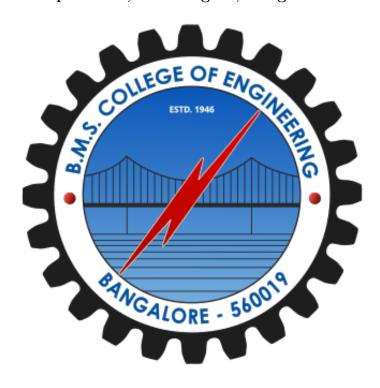


B.M.S. COLLEGE OF ENGINEERING

 ${\rm (Autonomous\ College\ under\ VTU)}$ Bull Temple Road, Basavangudi, Bangalore - 560019



Lab Observation

ON

Programming with R

Submitted by

R V Abhishek(1BM23CD047)

 $in\ fulfillment\ of\ mandatory\ observation\ submission\ for\ Lab\ assessment$

BACHELOR OF ENGINEERING

in

Computer Science & Engineering (Data Science)

Under the Guidance of

Dr. Kalyan N

Assistant Professor

 $\ \, \textbf{Department of CSE (Data Science)},$

B.M.S. College of Engineering

2025-2026

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Laboratory Certificate

This is to certify that Mr./Ms. $\underline{\mathbf{R}\ \mathbf{V}\ \mathbf{Abhishek}}$ has satisfactorily completed the course of experiments in practical $\underline{\mathbf{Programming}\ \mathbf{With}\ \mathbf{R}}$ prescribed by the Visvesvaraya Technology University for 5^{th} Semester Bachelor of Engineering course in the laboratory of the college in the year 2024 - 2025

Head of the Department

Staff Incharge of the Batch

Marks		Name of the Candidate:	R V Abhishek		
Maximum	Obtained	Branch:	CSE (Data Science)		
		USN:	1BM23CD047		

Date:

Signature of the Candidate

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Program to check what type of Triangle given 3 sides, and calculate its area

Date of Execution - 2025-08-13

This R program validates the sides of the triangle (taken as input from the user) and then if valid, calculates the area of the triangle using Heron's formula and checks what type of triangle it is

```
# Validating the triangle
is_valid_triangle <- function(a, b, c) {
  return ((a + b > c) & (b + c > a) & (a + c > b))
}
```

```
# Function to check the type of triangle
triangle_type <- function(a , b , c) {
   if (a == b && b == c) {
      return(" Equilateral ")
   } else if (a == b || b == c || a == c) {
      return(" Isosceles ")
   } else {
      return("Scalene")
   }
}</pre>
```

```
# Calculating Area using Heron's Formula
triangle_area <- function(a , b , c) {
   s <- (a + b + c) / 2 # Semi - perimeter
   # Heron 's formula
   area <- sqrt (s * (s - a) * (s - b) * (s - c))
   return (area)
}</pre>
```

```
# Validating inputs
validate_input <- function(x) {
   if (!is.numeric(x) || x <= 0) {
      stop("Error : Input must be a positive number.")
   }
   return(TRUE)
}</pre>
```

```
## Main Code Block
# 1. Defining 3 variables representing the 3 sides of the triangle
cat("Enter the lengths of the sides of the triangle :\n")
Enter the lengths of the sides of the triangle :
a <- as.numeric(readline(prompt = "Side A: "))</pre>
Side A:
b <- as.numeric(readline(prompt = "Side B: "))</pre>
Side B:
c <- as.numeric(readline(prompt = "Side C: "))</pre>
Side C:
# 2. Input Validation and implementation of all the functions.
# Input validation}
tryCatch ({
  validate_input(a)
 validate_input(b)
  validate_input(c)
  # Check if the inputs form a valid triangle
  if (!is_valid_triangle(a , b , c)) {
    stop("Error : The given sides do not form a valid triangle.")
  }
  # Determine the type of triangle
  type_of_triangle <- triangle_type(a , b , c)</pre>
  cat("The triangle is:", type_of_triangle, "\n")
  # Calculate the area of the triangle
  area_of_triangle <- triangle_area(a, b, c)</pre>
  cat("The area of the triangle is:", area_of_triangle, "\n")
}, error = function(e){
  cat(e$message, "\n")
})
```

missing value where TRUE/FALSE needed

Sample Output

```
Enter the lengths of the sides of the triangle:
Side a: 5
Side b: 5
Side c: 8

The triangle is: Isosceles
The area of the triangle is: 12

Enter the lengths of the sides of the triangle:**
Side a: 1
Side b: 2
Side c: 8

Error: The given sides do not form a valid triangle.
```

