

Ex.No. : 1 Vectors and List operations in R

Creating a vector

```
X <- c(1, 4, 5, 2, 6, 7)
print('using c function')
print(X)
Y <- seq(1, 10, length.out = 5)
print('using seq() function')
print(Y)
Z <- 5:10
print('using colon')
print(Z)
```

Accessing vector elements

```
X <- c(2, 5, 8, 1, 2)
print('using Subscript operator')
print(X[2])
Y <- c(4, 5, 2, 1, 7)
print('using c function')
print(Y[c(4, 1)])
Z <- c(5, 2, 1, 4, 4, 3)
print('Logical indexing')
print(Z[Z>3])
```

Modifying a vector

```
X <- c(2, 5, 1, 7, 8, 2)

# modify a specific element
X[3] <- 11
print('Using subscript operator')
print(X)
```

```
# Modify using different logics.
X[X>9] <- 0
print('Logical indexing')
print(X)
```

```
# Modify by specifying the position or elements.
X <- X[c(5, 2, 1)]
print('using c function')
print(X)
```

Deleting a vector

```
Creating a vector
X <- c(5, 2, 1, 6)
# Deleting a vector
X <- NULL
```

```
print('Deleted vector')
print(X)
```

Arithmetic operations

```
#Creating Vectors
```

```
X <- c(5, 2, 5, 1, 51, 2)
```

```
Y <- c(7, 9, 1, 5, 2, 1)
```

```
# Addition
```

```
Z <- X + Y
```

```
print('Addition')
```

```
print(Z)
```

```
# Subtraction
```

```
S <- X - Y
```

```
print('Subtraction')
```

```
print(S)
```

```
# Multiplication
```

```
M <- X * Y
```

```
print('Multiplication')
```

```
print(M)
```

```
# Division
```

```
D <- X / Y
```

```
print('Division')
```

```
print(D)
```

Sorting of Vectors

```
#Creating a Vector
```

```
X <- c(5, 2, 5, 1, 51, 2)
```

```
# Sort in ascending order
```

```
A <- sort(X)
```

```
print('sorting done in ascending order')
```

```
print(A)
```

```
# sort in descending order.
```

```
B <- sort(X, decreasing = TRUE)
```

```
print('sorting done in descending order')
```

```
print(B)
```

```
#II Creating List
```

```
#Creating Vectors
```

```
vec1 <- c(1, 2, 3)
```

```
vec2 <- c(TRUE, FALSE)
```

```
# Creating a list of Vectors
listt = list(vec1, vec2)
```

```
# Printing List
print (listt)
```

Adding elements to a list

```
# Creating Vectors
vec1 <- c(1, 2, 3)
vec2 <- c(TRUE, FALSE)
```

```
# Creating list of Vectors
lst = list(vec1, vec2)
```

```
# Creating a new Vector
vec3 <- c(1 + 3i)
```

```
# Adding Vector to list
lst[[3]]<- vec3
```

```
# Printing List
print (lst)
```

```
# determine the length of list
len <- length(lst)
```

```
# Creating new Vector
vec3 <- c(0.5, 2 + 2i)
```

```
# Using for loop to add elements
for( i in 1:2)
{
```

```
    # Adding vec to list
    lst[[len + i]]<- vec3
}
print (lst)
```

Removing elements from a list

```
lst[[2]]<-NULL
print ("Modified List")
print (lst)
```

Modifying elements in a list

```
# Creating Vectors
```

```
vec1 <- c(1, 2, 3)
```

```
vec2 <- c(TRUE, FALSE)
```

```
# Creating list of Vectors
```

```
lst = list(vec1, vec2)
```

```
print ("original list")
```

```
print (lst)
```

```
# Modifying List element
```

```
lst[[2]]<-c("TEACH", "CODING")
```

```
print ("Modified List")
```

```
print (lst)
```

Merging two lists

```
# R program to merge two lists of Vectors
```

```
# Creating 1st list
```

```
list_data1 <- list(c(1:3), c(TRUE, FALSE))
```

```
# Creating 2nd list
```

```
list_data2 <- list(c(0.1, 3.4))
```

```
print("First List")
```

```
print (list_data1)
```

```
print ("Second List")
```

```
print (list_data2)
```

```
print("Merged List")
```

```
# Merging Lists
```

```
merged_list <- c(list_data1, list_data2)
```

```
print (merged_list)
```

OUTPUT

#Creating Vector

```
[1] 1 4 5 2 6 7
```

```
[1] 1.00 3.25 5.50 7.75 10.00
```

```
[1] 5 6 7 8 9 10
```

#Accessing Vector Element

```
[1] 5
```

```
[1] 1 4
```

```
[1] 5 4 4
```

#Modifying Vector

```
[1] 2 5 11 7 8 2
```

```
[1] 2 5 0 7 8 2
```

```
[1] 8 5 2
```

#Deleting Vector

```
NULL
```

#Arithmetic Operation

```
[1] "Addition"
```

```
[1] 12 11 6 6 53 3
```

```
[1] "Subtraction"
```

```
[1] -2 -7 4 -4 49 1
```

```
[1] "Multiplication"
```

```
[1] 35 18 5 5 102 2
```

```
[1] "Division"
```

```
[1] 0.7142857 0.2222222 5.0000000 0.2000000 25.5000000 2.0000000
```

#Sorting Value

```
[1] "sorting done in ascending order"
```

```
[1] 1 2 2 5 5 51
```

```
[1] "sorting done in descending order"
```

```
[1] 51 5 5 2 2 1
```

#Adding element to the list

```
[[1]]
```

```
[1] 1 2 3
```

```
[[2]]
```

```
[1] TRUE FALSE
```

```
[[3]]
```

```
[1] 1+3i
```

```
[[4]]  
[1] 0.5+0i 2.0+2i
```

```
[[5]]  
[1] 0.5+0i 2.0+2i
```

#Modifying element in the list

```
[1] "Modified List"  
[[1]]  
[1] 1 2 3
```

```
[[2]]  
[1] "TEACH" "CODING"
```

