

Program No.: 1

Aim: Introduction to R in Data Science.

Theory:

R is a powerful and versatile open-source programming language primarily designed for statistical computing, data analysis, and visualization. It was created in the mid-1990s by Ross Ihaka and Robert Gentleman at the University of Auckland, New Zealand. Since then, R has evolved into one of the most widely used tools for researchers, data analysts, and statisticians worldwide.

Key Characteristics of R:

1. Statistical Computing:

R provides a rich environment for statistical techniques such as hypothesis testing, regression, clustering, classification, and time-series analysis.

2. Data Manipulation:

With built-in data structures like vectors, matrices, data frames, and lists, R allows efficient storage, transformation, and analysis of data. Packages like dplyr and tidyr further enhance these capabilities.

3. Data Visualization:

R has powerful tools for data visualization. Packages such as ggplot2 and lattice can generate simple charts to complex, multi-layered graphics that aid in understanding trends and patterns in data.

4. Extensibility:

R is open-source and supports thousands of user-contributed packages, which extend its functionality for specialized tasks like machine learning (caret), deep learning (keras), geospatial analysis (sf), and bioinformatics (Bioconductor).

5. Cross-Platform & Integration:

R can run on Windows, macOS, and Linux. It integrates with databases (SQL, MongoDB), other languages (Python, Java, C++), and big data tools (Hadoop, Spark).

6. Community & Support:

R has a large global community of developers and researchers. The CRAN (Comprehensive R Archive Network) hosts over 20,000 packages, ensuring support for a wide range of analytical tasks.

Applications of R in Data Science:

- **Data Cleaning & Preprocessing:** Handling missing values, filtering, and transforming data.
- **Statistical Analysis:** Performing tests, building models, and making predictions.
- **Visualization & Reporting:** Creating interactive dashboards and reports.
- **Machine Learning:** Implementing classification, regression, clustering, and neural networks.
- **Research & Academia:** R is widely used in scientific research for reproducible results.

In summary, R is not just a programming language but a complete environment for data-driven decision-making. Its combination of statistical power, visualization capabilities, and package ecosystem makes it a preferred choice in Data Science.

Program No.: 2

Aim: To study the installation of R and explore its basic syntax.

Theory:

R can be installed from CRAN (Comprehensive R Archive Network). After installation, RStudio (an IDE for R) can be used for writing and running R programs.

Some **basic syntax elements of R** are:

1. **Variables and Assignment:** Values can be assigned using <- or =.
2. **Data Types:** R supports numeric, character, logical, complex, and raw data types.
3. **Vectors:** A one-dimensional collection of data elements, created using c().
4. **Operators:** Arithmetic (+, -, *, /), relational (<, >, ==), and logical (&, |).
5. **Functions:** Built-in and user-defined functions are used to perform operations.

Source Code:

```
# Assigning values  
  
x <- 10  
  
y = 20  
  
print(x)  
  
print(y)  
  
# Creating a vector  
  
vec <- c(1, 2, 3, 4, 5)  
  
print(vec)  
  
# Arithmetic operations  
  
sum <- x + y
```

```
product <- x * y  
print(sum)  
print(product)  
  
# Logical comparison  
  
result <- x < y  
print(result)  
  
# Built-in function  
  
sqrt_val <- sqrt(25)  
print(sqrt_val)
```

Output:

```
[1] 10  
[1] 20  
[1] 1 2 3 4 5  
[1] 30  
[1] 200  
[1] TRUE  
[1] 5
```

Program No.: 3

Aim: To describe data, and perform viewing and manipulation operations using R.

Theory:

R provides powerful tools for data description and manipulation. Data can be imported, explored, and transformed efficiently.

- **Describing Data:** Functions like `summary()`, `str()`, and `dim()` provide insights into datasets.
- **Viewing Data:** The `head()`, `tail()`, and `View()` functions allow quick inspection.
- **Manipulating Data:** Using indexing, filtering, and packages like `dplyr` for operations such as `select()`, `filter()`, `mutate()`, `arrange()`.

