



SRI RAMACHANDRA

INSTITUTE OF HIGHER EDUCATION AND RESEARCH

(Category - I Deemed to be University) Porur, Chennai

SRI RAMACHANDRA ENGINEERING AND TECHNOLOGY

CSE 240 Data Science with R

STUDENT WORK BOOK

Name : M.Sathishkumar

Unique ID : E0119052

Year : II

Quarter : Q6

Department : B.Tech CSE (AI &ML)

Faculty Name : Prof.B.Nirmala

Academic Year : 2020-2021

Date:02/11/2020

Q.NO 1: Question

Consider 2 vectors c(9,10,11,12) and c(13,14,15,16).

Create a 4 by 2 matrix from these two vectors

Program:

```
#Matrix creation  
matrix<-matrix(c(c(9,10,11,12),c(13,14,15,16))),nrow = 4,ncol = 2)  
#combining two vertices and converting into 4 by 2 matrix  
print(matrix)
```

Output:

	[,1]	[,2]
[1,]	9	13
[2,]	10	14
[3,]	11	15
[4,]	12	16

Explanation:

- ★ Matrix is a two-dimensional data structure in R programming.
- ★ Matrix can be created using the matrix () function
- ★ Dimension of the matrix can be defined by passing appropriate value guments nrow and ncol

Date: 02/11/2020

Q.NO 2: Question

Program:

Write an R program to take input from the user (userID and Group/Branch) and display the values

```
#UserInput
Id = readline("Enter your UserID:")
#Prompts user for input(id)
Branch = readline(prompt="Enter you Branch/Group:")
#Prompts user for input(branch)
cat("Your UserID is",Id,"and you belong to",Branch,"group")
#Displaying Values
```

Output:

Enter your UserID:E0119052

Enter you Branch/Group:AI & ML

Your UserID is E0119052 and you belong to AI & ML group

Explanation:

- ★ readline reads a line from the terminal (in interactive use).
- ★ readline() lets the user enter a one-line string at the terminal.
- ★ The prompt argument is printed in front of the user input. It usually ends on ": ".

Date: 02/11/2020

Q.NO 3: Question

Create a data frame Write a R program to create a data frame from four given vectors.

a name b. Subject C. Score d. Rank

Program:

```
#Dataframe creation
Name <- c("Jhon", "Lee", "Suzan", "Abhinav", "Brain", "Emma", "David")
#Variables
Subject <- c("Data Science", "Machine Learning", "Deep Learning",
             "Data Structures", "Database Managemnt System",
             "Operating Systems", "Python Programming")

Score <- c(56, 76, 86, 96, 73, 87, 47)

Rank <- c(5,8,6,7,9,2,1)

df = data.frame(Name,Subject,Score,Rank)
print(df)
#Dataframe Created Successfully
```

Output:

	Name	Subject	Score	Rank
1	Jhon	Data Science	56	5
2	Lee	Machine Learning	76	8
3	Suzan	Deep Learning	86	6
4	Abhinav	Data Structures	96	7
5	Brain	Database Managemnt System	73	9
6	Emma	Operating Systems	87	2
7	David	Python Programming	47	1

Explanation:

- ★ A **data frame** is used for storing data tables. It is a list of vectors of equal length.
- ★ The top line of the table, called the **header**, contains the column names.
- ★ Each horizontal line afterward denotes a **data row**, which begins with header, and then followed by the actual data.
- ★ Each data member of a row is called a **cell**.

Date: 02/11/2020

Q.NO 4: Question

Write a R program to get the statistical summary and nature of the data of a given data frame. (use 3rd Question dataframe

Program:

```
#Dataframe creation
Name <- c("Jhon", "Lee", "Suzan", "Abhinav", "Brain", "Emma", "David")
#Variables
Subject <- c("Data Science", "Machine Learning", "Deep Learning",
             "Data Structures", "Database Managemnt System",
             "Operating Systems", "Python Programming")
Score <- c(56, 76, 86, 96, 73, 87, 47)
Rank <- c(5,8,6,7,9,2,1)
df = data.frame(Name,Subject,Score,Rank)
print(df)
#Dataframe Created Successfully
print(summary(df))
#Dataframe Stastical summary
```

Output:

	Name	Subject	Score	Rank
Abhinav:1	Data Science	:1	Min. :47.00	Min. :1.000
Brain :1	Data Structures	:1	1st Qu.:64.50	1st Qu.:3.500
David :1	Database Managemnt System	:1	Median :76.00	Median :6.000
Emma :1	Deep Learning	:1	Mean :74.43	Mean :5.429
Jhon :1	Machine Learning	:1	3rd Qu.:86.50	3rd Qu.:7.500
Lee :1	Operating Systems	:1	Max. :96.00	Max. :9.000
Suzan :1	Python Programming	:1		

Explanation:

Summary is a generic function used to produce result summaries of the results of various model fitting functions.

The function invokes particular methods which depend on the class of the first argument

The form of the value returned by summary depends on the class of its argument

Date: 02/11/2020

Q.NO 5: Question

Write a R program to extract specific column from a data frame using column name

Program:

```
#Dataframe creation
Name <- c("Jhon", "Lee", "Suzan", "Abhinav", "Brain", "Emma", "David")
#Variables
Subject <- c("Data Science", "Machine Learning", "Deep Learning",
             "Data Structures", "Database Managemnt System",
             "Operating Systems", "Python Programming")
Score <- c(56, 76, 86, 96, 73, 87, 47)
Rank <- c(5,8,6,7,9,2,1)
df = data.frame(Name,Subject,Score,Rank)
print(df)
#Dataframe Created Successfully
print(df["Subject"])
#Sliced column with column name
```

Output:

	Subject
1	Data Science
2	Machine Learning
3	Deep Learning
4	Data Structures
5	Database Managemnt System
6	Operating Systems
7	Python Programming

Explanation:

- ★ We retrieve a data frame column slice with the single square bracket "[" operator.
- ★ Here We can retrieve the same column slice by its name.
- ★ And we can pack the row names in an index vector in order to retrieve multiple rows

Date: 02/11/2020

Q.NO 6: Question

.Write a R program to extract first two rows from a given data frame

Program:

```
#Dataframe creation
Name <- c("Jhon", "Lee", "Suzan", "Abhinav", "Brain", "Emma", "David")
#Variables
Subject <- c("Data Science", "Machine Learning", "Deep Learning",
             "Data Structures", "Database Managemnt System",
             "Operating Systems", "Python Programming")
Score <- c(56, 76, 86, 96, 73, 87, 47)
Rank <- c(5,8,6,7,9,2,1)
df = data.frame(Name,Subject,Score,Rank)
#Dataframe Created Successfully
print(df[1:2,])
#splicing Dataframe with respect to index
```

Output:

	Name	Subject	Score	Rank
1	Jhon	Data Science	56	5
2	Lee	Machine Learning	76	8

Explanation:

- ★ We retrieve rows from a data frame with the single square bracket operator, just like what we did with columns.
- ★ However, in additional to an index vector of row positions, we append an extra comma character.
- ★ This is important, as the extra comma signals a wildcard match for the second coordinate for column positions.

Date: 03/11/2020

Q.NO 7: Question

Create an R script that calculates the square root of a given integer vector x of length one, if the value contained in x is negative it should return NA.

Program:

```
#Prompting user for input
x = c(as.integer(readline("Enter a number:")))
#Defining a function for finding square root
squareroot <- function(x){
  if (x < 0){
    print(NA)
  }else{
    print(paste("Square root of the given number :",sqrt(x)))
  }
}
squareroot(x)
#Invoking the function
```

Output:

Enter a number:57

```
[1] "Square root of the given number : 7.54983443527075"
```

Enter a number: -78

```
[1] NA
```

Explanation:

- ★ Missing values are represented by the symbol **NA** (not available)
- ★ If the numeric_Expression is a positive value, the sqrt function returns the square root of a given value.
- ★ If the numeric_Expression is a negative value, the sqrt function return **NaN**.
- ★ Numeric_Expression is not a number (**NaN**), or Negative Infinity, then sqrt in R returns **NaN**.

