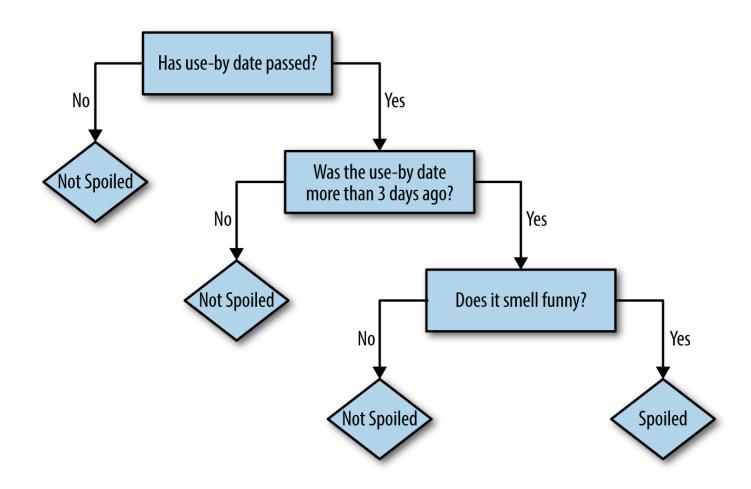


# **Decision Tree**

#### **Decision Trees**

- Set of rules that helps us make a decision
- set of if then examples

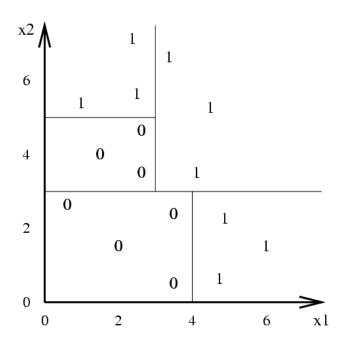


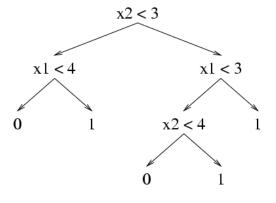


### We partition data into homogeneous rectangles

#### **Decision Tree Decision Boundaries**

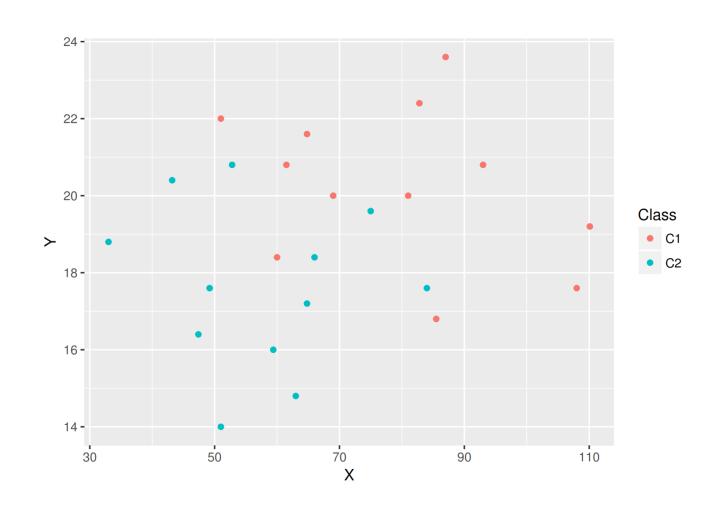
Decision trees divide the feature space into axis-parallel rectangles, and label each rectangle with one of the K classes.





