

# Trees

Applied Multivariate Statistics – Spring 2013

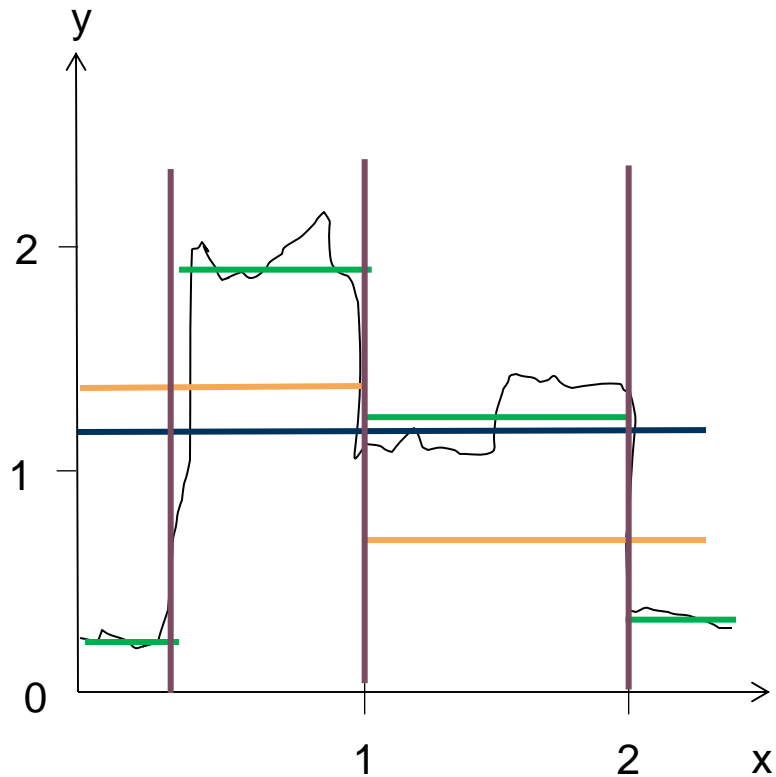


# Overview

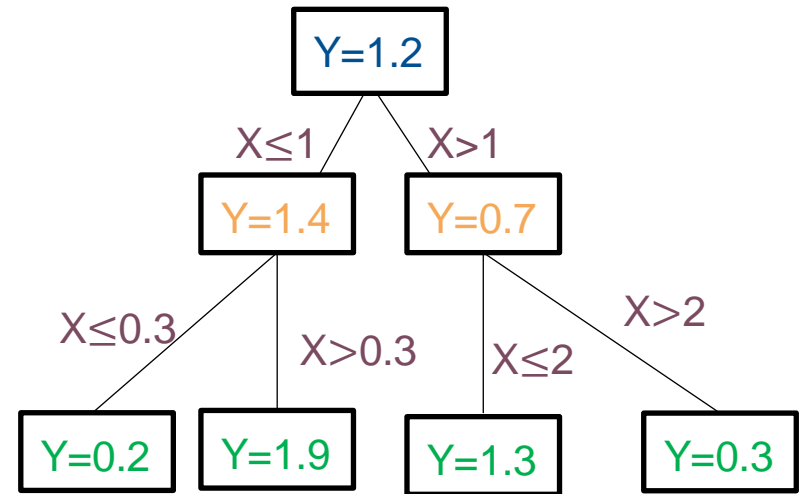
- Intuition for Trees
- Regression Trees
- Classification Trees

