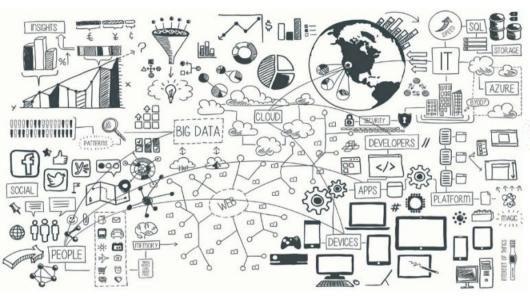
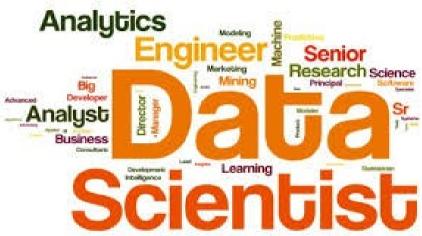
# **Data Mining (Minería de Datos)**

# **Decision Trees: Regression**





Sixto Herrera

Grupo de Meteorología Univ. de Cantabria – CSIC MACC / IFCA





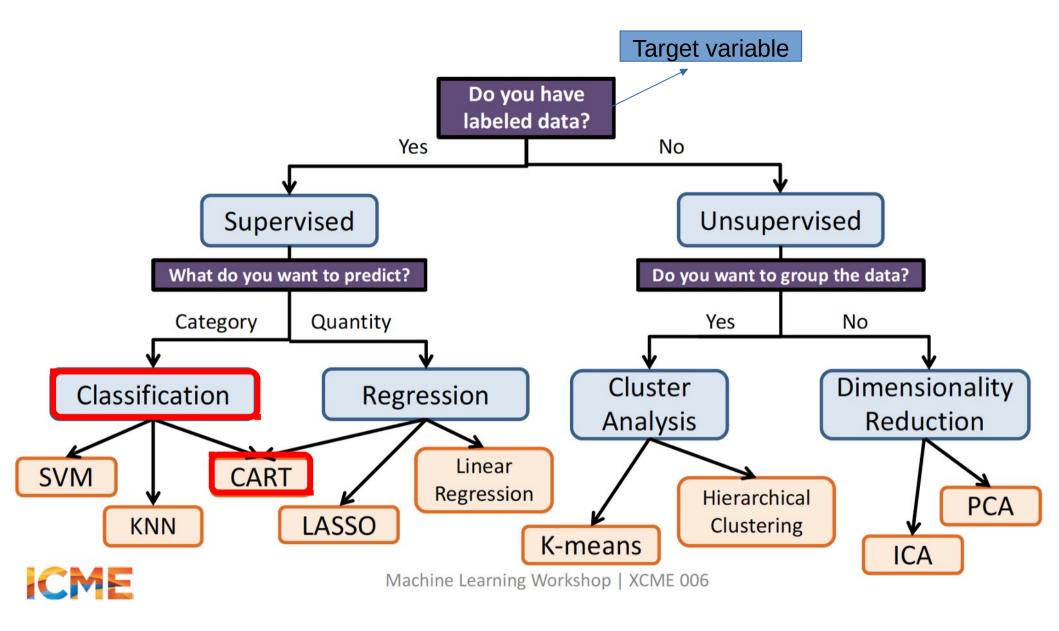


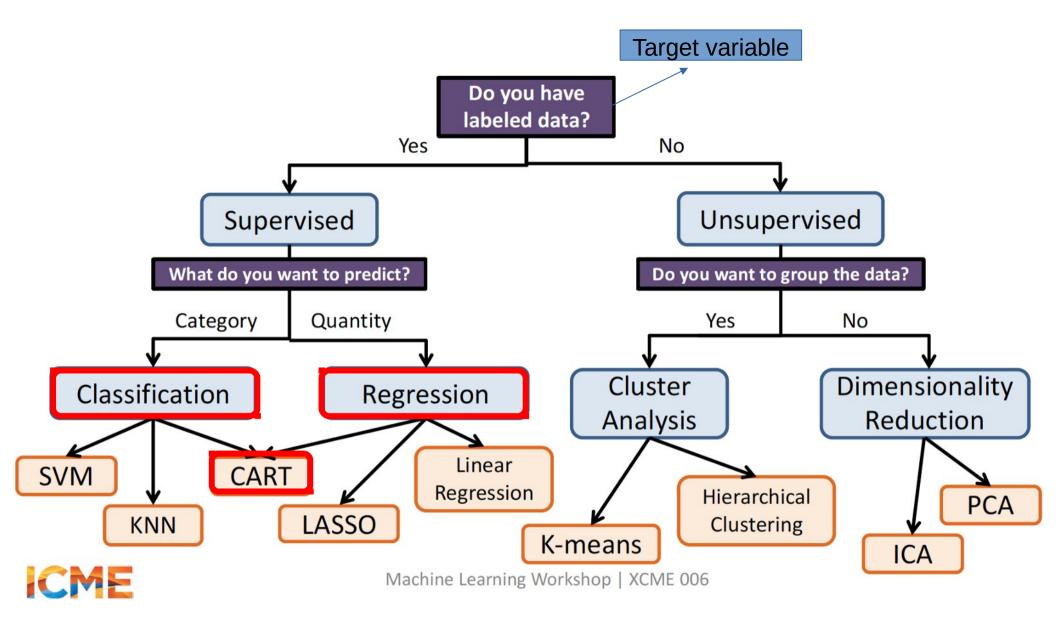
# **Trees Based Models**

Presentación, introducción y perspectiva histórica 31 Types of Machine Learning Nov Paradigmas, problemas canonicos y data challenges Reglas de asociación Machine Práctica: Reglas de asociación Learning Evaluación, sobrejuste y crossvalidacion 14 Práctica: Cross-validación 19 Árboles de clasificacion y decision Supervised Unsupervised Reinforcement Practica: Árboles de clasificación 26 T01. Datos discretos Task Driven Learn from Data Driven (Predict next value) (Identify Clusters) Mistakes 28 Técnicas de vecinos cercano (k-NN) Práctica: Vecinos cercanos Dic Reducción de dimensión no lineal Práctica: Reducción de dimensión no lineal 10 T02. Clasificación NOTA: Las líneas Árboles de clasificación y regresion (CART) de código de R Práctica: Árboles de clasificación y regresion (CART) 17 19 **Ensembles: Bagging and Boosting** en esta Ene Práctica Random Forests presentación se Práctica Gradient boosting muestran sobre T03. Prediccion Técnicas de agrupamiento un fondo gris 14 16 Practica: Técnicas de agrupamiento 21a Practica: El paquete CARET Master Universitario Oficial Data Science 21b Examen con el apoyo del CSIC

Oct

Aplazada (sesión de refuezo)





#### **Classification Trees**

#### Aim:

To classify a **categorical** target variable (R factor) based on a set of **categorical or continuous** predictors.

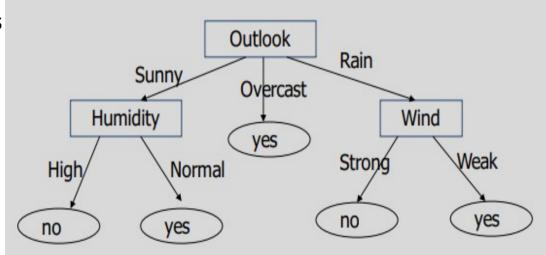
#### Structure:

- Each **node** corresponds to a test on an **attribute**
- Each **branch** corresponds to an **attribute value**
- Each leaf (terminal node) represents a final class
- Each **path** is a conjunction of attribute values

# **Key Points:**

- Due to their intuitive representation, they are **easy to assimilate** by humans
- They can be constructed **relatively fast** as compared to other methods
- In general, they provide as **good results** as other more complex methods

