

## PROGRAM 8

**Read the following file formats in Python/R:**

- **Comma-separated values**
- **XLSX**
- **ZIP**
- **Plain Text (txt)**
- **JSON**
- **XML**
- **HTML**
- **Images**
- **Hierarchical Data Format**
- **PDF**
- **DOCX**
- **MP3**

### **Execution code: -**

**# Comma-separated values**

```
import pandas as pd
df = pd.read_csv("/home/Loan_Prediction/train.csv")
```

**# XLSX**

```
import pandas as pd
df = pd.read_excel("/home/Loan_Prediction/world_city.xlsx", sheetname = "Invoice")
```

**# ZIP**

```
import zipfile
archive = zipfile.ZipFile('T.zip', 'r')
df = archive.read('train.csv')
```

**# Plain Text (txt)**

```
text_file = open("text.txt", "r")
lines = text_file.read()
```

**# JSON**

```
import pandas as pd
df = pd.read_json("/home/kunal/Downloads/Loan_Prediction/train.json")
```

**# XML**

```
import xml.etree.ElementTree as ET
tree = ET.parse('/home/sunilray/Desktop/2 sigma/train.xml')
root = tree.getroot()
print root.tag
```

## # **HTML**

```
url="https://en.wikipedia.org/wiki/Delhi"  
resp = requests.get(url)  
text = resp.text  
print(text)
```

## # **Images**

```
from scipy import misc  
f = misc.face()  
misc.imsave('face.png', f) # uses the  
Image module (PIL)  
import matplotlib.pyplot as plt  
plt.imshow(f)  
plt.show()
```

## # **Hierarchical Data Format**

```
Import pandas as pd  
t = pd.read_hdf('train.h5')
```

## # **PDF**

```
pdf2txt.py<pdf_file>.pdf
```

## # **DOCX**

```
pip install docx2txt  
import docx2txt  
text = docx2txt.process("file.docx")
```

## # **MP3**

```
from moviepy.editor import  
VideoFileClip  
clip=VideoFileClip('<video_file>.mp3')  
ipython_display(clip)
```

**Output: -**

	Id	Name	Job
0	1	Raju	Full Stack
1	2	Shyam	Frontend
2	3	Ghanshyam	Backend
3	4	Radheshyam	Data Science

	Id	City	Country
0	1	Delhi	India
1	2	Tokyo	Japan
2	3	Thimpu	Bhutan
3	4	Kathmandu	Nepal

.ipynb_checkpoints	19-Mar-20 3:59 PM	File folder	
Images	21-Mar-20 1:53 AM	File folder	
Importing files	21-Mar-20 1:12 AM	File folder	
AV-Importing data files.ipynb	21-Mar-20 1:46 AM	IPYNB File	2,823 KB
Importing files	21-Mar-20 1:12 AM	Compressed (zipp...	146 KB

*"In my previous article, I introduced you to the basics of (RDD / DataFrame / Dataset) and basics of operations (Transformation and Actions) from one of our past hackathons. In this my previous article. I will focus on manipulating RDD in Py (Transformation and Actions)."*

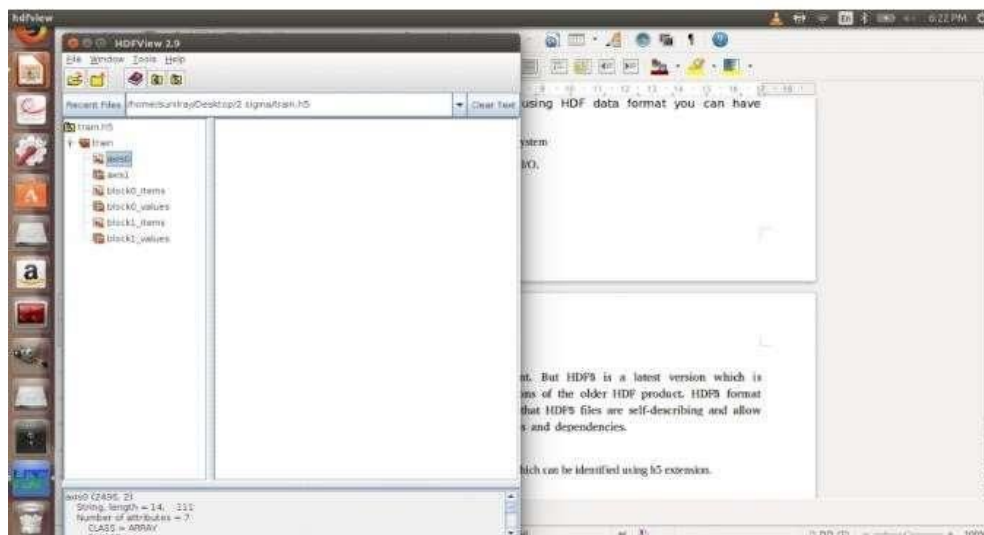
```
<class 'dict'>
```

	Name	Company	Job
0	Aniruddha	Analytics Vidhya	Intern
1	Jill	Google	Full time

```

<!DOCTYPE html>
<html class="client-nojs" lang="en" dir="ltr">
<head>
<meta charset="UTF-8"/>
<title>Delhi - Wikipedia</title>
<script>document.documentElement.className="client-nojs";RLCONF={wgBreakFrames:!1,"wgSeparatorTransformTable":["",""],"wgDigitTransformTable":["",""],"wgDefaultDateFormat":"dmy","wgMonthNames":["","January","February","March","April","May","June","July","August","September","October","November","December"],"wgRequestId":"Xn76cQpAICaAGoNFuaAAAE","wgCSRFnonce":!1,"wgCanonicalNamespace":"","wgCanonicalSpecialPageName":!1,"wgNamespaceNumber":0,"wgPageName":"Delhi","wgTitle":"Delhi","wgCurRevisionId":946313366,"wgRevisionId":946313366,"wgArticleId":37756,"wgIsArticle":!0,"wgIsRedirect":!1,"wgAction":"view","wgUserName":null,"wgUserGroups":[""],"wgCategories":["CS1 errors: missing periodical","Webarchive template wayback links","CS1 maint: archived copy as title","Articles with short description","Wikipedia indefinitely semi-protected pages","Wikipedia indefinitely move-protected pages","Use Indian English from October 2019","All Wikipedia articles written in Indian English"],"Use dmy dates from March 2020","Coordinates on Wikidata","Articles containing potentially dated statements from 2016","All articles containing potentially dated statements","Articles containing Persian-language text","Articles containing potentially dated statements from 2013","All articles with unourced statements","Articles with unourced statements from April 2018","Articles containing potentially dated statements from 2015","Pages using multiple image with auto scaled image

```





## PROGRAM 9

**Compute and print the mean and the standard deviation for each of the 4 measurement columns (i.e., sepal length and width, petal length and width. Compute and print the mean and the standard deviation for each of the 4 measurement columns, separately for each of the three Iris species**

### Execution code: -

```
library(tidyverse)
View(iris)
df <- data.frame(iris)
grp_spc <- group_by(iris,Species)
summarise(iris,cnt=n())
summarise(grp_spc,cnt=n())
summarize(iris,mn = mean(Sepal.Length),sd = sd(Sepal.Length))
summarize(iris,mn = mean(Sepal.Width),sd = sd(Sepal.Width))
summarize(iris,mn = mean(Petal.Length),sd = sd(Petal.Length))
summarize(iris,mn = mean(Petal.Width),sd = sd(Petal.Width))
summarize(grp_spc,mn = mean(Sepal.Length),sd = sd(Sepal.Length))
summarize(grp_spc,mn = mean(Sepal.Width),sd = sd(Sepal.Width))
summarize(grp_spc,mn = mean(Petal.Length),sd = sd(Petal.Length))
summarize(grp_spc,mn = mean(Petal.Width),sd = sd(Petal.Width))
```