# Idea of Trees: Regression Trees

## Continuous response



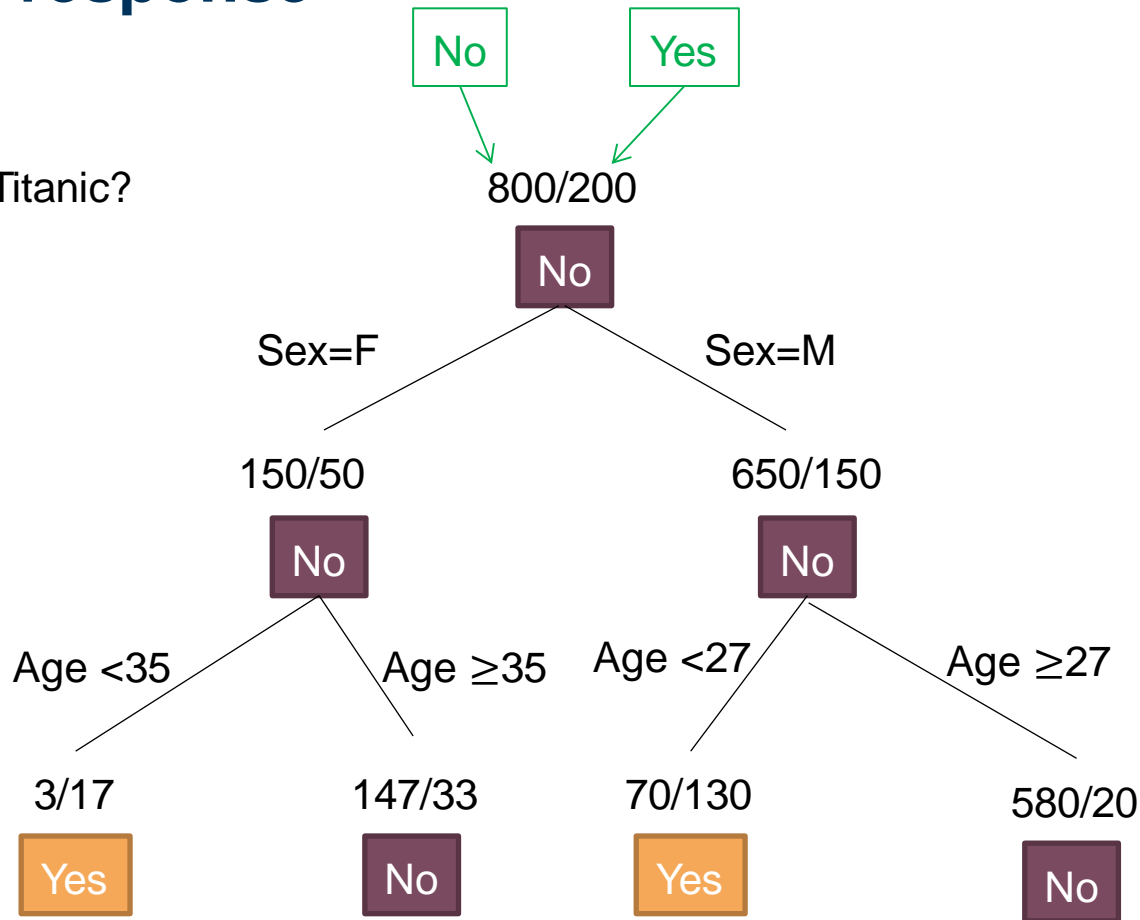
Binary Tree



# Idea of Trees: Classification Tree

## Discrete response

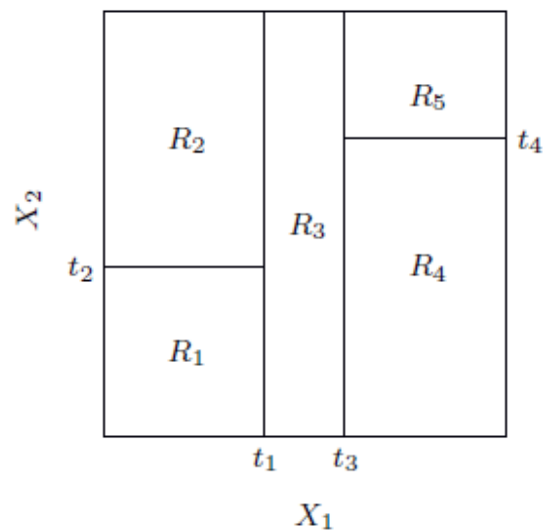
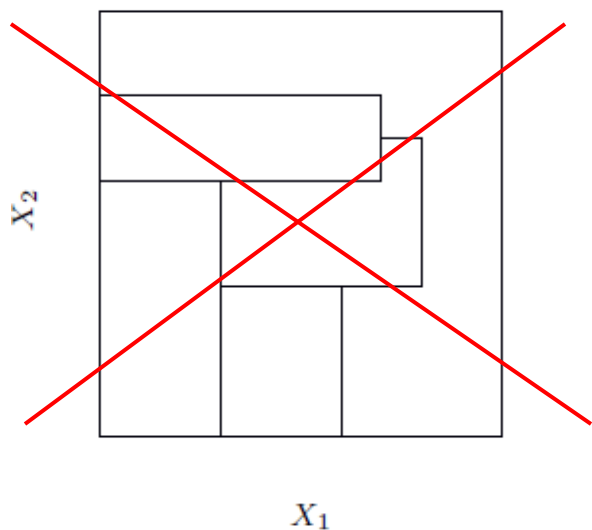
Survived in Titanic?