Day	Outlook	Temperature	Humidity	Wind	Play Tennis
1	Sunny	Hot	High	Weak	No
2	Sunny	Hot	High	Strong	No
3	Overcast	Hot	High	Weak	Yes
4	Rain	Mild	High	Weak	Yes
5	Rain	Cool	Normal	Weak	Yes
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7	Overcast	Cool	Normal	Strong	Yes
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11	Sunny	Mild	Normal	Strong	Yes
12	Overcast	Mild	High	Strong	Yes
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PlayTennis dataset: https://github.com/sjwhitworth/golearn/blob/master/examples/datasets/tennis.csv

# **Classification Trees**

# Aim:

To classify a **categorical** target variable (R factor) based on a set of **categorical or continuous** predictors.

#### **Structure:**

- Each **node** corresponds to a test on an **attribute**
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# **Key Points:**

- Due to their intuitive representation, they are **easy to assimilate** by humans
- They can be constructed **relatively fast** as compared to other methods
- In general, they provide as **good results** as other more complex methods

# Regression Trees

# Aim:

To predict a **continuous** target variable based on a set of **categorical or continuous** predictors.

## Structure:

- Each **node** corresponds to a test on an **attribute**
- Each **branch** corresponds to an **attribute value**
- Each leaf (terminal node) represents a final class
- Each **path** is a conjunction of attribute values

There are several algorithms to build up the tree. However, the idea of all of them is the same: evaluate attribute according to its **power of separation**.

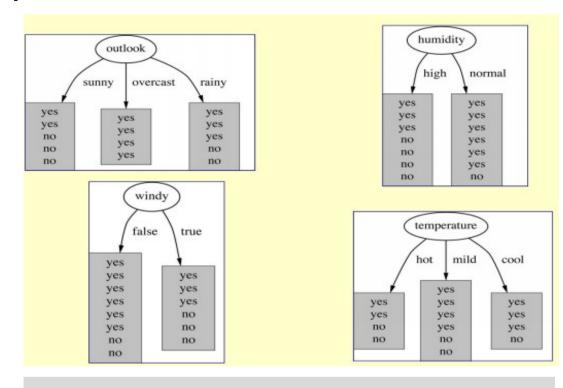
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#### Many variants for attribute selection:

- •from machine learning: **ID3** (Iterative Dichotomizer), **C4.5** and **C5.0** (Quinlan 86, 93)
- •from statistics: CART (Classification And Regression Trees) (Breiman et al. 84)
- •from pattern recognition: **CHAID** (**CH**i-squared **A**utomated **I**nteraction **D**etection) (Magidson 94)
- Their main difference is the criterion followed to perform the division of the node (splitting)

