightarrow $E = 1 \max_{k}(\hat{p}_{mk})$
 - 2. Gini index: $G = \sum_{k=1}^{K} \hat{p}_{mk} (1 \hat{p}_{mk})$
 - 3. Cross entropy: $D = -\sum_{i=1}^{\infty} \hat{p}_{mk} \log \hat{p}_{mk}$



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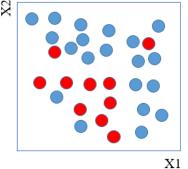
- \hat{p}_{mk} represents the proportion of training observations in the m^{th} region from the k^{th} class.
- Classification error rate is not sufficiently sensitive to node purity and in practice either Gini or Cross entropy is preferred.

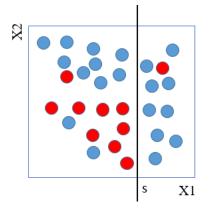


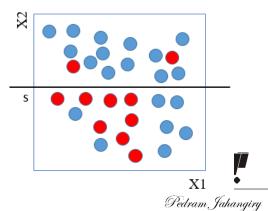


Decision Tree Metrics (Simple Example)

Node	$Gini \\ 1 - \sum_j p_j^2$	Cross entropy $-\sum_{j}p_{j}\log(p_{i})$	Error rate $1 - \max(p_i)$
Entire training data before split	$\left\{1 - \left(\left(\frac{10}{30}\right)^2 + \left(\frac{20}{30}\right)^2\right)\right\} = 0.44$	$-\left\{ \left(\frac{10}{30} \log \frac{10}{30} + \frac{20}{30} \log \frac{20}{30} \right) \right\} = 0.64$	$1 - \max\left\{\frac{10}{30}, \frac{20}{30}\right\}$ $= 1 - \frac{20}{30} = 0.333$





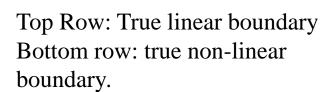


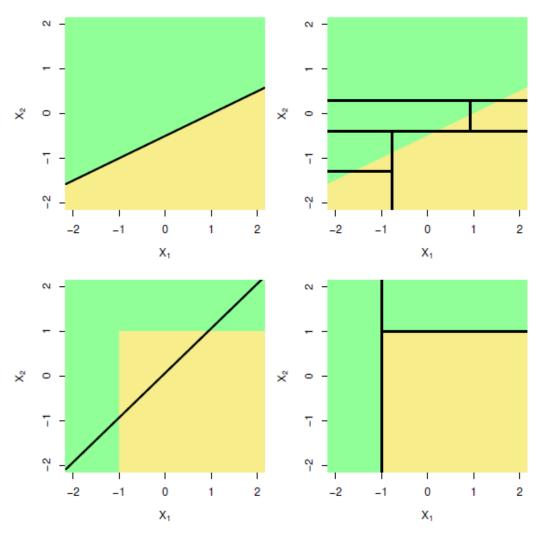




Trees Versus Linear Models

Left column: linear model; Right column: tree-based model









Part III

Pruning a tree
Tunning hyper parameters







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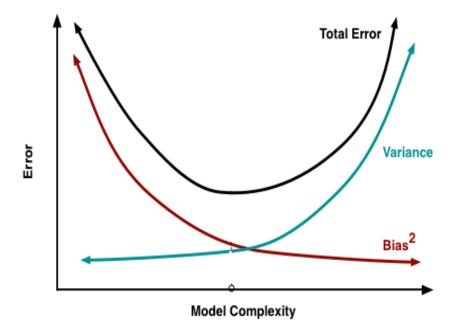
Pruning a tree

- A smaller tree with fewer splits may lead to lower variance and better interpretation (but) at the cost of higher bias.
- Smaller trees are too short-sighted:

"a seemingly worthless split early on in the tree might be followed by a very good split, a split that leads to a large reduction in RSS/impurity index later on"

- A better strategy is to grow a very large tree, this may produce good predictions on the training set, but is likely to overfit the data, leading to poor test set performance.
- So, we need to prune it back in order to obtain a subtree.
- Cost complexity pruning is used to do this.









Cost complexity pruning (weakest link pruning)

- Consider a sequence of trees indexed by a nonnegative tuning parameter α .
- For each value of α there corresponds a subtree $T \subset T_0$ such that the following objective function is minimized.

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

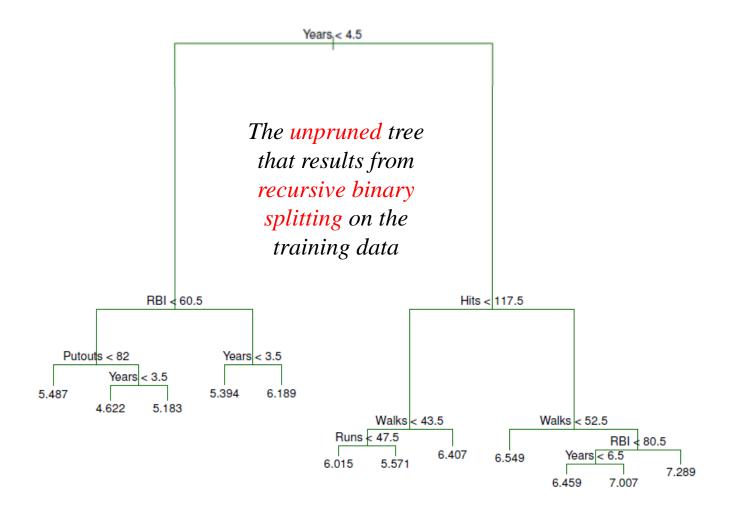
- |T| indicates the number of terminal nodes of the tree T
- R_m is the rectangle corresponding to m^{th} terminal node and
- \hat{y}_{Rm} is the mean of the training observations in R_m
- α controls the bias variance trade off and is determined by cross validation.
- Lastly, we return to full data set and obtain the subtree corresponding to α