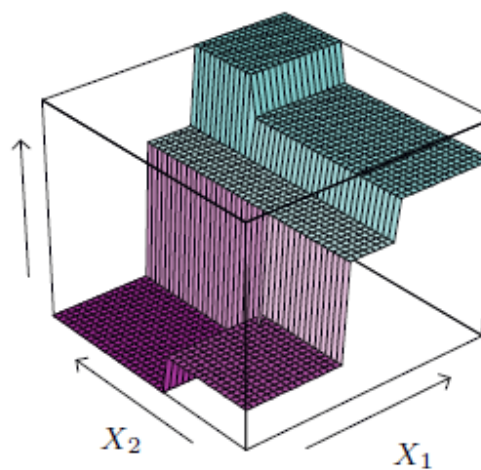
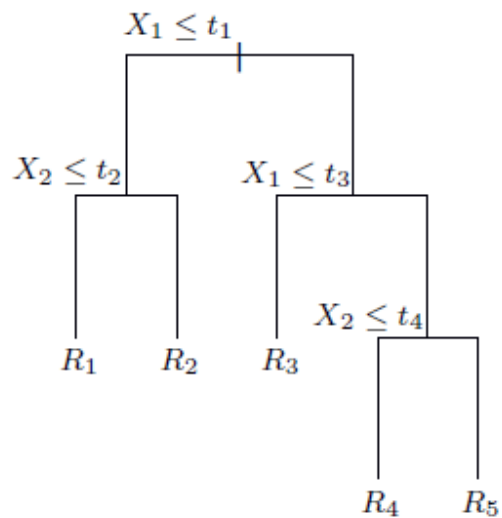
Missclassification rate:

- Total:  $(3+33+70+20) / 1000 = 0.126$
- "Yes"-class:  $53/200 = 0.26$
- "No"-class:  $73/800 = 0.09$