#Merging Two List

```
[1] "First List"  
[[1]]  
[1] 1 2 3
```

```
[[2]]  
[1] TRUE FALSE
```

```
[1] "Second List"  
[[1]]  
[1] 0.1 3.4
```

```
[1] "Merged List"  
[[1]]  
[1] 1 2 3
```

```
[[2]]  
[1] TRUE FALSE
```

```
[[3]]  
[1] 0.1 3.4
```

Ex.No. : 2 Matrices and Array operations in R

#Creation of an Array

```
vector1 <- c(1, 2, 3)
vector2 <- c(10, 15, 3, 11, 16, 12)
result <- array(c(vector1, vector2), dim = c(3, 3, 2))
print(result)
```

Operations on Arrays

Naming columns and rows

```
vector1 <- c(1, 2, 3)
vector2 <- c(10, 15, 3, 11, 16, 12)
column.names <- c("COL1", "COL2", "COL3")
row.names <- c("R1", "R2", "R3")
matrix.names <- c("Matrix.NO1", "Matrix.NO2")
result <- array(c(vector1, vector2), dim = c(3, 3, 2),
               dimnames = list(row.names, column.names, matrix.names))
print(result)
```

Manipulating array elements

```
vector1 <- c(1, 2, 3)
vector2 <- c(4, 6, 8, 0, 2, 4)
array1 <- array(c(vector1, vector2), dim = c(3, 3, 2))
vector3 <- c(3, 2, 1)
vector4 <- c(2, 4, 6, 8, 3, 5)
array2 <- array(c(vector3, vector4), dim = c(3, 3, 2))
matrix1 <- array1[,,2]
matrix2 <- array2[,,2]
result <- matrix1 + matrix2
print(result)
result <- matrix1 - matrix2
print(result)
result <- matrix1 * matrix2
result <- matrix1 / matrix2
```

Accessing Array elements

```
vector1 <- c(1, 2, 3)
vector2 <- c(10, 15, 3, 11, 16, 12)
column.names <- c("COLUMN1", "COLUMN2", "COLUMN3")
row.names <- c("ROW1", "ROW2", "ROW3")
matrix.names <- c("Matrix.NO1", "Matrix.NO2")
# taking vector as input
result <- array(c(vector1, vector2), dim = c(3, 3, 2),
               dimnames = list(row.names, column.names, matrix.names))
print(result)
```

```
# print third row of second matrix
print(result[3,,2])
```

Calculation across array element

```
vector1 <- c(3, 2, 1)
vector2 <- c(2, 4, 6, 8, 0, 1)
new.array <- array(c(vector1, vector2), dim = c(3, 3, 2))
print(new.array)
```

```
# using apply and calculate the sum of rows in matrices
result <- apply(new.array, c(1), sum)
print(result)
```

#Creation of Matrices:

Elements are arranged sequentially by row.

```
M <- matrix(c(3:14), nrow = 4, byrow = TRUE)
print(M)
```

Elements are arranged sequentially by column.

```
N <- matrix(c(3:14), nrow = 4, byrow = FALSE)
print(N)
```

Define the column and row names.

```
rownames = c("row1", "row2", "row3", "row4")
colnames = c("col1", "col2", "col3")
```

```
P <- matrix(c(3:14), nrow = 4, byrow = TRUE, dimnames = list(rownames, colnames))
print(P)
```

Operations on Matrices

Matrices Addition

Creating 1st Matrix

```
B = matrix(c(1, 2, 3, 4, 5, 6), nrow = 2, ncol = 3)
```

Creating 2nd Matrix

```
C = matrix(c(7, 8, 9, 10, 11, 12), nrow = 2, ncol = 3)
```

Getting number of rows and columns

```
num_of_rows = nrow(B)
```

```
num_of_cols = ncol(B)
```

Creating matrix to store results


```
sum = matrix(, nrow = num_of_rows, ncol = num_of_cols)
```

```
# Printing Original matrices
```

```
print(B)
```

```
print(C)
```

```
# Creating 1st Matrix
```

```
B = matrix(c(1, 2, 3, 4, 5, 6), nrow = 2, ncol = 3)
```

```
# Creating 2nd Matrix
```

```
C = matrix(c(7, 8, 9, 10, 11, 12), nrow = 2, ncol = 3)
```

```
# Getting number of rows and columns
```

```
num_of_rows = nrow(B)
```

```
num_of_cols = ncol(B)
```

```
# Creating matrix to store results
```

```
diff = matrix(, nrow = num_of_rows, ncol = num_of_cols)
```

```
# Printing Original matrices
```

```
print(B)
```

```
print(C)
```

```
# Calculating diff of matrices
```

```
for(row in 1:num_of_rows)
```

```
{
```

```
  for(col in 1:num_of_cols)
```

```
  {
```

```
    diff[row, col] <- B[row, col] - C[row, col]
```

```
  }
```

```
}
```

```
# Printing resultant matrix
```

```
print(diff)
```

```
# Calculating product of matrices
```

```
for(row in 1:num_of_rows)
```

```
{
```

```
  for(col in 1:num_of_cols)
```

```
  {
```

```
    prod[row, col] <- B[row, col] * C[row, col]
```

```
  }
```

```
}
```

```
# Printing resultant matrix
```

```
print(prod)
```

```
# R program for matrix multiplication
```

```
# using '*' operator

# Creating 1st Matrix
B = matrix(c(1, 2 + 3i, 5.4), nrow = 1, ncol = 3)

# Creating 2nd Matrix
C = matrix(c(2, 1i, 0.1), nrow = 1, ncol = 3)

# Printing the resultant matrix
print (B * C)
```

Matrices Division

```
# Creating 1st Matrix
B = matrix(c(1, 2, 3, 4, 5, 6), nrow = 2, ncol = 3)

# Creating 2nd Matrix
C = matrix(c(7, 8, 9, 10, 11, 12), nrow = 2, ncol = 3)

# Getting number of rows and columns
num_of_rows = nrow(B)
num_of_cols = ncol(B)

# Creating matrix to store results
div = matrix(, nrow = num_of_rows, ncol = num_of_cols)

# Printing Original matrices
print(B)
print(C)
```

```
# Calculating product of matrices
for(row in 1:num_of_rows)
{
  for(col in 1:num_of_cols)
  {
    div[row, col] <- B[row, col] / C[row, col]
  }
}

# Printing resultant matrix
print(div)

# Creating 1st Matrix
B = matrix(c(4, 6i, -1), nrow = 1, ncol = 3)
# Creating 2nd Matrix
C = matrix(c(2, 2i, 0), nrow = 1, ncol = 3)
# Printing the resultant matrix
print (B / C)
```