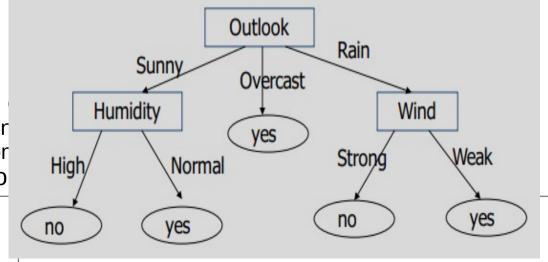
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Trees Based Models

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$$IG(X) = H(X) - H(X|Y) \begin{cases} H(X) = -\sum_{X} p(x) \log_2(p(x)) \\ H(X|Y) = -\sum_{X} \sum_{Y} p(x,y) \log_2(p(x|y)) \end{cases}$$

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$$GR = -\frac{IG}{Info} \qquad Info = -\sum_{i} \frac{|p_i|}{N} \log_2 \frac{|p_i|}{N}$$

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# What is the measure considered for Regression Trees?

Residual Sum of Squares (RSS) 
$$\longrightarrow \sum_{j=1}^{J} \sum_{i \in R_i} (y_i - \hat{y}_{R_j})^2$$
,

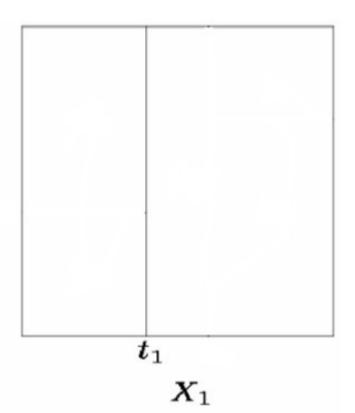




Generally we create the partitions by iteratively splitting one of the X variables into two regions.

First split on:  $X_1 = t_1$ 





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

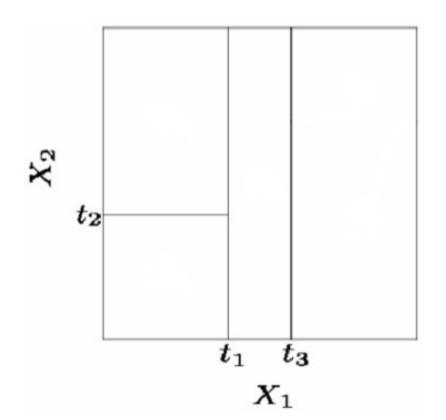
$$\sum_{i: x_i \in R_1(j,s)} (y_i - \hat{y}_{R_1})^2 + \sum_{i: x_i \in R_2(j,s)} (y_i - \hat{y}_{R_2})^2,$$

Generally we create the partitions by iteratively splitting one of the *X* variables into two regions.

First split on:  $X_1 = t_1$ 

If  $X_1 < t_1$  split on:  $X_2 = t_2$ 

If  $X_1 > = t_1$  split on:  $X_1 = t_3$ 



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

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

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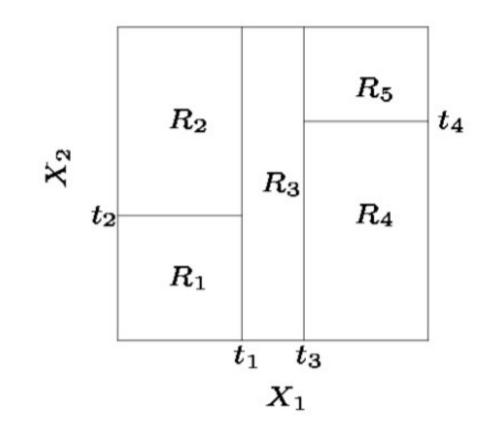
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If  $X_1 < t_1$  split on:  $X_2 = t_2$ 

If  $X_1 > = t_1$  split on:  $X_1 = t_2$ 

If  $X_1 > = t_3$  split on:  $X_2 = t_A$ 

until the tree correctly predicts the sample ...



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

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Generally we create the partitions by iteratively splitting one of the *X* variables into two regions.

First split on:  $X_1 = t_1$ 

If  $X_1 < t_1$  split on:  $X_2 = t_2$ 

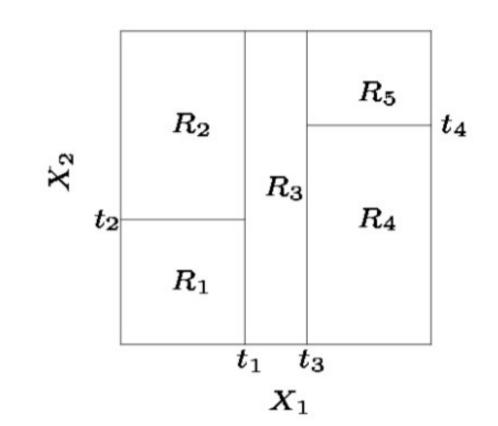
If  $X_1 > = t_1$  split on:  $X_1 = t_3$ 

If  $X_1 > = t_3$  split on:  $X_2 = t_4$ 

. . .

