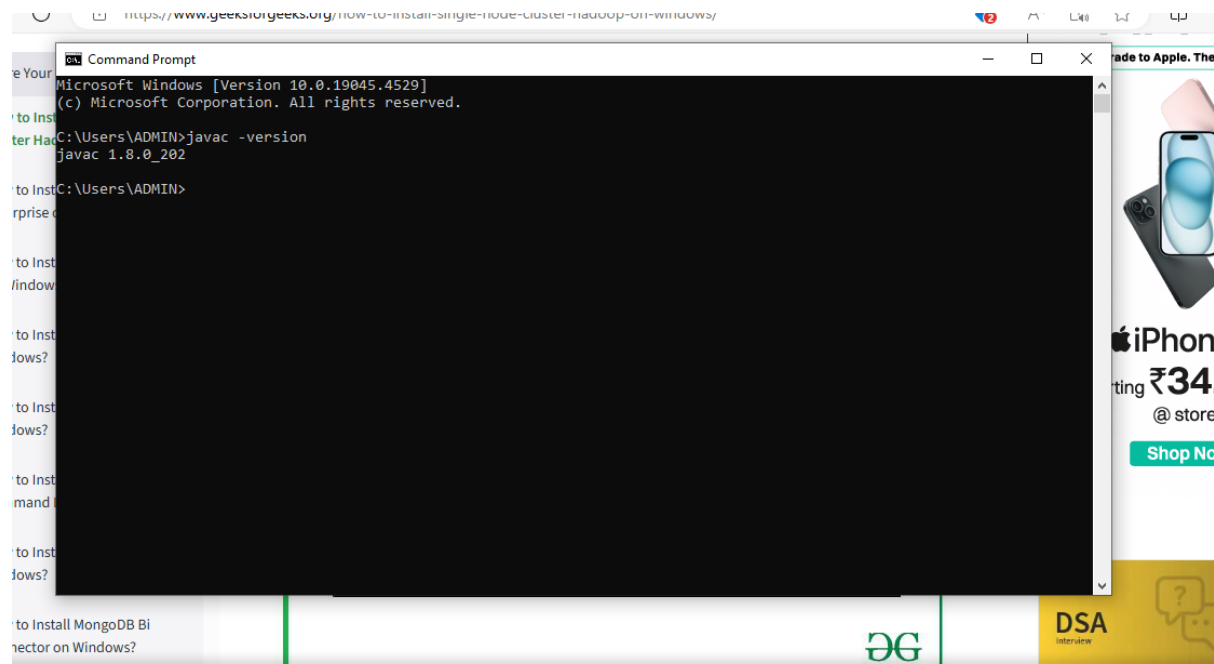
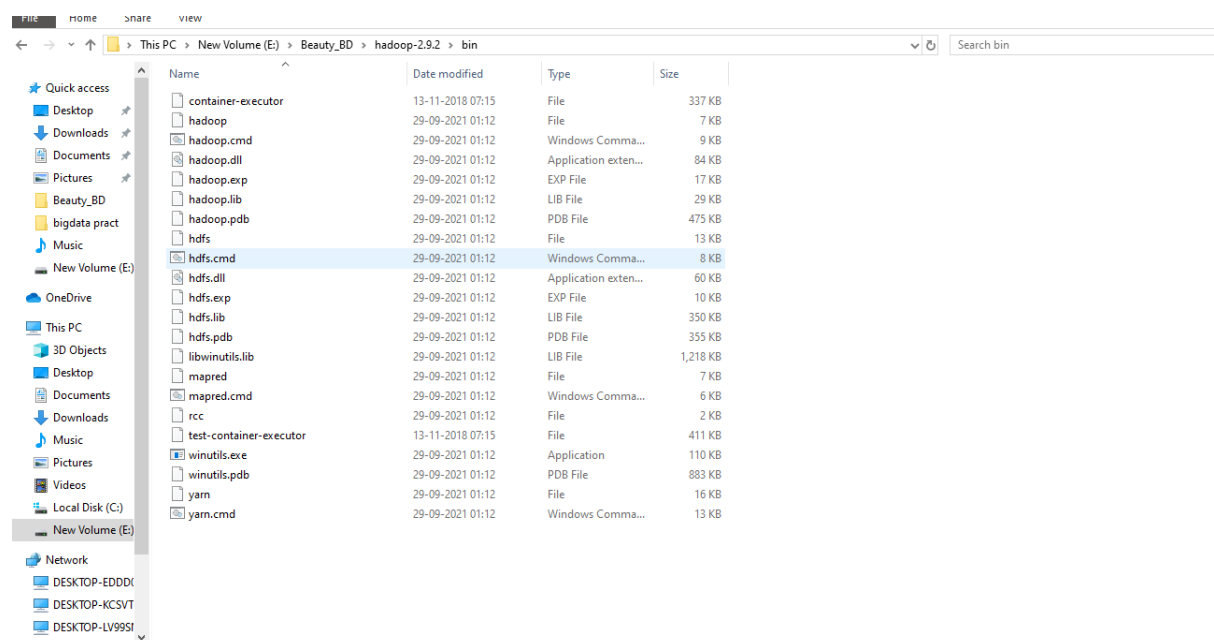