# Intuition of Trees: Recursive Partitioning

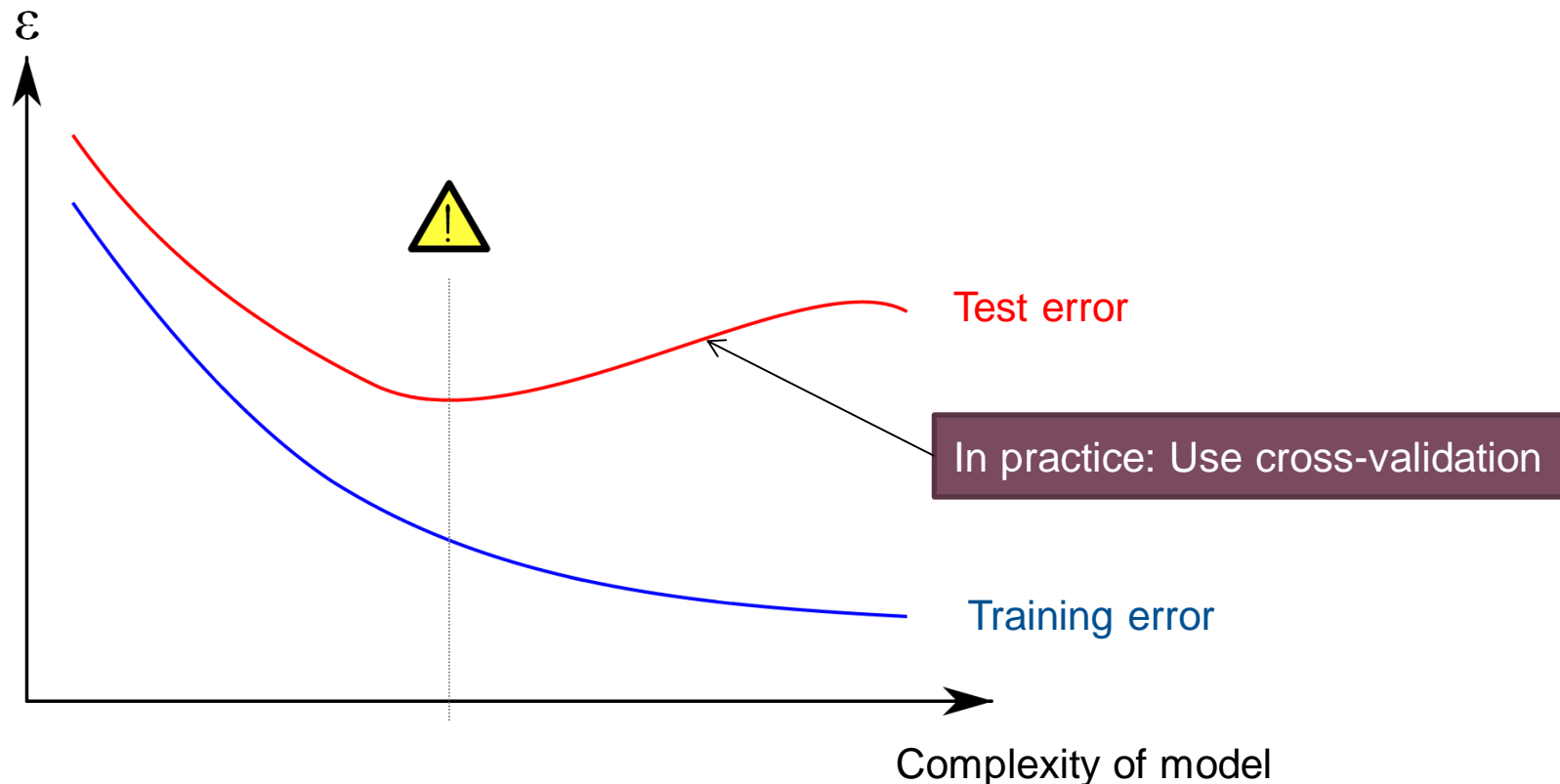


For simplicity:  
Restrict to  
recursive  
binary splits



# Fighting overfitting: Cost-complexity pruning

Overfitting: Fitting the training data **perfectly** might not be good for predicting future data



For trees:

1. Fit a very detailed model
2. Prune it using a complexity penalty to optimize cross-validation performance

# Building Regression Trees 1/2

- Assume given partition of space  $R_1, \dots, R_M$

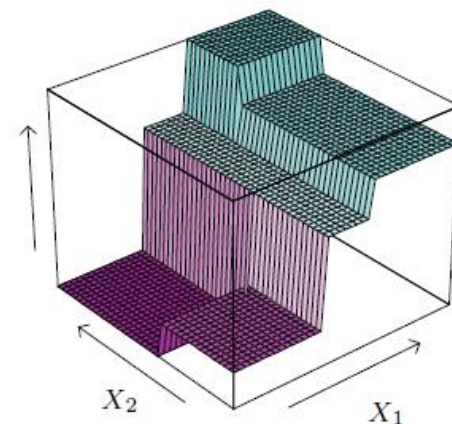
Tree model: 
$$f(x) = \sum_{m=1}^M c_m I(x \in R_m)$$

- Goal is to minimize sum of squared residuals:

$$\sum (y_i - f(x_i))^2$$

- Solution: Average of data points in every region

$$\hat{c}_m = \text{ave}(y_i | x_i \in R_m)$$



## Building Regression Trees 2/2

- Finding the best binary partition is computationally infeasible
- Use greedy approach: For variable  $j$  and split point  $s$  define the two generated regions:

$$R_1(j, s) = \{X | X_j \leq s\} \quad \text{and} \quad R_2(j, s) = \{X | X_j > s\}$$

- Choose splitting variable  $j$  and split point  $s$  that solve:

$$\min_{j, s} \left[ \min_{c_1} \sum_{x_i \in R_1(j, s)} (y_i - c_1)^2 + \min_{c_2} \sum_{x_i \in R_2(j, s)} (y_i - c_2)^2 \right]$$

inner minimization is solved by

$$\hat{c}_1 = \text{ave}(y_i | x_i \in R_1(j, s)) \quad \text{and} \quad \hat{c}_2 = \text{ave}(y_i | x_i \in R_2(j, s))$$

