

List of Programs:

1. Download and install R-Programming environment and install basic packages using install.Packages () command in R.
2. Learn all the basics of R-Programming (Data types ,Variables , Operators ,Control statements)
3. Write a R program to create a two-dimensional 5x3 array of sequence of even integers greater than 50.
4. a)Create a Scatter plot from CSV in R
b) Create a Json Files
5. Implement data frames in R. Write a program to join columns and rows in a data frame using c bind () and r bind () in R.
6. a)Write a R program to find factorial of a number using recursion
b)Write a R program to mean, variance, standard deviation for the given probability distribution
7. Implement Apriori algorithm to extract association rule of datamining.
8. Implement k-means clustering technique.
9. Implement Classification algorithm.
10. Create pie charts and bar charts using R.

Introduction to R programming:

R is a programming language and free software developed by Ross Ihaka and Robert Gentleman in 1993. R possesses an extensive catalog of statistical and graphical methods. It includes machine learning algorithms, linear regression, time series, statistical inference to name a few. Most of the R libraries are written in R, but for heavy computational tasks, C, C++ and Fortran codes are preferred. R is not only entrusted by academic, but many large companies also use R programming language, including Uber, Google, Airbnb, Facebook and so on.

Data analysis with R is done in a series of steps; programming, transforming, discovering, modeling and communicate the results.

Program: R is a clear and accessible programming tool

Transform: R is made up of a collection of libraries designed specifically for data science

Discover: Investigate the data, refine your hypothesis and analyze them

Model: R provides a wide array of tools to capture the right model for your data

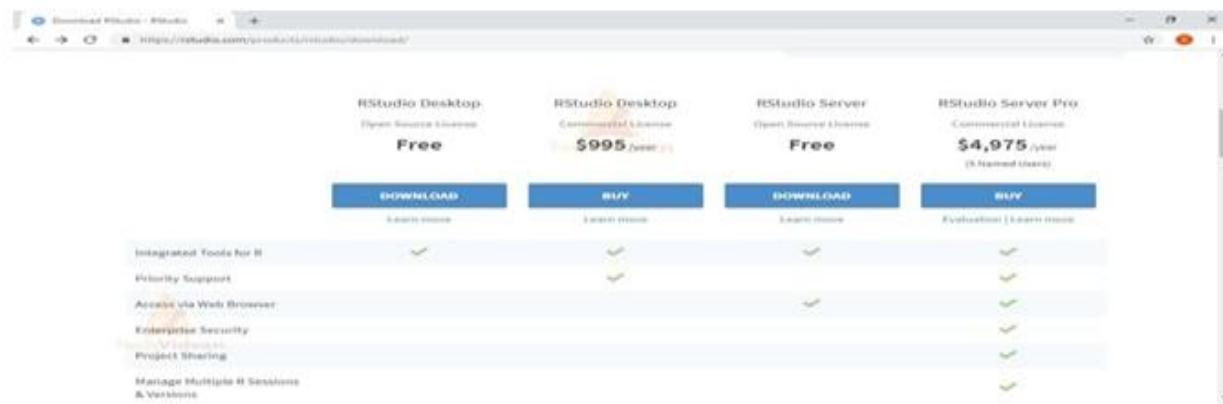
Communicate: Integrate codes, graphs, and outputs to a report with R Markdown or build Shiny apps to share with the world

What is R used for?

- Statistical inference
- Data analysis
- Machine learning algorithm

Installation of R-Studio on windows:

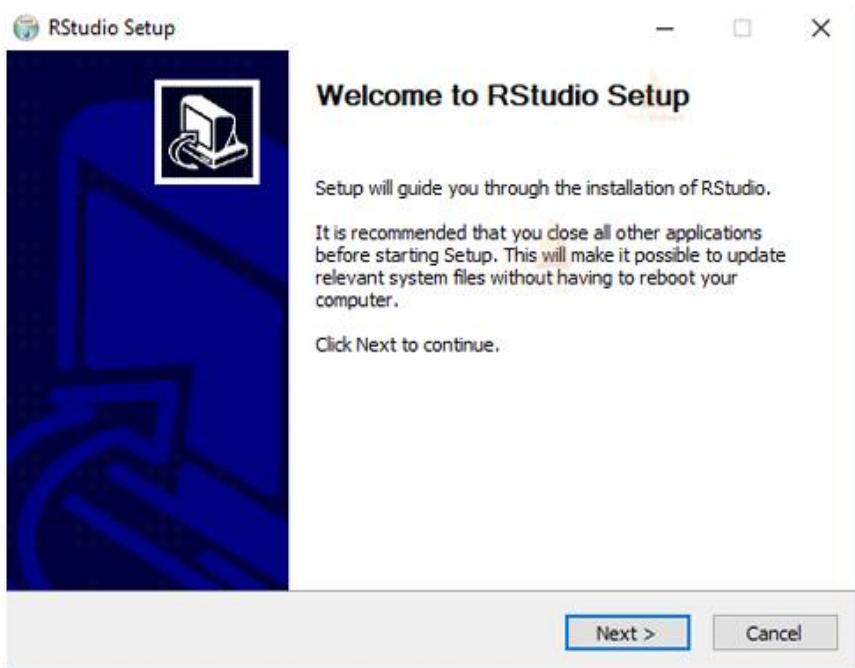
Step – 1: With R-base installed, let's move on to installing RStudio. To begin, go to download RStudio and click on the download button for RStudio desktop.



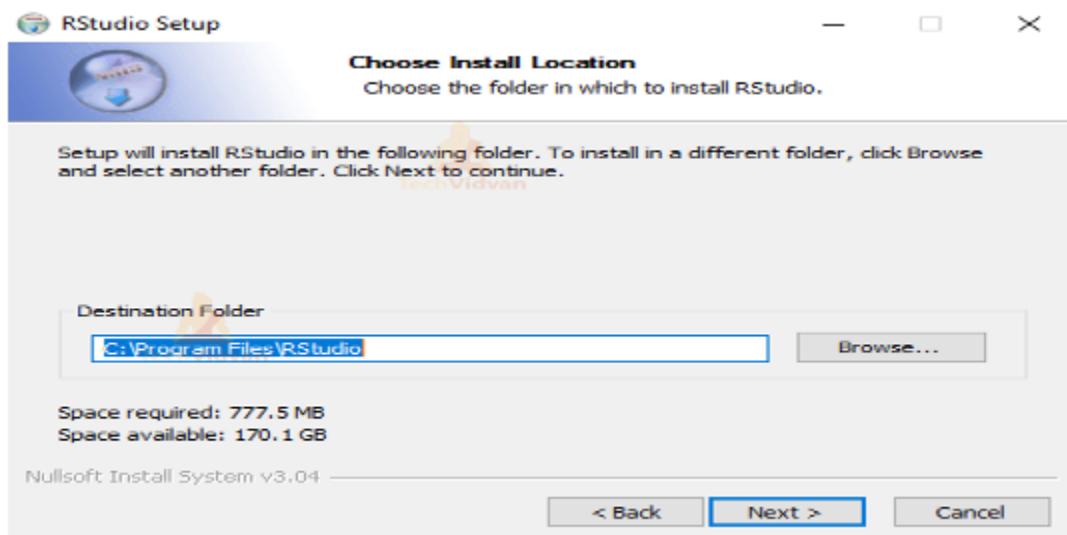
Step – 2: Click on the link for the windows version of RStudio and save the .exe file.

Step – 3: Run the .exe and follow the installation instructions.

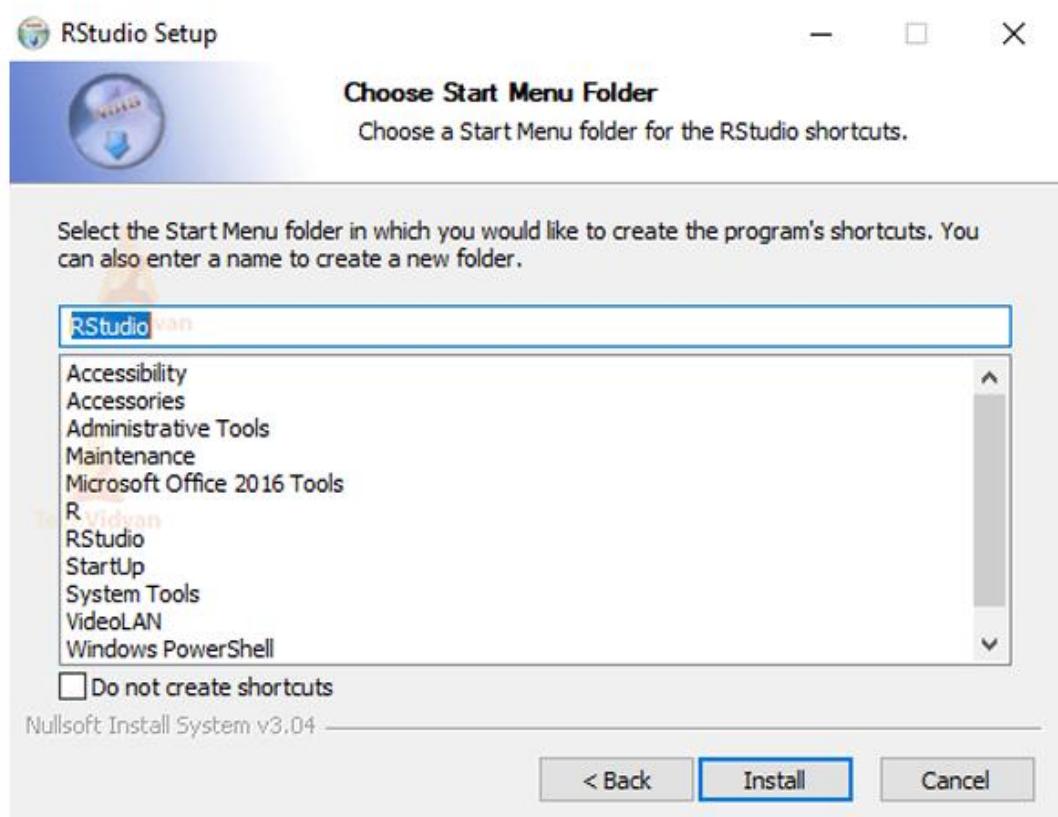
3. Click Next on the welcome window.