## Practical.1

### Step 1 :-Verify the Java installed

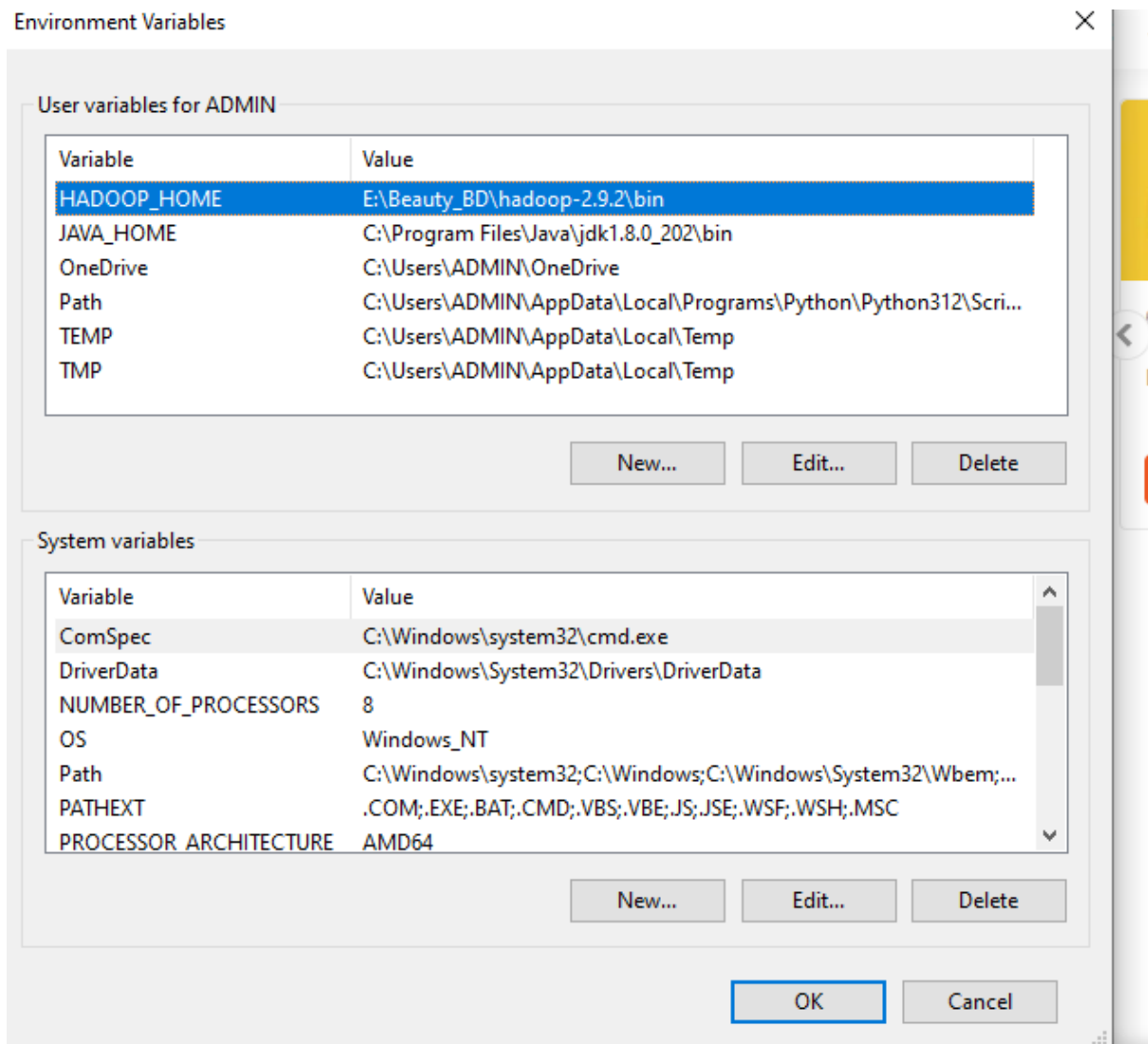


### Step 2: Extract Hadoop at C:\Hadoop



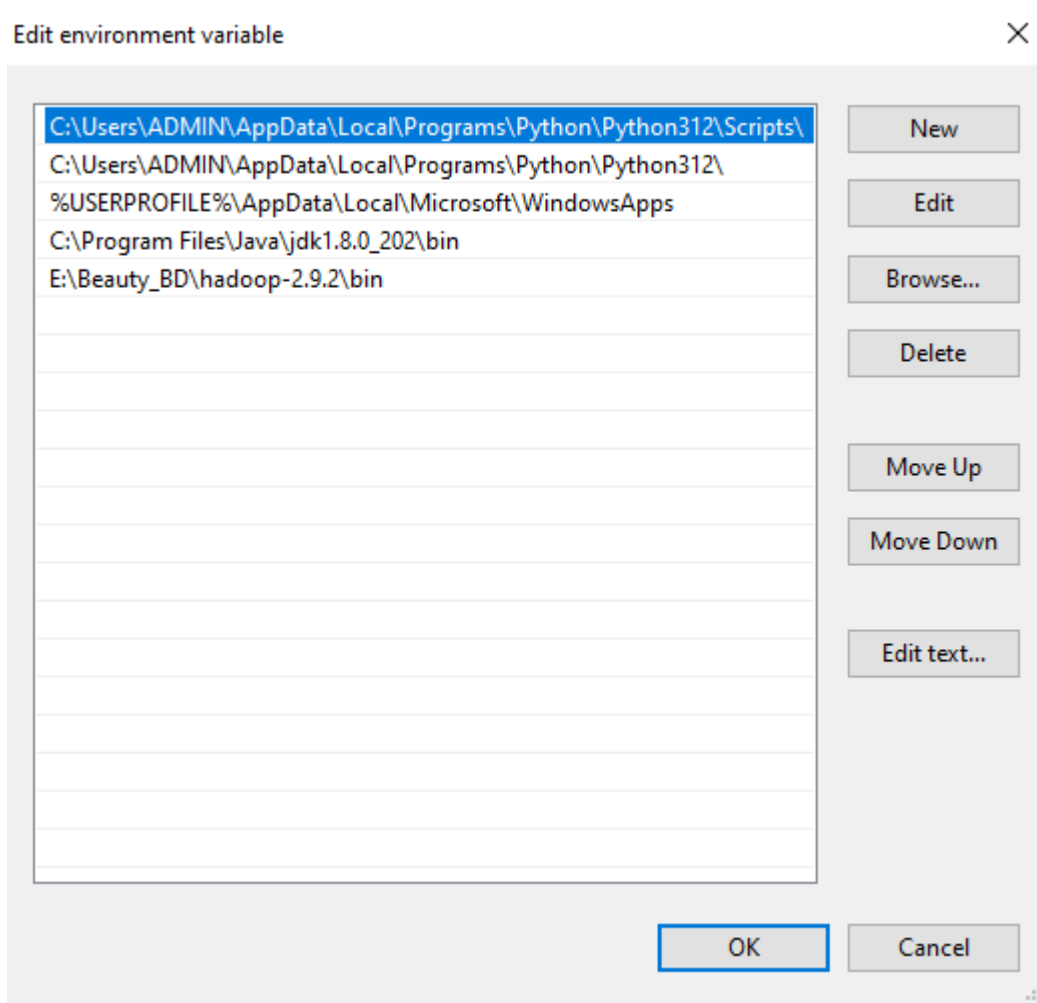
### Step 3: Setting up the HADOOP\_HOME variable And

### Step 4: Set JAVA\_HOME variable



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## Step 5: Set Hadoop and Java bin directory path



## Step 6: Hadoop Configuration :

1. Core-site.xml
2. Mapred-site.xml
3. Hdfs-site.xml
4. Yarn-site.xml
5. Hadoop-env.cmd
6. Create two folders datanode and namenode

### Step 6.1: Core-site.xml configuration

```
<configuration>
  <property>
    <name>fs.defaultFS</name>
    <value>hdfs://localhost:9000</value>
  </property>
</configuration>
```

### Step 6.2: Mapred-site.xml configuration

```
<configuration>
  <property>
    <name>mapreduce.framework.name</name>
    <value>yarn</value>
  </property>
</configuration>
```

### Step 6.3: Hdfs-site.xml configuration

```
<configuration>
  <property>
    <name>dfs.replication</name>
    <value>1</value>
  </property>
  <property>
    <name>dfs.namenode.name.dir</name>
    <value>C:\hadoop-2.8.0\data\namenode</value>
  </property>
  <property>
    <name>dfs.datanode.data.dir</name>
    <value>C:\hadoop-2.8.0\data\datanode</value>
  </property>
</configuration>
```

### Step 6.4: Yarn-site.xml configuration

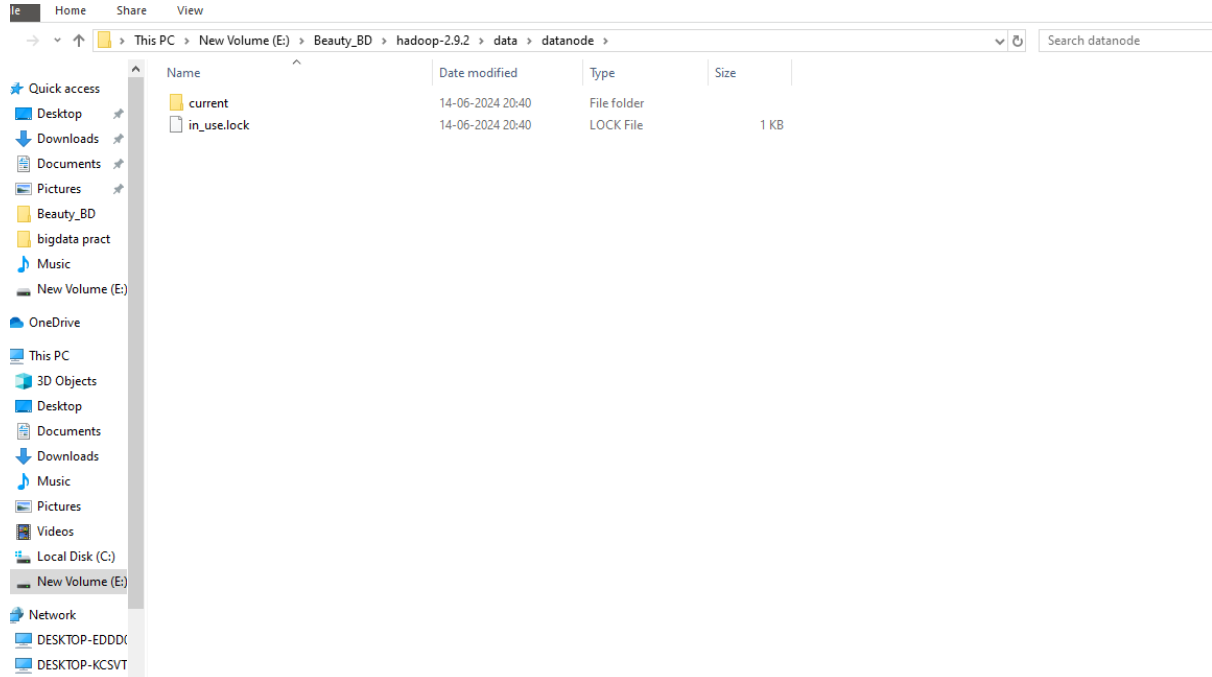
```
<configuration>
  <property>
    <name>yarn.nodemanager.aux-services</name>
    <value>mapreduce_shuffle</value>
  </property>
  <property>
    <name>yarn.nodemanager.auxservices.mapreduce.shuffle.class</name>
    <value>org.apache.hadoop.mapred.ShuffleHandler</value>
  </property>
```

```
</configuration>
```

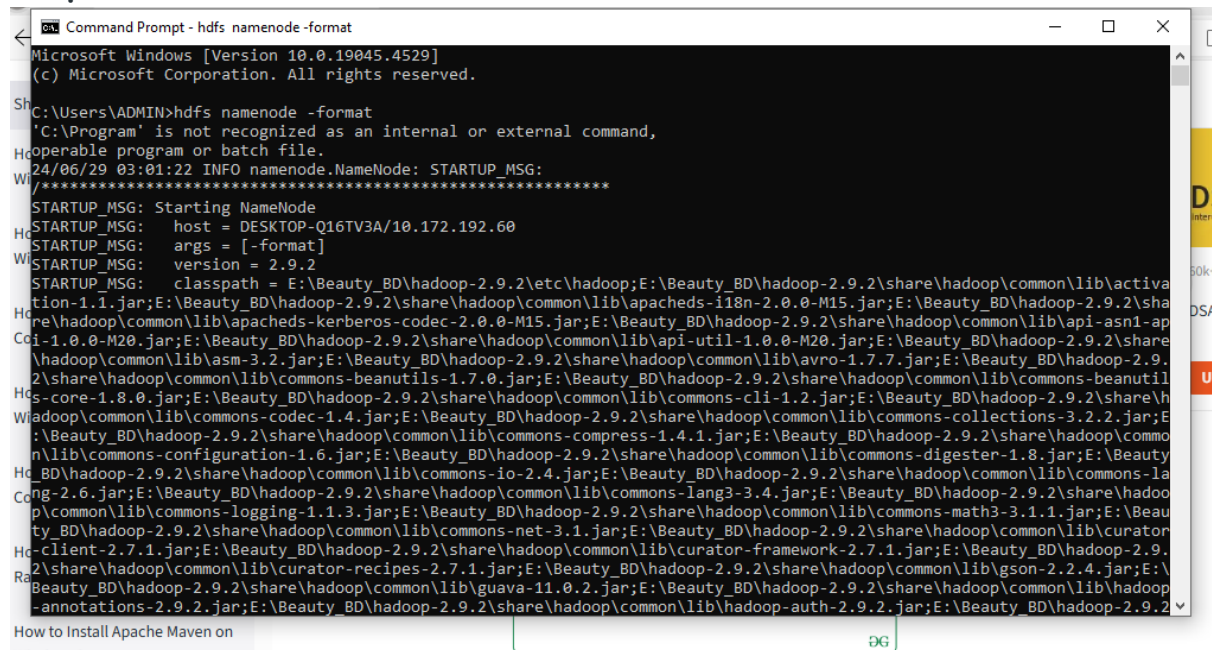
## Step 6.5: Hadoop-env.cmd configuration

Set "JAVA\_HOME=C:\Java" (On C:\java this is path to file jdk.18.0)