Source Code:

```
# Creating a sample data frame
```

```
data <- data.frame(
```

```
  ID = 1:5,
```

```
  Name = c("A", "B", "C", "D", "E"),
```

```
  Age = c(21, 25, 19, 23, 22),
```

```
  Score = c(85, 90, 78, 88, 95)
```

```
)
```

```
# Describing the data
```

```
print("Summary of Data:")
```

```
print(summary(data))
```

```
print("Structure of Data:")
```

```

print(str(data))

# Viewing data

print("First few rows:")

print(head(data, 3))

print("Last few rows:")

print(tail(data, 2))

# Manipulating data

print("Selecting Age column:")

print(data$Age)

print("Filtering rows where Score > 85:")

print(subset(data, Score > 85))

```

Output:

```

[1] "Summary of Data:"
      ID      Name       Age      Score
Min.   :1  Length:5      Min.   :19  Min.   :78.0
1st Qu.:2  Class :character  1st Qu.:21  1st Qu.:85.0
Median  :3  Mode   :character  Median :22  Median :88.0
Mean    :3                      Mean   :22  Mean   :87.2
3rd Qu.:4                      3rd Qu.:23  3rd Qu.:90.0
Max.    :5                      Max.   :25  Max.   :95.0

[1] "Structure of Data:"
'data.frame': 5 obs. of 4 variables:
 $ ID   : int 1 2 3 4 5
 $ Name : chr "A" "B" "C" "D" ...

```

```
$ Age  : num  21 25 19 23 22
$ Score: num  85 90 78 88 95
NULL
[1] "First few rows:"
  ID Name Age Score
1  1    A   21    85
2  2    B   25    90
3  3    C   19    78
[1] "Last few rows:"
  ID Name Age Score
4  4    D   23    88
5  5    E   22    95
```

```
[1] "Selecting Age column:"
[1] 21 25 19 23 22
[1] "Filtering rows where Score > 85:"
  ID Name Age Score
2  2    B   25    90
4  4    D   23    88
5  5    E   22    95
```

Program No.: 4

Aim: To visualize data using tables, charts, and plots in R.

Theory:

R is widely known for its strong visualization capabilities. Some methods include:

- **Tables:** Using `table()` for frequency distributions.
- **Charts and Plots:** Functions like `barplot()`, `hist()`, `pie()`, and `plot()`.
- **Advanced Visualizations:** Using `ggplot2` for layered and customizable graphics.

Source Code:

```
marks <- c(85, 90, 78, 88, 95, 70, 80, 92)

# Frequency table
print("Frequency Table:")
print(table(marks))

# Histogram
hist(marks, col="lightblue", main="Marks Distribution", xlab="Marks")

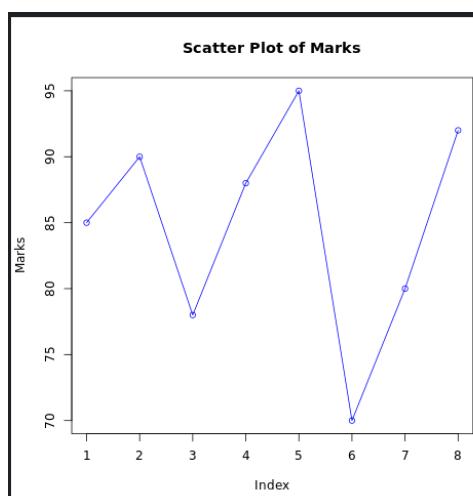
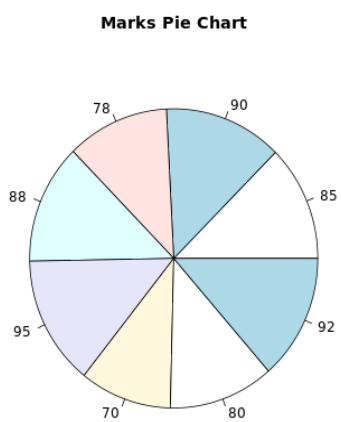
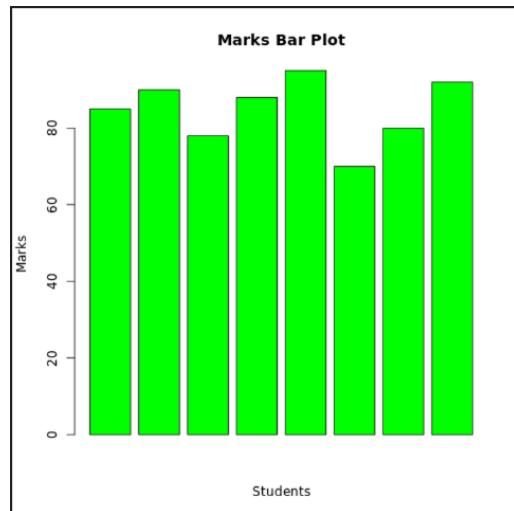
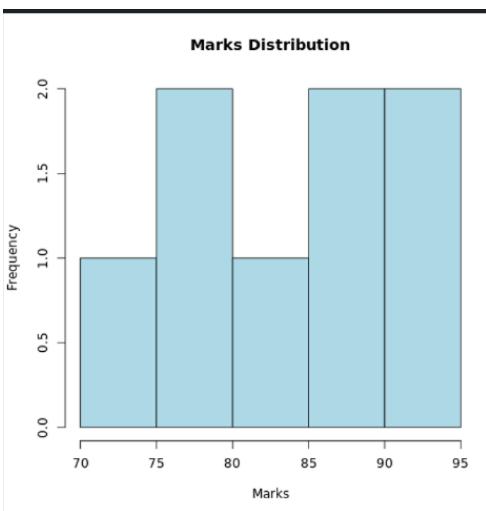
# Barplot
barplot(marks, col="green", main="Marks Bar Plot", xlab="Students", ylab="Marks")

# Pie chart
pie(marks, labels = marks, main="Marks Pie Chart")

# Scatter plot
plot(marks, type="o", col="blue", main="Scatter Plot of Marks", xlab="Index", ylab="Marks")
```