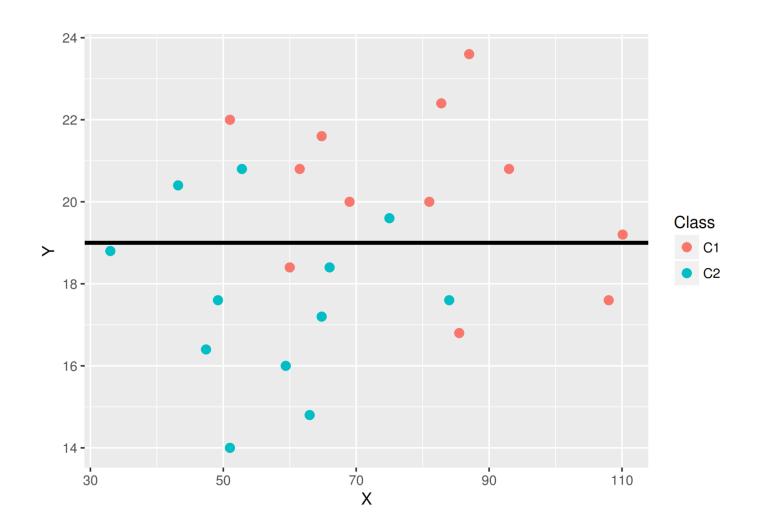


Two variables: X and Y and variable Class – needs to be predicted





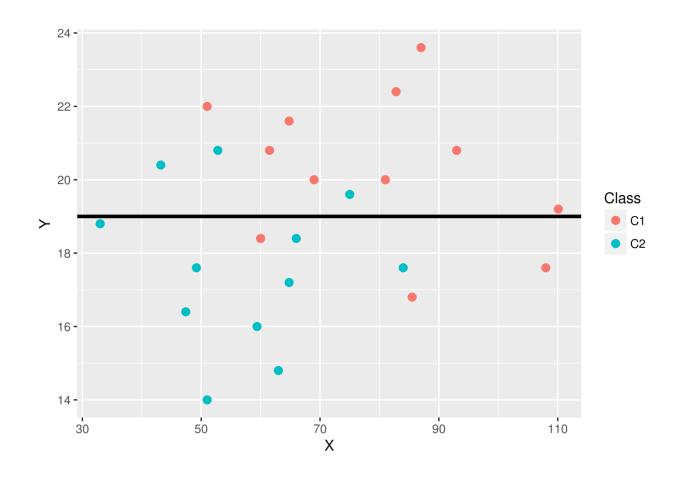
 $Y \ge 19$  and Y < 19





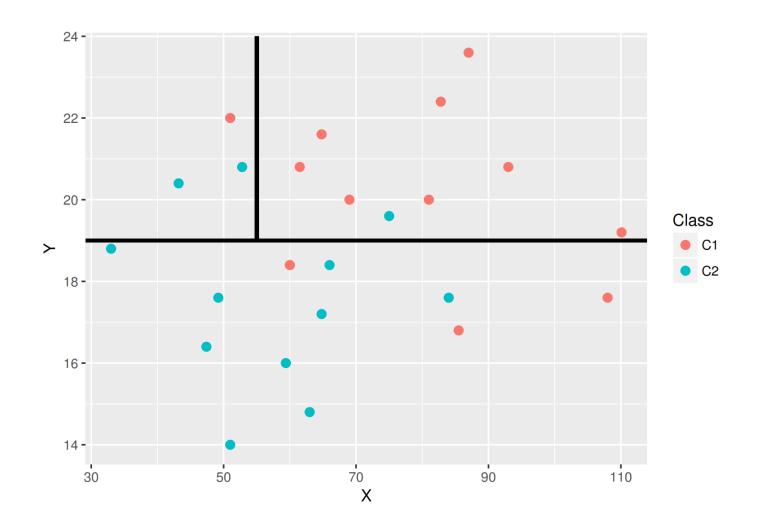
Example: Split N 1

Split N1:  $Y \ge 19$  and Y < 19. Lets look at the prevailing class at each rectangle IF  $Y \ge 19$  THEN C1 IF Y < 19 THEN C2