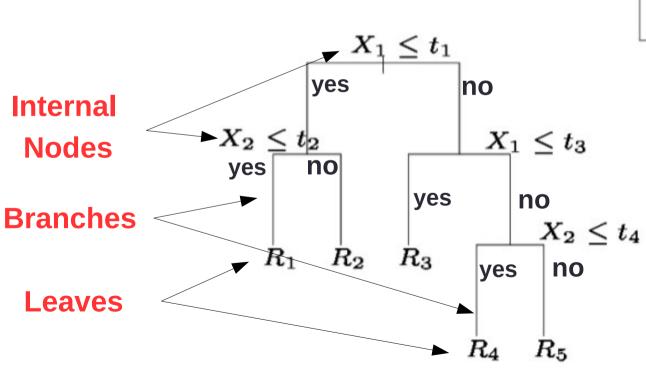
until the tree correctly predicts the sample or some stop criteria has been reached:

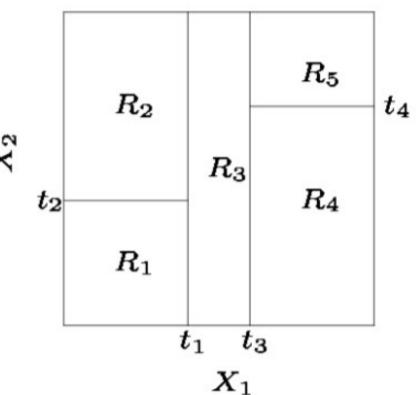
- Reduction cost too small vs Complexity
- Max. depth
- Purity
- Sub-sample too small



# Types of trees:

- Classification: discrete outputs
  - CLS, ID3, C4.5, ID4, ....
- Regression: continuous outputs
  - CART, M5, M5', ....







# **Housing Values in Suburbs of Boston**

## **Description:**

The Boston data frame has 506 rows and 14 columns.

#### **Format:**

This data frame contains the following columns:

**crim** - per capita crime rate by town.

**zn** - proportion of residential land zoned for lots over 25000 sq.ft.

indus - proportion of non-retail business acres per town.

**chas** - Charles River dummy variable (= 1 if tract bounds river; 0 otherwise).

**nox** - nitrogen oxides concentration (parts per 10 million).

**rm** - average number of rooms per dwelling.

age - proportion of owner-occupied units built prior to 1940.

**dis** - weighted mean of distances to five Boston employment centres.

rad - index of accessibility to radial highways.

tax - full-value property-tax rate per 10000 \$.

**ptratio** - pupil-teacher ratio by town.

**black** - 1000(Bk - 0.63)<sup>2</sup> where Bk is the proportion of blacks by town.

**Istat** - lower status of the population (percent).

medv - median value of owner-occupied homes in 1000s \$.





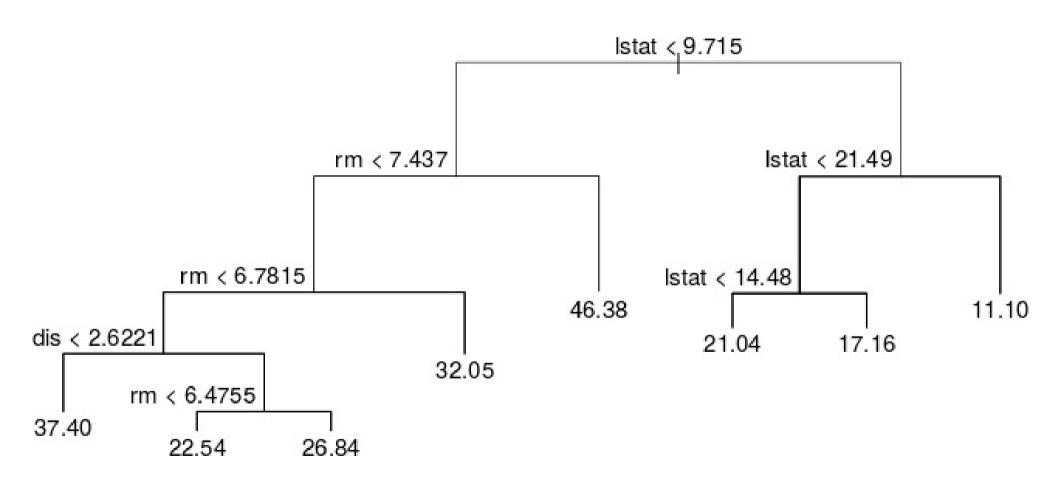
```
library(tree)
set.seed(1)
indTrain <- sample(1:nrow(Boston), nrow(Boston)/2) ## Splitting the sample
tree.boston <- tree(medv~., Boston, subset = indTrain) ## Adjusting the regression tree.
summary(tree.boston)
```

```
## Regression tree:
## tree(formula = medv ~ ., data = Boston, subset = train)
## Variables actually used in tree construction:
## [1] "lstat" "rm"
                      "dis"
## Number of terminal nodes: 8
## Residual mean deviance: 12.65 = 3099 / 245
## Distribution of residuals:
       Min. 1st Ou.
                         Median
                                            3rd Ou.
##
                                     Mean
                                                         Max.
## -14.10000 -2.04200 -0.05357 0.00000
                                            1.96000 12.60000
```



```
library(tree)
set.seed(1)
indTrain <- sample(1:nrow(Boston), nrow(Boston)/2) ## Splitting the sample
tree.boston <- tree(medv~., Boston, subset = indTrain) ## Adjusting the regression tree.
summary(tree.boston)

plot(tree.boston)
text(tree.boston, pretty = 0)</pre>
```