Creating and Manipulating Data Structures

Date of Execution - 2025-08-20

Is 50 present in the vector? FALSE

Objective - This program evaluates the student's understanding of different data structures (vectors, matrices, lists, and data frames) in R and how to manipulate them.

```
# 1. Create a vector of random numbers and apply operations such as sorting and searching
set.seed(42) # For reproducibility
random_vector <- runif(20, min = 1, max = 100)</pre>
cat("Original random vector:\n")
Original random vector:
print(random_vector)
 [1] 91.56580 93.77047 29.32781 83.21431 64.53281 52.39050 73.92224
 [8] 14.33199 66.04224 70.80141 46.31644 72.19211 93.53255 26.28745
[15] 46.76699 94.06144 97.84442 12.63125 48.02471 56.47294
# Sort the vector
sorted_vector <- sort(random_vector)</pre>
cat("Sorted vector:\n")
Sorted vector:
print(sorted_vector)
 [1] 12.63125 14.33199 26.28745 29.32781 46.31644 46.76699 48.02471
 [8] 52.39050 56.47294 64.53281 66.04224 70.80141 72.19211 73.92224
[15] 83.21431 91.56580 93.53255 93.77047 94.06144 97.84442
# Search for a specific value (check if a number is present)
search_value <- 50
is_value_present <- any(random_vector == search_value)</pre>
cat("Is", search_value, "present in the vector?", is_value_present, "\n")
```

```
# Find values in the vector greater than 60
values_greater_than_60 <- random_vector[random_vector > 60]
cat("Values greater than 60:\n")
Values greater than 60:
print(values_greater_than_60)
 [1] 91.56580 93.77047 83.21431 64.53281 73.92224 66.04224 70.80141
 [8] 72.19211 93.53255 94.06144 97.84442
# 2. Convert the vector into a matrix and perform matrix multiplication
# Create a 4x5 matrix from the vector
matrix_from_vector <- matrix(random_vector, nrow = 4, ncol = 5)</pre>
cat("Matrix from vector:\n")
Matrix from vector:
print(matrix_from_vector)
         [,1]
                  [,2]
                            [,3]
                                     [,4]
                                              [,5]
[1,] 91.56580 64.53281 66.04224 93.53255 97.84442
[2,] 93.77047 52.39050 70.80141 26.28745 12.63125
[3,] 29.32781 73.92224 46.31644 46.76699 48.02471
[4,] 83.21431 14.33199 72.19211 94.06144 56.47294
# Perform matrix multiplication (matrix with its transpose)
matrix_transpose <- t(matrix_from_vector)</pre>
matrix_multiplication_result <- matrix_from_vector %*% matrix_transpose</pre>
cat("Matrix multiplication result:\n")
Matrix multiplication result:
print(matrix_multiplication_result)
         [,1]
                  [,2]
                            [,3]
                                     [,4]
[1,] 35232.22 20337.59 19587.86 27635.57
[2,] 20337.59 17401.08 11738.17 16851.17
[3,] 19587.86 11738.17 12963.36 13954.70
[4,] 27635.57 16851.17 13954.70 24378.48
```

```
# Element-wise matrix multiplication (Hadamard product)
elementwise_multiplication_result <- matrix_from_vector * matrix_from_vector
cat("Element-wise matrix multiplication result:\n")
Element-wise matrix multiplication result:
print(elementwise_multiplication_result)
          [,1]
                   [,2] [,3]
                                     [,4]
                                               [,5]
[1,] 8384.2954 4164.483 4361.577 8748.3384 9573.5298
[2,] 8792.9003 2744.764 5012.840 691.0302 159.5484
[3,] 860.1207 5464.498 2145.212 2187.1513 2306.3729
[4,] 6924.6222 205.406 5211.701 8847.5541 3189.1932
# 3. Create a list containing different types of elements and perform subsetting
my_list <- list(</pre>
 numbers = random_vector,
 characters = c("A", "B", "C", "D"),
 logical_values = c(TRUE, FALSE, TRUE),
 matrix = matrix_from_vector
)
cat("List:\n")
List:
print(my_list)
$numbers
 [1] 91.56580 93.77047 29.32781 83.21431 64.53281 52.39050 73.92224
 [8] 14.33199 66.04224 70.80141 46.31644 72.19211 93.53255 26.28745
[15] 46.76699 94.06144 97.84442 12.63125 48.02471 56.47294
$characters
[1] "A" "B" "C" "D"
$logical_values
[1] TRUE FALSE TRUE
$matrix
         [,1] [,2] [,3] [,4]
                                            [,5]
```

```
[1,] 91.56580 64.53281 66.04224 93.53255 97.84442
[2,] 93.77047 52.39050 70.80141 26.28745 12.63125
[3,] 29.32781 73.92224 46.31644 46.76699 48.02471
[4,] 83.21431 14.33199 72.19211 94.06144 56.47294
# Subsetting the list (extracting numeric and logical parts)
subset_numeric <- my_list$numbers</pre>
cat("Subset (numeric part of the list):\n")
Subset (numeric part of the list):
str(subset_numeric)
 num [1:20] 91.6 93.8 29.3 83.2 64.5 ...
cat("\n")
subset_logical <- my_list$logical_values</pre>
cat("Subset (logical part of the list):\n", subset_logical, "\n")
Subset (logical part of the list):
 TRUE FALSE TRUE
# Modify elements in the list (replace the second character with "Z")
my_list$characters[2] <- "Z"</pre>
cat("Modified list of characters:\n", my_list$characters, "\n")
Modified list of characters:
 AZCD
# Apply a function to the numeric part of the list
# (e.g., calculate the square of the numbers)
squared_numbers <- my_list$numbers ^ 2</pre>
cat("Squared numbers:\n")
Squared numbers:
str(squared_numbers)
 num [1:20] 8384 8793 860 6925 4164 ...
```

```
cat("\n")
# 4. Create a data frame and perform operations such as filtering,
# summarizing, and handling missing values
\# Create a data frame
df <- data.frame(</pre>
 ID = 1:20.
 Age = sample(18:65, 20, replace = TRUE),
 Score = runif(20, min = 50, max = 100),
 Passed = sample(c(TRUE, FALSE), 20, replace = TRUE)
cat("Data frame:\n")
Data frame:
print(df)
   ID Age
            Score Passed
   1 64 71.78858 FALSE
1
  2 20 51.87155 FALSE
2
3
  3 58 98.67700 FALSE
4
  4 42 71.58756 FALSE
  5 44 97.87883 FALSE
5
  6 53 94.38775 FALSE
6
  7 54 81.99894
7
                   TRUE
8
   8 48 98.54833 FALSE
   9 62 80.94191
                   TRUE
9
10 10 22 66.67136 FALSE
11 11 37 67.33741
                   TRUE
12 12 51 69.92427 FALSE
13 13 45 89.23464 FALSE
14 14 57 51.94682 FALSE
15 15 20 87.43977 FALSE
16 16 50 83.86384
                    TRUE
17 17 59 58.56322
                    TRUE
18 18 41 63.05440
                    TRUE
19 19 47 75.72065
                    TRUE
```