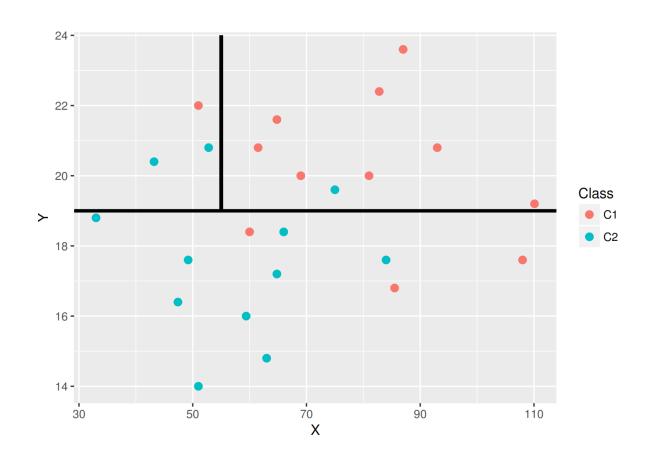


Split N2:  $X \le 55$  and X > 55





Split N2:  $X \le 55$  and X > 55





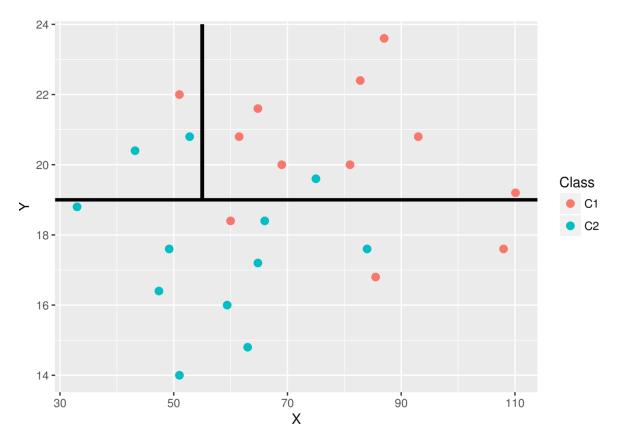
### Split N 2

#### Classification rules:

IF  $Y \le 19$  THEN C2

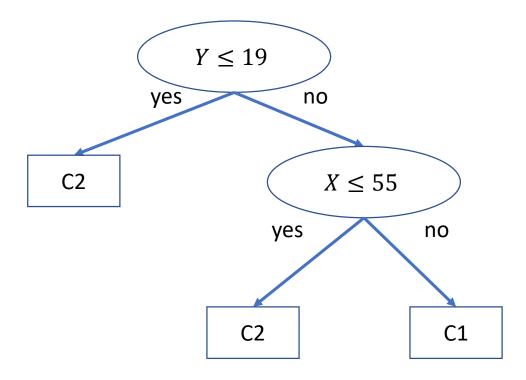
IF Y > 19 and  $X \le 55$  THEN C2

IF IF Y > 19 and X > 55 THEN C1





### The Decision Tree



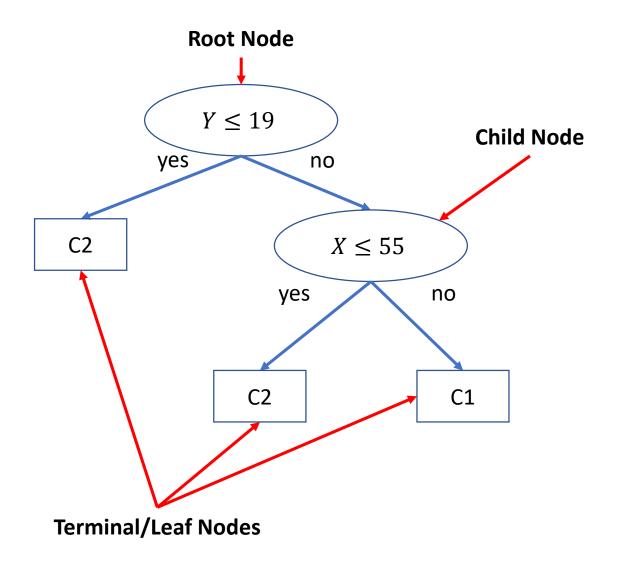


- Decision Tree has three types of nodes:
  - Root Node: top (or left-most) node with no incoming edges and zero or more outgoing edges.
  - Child or Internal Node: descendent node which has exactly one incoming edge and two or more outgoing edges.
  - Leaf Node: terminal node which has exactly one incoming edge and no outgoing edges.
- In Decision Tree, each leaf node is assigned a class label.
- The rules or branches are the unique path (edges) with a set of conditions (attribute) that divide the observations into smaller subset.

#### Read down tree to derive rule

E.g., If lot size < 19, and if income > 84.75, then class = "owner"







Recursive partitioning: Repeatedly split the records into two parts so as to achieve maximum homogeneity/purity within the new parts, or decrease the impurity.

**Pruning the tree:** Simplify the tree by pruning peripheral branches to avoid overfitting



## **Recursive Partitioning Steps**

- Pick one of the predictor variables,  $x_i$
- Pick a value of  $x_{i_j}$  say  $s_i$ , that divides the training data into two (not necessarily equal) portions
- Measure how "pure" or homogeneous each of the resulting portions are
  - "Pure" = containing records of mostly one class
- Algorithm tries different values of  $x_{i,j}$  and  $s_{i,j}$  to maximize purity in initial split
- After you get a "maximum purity" split, repeat the process for a second split, and so on



# What is impurity

Saying that the split is pure we mean that all records belong to the same class.

Two main measures of impurity

- Gini index
- Entropy

### Gini Index

Gini Index for rectangle A containing m records

$$I(A) = 1 - \sum_{k=1}^{m} p_k^2$$

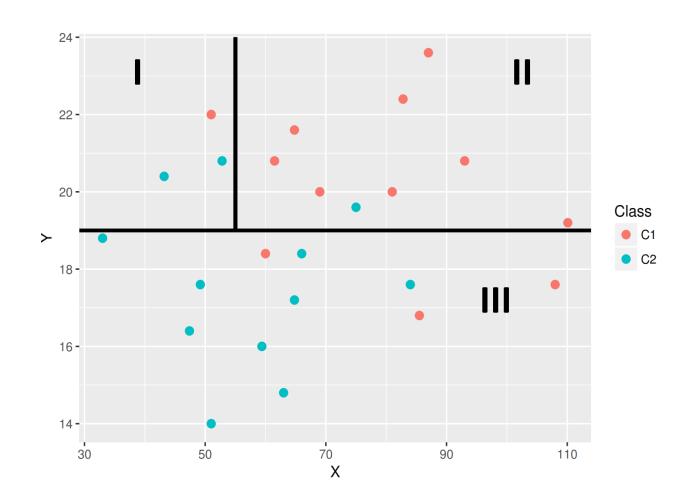
p = proportion of cases in rectangle A that belong
to class k