OUTPUT

#Naming Column and Rows

Matrix.NO1

```
COL1 COL2 COL3
R1  1  10  11
R2  2  15  16
R3  3   3  12
```

, , Matrix.NO2

```
COL1 COL2 COL3
R1  1  10  11
R2  2  15  16
R3  3   3  12
```

#Manipulating array Elements

```
[,1] [,2] [,3]
[1,]  4   6   8
[2,]  4  10   5
[3,]  4  14   9
```

```
[,1] [,2] [,3]
[1,] -2   2 -8
[2,]  0   2 -1
[3,]  2   2 -1
```

#Accessing Array Elements

print third row of second matrix

```
COLUMN1 COLUMN2 COLUMN3
      3         3        12
```

#Calculation across array elements

using apply and calculate the sum of rows in matrices

```
[1] 26 12 16
```

#Matrices Operation

#Addition

```
[,1] [,2] [,3]
[1,]  8  12  16
[2,] 10  14  18
```

#Subtraction

```
[,1] [,2] [,3]
[1,]  6   6   6
[2,]  6   6   6
```

#Multiplication

```
[,1] [,2] [,3]
[1,]  7  27  55
```

[2,] 16 40 72

#Division

	[,1]	[,2]	[,3]
[1,]	0.1428571	0.3333333	0.4545455
[2,]	0.2500000	0.4000000	0.5000000

	[,1]	[,2]	[,3]
[1,]	2+0i	3+0i	-Inf+NaNi

Ex.No. : 3 Saving, loading and removing R datastructures

#Method 1: Using save. image and load method

#Syntax:

```
save.image(file = ".RData")
```

Arguments :

file – name of the file where the R object is saved to or read from.

```
obj1<-c(1:15)
```

```
obj2<-FALSE
```

```
Obj3<-"Welcome to R Environment"
```

```
save.image("ex1.Rdata")
```

#Syntax:

```
Load(path)
```

```
load("ex1.Rdata")
```

#Method 2: Using saveRDS and readRDS method

Syntax:

```
saveRDS(object, file = "")
```

```
obj1<-c(1:15)
```

```
obj2<-FALSE
```

```
Obj3<-"Welcome to R Environment"
```

```
saveRDS(obj1,file="ex1int.Rdata")
```

```
print("Data object")
```

```
readRDS("ex1int.Rdata")
```

#Method 3: Using the save and load method

#Syntax:

```
save(objects, file)
```

```
save(obj1, obj3, file ="tempworkspaceobj.RData")
```

```
load("tempworkspaceobj.RData")
```

OUTPUT

```
[1] "Data Object"
```

```
[1] 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15
```

Ex.No. : 4 Visualizing numeric data – scatterplot, boxplot, piechart, histograms

```
us_cars<-read.csv("S:/ML_Dataset/usedcars.csv",stringsAsFactors = FALSE)
str(us_cars)

#Box Plot
boxplot(us_cars$price,main="Used cars price",ylab="price($)",col="blue",border="red")
boxplot(us_cars$mileage,main="Used cars mileage", ylab=" ",col="red",border="brown")
boxplot(us_cars[,0:6],main='Used Cars')

#Bar plot
barplot(us_cars$mileage,main='Used Cars',
        xlab="Mileage",horiz=FALSE,col="blue")

#Histogram
hist(us_cars$mileage,main="Used Cars Mileage",
     xlab='Mileage',col="yellow",border="red")
hist(us_cars$price,main="Used Cars Price",
     xlab="price",col="Green", border="red")

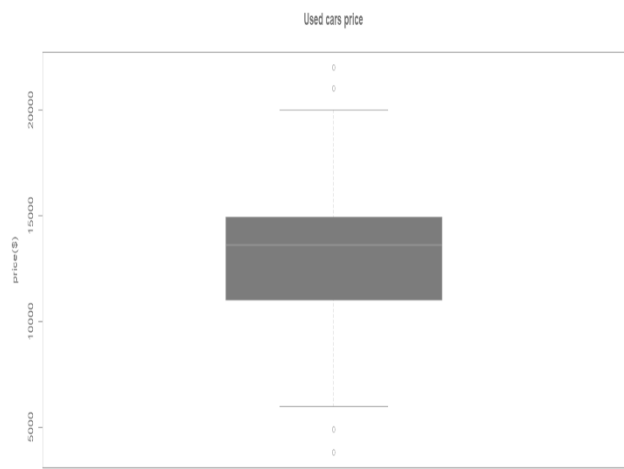
#scatter plot
plot(us_cars$price,us_cars$mileage,main="Scatterplot",
     xlab="Price",ylab="Mileage",pch=12,col='green')

#pie
n=12
pie(rep(1,n),col=rainbow(n))
pie(rep(1,n),col=heat.colors(n))
pie(rep(1,n),col=terrain.colors(n))

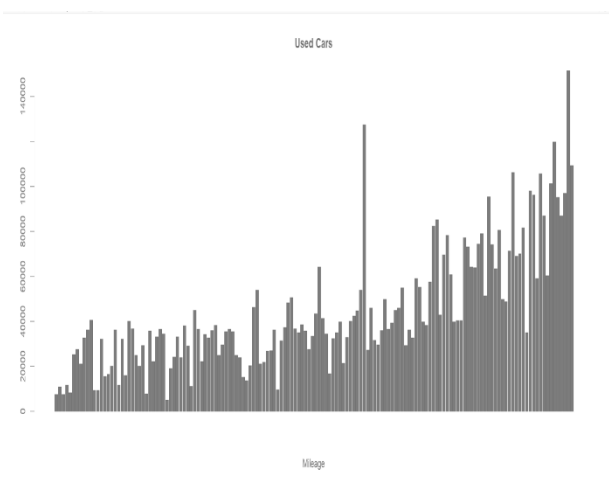
#3d Graphs
cone<-function(x,y)
{
  sqrt(x^2 + y^2)
}
#prepare variables.
x<-y<-seq(-1,1,length=30)
z<-outer(x,y,cone)
persp(x,y,z,main="Perspective Plot of a Cone",
     zlab="Height",
     theta=30,phi=15,
     col="green",shade=0.4)
```

