Growing decision trees STAT 471

Rolling into a new unit!

Unit 1: Intro to modern data mining

Unit 2: Tuning predictive models

Unit 3: Regression-based methods

Unit 4: Tree-based methods

Unit 5: Deep learning

Lecture 1: Growing decision trees

Lecture 2: Tree pruning and bagging

Lecture 3: Random forests

Lecture 4: Boosting

Lecture 5: Unit review and quiz in class

Homework 4 due the following Wednesday.

Leaving the land of linearity

Most methods covered so far based on $\hat{\beta}_0 + \hat{\beta}_1 X_1 + \cdots + \hat{\beta}_p X_p$ in some way:

- Linear regression
- Logistic regression
- Ridge, lasso, elastic net

Notable exception: K-nearest neighbors (recall Unit 2)

In Unit 4 we will leave the land of linearity.

Entering the land of trees and forests

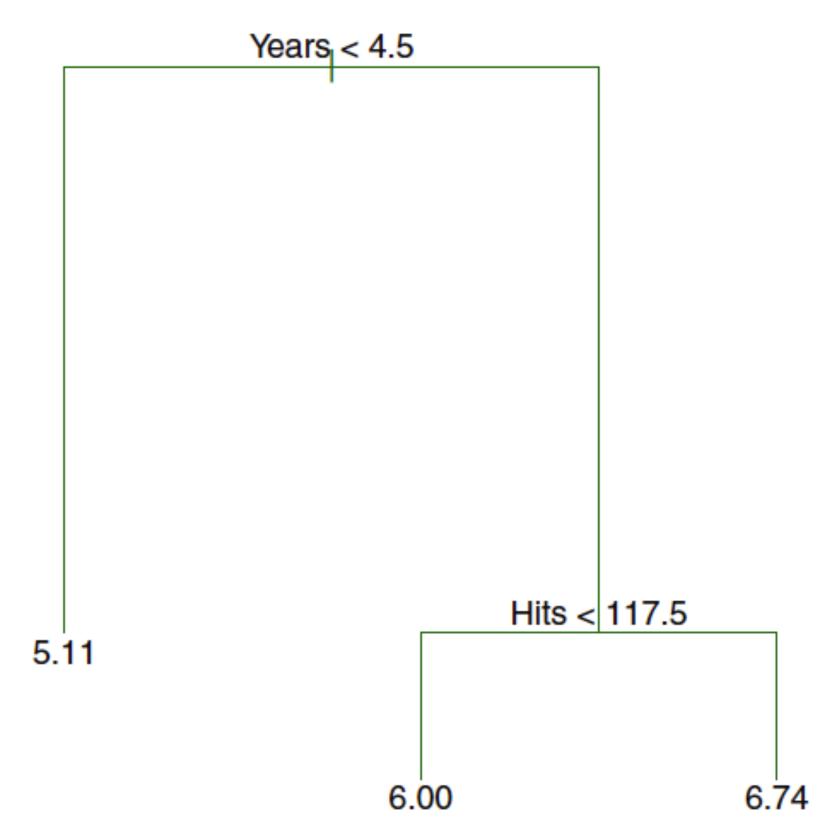
Decision trees (lectures 1 and 2) are predictive models based on recursively partitioning the feature space.

Their prediction rules can be nicely illustrated and are very interpretable.

However, trees are somewhat unstable and do not give the best prediction performance.

Nevertheless, trees can be used as building blocks for state-of-the-art prediction performance:

- Random forests (lecture 3)
- Boosting (lecture 4)



Predicting baseball players' salaries based on years played and number of hits in the previous year.