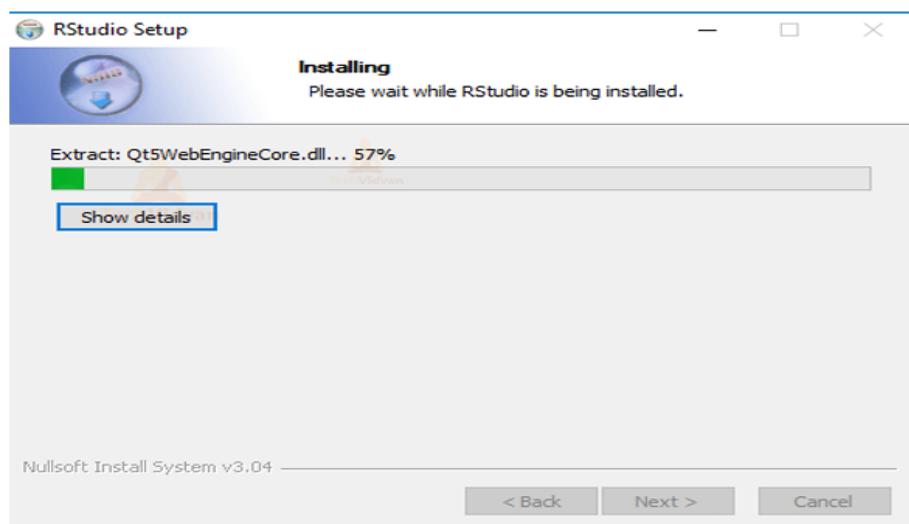
Enter/browse the path to the installation folder and click Next to proceed.



Select the folder for the start menu shortcut or click on do not create shortcuts and then click Next.



Wait for the installation process to complete.



Click Finish to end the installation.



Install the R Packages:-

In RStudio, if you require a particular library, then you can go through the following instructions:

First, run R Studio.

After clicking on the packages tab, click on install. The following dialog box will appear.

In the Install Packages dialog, write the package name you want to install under the Packages field and then click install. This will install the package you searched for or give you a list of matching packages based on your package text.

2.LEARN ALL THE BASICS OF R PROGRAMMING (Data Types, variables, operators, control statements)

Aim

The aim of the code is to demonstrate the use of basic data structures in R programming, such as vectors, lists, data frames, and matrices.

Algorithm

1. Create Vectors:

- Create a numeric vector int_vector containing integers.
- Create a character vector char_vector containing strings (fruit names).

2. Create a List:

- Create a list my_list which contains the previously created vectors (int_vector and char_vector), a numeric value (3.14), and a boolean value (TRUE).

3. Access Elements in the List:

- Access the second element of the list (the character vector char_vector) and print it.

4. Create a Data Frame:

- Create a data frame data containing three columns: Name, Age, and Score.
- Populate the columns with appropriate data (names, ages, and scores).

5. Access Columns in the Data Frame:

- Access the Name column of the data frame using the \$ operator and print it.

6. Create a Matrix:

- Create a matrix my_matrix with 2 rows and 3 columns, containing the numbers 1 to 6.

7. Access Elements in the Matrix:

- Access the element located at row 1, column 2 in the matrix and print it.

8. Operations on Data Structures:

- **Vector Operation:** Calculate the sum of the integer vector int_vector and print the result.
- **Matrix Operation:** Compute the transpose of the matrix my_matrix and print the transposed matrix.

9. Add a New Column to the Data Frame:

- Add a new column City to the data data frame, assigning each row a city name.
- Print the updated data frame to show the new column.

A.PROGRAM:

```
int_vector<-c(1,2,3,4,5)
print("Integer vector:")
print(int_vector)

char_vector<-c("apple","banana","Cherry")
print('character vector:')
print(char_vector)

my_list<-list(int_vector,char_vector,3.14,TRUE)
print("list:")
print(my_list)

print("Second element of the list(char_vector):")
print(my_list[[2]])

data<-data.frame(
  Name=c("john","mary","steve"),
  Age=c(23,25,30),
  Score=c(88.5,91.3,85.0)
)
print("Data Frame:")
print(data)

print("Accessing the 'Name'column of the data frame:")
print(data$Name)

my_matrix<-matrix(1:6,nrow=2,ncol=3)
print("matrix:")
```

```
print(my_matrix)
print("Element at row1,column2 of the matrix:")
print(my_matrix[1,2])
sum_vector<-sum(int_vector)
print("sum of the integer vector:")
print(sum_vector)
matrix_transpose<-t(my_matrix)
print("Transpose of matrix:")
print(matrix_transpose)
data$city<-c("New york","Los angeles","chicago")
print("updated data frame with 'city' column:")
print(data)
```

R-OPERATORS:

```
A<-matrix(c(1,2,5,7,2,4,6,8),nrow=4,ncol=2)
B<-matrix(c(9,10,11,12,13,14,15,16),nrow=2,ncol=4)
cat("matrix A:\n")
print(A)
cat("\n matrix B:\n")
print(B)
cat("\n matrix Multiplication of A and B(%*%):\n")
result_matrix_multiply<-A%*%B
print(result_matrix_multiply)
df<-data.frame(
  ID=c(1,2,3,4,5),
  Age=c(25,30,22,35,28),
  Salary=c(50000,60000,45000,70000,52000),
  Department=c("HR","IT","Finance","IT","HR")
```

```
)  
cat("\n Data Frame:\n")  
print(df)  
filtered_data<-df[df$Age>=30&df$salary>=50000,]  
cat("\n filtered Data(Age>=30 and salary>=50000):\n")  
print(filtered_data)  
is_high_salary<-df$salary>55000  
is_in_IT<-df$Department=="IT"  
cat("\n Salary>55000(high salary):\n")  
print(is_high_salary)  
cat("\n Department is IT:\n")  
print(is_in_IT)  
cat("\n High salary AND in IT Department:\n")  
print(is_high_salary&is_in_IT)  
cat("\n High salary OR in IT Department:\n")  
print(is_high_salary|is_in_IT)  
vector1<-c(5,10,15,20)  
vector2<-c(2,4,6,8)  
cat("\n Element-wise Addition of vectors:\n")  
print(vector1+vector2)  
cat("\n Element-wise multiplication of vectors:\n")  
print(vector1*vector2)  
cat("\n Advanced sequence generation and modulus operations:\n")  
for (i in 1:15){  
  seq_val<-i*3  
  modulus_val<-seq_val%%4  
  cat("multiple of 3:",seq_val,",modulus with 4:",modulus_val,"\n")
```

```

}

x<-5

y<-10

z<-20

cat("\n Assignment with nested Expressions:\n")

result<-(x<-x+2)*(y<-y-3)+(z<-z/2)

cat("final result with complex assignment and expression:",result,"\n")

fruits<-c("apple","banana","cherry","data")

search_fruits<-c("banana","fig","data")

cat("\n check if search fruits are in the fruit list(%in%):\n")

print(search_fruits %in% fruits)

na_vector<-c(1,NA,3,4,NA)

cat("\n Relational operators with NA-safe comparisons:\n")

cat("Is NA equal to 3:",NA==3,"\n")

cat("Is NA not equal to 3:",NA!=3,"\n")

cat("Is NA less than 3:",NA<3,"\n")

cat("\n checking NA values using is.na():\n")

print(is.na(na_vector))

matrix_result<-(A*2)> 10

cat("\n complex matrix Element-wise operations with relational operations:\n")

print(matrix_result)

```

CONTROL-STATEMENT:

```

x<-10

if(x>5){

  print("x is greater than 5")

}else{

```

```
print("x is less than or equal to 5")
}

y<-7

if(y>10){

  print("y is greater than 10")

}else if(y==7){

  print("y is equal to 7")

}else{

  print("y is less than 7")

}

for(i in 1:5){

  print(paste("current value of i:",i))

}

count<-1

while(count<=3){

  print(paste("current count is:",count))

  count<-count+1

}

day<-3

result<-switch(day,

  "sunday"="weekend",
  "monday"="weekday",
  "tuesday"="weekday",
  "wednesday"="weekday",
  "thursday"="weekday",
  "friday"="weekday",
  "saturday"="weekend",
  "Invalid day")

print(paste("The day is:",result))
```