- Repeat splitting process on each of the two resulting regions



# Pruning Regression Trees

- Stop splitting when some minimal node size (= nmb. of samples per node) is reached (e.g. 5)
- Then, cut back the tree again (“pruning”) to optimize the cost-complexity criterion:

$$N_m = \#\{x_i \in R_m\},$$

$$\hat{c}_m = \frac{1}{N_m} \sum_{x_i \in R_m} y_i,$$

$$Q_m(T) = \frac{1}{N_m} \sum_{x_i \in R_m} (y_i - \hat{c}_m)^2$$

“Impurity measure”



$$C_\alpha(T) = \sum_{m=1}^{|T|} N_m Q_m(T) + \alpha |T|$$

Goodness of fit

Complexity

- Tuning parameter  $\alpha$  is chosen by cross-validation

# Classification Trees

- Regression Tree:  
Quality of split measured by “Squared error”
- Classification Tree:  
Quality of split measured by general “Impurity measure”

# Classification Trees: Impurity Measures

- Proportion of class k observations in node m:

$$\hat{p}_{mk} = \frac{1}{N_m} \sum_{x_i \in R_m} I(y_i = k)$$

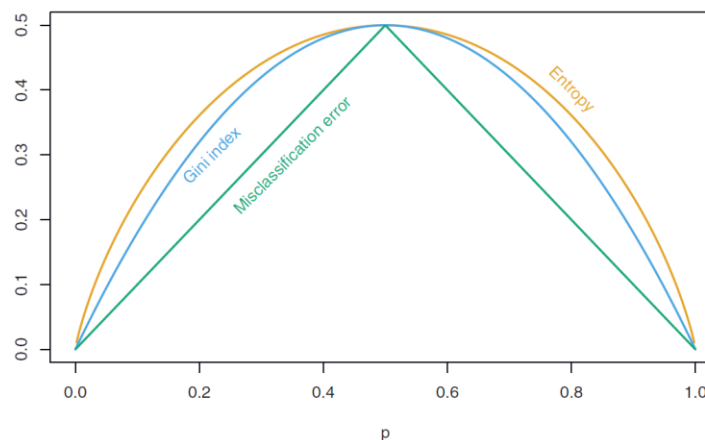
- Define majority class in node m:  $k(m)$
- Common impurity measures  $Q_m(T)$ :

Misclassification error:  $\frac{1}{N_m} \sum_{i \in R_m} I(y_i \neq k(m)) = 1 - \hat{p}_{mk(m)}$

Gini index:  $\sum_{k \neq k'} \hat{p}_{mk} \hat{p}_{mk'} = \sum_{k=1}^K \hat{p}_{mk} (1 - \hat{p}_{mk})$

Cross-entropy or deviance:  $-\sum_{k=1}^K \hat{p}_{mk} \log \hat{p}_{mk}$

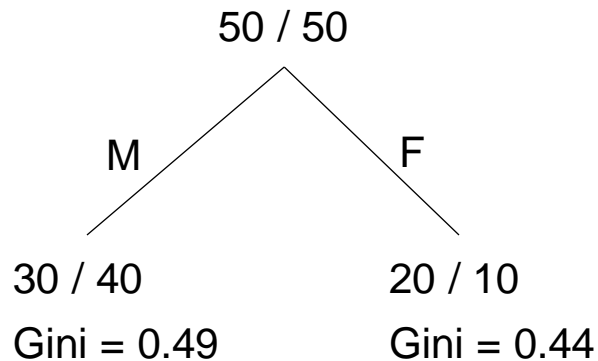
- For just two classes:



# Example: Gini Index

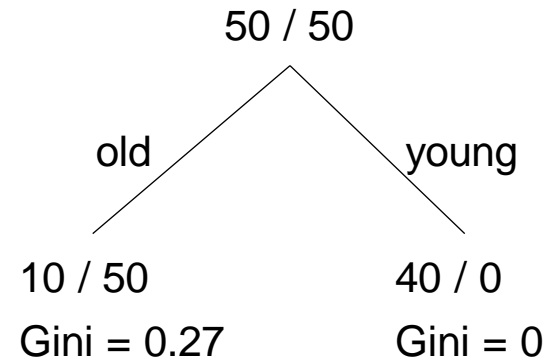
Side effects after treatment? 100 persons, 50 with and 50 without side effects:  
50 / 50 (No / Yes)