Tree-based models versus linear models

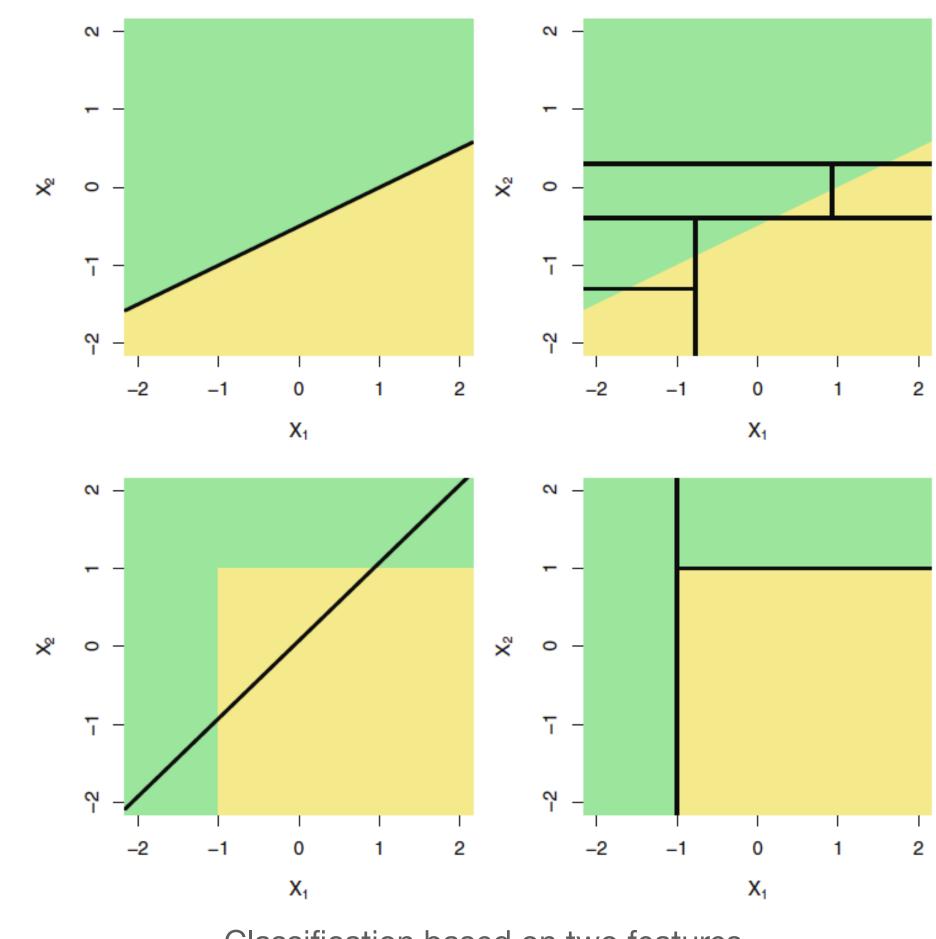
Which perform better?

Neither tree-based nor linear models dominate the other.

Each prediction method works better when the underlying trend in the data matches its modeling choice.

E.g. for classification:

- Linear model → linear decision boundary
- Decision tree → unions of rectangles



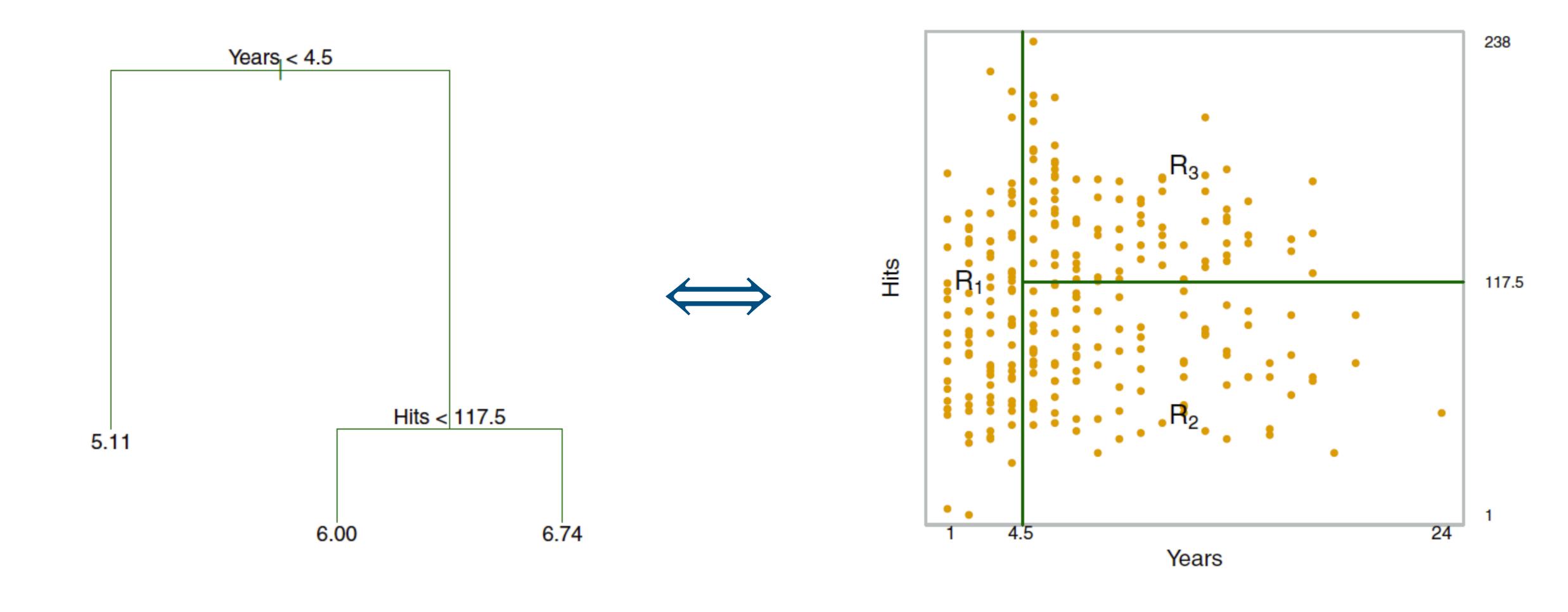
Classification based on two features (colors indicate the two classes).