OUTPUT

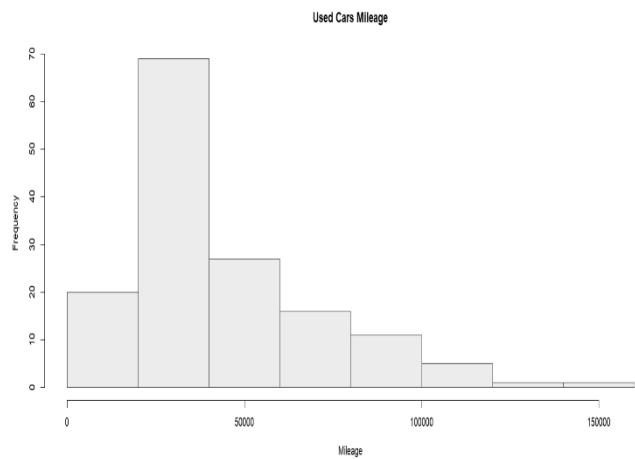
BoxPlot



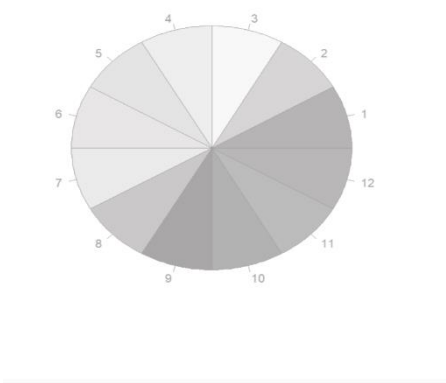
BarPlot



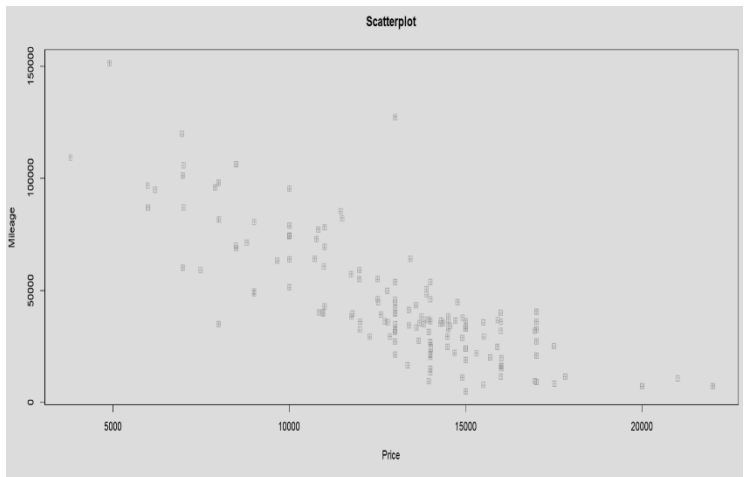
Histogram



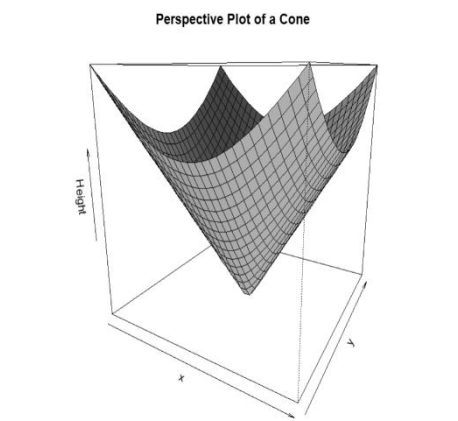
Piechart



ScatterPlot



3D Graphs



Ex.No. : 5 Measuring Central Tendency

```
usedcars <- read.csv("D:/RPractical_progmas/usedcars.csv", stringsAsFactors = FALSE)
```

```
str(usedcars)
print(head(usedcars))
max1=max(usedcars$price)
print(max1)
```

```
min1=min(usedcars$price)
print(min1)
```

```
range=max1-min1
print(range)
r=range(usedcars$price)
print(r)
```

```
variance=var(usedcars$price)
print(variance)
```

```
std=sd(usedcars$price)
```

```
quartiles=quantile(usedcars$price)
print(quartiles)
```

```
IQR=IQR(usedcars$price)
print(IQR)
```

```
summary=summary(usedcars$price)
print(summary)
```

```
summary(usedcars)
```

OUTPUT

	year	model	price	mileage	color	transmission
1	2011	SEL	21992	7413	Yellow	AUTO
2	2011	SEL	20995	10926	Gray	AUTO
3	2011	SEL	19995	7351	Silver	AUTO
4	2011	SEL	17809	11613	Gray	AUTO
5	2012	SE	17500	8367	White	AUTO
6	2010	SEL	17495	25125	Silver	AUTO

```
#Max
[1] 21992
```


#Min

[1] 3800

#Range

[1] 18192

[1] 3800 21992

#variance

[1] 9749892

#Standard Deviation

[1] 3122.482

#Quartiles

0%	25%	50%	75%	100%
3800.0	10995.0	13591.5	14904.5	21992.0

#IQR

[1] 3909.5

#Summary

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
3800	10995	13592	12962	14904	21992

#Summary used Cars

year	model	price	mileage
Min. :2000	Length:150	Min. : 3800	Min. : 4867
1st Qu.:2008	Class :character	1st Qu.:10995	1st Qu.: 27200
Median :2009	Mode :character	Median :13592	Median : 36385
Mean :2009		Mean :12962	Mean : 44261
3rd Qu.:2010		3rd Qu.:14904	3rd Qu.: 55125
Max. :2012		Max. :21992	Max. :151479
color	transmission		
Length:150	Length:150		
Class :character	Class :character		
Mode :character	Mode :character		