Date: 03/11/2020

Q.NO 8: Question

Demonstrate and examine the output of letter and LETTER

Program:

```
a = "letter"  
b = "LETTER"  
print(a)           #printing letter  
print(b)           #printing LETTER
```

Output:

```
[1] "letter"  
[1] "LETTER"
```

Explanation:

- ★ This is useful because you can always view your data: just print it.
- ★ You needn't write special printing logic, even for complicated data structures.
- ★ The print function has a significant limitation, it prints only one object at a time.
- ★ Trying to print multiple items gives this mind-numbing error message

Date: 03/11/2020

Q.NO 9: Question

Create an R script that, given a numeric vector x with length 3, will print the elements by order from high to low.

Program:

```
desc <- function(x){  
  x<-c(5,7,8)  
  }  
print(sort(desc(x),decreasing = TRUE))
```

Output:

```
[1] 8 7 5
```

Explanation:

- ★ Sort (or *order*) a vector or factor (partially) into ascending or descending order. For ordering along more than one variable, e.g., for sorting data frame
- ★ To sort a data frame in R, use the **sort()** function. By default, sorting is ASCENDING.
- ★ Prepend the sorting variable by a minus sign to indicate DESCENDING order

Date: 03/11/2020

Q.NO 10: Question

Create an R script that returns the amount of values that are larger than the mean of a vector.
You are allowed to use mean(). (Use function)

Program:

```
#function
nums<- function(vec)
{
  #returning values greater then mean value
  return(vec[vec>mean(vec)])
}
#Assigning values
nums(c(78,12,459,175,451,178,587,24,985))
```

Output:

459 451 587 985

Explanation:

- ★ It is calculated by taking the sum of the values and dividing with the number of values in a data series.
- ★ The function **mean()** is used to calculate and decision operator ">" is used to compare the giving condition and take decision

Date: 03/11/2020

Q.NO 11: Question

Write a double for loop which prints 30 numbers (1:10, 2:11, 3:12). Those are three clusters of ten numbers each. The first loop determines the number of clusters (3) via its length; the second loop the numbers to be printed (1 to 10 at the beginning). Each cluster starts one number higher than the previous one

Program:

```
#Nested looping
for(i in seq(1,3)){
  #for every single run in outside loop inside loop gets executed 10 times
  for(j in seq(i,i+10L)){
    print(j)
  }
}
```

Output:

1 2 3 4 5 6 7 8 9 10 11 2 3 4 5 6 7 8 9 10 11 12 3 4 5 6 7 8 9 10 11 12 13

Explanation:

- ★ In Nested for Loop, it makes use of the control structures to manage the execution of the expression, one such control structure is Nested for Loop a similar to basic 'for' loop executes.
- ★ It can be defined as placing one 'for' loop inside the first 'for' loop is called as nesting or loop of loops in some terms, which takes the responsibility of two loops such that the outer loop controls the number of repetition of the whole inner detailed information until it is false,
- ★ in other words, the inner loop executes n-times of every execution of the outer for loop and also, it's a great tool to work with R Programming Language.

Date: 03/11/2020

Q.NO 12: Question

a. You have the data.frame 'mydf' with four columns like below

a = c(3,7,NA, 9) b = c(2,NA,9,3) f = c(5,2,5,6) d = c(NA,3,4,NA)

You want to add another column '5': the 5th column contains the value of col 2 if col 1 is NA; the 5th column contains the value of col 4 if col 2 is NA; the 5th column contains the value of col 3 in all other cases.

Program:

```
#Creating a dataframe
mydf = data.frame(a=c(3,7,NA, 9),b=c(2,NA,9,3),f=c(5,2,5,6),d=c(NA,3,4,NA))
#Applying conditions as required to create 5th column
mydf[,5] <- ifelse(is.na(mydf[,1]) & !is.na(mydf[,2]),mydf[,2],
                  ifelse(is.na(mydf[,2]) & !is.na(mydf[,4]),mydf[,4], mydf[,3]))

print(mydf)
```

Output:

	a	b	f	d	V5
1	3	2	5	NA	5
2	7	NA	2	3	3
3	NA	9	5	4	9
4	9	3	6	NA	6

Explanation:

- ★ An if statement can be followed by an optional else statement which executes when the Boolean expression is false.
- ★ An if statement can be followed by an optional else if...else statement, which is very useful to test various conditions using single if...else if statement

Date: 03/11/2020

Q.NO 13: Question

Write a while loop starting with x = 0. The loop prints all numbers up to 35 but it skips number 7.
Condition: If x== 7 next

Program:

```
#While looping
x<-0
while (x<=35) {
  if(x==7){
    x<-x+1
    #Applying next function to bypass if x equals 7
    next
  }
  cat(x,"")
  x=x+1
}
```

Output:

0 1 2 3 4 5 6 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35

Explanation:

- ★ The **next** statement is useful when we want to skip the current iteration of a loop without terminating it.
- ★ On encountering next, the R parser skips further evaluation and starts next iteration of the loop. In other words, the inner loop executes n-times of every execution of the outer for loop.

Date: 03/11/2020

Q.NO 14: Question

Examine the difference between typeof and class () method using R program

Program:

```
#Variable  
num<-458  
class(num)    #It gives numeric  
typeof(num)   #It gives double
```

Output:

'numeric'

'double'

Explanation:

- ★ 'class' is a property assigned to an object that determines how generic functions operate with it. It is not a mutually exclusive classification. If an object has no specific class assigned to it, such as a simple numeric vector, its class is usually the same as its mode, by convention.
- ★ Typeof determines the (R internal) type or storage mode of any object

Date: 03/11/2020

Q.NO 15: Question

Create a function and demonstrate their features like required, keyword, default.

Program:

```
#Keyword Argument
keyword<-function(a,b){
  return(a+b)
}
keyword(b=1,a=3)

#default Argument
def <- function(a=0) {
  return(a+1)
}
def(5)

#required Argument
norma<-function(x){
  return(x+1)
}
norma(1)
```

Output:

4
6
2

Explanation:

- ★ *Arguments are always named when you define a function.* When you call a function, you do not have to specify the name of the argument.
- ★ Arguments are optional; you do not have to specify a value for them. They can have a default value, which is used if you do not specify a value for that argument yourself.
- ★ You can use as many arguments as you like, there is no limit to the number of arguments. An argument list comprises of comma-separated values that contain the various formal arguments.

Date: 03/11/2020

Q.NO 16: Question

Create a dataframe and delete the row and column. (Use the own data values to create frame)

Program:

```
#Dataframe creation
Name <- c("Jhon", "Lee", "Suzan", "Abhinav", "Brain", "Emma", "David")
#Variables
Subject <- c("Data Science", "Machine Learning", "Deep Learning", "Data Structures",
             "Database Managemnt System", "Operating Systems", "Python Programming")

Score <- c(56, 76, 86, 96, 73, 87, 47)

Rank <- c(5,8,6,7,9,2,1)

df = data.frame(Name,Subject,Score,Rank)
del_row_df<-df[-c(2),] # row deletion
del_col_df<-within(df, rm("Subject")) #Colmn deletion
print("After deleting Row 2")
del_row_df
print("After Deleting Subject Column")
del_col_df
```

Output:

[1] "After deleting Row 2"

	Name	Subject	Score	Rank
1	Jhon	Data Science	56	5
3	Suzan	Deep Learning	86	6
4	Abhinav	Data Structures	96	7
5	Brain	Database Managemnt System	73	9
6	Emma	Operating Systems	87	2
7	David	Python Programming	47	1

[1] "After Deleting Subject Column"

Name	Score	Rank
Jhon	56	5
Lee	76	8
Suzan	86	6
Abhinav	96	7
Brain	73	9
Emma	87	2
David	47	1

Explanation:

- ★ -c(rowNum) is used to return a slice of a dataframe without the mentioned row i.e. row deletion
- ★ The within function returns a subset of a dataframe with the second argument as a function to apply to all the column rm function removes the "Subject" column out of the data frame typeof determines the (R internal) type or storage mode of any object.

Date: 04/11/2020

Q.NO 17: Question

Write a function that turns (e.g.) a vector `c("a", "b", "c")` into the string "a, b, and c". Think carefully about what it should do if given a vector of length 0, 1, or 2.