- I(A) = 0 when all cases belong to same class
- Max value when all classes are equally represented (= 0.50 in binary case)



# Gini Index

### Calculate Gini index for rectangles I, II and III



# Entropy

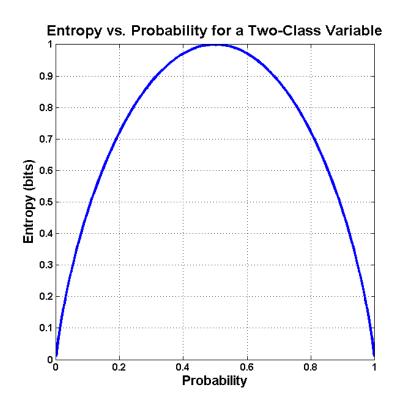
$$entropy(A) = -\sum_{k=1}^{m} p_k \log_2(p_k)$$

p = proportion of cases (out of m) in rectangle A that belong to class k

• Entropy ranges between 0 (most pure) and  $log_2(m)$  (equal representation of classes)



# Measures of impurity: Entropy

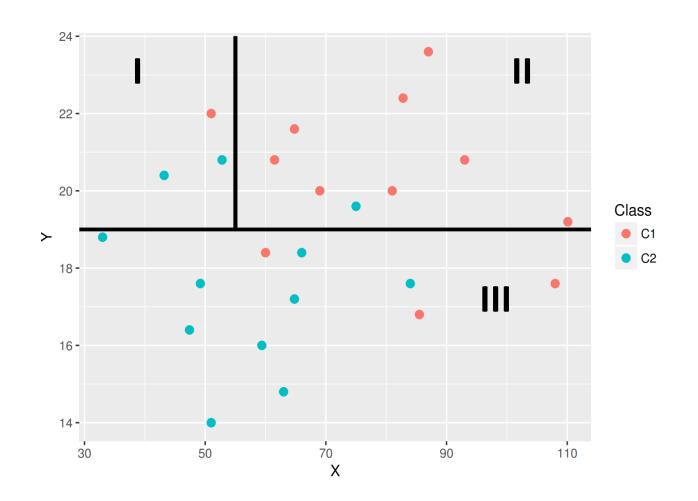


•Entropy ranges between 0 (most pure) and  $log_2(0.5)$  (equal representation of classes)



# Entropy

### Calculate entropy for rectangles I, II and III

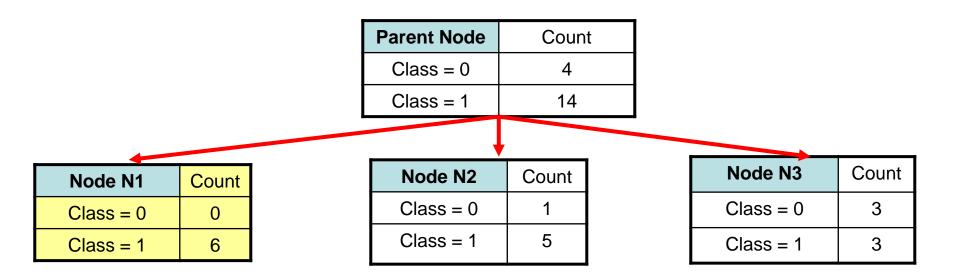


# Impurity and Recursive Partitioning

- Obtain overall impurity measure (weighted avg. of individual rectangles)
- At each successive stage, compare this measure across all possible splits in all variables
- Choose the split that reduces impurity the most
- Chosen split points become nodes on the tree



# Decision tree partitioning: Measuring impurity



Calculate Gini index and entropy for each node



# Decision tree partitioning: Measuring impurity

Parent Node	Count
Class = 0	4
Class = 1	14

Node N1	Count
Class = 0	0
Class = 1	6
Gini	0
Entropy	0

Node N2	Count
Class = 0	1
Class = 1	5
Gini	0.278
Entropy	0.650

Node N3	Count
Class = 0	3
Class = 1	3
Gini	0.5
Entropy	1



# Information gain

- To determine how well a test condition performs, we need to compare the degree of impurity of the parent node (before splitting) and the child node (after splitting).
- The larger the different, the better the test condition.
- The **gain**  $\Delta$ , is a criterion that can be used to determine the goodness of a split.