Ex.No. : 6 Data Pre-Processing Methods

```
mydata<-read.csv("S:/ML_Dataset/mysampleddata.csv")
mydata
```

```
#dealing with missing values
```

```
mydata$age<-ifelse(is.na(mydata$age),ave(mydata$age,FUN = function(x)
mean(x,na.rm=TRUE)),mydata$age)
mydata
mydata$salary<-ifelse(is.na(mydata$salary),ave(mydata$salary, FUN = function(x)
mean(x,na.rm=TRUE)),mydata$salary)
mydata
mydata$age<-as.numeric(format(round(mydata$age,0)))
```

```
#Dealing with categorical Data
```

```
mydata$nation<-factor(mydata$nation,levels=c('India','Russia','Germany'),labels=c(1,2,3))
mydata$purchased_item<-factor(mydata$purchased_item,levels = c('No','Yes'),labels=c(0,1))
mydata
```

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

```
library(caTools)
```

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

```
Split<-sample.split(mydata$purchased_item,SplitRatio=0.8)
```

```
training_set<-subset(mydata,Split==TRUE)
```

```
test_set<-subset(mydata,Split==FALSE)
```

```
training_set[,3:4]<-scale(training_set[,3:4])
```

```
test_set[,3:4]<-scale(test_set[,3:4])
```

OUTPUT

#Dealing with missing values

```
> mydata
  nation purchased_item age salary
1   India             No  25  35000
2  Russia             Yes NA  40000
3 Germany             No  50  54000
4  Russia             No  35     NA
5 Germany             Yes  40  60000
6   India             Yes  35  58000
7  Russia             No  NA  52000
8   India             Yes  48     NA
9 Germany             No  50  83000
10  India             Yes  37     NA
11 Germany            No  21  24000
12  India             Yes  NA  60000
13 Russia             No  63  70000
14 Germany            Yes  26  36000
15  India             No  45  40000
> mydata$age<-ifelse(is.na(mydata$age),ave(n
> mydata
  nation purchased_item age salary
1   India             No 25.00000  35000
2  Russia             Yes 39.58333  40000
3 Germany             No 50.00000  54000
4  Russia             No 35.00000     NA
5 Germany             Yes 40.00000  60000
6   India             Yes 35.00000  58000
7  Russia             No 39.58333  52000
8   India             Yes 48.00000     NA
9 Germany             No 50.00000  83000
10  India             Yes 37.00000     NA
11 Germany            No 21.00000  24000
12  India             Yes 39.58333  60000
13 Russia             No 63.00000  70000
14 Germany            Yes 26.00000  36000
15  India             No 45.00000  40000
```

#Dealing with categorical Data

```
> mydata
  nation purchased_item age salary
1      1             0  25  35000
2      2             1  40  40000
3      3             0  50  54000
4      2             0  35     NA
5      3             1  40  60000
6      1             1  35  58000
7      2             0  40  52000
8      1             1  48     NA
9      3             0  50  83000
10     1             1  37     NA
11     3             0  21  24000
12     1             1  40  60000
13     2             0  63  70000
14     3             1  26  36000
15     1             0  45  40000
>
```

Ex.No. : 7 Build model using K-Nearest Neighbour

```
loan <- read.csv("credit_data.csv")
```

```
str(loan)
```

#Data Cleaning

```
loan.subset <-
```

```
loan[c('Creditability','Age..years.','Sex...Marital.Status','Occupation','Account.Balance','Credit  
.Amount','Length.of.current.employment','Purpose')]
```

```
str(loan.subset)
```

#Data Normalization

```
head(loan.subset)
```

#Normalization

```
normalize <- function(x) {
```

```
  return ((x - min(x)) / (max(x) - min(x))) }
```

```
loan.subset.n <- as.data.frame(lapply(loan.subset[,2:8], normalize))
```

```
head(loan.subset.n)
```

#Data Splicing

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

```
dat.d <- sample(1:nrow(loan.subset.n),size=nrow(loan.subset.n)*0.7,replace = FALSE)
```

```
train.loan <- loan.subset[dat.d,] # 70% training data
```

```
test.loan <- loan.subset[-dat.d,] # remaining 30% test data
```

```
#Creating separate dataframe for 'Creditability' feature which is our target.
```

```
train.loan_labels <- loan.subset[dat.d,1]
```

```
test.loan_labels <- loan.subset[-dat.d,1]
```

#Building a Machine Learning model

```
install.packages('class')
```

```
library(class)
```

```
NROW(train.loan_labels)
```

```
knn.26 <- knn(train=train.loan, test=test.loan, cl=train.loan_labels, k=26)
```

```
knn.27 <- knn(train=train.loan, test=test.loan, cl=train.loan_labels, k=27)
```

#Model Evaluation

```
ACC.26 <- 100 * sum(test.loan_labels == knn.26)/NROW(test.loan_labels)
```

```
ACC.27 <- 100 * sum(test.loan_labels == knn.27)/NROW(test.loan_labels)
```

```
ACC.26
```

```
ACC.27
```

```
table(knn.26 ,test.loan_labels)
```

```
table(knn.27 ,test.loan_labels)
```

```
install.packages('caret')
```

```
library(caret)
```

```
confusionMatrix(table(knn.26 ,test.loan_labels))
```

#Optimization

```
i=1
```

```
k.optm=1
```

```
for (i in 1:28){
```

```
  knn.mod <- knn(train=train.loan, test=test.loan, cl=train.loan_labels, k=i)
```

```
  k.optm[i] <- 100 * sum(test.loan_labels == knn.mod)/NROW(test.loan_labels)
```

```
  k=i
```

```
  cat(k, '=', k.optm[i], "\n")
```

```
}
```

#Accuracy plot

```
plot(k.optm, type="b", xlab="K- Value", ylab="Accuracy level")
```

OUTPUT

#DataCleaning

```
> str(loan.subset)
'data.frame':  1000 obs. of  8 variables:
 $ Creditability      : int  1 1 1 1 1 1 1 1 1 1 ...
 $ Age..years.        : int  21 36 23 39 38 48 39 40 65 23 ...
 $ Sex...Marital.Status : int  2 3 2 3 3 3 3 3 2 2 ...
 $ Occupation         : int  3 3 2 2 2 2 2 2 1 1 ...
 $ Account.Balance     : int  1 1 2 1 1 1 1 1 4 2 ...
 $ Credit.Amount       : int  1049 2799 841 2122 2171 2241 3398 1361 1098 3758 ...
 $ Length.of.current.employment: int  2 3 4 3 3 2 4 2 1 1 ...
 $ Purpose            : int  2 0 9 0 0 0 0 0 3 3 ...
```