### Output: -

```
> library(tidyverse)
> view(iris)
> df <- data.frame(iris)
> grp_spc <- group_by(iris,Species)
> summarise(iris,cnt=n())
  cnt
1 150
> summarise(grp_spc,cnt=n())
# A tibble: 3 × 2
  Species      cnt
  <fct>    <int>
1 setosa      50
2 versicolor  50
3 virginica   50
> summarize(iris,mn = mean(Sepal.Length),sd = sd(Sepal.Length))
      mn      sd
1 5.843333 0.8280661
> summarize(iris,mn = mean(Sepal.Width),sd = sd(Sepal.Width))
      mn      sd
1 3.057333 0.4358663
> summarize(iris,mn = mean(Petal.Length),sd = sd(Petal.Length))
      mn      sd
1 3.758 1.765298
> summarize(iris,mn = mean(Petal.Width),sd = sd(Petal.Width))
      mn      sd
1 1.199333 0.7622377
```

```

> summarize(grp_spc, mn = mean(Sepal.Length), sd = sd(Sepal.Length))
# A tibble: 3 × 3
  Species      mn      sd
  <fct>      <dbl> <dbl>
1 setosa      5.01 0.352
2 versicolor  5.94 0.516
3 virginica   6.59 0.636
> summarize(grp_spc, mn = mean(Sepal.Width), sd = sd(Sepal.Width))
# A tibble: 3 × 3
  Species      mn      sd
  <fct>      <dbl> <dbl>
1 setosa      3.43 0.379
2 versicolor  2.77 0.314
3 virginica   2.97 0.322
> summarize(grp_spc, mn = mean(Petal.Length), sd = sd(Petal.Length))
# A tibble: 3 × 3
  Species      mn      sd
  <fct>      <dbl> <dbl>
1 setosa      1.46 0.174
2 versicolor  4.26 0.470
3 virginica   5.55 0.552
> summarize(grp_spc, mn = mean(Petal.Width), sd = sd(Petal.Width))
# A tibble: 3 × 3
  Species      mn      sd
  <fct>      <dbl> <dbl>
1 setosa      0.246 0.105
2 versicolor  1.33 0.198
3 virginica   2.03 0.275
> |

```



## Program 12

**Find the correlation matrix:**

**A:-Plot the correlation plot on dataset and visualize giving an overview of relationship among variable on dataset.**

**Input: -**

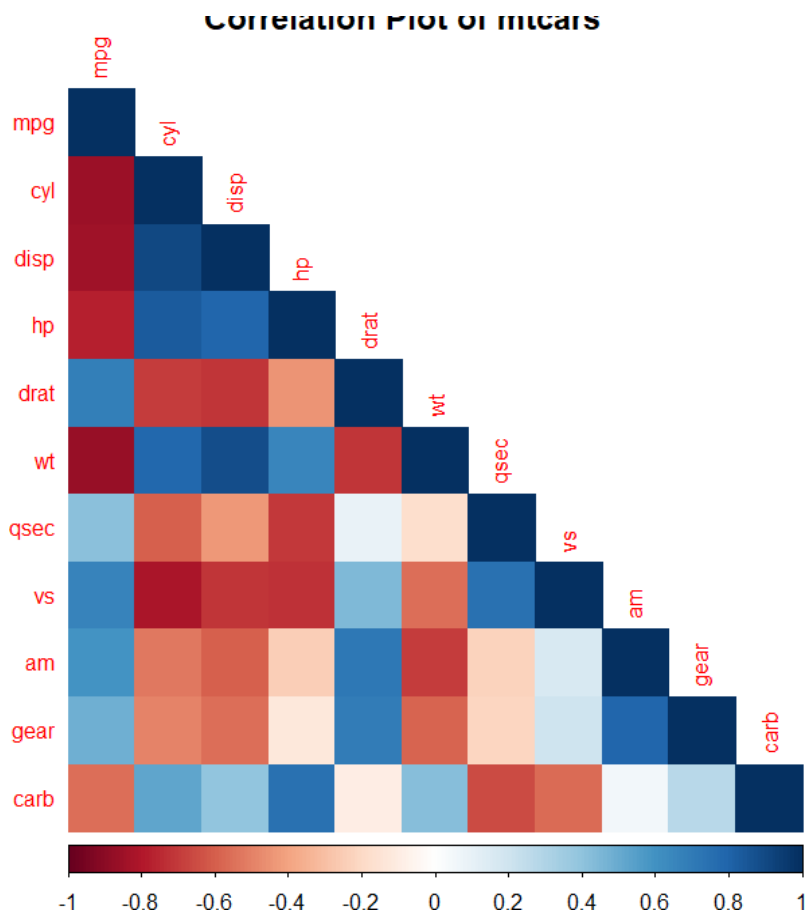
```
library(corrplot)
```

```
data(mtcars)
```

```
correlation_matrix <- cor(mtcars)
```

```
corrplot(correlation_matrix, method = "color", type = "lower", tl.cex = 0.8, title = "Correlation Plot of mtcars")
```

**Output: -**







**B:-Analysis of covariance :variance (ANOVA) if data have categorical variables on data set.**

**Input:**

```
data(mtcars)
```

```
model <- aov(mpg ~ am + hp, data = mtcars)
```

```
summary(model)
```

**Output:**

```
> summary(model)
      Df Sum Sq Mean Sq F value    Pr(>F)
am      1  405.2    405.2    47.87 1.33e-07 ***
hp      1  475.5    475.5    56.18 2.92e-08 ***
Residuals 29  245.4      8.5
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
> |
```

---

## Program 14

**a.Create a data frame from the sample data set**

**b.Create a table with the needed variables**

**c.perform the chi-Square test**

**Input: -**

```
df <- mtcars
df
name <- c("Alice", "Bob", "Charlie", "David", "Eva")
age <- c(25, 30, 22, 35, 28)
score <- c(85, 92, 78, 88, 95)
df <- data.frame(Name = name, Age = age, Score = score)
df
name <- c("Alice", "Bob", "Charlie", "David", "Eva")
age <- c(25, 30, 22, 35, 28)
score <- c(85, 92, 78, 88, 95)
df <- data.frame(Name = name, Age = age, Score = score)
library(knitr)
kable(df, caption = "Sample Data Table")
name <- c("Alice", "Bob", "Charlie", "David", "Eva")
age <- c(25, 30, 22, 35, 28)
age_group <- cut(age, breaks = c(20, 25, 30, 35, 40), labels = c("20-25", "26-30", "31-35", "36-40"))
contingency_table <- table(name, age_group)
chisq_test <- chisq.test(contingency_table)
print(chisq_test)
```

**Output: -**

```
Console Terminal x Background Jobs x
R 4.3.2 · C:/Users/Harsh Raj/AppData/Local/Temp/354232b0-9aeb-4565-b7aa-7ccc920c8701_programR[1].zip.701/
> df <- mtcars
> df
```