•  $\Delta = I(Parent) - I(Split)$ 



# Gain from splitting

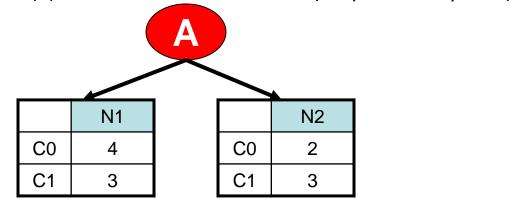
### Example:

Parent
C0 6
C1 6
Gini = 0.5

**Gini**:  $1 - (6/12)^2 - (6/12)^2$ 

= 0.5

Suppose there are two ways (A and B) to split the data into smaller subset.



 N1
 N2

 C0
 1

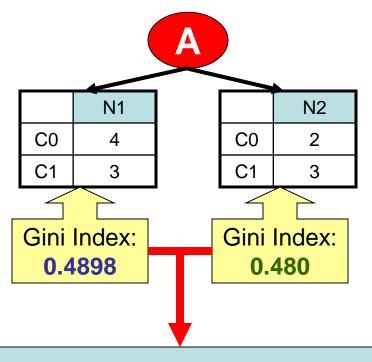
 C1
 4

 C1
 2

Which one is better?



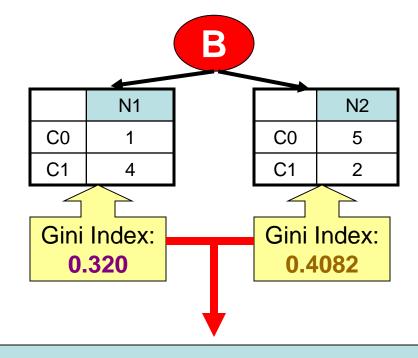
### Information Gain



### **Weighted Average of Gini Index:**

$$[(7/12) \times \mathbf{0.4898}] + [(5/12) \times \mathbf{0.480}]$$
  
= **0.486**

**Gain,** 
$$\Delta = 0.5 - 0.486 = 0.014$$



### **Weighted Average of Gini Index:**

$$[(5/12) \times 0.320] + [(7/12) \times 0.4082]$$
  
= 0.3715

**Gain**, 
$$\Delta = 0.5 - 0.3715 = 0.1285$$

Splitting by option B is better



# Doing in R



# we will use the following libraries

- rpart
- rpart.plot to plot the trees



```
weather<-read.csv("weather.csv")</pre>
str(weather)
## 'data.frame': 366 obs. of 21 variables:
    $ MinTemp
                        8 14 13.7 13.3 7.6 6.2 6.1 8.3 8.8 8.4 ...
##
                  : num
##
   $ MaxTemp
                         24.3 26.9 23.4 15.5 16.1 16.9 18.2 17 19.5 22.8 ...
                  : num
   $ Rainfall : num
##
                        0 3.6 3.6 39.8 2.8 0 0.2 0 0 16.2 ...
   $ Evaporation : num 3.4 4.4 5.8 7.2 5.6 5.8 4.2 5.6 4 5.4 ...
##
   $ Sunshine
##
                        6.3 9.7 3.3 9.1 10.6 8.2 8.4 4.6 4.1 7.7 ...
   $ WindGustDir : Factor w/ 16 levels "E", "ENE", "ESE", ...: 8 2 8 8 11 10 10 1 9 1 ...
##
   $ WindGustSpeed: int 30 39 85 54 50 44 43 41 48 31 ...
##
   $ WindDir9am : Factor w/ 16 levels "E", "ENE", "ESE", ...: 13 1 4 15 11 10 10 10 1 9 ...
##
   $ WindDir3pm : Factor w/ 16 levels "E", "ENE", "ESE", ...: 8 14 6 14 3 1 3 1 2 3 ...
##
   $ WindSpeed9am : int 6 4 6 30 20 20 19 11 19 7 ...
##
   $ WindSpeed3pm : int 20 17 6 24 28 24 26 24 17 6 ...
##
   $ Humidity9am
                 : int 68 80 82 62 68 70 63 65 70 82 ...
##
   $ Humidity3pm : int
                        29 36 69 56 49 57 47 57 48 32 ...
##
   $ Pressure9am : num
##
                         1020 1012 1010 1006 1018 ...
   $ Pressure3pm : num
                        1015 1008 1007 1007 1018 . . .
##
   $ Cloud9am
                  : int 7582774677...
##
   $ Cloud3pm : int 7 3 7 7 7 5 6 7 7 1 ...
##
    $ Temp9am
                         14.4 17.5 15.4 13.5 11.1 10.9 12.4 12.1 14.1 13.3 ...
##
                  : num
##
    $ Temp3pm
                         23.6 25.7 20.2 14.1 15.4 14.8 17.3 15.5 18.9 21.7 ...
                  : num
##
   $ RainToday
                  : Factor w/ 2 levels "No", "Yes": 1 2 2 2 2 1 1 1 1 2 ...
##
    $ RainTomorrow : Factor w/ 2 levels "No", "Yes": 2 2 2 2 1 1 1 1 2 1 ...
```