Data Normalization

```
> head(loan.subset)
  Creditability Age..years. Sex...Marital.Status Occupation Account.Balance Credit.Amount Length.of.current.employment Purpose
1             1           21                2          3             1           1049                2             2
2             1           36                3          3             1           2799                3             0
3             1           23                2          2             2            841                4             9
4             1           39                3          2             1           2122                3             0
5             1           38                3          2             1           2171                3             0
6             1           48                3          2             1           2241                2             0

> normalize<-function(x){
+   return((x-min(x))/(max(x)-min(x)))
+ }
> loan.subset.n<-as.data.frame(lapply(loan.subset[,2:8],normalize))
> head(loan.subset.n)
  Age..years. Sex...Marital.Status Occupation Account.Balance Credit.Amount Length.of.current.employment Purpose
1 0.03571429 0.3333333 0.6666667 0.0000000 0.04396390 0.25 0.2
2 0.30357143 0.6666667 0.6666667 0.0000000 0.14025531 0.50 0.0
3 0.07142857 0.3333333 0.3333333 0.3333333 0.03251898 0.75 0.9
4 0.35714286 0.6666667 0.3333333 0.0000000 0.10300429 0.50 0.0
5 0.33928571 0.6666667 0.3333333 0.0000000 0.10570045 0.50 0.0
6 0.51785714 0.6666667 0.3333333 0.0000000 0.10955211 0.25 0.0
```

Building a Machine Learning model

```
[1] 700
```

#Model Evaluation

```
[1] 69
```

```
[1] 69
```

```
test.loan_labels
```

```
knn.26 0 1
       0 8 6
       1 87 199
```

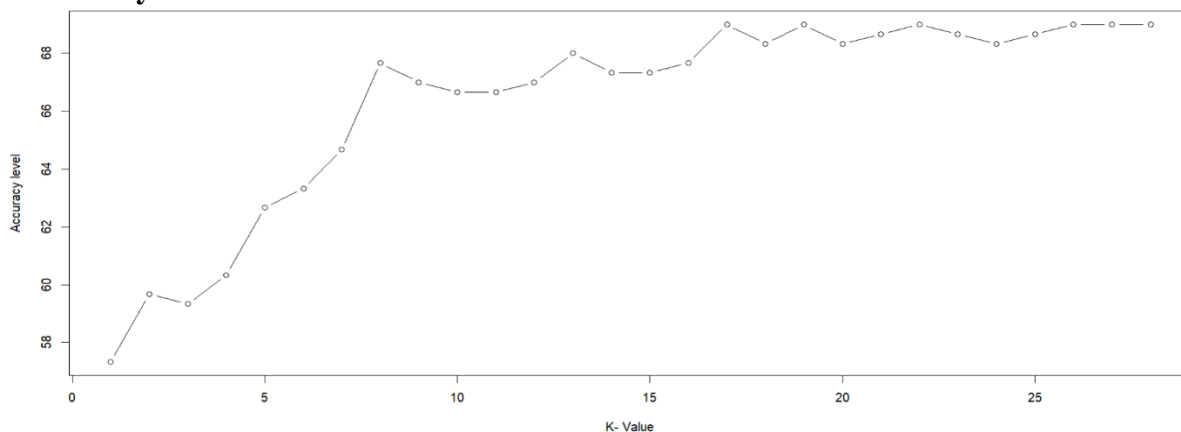
```
test.loan_labels
```

```
knn.27 0 1
       0 8 6
       1 87 199
       2
```

#Optimization

1 = 57.33333 2 = 59.66667 3 = 59.33333 4 = 60.33333 5 = 62.66667 6 = 63.33333 7 = 64.66667 8 = 67.66667 9 = 67 10 = 66.66667 11 = 66.66667 12 = 67 13 = 68 14 = 67.33333 15 = 67.33333 16 = 67.66667 17 = 69 18 = 68.33333 19 = 69 20 = 68.33333 21 = 68.66667 22 = 69 23 = 68.66667 24 = 68.33333 25 = 68.66667 26 = 69 27 = 69 28 = 69