OUTPUT:

```
[1] "Integer vector:"  
[1] 1 2 3 4 5  
[1] "character vector:"  
[1] "apple" "banana" "Cherry"  
[1] "list:"  
[[1]]  
[1] 1 2 3 4 5  
[[2]]  
[1] "apple" "banana" "Cherry"  
[[3]]  
[1] 3.14  
[[4]]  
[1] TRUE  
[1] "Second element of the list(char_vector):"  
[1] "apple" "banana" "Cherry"  
[1] "Data Frame:"  
    Name Age Score  
1  john 23 88.5  
2  mary 25 91.3  
3 steve 30 85.0  
[1] "Accessing the 'Name' column of the data frame:"  
[1] "john" "mary" "steve"  
[1] "matrix:"  
    [,1] [,2] [,3]  
[1,] 1 3 5  
[2,] 2 4 6  
[1] "Element at row1,column2 of the matrix:"  
[1] 3  
[1] "sum of the integer vector:"  
[1] 15  
[1] "Transpose of matrix:"  
    [,1] [,2]  
[1,] 1 2  
[2,] 3 4  
[3,] 5 6  
[1] "updated data frame with 'city' column:"  
    Name Age Score      city  
1  john 23 88.5  New york  
2  mary 25 91.3 Los angeles
```

```
3 steve 30 85.0 chicago
```

matrix A:

```
[,1] [,2]  
[1,] 1 2  
[2,] 2 4  
[3,] 5 6  
[4,] 7 8
```

matrix B:

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

matrix Multiplication of A and B(%*%):

```
[,1] [,2] [,3] [,4]  
[1,] 29 35 41 47  
[2,] 58 70 82 94  
[3,] 105 127 149 171  
[4,] 143 173 203 233
```

Data Frame:

ID	Age	Salary	Department
1	25	50000	HR
2	30	60000	IT
3	22	45000	Finance
4	35	70000	IT
5	28	52000	HR

filtered Data(Age>=30 and salary>=55000):

```
[1] ID      Age      Salary    Department  
<0 rows> (or 0-length row.names)
```

Salary>55000(high salary):

logical(0)

Department is IT:

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

High salary AND in IT Department:

logical(0)

High salary OR in IT Department:

logical(0)

Element-wise Addition of vectors:

```
[1] 7 14 21 28
```

Element-wise multiplication of vectors:

```
[1] 10 40 90 160
```

Advanced sequence generation and modulus operations:

```
multiple of 3: 3 ,modulus with 4: 3
multiple of 3: 6 ,modulus with 4: 2
multiple of 3: 9 ,modulus with 4: 1
multiple of 3: 12 ,modulus with 4: 0
multiple of 3: 15 ,modulus with 4: 3
multiple of 3: 18 ,modulus with 4: 2
multiple of 3: 21 ,modulus with 4: 1
multiple of 3: 24 ,modulus with 4: 0
multiple of 3: 27 ,modulus with 4: 3
multiple of 3: 30 ,modulus with 4: 2
multiple of 3: 33 ,modulus with 4: 1
multiple of 3: 36 ,modulus with 4: 0
multiple of 3: 39 ,modulus with 4: 3
multiple of 3: 42 ,modulus with 4: 2
multiple of 3: 45 ,modulus with 4: 1
```

Assignment with nested Expressions:

```
final result with complex assignment and expression: 59
```

check if search fruits are in the fruit list(%in%):

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

Relational operators with NA-safe comparisons:

```
Is NA equal to 3: NA
```

```
Is NA not equal to 3: NA
```

```
Is NA less than 3: NA
```

checking NA values using is.na():

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

complex matrix Element-wise operations with relational operations:

```
[,1] [,2]
```

```
[1,] FALSE FALSE
```

```
[2,] FALSE FALSE
```

```
[3,] FALSE TRUE
```

```
[4,] TRUE TRUE
```

```
[1] "x is greater than 5"
```

```
[1] "y is equal to 7"
```

```
[1] "current value of i: 1"
```

```
[1] "current value of i: 2"
```

```
[1] "current value of i: 3"
```

```
[1] "current value of i: 4"
```

```
[1] "current value of i: 5"
```

```
[1] "current count is: 1"
[1] "current count is: 2"
[1] "current count is: 3"
[1] "The day is: weekday"
```

B.PROGRAM

```
num_var<-42
char_var<-"Hello"
log_var<-TRUE
vec_var<-c(1,2,3,4,5)
cat("Numeric variable:",num_var,"\n")
cat("Character variable:",char_var,"\n")
cat("Logical variable:",log_var,"\n")
cat("Vector variable:",vec_var,"\n\n")
x<-10
y<-5
cat("Addition :",x+y,"\n")
cat("Greater than comparison:",x>y," \n")
cat("Logical AND(x>5 AND y>5):",x>5 && y>5," \n\n")
if(x>y)
{
  cat("x is greater than y \n")
} else {
  cat("x is not greater than y \n")
}
cat("\n using for loop:\n")
for(i in vec_var)
{
  cat("value:",i," \n")
}
cat("\n using while loop:\n")
i<-1
while(i<=2)
{
  cat("Iteration:",i," \n")
  i<-i+1
}
```

OUTPUT:

Numeric variable: 42

Character variable: Hello

Logical variable: TRUE

Vector variable: 1 2 3 4 5

Addition : 15

Greater than comparison: TRUE

Logical AND(x>5 AND y>5): FALSE

x is greater than y

value: 1

value: 2

value: 3

value: 4

value: 5

Iteration: 1

Iteration: 2

3.To Create a two-dimensional 5x3 array of sequence of even integers greater than 50

Aim :

To create a R program to generate and display a two-dimensional array (5x3) filled with even integers, each of which is greater than 50.

Algorithm:

1. Initialize an empty 5x3 array.
2. Start with the first even integer greater than 50 (which is 52).
3. Iterate and place the next even integers (increment by 2) in the array until the array is filled.

A.PROGRAM

```
a<-array(seq(from=50,length.out=15,by=2),c(5,3))
print("content of the array")
print("5*3 array of sequence of even integer greater than 50:")
print(a)
```

OUTPUT:

```
[1] "content of the array"
[1] "5*3 array of sequence of even integer greater than 50:"
 [,1] [,2] [,3]
[1,] 50 60 70
[2,] 52 62 72
[3,] 54 64 74
[4,] 56 66 76
[5,] 58 68 78
```

B.PROGRAM:

```
x<-array(seq(from=100,length.out=16,by=13),c(4,4))
print("Content of the array")
print("4*4 array of sequence of even integer greater than 100:")
print(x)
```

OUTPUT:

```
[1] "Content of the array"
[1] "4*4 array of sequence of even integer greater than 100:"
[1] [2] [3] [4]
[1,] 100 152 204 256
[2,] 113 165 217 269
[3,] 126 178 230 282
[4,] 139 191 243 295
```

4. Scatter Plot from CSV in R

Aim:

To create a scatter plot using data from a CSV file in R programming. The program will read the data from a CSV file, extract the relevant variables for plotting, and generate a scatter plot to visually represent the relationship between two numerical variables.

Algorithm:

1. Load Required Libraries: Use necessary libraries such as ggplot2 for visualization and readr or utils to read the CSV file.
2. Read the CSV File: Import the data from a CSV file into a data frame.
3. Extract Data: Identify and extract the two columns of numerical data that you want to plot along the X and Y axes.
4. Plot the Data: Use the ggplot() function or base plotting methods in R to generate a scatter plot.
5. Customize the Plot: Add labels, titles, and customize the appearance of the plot.
6. Display the Plot: Display the scatter plot to the user.

4A1.PROGRAM:

```
data<-read.csv("D:/sample R/record.csv")
print(data)
print(is.data.frame(data))
print(ncol(data))
print(nrow(data))
max_sal<-max(data$SALARY)
print(max_sal)
details<-subset(data,DEPARTMENT=="IT")
print(details)
details<-subset(data,DEPARTMENT=="HR"&SALARY>600)
print(details)
details1<-subset(data,as.Date(START_DATE)>as.Date("02/06/2024"))
print(details1)
details<-subset(data,as.Date(START_DATE)>as.Date("25/06/2023"))
write.csv(details,"output.csv")
new_details<-read.csv("output.csv")
print(new_details)
plot(x=data$YEAR.OF.EXPERIENCE,
```

```

y=data$SALARY,
xlim=c(0,20),
xlab="NAME",
ylab="SALARY",
col="red",
main="Sactter plot")

```

Data set:

ID	NAME	SALARY	START_DATE	DEPARTMENT	YEAR OF EXPERIENCE
1	AARAV	80000	25-11-2023	IT	5
2	ANANYA	45000	10/5/2023	OPERATOR	3
3	ISHAAN	650000	7/8/2024	HR	10
4	PRIYA	56000	1/9/2024	MANAGER	6
5	RAHUL	90000	11/11/2024	DOCTOR	5
6	SAANVI	50000	5/8/2023	SI	4
7	ARJUN	85000	12/12/2023	PROGRAMMER	5
8	DIYA	45000	4/7/2024	ASST.PROFESSOR	4
9	VISHAL	100000	17-08-2024	DIRECTOR	9
10	ADITI	520000	15-04-2023	DEAN	12
11	RAVI	120000	2/6/2024	IT	13
12	NEHA	45000	3/9/2023	OPERATOR	3
13	KARAN	650000	14-11-2023	HR	10
14	SIMRAN	56000	8/2/2023	MANAGER	6
15	RITIKA	90000	6/8/2024	DOCTOR	5
16	SIVANI	50000	8/12/2023	SI	4
17	ARUN	85000	21-06-2024	PROGRAMMER	6
18	SIVARAJ	4500	25-11-2023	ASST.PROFESSOR	1
19	RAJU	100000	10/8/2023	DIRECTOR	10
20	ARJUN	5200000	7/8/2024	DEAN	12