Salary example continued









Finding the optimal α or T

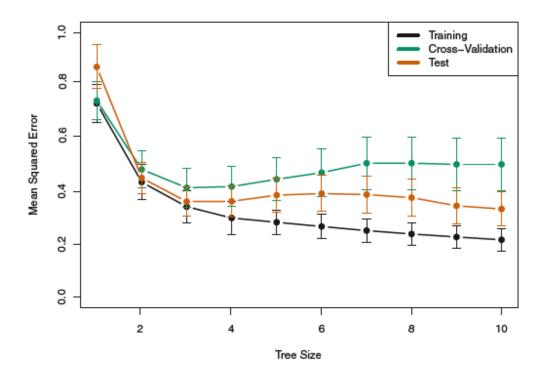


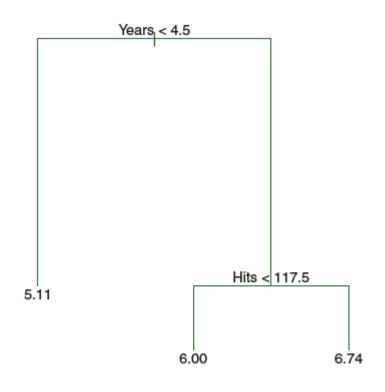
FIGURE 8.5. Regression tree analysis for the Hitters data. The training, cross-validation, and test MSE are shown as a function of the number of terminal nodes in the pruned tree. Standard error bands are displayed. The minimum cross-validation error occurs at a tree size of three.

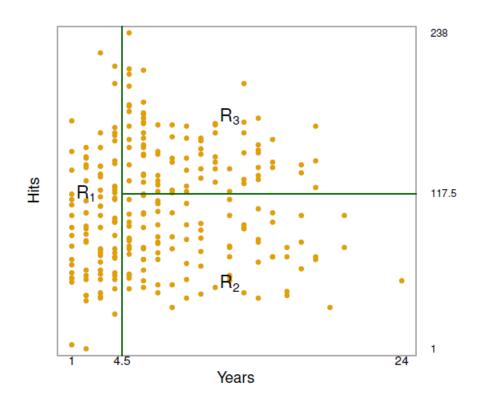






The optimal (pruned) tree





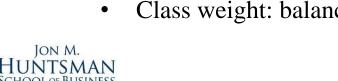




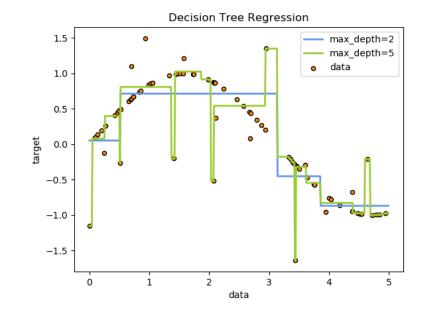


Other hyperparameters

- ✓ To avoid overfitting, regularization parameters can be added to the model such as:
 - Maximum depth of the tree
 - Minimum population at a node
 - Maximum number of decision nodes
 - Minimum impurity decrease (info gain)
 - Alpha (complexity parameter)
- Other hyperparameters are:
 - Criterion: gini, entropy
 - Splitter: best, random
 - Class weight: balanced, none



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Part IV Pros and Cons Applications in finance







DTs' Pros and Cons

Pros:

- Easy to interpret and visualize
- Can easily handle categorical data without the need to create dummy variables
- Can easily capture Non-linear patterns
- Can handle data in its raw form (no preprocessing needed). Why?
- Has no assumptions about distribution because of the non-parametric nature of the algorithm

Cons:

- Poor level of predictive accuracy.
- Sensitive to noisy data. It can overfit noisy data. Small variations in data can result in the different decision tree*.
 - *This can be reduced by bagging and boosting algorithms.









DTs' Applications in finance

- Enhancing detection of fraud in financial statements,
- Generating consistent decision processes in equity and fixed-income selection
- Simplifying communication of investment strategies to clients.
- Portfolio allocation problems.





Parameter	Description	Default	Options
max_depth	The maximum number of levels: split the nodes until max_depth has been reached. All leaves are pure or contain fewer samples than min_samples_split.	None	int
max_features	Number of features to consider for a split.	None	None: all features int: # features float: fraction auto, sqrt: sqrt(n_ features) log2: log2(n_ features)
max_leaf_nodes	Split nodes until creating this many leaves.	None	None: unlimited int
min_impurity_decrease	Split node if impurity decreases by at least this value.	0	float
min_samples_leaf	A split will only be considered if there are at least min_samples_ leaf training samples in each of the left and right branches.	1	int; float (as a percent of N)
min_samples_split	The minimum number of samples required to split an internal node.	2	int; float (percent of N)
min_weight_fraction_ leaf	The minimum weighted fraction of the sum total of all sample weights needed at a leaf node. Samples have equal weight unless sample_weight is provided in the fit method.	0	

Appendix A

Scikit-Learn decision tree parameters.



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