20 20 60 83.78036 FALSE

```
\# Filter the data frame (rows where Age > 30 and Score > 70)
filtered_df <- subset(df, Age > 30 & Score > 70)
cat("Filtered data frame (Age > 30 and Score > 70):\n")
Filtered data frame (Age > 30 and Score > 70):
print(filtered_df)
   ID Age
             Score Passed
   1 64 71.78858 FALSE
1
   3 58 98.67700 FALSE
3
4
  4 42 71.58756 FALSE
5 5 44 97.87883 FALSE
  6 53 94.38775 FALSE
  7 54 81.99894
7
                    TRUE
  8 48 98.54833 FALSE
8
   9 62 80.94191
                    TRUE
13 13 45 89.23464 FALSE
16 16 50 83.86384
                    TRUE
19 19 47 75.72065
                    TRUE
20 20 60 83.78036 FALSE
# Calculate mean, sum, and variance of numerical columns (Age and Score)
mean_age <- mean(df$Age)</pre>
sum_age <- sum(df$Age)</pre>
var_age <- var(df$Age)</pre>
mean_score <- mean(df$Score)</pre>
sum_score <- sum(df$Score)</pre>
var_score <- var(df$Score)</pre>
cat("Summary statistics for Age column:\n")
Summary statistics for Age column:
cat("Mean Age:", mean_age, "\n")
```

