HW 04

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```
# Company
             : Stevens
# Project
             : HW_04
# Purpose
             : Perform Naïve Bayes Classification
# First Name : Preet
# Last Name : Dabhi
# Id
                  : 10459151
# Date
             : 03/6/2022
## Delete all the objects from your R- environment.
rm(list=ls())
#Import package 'e1071' for Naive Bayes Classifier and class package
#install.packages("e1071")
#install.packages(class)
library(e1071)
library(class)
dataFrame <- read.csv("D:/SEM 3/CS 513/HW_02/breast-cancer-wisconsin.csv", header=TRUE, sep=",")
 summary(dataFrame)
##
       Sample
                          F1
                                          F2
                                                          F3
                     Min. : 1.000
                                    Min. : 1.000
                                                         : 1.000
##
   Min. :
             61634
                                                    Min.
   1st Qu.: 870688
                     1st Qu.: 2.000
                                     1st Qu.: 1.000
                                                    1st Qu.: 1.000
   Median : 1171710
                     Median : 4.000
                                     Median : 1.000
                                                    Median : 1.000
  Mean
         : 1071704
                     Mean
                          : 4.418
                                    Mean
                                          : 3.134
                                                    Mean
                                                          : 3.207
   3rd Qu.: 1238298
                     3rd Qu.: 6.000
                                     3rd Qu.: 5.000
                                                    3rd Qu.: 5.000
##
         :13454352
                                                           :10.000
##
   Max.
                     Max.
                           :10.000
                                    Max.
                                           :10.000
                                                    Max.
##
         F4
                                       F6
                        F5
                                                          F7
##
  Min. : 1.000
                   Min. : 1.000
                                   Length:699
                                                    Min. : 1.000
   1st Qu.: 1.000
                   1st Qu.: 2.000
                                                    1st Qu.: 2.000
##
                                   Class :character
## Median : 1.000
                   Median : 2.000
                                   Mode :character
                                                    Median : 3.000
## Mean
        : 2.807
                   Mean : 3.216
                                                    Mean
                                                         : 3.438
##
  3rd Qu.: 4.000
                   3rd Qu.: 4.000
                                                    3rd Qu.: 5.000
##
  Max.
         :10.000
                   Max.
                         :10.000
                                                    Max. :10.000
##
         F8
                        F9
                                      Class
## Min. : 1.000
                   Min. : 1.000
                                   Min.
                                         :2.00
## 1st Qu.: 1.000
                   1st Qu.: 1.000
                                   1st Qu.:2.00
```

Median:2.00

Median : 1.000

Median : 1.000

```
## Mean : 2.867 Mean : 1.589 Mean :2.69
## 3rd Qu.: 4.000 3rd Qu.: 1.000 3rd Qu.:4.00
## Max. :10.000 Max. :10.000 Max. :4.00
#Here we can see that by running summary on the dataframe F6 column there are some missing values in it
n <- as.numeric(as.character(dataFrame$F6))</pre>
## Warning: NAs introduced by coercion
dataFrame$F6 <- n
#Remove the rows with missing values
dataFrame <- na.omit(dataFrame)</pre>
View(dataFrame)
#Replacing class column 2 and 4 with Benign and Malignant
dataFrame$Class<- factor(dataFrame$Class , levels = c("2","4") , labels = c("Benign", "Malignant"))
is.factor(dataFrame$Class)
## [1] TRUE
newData<- dataFrame[2:11]
View(newData)
#70% of the sample size
sample_size <- floor(0.70 * nrow(newData))</pre>
#Set the seed to make your partition reproducible
set.seed(123)
trainData <- sample(seq_len(nrow(newData)), size = sample_size)</pre>
#Loading 70% Breast cancer record in training dataset
training <- newData[trainData, ]</pre>
#Loading 30% Breast cancer in test dataset
test <- newData[-trainData, ]</pre>
#Implementing NaiveBayes
naive_bayes_model<- naiveBayes(Class ~ ., data = training)</pre>
#Predicting target class for the Validation set
naive_bayes_predict <- predict(naive_bayes_model, test)</pre>
conf_matrix <- table(predict_nb=naive_bayes_predict,class=test$Class)</pre>
print(conf_matrix)
##
              class
## predict_nb Benign Malignant
## Benign
                 132
```

Malignant

7

```
#Output of Naive Bayes Classifier
accuracy <- function(x){sum(diag(x)/(sum(rowSums(x)))) * 100}
accuracy(conf_matrix)</pre>
```

[1] 96.09756