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Trees Based Models

**Regression Trees: Example** 

library(tree) set.seed(1) indTrain <- sample(1:nrow(Boston), nrow(Boston)/2) ## Splitting the sample tree.boston <- tree(medv~., Boston, subset = indTrain) ## Adjusting the regression tree. summary(tree.boston) ? tree.control

#### tree.control

#### Use:

tree.control(nobs, mincut =  $\frac{5}{10}$ , minsize =  $\frac{10}{10}$ , mindev =  $\frac{0.01}{10}$ )

#### **Arguments:**

**nobs** - the number of observations in the training set.

**mincut** - the minimum number of observations to include in either child node.

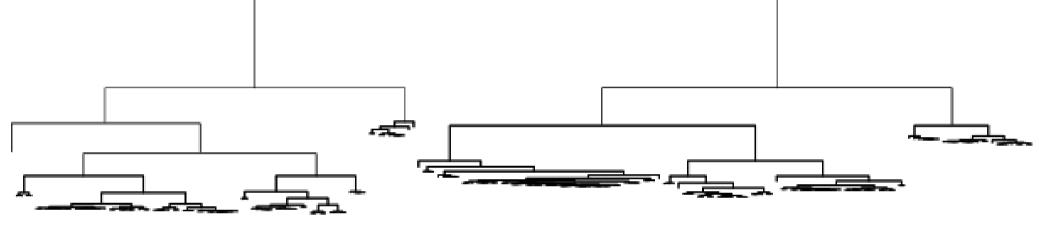
minsize - the smallest allowed node size.

**mindey** - the within-node deviance must be at least this times that of the root node for the node to be split.





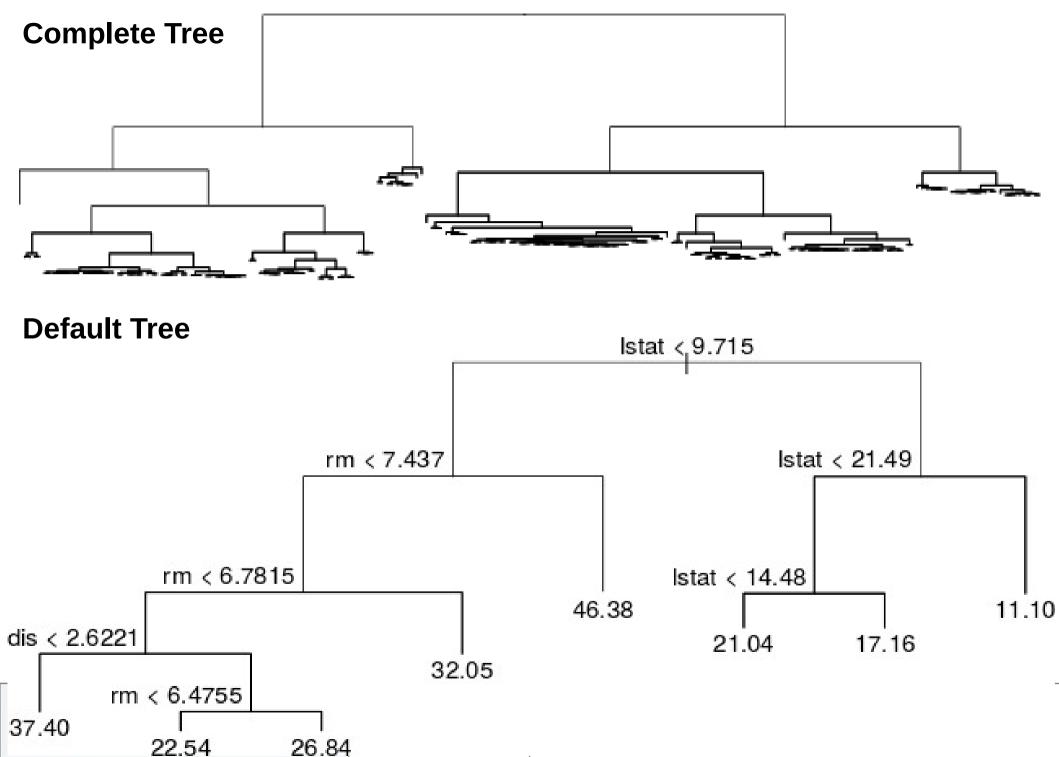
```
library(tree)
set.seed(1)
indTrain <- sample(1:nrow(Boston), nrow(Boston)/2) ## Splitting the sample
## Building the deepest tree.
deepTree.boston <- tree(medv~., Boston, subset = indTrain, split = "deviance", control =
tree.control(length(indTrain), mincut = 1, minsize = 2, mindev = 0))
plot(deepTree.boston)</pre>
```