Hitters data

Major League Baseball Data from the 1986 and 1987 seasons.

- Observations: 322 MLB players
- Response: Salary (1987 annual salary on opening day in thousands of dollars)
- Features: Assists, AtBat,...,Hits,...,Years (19 total)

Tree \iff Partition into nested, axis-aligned rectangles



Mathematical expression of the prediction rule

A trained tree consists of:

- M regions $\widehat{R}_1, ..., \widehat{R}_M$
- response values $\hat{c}_1, ..., \hat{c}_M$

For a new feature vector X^{test} , predict the constant value \widehat{c}_m for region \widehat{R}_m :

$$\widehat{Y}^{\text{test}} = \sum_{m=1}^{M} \widehat{c}_m \cdot I(X^{\text{test}} \in \widehat{R}_m).$$

 R_2 R_3 R_4 R_4 R_5 R_4 R_4 R_4 R_4 R_4 R_4 R_4 R_5 R_4 R_4 R_4 R_4 R_5 R_4 R_4 R_4 R_5 R_6 R_7 R_8 R_8 R_8 R_8

(continuous or categorical response)

Partitioning for continuous and categorical features

Suppose we partition on X_i .

- If X_j is continuous, we just find a split point s and split into $\{X: X_j < s\}$ and $\{X: X_j \geq s\}$.
- If X_j is categorical, e.g. with levels $\{a,b,c,d,e\}$, then we need to split the levels into two groups, e.g. $\{a,c\}$ and $\{b,d,e\}$, giving the partitions $\{X:X_j\in\{a,c\}\}$ and $\{X:X_j\in\{b,d,e\}\}$.

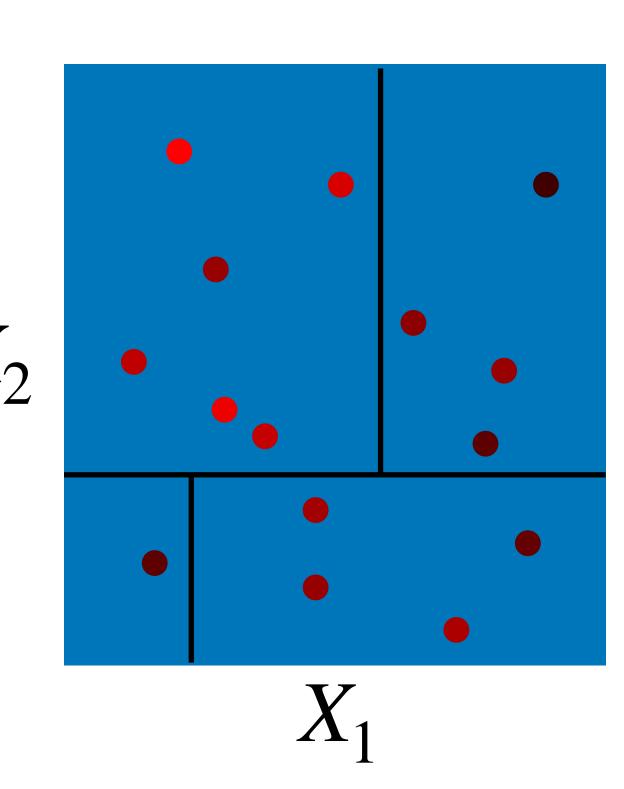
The squared error objective

As usual, we are given a training dataset $(X_1, Y_1), \ldots, (X_n, Y_n)$.