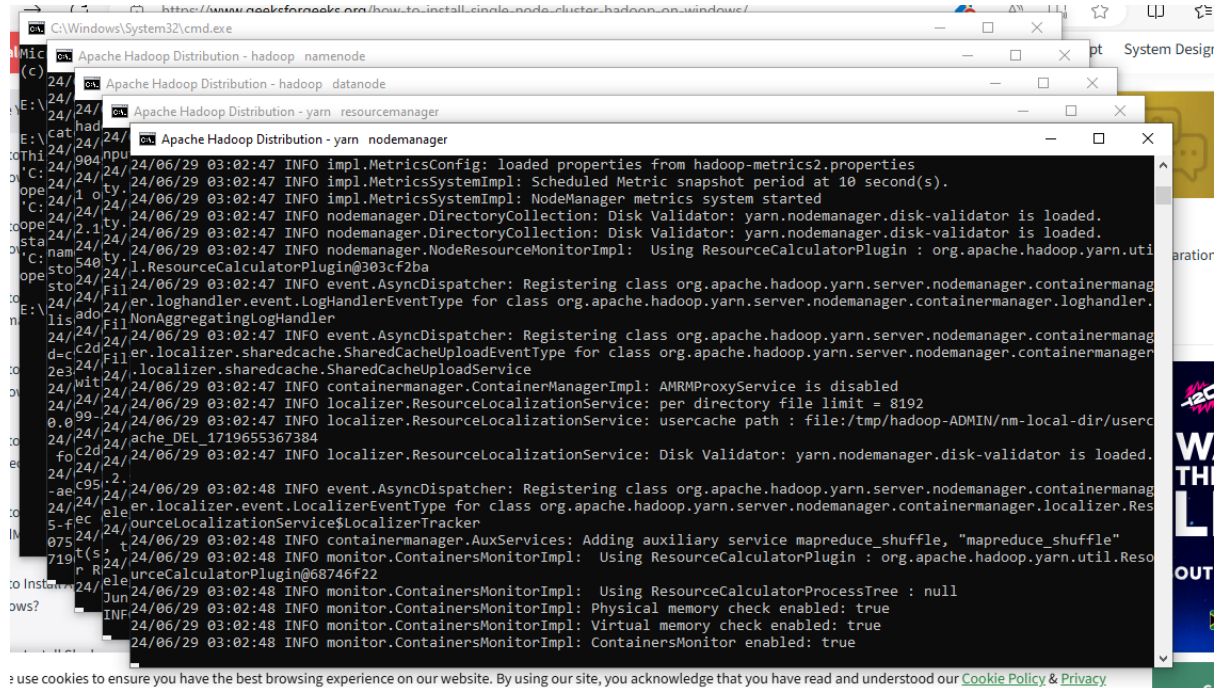
## Step 6.6: Create datanode and namenode folders



## Step 7: Format the namenode folder



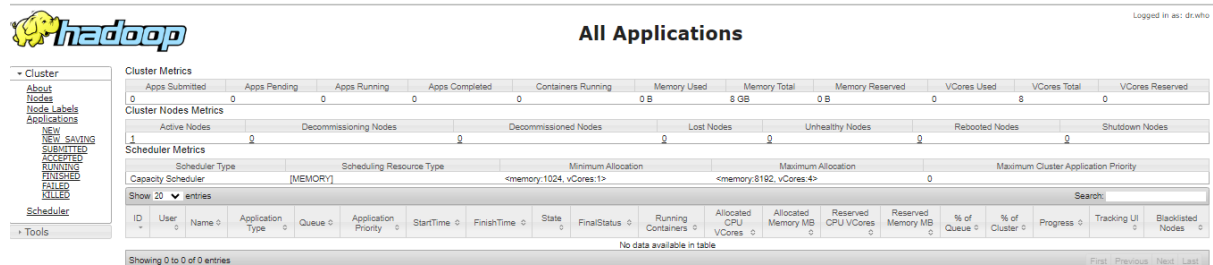
## Step 8: Testing the setup



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### Step 8.1: Testing the setup:

### Step 9: Open: <http://localhost:8088>



Step 10: <http://localhost:50070>

HadoopOverviewDatanodesDatanode Volume FailuresSnapshotStartup ProgressUtilities

Overview 'localhost:9000' (active)

Started:	Sat Jun 29 03:02:46 -0700 2024
Version:	2.9.2, r826afbeae31ca687bc2f8471dc841b66ed2c6704
Compiled:	Tue Nov 13 04:42:00 -0800 2018 by ajisaka from branch-2.9.2
Cluster ID:	CID-f0ac2e34-c5ea-4b71-815c-46ab2d3b2027
Block Pool ID:	BP-366826651-10.172.192.60-1719049540241

Summary

Security is off.  
Safemode is off.  
1 files and directories, 0 blocks = 1 total filesystem object(s).  
Heap Memory used 96.34 MB of 182.5 MB Heap Memory. Max Heap Memory is 889 MB.  
Non Heap Memory used 41.84 MB of 43.38 MB Committed Non Heap Memory. Max Non Heap Memory is <unbounded>.

## **Practical 2:**

**Aim:** Classification using SVM

**Requirement:**

R tool

**Code:**

```
getwd()
```

```
read.csv()
```

```
ds=read.csv("E:/Rajdeep/bigdata pract/dataset/social.csv",TRUE,"")
```

```
ds
```

```
ds=ds[3:5]
```

```
ds
```

```
install("catools")
```

```
library(caTools)
```

```
set.seed(123)
```

```
split=sample.split(ds$Purchased, SplitRatio=0.75)
```

```
training_set=(subset(ds, split == TRUE))
```

```
test_set =(subset(ds, split == FALSE))
```

```
ds
```

```
test_set[-3]=scale(test_set[-3])
```

```
training_set[-3]=scale(training_set[-3])
```

```
test_set[-3]
```

```
training_set[-3]
```

```
install.packages('e1071')
```

```
library('e1071')
```

```
classifier=svm(formula=Purchased ~ ., data= training_set , type='C-classification',kernal='linear')
```

```
classifier
```

```
y_pred=predict(classifier, newdata=test_set[-3])
```

```
y_pred
```

```
cm=table(test_set[, 3],y_pred)
```

```
cm
```



```

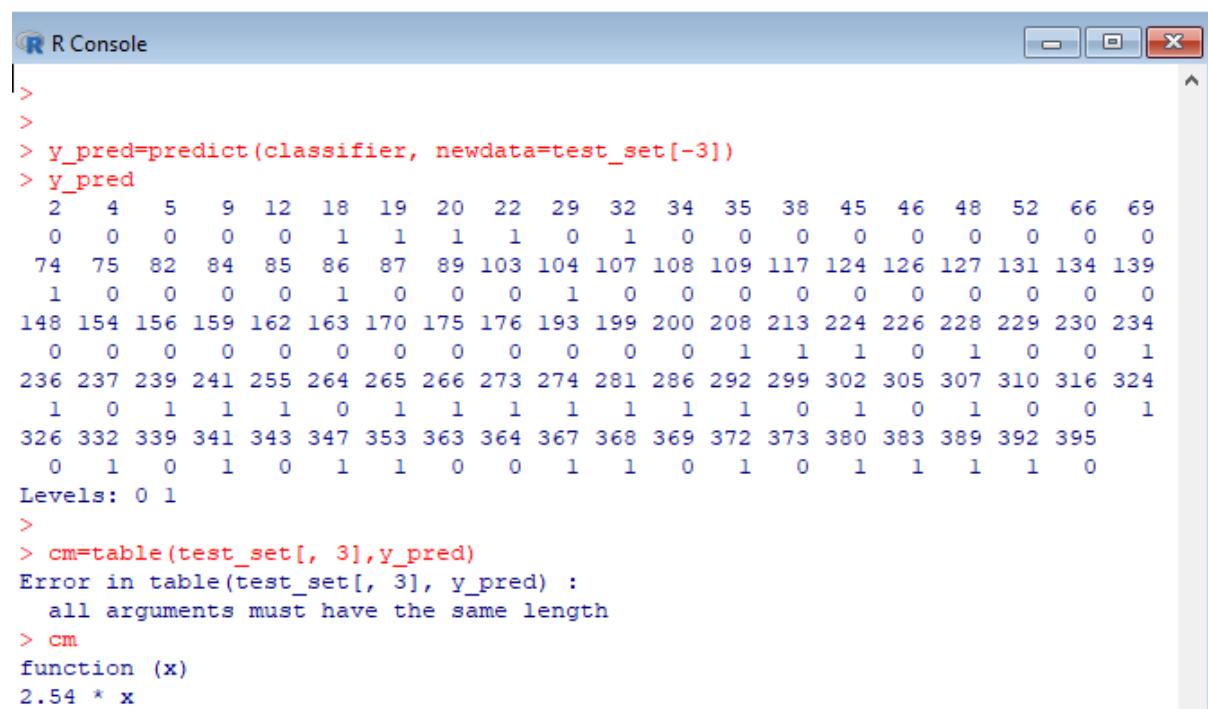
> set = training_set
> x1 = seq(min(set[, 1]) - 1, max(set[, 1]) + 1, by = 0.01)
> x2 = seq(min(set[, 2]) - 1, max(set[, 2]) + 1, by = 0.01)

> grid_set = expand.grid(x1, x2)
> colnames(grid_set) = c('Age', 'EstimatedSalary')
> y_grid = predict(classifier, newdata = grid_set)
> plot(set[, -3],
+       main = 'SVM (Training set)',
+       xlab = 'Age', ylab = 'Estimated Salary',
+       xlim = range(x1), ylim = range(x2))

> contour(x1, x2, matrix(as.numeric(y_grid), length(x1), length(x2)), add = TRUE)
> points(grid_set, pch = '.', col = ifelse(y_grid == 1, 'coral1', 'aquamarine'))
> points(set, pch = 21, bg = ifelse(set[, 3] == 1, 'green4', 'red3'))

```

### Output:



```

R Console
>
>
> y_pred=predict(classifier, newdata=test_set[-3])
> y_pred
 2   4   5   9  12  18  19  20  22  29  32  34  35  38  45  46  48  52  66  69
0   0   0   0   0   1   1   1   1   0   1   0   0   0   0   0   0   0   0
74  75  82  84  85  86  87  89 103 104 107 108 109 117 124 126 127 131 134 139
1   0   0   0   0   1   0   0   0   1   0   0   0   0   0   0   0   0   0
148 154 156 159 162 163 170 175 176 193 199 200 208 213 224 226 228 229 230 234
0   0   0   0   0   0   0   0   0   0   0   0   1   1   1   0   1   0   0   1
236 237 239 241 255 264 265 266 273 274 281 286 292 299 302 305 307 310 316 324
1   0   1   1   1   0   1   1   1   1   1   1   1   0   1   0   1   0   0   1
326 332 339 341 343 347 353 363 364 367 368 369 372 373 380 383 389 392 395
0   1   0   1   0   1   1   0   0   1   1   0   1   0   1   1   1   1   0
Levels: 0 1
>
> cm=table(test_set[, 3],y_pred)
Error in table(test_set[, 3], y_pred) :
  all arguments must have the same length
> cm
function (x)
2.54 * x