#Accuracy Plot



Ex.No. : 8 Decision Tree

```
mushrooms <- read.csv("mushroom.csv", header = FALSE)

head(mushrooms)

colnames(mushrooms) <-
c("Class", "cap.shape", "cap.surface", "cap.color", "bruises", "odor", "gill.attachment", "gill.spacing",
  "gill.size", "gill.color", "stalk.shape", "stalk.root", "stalk.surface.above.ring", "stalk.surface.below.ring",
  "stalk.color.above.ring", "stalk.color.below.ring", "veil.type", "veil.color", "ring.number", "ring.type", "print",
  "population", "habitat")

head(mushrooms)

# Define the factor names for "Class"
levels(mushrooms$Class) <- c("Edible", "Poisonous")

# Define the factor names for "odor"
levels(mushrooms$odor) <-
c("Almonds", "Anise", "Creosote", "Fishy", "Foul", "Musty", "None", "Pungent", "Spicy")

# Define the factor names for "print"
levels(mushrooms$print) <-
c("Black", "Brown", "Buff", "Chocolate", "Green", "Orange", "Purple", "White", "Yellow")

head(mushrooms)

# Import our required libraries
library(rpart)
library(rpart.plot)

# Create a classification decision tree using "Class" as the variable we want to predict and
everything else as its predictors.
myDecisionTree <- rpart(Class ~ ., data = mushrooms, method = "class")

# Print out a summary of our created model.
print(myDecisionTree)

rpart.plot(myDecisionTree, type = 3, extra = 2, under = TRUE, faclen=5, cex = .75)

newCase <- mushrooms[10,-1]

newCase

predict(myDecisionTree, newCase, type = "class")
```

```

train_ind <- sample(c(1:nrow(mushrooms)), size = 10)

## 75% of the sample size

n <- nrow(mushrooms)

smp_size <- floor(0.75 * n)

## set the seed to make your partition reproducible

set.seed(123)

train_ind <- sample(c(1:n), size = smp_size)

mushrooms_train <- mushrooms[train_ind, ]
mushrooms_test <- mushrooms[-train_ind, ]

newDT <- rpart(Class ~ ., data = mushrooms_train, method = "class")
result <- predict(newDT, mushrooms_test[, -1], type = "class")

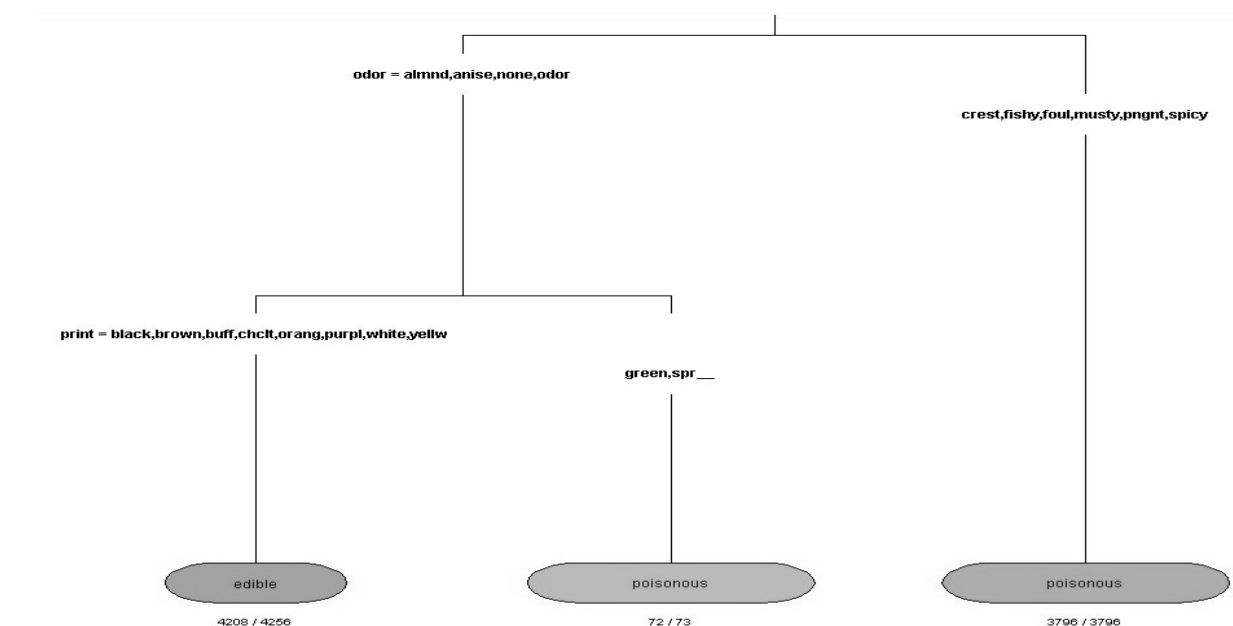
head(result)

head(mushrooms_test$Class)

table(mushrooms_test$Class, result)

```

OUTPUT



Ex.No. : 9 Develop a model using logistic regression

```
library(tidyverse)
library(modelr)
library(broom)

#Install ISLR Package
install.packages('ISLR')

#Load ISLR Package
library('ISLR')

# Load data
(mydata <- as_tibble(ISLR::Default))

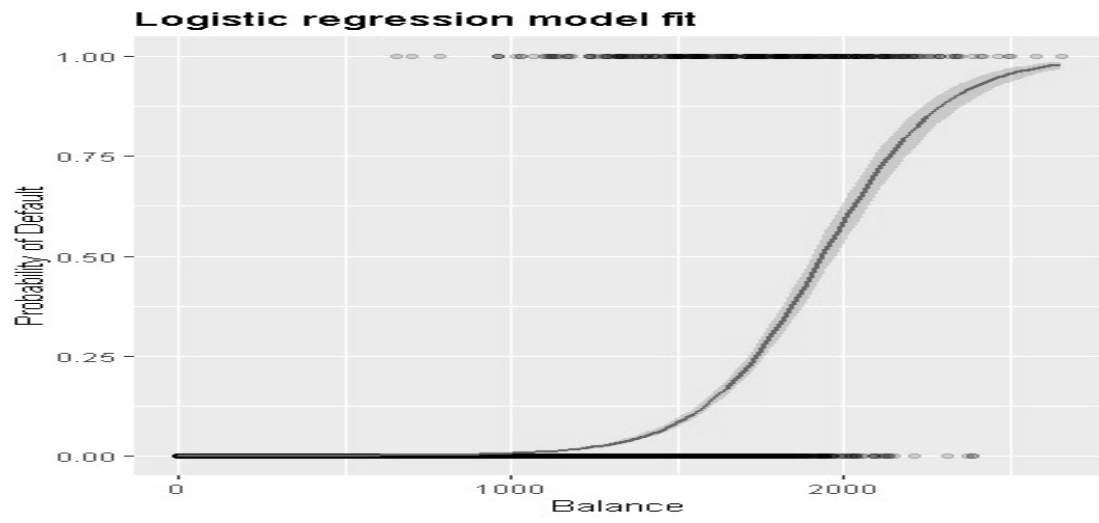
#Checking for NA values
sum(is.na(mydata))

#Creating the Training and Testing data set
sample <- sample(c(TRUE, FALSE), nrow(mydata), replace = T, prob = c(0.6,0.4))
train <- mydata[sample, ]
test <- mydata[!sample, ]

#Fitting a logistic regression model
logmodel <- glm(default ~ balance, family = "binomial", data = train)
#Plotting a graph: Probability of default Vs Balance
mydata %>%
mutate(prob = ifelse(default == "Yes", 1, 0)) %>%
ggplot(aes(balance, prob)) +
geom_point(alpha = .15) +
geom_smooth(method = "glm", method.args = list(family = "binomial")) +
ggtitle("Logistic regression model fit") +
xlab("Balance") +
ylab("Probability of Default")

#Summary of the Logistic Regression
summary(logmodel)
```

