Decision Tree

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 ${\bf Importing\ libraries}$

library(party)

```
## Loading required package: grid
## Loading required package: mvtnorm
## Loading required package: modeltools
## Loading required package: stats4
## Loading required package: strucchange
## Loading required package: zoo
## Attaching package: 'zoo'
## The following objects are masked from 'package:base':
##
##
       as.Date, as.Date.numeric
## Loading required package: sandwich
library(plyr)
## Attaching package: 'plyr'
## The following object is masked from 'package:modeltools':
##
##
       empty
library(readr)
```

Importing dataset

```
dataset <- read.csv("D:/Internship/Task 3/iris.csv")
#converting Species from character to factor
dataset$Species=as.factor(dataset$Species)
head(dataset)</pre>
```

```
{\tt Id SepalLengthCm SepalWidthCm PetalLengthCm PetalWidthCm}\\
##
                                                                    Species
## 1
                  5.1
                               3.5
                                              1.4
                                                           0.2 Iris-setosa
## 2 2
                               3.0
                  4.9
                                              1.4
                                                           0.2 Iris-setosa
## 3 3
                  4.7
                               3.2
                                              1.3
                                                           0.2 Iris-setosa
## 4 4
                  4.6
                               3.1
                                              1.5
                                                           0.2 Iris-setosa
                  5.0
                               3.6
                                                           0.2 Iris-setosa
## 5 5
                                              1.4
## 6 6
                  5.4
                               3.9
                                              1.7
                                                           0.4 Iris-setosa
```

```
#Removing Id column
dataset <- dataset[-1]
head(dataset)</pre>
```

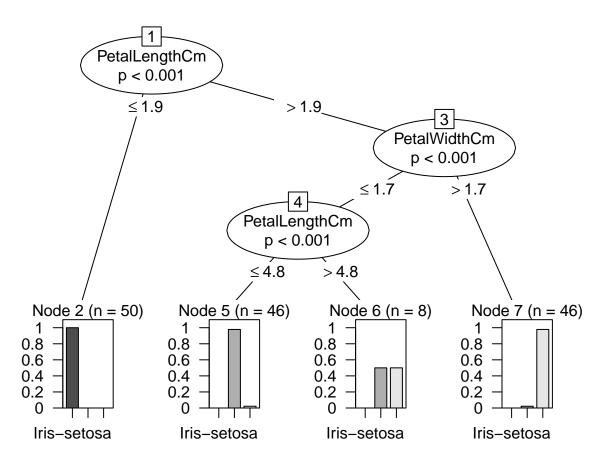
```
{\tt SepalLengthCm\ SepalWidthCm\ PetalLengthCm\ PetalWidthCm}
##
                                                                  Species
## 1
               5.1
                             3.5
                                            1.4
                                                         0.2 Iris-setosa
## 2
               4.9
                             3.0
                                            1.4
                                                         0.2 Iris-setosa
## 3
               4.7
                             3.2
                                            1.3
                                                         0.2 Iris-setosa
                                                         0.2 Iris-setosa
## 4
               4.6
                             3.1
                                            1.5
## 5
               5.0
                             3.6
                                            1.4
                                                         0.2 Iris-setosa
## 6
               5.4
                             3.9
                                            1.7
                                                         0.4 Iris-setosa
```

Defining The Decision Tree Algorithm

```
model <- ctree(Species ~., data = dataset)</pre>
```

Visualizing Decision Tree

```
plot(model)
```



Prediction

```
table(predict(model, dataset[,1:4]), dataset$Species)
```

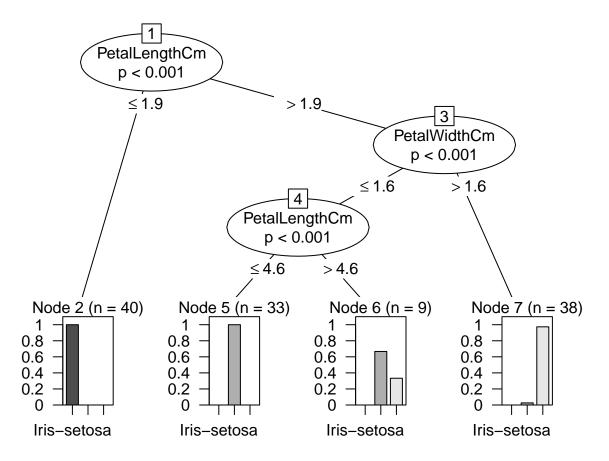
```
##
##
                      Iris-setosa Iris-versicolor Iris-virginica
##
     Iris-setosa
                                50
                                 0
                                                 49
##
     Iris-versicolor
                                                                  5
                                 0
                                                                 45
##
     Iris-virginica
                                                  1
```

Decision Tree Classifier: 70:30 Train & Test samples

```
library("caTools")
set.seed(100)
splitm <-sample.split(dataset$Species, SplitRatio = 0.8)
train=subset(dataset,splitm==TRUE)
test=subset(dataset,splitm==FALSE)
model1 <- ctree(Species ~ .,data=train)</pre>
```

Visualising

```
plot(model1)
```



Prediction

table(predict(model1,test[,1:4]),test\$Species)

```
##
##
                      Iris-setosa Iris-versicolor Iris-virginica
##
     Iris-setosa
                               10
                                0
                                                 9
##
     Iris-versicolor
                                                                 1
##
     Iris-virginica
                                0
                                                 1
                                                                 9
```