For a fixed M, we seek rectangles $\widehat{R}_1, \ldots, \widehat{R}_M$ and values $\widehat{c}_1, \ldots, \widehat{c}_M$ to minimize the residual sum of squares:

$$\widehat{R}_{1},...,\widehat{R}_{M},\widehat{c}_{1},...,\widehat{c}_{M} = \underset{R_{1},...,R_{M},c_{1},...,c_{M}}{\operatorname{arg min}} \sum_{i=1}^{n} (Y_{i} - \widehat{Y}_{i})^{2};$$

$$\widehat{Y}_i = \sum_{m=1}^{M} c_m \cdot I(X_i \in R_m).$$



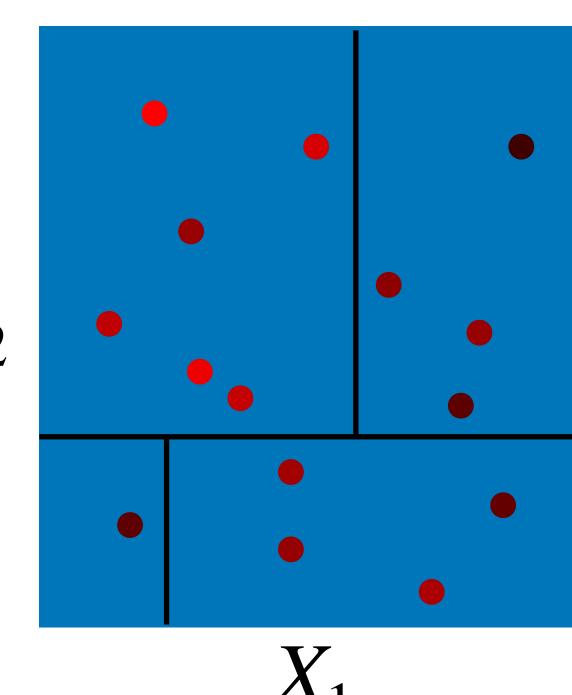
Optimal \hat{c}_m given \widehat{R}_m

First let's consider a simpler problem, where rectangles $\widehat{R}_1, ..., \widehat{R}_M$ are given:

$$\hat{c}_1, \dots, \hat{c}_M = \arg\min_{c_1, \dots, c_M} \sum_{i=1}^n (Y_i - \hat{Y}_i)^2; \quad \hat{Y}_i = \sum_{m=1}^M c_m \cdot I(X_i \in \widehat{R}_m).$$

We're fitting a constant to each region, so the solution is

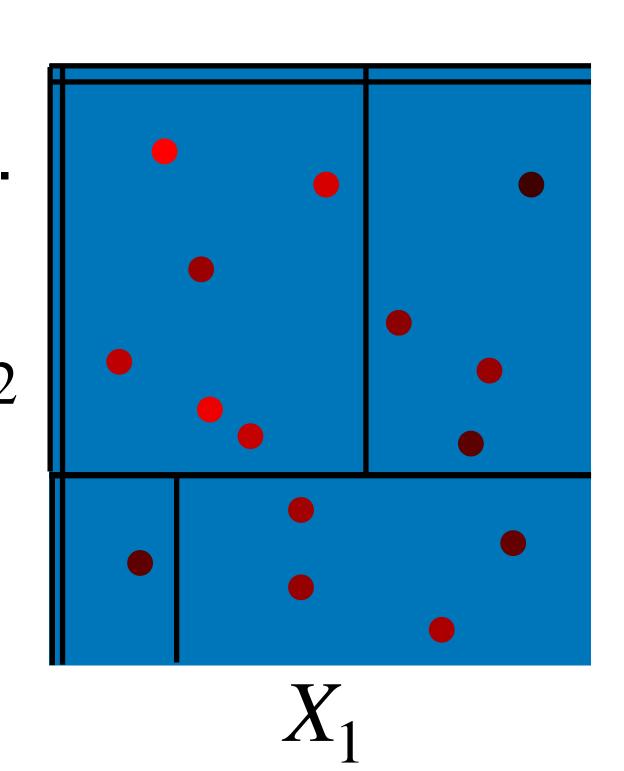
$$\widehat{c}_m = \operatorname{mean}\left(\{Y_i : X_i \in \widehat{R}_m\}\right).$$