### Doing in R

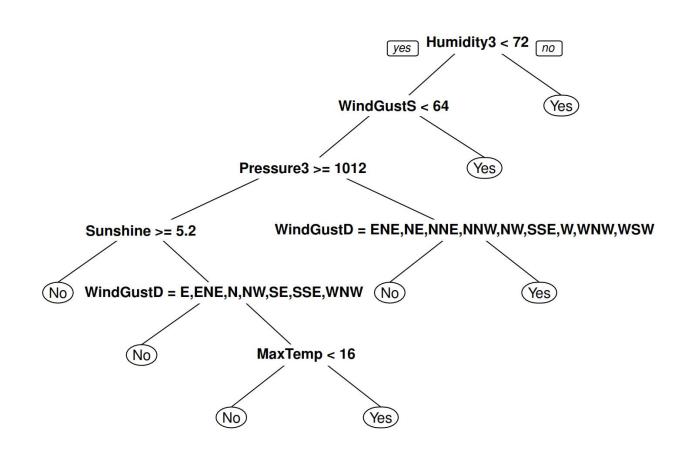
Specify the formula, with independent and dependent variables

```
library(rpart)
model<-rpart(RainTomorrow~., data=weather)</pre>
```



### Plot using rpart.plot library

```
library(rpart.plot)
prp(model)
```

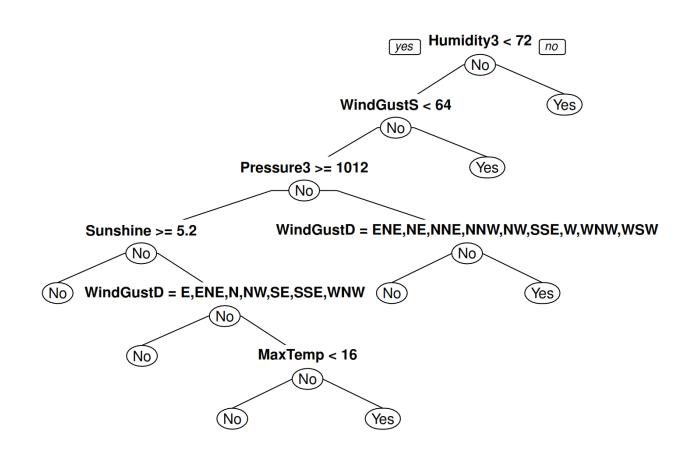




### Doing in R

You can change the type argument to get different layouts for the tree. Look for help for more info: ?rpart.plot::prp

```
prp(model, type=1)
```

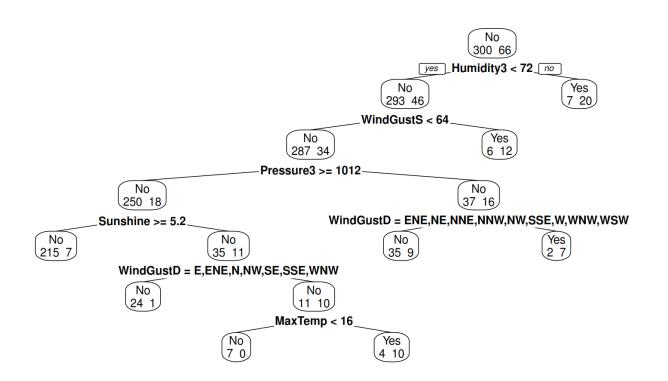




### You can add extra information to the plot using argument extra

prp(model, type=2, extra=1, main="Number of observations that fall in the node per class")

#### Number of observations that fall in the node per class

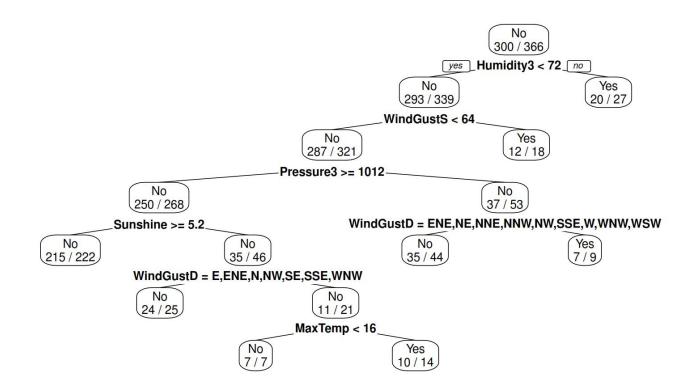




#### extra=2

prp(model, type=2, extra=2, main="Classification rate at the node")