Mean Age: 46.7

```
cat("Sum of Age:", sum_age, "\n")
Sum of Age: 934
cat("Variance of Age:", var_age, "\n")
Variance of Age: 179.6947
cat("Summary statistics for Score column:\n")
Summary statistics for Score column:
cat("Mean Score:", mean_score, "\n")
Mean Score: 77.26086
cat("Sum of Score:", sum_score, "\n")
Sum of Score: 1545.217
cat("Variance of Score:", var_score, "\n")
Variance of Score: 219.2162
# 5. Handling missing values in the data frame
# Introduce some NA values in the Score column
df$Score[sample(1:20, 5)] <- NA</pre>
cat("Data frame with missing values:\n")
Data frame with missing values:
print(df)
  ID Age
            Score Passed
  1 64 71.78858 FALSE
   2 20 51.87155 FALSE
3 3 58 98.67700 FALSE
  4 42
               NA FALSE
5
   5 44 97.87883 FALSE
  6 53 94.38775 FALSE
6
```

```
7
   7 54 81.99894
                   TRUE
   8 48 98.54833 FALSE
   9 62
                   TRUE
               NA
10 10 22
               NA FALSE
11 11 37 67.33741
                   TRUE
12 12 51
               NA FALSE
13 13 45
               NA
                  FALSE
14 14 57 51.94682
                  FALSE
15 15 20 87.43977
                  FALSE
16 16 50 83.86384
                   TRUE
17 17 59 58.56322
                   TRUE
18 18 41 63.05440
                   TRUE
19 19 47 75.72065
                   TRUE
20 20 60 83.78036 FALSE
```

```
# Replace NA values with the mean of the Score column

df$Score[is.na(df$Score)] <- mean(df$Score, na.rm = TRUE)

cat("Data frame after imputation of missing values:\n")</pre>
```

Data frame after imputation of missing values:

print(df)

```
ID Age
            Score Passed
   1 64 71.78858 FALSE
1
2
  2 20 51.87155 FALSE
3
   3 58 98.67700 FALSE
   4 42 77.79050 FALSE
4
   5 44 97.87883 FALSE
5
6
   6 53 94.38775 FALSE
7
   7 54 81.99894
                   TRUE
   8 48 98.54833 FALSE
8
   9 62 77.79050
9
                   TRUE
10 10 22 77.79050 FALSE
      37 67.33741
                   TRUE
12 12 51 77.79050 FALSE
13 13 45 77.79050 FALSE
14 14 57 51.94682 FALSE
15 15 20 87.43977
                  FALSE
16 16 50 83.86384
                   TRUE
17 17 59 58.56322
                   TRUE
```

```
18 18 41 63.05440
                    TRUE
19 19 47 75.72065
                    TRUE
20 20 60 83.78036 FALSE
# Grouping the data by Passed status and calculating group-wise statistics
library(dplyr)
Attaching package: 'dplyr'
The following objects are masked from 'package:stats':
    filter, lag
The following objects are masked from 'package:base':
    intersect, setdiff, setequal, union
grouped_stats <- df %>%
  group_by(Passed) %>%
  summarise(
   mean_score = mean(Score, na.rm = TRUE),
   mean_age = mean(Age)
cat("Grouped statistics by Passed status:\n")
Grouped statistics by Passed status:
print(grouped_stats)
# A tibble: 2 x 3
  Passed mean_score mean_age
  <1g1>
              <dbl>
                       <dbl>
1 FALSE
              80.6
                        44.9
2 TRUE
              72.6
                        50
```