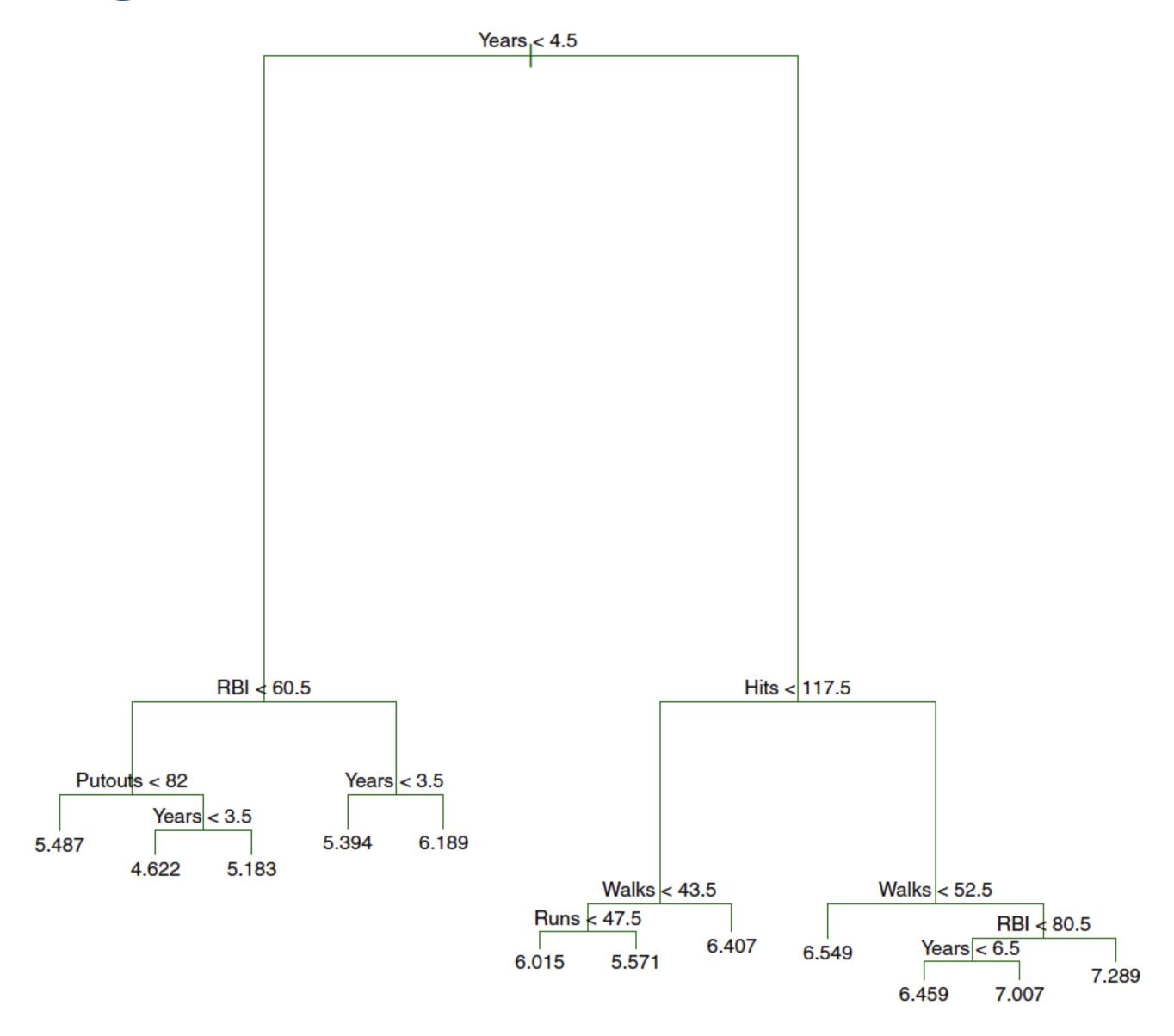
Finding the rectangles \widehat{R}_m

The optimal set of rectangles is computationally intractable to find. In practice, we employ a greedy top-down algorithm:

- 1. Fit constant model to the entire space and calculate RSS.
- 2. Find split of the whole region that decreases RSS the most.
- 3. Find next split that decreases the RSS the most.
- 4. Repeat until there are M regions.