OUTPUT



Ex.No. : 10 Identify patterns using Association Rules

```
# load the grocery data into a sparse matrix
library(arules)
groceries <- read.transactions("groceries.csv", sep = ",")
summary(groceries)

# look at the first five transactions
inspect(groceries[1:5])

# examine the frequency of items
itemFrequency(groceries[, 1:3])

# plot the frequency of items
itemFrequencyPlot(groceries, support = 0.1)
itemFrequencyPlot(groceries, topN = 20)

# a visualization of the sparse matrix for the first five transactions
image(groceries[1:5])

# visualization of a random sample of 100 transactions
image(sample(groceries, 100))

## Step 3: Training a model on the data ----
library(arules)

# default settings result in zero rules learned
apriori(groceries)

# set better support and confidence levels to learn more rules
groceryrules <- apriori(groceries, parameter = list(support =
  0.006, confidence = 0.25, minlen = 2))
groceryrules

## Step 4: Evaluating model performance ----
# summary of grocery association rules
summary(groceryrules)

# look at the first three rules
inspect(groceryrules[1:3])

## Step 5: Improving model performance ----

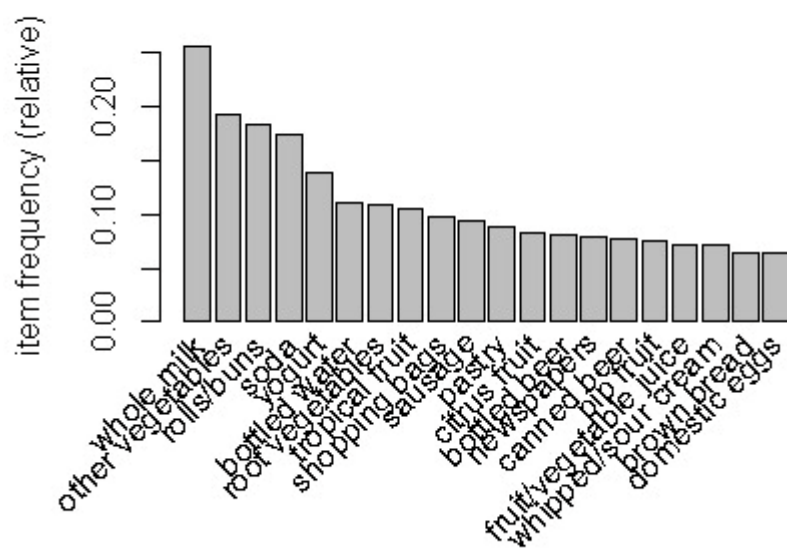
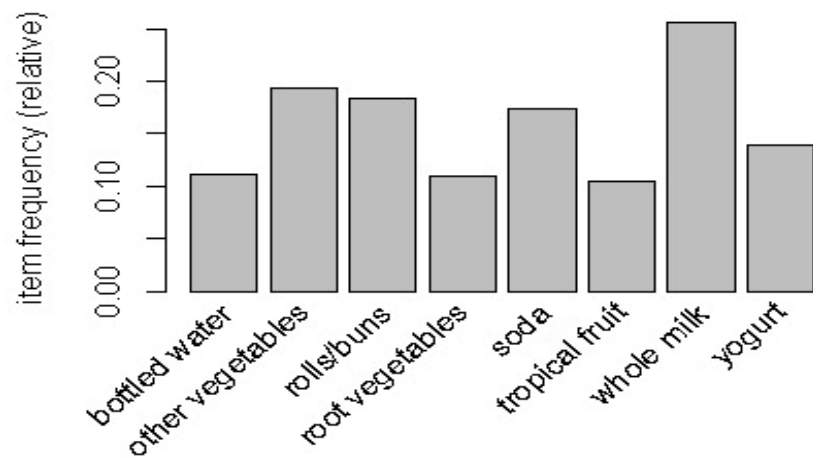
# sorting grocery rules by lift
inspect(sort(groceryrules, by = "lift")[1:5])

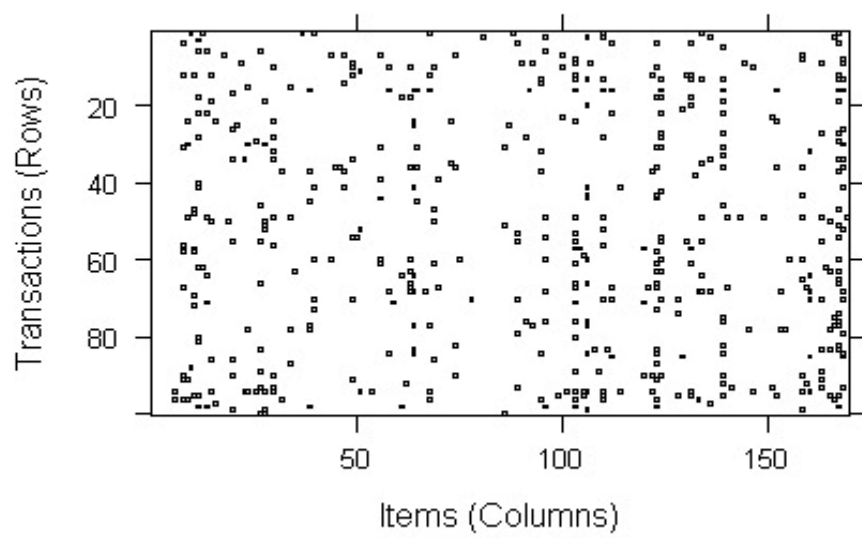
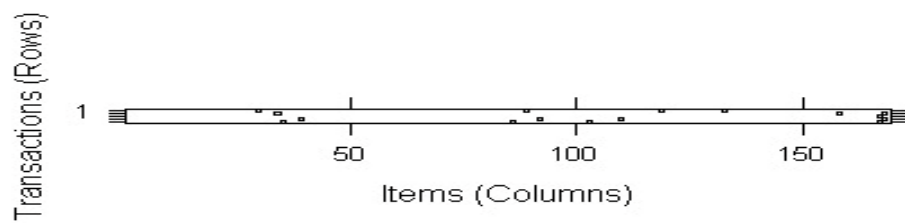
# finding subsets of rules containing any berry items
berryrules <- subset(groceryrules, items %in% "berries")
inspect(berryrules)
```

```
# writing the rules to a CSV file
write(groceryrules, file = "groceryrules.csv",
      sep = ",", quote = TRUE, row.names = FALSE)
```

```
# converting the rule set to a data frame
groceryrules_df <- as(groceryrules, "data.frame")
str(groceryrules_df)
```

OUTPUT





Ex.No. : 11 Identify patterns using Association Rules