```

### Practical 3:

**Aim:** write program in R of Naive bayes's theorem

**Requirement:**

R tool

**Code:**

```
data(iris)
```

```
str(iris)
```

```
install.packages("e1071")
```

```
install.packages("caTools")
```

```
install.packages("caret")
```

```
library(e1071)
```

```
library(caTools)
```

```
library(caret)
```

```
split <- sample.split(iris, SplitRatio=0.7)
```

```
train_c1 <- subset(iris, split=="TRUE")
```

```
test_c1 <- subset(iris, split == "FALSE")
```

```
train_scale <- scale(train_c1[, 1:4])
```

```
test_scale <- scale(test_c1[, 1:4])
```

```
set.seed(120)
```

```
classifier_c1 <- naiveBayes(Species ~ ., data = train_c1)
```

```
classifier_c1
```

```
y_pred <- predict(classifier_c1, newdata= test_c1)
```

```
cm <- table(test_c1$Species, y_pred)
```

```
cm
```

```
confusionMatrix(cm)
```

## Output:

```
R Console
virginica      0      1      19

Overall Statistics

      Accuracy : 0.9333
      95% CI   : (0.838, 0.9815)
No Information Rate : 0.3667
P-Value [Acc > NIR] : < 2.2e-16

      Kappa : 0.9

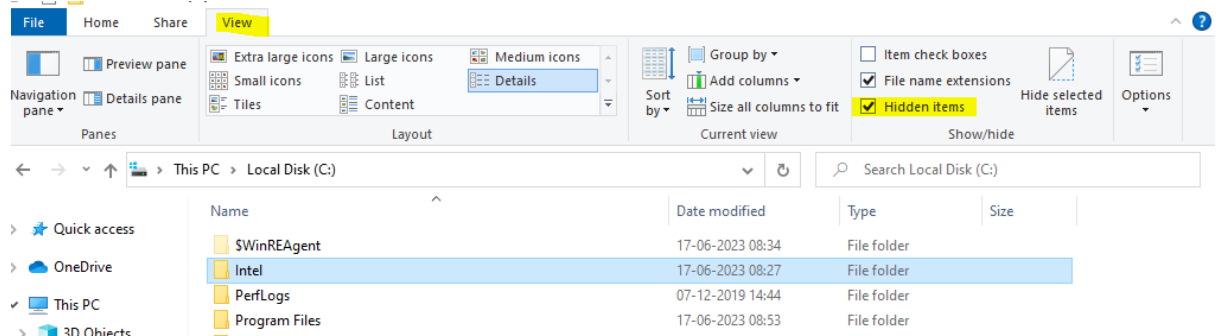
McNemar's Test P-Value : NA

Statistics by Class:

      Class: setosa Class: versicolor Class: virginica
Sensitivity          1.0000          0.9444          0.8636
Specificity          1.0000          0.9286          0.9737
Pos Pred Value       1.0000          0.8500          0.9500
Neg Pred Value       1.0000          0.9750          0.9250
Prevalence           0.3333          0.3000          0.3667
Detection Rate       0.3333          0.2833          0.3167
Detection Prevalence 0.3333          0.3333          0.3333
Balanced Accuracy     1.0000          0.9365          0.9187
> |
```

## Install python package:

1. You will need to make the hidden folder visible: go to "C:" drive on top click on tab "view"
2. Select "hidden Items" option:



3. Go to the below path:  
C:\Users\Your Name\AppData\Local\Programs\Python\Python36-32\Scripts
4. Set the below path in command prompt and then use the below command:  
python -m pip install pymongo

```
Microsoft Windows [Version 10.0.19045.2965]
(c) Microsoft Corporation. All rights reserved.

C:\Users\RPIMS>cd\

C:\>cd C:\Users\RPIMS\AppData\Local\Programs\Python\Python310\Scripts

C:\Users\RPIMS\AppData\Local\Programs\Python\Python310\Scripts>python -m pip install pymongo
Defaulting to user installation because normal site-packages is not writeable
Requirement already satisfied: pymongo in c:\users\rpims\appdata\roaming\python\python310\site-packages (4.3.3)
Requirement already satisfied: dnspython<3.0.0,>=1.16.0 in c:\users\rpims\appdata\roaming\python\python310\site-packages
(from pymongo) (2.3.0)

C:\Users\RPIMS\AppData\Local\Programs\Python\Python310\Scripts>
```

## **Practical :4**

**Aim:** Implement an application that stores big data in Hbase / MongoDB and manipulate it using R / Python

### **Requirement:**

- a. Python Package: PyMongo
- b. Mongo Database

### **Step A: Install Mongo database**

**Step 1)** Go to (<https://www.mongodb.com/download-center/community>) and Download MongoDB Community Server. We will install the 64-bit version for Windows.

Select the server you would like to run:

**MongoDB Community Server**  
FEATURE RICH. DEVELOPER READY.

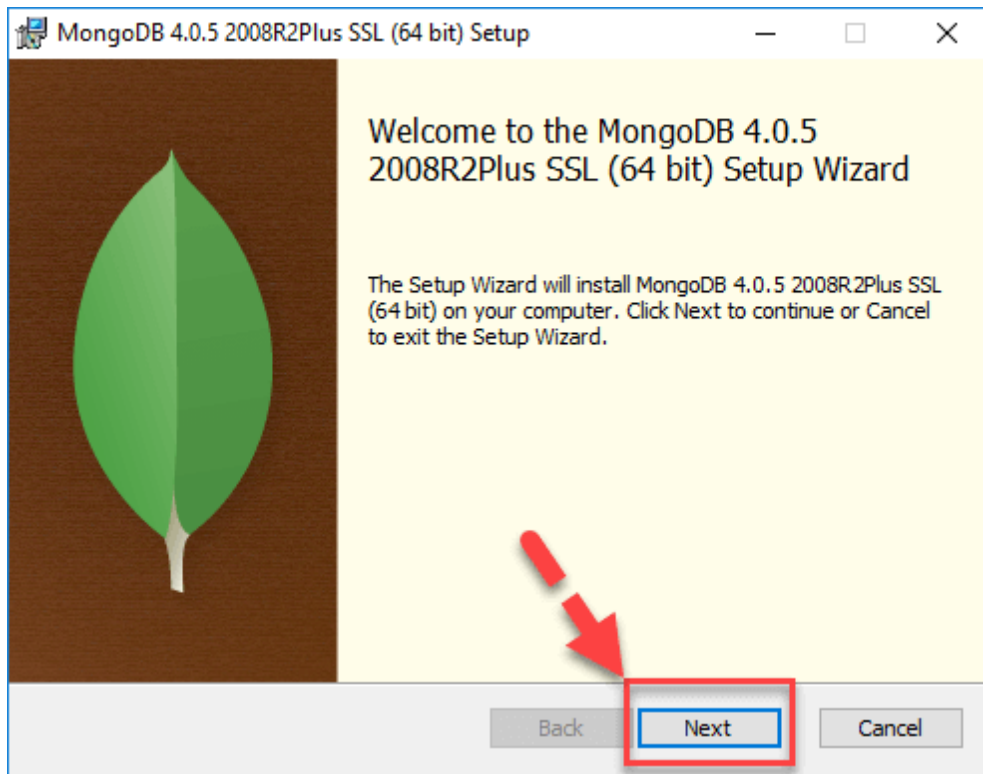
Version: 4.0.5 (current release) OS: Windows 64-bit x64

Package: MSI

**Download**

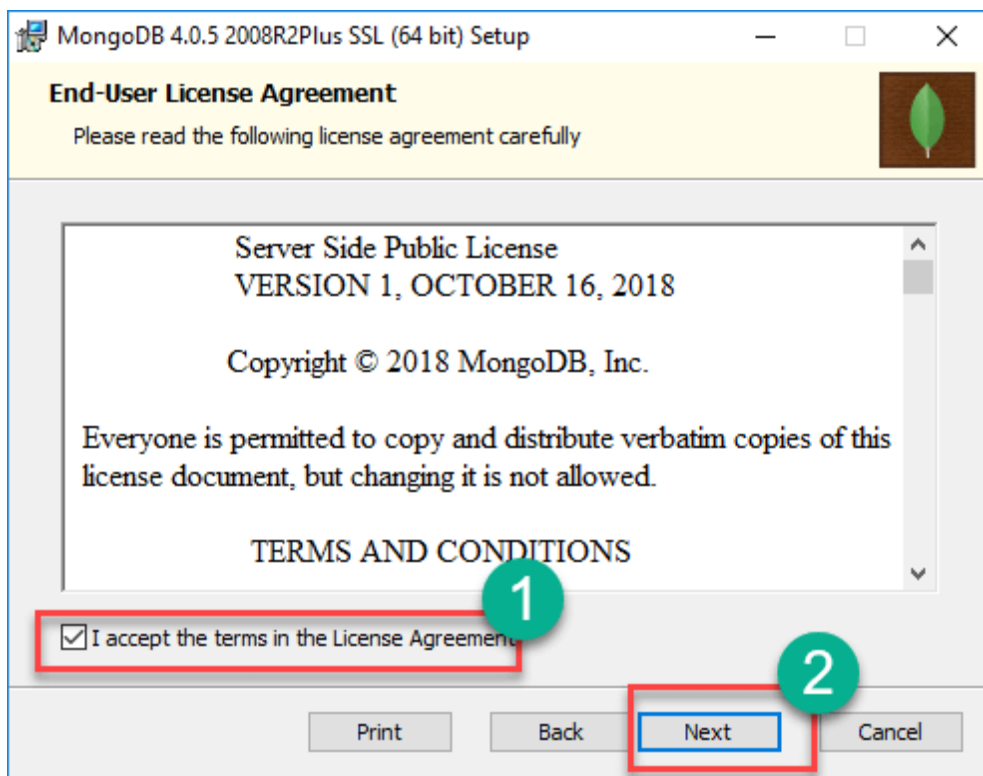
[https://fastdl.mongodb.org/win32/mongodb-win32-x86\\_64-2008plus-ssl-4.0.5-signed.msi](https://fastdl.mongodb.org/win32/mongodb-win32-x86_64-2008plus-ssl-4.0.5-signed.msi)