Final output



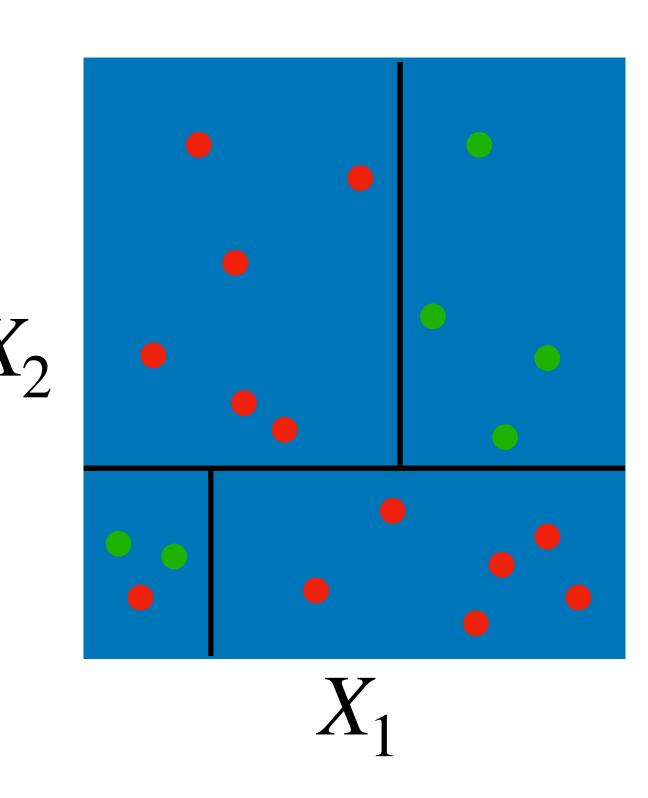
The misclassification error objective

As usual, we are given a training dataset $(X_1, Y_1), \ldots, (X_n, Y_n)$, where the response Y is binary.

For a fixed M, we seek rectangles $\widehat{R}_1, \ldots, \widehat{R}_M$ and values $\widehat{c}_1, \ldots, \widehat{c}_M$ to minimize the misclassification loss:

$$\widehat{R}_{1},...,\widehat{R}_{M},\widehat{c}_{1},...,\widehat{c}_{M} = \underset{R_{1},...,R_{M},c_{1},...,c_{M}}{\operatorname{arg min}} \frac{1}{n} \sum_{i=1}^{n} I(Y_{i} \neq \widehat{Y}_{i});$$

$$\widehat{Y}_i = \sum_{m=1}^{M} c_m \cdot I(X_i \in R_m).$$



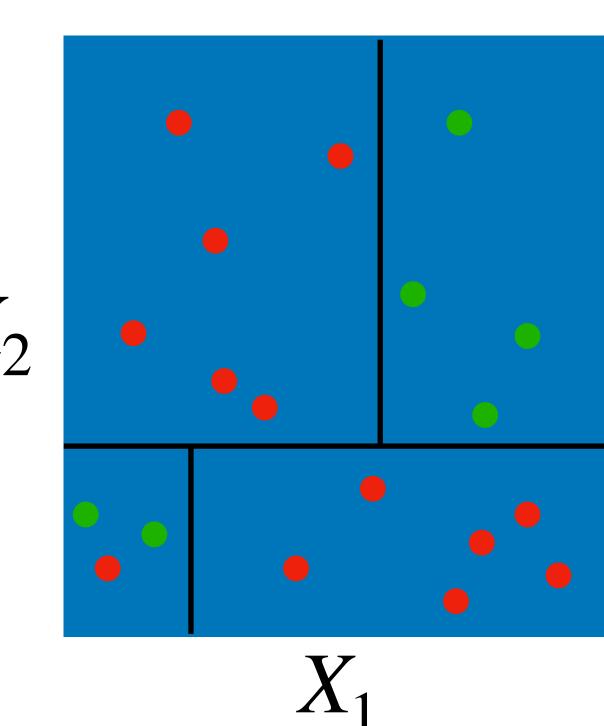
Optimal \hat{c}_m given \widehat{R}_m

First let's consider a simpler problem, where rectangles $\widehat{R}_1, \ldots, \widehat{R}_M$ are given:

$$\hat{c}_1, \dots, \hat{c}_M = \underset{c_1, \dots, c_M}{\operatorname{arg min}} \frac{1}{n} \sum_{i=1}^n I(Y_i \neq \hat{Y}_i); \quad \hat{Y}_i = \sum_{m=1}^M c_m \cdot I(X_i \in \widehat{R}_m).$$

We're fitting the same category to each region, so the solution is the majority vote:

$$\widehat{c}_m = \text{mode}\left(\{Y_i : X_i \in \widehat{R}_m\}\right).$$



Finding the rectangles \widehat{R}_m

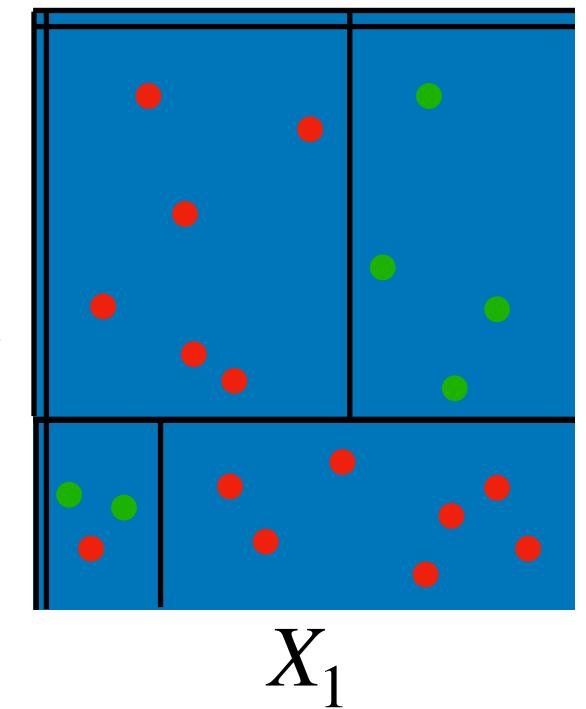
We use a very similar greedy recursive splitting algorithm.

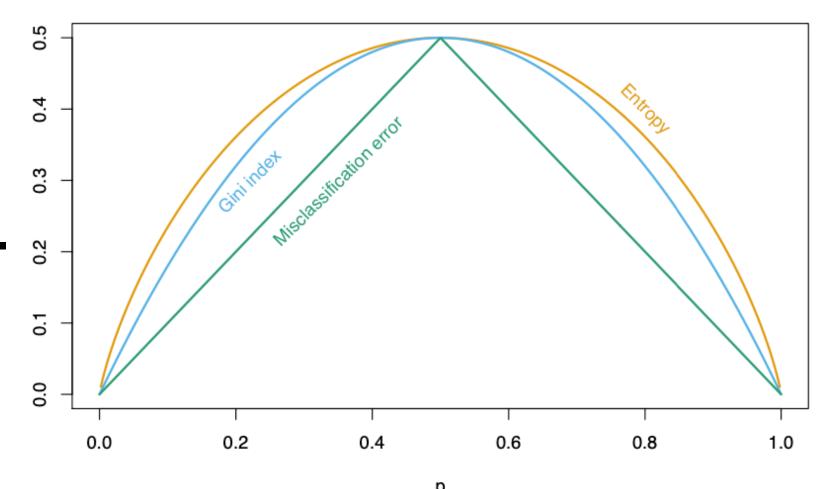
Let \widehat{p}_m be proportion of class 1 in region m. The misclassification error in that region is $\min(\widehat{p}_m, 1 - \widehat{p}_m)$.

Misclassification error not sensitive enough to find good split points at each step; instead, evaluate impurity using

Gini index =
$$2\hat{p}_m(1 - \hat{p}_m)$$
.