	mpg	cyl	dis	hp	drat	wt	qsec	vs	am	gear	carb
Mazda RX4	21.0	6	160.0	110	3.90	2.620	16.46	0	1	4	4
Mazda RX4 Wag	21.0	6	160.0	110	3.90	2.875	17.02	0	1	4	4
Datsun 710	22.8	4	108.0	93	3.85	2.320	18.61	1	1	4	1
Hornet 4 Drive	21.4	6	258.0	110	3.08	3.215	19.44	1	0	3	1
Hornet Sportabout	18.7	8	360.0	175	3.15	3.440	17.02	0	0	3	2
Valiant	18.1	6	225.0	105	2.76	3.460	20.22	1	0	3	1
Duster 360	14.3	8	360.0	245	3.21	3.570	15.84	0	0	3	4
Merc 240D	24.4	4	146.7	62	3.69	3.190	20.00	1	0	4	2
Merc 230	22.8	4	140.8	95	3.92	3.150	22.90	1	0	4	2
Merc 280	19.2	6	167.6	123	3.92	3.440	18.30	1	0	4	4
Merc 280C	17.8	6	167.6	123	3.92	3.440	18.90	1	0	4	4
Merc 450SE	16.4	8	275.8	180	3.07	4.070	17.40	0	0	3	3
Merc 450SL	17.3	8	275.8	180	3.07	3.730	17.60	0	0	3	3
Merc 450SLC	15.2	8	275.8	180	3.07	3.780	18.00	0	0	3	3
Cadillac Fleetwood	10.4	8	472.0	205	2.93	5.250	17.98	0	0	3	4
Lincoln Continental	10.4	8	460.0	215	3.00	5.424	17.82	0	0	3	4
Chrysler Imperial	14.7	8	440.0	230	3.23	5.345	17.42	0	0	3	4
Fiat 128	32.4	4	78.7	66	4.08	2.200	19.47	1	1	4	1
Honda Civic	30.4	4	75.7	52	4.93	1.615	18.52	1	1	4	2
Toyota Corolla	33.9	4	71.1	65	4.22	1.835	19.90	1	1	4	1
Toyota Corona	21.5	4	120.1	97	3.70	2.465	20.01	1	0	3	1
Dodge Challenger	15.5	8	318.0	150	2.76	3.520	16.87	0	0	3	2
AMC Javelin	15.2	8	304.0	150	3.15	3.435	17.30	0	0	3	2
Camaro Z28	13.3	8	350.0	245	3.73	3.840	15.41	0	0	3	4
Pontiac Firebird	19.2	8	400.0	175	3.08	3.845	17.05	0	0	3	2
Fiat X1-9	27.3	4	79.0	66	4.08	1.935	18.90	1	1	4	1
Porsche 914-2	26.0	4	120.3	91	4.43	2.140	16.70	0	1	5	2
Lotus Europa	30.4	4	95.1	113	3.77	1.513	16.90	1	1	5	2
Ford Pantera L	15.8	8	351.0	264	4.22	3.170	14.50	0	1	5	4
Ferrari Dino	19.7	6	145.0	175	3.62	2.770	15.50	0	1	5	6
Maserati Bora	15.0	8	301.0	335	3.54	3.570	14.60	0	1	5	8
Volvo 142E	21.4	4	121.0	109	4.11	2.780	18.60	1	1	4	2

```
> name <- c("Alice", "Bob", "Charlie", "David", "Eva")
> age <- c(25, 30, 22, 35, 28)
> score <- c(85, 92, 78, 88, 95)
> df <- data.frame(Name = name, Age = age, Score = score)
> df
```

```
> df
  Name Age Score
1  Alice  25   85
2   Bob   30   92
3 Charlie  22   78
4  David  35   88
5   Eva   28   95
> name <- c("Alice", "Bob", "Charlie", "David", "Eva")
> age <- c(25, 30, 22, 35, 28)
> score <- c(85, 92, 78, 88, 95)
> df <- data.frame(Name = name, Age = age, Score = score)
> library(knitr)
> kable(df, caption = "Sample Data Table")
```

Table: Sample Data Table

Name	Age	Score
Alice	25	85
Bob	30	92
Charlie	22	78
David	35	88
Eva	28	95

```
> name <- c("Alice", "Bob", "Charlie", "David", "Eva")
> age <- c(25, 30, 22, 35, 28)
> age_group <- cut(age, breaks = c(20, 25, 30, 35, 40), labels = c("20-25", "26-30", "31-35", "36-40"))
> contingency_table <- table(name, age_group)
> chisq_test <- chisq.test(contingency_table)
Warning message:
In chisq.test(contingency_table) :
  Chi-squared approximation may be incorrect
> print(chisq_test)

Pearson's Chi-squared test

data: contingency_table
X-squared = NaN, df = 12, p-value = NA
```

## **Program-15**

**Perform complete steps of exploratory data analysis on standard dataset (iris flowers, Wine Quality Dataset etc.)**

### **Input:**

```
data(iris)
head(iris)
summary(iris)
hist(iris$Sepal.Length, main = "Histogram of Sepal Length", xlab = "Sepal Length (cm)")
hist(iris$Sepal.Width, main = "Histogram of Sepal Width", xlab = "Sepal Width (cm)")
hist(iris$Petal.Length, main = "Histogram of Petal Length", xlab = "Petal Length (cm)")
hist(iris$Petal.Width, main = "Histogram of Petal Width", xlab = "Petal Width (cm)")

boxplot(iris$Sepal.Length, main = "Boxplot of Sepal Length")
boxplot(iris$Sepal.Width, main = "Boxplot of Sepal Width")
boxplot(iris$Petal.Length, main = "Boxplot of Petal Length")
boxplot(iris$Petal.Width, main = "Boxplot of Petal Width")

pairs(iris[, 1:4], pch = 19, col = iris$Species)

cor(iris[, 1:4])
```

### **Output:**

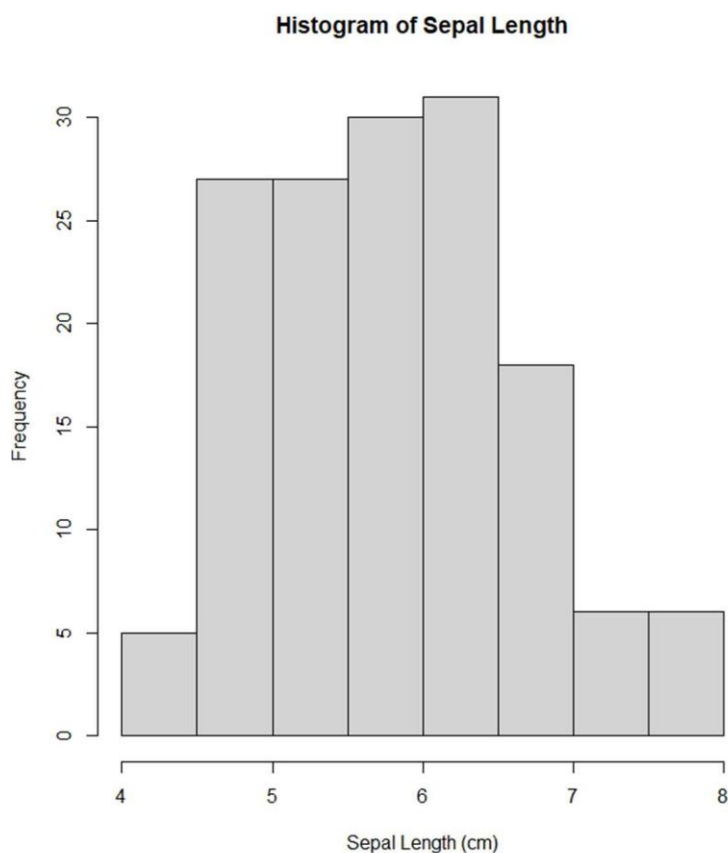
```

Console Terminal Jobs
R 4.3.1 ~ /
> data(iris)
> head(iris)
  Sepal.Length Sepal.width Petal.Length Petal.width Species
1          5.1         3.5         1.4         0.2  setosa
2          4.9         3.0         1.4         0.2  setosa
3          4.7         3.2         1.3         0.2  setosa
4          4.6         3.1         1.5         0.2  setosa
5          5.0         3.6         1.4         0.2  setosa
6          5.4         3.9         1.7         0.4  setosa
> summary(iris)
      Sepal.Length      Sepal.width      Petal.Length      Petal.width      Species
Min.   :4.300   Min.   :2.000   Min.   :1.000   Min.   :0.100   setosa   :50
1st Qu.:5.100   1st Qu.:2.800   1st Qu.:1.600   1st Qu.:0.300   versicolor:50
Median :5.800   Median :3.000   Median :4.350   Median :1.300   virginica :50
Mean   :5.843   Mean   :3.057   Mean   :3.758   Mean   :1.199
3rd Qu.:6.400   3rd Qu.:3.300   3rd Qu.:5.100   3rd Qu.:1.800
Max.   :7.900   Max.   :4.400   Max.   :6.900   Max.   :2.500
> hist(iris$Sepal.Length, main = "Histogram of Sepal Length", xlab = "Sepal Length (cm)")
> hist(iris$Sepal.width, main = "Histogram of Sepal width", xlab = "Sepal width (cm)")
> hist(iris$Petal.Length, main = "Histogram of Petal Length", xlab = "Petal Length (cm)")
> hist(iris$Petal.width, main = "Histogram of Petal width", xlab = "Petal width (cm)")
> boxplot(iris$Sepal.Length, main = "Boxplot of Sepal Length")
> boxplot(iris$Sepal.width, main = "Boxplot of Sepal width")
> boxplot(iris$Petal.Length, main = "Boxplot of Petal Length")
> boxplot(iris$Petal.width, main = "Boxplot of Petal width")
> pairs(iris[, 1:4], pch = 19, col = iris$Species)
> cor(iris[, 1:4])
      Sepal.Length Sepal.width Petal.Length Petal.width
Sepal.Length  1.0000000 -0.1175698  0.8717538  0.8179411
Sepal.width   -0.1175698  1.0000000 -0.4284401 -0.3661259
Petal.Length   0.8717538 -0.4284401  1.0000000  0.9628654
Petal.width    0.8179411 -0.3661259  0.9628654  1.0000000
> |