**Step 2)** Once download is complete open the msi file. Click Next in the start up screen

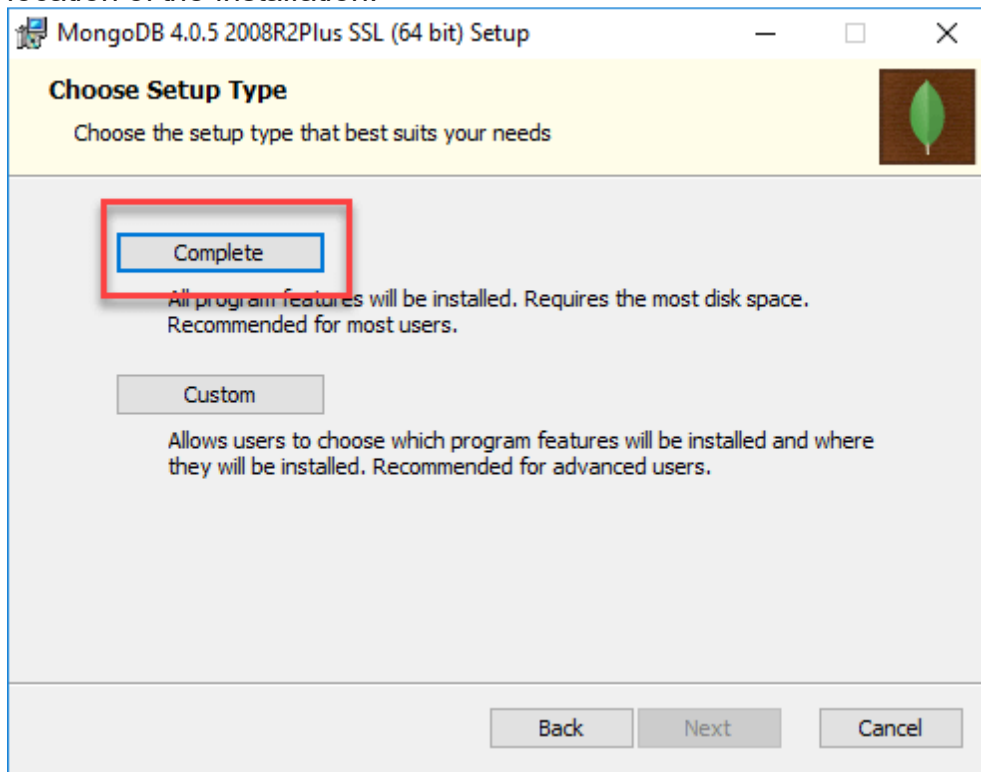


### Step 3)

1. Accept the End-User License Agreement
2. Click Next



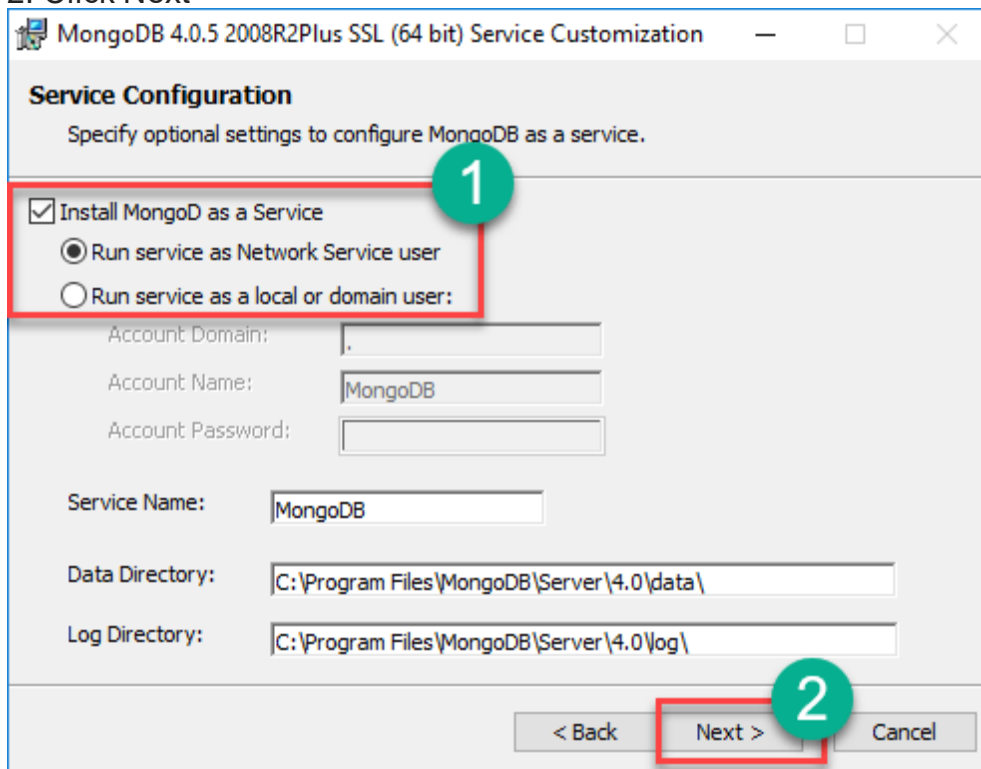
**Step 4)** Click on the "complete" button to install all of the components. The custom option can be used to install selective components or if you want to change the location of the installation.