OUTPUT:

ID	NAME	SALARY	START_DATE	DEPARTMENT	YEAR.OF.EXPERIENCE
1 1	AARAV	80000	25-11-2023	IT	5
2 2	ANANYA	45000	10/5/2023	OPERATOR	3
3 3	ISHAAN	650000	7/8/2024	HR	10
4 4	PRIYA	56000	1/9/2024	MANAGER	6
5 5	RAHUL	90000	11/11/2024	DOCTOR	5
6 6	SAANVI	50000	5/8/2023	SI	4
7 7	ARJUN	85000	12/12/2023	PROGRAMMER	5
8 8	DIYA	45000	4/7/2024	ASST.PROFESSOR	4
9 9	VISHAL	100000	17-08-2024	DIRECTOR	9
10 10	ADITI	520000	15-04-2023	DEAN	12
11 11	RAVI	120000	2/6/2024	IT	13
12 12	NEHA	45000	3/9/2023	OPERATOR	3
13 13	KARAN	650000	14-11-2023	HR	10
14 14	SIMRAN	56000	8/2/2023	MANAGER	6
15 15	RITIKA	90000	6/8/2024	DOCTOR	5
16 16	SIVANI	50000	8/12/2023	SI	4
17 17	ARUN	85000	21-06-2024	PROGRAMMER	6
18 18	SIVARAJ	4500	25-11-2023	ASST.PROFESSOR	1
19 19	RAJU	100000	10/8/2023	DIRECTOR	10
20 20	ARJUN	5200000	7/8/2024	DEAN	12

[1] TRUE

[1] 6

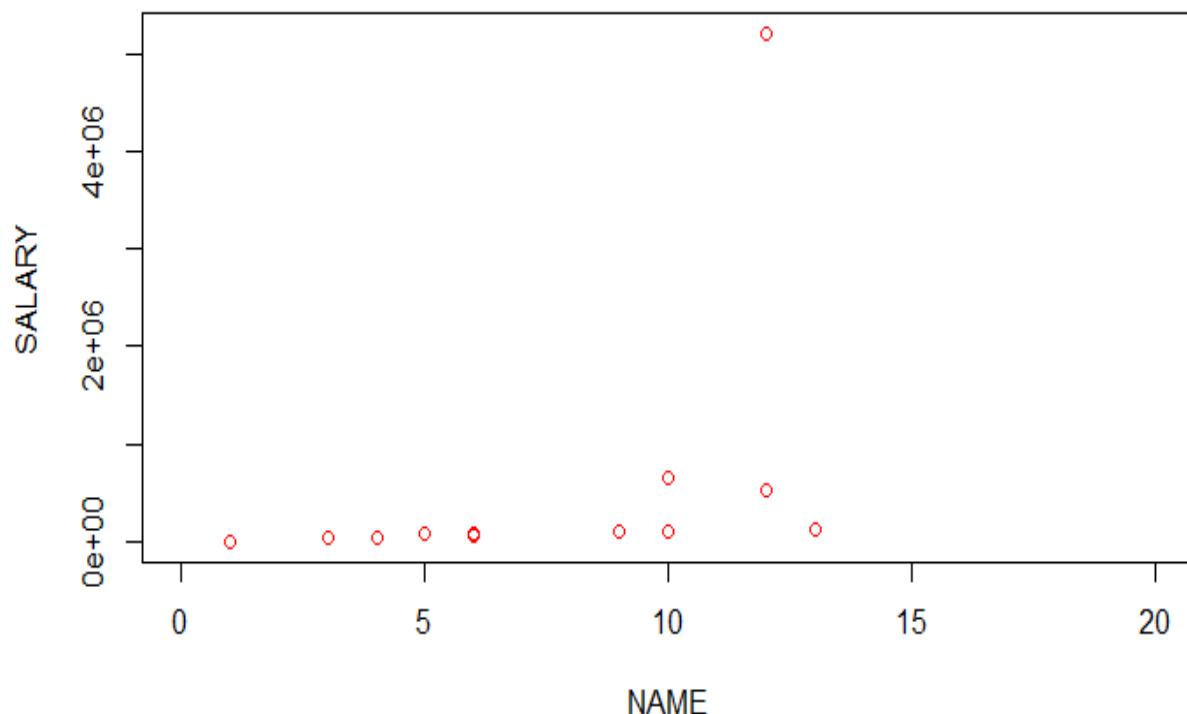
[1] 20

[1] 5200000

ID	NAME	SALARY	START_DATE	DEPARTMENT	YEAR.OF.EXPERIENCE
1 1	AARAV	80000	25-11-2023	IT	5
11 11	RAVI	120000	2/6/2024	IT	13
ID	NAME	SALARY	START_DATE	DEPARTMENT	YEAR.OF.EXPERIENCE
3 3	ISHAAN	650000	7/8/2024	HR	10
13 13	KARAN	650000	14-11-2023	HR	10
ID	NAME	SALARY	START_DATE	DEPARTMENT	YEAR.OF.EXPERIENCE
1 1	AARAV	80000	25-11-2023	IT	5

X	ID	NAME	SALARY	START_DATE	DEPARTMENT	YEAR.OF.EXPERIENCE
9	9	VISHAL	100000	17-08-2024	DIRECTOR	9
10	10	ADITI	520000	15-04-2023	DEAN	12
13	13	KARAN	650000	14-11-2023	HR	10
17	17	ARUN	85000	21-06-2024	PROGRAMMER	6
18	18	SIVARAJ	4500	25-11-2023	ASST.PROFESSOR	1
1	1	AARAV	80000	25-11-2023	IT	5
2	18	SIVARAJ	4500	25-11-2023	ASST.PROFESSOR	1

Sactter plot



4 1) B) PROGRAM:

PROGRAM:

```
data<-read.csv("D:/sample R/students_data.csv")
print(data)
file_path<-"D:/sample R/students_data.csv"
students_data<-read.csv("D:/sample R/students_data.csv")
head(students_data)
str(students_data)
summary(students_data)
```

NAME	AGE	SCORE
Alice	23	88
Bob	22	95
Charlie	21	78
David	24	85
Eve	22	92

OUTPUT:

	NAME	AGE	SCORE
1	Alice	23	88
2	Bob	22	95
3	Charlie	21	78
4	David	24	85
5	Eve	22	92

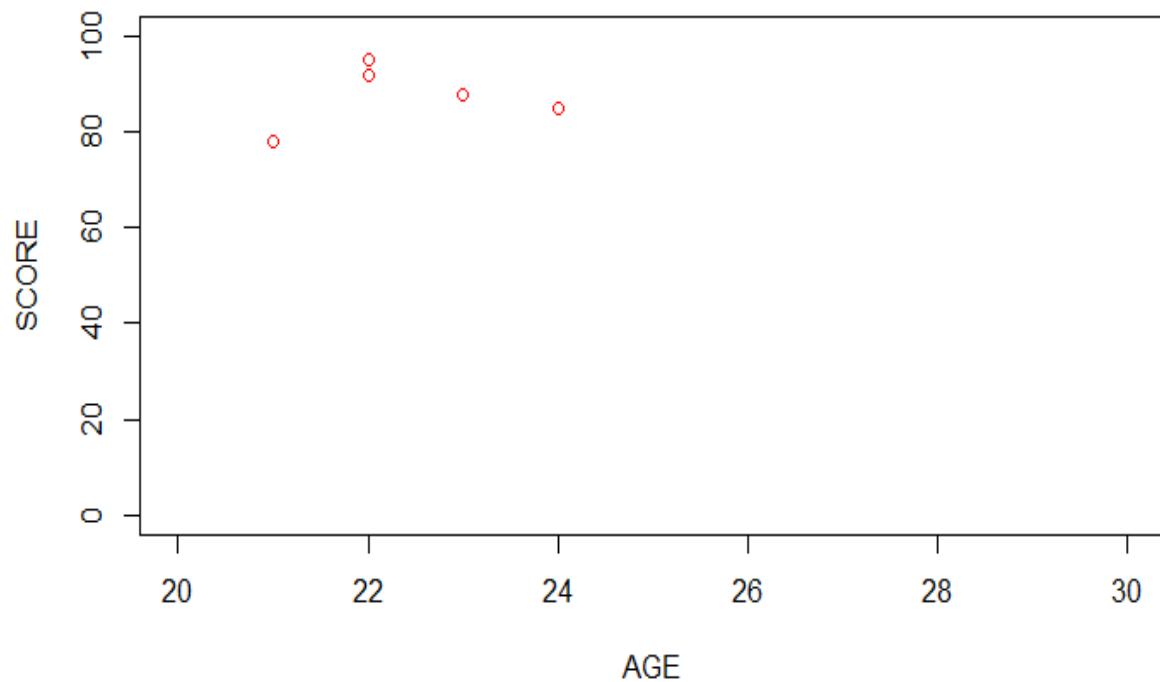
	NAME	AGE	SCORE
1	Alice	23	88
2	Bob	22	95
3	Charlie	21	78
4	David	24	85
5	Eve	22	92

'data.frame': 5 obs. of 3 variables:

```
$ NAME : chr "Alice" "Bob" "Charlie" "David" ...
$ AGE  : int 23 22 21 24 22
$ SCORE: int 88 95 78 85 92
```

	NAME	AGE	SCORE
Length:5		Min. :21.0	Min. :78.0
Class :character		1st Qu.:22.0	1st Qu.:85.0
Mode :character		Median :22.0	Median :88.0
		Mean :22.4	Mean :87.6
		3rd Qu.:23.0	3rd Qu.:92.0
		Max. :24.0	Max. :95.0

Scatter Plot



4 b. JSON FILE

Aim:

To demonstrate how to handle JSON (JavaScript Object Notation) data in R programming. JSON is a widely used data format for representing structured data, commonly used in APIs and web applications. In this program, we will explore how to read, write, and manipulate JSON data using R.