Program:

```
#collapse function
convert=function(x) {
  #Using collapse for concatenating into one string
  y=paste(x,collapse=",")
  print(y)
}
#Checking for 3 values
convert(c("sathish","kumar","good"))
convert(c("sathish","kumar"))
#Checking for 1 value
convert(c("sathish"))
```

Output:

```
[1] "sathish,kumar,good"
[1] "sathish,kumar"
[1] "sathish"
```

Explanation:

- ★ **'collapse'** is a property Collapses a character vector of any length into a length 1 vector.
- ★ Syntax - `collapse(x, sep = "", width = Inf, last = "")`
- ★ **'paste'** Concatenate vectors after converting to character.

Date: 04/11/2020

Q.NO 18: Question

Consider a data frame

Create a function that, given a data frame and two indexes, exchanges two values of the Code variable with each other.

For example, if the index is 1 and 3, you assign: `df[1,'Code']=df[3,'Code']` `df[3,'Code']=df[1,'Code']`

Program:

```
#Values for dataframe
Id=c(1:10)
Age=c(14,12,15,10,23,21,41,56,78,12)
Sex=c('F','M','M','F','M','F','M','M','F','M')
Code=letters[1:10]
#Creating a dataframe
df=data.frame(Id,Age,Sex,Code)

#function that, given a data frame and two indexes, exchanges two values of the Code with each other.
change_values=function(df,firstindex,secondindex)
{
  first_value=df[firstindex,'Code']
  df[firstindex,'Code']=df[secondindex,'Code']
  df[secondindex,'Code']=first_value
  return(df)
}
#Interchanging values
df=change_values(df,4,8)
df=change_values(df,9,1)
df=change_values(df,3,10)
df
```

Output:

Id	Age	Sex	Code
1	14	F	i
2	12	M	b
3	15	M	j
4	10	F	h
5	23	M	e
6	21	F	f
7	41	M	g
8	56	M	d
9	78	F	a
10	12	M	c

Explanation:

- ★ A data frame is a table or a two-dimensional array-like structure in which each column contains values of one variable and each row contains one set of values from each column.

Date: 03/11/2020

Q.NO 19: Question

Create a function that given a numeric vector, sort this in ascending order and duplicate it by two.

Program:

```
#Ascending values
sort_asc=function(vec)
{
  #Mutlipying by 2
  vec=sort(vec)*2
  return(vec)
}
#input
vec=c(1,5,9,2,5,1,6,3)
#Invoking function
sort_asc(vec)
```

Output:

2 2 4 6 10 10 12 18

Explanation:

- ★ Sort (or *order*) a vector or factor (partially) into ascending or descending order. For ordering along more than one variable
- ★ By default, sorting is ASCENDING. Prepend the sorting variable by a minus sign to indicate DESCENDING order

Date: 04/11/2020

Q.NO 20: Question

Create a function that given a numeric vector X returns the digits 0 to 9 that are not in X. If X=0 2 4 8 the function return 1 3 5 6 7 9

Program:

```
#Fuction to find missing numbers in the limit
nums=function(input)
{ #limit to check
  lim=0:9
  #Checking and saving the values missing
  output=lim[!lim %in% input]
  ##in% gives the present postion
  return(output)
}
#Assigning input
input=c(1,5,3,8)
#Invoking function
nums(input)
```

Output:

0 2 4 6 7 9

Explanation:

- ★ %in% operator is used to identify if an element belongs to a vector or Dataframe.
- ★ We can
 - select column of a dataframe in R using %in% operator.
 - create new variable of a column using %in% operator
 - drop column of a dataframe in R using %in% operator.

Date: 04/11/2020

Q.NO 21: Question

Create a function that given one word, return the position of word's letters on letters vector. For example, if the word is 'abc', the function will return 1 2 3.

Program:

```
#function to convert string to corresponding number
convert_str_num<-function(str)
{
  #extracting letters seperately
  extract = stri_extract_all(str, regex=c('\\p{L}'))
  #converts letters to corresponding numbers
  converted = letters%in%unlist(extract)
  #Returning Value
  return(which(converted))
}
str='abc'
#Invoking function
convert(str)
```

Output:

1 2 3

Explanation:

- ★ Regular expressions are the default pattern engine in stringr. That means when you use a pattern matching function with a bare string, it's equivalent to wrapping it in a call to REGEX()
- ★ Regular expressions are a concise and flexible tool for describing patterns in strings.

Date: 04/11/2020

Q.NO 22: Question

Write a code to check the given string is anagram or not

Program:

```
#Creating a function to check for anagram
anagram<-function(str1,str2)
{
  #Splicing word into letter with regex
  str1=unlist(str_extract_all(str1, regex=c('\\p{L}'))))
  str2=unlist(str_extract_all(str2, regex=c('\\p{L}'))))
  #First Checking for length
  if (length(str1)==length(str2))
  {
    #Reverse and check and return true
    match=unique(str1%in%str2==str2%in%str1)
    return( ifelse(length(str1)==length(str2) & length(match)==1,ifelse(match==TRUE,TRUE,FALSE),FALSE))
  }
  #If length is not equal diirectly returs false
  if (length(str1)!=length(str2))
    return(FALSE)
}
#Input
str1='rotator'
str2='rotator'
#Invoking function
print(anagram(str1,str2))
```

Output:

```
[1] TRUE
```

Explanation:

- ★ Regular expressions are the default pattern engine in stringr. That means when you use a pattern matching function with a bare string, it's equivalent to wrapping it in a call to REGEX()
- ★ Regular expressions are a concise and flexible tool for describing patterns in strings.
- ★ Basically, we're splitting the word up into letters. Then using the unlist function to convert this into a vector. Then sorting the vector into alphabetical order.

Date: 05/11/2020

Q.NO 23: Question

List all example files available with the `readr` library.

Program:

```
#Listing all example files available with the readr library.  
# install.packages('readr')  
print(library(readr, help, pos = 2, lib.loc = NULL))  
files=readr_example()  
print(files)
```

Output:

```
[1] "readr"      "stats"      "graphics"   "grDevices"  "utils"      "datasets"  
[7] "methods"    "base"  
[1] "challenge.csv"      "epa78.txt"      "example.log"  
[4] "fwf-sample.txt"     "massey-rating.txt" "mtcars.csv"  
[7] "mtcars.csv.bz2"     "mtcars.csv.zip"
```

Explanation:

- ★ The goal of `readr` is to provide a fast and friendly way to read rectangular data (like csv, tsv, and fwf).
- ★ It is designed to flexibly parse many types of data found in the wild, while still cleanly failing when data unexpectedly changes.
- ★ If you are new to `readr`, the best place to start is the data import chapter in R for data science.

Date: 05/11/2020

Q.NO 24: Question

Read the `mtcars.csv` file.

Program:

```
#Reading the mtcars.csv files  
print(mtcars)
```

Output:

	mpg	cyl	disp	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

Explanation:

- ★ We used the **mtcars** data set that is built-in to the **R** distribution .
- ★ **mtcars** data comes from the 1974 Motor Trend magazine.
- ★ The data includes fuel consumption data, and ten aspects of car design for then-current car models

Date: 05/11/2020

Q.NO 25: Question

Read the first 10 lines from the `mtcars.csv` file.