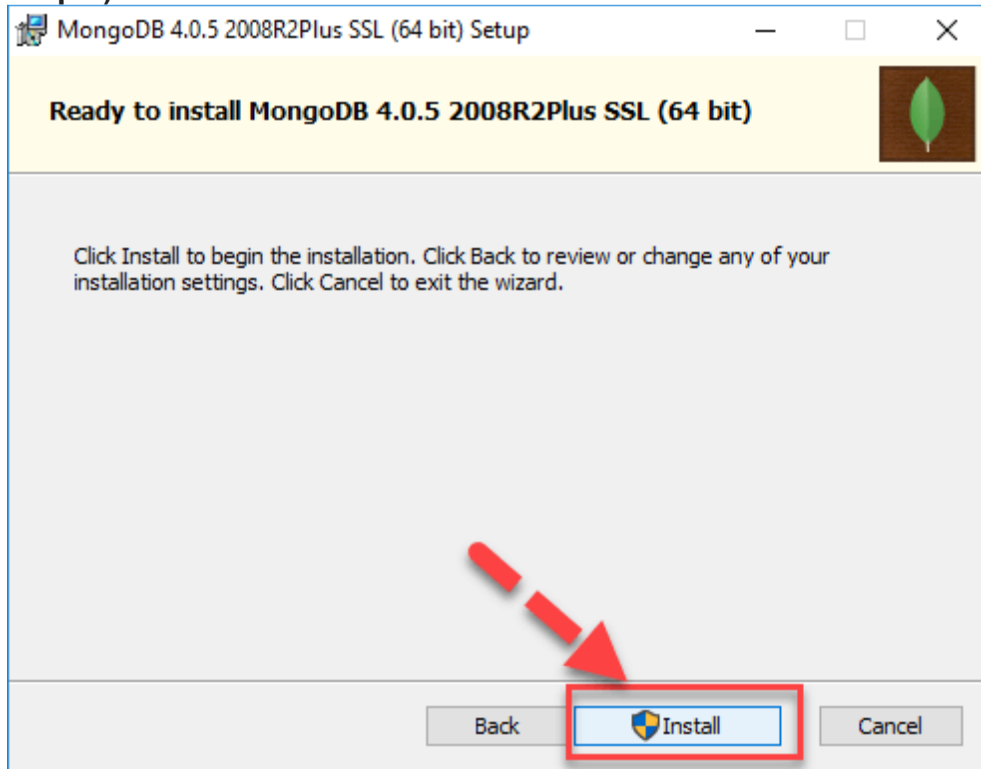
**Step 5)**

1. Select "Run service as Network Service user". make a note of the data directory, we'll need this later.

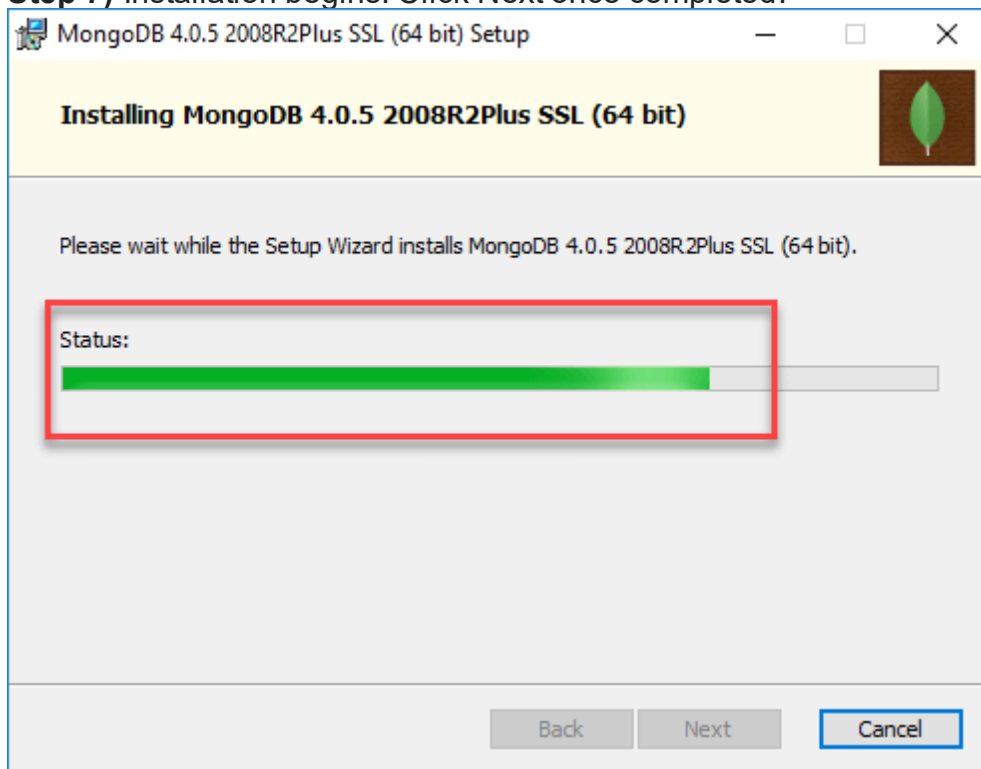
2. Click Next



**Step 6)** Click on the Install button to start the installation.

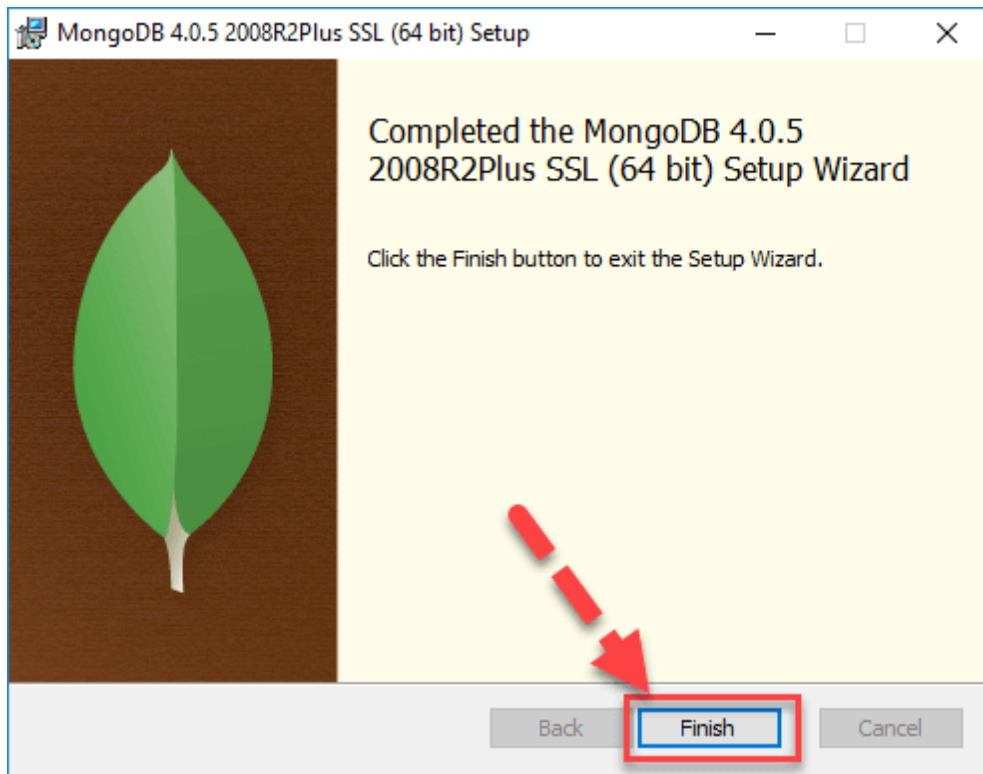


**Step 7)** Installation begins. Click Next once completed.



**Step 8)** Click on the Finish button to complete the installation





**Program 1:** Displaying the database name:

```
import pymongo
myclient = pymongo.MongoClient("mongodb://localhost:27017/")
mydb = myclient["mybigdata"]
print(myclient.list_database_names())
```

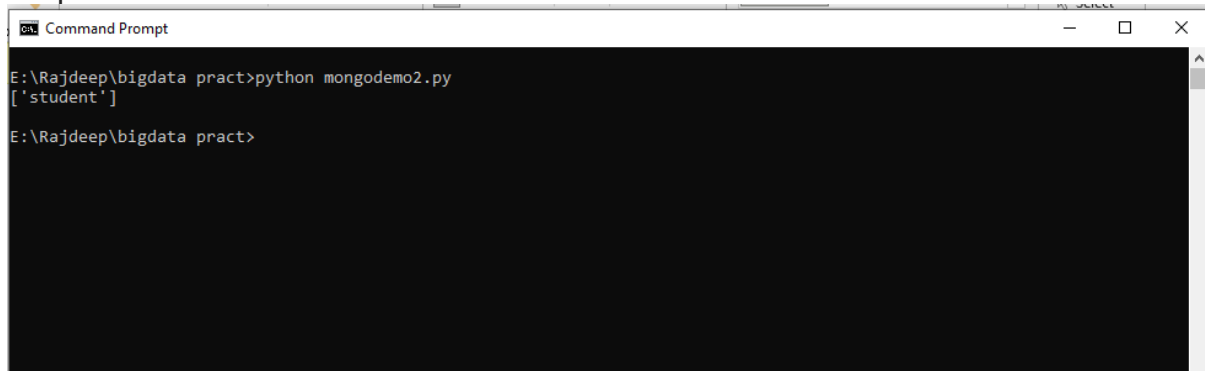
Output:

```
Command Prompt
E:\Rajdeep\bigdata pract>python mongodemo1.py
['admin', 'config', 'local', 'mybigdata']
E:\Rajdeep\bigdata pract>
```

**Program 2:** Creating collection:

```
import pymongo
myclient = pymongo.MongoClient("mongodb://localhost:27017/")
mydb = myclient["mybigdata"]
mycol=mydb["student"]
print(mydb.list_collection_names())
```

Output:

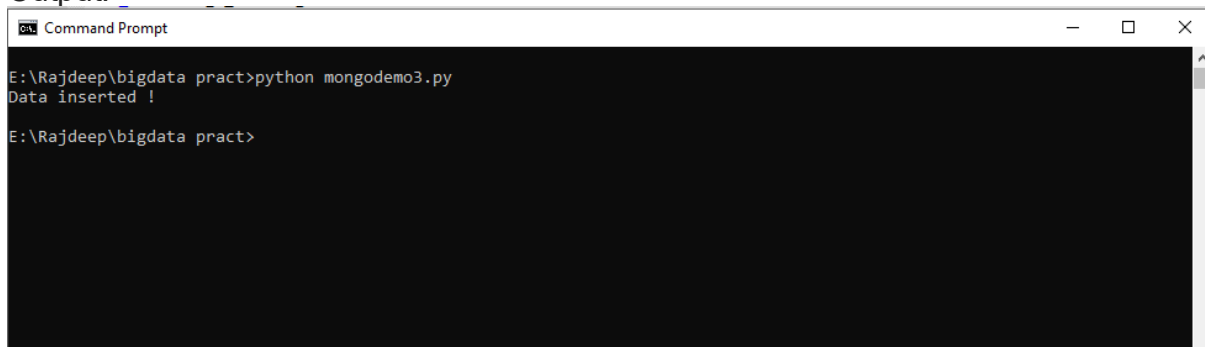


```
Command Prompt
E:\Rajdeep\bigdata pract>python mongodemo2.py
['student']
E:\Rajdeep\bigdata pract>
```

### **Program 3:** Inserting Data

```
import pymongo
myclient = pymongo.MongoClient("mongodb://localhost:27017/")
mydb = myclient["mybigdata"]
mycol=mydb["student"]
mydict={"name":"vai", "address":"bhy"}
x=mycol.insert_one(mydict)
print("Data inserted !")
```

Output:

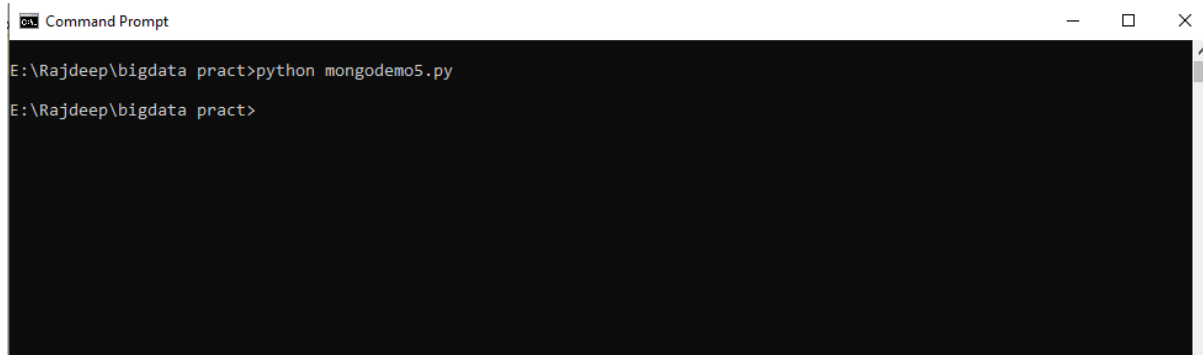


```
Command Prompt
E:\Rajdeep\bigdata pract>python mongodemo3.py
Data inserted !
E:\Rajdeep\bigdata pract>
```

### **Program 4:** Insert Multiple data into Collection

```
import pymongo
myclient = pymongo.MongoClient("mongodb://localhost:27017/")
mydb = myclient["mybigdata"]
mycol=mydb["student"]
mylist=[{"name":"Ganesh", "address":"Mumbai"}, {"name":"Varun",
"address":"Mumbai"},
{"name":"Prasoon", "address":"Pune"}, {"name":"Satish", "address":"Pune"},]
x=mycol.insert_many(mylist)
print("Data inserted !")
```

Output:



```
Command Prompt
E:\Rajdeep\bigdata pract>python mongodemo5.py
E:\Rajdeep\bigdata pract>
```

**Program 5:** Displaying the collection data:

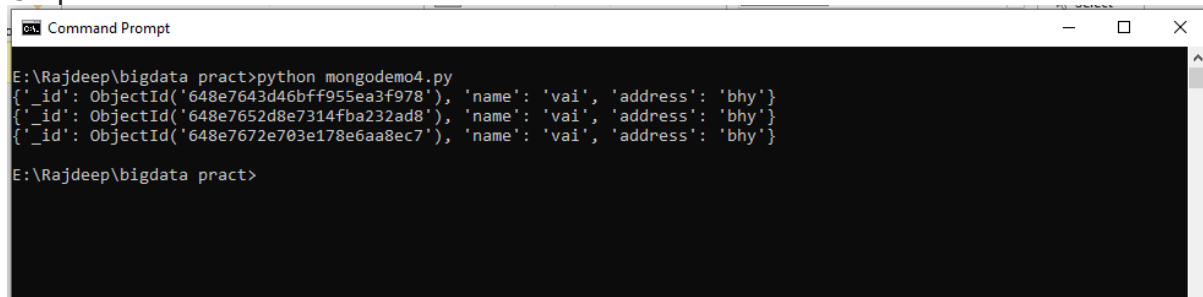
```
import pymongo
myclient = pymongo.MongoClient("mongodb://localhost:27017/")
mydb = myclient["mybigdata"]
mycol = mydb["student"]

myquery = { "name": "Vai" }

mydoc = mycol.find(myquery)

for x in mydoc:
    print(x)
```

Output:



```
Command Prompt
E:\Rajdeep\bigdata pract>python mongodemo4.py
{'_id': ObjectId('648e7643d46bff955ea3f978'), 'name': 'vai', 'address': 'bhy'}
{'_id': ObjectId('648e7652d8e7314fba232ad8'), 'name': 'vai', 'address': 'bhy'}
{'_id': ObjectId('648e7672e703e178e6aa8ec7'), 'name': 'vai', 'address': 'bhy'}
E:\Rajdeep\bigdata pract>
```

### **Practical 5:**

K means clustering.