Algorithm:

1. Install and Load the JSON Package:

- First, ensure that the jsonlite package is installed and loaded.

2. Reading JSON Data:

- Use the fromJSON() function to parse JSON data from a string or a file into an R object (e.g., a list, data frame).

3. Writing JSON Data:

- Use the toJSON() function to convert an R object into JSON format, and optionally write it to a file.

4. Manipulating JSON Data:

- Once the JSON data is read into an R object (typically a list or data frame), it can be manipulated as needed (e.g., adding/removing elements, modifying values).

5. Saving JSON Data:

- After making modifications, the R object can be converted back into JSON format and saved to a file.

JSON CODE:

```
{  
  "ID": ["1", "2", "3", "4", "5"],  
  "NAME": ["sivaranjani", "snegha", "kavya", "lakshmi", "soniya"],  
  "SALARY": [722.5, 815.2, 1611, 2829, 843.25],  
  "STARTDATE": ["6/17/2014", "1/1/2012", "11/15/2024", "9/23/2022", "10/12/2023"],  
  "DEPT": ["IT", "IT", "HR", "OPERATION", "FINANCE"]}
```

}

A.PROGRAM:

```
Library("rjson")
Result<-fromJSON(file="D:/sample R/j2.json")
Print(result)
```

OUTPUT:

ID

```
[1] "1","2","3","4","5"
```

NAME

```
[1] "sivaranjani","snegha","kavya","lakshmi","soniya"
```

SALARY

```
[1] "722.5","815.2","1611","2829","843.25"
```

STARTDATE

```
[1] "6/17/2014","1/1/2012","11/15/2024","9/23/2022","10/12/2023"
```

DEPT

```
[1] "IT","IT","HR","OPERATION","FINANCE"
```

B.PROGRAM

JSON CODE:

```
{  
"ID":["1","2","3","4","5"],  
"NAME":["sivaranjani","snegha","kavya","lakshmi","soniya"],  
"SALARY":["722.5","815.2","1611","2829","843.25"],  
"STARTDATE":["6/17/2014","1/1/2012","11/15/2024","9/23/2022","10/12/2023"],  
"DEPT":["IT","IT","HR","OPERATION","FINANCE"]  
}
```

CODE:

```
library(rjson)  
  
json_data <- fromJSON(file="D:/sample R/j2.json")  
  
print(json_data)  
  
if ("name" %in% names(json_data)) {  
  print(paste("Name: ", json_data$name))  
}  
else {  
  print("Name not found in the JSON data.")  
}  
  
if ("address" %in% names(json_data)) {  
  address <- json_data$address  
  print(paste("City: ", address$city))  
  print(paste("Zip Code: ", address$zip_code))  
}  
else {  
  print("Address not found in the JSON data.")
```

```
}

if ("items" %in% names(json_data)) {

  items <- json_data$items

  for (item in items) {

    print(paste("Item Name: ", item$name))

    print(paste("Item Price: ", item$price))

  }

} else {

  print("Items array not found in the JSON data.")

}
```

OUTPUT:

\$ID

```
[1] "1" "2" "3" "4" "5"
```

\$NAME

```
[1] "sivaranjani" "snegha"    "kavya"     "lakshmi"   "soniya"
```

\$SALARY

```
[1] "722.5"  "815.2"  "1611"   "2829"   "843.25"
```

\$STARTDATE

```
[1] "6/17/2014" "1/1/2012" "11/15/2024" "9/23/2022" "10/12/2023"
```

\$DEPT

```
[1] "IT"      "IT"      "HR"      "OPERATION" "FINANCE"
```

```
[1] "Name not found in the JSON data."
```

```
[1] "Address not found in the JSON data."
```

```
[1] "Items array not found in the JSON data."
```

5.Data Frame using CBind() and RBind() in R

Aim:

To demonstrate how to use the cbind() and rbind() functions in R to create and manipulate data frames. These functions allow for column-wise (cbind()) and row-wise (rbind()) binding of data, which is useful for combining multiple datasets into a single data frame.

Definitions:

- **cbind()**: This function is used to combine objects by columns. It binds two or more vectors, matrices, or data frames side by side.
- **rbind()**: This function is used to combine objects by rows. It stacks two or more vectors, matrices, or data frames on top of each other.

Key Concepts:

- **Column Binding (cbind())**: When you combine data by columns, the number of rows must be the same across all objects.
- **Row Binding (rbind())**: When you combine data by rows, the number of columns must be the same across all objects.

Algorithm:

1. **Create Data**: Start by creating individual vectors or data frames that you want to combine.
2. **Column Binding (cbind())**: Use cbind() to combine the data objects by columns.
3. **Row Binding (rbind())**: Use rbind() to combine the data objects by rows.
4. **Create a Data Frame**: After combining the data using either cbind() or rbind(), ensure the data is organized into a data frame.
5. **Display the Result**: Print or view the resulting data frame to verify the changes.

A.PROGRAM:

```
Name <- c("shubham Rastogi","nishka jain","Gunjan garg","sumit chaudhary")
Address <- c("Moradabad","etah","sambhal","Khurja")
Marks <- c(255,355,455,655)
info <- cbind(Name, Address, Marks)
print(info)
new.stuinfo <- data.frame(
  Name = c("Deepmala","Arun"),
  Address = c("khurja","moradabad"),
  Marks = c("755","855"),
  stringsAsFactors = FALSE
)
cat("### The Second data frame \n")
print(new.stuinfo)
all.info<- rbind(info,new.stuinfo)
```

```
cat("### The Combined data frame \n")
print(all.info)
```

OUTPUT:

	Name	Address	Marks
[1,]	"shubham Rastogi"	"Moradabad"	"255"
[2,]	"nishka jain"	"etah"	"355"
[3,]	"Gunjan garg"	"sambhal"	"455"
[4,]	"sumit chaudhary"	"Khurja"	"655"
### The Second data frame			
	Name	Address	Marks
1	Deepmala	khurja	755
2	Arun	moradabad	855
### The Combined data frame			
	Name	Address	Marks
1	shubham Rastogi	Moradabad	255
2	nishka jain	etah	355
3	Gunjan garg	sambhal	455
4	sumit chaudhary	Khurja	655
5	Deepmala	khurja	755
6	Arun	moradabad	855

B.PROGRAM:

```
df1<-data.frame(  
  Name=c("priya","mano","Ram"),  
  Age=c(35,26,30)  
)  
df2<-data.frame(  
  Gender=c("Female","male","male"),  
  occupation=c("programmer","doctor","engineer")  
)  
df_combined_columns<-cbind(df1,df2)  
print("Data frame after column-wise binding:")  
print(df_combined_columns)  
df3<-data.frame(  
  Name=c("Raja","kavitha"),  
  Age=c(40,45)  
)  
df4<-data.frame(  
  Gender=c("male","female"),  
  occupation=c("Lawyer","Scientist")  
)  
df_combined_rows<-rbind(df_combined_columns,cbind(df3,df4))  
print("Data frame after row-wise binding:")  
print(df_combined_rows)
```

OUTPUT:

[1] "Data frame after column-wise binding:"

	Name	Age	Gender	occupation
1	priya	35	Female	programmer
2	mano	26	male	doctor
3	Ram	30	male	engineer

[1] "Data frame after row-wise binding:"

	Name	Age	Gender	occupation
1	priya	35	Female	programmer
2	mano	26	male	doctor
3	Ram	30	male	engineer
4	Raja	40	male	Lawyer
5	kavitha	45	female	Scientist

6A1.Find the factorial of a number using recursion.

Aim:

The aim of this program is to find the factorial of a given number using recursion in R programming. The factorial of a number n (denoted as $n!$) is the product of all positive integers less than or equal to n . Recursion is a programming technique where a function calls itself to solve smaller instances of the same problem.

Factorial Definition:

- Factorial of 0: $0! = 10! = 10! = 1$
- Factorial of a positive integer n : $n! = n \times (n-1) \times (n-2) \times \dots \times 1$ $= n \times (n-1) \times (n-2) \times \dots \times 1$

For example:

- $5! = 5 \times 4 \times 3 \times 2 \times 1 = 120$
- $3! = 3 \times 2 \times 1 = 6$

Key Concept:

Recursion works by breaking down the problem into smaller subproblems. The recursive function for factorial can be defined as:

- Base Case: If $n=0$, the factorial is 1.
- Recursive Case: $n! = n \times (n-1) \times (n-2) \times \dots \times 1$

Algorithm:

1. Input: Read the number n for which the factorial needs to be calculated.
2. Base Case Check: If $n=0$, return 1 (because $0! = 10! = 1$).
3. Recursive Case: If $n > 0$, recursively call the factorial function for $n-1$ and multiply the result by n .
4. Output: Return the computed factorial of n .