Program:

```
#Reading a the first 10 lines from the mtcars.csv file.  
head(mtcars,10)
```

Output:

	mpg	cyl	disp	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

Explanation:

- ★ We used the **mtcars** data set that is built-in to the **R** distribution .
- ★ **mtcars** data comes from the 1974 Motor Trend magazine.
- ★ The data includes fuel consumption data, and ten aspects of car design for then-current car models.
- ★ Head function with specified parameter gives top rows of specified datax

Date: 06/11/2020

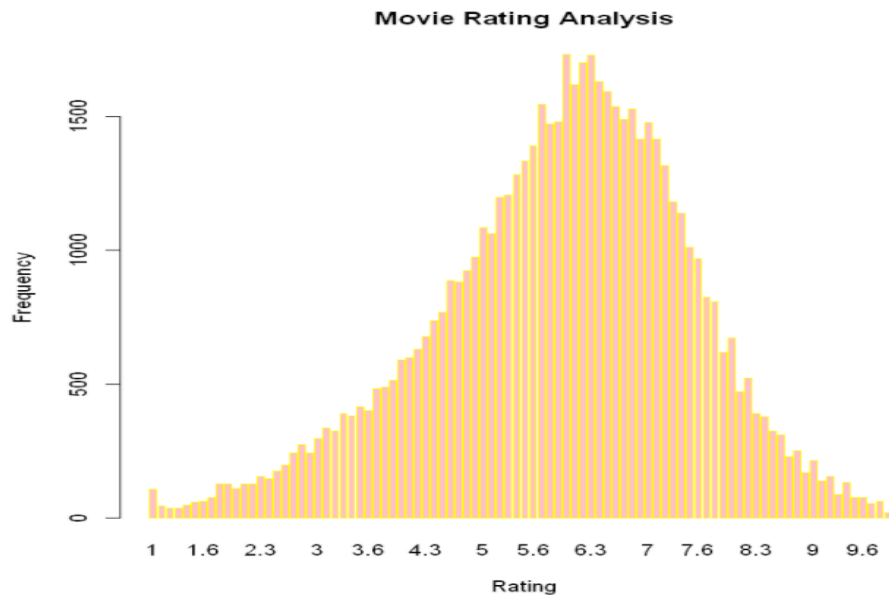
Q.NO 26: Question

Program:

Bar plot – Movies Dataset

```
#Bar Chart
#Importing Required Packages
library(ggplot2movies)
#taking rating column values from movies dataset
rating=movies$rating
#changing raw data to frequency values
table=table(rating)
#barplot(table name,horiz = true)
#default horiz is false
barplot(table,                                #tablename
        main ="Movie Rating Analysis",        #header_name
        xlab ="Rating",                      #xlabel
        ylab ="Frequency",                   #ylabel
        border="yellow",                     #bordercolour
        col = "pink"                         #colours
)
```

Output:



Explanation:

- ★ A bar chart represents data in rectangular bars with length of the bar proportional to the value of the variable. R uses the function **barplot()** to create bar charts.

Date: 06/11/2020

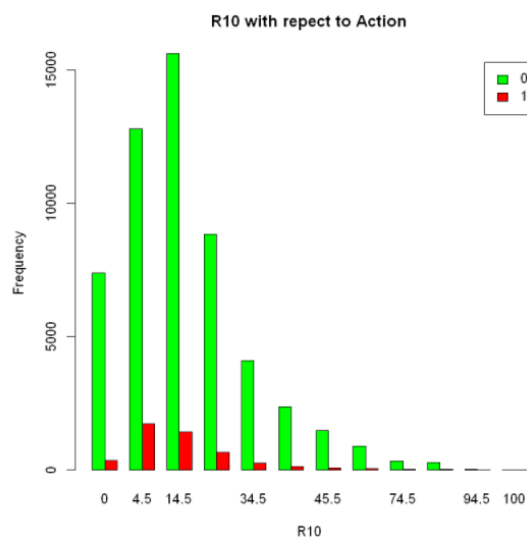
Q.NO 27: Question

Program:

Group BarGraph – Movies Dataset

```
#Bar Chart
#Importing Required Packages
library(ggplot2movies)
data=table(head(movies$r4,100),head(movies$r8,100))
# GROUPED BARGRAPH
barplot(data,
        ylab="Frequency",
        main="Double Bar Plot",
        xlab = "Rating",beside=TRUE,
        legend=rownames(data),
        col=c("green","red"))
.
```

Output:



Explanation:

- ★ A bar chart represents data in rectangular bars with length of the bar proportional to the value of the variable. R uses the function **barplot()** to create bar charts.

Date: 06/11/2020

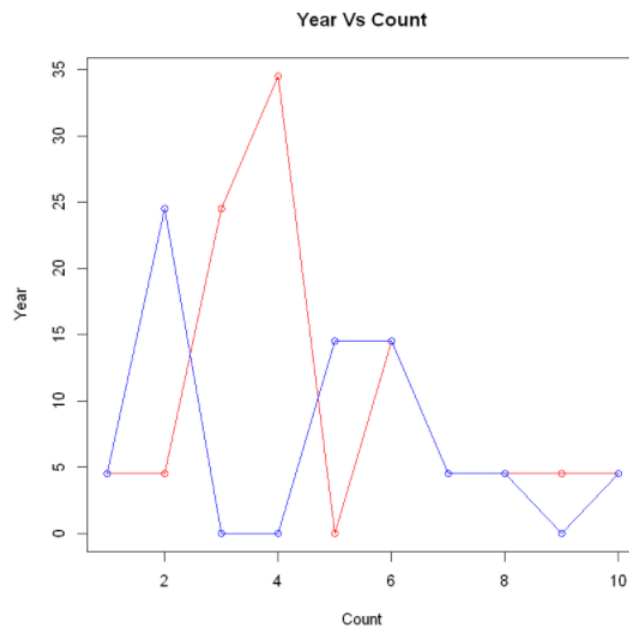
Q.NO 28: Question

Program:

Line Chart – Movies Dataset

```
#Line plot
#Importing Required Packages
library(ggplot2movies)
#Line chart only accepts numbers(no string)
plot(head(movies$r9,10),
      type = "o",
      col = "red",
      xlab = "Count",
      ylab = "Year",
      main = "Year Vs Count")
#Plotting multiple lines for comparison
lines(head(movies$r4,10), type = "o", col = "blue")
```

Output:



Explanation

- ★ A line chart is a graph that connects a series of points by drawing line segments between them. These points are ordered in one of their coordinate (usually the x-coordinate) ".

Date: 06/11/2020

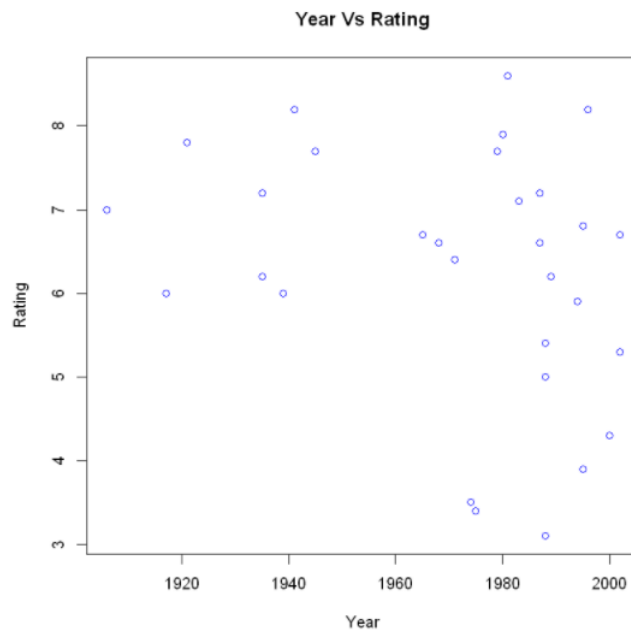
Q.NO 29: Question

Program:

Scatter plot – Movies Dataset

```
#Scatter plot
#Importing Required Packages
library(ggplot2movies)
#plot(x,y)
plot(head(movies$year,30),head(movies$rating,30),
      xlab = "Year",
      ylab = "Rating",
      main = "Year Vs Ratin",
      col = "Blue"
    )
```

Output:



Explanation:

- ★ Scatterplots show many points plotted in the Cartesian plane. Each point represents the values of two variables. One variable is chosen in the horizontal axis and another in the vertical axis.