Split on sex



$$\begin{aligned} \text{Total Gini} &= 0.49 + 0.44 = \\ &= 0.93 \end{aligned}$$

Split on age



$$\begin{aligned} \text{Total Gini} &= 0.27 + 0 = \\ &= 0.27 \end{aligned}$$

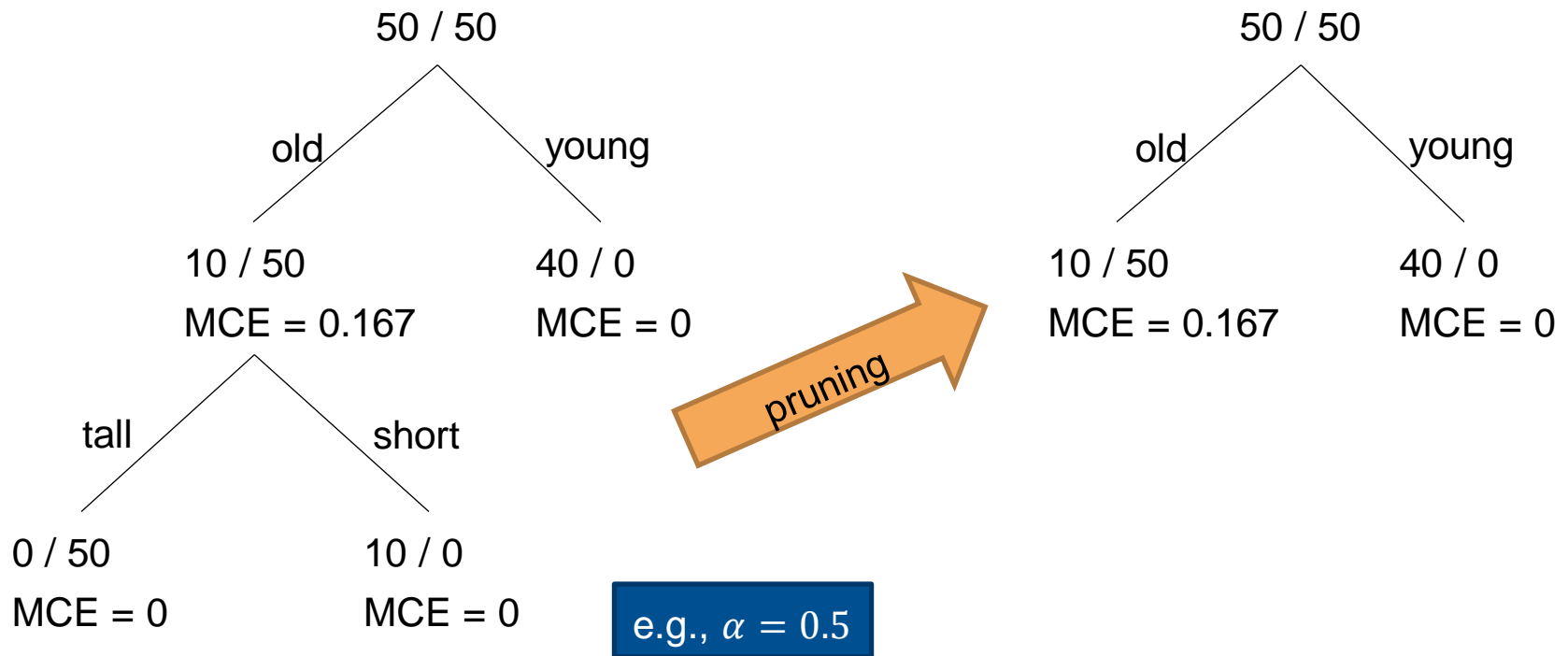
0.27 < 0.93, therefore:  
Choose split on age

# Classification Trees: Impurity Measures

- Usually:
  - Gini Index used for building
  - Misclassification error used for pruning

# Example: Pruning using Misclass. Error (MCE)

$$C_\alpha(T) = \sum_{m=1}^{|T|} N_m Q_m(T) + \alpha |T|$$



$$C_\alpha(T) = 50 * 0 + 10 * 0 + 40 * 0 + 0.5 * 3 = 1.5$$

$$C_\alpha(T) = 60 * 0.167 + 40 * 0 + 0.5 * 2 = 11.0$$

Smaller  $C_\alpha(T)$ , therefore don't prune

# Trees in R

- Function “rpart” (recursive partitioning) in package “rpart” together with “print”, “plot”, “text”
- ***Function “rpart” automatically prunes using optimal  $\alpha$  based on 10-fold CV***
- Functions “plotcp” and “printcp” for cost-complexity information
- Function “prune” for manual pruning

# Concepts to know

- Trees as recursive partitionings
- Concept of cost-complexity pruning
- Impurity measures



# R functions to know

- From package “rpart”: “rpart”, “print”, “plot”, “text”, “plotcp”, “printcp”, “prune”