```

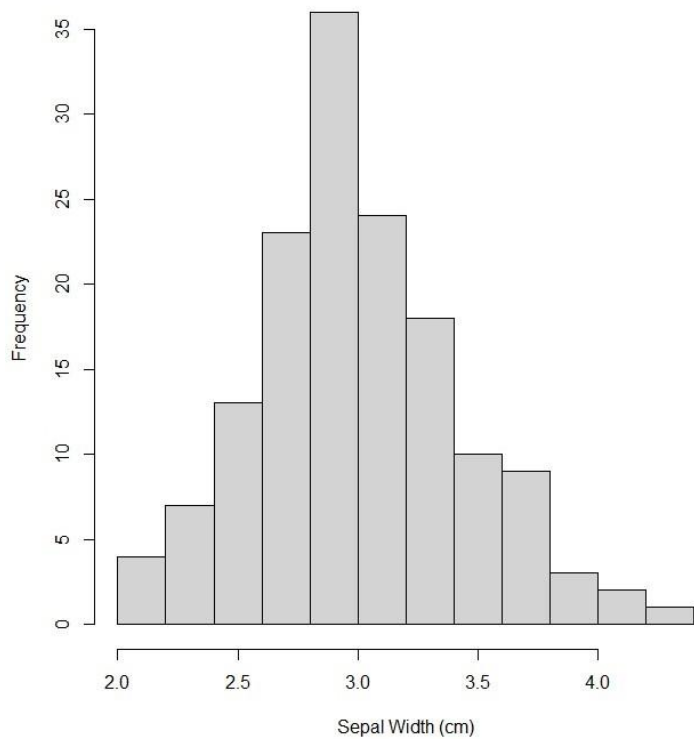
R Graphics: Device 2 (ACTIVE)