Date: 06/11/2020

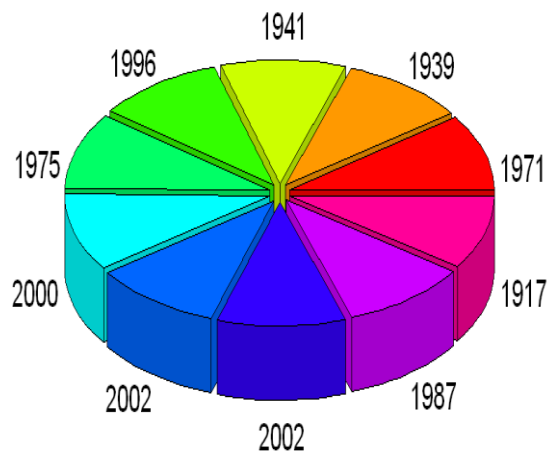
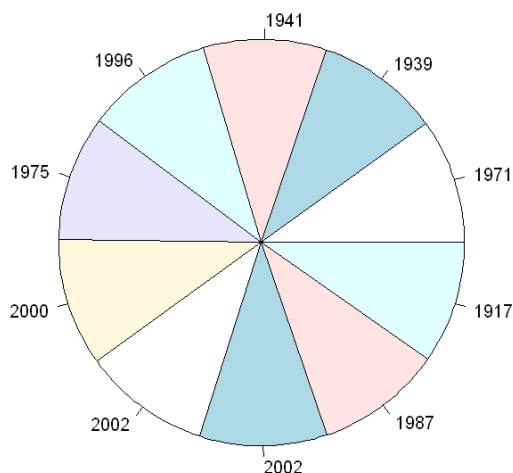
Q.NO 29: Question

Program:

Pie Chart – Movies Dataset

```
#Scatter plot
#Importing Required Packages
library(ggplot2movies)
library(plotrix)
#pie chart
pie(head(movies$year,10),labels = head(movies$year,10))
#3d pie chart
pie3D(head(movies$year,10),explode = 0.05,labels = head(movies$year,10))
```

Output:



Explanation:

- ★ pie-chart is a representation of values as slices of a circle with different colors. The slices are labeled and the numbers corresponding to each slice is also represented in the chart.

Date: 06/11/2020

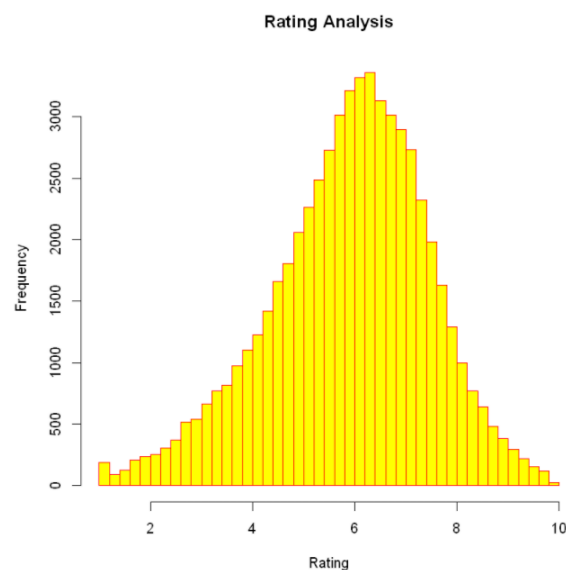
Q.NO 30: Question

Program:

Hist plot – Movies Dataset

```
#Hist plot
#Importing Required Packages
library(ggplot2movies)
#accessing ratings from movies dataset
rating=(movies$rating)
hist(rating,
     xlab = "Rating",
     col = "yellow",
     border = "red",
     breaks = 40,
     main="Rating Analysis")
```

Output:



Explanation:

- ★ A histogram represents the frequencies of values of a variable bucketed into ranges. Histogram is similar to bar chart but the difference is it groups the values into continuous ranges.

Date: 06/11/2020

Q.NO 31: Question

Program:

difference plot – Movies dataset

```
#Hist plot
#Importing Required Packages
library(ggplot2movies)
data=density(movies$rating)
plot(data,main="Movies Frequency", xlab="Rating")
#kde plotting
polygon(data,col='yellow',border="red")
|
```

Output:



Explanation:

- ★ polygon draws the polygons whose vertices are given in x and y.

Date: 06/11/2020

Q.NO 32: Question

Program:

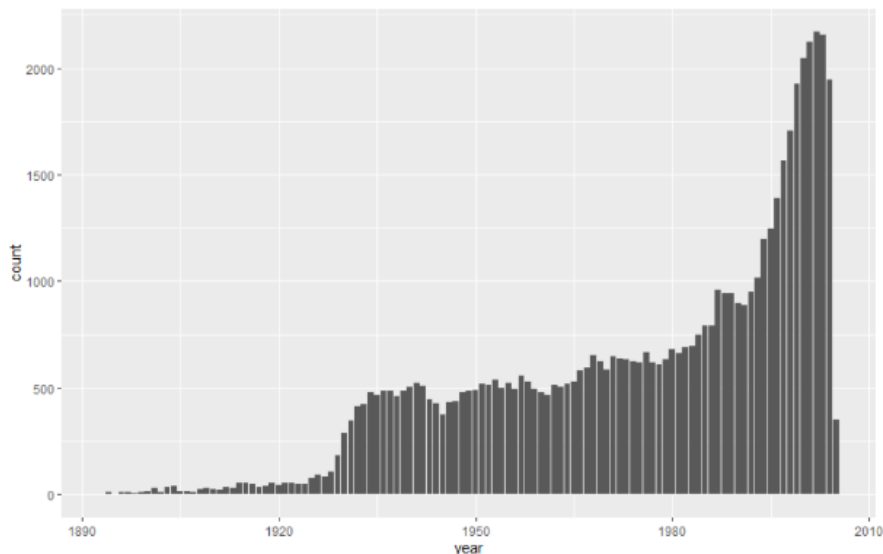
Demonstrate the ggplot2 layer

```
library(ggplot2)
library(ggplot2movies)
dplot <- ggplot(movies, aes(year))

dplot + geom_bar()
dplot + geom_bar(position = "fill")

dplot + geom_bar(position = "dodge")
```

Output:



Explanation:

- ★ ggplot() initializes a ggplot object. It can be used to declare the input data frame for a graphic and to specify the set of plot aesthetics intended to be common throughout all subsequent layers unless specifically overridden.

Date: 06/11/2020

Q.NO 33: Question

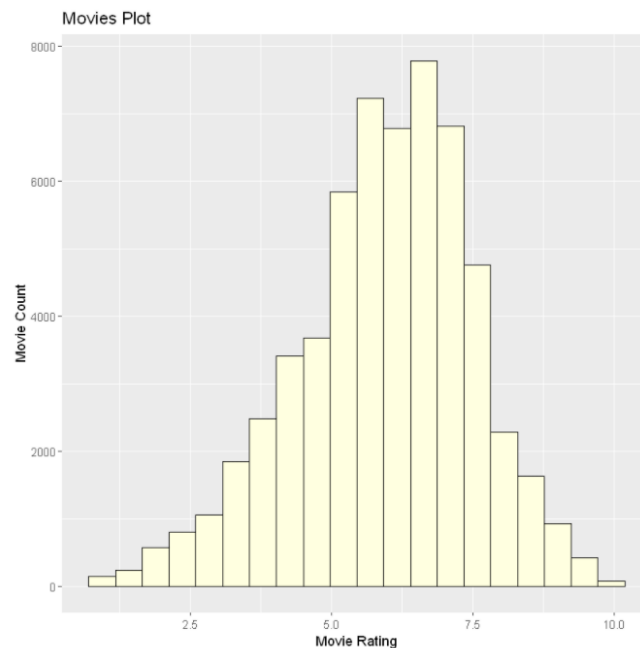
Program:

Ggplot – histogram plotting

```
library(ggplot2)
library(ggplot2movies)
movies<-ggplot(data=movies,aes(x=rating))
movies<-movies+geom_histogram(bins = 20,
                              color = "black",
                              fill="light yellow",
                              alpha=1)+xlab("Movie Rating")+
                              ylab("Movie Count")+
                              ggtitle("Movies Plot")

print(movies)
```

Output:



Explanation:

- ★ `ggplot()` initializes a ggplot object. It can be used to declare the input data frame for a graphic and to specify the set of plot aesthetics intended to be common throughout all subsequent layers unless specifically overridden.

Date: 06/11/2020

Q.NO 34: Question

Program:

Word Cloud

```
#Importing Required Packages
library(wordcloud)
data=read.csv("E:/R Programs For Practice/DAY - 5/Class/Data/file.csv",
              header= TRUE)
head(data)
#Creates a word cloud with parameters value and frequency
#wordcloud(word = dataframe_name$words,dataframe_name$words,min.freq="Le:
wordcloud(words=data$word,
          freq=data$freq,
          min.freq=2,
          max.words = 50,
          random.order = FALSE)
```

Output:



Explanation:

- ★ wordcount() counts words. Currently a "word" is a clustering of characters separated from another clustering of characters by at least 1 space. That is the law.