Output:

```
[1] "Frequency Table:"  
marks  
70 78 80 85 88 90 92 95  
1 1 1 1 1 1 1 1
```



Program No.: 5

Aim: To implement binomial distribution and plot its density and distribution functions in R.

Theory:

The **Binomial distribution** is a discrete probability distribution representing the number of successes in a fixed number of independent trials, each with the same probability of success.

- Probability Mass Function: $P(X = k) = \binom{n}{k} p^k (1 - p)^{n-k}$
- In R:
 - `dbinom(k, n, p)` → Probability at k .
 - `pbinom(k, n, p)` → Cumulative probability up to k .
 - `rbinom(n, size, prob)` → Generates random binomial values.

Source Code:

```
# Parameters  
  
n <- 10 # number of trials  
  
p <- 0.5 # probability of success  
  
# Probability distribution  
  
x <- 0:n  
  
prob <- dbinom(x, n, p)  
  
# Print probabilities  
  
print("Binomial Probabilities:")  
  
print(prob)  
  
# Plotting probability mass function
```

```

barplot(prob, names.arg=x, col="lightblue",
        main="Binomial Distribution (n=10, p=0.5)",
        xlab="Number of Successes", ylab="Probability")

# Cumulative distribution

cum_prob <- pbinom(x, n, p)

print("Cumulative Probabilities:")

print(cum_prob)

# Plot cumulative distribution

plot(x, cum_prob, type="o", col="red",
      main="Cumulative Distribution Function",
      xlab="Number of Successes", ylab="Cumulative Probability")

```

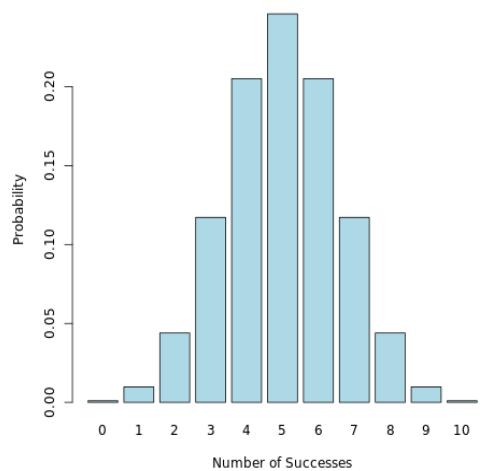
Output:

```

[1] "Binomial Probabilities:"
[1] 0.0009765625 0.0097656250 0.0439453125 0.1171875000 0.2050781250
[6] 0.2460937500 0.2050781250 0.1171875000 0.0439453125 0.0097656250
[11] 0.0009765625
[1] "Cumulative Probabilities:"
[1] 0.0009765625 0.0107421875 0.0546875000 0.1718750000 0.3769531250
[6] 0.6230468750 0.8281250000 0.9453125000 0.9892578125 0.9990234375
[11] 1.0000000000

```

Binomial Distribution (n=10, p=0.5)



Cumulative Distribution Function