**Aim:** Read a datafile grades\_km\_input.csv and apply k-means clustering.

**Requirement:**

R tool

#### **Code:**

```
install.packages("plyr")
install.packages("ggplot2")
install.packages("cluster")
install.packages("lattice")
install.packages("grid")
install.packages("gridExtra")
```

```
library(plyr)
library(ggplot2)
library(cluster)
library(lattice)
library(grid)
library(gridExtra)
```

```
grade_input=as.data.frame(read.csv("E:/Rajdeep/bigdata
pract/dataset/grades_km_input.csv"))
```

```
kmdata_orig=as.matrix(grade_input[, c ("Student","English","Math","Science")])
kmdata=kmdata_orig[,2:4]
kmdata[1:10,]
wss=numeric(15)
```

```
for(k in 1:15)wss[k]=sum(kmeans(kmdata,centers=k,nstart=25)$withinss)
plot(1:15,wss,type="b",xlab="Number of Clusters",ylab="Within sum of square")
km = kmeans(kmdata,3,nstart=25)
km
```

```
c( wss[3] , sum(km$withinss))
df=as.data.frame(kmdata_orig[,2:4])
df$cluster=factor(km$cluster)
centers=as.data.frame(km$centers)
```

```
g1=ggplot(data=df, aes(x=English, y=Math, color=cluster )) +
geom_point() + theme(legend.position="right") +
geom_point(data=centers,aes(x=English,y=Math, color=as.factor(c(1,2,3))),size=10,
alpha=.3, show.legend =FALSE)
```

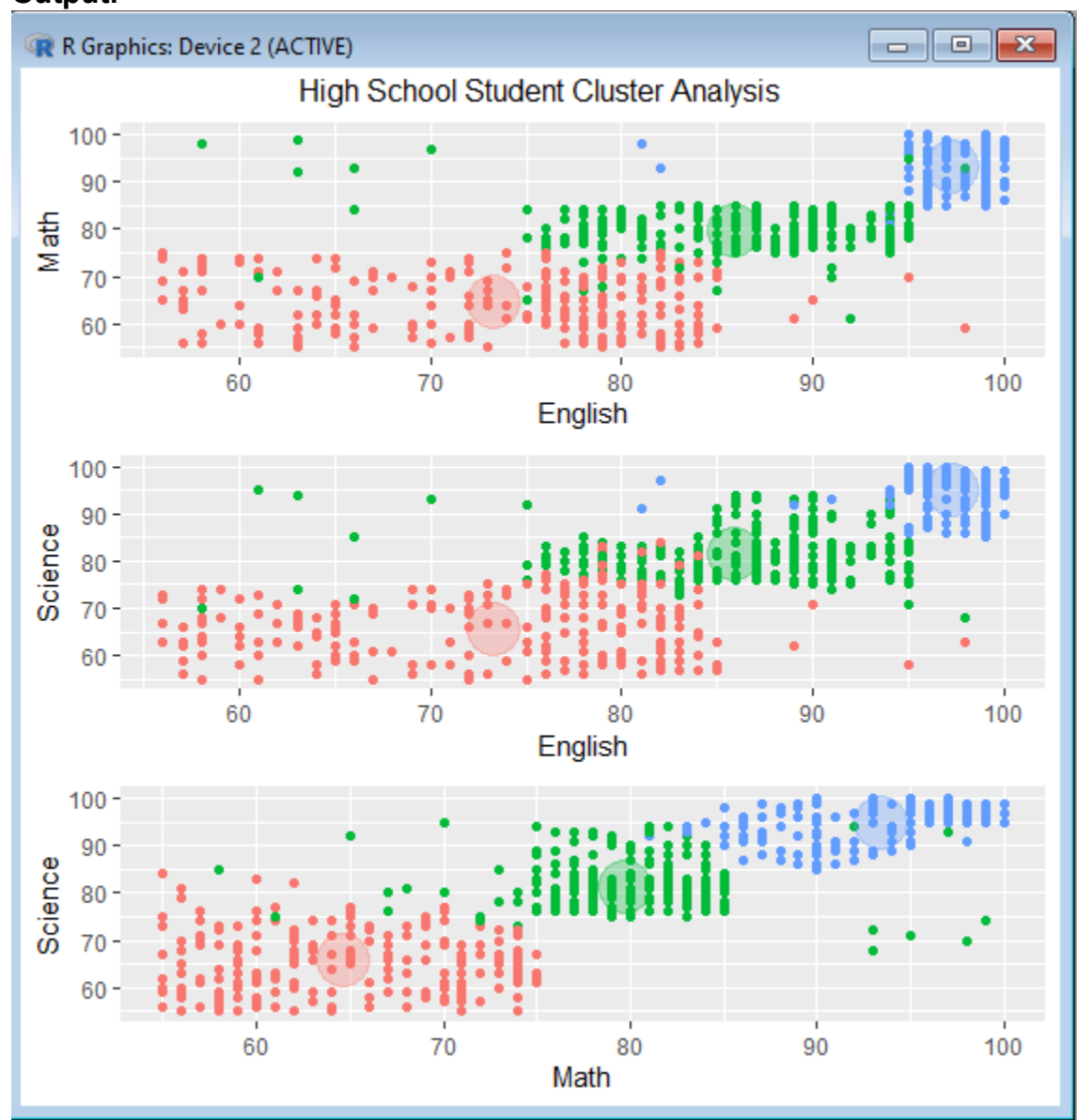
```
g2=ggplot(data=df, aes(x=English, y=Science, color=cluster )) +
geom_point () +geom_point(data=centers,aes(x=English,y=Science,
color=as.factor(c(1,2,3))),size=10, alpha=.3, show.legend=FALSE)
```

```
g3 = ggplot(data=df, aes(x=Math, y=Science, color=cluster )) +
```

```
geom_point() + geom_point(data=centers,aes(x=Math,y=Science,
color=as.factor(c(1,2,3))),size=10, alpha=.3, show.legend=FALSE)
```

```
tmp=ggplot_gtable(ggplot_build(g1))
grid.arrange(arrangeGrob(g1 + theme(legend.position="none"),g2 +
theme(legend.position="none"),g3 + theme(legend.position="none"),top ="High
School Student Cluster Analysis" ,ncol=1))
```

### Output:



### **Practical 6:**

a. Simple Linear regression

**Aim:** Create your own data for years of experience and salary in lakhs and apply linear regression model to predict the salary

**Requirement:**

R tool

Code:

```
years_of_exp = c(7,5,1,3)
```

```
salary_in_lakhs = c(21,13,6,8)
```

```
employee.data = data.frame(years_of_exp, salary_in_lakhs)
```

```
employee.data
```

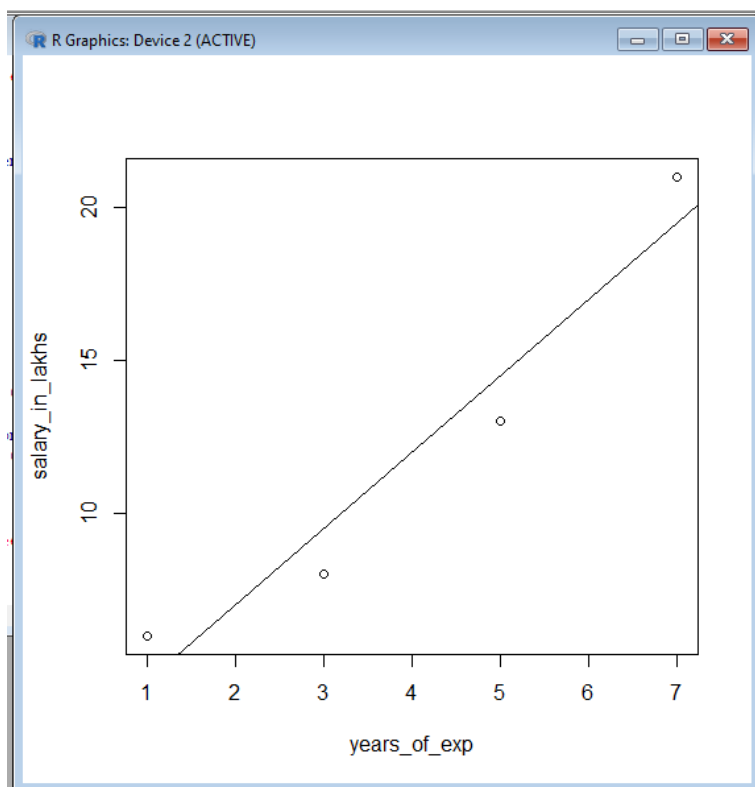
```
model <- lm(salary_in_lakhs ~ years_of_exp, data = employee.data)
```

```
summary(model)
```

```
plot(salary_in_lakhs ~ years_of_exp, data = employee.data)
```

```
abline(model)
```

**Output:**



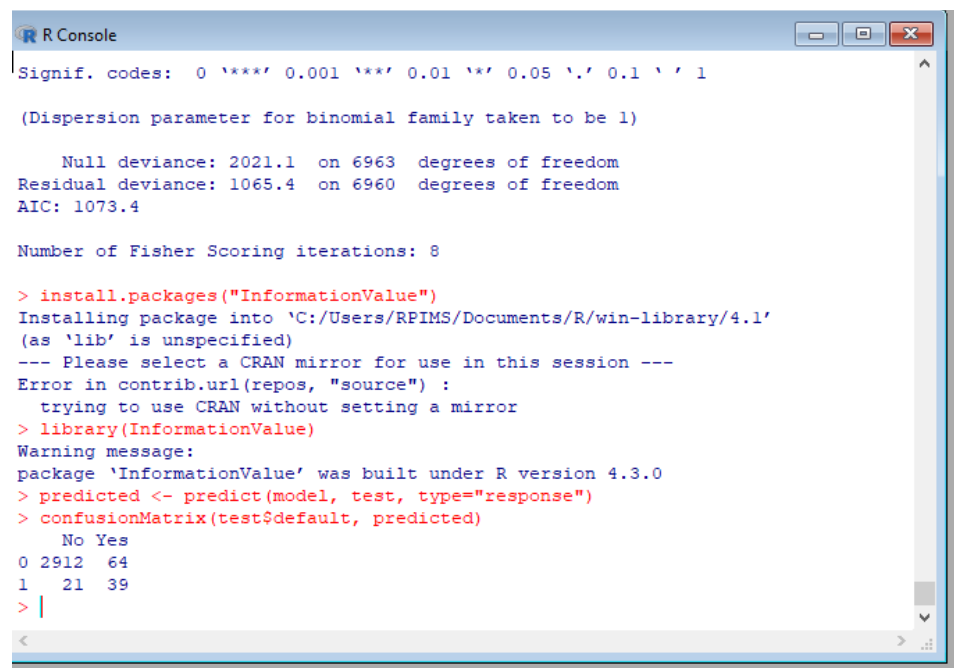
b.: Logistic regression:

**Aim:** Take the in-built data from ISLR package and apply generalized logistic regression to find whether a person would be defaulter or not; considering input as student, income and balance.