Find split that minimizes the total impurity of the regions. 3

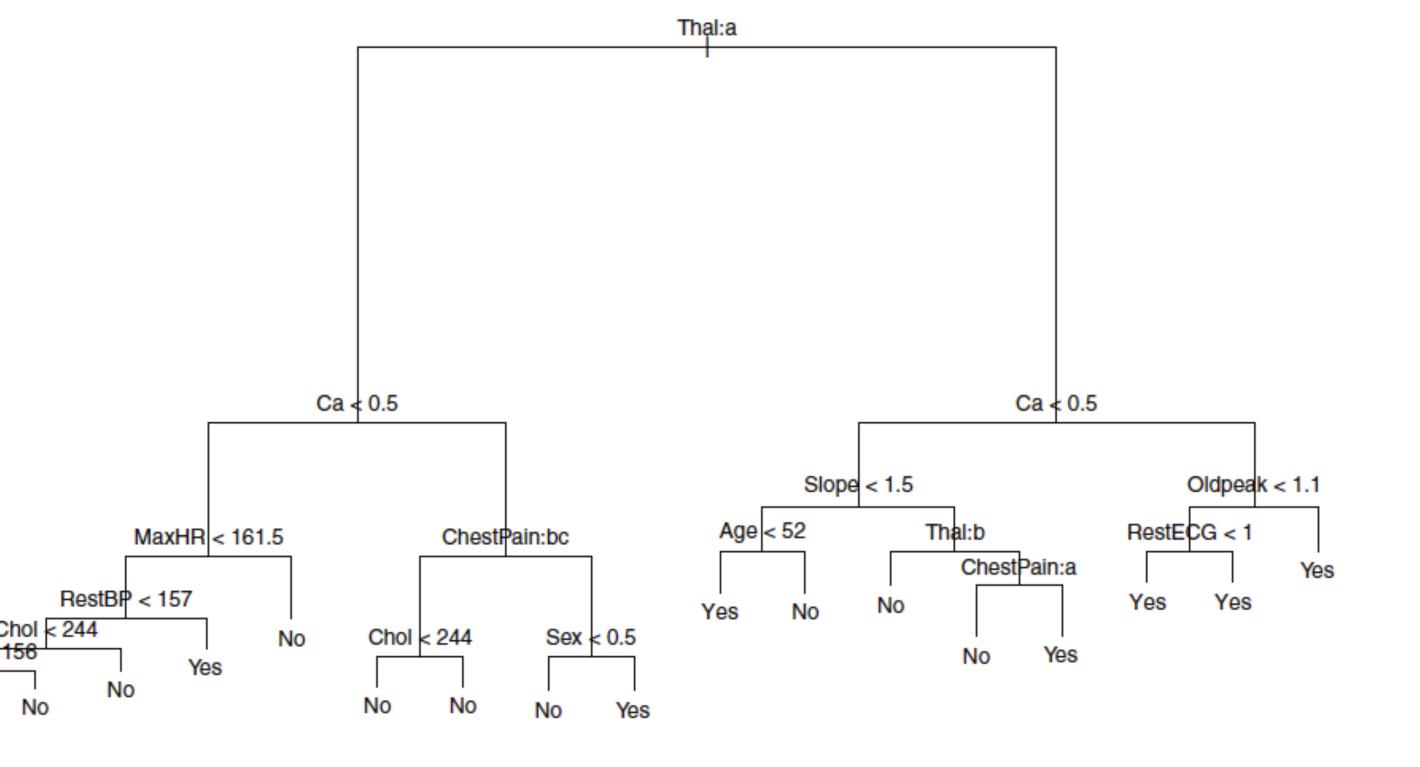




Final output

Example: Heart disease data set.

- 303 patients with chest pain
- Binary response HD (heart disease)
- 13 demographic and clinical features



Note: Classification trees extend seamlessly to more than two classes!

Summary

- Decision trees partition the feature space into axis-aligned nested rectangles, producing a constant prediction for feature vectors in each rectangle.
- Decision trees are built by recursively choosing
 - The optimal rectangle to split
 - The optimal feature to split that rectangle on
 - The optimal split-point for that feature
- Regression and classification trees aim to minimize squared error and misclassification losses, respectively.