A.PROGRAM:

```
recur_fact<-function(num){  
  if(num<=1){  
    return(1)  
  }else{  
    return(num*recur_fact(num-1))  
  }  
}  
  
print(paste("The factorial of 5 is",recur_fact(5)))  
  
findfactorial<-function(n){  
  factorial<-1  
  if((n==0)|(n==1))  
    factorial<-1  
  else{  
    for(i in 1:n)  
      factorial<-factorial*i  
  }  
  return(factorial)  
}  
  
print(paste(findfactorial(0)))  
print(paste(findfactorial(3)))  
  
num = as.integer(readline(prompt="Enter a number:"))  
  
factorial = 1  
  
if (num < 0){  
  print("Sorry; factorial does not exist for negative numbers")  
} else if (num == 0) {
```

```
print("The factorial of 0 is 1")
} else {
  for (i in 1:num) {
    factorial = factorial * i
  }
  print(paste("The factorial of", num, "is", factorial))
}
```

OUTPUT:

```
[1] "The factorial of 5 is 120"
[1] "1"
[1] "6"
Enter a number:8
"The factorial of 8 is 40320
```

B.PROGRAM:

```
factorial<-function(n){  
  if(n==0){  
    return(1)  
  }else{  
    return(n*factorial(n-1))  
  }  
}  
  
number<-7  
  
result<-factorial(number)  
  
print(paste("Factorial of",number,"is",result))
```

output:

```
[1] "Factorial of 7 is 5040"
```

6B . Mean,Variance,Standard deviation for the given probability distribution

Aim:

The aim of this program is to calculate the **mean**, **variance**, and **standard deviation** for a given probability distribution in **R programming**. **Mean (Expected Value)**: The average value of a random variable, weighted by its probabilities.

- **Variance**: Measures the spread of the values in the probability distribution, indicating how much the values deviate from the mean.
- **Standard Deviation**: The square root of the variance, which provides a measure of the dispersion of the distribution in the same units as the data.

Formulae:

1. Mean (Expected Value):

$$E(X) = \sum_{i=1}^n x_i \cdot p_i$$

Where x_i is the value, and p_i is the probability of x_i .

2. Variance:

$$\text{Var}(X) = \sum_{i=1}^n p_i \cdot (x_i - E(X))^2$$

Where x_i is the value, p_i is the probability, and $E(X)$ is the mean.

3. Standard Deviation:

$$\sigma(X) = \sqrt{\text{Var}(X)}$$

Algorithm:

1. Input:

- A set of values x_1, x_2, \dots, x_n representing the outcomes of a random variable.
- Corresponding probabilities p_1, p_2, \dots, p_n representing the probability of each outcome.

2. Calculate the Mean:

- Compute the mean (expected value) of the distribution using the formula:
$$E(X) = \sum_{i=1}^n x_i \cdot p_i$$

3. Calculate the Variance:

- Compute the variance using the formula: $\text{Var}(X) = \sum_{i=1}^n p_i \cdot (x_i - E(X))^2$

4. Calculate the Standard Deviation:

- Compute the standard deviation by taking the square root of the variance:
 $\sigma(X) = \sqrt{\text{Var}(X)}$

5. **Output:** Return the values for the mean, variance, and standard deviation.

A.PROGRAM:

```
x<-c(1,2,3,4,5,1,2,3,1,2,4,5,2,3,1,1,2,3,5,6)  
mean.result=mean(x)  
print(mean.result)  
median.result=median(x)  
print(median.result)  
variance.result=var(x)  
print(variance.result)  
sd.result=sqrt(var(x))  
print(sd.result)
```

output:

```
[1] 2.8  
[1] 2.5  
[1] 2.484211  
[1] 1.576138
```

B.PROGRAM:

```
outcomes <- c(1, 2, 3)
probabilities <- c(0.2, 0.5, 0.3)
mean_value <- sum(outcomes * probabilities)
cat("Mean (E(X)):", mean_value, "\n")
variance_value <- sum((outcomes - mean_value)^2 * probabilities)
cat("Variance (Var(X)):", variance_value, "\n")
std_dev_value <- sqrt(variance_value)
cat("Standard Deviation (SD):", std_dev_value, "\n")
```

output:

```
Mean (E(X)): 2.1
Variance (Var(X)): 0.49
Standard Deviation (SD): 0.7
```

7.Apriori Algorithm to extract Association rule of datamining

Aim:

To implement the Apriori algorithm in R programming to extract association rules from a given transactional dataset. The Apriori algorithm is a popular method in data mining that is used for discovering frequent itemsets and generating association rules from large datasets.

Algorithm:

1. Input:

- A transactional dataset with items bought by customers.
- Minimum support threshold and confidence threshold.

2. Generate Frequent Itemsets:

- Start with individual items and count their frequency.
- Find itemsets that meet the minimum support threshold.
- Use these frequent itemsets to generate candidate itemsets of length 2, then 3, and so on.
- Continue until no more frequent itemsets are found.

3. Generate Association Rules:

- For each frequent itemset, generate rules.
- Calculate the confidence of each rule.
- Keep the rules that meet the minimum confidence threshold.

4. Output:

- A list of association rules with support, confidence, and lift values.

A.PROGRAM:

```
library(arules)
library(arulesViz)
library(RColorBrewer)
data("Groceries")
```

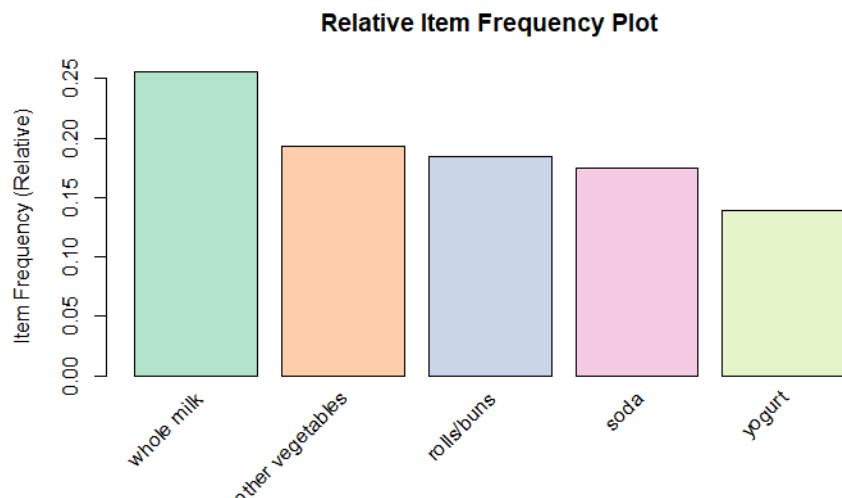
```

rules <- apriori(Groceries,
  parameter = list(supp = 0.01, conf = 0.2))
inspect(rules[1:10])
arules::itemFrequencyPlot(Groceries, topN = 5,
  col = brewer.pal(8, 'Pastel2'),
  main = 'Relative Item Frequency Plot',
  type = "relative",
  ylab = "Item Frequency (Relative)")

```

OUTPUT

	lhs	rhs	support	confidence	coverage	lift	count
[1]	{}	=> {whole milk}	0.255516	0.255516	0.1000000	1.0000	513
[2]	{hard cheese}	=> {whole milk}	0.010066	0.4107884	0.024504	1.60768	99
[3]	{butter milk}	=> {other vegetable}	0.01037	0.3709091	0.0279613	1.91691	102
[4]	{butter milk}	=> {whole milk}	0.011591	0.4145455	0.0279613	1.62238	114
[5]	{ham}	=> {whole milk}	0.011489	0.4414062	0.026029	1.7275	113
[6]	{sliced cheese}	=> {whole milk}	0.010777	0.4398340	0.0245043	1.7213	106
[7]	{oil}	=> {whole milk}	0.011286	0.402173	0.02806304	1.5739	111
[8]	{onions}	=> {other vegetable}	0.014234	0.459016	0.03101169	2.3722	140
[9]	{onions}	=> {whole milk}	0.012099	0.39016	0.03101169	1.5269	119
[10]	{berries}	=> {yogurt}	0.010574	0.31804	0.03324860	2.2798	104



B.PROGRAM:

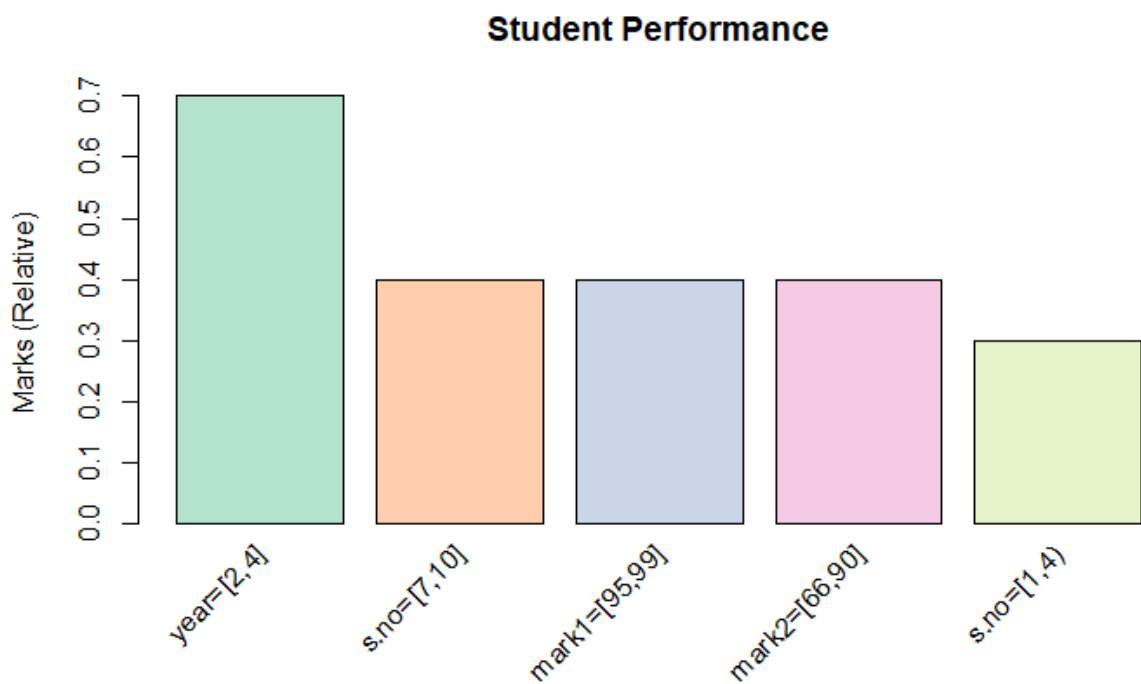
```
library(arules)
library(arulesViz)
library(RColorBrewer)
stud_details <- read.csv("D:/sample R/stud_details.csv")
stud_details_trans <- as(stud_details, "transactions")
rules <- apriori(stud_details_trans, parameter = list(supp = 0.01, conf = 0.2))
inspect(rules[1:10])
itemFrequencyPlot(stud_details_trans, topN = 5,
                  col = brewer.pal(8, 'Pastel2'),
                  main = 'Student Performance',
                  type = "relative",
                  ylab = "Marks (Relative)")
```