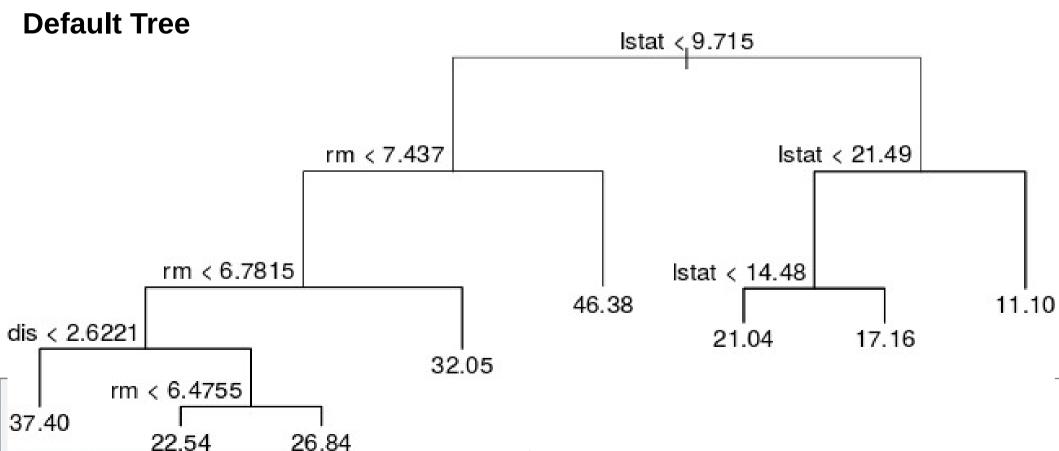




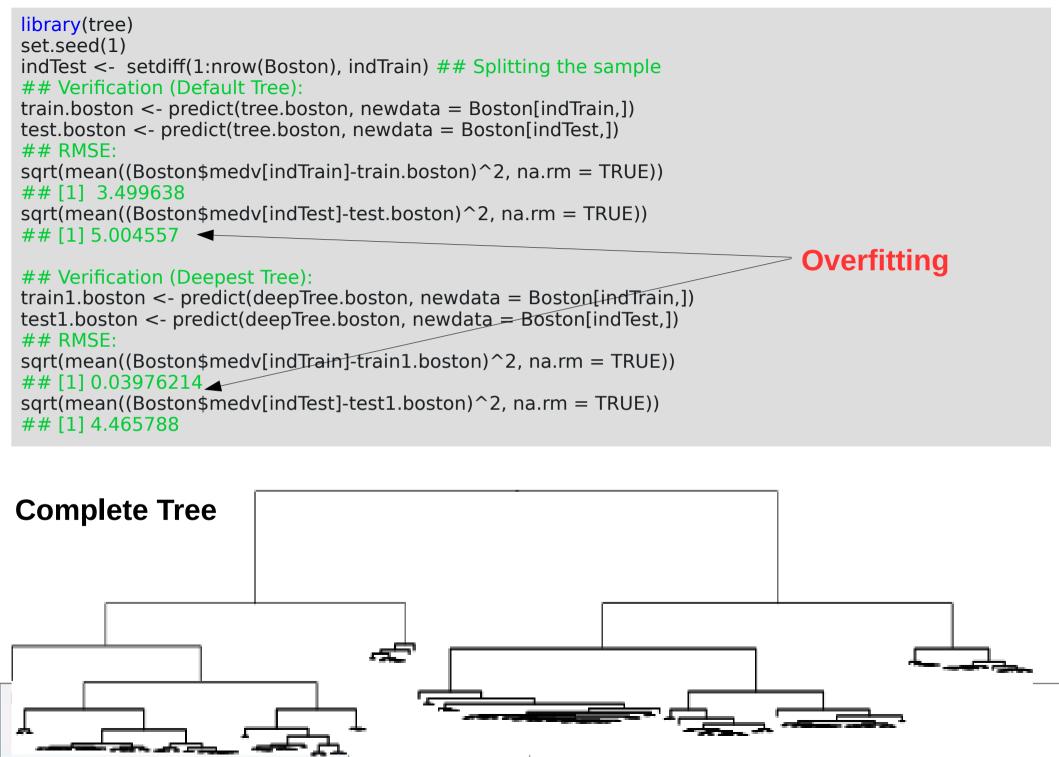




```
library(tree)
set.seed(1)
indTest <- setdiff(1:nrow(Boston), indTrain) ## Splitting the sample
## Verification (Default Tree):
train.boston <- predict(tree.boston, newdata = Boston[indTrain,])
test.boston <- predict(tree.boston, newdata = Boston[indTest,])
## RMSE:
sqrt(mean((Boston$medv[indTrain]-train.boston)^2, na.rm = TRUE))
## [1] 3.499638
sqrt(mean((Boston$medv[indTest]-test.boston)^2, na.rm = TRUE))
## [1] 5.004557</pre>
```