File History Resize

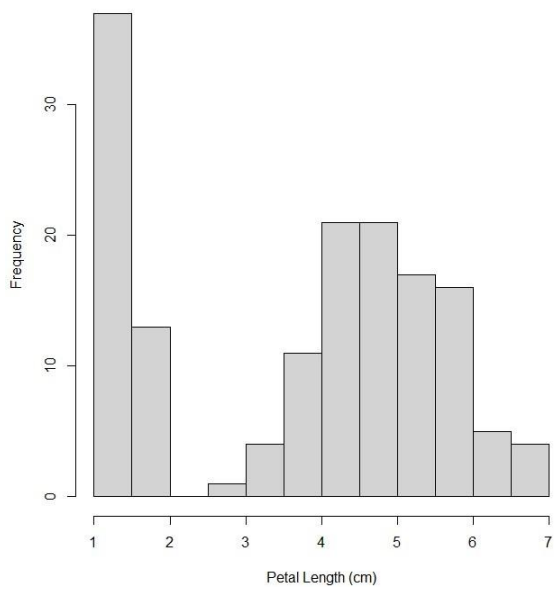




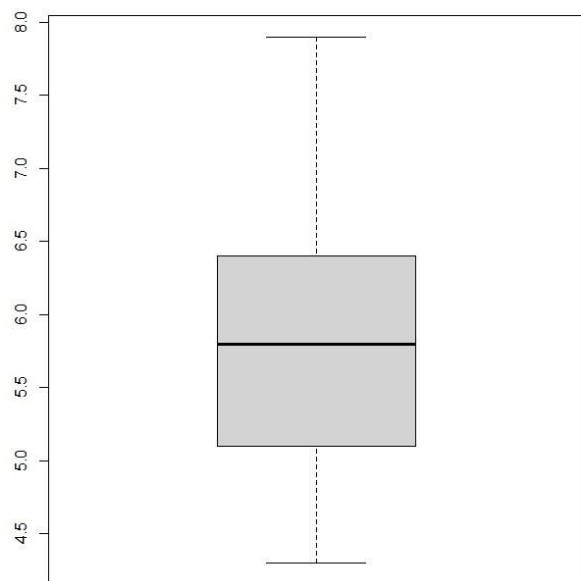
**Histogram of Sepal Width**



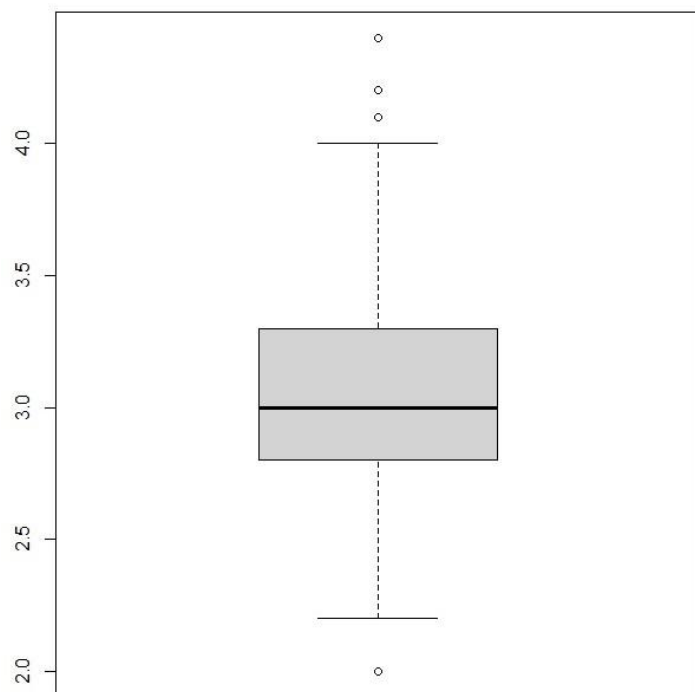
**Histogram of Petal Length**



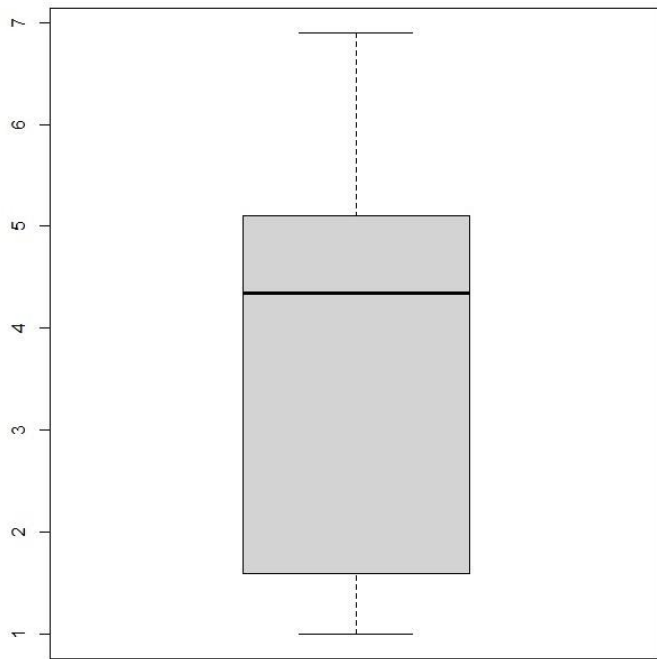
**Boxplot of Sepal Length**



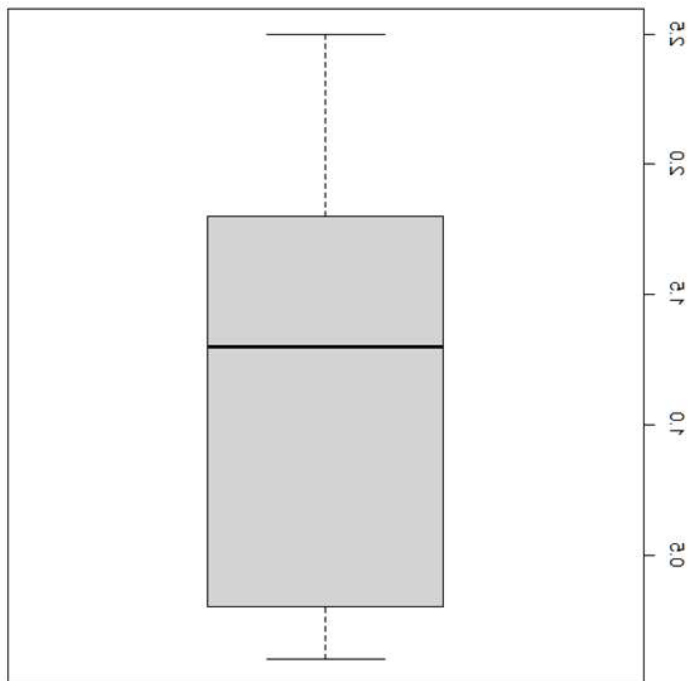
**Boxplot of Sepal Width**



Boxplot of Petal Length



Boxplot of Petal Width





## **Program :- 10**

- 1. a. Find the data distributions using box and scatter plot.**
- b. Find the outliers using box plot**
- c. Plot the histogram, bar chart and pie chart on sample data**
- d. Plot Pie chart, Histogram(3D) [Including colourful ones]**

### **Input Code:**

**a.**

```
# Sample data generation
set.seed(123) # Set a seed for reproducibility
data <- data.frame(
  Group = rep(1:3, each = 50), # Three groups
  Value = c(rnorm(50), rnorm(50, mean = 2), rnorm(50, mean = 4)) # Three sets of
random values
)

# Create a boxplot
boxplot(data$Value ~ data$Group,
  main = "Boxplot of Data Distributions",
  xlab = "Group",
  ylab = "Value",
  col = "lightblue",
  border = "blue")

# Create a scatter plot
plot(data$Group, data$Value,
  col = data$Group,
  pch = 19,
  main = "Scatter Plot of Data Distributions",
  xlab = "Group",
  ylab = "Value")

legend("topright", legend = unique(data$Group), col = 1:3, pch = 19)
```

**b.**

```
# Sample data generation
set.seed(123) # Set a seed for reproducibility
data <- data.frame(
  Value = c(rnorm(100), 5, 6, 15, 18) # 100 normal values and 4 outliers
)

# Create a boxplot
boxplot(data$Value,
  main = "Boxplot with Outliers")

# Identify potential outliers
outliers <- boxplot(data$Value)$out

# Print the potential outliers
cat("Potential Outliers:", outliers, "\n")
```