Stud_details:

s.no	s.name	mark1	mark2	dept	year
1	priya	60	56	mca	1
2	mano	67	66	bca	3
3	ram	68	67	mba	2
4	raja	58	78	bba	1
5	ravi	79	65	bcom	2
6	malathi	78	45	mcom	3
7	snegha	96	55	msc	2
8	ashwin	95	90	bsc	2
9	kavya	98	61	ba	1
10	siva	99	62	be	4

OUTPUT:

	lhs	rhs	support	confidence	coverage	lift	count
[1]	{ } => {s.no=[4,7)}		0.3	0.3	1	1	3
[2]	{ } => {mark1=[58,68)}		0.3	0.3	1	1	3
[3]	{ } => {mark2=[45,61)}		0.3	0.3	1	1	3
[4]	{ } => {year=[1,2)}		0.3	0.3	1	1	3
[5]	{ } => {mark2=[61,66) }		0.3	0.3	1	1	3
[6]	{ } => {mark1=[68,95)}		0.3	0.3	1	1	3
[7]	{ } => {s.no=[1,4)}		0.3	0.3	1	1	3
[8]	{ } => {mark1=[95,99]}		0.4	0.4	1	1	4
[9]	{ } => {mark2=[66,90]}		0.4	0.4	1	1	4
[10]	{ } => {s.no=[7,10]}		0.4	0.4	1	1	4



8.K-Means Clustering Technique

Aim:

To implement the K-Means clustering technique in R programming. K-Means is an unsupervised machine learning algorithm used to partition a dataset into K distinct, non-overlapping subsets (clusters).

Algorithm:

1. Input:

- A dataset with features (numerical values).
- The number of clusters, KKK.

2. Initialization:

- Select KKK initial centroids randomly from the data points.

3. Cluster Assignment:

- For each data point, calculate the distance to each of the KKK centroids.
- Assign the data point to the cluster with the nearest centroid.

4. Centroid Update:

- After all points are assigned, recalculate the centroids as the mean of the data points in each cluster.

5. Repeat:

- Repeat the cluster assignment and centroid update steps until the centroids no longer change or a maximum number of iterations is reached.

6. Output:

- Final clusters with their corresponding data points.
- The final centroids.

A.PROGRAM:

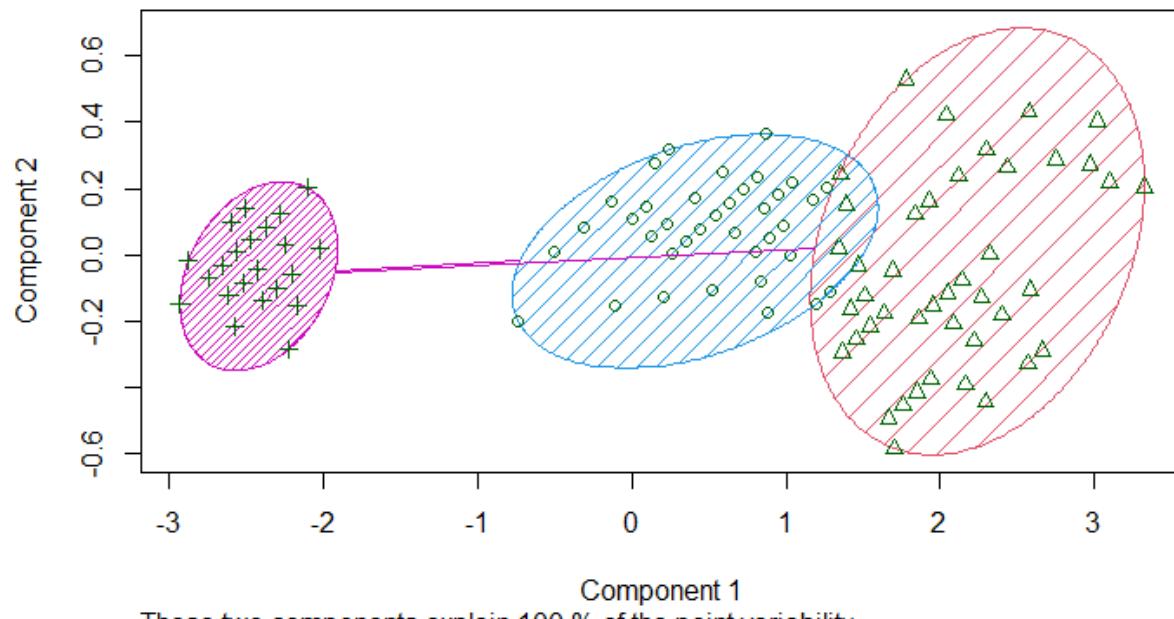
```
data("iris")
head(iris)
x=iris[,3:4]
head(x)
model=kmeans(x,3)
library(cluster)
clusplot(x,model$cluster)
clusplot(x,model$cluster,color=T,shade=T)
```

OUTPUT:

	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

	Petal.Length	Petal.Width
1	1.4	0.2
2	1.4	0.2
3	1.3	0.2
4	1.5	0.2
5	1.4	0.2
6	1.7	0.4

CLUSPLOT(x)



Component 1

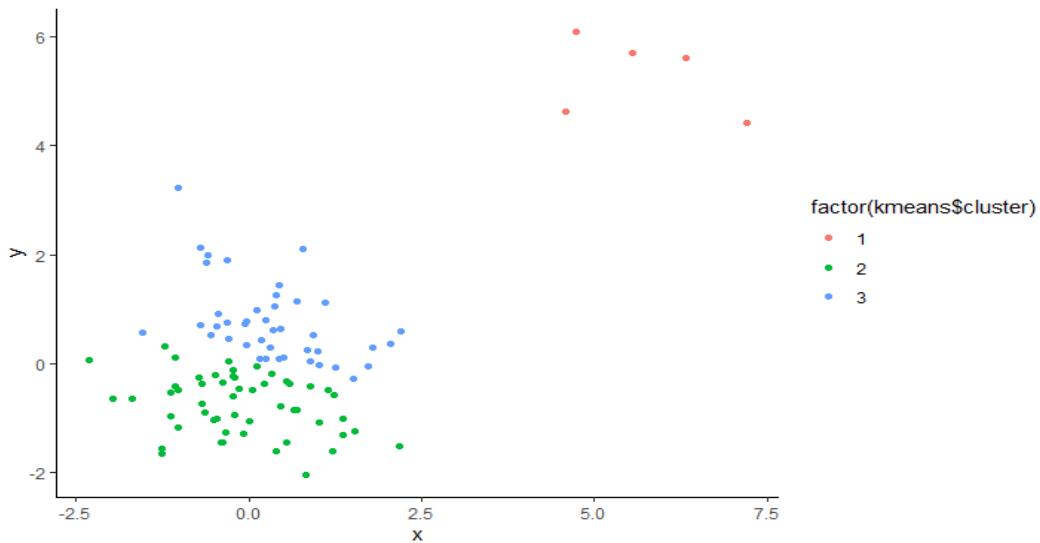
These two components explain 100 % of the point variability.

8B.PROGRAM:

```
library(ggplot2)
set.seed(123)
data<-data.frame(x=rnorm(100,mean=0,sd=1),
                  y=rnorm(100,mean=0,sd=1))
data$x[1:5]<-rnorm(20,mean=5,sd=1)
data$y[1:5]<-rnorm(20,mean=5,sd=1)
set.seed(123)
kmeans<-kmeans(data,centers=3)
print(kmeans$cluster)
ggplot(data,aes(x=x,y=y,color=factor(kmeans$cluster)))+geom_point()+theme_classic()
```

OUTPUT:

```
[1] 1 1 1 1 1 3 2 2 2 3 2 3 2 3 3 2 2 2 2 2 2 3 2 3 3 3 3 2 2 3 2 3 3 2 3 2 2 2 2 2 2  
[48] 3 3 2 3 3 3 2 2 3 3 2 3 2 2 3 2 3 3 2 2 3 3 2 2 2 2 3 2 2 2 3 3 3 2 3 2 3 2  
[95] 2 3 3 2 2 2
```



9.CLASSIFICATION ALGORITHM

Aim:

The aim of this program is to implement a Classification algorithm in R programming to predict the class labels of a dataset based on input features.