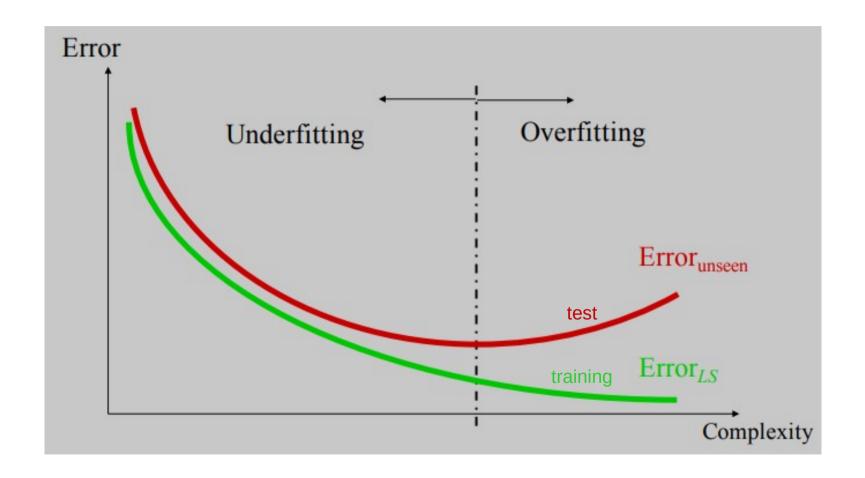


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test.boston <- predict(tree.boston, newdata = Boston[indTest,])
## RMSF:
sqrt(mean((Boston$medv[indTrain]-train.boston)^2, na.rm = TRUE))
## [1] 3.499638
sgrt(mean((Boston\$medv[indTest]-test.boston)^2, na.rm = TRUE))
## [1] 5.004557
## Verification (Deepest Tree):
train1.boston <- predict(deepTree.boston, newdata = Boston[indTrain,])
test1.boston <- predict(deepTree.boston, newdata = Boston[indTest,])
## RMSE:
sgrt(mean((Boston$medv[indTrain]-train1.boston)^2, na.rm = TRUE))
## [1] 0.03976214
sgrt(mean((Boston$medv[indTest]-test1.boston)^2, na.rm = TRUE))
## [1] 4.465788
Complete Tree
```



# **Overfitting**

A large tree (i.e. with many terminal nodes) may tend to overfit the training data, leading to poor performance in the test set. Generally, we can improve this behavior by pruning the tree, i.e., cutting off some of the terminal nodes.







#### How can we avoid overfitting?

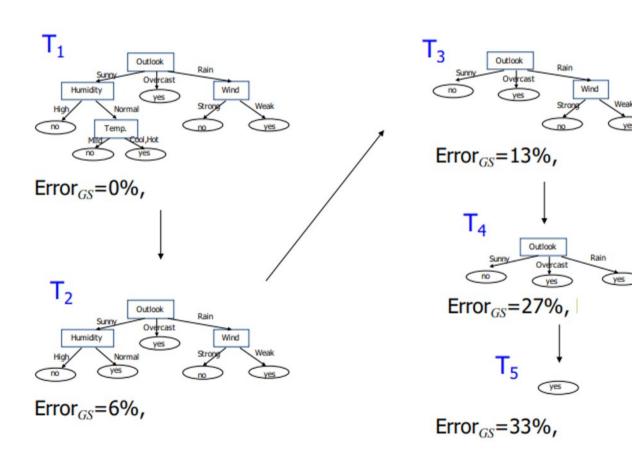
**Pre-pruning:** stop growing the tree before it reaches the point at which it perfectly classifies the learning sample. *Procedure:* 

Stop splitting a node if:

- The number of objects is too small
- The impurity is low enough
- This approach leads to small trees but can remove relevant splits

Post-pruning: allow the tree to overfit and then, one finalized, remove the less useful nodes. In general, this is preferred option. *Procedure:*Compute a sequence of trees
{T1, T2, ...} where T1 is the complete tree. T2 is obtained by removing from T1 the node that less increases the error. Sometimes, this process is guided based on some cost-complexity criterion (e.g. in medicine)

The question is: where to stop? In practice, it is usual to split the learning dataset into two subsets: a training sample for growing the tree and a test sample for evaluating its generalization error (e.g. hold-out cross-validation).