#### Classification rate at the node

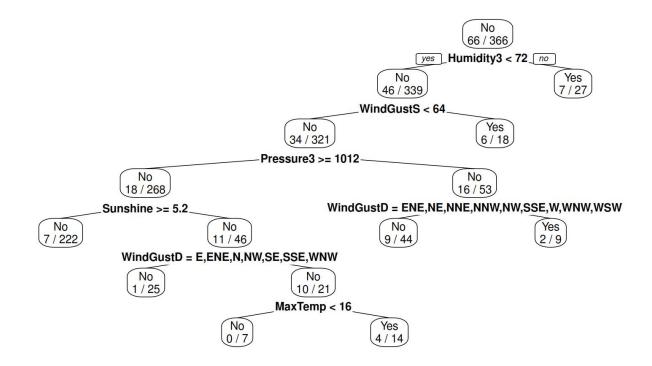




#### extra=3

```
prp(model, type=2, extra=3, main="Misclassification rate at the node")
```

#### Misclassification rate at the node

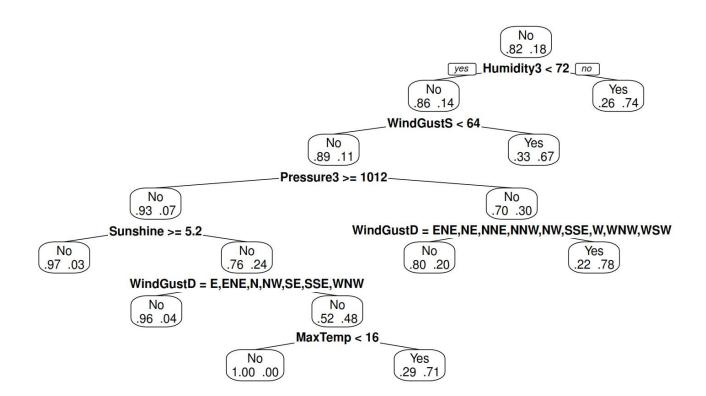




#### extra=4

prp(model, type=2, extra=4, main="Probabilities per class")

#### Probabilities per class





### Look at the decision rules using rattle package

```
library(rattle)
## Rattle: A free graphical interface for data mining with R.
## Version 4.1.0 Copyright (c) 2006-2015 Togaware Pty Ltd.
## Type 'rattle()' to shake, rattle, and roll your data.
## Attaching package: 'rattle'
## The following object is masked _by_ '.GlobalEnv':
##
##
       weather
asRules(model)
##
   Rule number: 19 [RainTomorrow=Yes cover=9 (2%) prob=0.78]
      Humidity3pm< 71.5
     WindGustSpeed< 64
##
      Pressure3pm< 1012
##
      WindGustDir=E,ESE,N,S,SW
##
    Rule number: 3 [RainTomorrow=Yes cover=27 (7%) prob=0.74]
##
      Humidity3pm>=71.5
##
    Rule number: 71 [RainTomorrow=Yes cover=14 (4%) prob=0.71]
##
      Humidity3pm< 71.5
      WindGustSpeed< 64
      Pressure3pm>=1012
      Sunshine < 5.15
      WindGustDir=ESE, NE, NNE, NNW, S
##
      MaxTemp>=15.8
##
   Rule number: 5 [RainTomorrow=Yes cover=18 (5%) prob=0.67]
##
      Humidity3pm< 71.5
      WindGustSpeed>=64
##
```



### Rules

- Ignore the rule number
- The predicted class is Yes, it covers 2% of the data, overall 9 cases in the terminal node, probability of Yes is 0.78

```
Rule number: 19 [RainTomorrow=Yes cover=9 (2%) prob=0.78]
Humidity3pm< 71.5
WindGustSpeed< 64
Pressure3pm< 1012
WindGustDir=E,ESE,N,S,SW
```



- The terminal node predicts No,
- covers 7% of the data (25 cases)
- prob=0.04, probability of **Yes** is 0.04, for **No** is 0.96

```
Rule number: 34 [RainTomorrow=No cover=25 (7%) prob=0.04]
Humidity3pm< 71.5
WindGustSpeed< 64
Pressure3pm>=1012
Sunshine< 5.15
WindGustDir=E,ENE,N,NW,SE,SSE,WNW</pre>
```



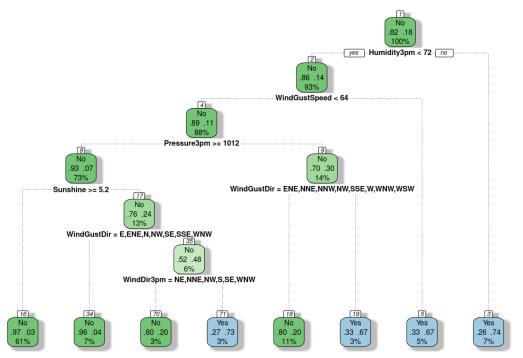
### Controlling the tree

- by default Gini coefficient is the impurity measure
- Tree is pruned using Complexity parameter
- Other parameters to control tree growth
  - minsplit: the minimum number of observations that must exist in a node in order for a split to be attempted
  - minbucket: the minmum number of observations in any terminal node,
  - look for more at help: ??rpart.control