Algorithm:

1. **Input:** A dataset with features and class labels.
2. **Preprocessing:**
 - o Handle missing values.
 - o Normalize or scale features if required.
 - o Split the dataset into training and testing sets.
3. **Choose a Classification Algorithm:**
 - o In this case, we will use a **Decision Tree** algorithm as an example.
4. **Train the Model:** Use the training data to train the model.
5. **Test the Model:** Evaluate the model on the test data using performance metrics.
6. **Output:** The model's classification results (predictions) and performance metrics.

A.PROGRAM:

```
library(rpart)
library(rpart.plot)
library(caret)
data(iris)
set.seed(123)
train_index<-createDataPartition(iris$Species,p=0.8,list=FALSE)
train_data<-iris[train_index,]
test_data<-iris[-train_index,]
```

```

tree_model<-rpart(Species ~.,
                    data=train_data,
                    method="class",
                    control=rpart.control(minsplit=10,cp=0.01))

rpart.plot(tree_model,box.palette="auto",nn=TRUE)

predictions<-predict(tree_model,test_data,type="class")

confusionMatrix(predictions,test_data$Species)

```

OUTPUT:

Confusion Matrix and Statistics

		Reference		
Prediction	setosa	versicolor	virginica	
setosa	10	0	0	
versicolor	0	10	1	
virginica	0	0	9	

Overall Statistics

Accuracy : 0.9667
 95% CI : (0.8278, 0.9992)
 No Information Rate : 0.3333
 P-Value [Acc > NIR] : 2.963e-13

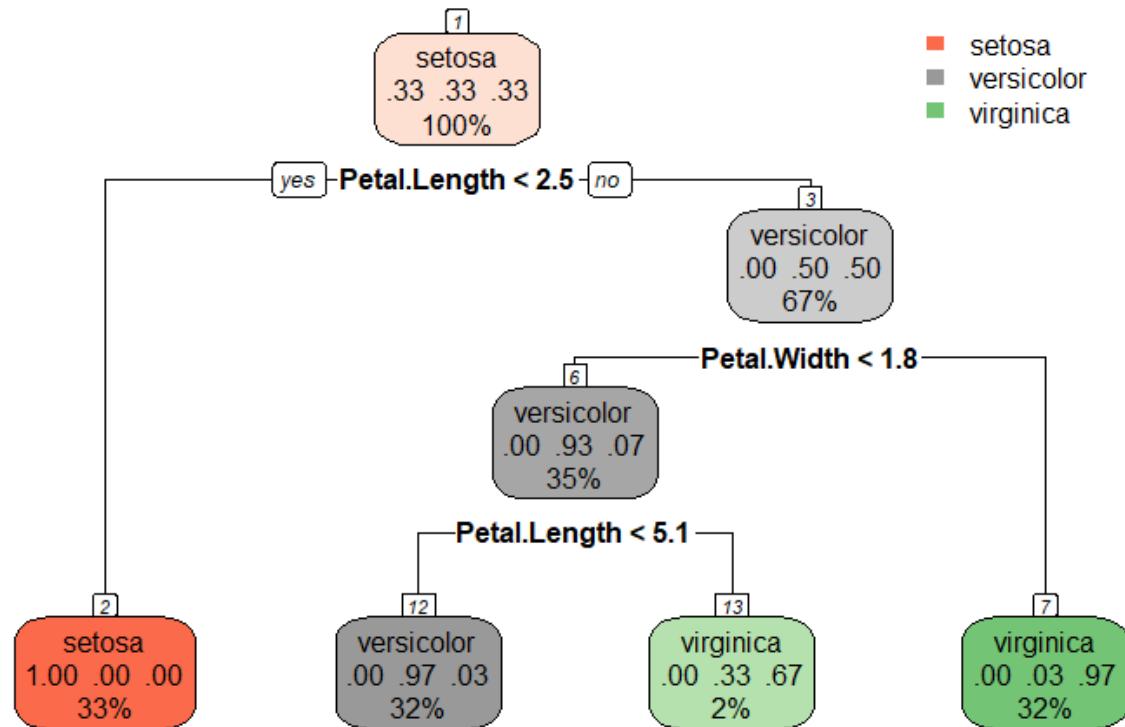
Kappa : 0.95

Mcnemar's Test P-Value : NA

Statistics by Class:

	Class: setosa	Class: versicolor	Class: virginica
Sensitivity	1.0000	1.0000	0.9000
Specificity	1.0000	0.9500	1.0000
Pos Pred Value	1.0000	0.9091	1.0000
Neg Pred Value	1.0000	1.0000	0.9524

Prevalence	0.3333	0.3333	0.3333
Detection Rate	0.3333	0.3333	0.3000
Detection Prevalence	0.3333	0.3667	0.3000
Balanced Accuracy	1.0000	0.9750	0.9500



9B.PROGRAM:

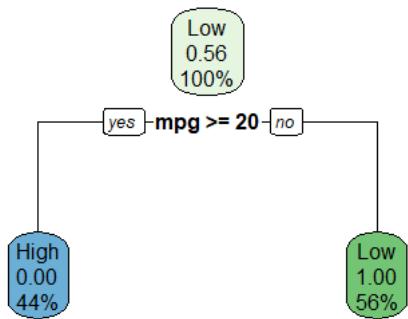
```
library(rpart)
library(rpart.plot)
library(caret)
data(mtcars)

mtcars$mpg_category <- ifelse(mtcars$mpg > 20, "High", "Low")
set.seed(123)

train_data <- mtcars[createDataPartition(mtcars$mpg_category, p=0.8, list=FALSE), ]
test_data <- mtcars[-createDataPartition(mtcars$mpg_category, p=0.8, list=FALSE), ]
tree_model <- rpart(mpg_category ~ ., data=train_data)
rpart.plot(tree_model)

predictions <- predict(tree_model, test_data, type="class")
confusionMatrix(predictions, test_data$mpg_category)
```

OUTPUT:



10.Pie Charts and Bar Charts Using R

Aim:

The aim of this program is to demonstrate how to create Pie Charts and Bar Charts in R programming for data visualization. These charts are used to represent categorical data in a way that is easy to interpret.

Algorithm:

1. Pie Chart Algorithm:

1. **Input:** A categorical dataset with counts or percentages for each category.
2. **Preprocessing:**
 - o Convert the data into a vector of counts or percentages.
3. **Generate Pie Chart:**
 - o Use the pie() function in R to create the pie chart.
 - o Specify the categories and corresponding values.
4. **Output:** A pie chart displaying the distribution of categories.

2. Bar Chart Algorithm:

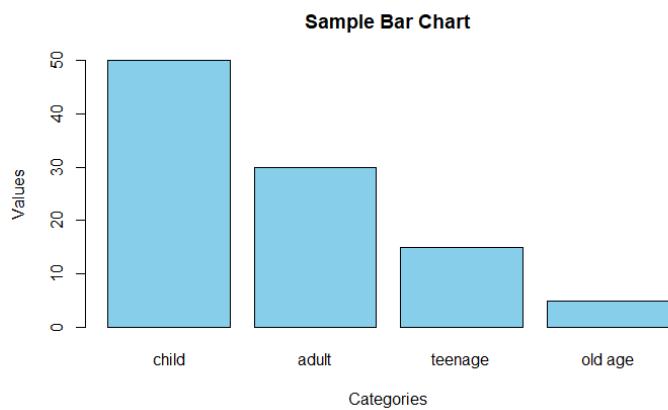
1. **Input:** A categorical dataset with counts or values for each category.
2. **Preprocessing:**
 - o Convert the data into a vector of counts or values.
3. **Generate Bar Chart:**
 - o Use the barplot() function in R to create the bar chart.
 - o Specify the categories and their corresponding values.
4. **Output:** A bar chart displaying the comparison of categories.

A.PROGRAM

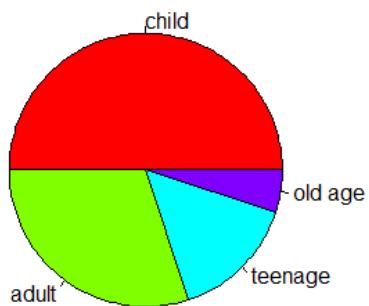
```
patient <- c(50, 30, 15, 5)
count <- c("child", "adult", "teenage", "old age")
pie(patient, labels = count, col = rainbow(length(patient)),
     main = "Pie Chart of patient")

patient <- c(50, 30, 15, 5)
count <- c("child", "adult", "teenage", "old age")
barplot(patient, names.arg = count, col = "skyblue",
        main = "Sample Bar Chart",
        xlab = "Categories", ylab = "Values")
```

OUTPUT:



Pie Chart of patient



B.PROGRAM

```
data <- read.csv("D:/DV/Book1.csv")
print(data)
barplot(data$value,
        names.arg = data$category,
        col = "blue",
        main = "Bar Chart",
        xlab = "Category",
        ylab = "Value",
        border = "white")
percentages <- round(data$value / sum(data$Value) * 100, 1)
pie(data$value,
    labels = paste(data$category, "(", percentages, "%)", sep = ""),
    col = rainbow(length(data$value)),
    main = "Pie Chart")
```