```
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set.seed(1)
indTrain <- sample(1:nrow(Boston), nrow(Boston)/2) ## Splitting the sample
tree.boston \leftarrow tree(medv\sim., Boston, subset = indTrain) ## Adjusting the regression tree.
summary(tree.boston)
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## Regression tree:
## tree(formula = medv ~ ., data = Boston, subset = train)
## Variables actually used in tree construction:
## [1] "lstat" "rm"
## Number of terminal nodes: 8
                                            Regression trees: CART
## Residual mean deviance: 12.65 = 3099 / 245
## Distribution of residuals:
                                                Output corresponds to
                                                                                         the
                      Median
      Min. 1st Ou.
                                Mean
## -14.10000 -2.04200 -0.05357
                              0.00000
                                      1.9
                                            response mean value.
```

Valid for any kind of attributes





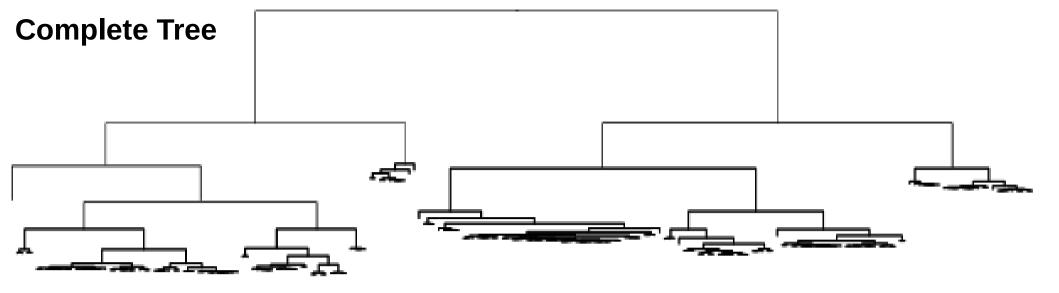
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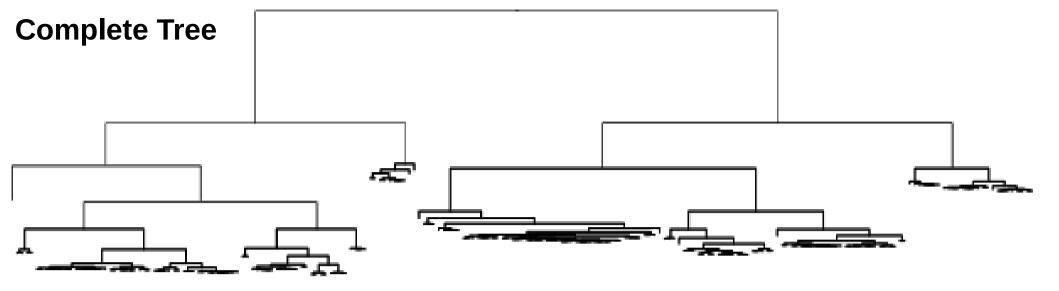




Given a subtree, we can estimate the test error rate

Choose the subtree with the lowest test error rate

Residual Sum of Squares (RSS) 
$$\longrightarrow \sum_{j=1}^{J} \sum_{i \in R_j} (y_i - \hat{y}_{R_j})^2$$
,



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**Computationally expensive iii** 

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,

# **Cost complexity Pruning** - Regularization

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

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

# **Cost complexity Pruning** - Regularization

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

Controls a trade-off between the subtree's complexity and its fit to the training data

Residual Sum of Squares (RSS) 
$$\longrightarrow \sum_{j=1}^{J} \sum_{i \in R_j} (y_i - \hat{y}_{R_j})^2$$
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For each value of the regularization parameter exists a subtree minimizing the expression above.

**Minimization problem** ← **Lagrange Multipliers** 

Residual Sum of Squares (RSS)

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

#### Pros:

- Trees are very easy to explain to people (probably even easier than linear regression)
- Trees can be plotted graphically, and are easily interpreted even by nonexpert
- Trees can easily handle qualitative predictors without the need to create dummy variables.
- They work fine on both classification and regression problems

#### Cons:

- Trees don't have the same prediction accuracy as some of the more complicated approaches that we examine in this course.
- Trees tend to overfit.





```
library(tree)
set.seed(1)
indTrain <- sample(1:nrow(Boston), nrow(Boston)/2) ## Splitting the sample
tree.boston <- tree(medv~., Boston, subset = indTrain) ## Adjusting the regression tree.
summary(tree.boston)
```

```
## Regression tree:
## tree(formula = medv ~ ., data = Boston, subset = train)
## Variables actually used in tree construction:
## [1] "lstat" "rm"
## Number of terminal nodes: 8
                                           Model trees: M5, M5', ...
## Residual mean deviance: 12.65 = 3099 / 245
## Distribution of residuals:
                                             Output corresponds to a linear
                      Median
      Min. 1st Ou.
                                Mean
## -14.10000 -2.04200 -0.05357
                             0.00000
                                      1.9
                                           regression model of the instances
```

 $\longrightarrow \sum_{j=1}^{J} \sum_{i \in R_j} (y_i - \hat{y}_{R_j})^2,$ **Residual Sum of Squares (RSS)** -



that reach the leaf.

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## -14.10000 -2.04200 -0.05357 0.00000
                                     1.9
                                          regression model of the instances
                                          that reach the leaf.
```

# M5 is included in the caret R-package (Cubist-implementation)

```
library(caret)
if (!require(Cubist)) install.packages("Cubist")
? models
autoTree <- train(form = medv~., data= Boston, subset = indTrain, method = "cubist")
summary(autoTree)
```

#### **Rules** ↔ **Decision Trees**







**Trees Based Models** 

**Regression Trees: Example** 

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- Trees can be plotted graphically, and are easily interpreted even by nonexpert
- Trees can easily handle qualitative predictors without the need to create dummy variables.
- They work fine on both classification and regression problems

#### Cons:

Trees don't have the same prediction accuracy as some of the more complicated approaches that we examine in this course.

By aggregating many decision trees, the predictive performance of trees can be substantially improved.



Bagging, Random Forest, Boosting





# Which model is better?

- If the relationship between the predictors and response is linear, then classical linear models such as linear regression would outperform regression trees
- On the other hand, if the relationship between the predictors is non-linear, then decision trees would outperform classical approaches
- Categorical variables can be used for trees.
- Trees are more easily interpretable than linear models.
- Visualization. The graphical representation of the trees is very useful.

$$f(X) = \beta_0 + \sum_{j=1}^{p} X_j \beta_j, \qquad f(X) = \sum_{m=1}^{M} c_m \cdot 1_{(X \in R_m)}$$

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- Categorical variables can be used for trees.
- Trees are more easily interpretables than linear models.
- Visualization. The graphical representation of the trees is very useful.
- Top row: the true decision boundary is linear
  - Left: linear model (good)
  - Right: decision tree
- Bottom row: the true decision boundary is non-linear
  - Left: linear model
  - Right: decision tree (good)