Code:

```
install.packages("ISLR")
library(ISLR)
data <- ISLR::Default
print(head(ISLR::Default))
summary(data)
nrow(data)
set.seed(1)
sample <- sample(c(TRUE, FALSE), nrow(data), replace=TRUE, prob=c(0.7,0.3))
print(sample)
train <- data[sample, ]
test <- data[!sample, ]
nrow(train)
nrow(test)
model <- glm(default~student+balance+income, family="binomial", data=train)
summary(model)
install.packages("InformationValue")
library(InformationValue)
predicted <- predict(model, test, type="response")
confusionMatrix(test$default, predicted)
```

Output:



```
R Console
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

    Null deviance: 2021.1  on 6963  degrees of freedom
Residual deviance: 1065.4  on 6960  degrees of freedom
AIC: 1073.4

Number of Fisher Scoring iterations: 8

> install.packages("InformationValue")
Installing package into 'C:/Users/RPIMS/Documents/R/win-library/4.1'
(as 'lib' is unspecified)
--- Please select a CRAN mirror for use in this session ---
Error in contrib.url(repos, "source") :
  trying to use CRAN without setting a mirror
> library(InformationValue)
Warning message:
package 'InformationValue' was built under R version 4.3.0
> predicted <- predict(model, test, type="response")
> confusionMatrix(test$default, predicted)
      No Yes
0 2912  64
1   21  39
> |
```

## **Practical 7:**

**Aim:** Implement Decision tree classification techniques

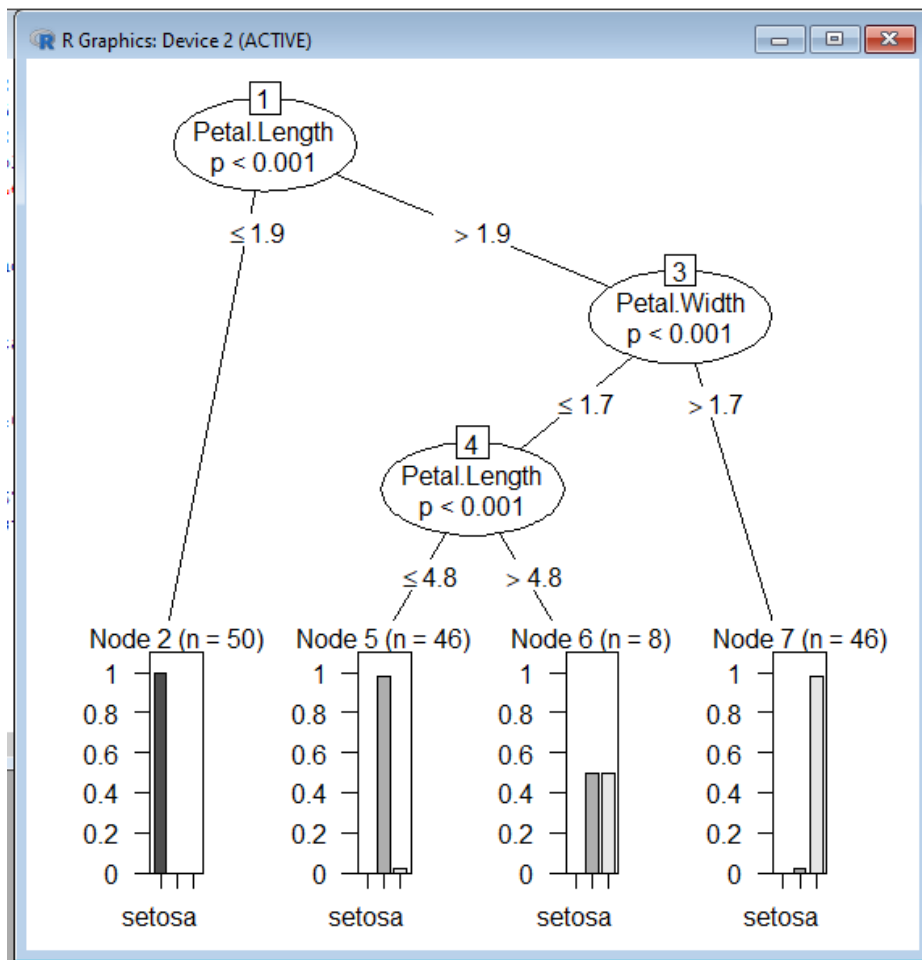
## **Requirement:**

R tool

## **Code:**

```
library("party")  
print(head(readingSkills))  
  
str(iris)  
iris_ctree <- ctree(Species ~ Sepal.Width + Sepal.Length + Petal.Length +  
Petal.Width, data=iris)  
  
print (iris_ctree)  
plot(iris_ctree)
```

## **Output:**





## **Practical 8:**

Apriori algorithm

**Aim:** Perform Apriori algorithm using Groceries dataset from the R arules package.

### **Requirement:**

R tool

### **Code:**

```
library(arules)
library(arulesViz)
library(RColorBrewer)
```

```
data(Groceries)
Groceries
```

```
summary(Groceries)
class(Groceries)
```

```
rules = apriori(Groceries, parameter = list(supp = 0.02, conf = 0.2))
summary(rules)
```

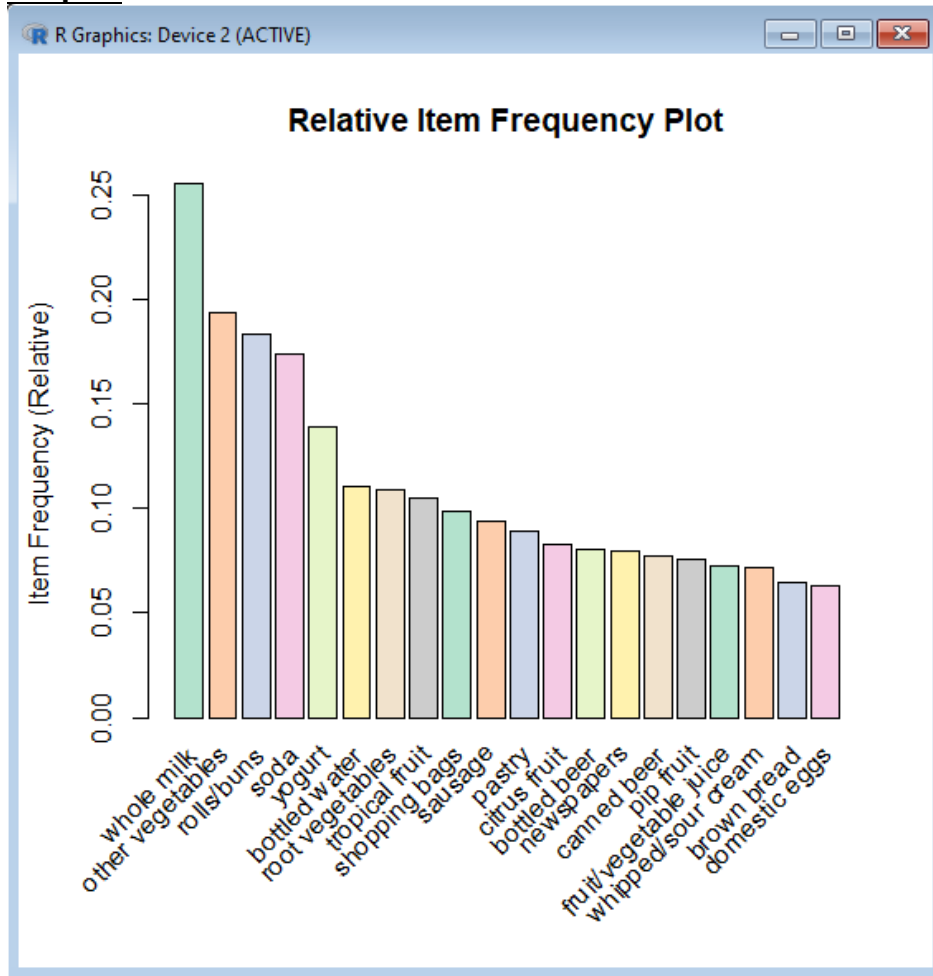
```
inspect(rules[1:10])
```

```
arules::itemFrequencyPlot(Groceries, topN = 20,
col = brewer.pal(8, 'Pastel2'),
main = 'Relative Item Frequency Plot',
type = "relative",
ylab = "Item Frequency (Relative)")
```

```
itemsets = apriori(Groceries, parameter = list(minlen=2, maxlen=2,support=0.02,
target="frequent itemsets"))
summary(itemsets)
inspect(itemsets)
itemsets_3 = apriori(Groceries, parameter = list(minlen=3, maxlen=3,support=0.02,
target="frequent itemsets"))
summary(itemsets_3)
```

```
inspect(itemsets_3)
```

## Output:



R Console

```
3      3      3      3      3      3

summary of quality measures:
  support      count
Min.   :0.02227   Min.   :219.0
1st Qu.:0.02250   1st Qu.:221.2
Median :0.02272   Median :223.5
Mean   :0.02272   Mean   :223.5
3rd Qu.:0.02295   3rd Qu.:225.8
Max.   :0.02318   Max.   :228.0

includes transaction ID lists: FALSE

mining info:
  data ntransactions support confidence
Groceries      9835      0.02          1

apriori(data = Groceries, parameter = list(minlen = 3, maxlen = 3, support = 0$
>
>
> inspect(itemsets_3)
  items                                     support      count
[1] {root vegetables, other vegetables, whole milk} 0.02318251 228
[2] {other vegetables, whole milk, yogurt}          0.02226741 219
> |
```