**c.**

**d.**

```
# Sample data generation
set.seed(123) # Set a seed for reproducibility
sample_data <- data.frame(
  Category = rep(letters[1:5], each = 20), # Five categories
  Value = c(rnorm(20, mean = 20, sd = 5), rnorm(20, mean = 40, sd = 8),
            rnorm(20, mean = 30, sd = 7), rnorm(20, mean = 25, sd = 6),
            rnorm(20, mean = 35, sd = 9))
)

# Create a histogram
hist(sample_data$Value, breaks = 10, main = "Histogram of Sample Data", xlab =
"Value", col = "lightblue")

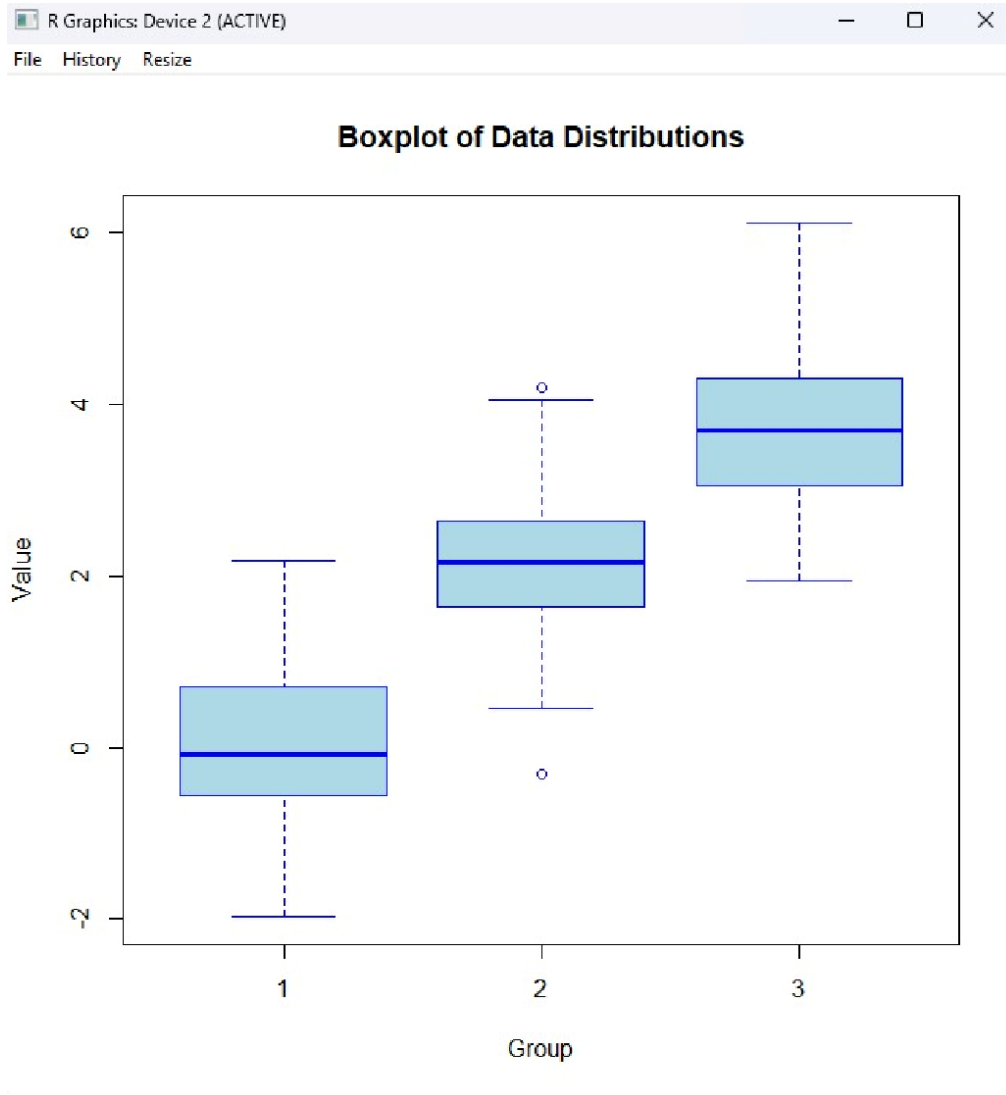
# Create a Histogram
hist(sample_data$Value,
      breaks = 10,
      main = "Histogram of Sample Data",
      xlab = "Value",
      col = "lightblue")

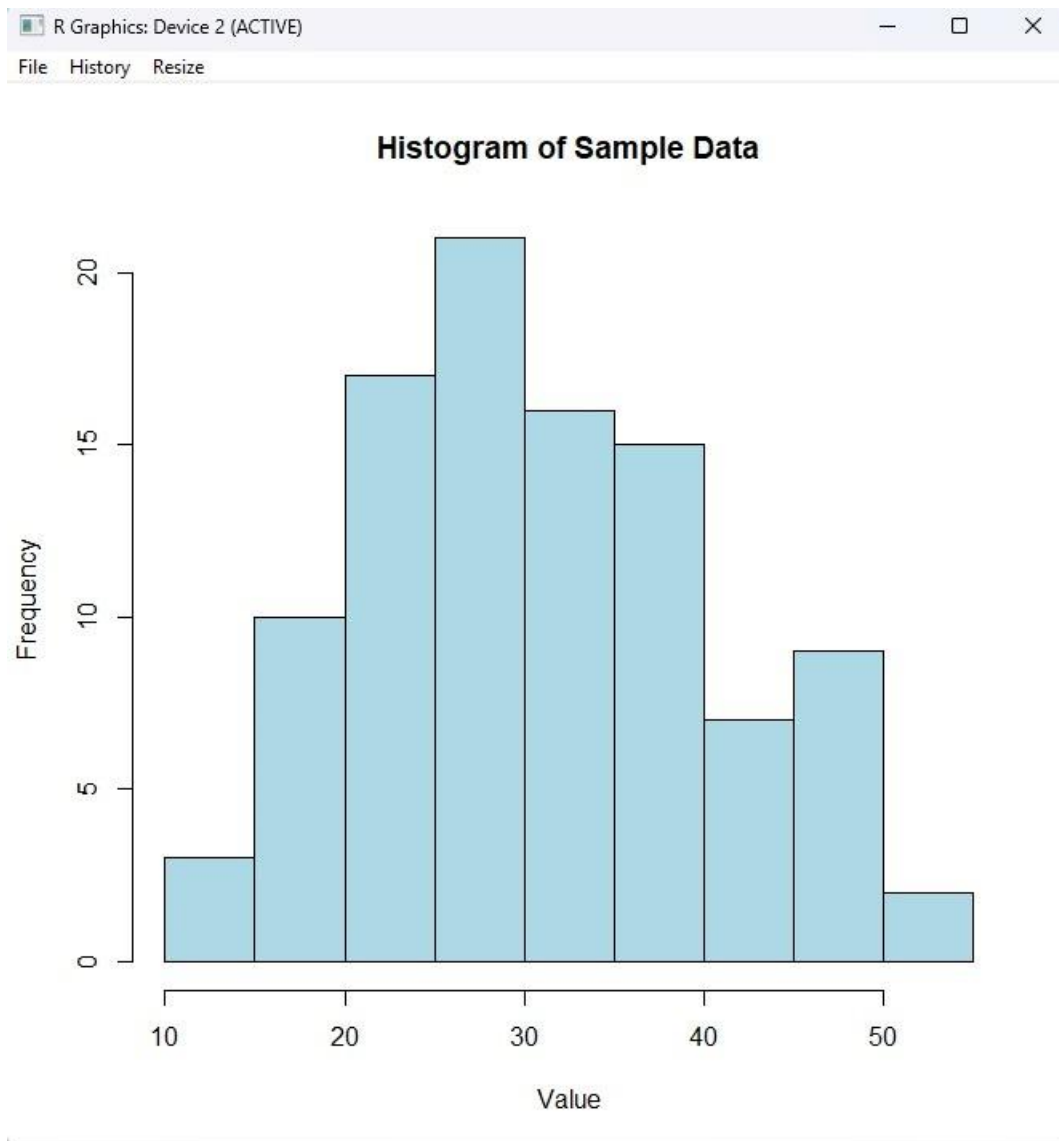
# Create a Bar Chart
category_counts <- table(sample_data$Category)
barplot(category_counts,
        main = "Bar Chart of Sample Data",
        xlab = "Category",
        ylab = "Count",
        col = "lightblue")

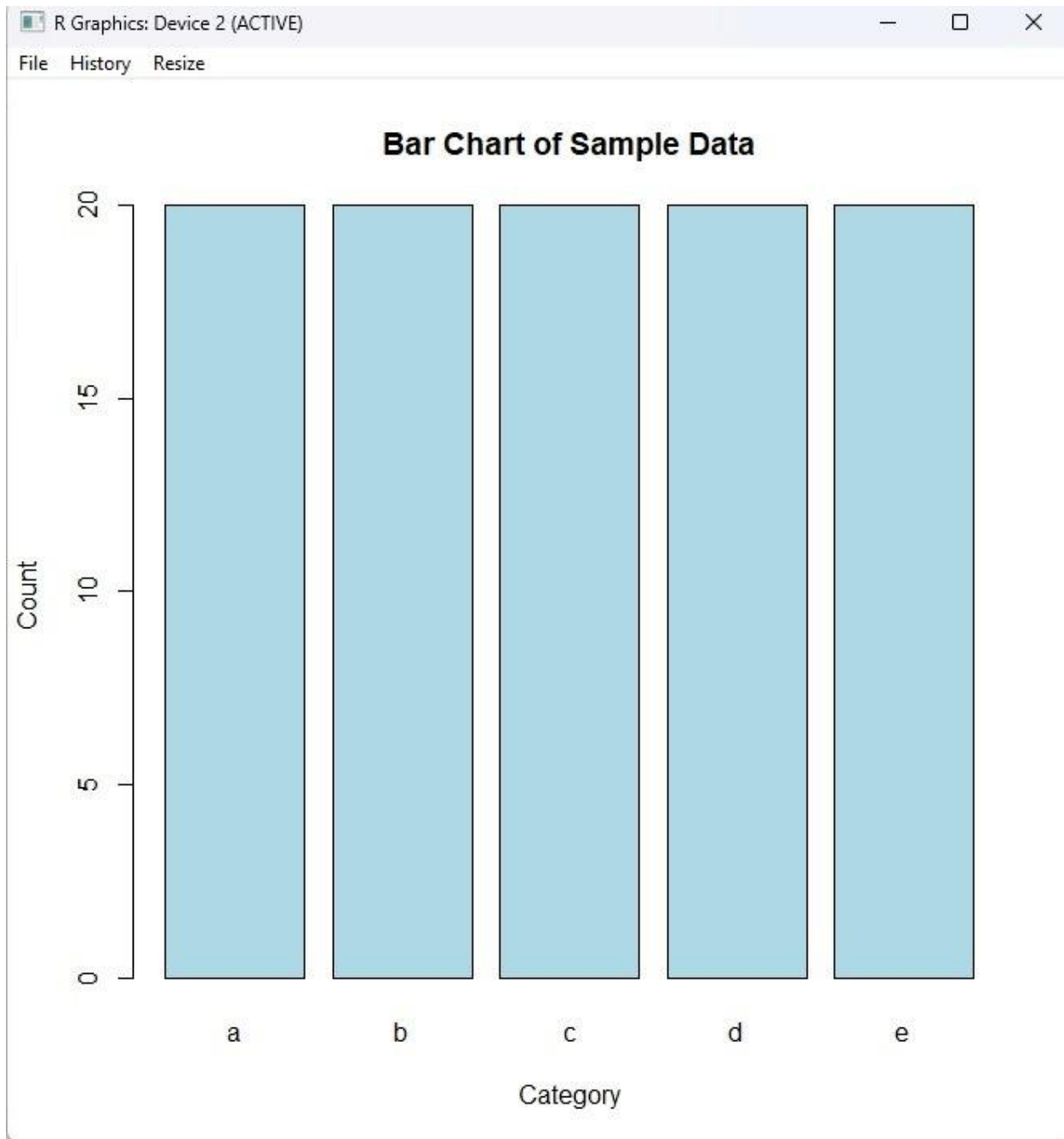
# Create a Pie Chart
category_proportions <- prop.table(category_counts)
colors <- rainbow(length(category_proportions))

pie(category_proportions,
    labels = category_counts,
    main = "Pie Chart of Sample Data",
    col = colors)
```