### fancyRpartPlot from library rattle

```
fancyRpartPlot(model)
```





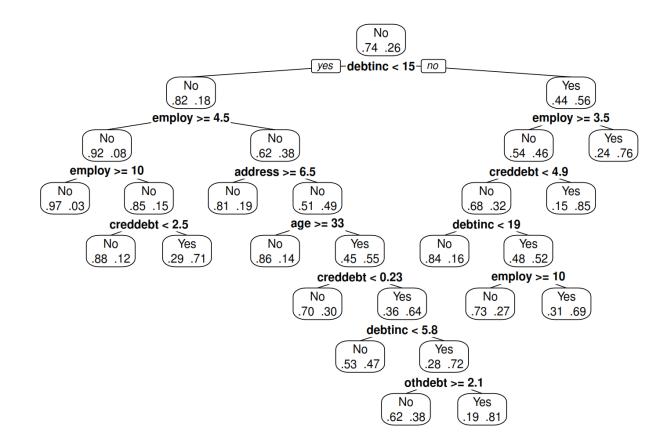


```
credit<-read.csv("credit.csv")
library(caret)

## Loading required package: lattice
index<-createDataPartition(credit$default, p=0.8, list=F)
Train<-credit[index,]
Test<-credit[-index,]</pre>
```



```
model_c<-rpart(default~., data=Train)
prp(model_c, type=2, extra=4)</pre>
```





```
pred_class<-predict(model_c, Test, type="class")</pre>
pred_class[1:20]
    10
       15
           16
               25
                  36 38
                           40
                               43 44
                                      48
                                          53
                                              62
                                                  66
                                                                  82
                                                                     84
## Yes
       No No Yes Yes No No No No
                                      No
                                          No
                                             No Yes
                                                     No No
                                                             No
                                                                 No
                                                                     No
   86
       99
   No
       No
##
## Levels: No Yes
```



```
confusionMatrix(pred_class, Test$default, positive="Yes")
## Confusion Matrix and Statistics
##
##
             Reference
## Prediction No Yes
##
         No 93 15
         Yes 10 21
##
##
##
                  Accuracy : 0.8201
                    95% CI: (0.7461, 0.8801)
##
       No Information Rate: 0.741
##
##
      P-Value [Acc > NIR] : 0.01823
##
##
                     Kappa : 0.5093
    Mcnemar's Test P-Value: 0.42371
##
##
##
               Sensitivity: 0.5833
               Specificity: 0.9029
##
            Pos Pred Value: 0.6774
##
##
            Neg Pred Value: 0.8611
##
                Prevalence: 0.2590
            Detection Rate: 0.1511
##
      Detection Prevalence: 0.2230
##
##
         Balanced Accuracy: 0.7431
##
##
          'Positive' Class : Yes
##
```

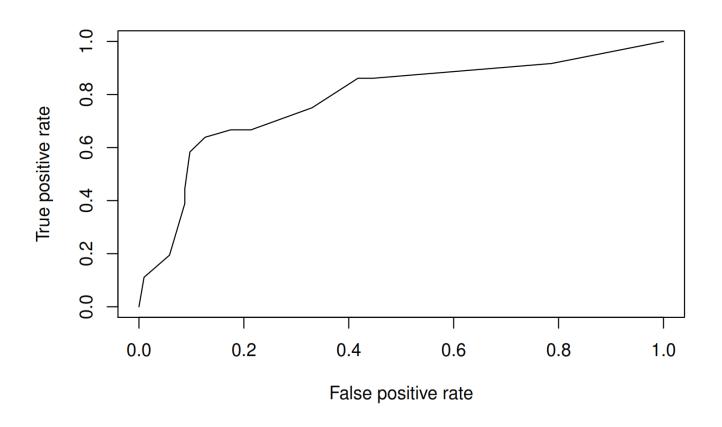


```
pred_prob<-predict(model_c, Test)
pred_prob[1:10,]

## No Yes
## 10 0.2439024 0.75609756
## 15 0.9736842 0.02631579
## 16 0.8823529 0.11764706
## 25 0.2439024 0.75609756
## 36 0.1500000 0.85000000
## 38 0.9736842 0.02631579
## 40 0.9736842 0.02631579
## 43 0.8823529 0.11764706
## 44 0.9736842 0.02631579
## 48 0.8571429 0.14285714</pre>
```



```
P_Test <- prediction(pred_prob[,2], Test$default)
perf <- performance(P_Test,"tpr","fpr")
plot(perf)</pre>
```





```
performance(P_Test, "auc")@y.values
## [[1]]
## [1] 0.7815534
```

