# Laboratory Practice I

## Data Analytics

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## Practical 2

*Download Pima Indians Diabetes dataset. Use Naive Bayes‟ Algorithm for classification*

- Load the data from CSV file and split it into training and test datasets.
- Summarize the properties in the training dataset so that we can calculate probabilities and make predictions.
- *Classify samples from a test dataset and a summarized training dataset.*

Dataset Link - https://www.kaggle.com/uciml/pima-indians-diabetes-database

- ***Installing Libraries and Importing Data Set***

```
#Installing necessary Libraries
> install.packages('e1071')
> install.packages('caTools')

#Checking that the libraries are successfully installed
> library(caTools)
> library(e1071)

#Importing The Dataset
> mydata <- read.csv("~/Documents/BE/LP1/diabetes.csv")
>   View(mydata)
```

- ***Spiting the Dataset into training and testing Data***

```
> temp_field<-sample.split(mydata,SplitRatio=0.7)
> #70% will b in training
> train<-subset(mydata, temp_field==TRUE)
> #30% will be in testing
> test<-subset(mydata, temp_field == FALSE)

#Checking the
> head(train)
```

```
  Pregnancies Glucose BloodPressure SkinThickness Insulin  BMI DiabetesPedigreeFunction Age
Outcome
3           8     183           64           0       0 23.3                  0.672  32       1
4           1      89           66          23      94 28.1                  0.167  21       0
5           0     137           40          35     168 43.1                  2.288  33       1
6           5     116           74           0       0 25.6                  0.201  30       0
8          10     115            0           0       0 35.3                  0.134  29       0
9           2     197           70          45     543 30.5                  0.158  53       1
```

```
> head(test)
  Pregnancies Glucose BloodPressure SkinThickness Insulin  BMI DiabetesPedigreeFunction Age
Outcome
1           6     148           72          35       0 33.6                  0.627  50       1
2           1      85           66          29       0 26.6                  0.351  31       0
7           3      78           50          32      88 31.0                  0.248  26       1
10          8     125           96           0       0  0.0                  0.232  54       1
11          4     110           92           0       0 37.6                  0.191  30       0
16          7     100            0           0       0 30.0                  0.484  32       1
```

- ***> #Using Naive Bayes Algorithm, training the train Data Set***

```
> my_model<-naiveBayes(as.factor(train$Outcome)~.,train)
> my_model

Naive Bayes Classifier for Discrete Predictors

Call:
naiveBayes.default(x = X, y = Y, laplace = laplace)

A-priori probabilities:
Y
       0        1
0.6497065 0.3502935

Conditional probabilities:
  Pregnancies
Y       [,1]     [,2]
  0 3.253012 2.975604
  1 4.636872 3.662286

  Glucose
Y       [,1]     [,2]
  0 109.1928 26.20657
  1 142.4916 33.87259
```

```
  BloodPressure
Y       [,1]     [,2]
  0 67.91265 18.21095
  1 70.83799 21.18993

  SkinThickness
Y       [,1]     [,2]
  0 19.29819 15.03807
  1 22.16201 18.07387

  Insulin
Y       [,1]      [,2]
  0  65.10542  98.29565
  1 100.30168 142.80693

  BMI
Y       [,1]     [,2]
  0 30.44277 7.229345
  1 34.88994 6.879959

  DiabetesPedigreeFunction
Y       [,1]      [,2]
  0 0.4342139 0.3019496
  1 0.5815140 0.3794261

  Age
Y       [,1]     [,2]
  0 30.96687 11.35298
  1 36.95531 11.01981
```

- **#Now predicting the data remaining Split data using Trained dataset**

```
> #predicting, try putting type="class" or type="raw" after the test data
> pred1<-predict(my_model,test[,-9])
> pred1
 [1] 1 0 0 0 0 0 0 0 1 0 1 0 1 0 0 1 0 0 1 0 0 0 0 0 1 0 0 0 0 0 0 0 0 1 1 0 0 0 1 0 0 0 0 0 1 0 0 0 1 0 0 1 1 1
0 0 0 0 0 1 0 0 1 0 0 1 0 1 0 1
 [71] 0 0 0 0 0 0 0 1 0 1 0 1 0 0 0 0 1 0 0 1 1 0 0 0 0 0 0 0 0 0 1 0 1 0 0 0 0 0 0 0 1 0 0 1 0 0 0 0 0 1 1 1 0
0 1 1 1 1 0 1 1 0 0 0 0 0 0 0 1
[141] 0 0 1 0 0 0 0 0 0 0 0 0 0 1 0 0 0 1 0 0 0 0 1 1 0 1 0 0 0 0 0 0 0 1 0 1 0 0 0 0 1 0 1 1 0 0 1 0 0 0 0 0 0
0 1 0 1 0 1 1 0 1 1 0 1 0 1 1 1 0
[211] 0 0 0 0 0 1 1 0 0 1 1 1 1 0 0 1 1 1 0 0 0 1 0 0 1 1 1 0 1 0 0 0 0 0 0 0 0 0 0 1 0 1 1 1 0 0 0 0
Levels: 0 1
```

- **#Creating Confusion Matrix.**

```
> table(pred1, test$Outcome, dnn=c("predicted", "Actual"))
          Actual
predicted   0   1
        0 140  37
        1  28  52

> #To save the prediction
> output<-cbind(test, pred1)
> View(output)
```