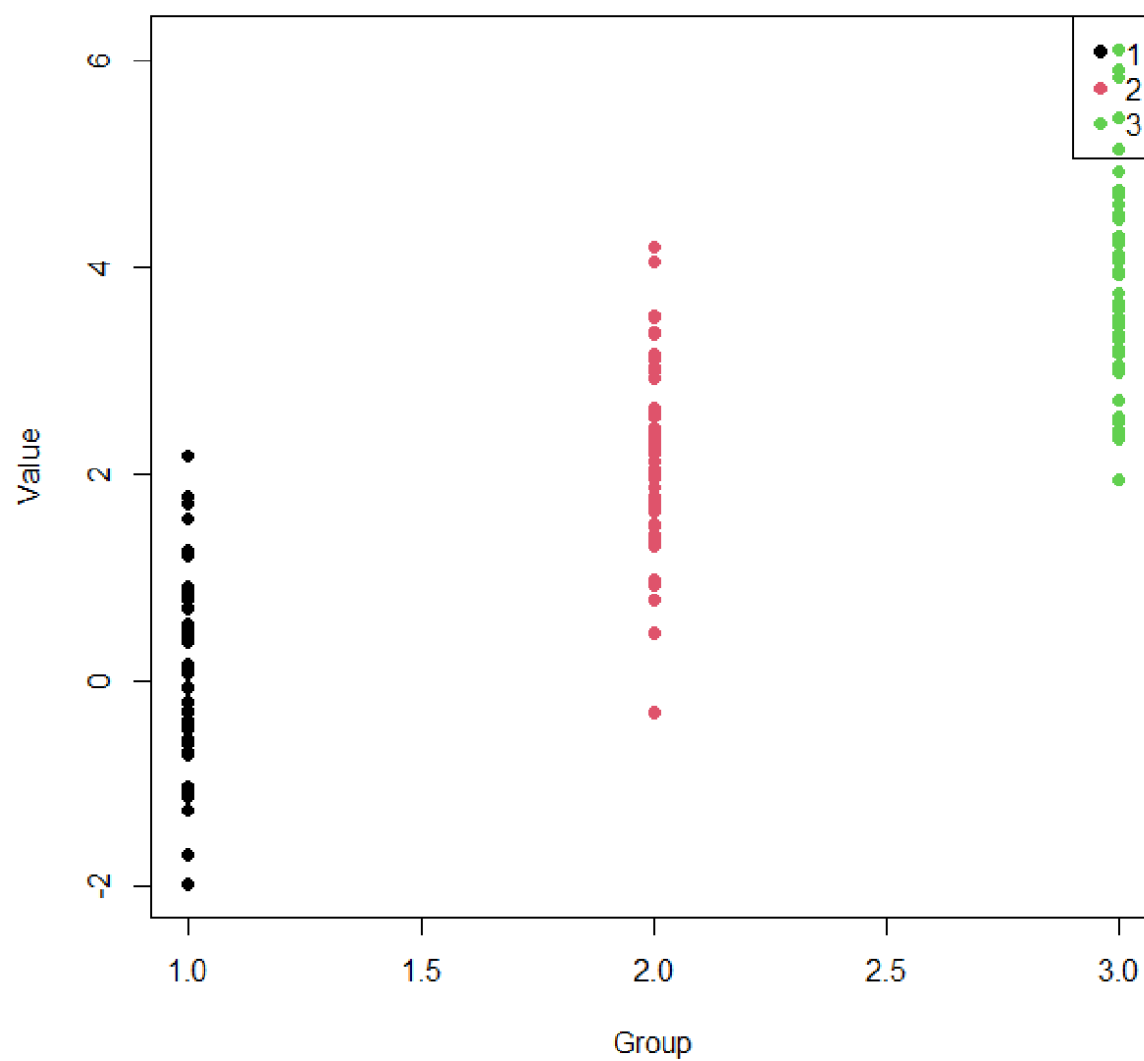


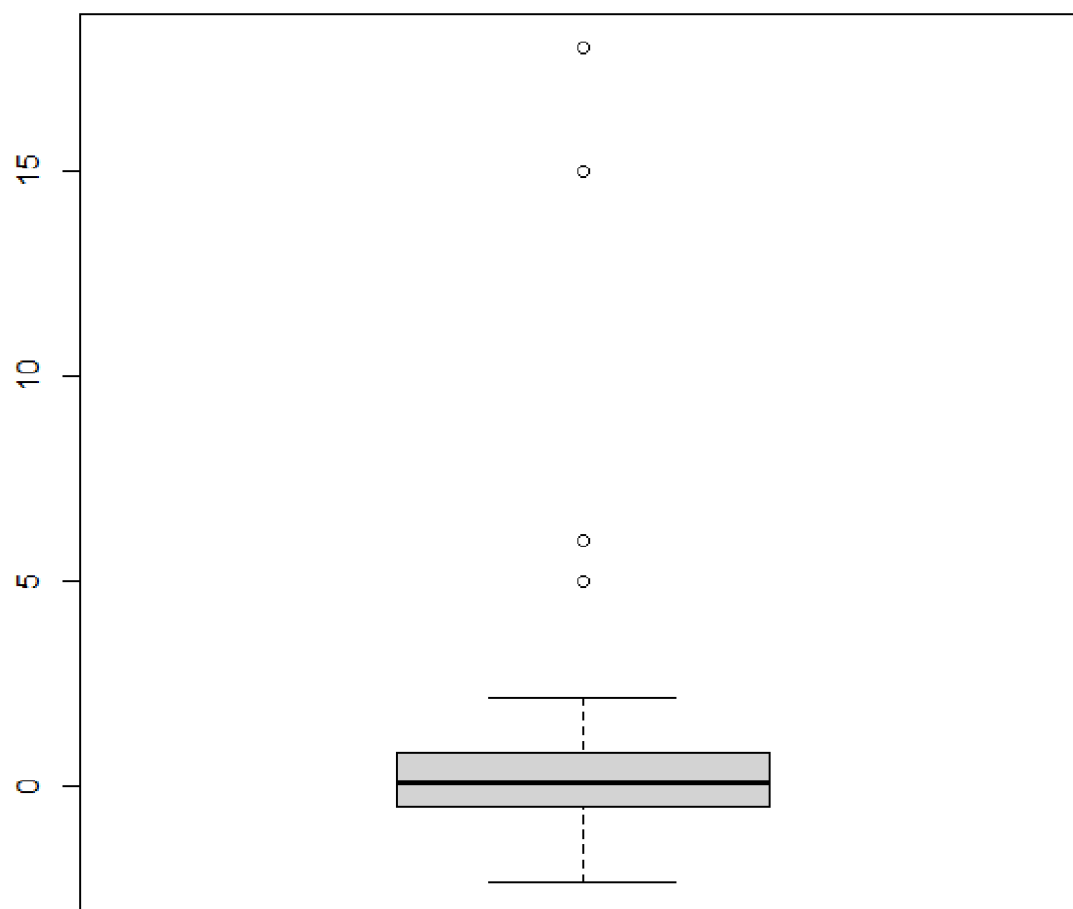




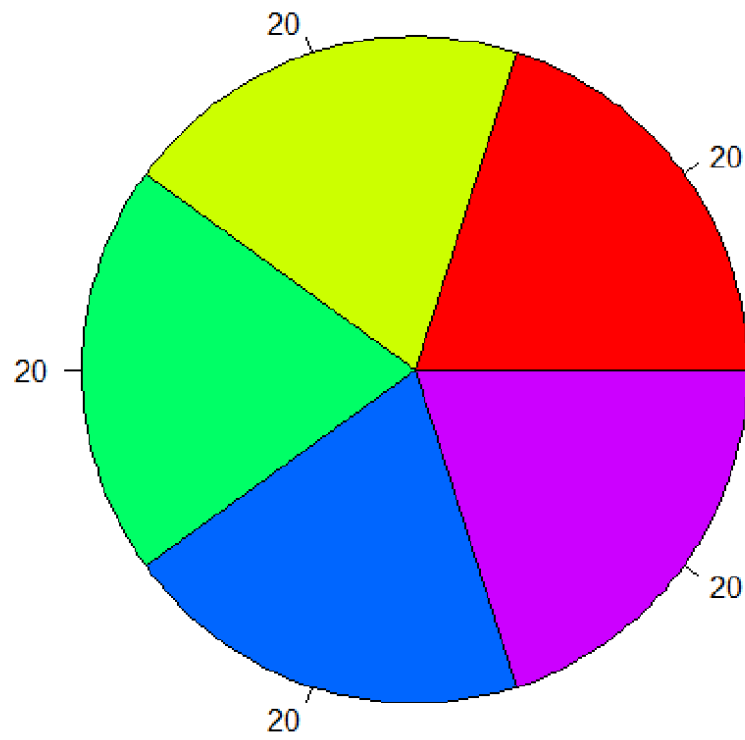


**Scatter Plot of Data Distributions**





**Pie Chart of Sample Data**







## **Program :- 11**

**1. Import a sample dataset and perform Regression techniques to find out relation between variables.**

### **Input Code:**

```
# Load the dataset
data(mtcars)

# View the first few rows of the dataset
head(mtcars)

# Perform a linear regression
linear_model <- lm(mpg ~ wt, data = mtcars)

# Summary of the linear regression
summary(linear_model)

# Plot the regression line
plot(mtcars$wt, mtcars$mpg, main = "Scatterplot of mpg vs. wt", xlab = "Weight
(wt)", ylab = "Miles per Gallon (mpg)")
abline(linear_model, col = "red")

# Perform a multiple linear regression
multi_linear_model <- lm(mpg ~ wt + hp + qsec, data = mtcars)

# Summary of the multiple linear regression
summary(multi_linear_model)
```