Date: 07/11/2020

Q.NO 35: Question

Perform the following operation:

1. T-Test (score 1 and score 2)

Program:

```
#test
Score1 <- c(3,3,3,12,15,16,17,19,23,24,32)
Score2 <- c(20,13,13,20,29,32,23,20,25,15,30)
#t.test()
print(t.test(Score1,Score2))
```

Output:

```
Welch Two Sample t-test

data:  Score1 and Score2
t = -1.9005, df = 17.977, p-value = 0.07353
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
 -13.9732668  0.7005395
sample estimates:
mean of x mean of y
 15.18182  21.81818
```

Explanation:

- ★ A t-test allows us to compare the average values of the two data sets and determine if they came from the same population.
- ★ In the above examples, if we were to take a sample of students from class A and another sample of students from class B, we would not expect them to have exactly the same mean and standard deviation.

Manual calculation for the T-value

MAR

M	T	W	T	F	S	S
•	•	1	2	3	4	•
5	6	7	8	9	10	11
12	13	14	15	16	17	18
19	20	21	22	23	24	25
26	27	28	29	30	31	•

APR

M	T	W	T	F	S	S
30	•	•	•	•	•	1
2	3	4	5	6	7	8
9	10	11	12	13	14	15
16	17	18	19	20	21	22
23	24	25	26	27	28	29

20-02-2018
Tuesday
FEBRUARY

20
Day 051-314 • Week-08

Appointments / Meetings

8 am

9

10

11

12

1 pm

2

3

4

5

6

7

Evening

☒

T-Test

Formula

$$= \frac{|\bar{X}_1 - \bar{X}_2|}{\sqrt{\frac{S_1^2}{n_1} + \frac{S_2^2}{n_2}}}$$

$\bar{X}_1 \rightarrow$ Sample 1 Mean
 $\bar{X}_2 \rightarrow$ Sample 2 Mean
 $n_1 \rightarrow$ Sample 1 Count
 $n_2 \rightarrow$ Sample 2 Count
 $S_1 \rightarrow$ Variance 1
 $S_2 \rightarrow$ Variance 2

Data

Score 1 = 3, 3, 3, 12, 15, 16, 17, 19, 23, 24, 32

Score 2 = 20, 13, 13, 20, 29, 32, 23, 20, 25, 15, 30

$\bar{X}_1 = \frac{3+3+3+12+15+16+17+19+23+24+32}{11}$
 $= 15.1818$
 $\bar{X}_2 = \frac{20+13+13+20+29+32+23+20+25+15+30}{11}$
 $= 21.8181$
 $S_1 = \frac{\sum (x_i - \bar{x})^2}{n}$
 $= 89.56364$
 $S_2 = \frac{\sum (x_i - \bar{x})^2}{n_2}$
 $= 44.56364$
 $n_1 = 11, n_2 = 11$

MAR

M	T	W	T	F	S	S
•	•	•	1	2	3	4
5	6	7	8	9	10	11
12	13	14	15	16	17	18
19	20	21	22	23	24	25
26	27	28	29	30	31	•

APR

M	T	W	T	F	S	S
30	•	•	•	•	•	1
2	3	4	5	6	7	8
9	10	11	12	13	14	15
16	17	18	19	20	21	22
23	24	25	26	27	28	29

22-02-2018

Thursday

FEBRUARY

Day 053-312 ♦ Week-08

22

Appointments / Meetings

8 am $t \text{ value} = \frac{|15.18182 - 21.818|}{\sqrt{\frac{89.564}{11} + \frac{44.564}{11}}}$

9 $= \frac{6.63618}{\sqrt{8.1421 + 4.05127}}$

10 $= \frac{6.63618}{\sqrt{12.1933}}$

11 $= \frac{6.63618}{3.49189}$

12 $= 1.9005$

1 pm

2

3

4

Excel sheet R- Program

Subject	Score1	Score2
1	3	20
2	3	13
3	3	13
4	12	20
5	15	29
6	16	32
7	17	23
8	19	20
9	23	25
10	24	15
11	32	30
Mean	15.18182	21.8182
Std	9.463807	6.6756
Variance	89.56364	44.5636
Count	11	11
T-Test (P value)	0.071879	
T Value manually	0.19004	
Degree of freedom	20	
T Value for 20 with 5 percent error (critical value)	1.725	
Critical value is higher than calculated T Value. So we can reject NULL HYPOTHESIS		

R Program

Code:

```
#Anova Test
A<-c(0,2,3,5,8,10,12)
B<-c(1,2,3,9,10,10,11)
C<-c(1,4,5,5,8,9,10)
#Making data as dataframe
combined_groups <-data.frame(cbind(A,B,C))
print(combined_groups)
#stacking into one column
stack_group = stack(combined_groups)
print(stack_group)
#applying aov anova function
anova = aov(values~ind,data = stack_group)
print(summary(anova))
```

Output:

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
ind	2	2.67	1.333	0.082	0.921
Residuals	18	291.14	16.175		

Explanation:

- ★ An **ANOVA test** is a way to find out if survey or experiment results are significant.
- ★ In other words, they help you to figure out if you need to reject the null hypothesis or accept the alternate hypothesis.
- ★ Basically, you're **testing** groups to see if there's a difference between them.

Date: 06/11/2020

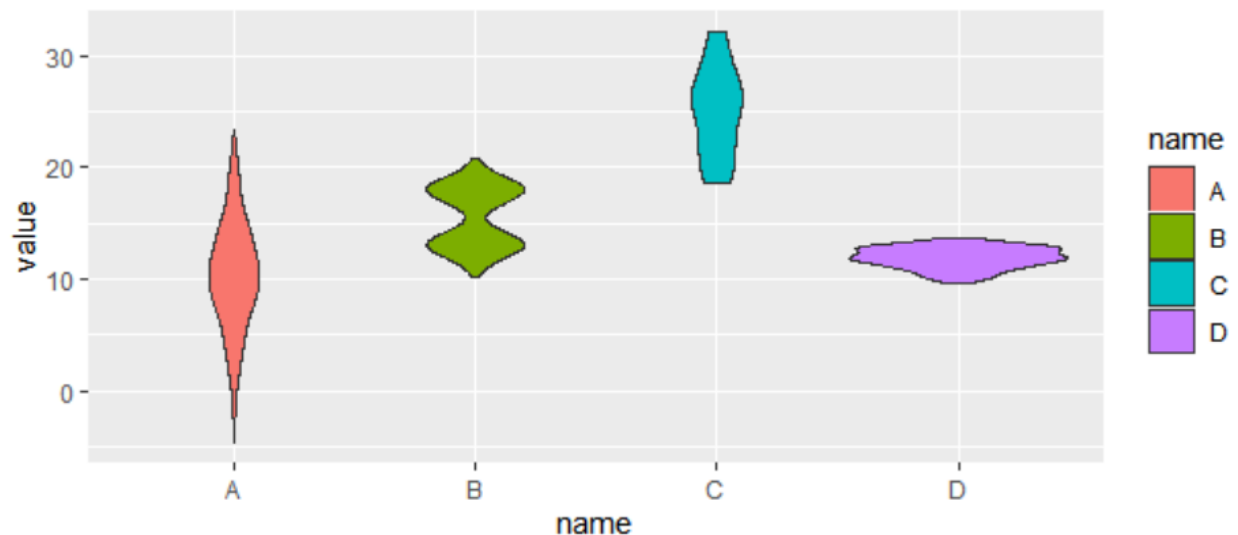
Q.NO 36: Question

Program:

Draw the Violin plot

```
#Importing Required Packages
library(ggplot2)
data <- data.frame(
  name=c( rep("A",500), rep("B",500),
          rep("B",500), rep("C",20), rep('D', 100) ),
  value=c( rnorm(500, 10, 5), rnorm(500, 13, 1),
           rnorm(500, 18, 1), rnorm(20, 25, 4), rnorm(100, 12, 1) )
)
#Ggplot()
p <- ggplot(data, aes(x=name, y=value, fill=name)) +
  geom_violin()
print(p)
```

Output:



Explanation:

- ★ A **violin plot** is a method of **plotting** numeric data. It is similar to a box **plot**, with the addition of a rotated kernel density **plot** on each side.
- ★ While a box **plot** only shows summary statistics such as mean/median and interquartile ranges, the **violin plot** shows the full distribution of the data.