## Output:

```
> # Load the dataset
> data(mtcars)
> # View the first few rows of the dataset
> head(mtcars)
```

	mpg	cyl	disp	hp	drat	wt	qsec	vs	am	gear	carb
Mazda RX4	21.0	6	160	110	3.90	2.620	16.46	0	1	4	4
Mazda RX4 Wag	21.0	6	160	110	3.90	2.875	17.02	0	1	4	4
Datsun 710	22.8	4	108	93	3.85	2.320	18.61	1	1	4	1
Hornet 4 Drive	21.4	6	258	110	3.08	3.215	19.44	1	0	3	1
Hornet Sportabout	18.7	8	360	175	3.15	3.440	17.02	0	0	3	2
Valiant	18.1	6	225	105	2.76	3.460	20.22	1	0	3	1

```
> # Perform a linear regression
> linear_model <- lm(mpg ~ wt, data = mtcars)
> Summary of the linear regression
> summary(linear_model)
```

Call:  
lm(formula = mpg ~ wt, data = mtcars)

Residuals:

	Min	1Q	Median	3Q	Max
	-4.5432	-2.3647	-0.1252	1.4096	6.8727

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	37.2851	1.8776	19.858	< 2e-16 ***
wt	-5.3445	0.5591	-9.559	1.29e-10 ***

---  
Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 3.046 on 30 degrees of freedom  
Multiple R-squared: 0.7528, Adjusted R-squared: 0.7446  
F-statistic: 91.38 on 1 and 30 DF, p-value: 1.294e-10

```
> # Plot the regression line
> plot(mtcars$wt, mtcars$mpg, main = "Scatterplot of mpg vs. wt", xlab = "Weight (wt)", ylab = "Miles per Gallon (mpg)")
> abline(linear_model, col = "red")
> # Perform a multiple linear regression
> multi_linear_model <- lm(mpg ~ wt + hp + qsec, data = mtcars)
> Summary of the multiple linear regression
> summary(multi_linear_model)
```

Call:  
lm(formula = mpg ~ wt + hp + qsec, data = mtcars)

Residuals:

	Min	1Q	Median	3Q	Max
	-3.8591	-1.6418	-0.4636	1.1940	5.6092

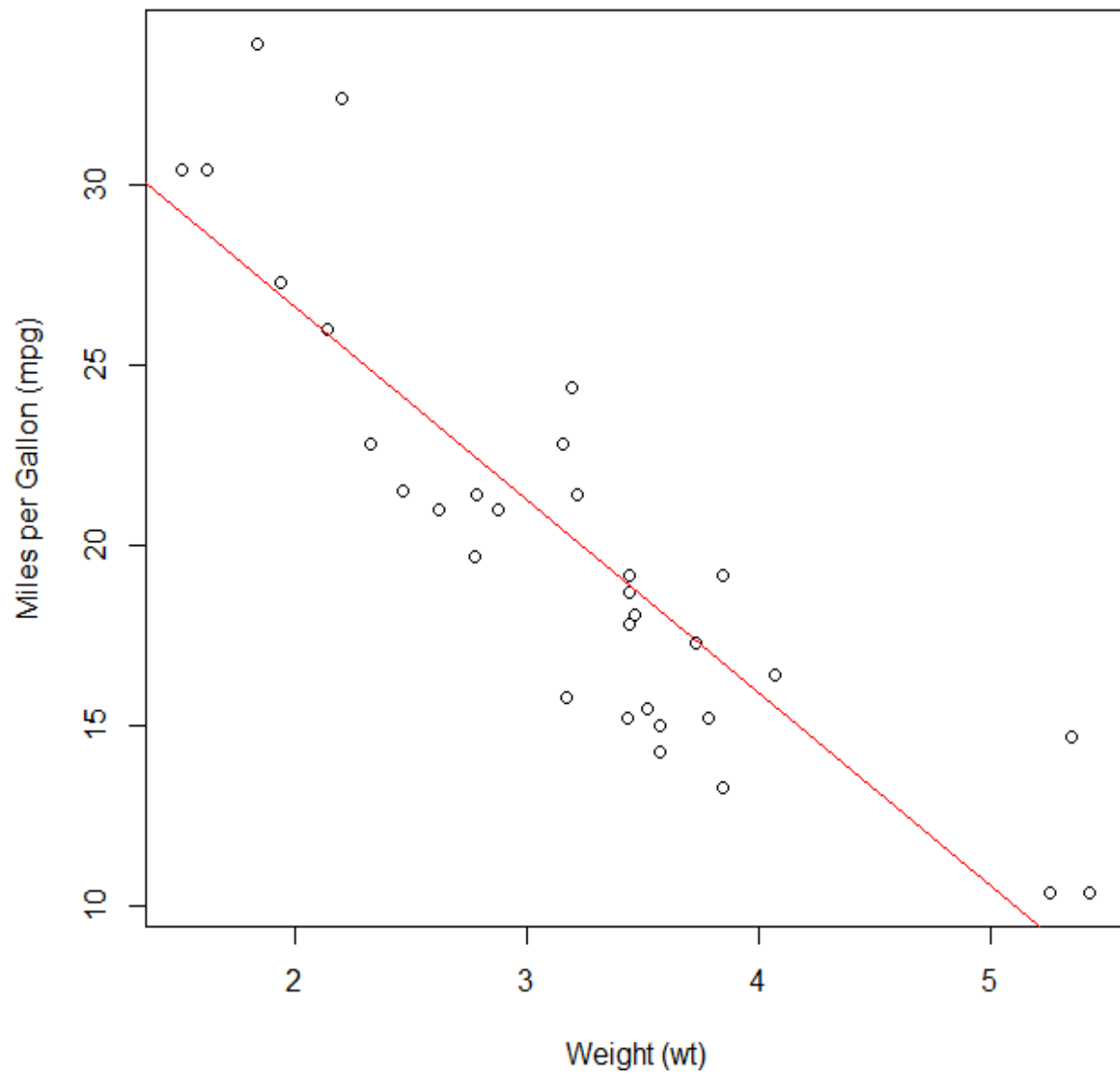
Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	27.61053	8.41993	3.279	0.00278 **
wt	-4.35880	0.75270	-5.791	3.22e-06 ***
hp	-0.01782	0.01498	-1.190	0.24418
qsec	0.51083	0.43922	1.163	0.25463

---  
Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 2.578 on 28 degrees of freedom  
Multiple R-squared: 0.8348, Adjusted R-squared: 0.8171  
F-statistic: 47.15 on 3 and 28 DF, p-value: 4.506e-11

**Scatterplot of mpg vs. wt**



### **Program :- 13**

**1. Write a program to create 3D plot , to add title, change viewing direction, add color and shade to the plot.**

#### **Input Code:**

```
# Create sample data
x <- seq(-10, 10, length.out = 50)
y <- seq(-10, 10, length.out = 50)
z <- outer(x, y, function(x, y) sin(sqrt(x^2 + y^2)) / (sqrt(x^2 + y^2)))

# Create the 3D plot
persp(x, y, z,
      main = "3D Plot with Title",
      theta = 30, phi = 20, # Change viewing angles
      col = "skyblue",     # Set color
      shade = 0.5)         # Add shading
```

**Output:**

**3D Plot with Title**