Date: 07/11/2020

Q.NO 37: Question

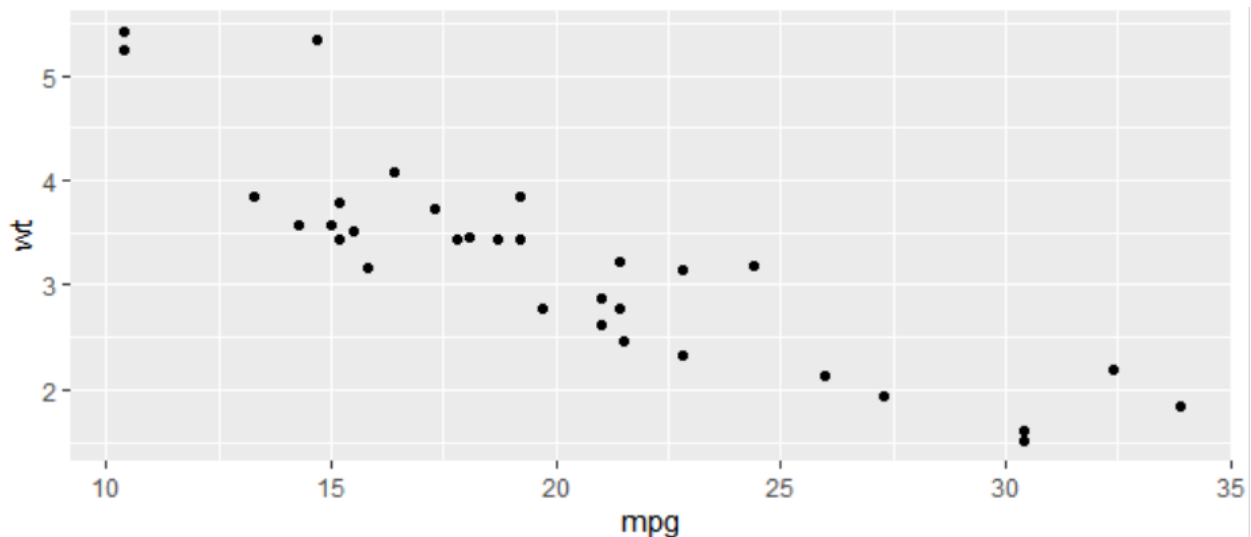
Program:

Draw the Qplot for 5 Types

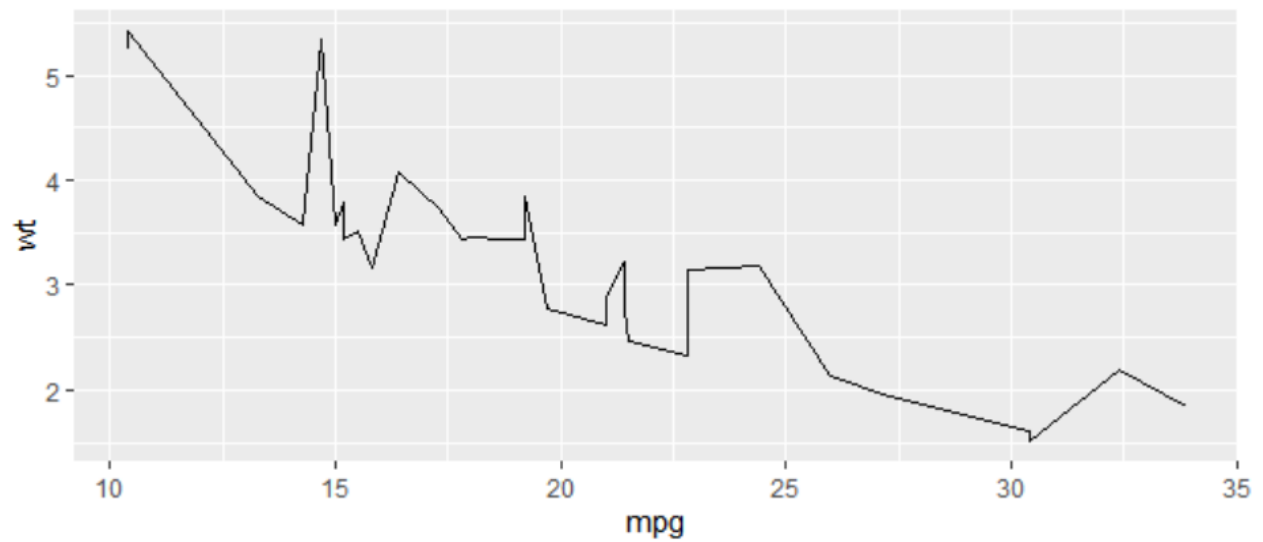
```
#qqplot
#Plotting with 5 geom
#geom = point
qplot(mpg, wt, data = mtcars, geom = "point")
#geom = line
qplot(mpg, wt, data = mtcars, geom = "line")
#geom = histogram
qplot(mpg, data = mtcars, geom = "histogram", bins = 30)
#geom = path
qplot(mpg, wt, data = mtcars, geom = "path")
#geom = smooth
qplot(mpg, wt, data = mtcars, geom = "smooth")
```

Output:

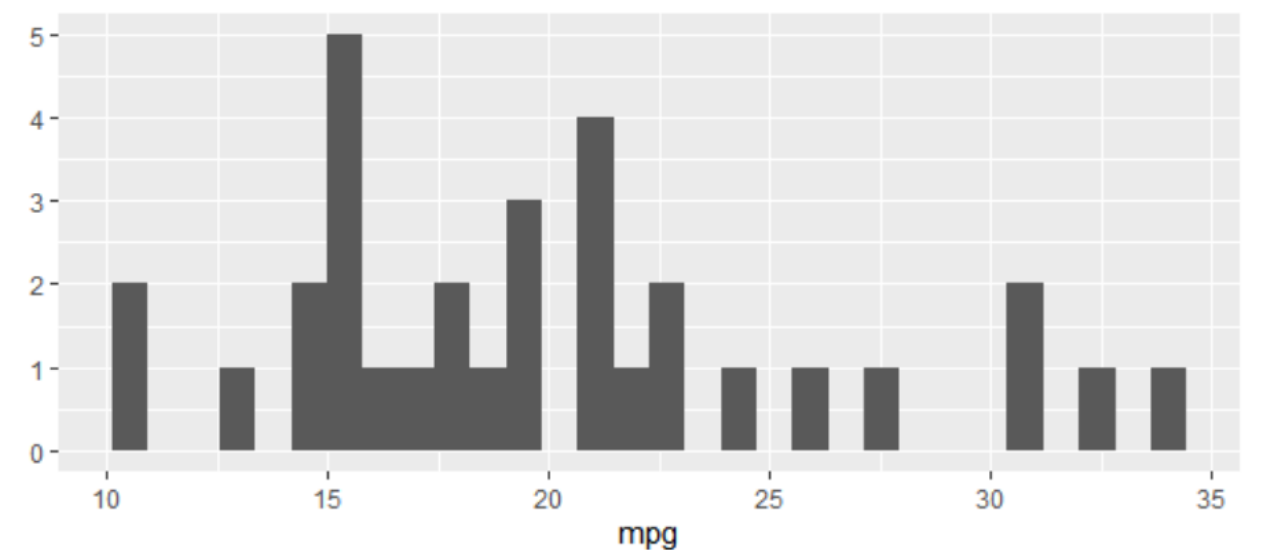
Point plot



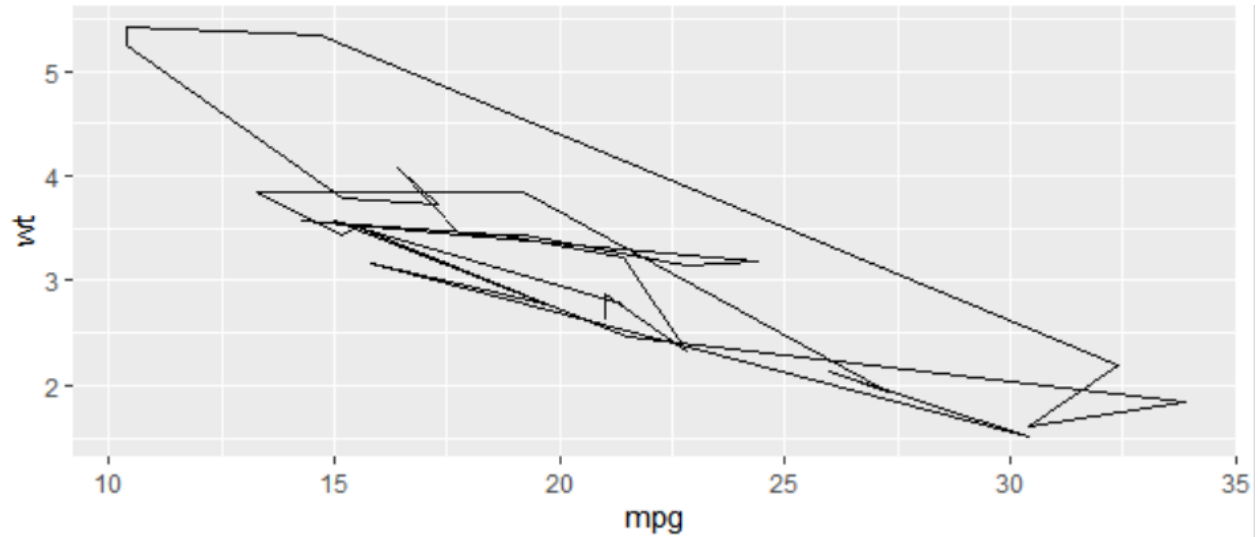
Line plot



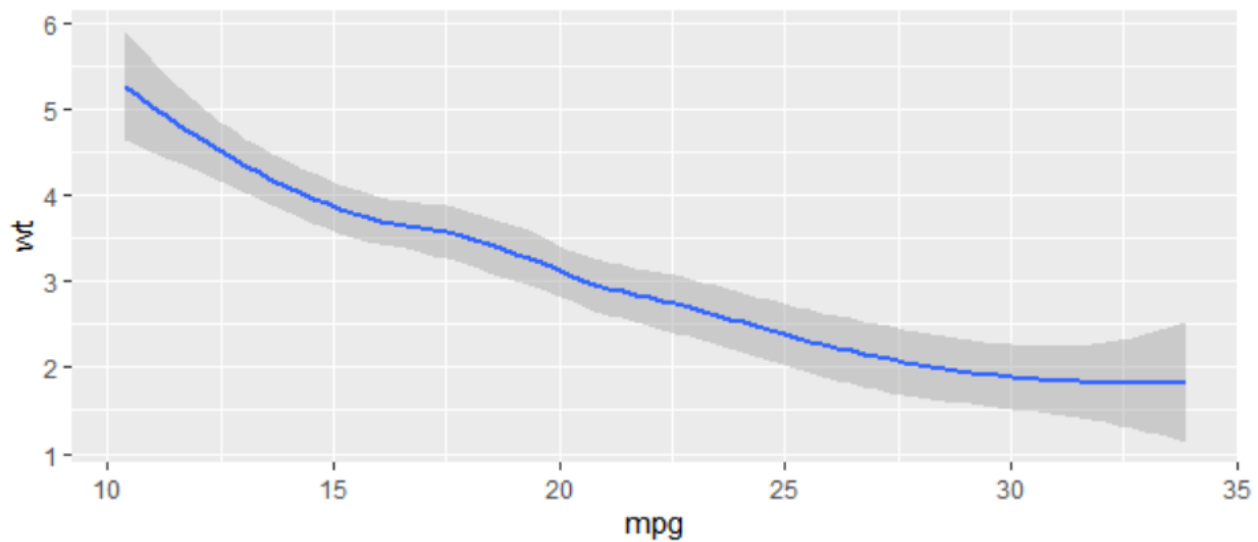
Histogram



Path



Smooth



Explanation:

- ★ **qplot()** is a shortcut designed to be familiar if you're used to base `plot()` .
- ★ It's a convenient wrapper for creating a number of different types of plots using a consistent calling scheme.

Date: 07/11/2020

Q.NO 38: Question

Program:

Demonstrate the skewness and kurtosis using built-in dataset or your own dataset to get the data summary.

```
#Importing Required Packages
library(e1071)
#Reading Csv file
df = read.csv("E:/R Programs For Practice/DAY - 6/Data/mark.csv")
#skewness
cat("Skewness",skewness(df$mark),"\n")
plot(density(df$mark))
polygon(density(df$mark), col="red", border="blue")
#kurtosis
cat("kurtosis: ",kurtosis(df$mark))|
```

Output:

```
Skewness -0.4277135
kurtosis:  -0.7261525
```

Explanation:

- ★ **Skewness** is a measure of symmetry, or more precisely, the lack of symmetry.
- ★ **Kurtosis** is a measure of whether the data are heavy-tailed or light-tailed relative to a normal distribution.

Date: 10/11/2020

Q.NO 39: Question

Program:

The coin is flipped ten times. Find the probability of 7 heads occurring.

```
#The coin is flipped ten times. Find the probability of 7 heads occurring.  
#BinomialDistribution  
cat("probability of 7 heads occurring in 10 times = ",pbinom(7,size=10,prob=0.5))  
|
```

Output:

```
> cat("probability of 7 heads occurring in 10 times = ",pbinom(7,size=10,prob=0.5))  
probability of 7 heads occurring in 10 times = 0.9453125
```

Explanation:

- ★ The **binomial distribution** is a discrete probability distribution. It describes the outcome of n independent trials in an experiment. Each trial is assumed to have only two outcomes, either success or failure.
- ★ This function gives the cumulative probability of an event. It is a single value representing the probability.

Date: 10/11/2020

Q.NO 40: Question

Program:

A card is selected three times (and replaced). Find the probability of 2 face cards occurring.

```
# A card is selected three times (and replaced). Find the probability of 2 face cards occurring
cat("probability of 2 face cards occurring in three times = ",pbinom(2,size=3,prob=1/52))
```

Output:

```
> cat("probability of 2 face cards occurring in three times = ",pbinom(2,size=3,prob=1/52))
probability of 2 face cards occurring in three times = 0.9999929
```

Explanation:

- ★ The **binomial distribution** is a discrete probability distribution. It describes the outcome of n independent trials in an experiment. Each trial is assumed to have only two outcomes, either success or failure.
- ★ This function gives the cumulative probability of an event. It is a single value representing the probability.

Date: 10/11/2020

Q.NO 41: Question

Program:

A student decides to guess on a section of his ACT test. The section contains 50 multiple choice questions and each question has 5 possible answers. Find the expected number of correct responses.

```
# A student decides to guess on a section of his ACT test.  
# The section contains 50 multiple choice questions and each question has 5 possible answers.  
# Find the expected number of correct responses.  
cat("probability of correct responses = ", pbinom(4, size=50, prob=0.2))  
|
```

Output:

```
> cat("probability of correct responses = ", pbinom(4, size=50, prob=0.2))  
probability of correct responses = 0.01849602  
> |
```

Explanation:

- ★ The **binomial distribution** is a discrete probability distribution. It describes the outcome of n independent trials in an experiment. Each trial is assumed to have only two outcomes, either success or failure.
- ★ This function gives the cumulative probability of an event. It is a single value representing the probability.

Date: 10/11/2020

Q.NO 42: Question

Program:

A company ships 5000 cell phones. They are expected to last an average of 10,000 hours before needing repair; with a standard deviation of 500 hours.

Assume the survival time of the phones are normally distributed. If a phone is randomly selected to be tracked for repairs find the expected number that needs repair,

a) after 11,000 hours

```
#A company ships 5000 cell phones.  
#They are expected to last an average of 10,000 hours before needing repair;  
#with a standard deviation of 500 hours.  
#Assume the survival time of the phones are normally distributed.  
#If a phone is randomly selected to be tracked for repairs find the expected number that ne  
#a) after 11,000 hours  
cat("Expected Number of Phones to be repaired = ",  
    pnorm(11000,mean=10000,sd=500,lower.tail=FALSE))
```

Output:

```
> cat("probability of correct responses = ",pbinom(4,size=50,prob=0.2))  
probability of correct responses = 0.01849602  
.
```

Explanation:

- ★ **pnorm** is the **R** function that calculates the c. d. f. where X is normal. Optional arguments described on the on-line documentation specify the parameters of the particular normal distribution.