Basic Statistical Operations on Open-Source Datasets

Date of Execution - 2025-08-26

Objective: This program emphasizes the application of statistical concepts on real-world datasets and visualization of the data.

```
# Load necessary
library(dplyr) # For data manipulation
library(ggplot2) # For visualization
library(moments) # For skewness and kurtosis
library(palmerpenguins) # For Palmer Penguins dataset
data(iris) # Load Iris dataset
data(penguins) # Load Palmer Penguins
# Function to calculate mode
calc_mode <- function(x) {</pre>
  return (as.numeric (names (sort (table (x), decreasing = TRUE)) [1] ))
# Perform Statistical Analysis on Iris Dataset
print("---- Iris Dataset Analysis ----")
[1] "---- Iris Dataset Analysis ----"
# Mean
iris_mean <- sapply (iris[, 1:4], mean, na.rm = TRUE )</pre>
print(paste("Mean of Iris dataset : ", iris_mean))
[1] "Mean of Iris dataset: 5.843333333333333"
[2] "Mean of Iris dataset : 3.05733333333333"
[3] "Mean of Iris dataset: 3.758"
[4] "Mean of Iris dataset: 1.199333333333333"
#Median
iris_median <- sapply(iris[, 1:4], median, na.rm = TRUE )</pre>
print(paste("Median of Iris dataset : ", iris_median))
```

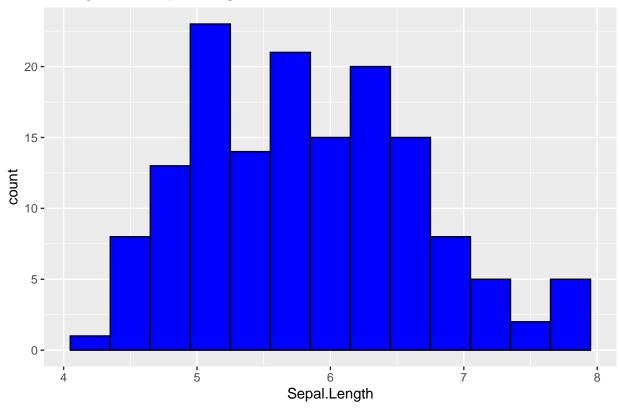
```
[1] "Median of Iris dataset: 5.8" "Median of Iris dataset: 3"
[3] "Median of Iris dataset: 4.35" "Median of Iris dataset: 1.3"
iris_mode <- sapply(iris[, 1:4], calc_mode )</pre>
print(paste("Mode of Iris dataset : ", iris_median))
[1] "Mode of Iris dataset : 5.8" "Mode of Iris dataset : 3"
[3] "Mode of Iris dataset : 4.35" "Mode of Iris dataset : 1.3"
#Variance
iris_variance <- sapply(iris[, 1:4], var, na.rm = TRUE )</pre>
print(paste("Variance of Iris dataset : ", iris_variance))
[1] "Variance of Iris dataset: 0.685693512304251"
[2] "Variance of Iris dataset : 0.189979418344519"
[3] "Variance of Iris dataset: 3.11627785234899"
[4] "Variance of Iris dataset: 0.581006263982103"
#Standard Deviation
iris_sd <- sapply(iris[, 1:4], sd, na.rm = TRUE )</pre>
print(paste("Standard Deviation of Iris dataset : ", iris_sd))
[1] "Standard Deviation of Iris dataset: 0.828066127977863"
[2] "Standard Deviation of Iris dataset : 0.435866284936698"
[3] "Standard Deviation of Iris dataset: 1.76529823325947"
[4] "Standard Deviation of Iris dataset: 0.762237668960347"
#Skewness
iris_skewness <- sapply(iris[, 1:4], skewness, na.rm = TRUE )</pre>
print(paste("Skewness of Iris dataset : ", iris_skewness))
[1] "Skewness of Iris dataset: 0.311753058502296"
[2] "Skewness of Iris dataset : 0.315767106338938"
[3] "Skewness of Iris dataset: -0.272127666456721"
[4] "Skewness of Iris dataset: -0.101934206565599"
# Hypothesis Testing (t-test) between Sepal.Length of Setosa and Versicolor
setosa <- subset(iris, Species == "setosa")$Sepal.Length</pre>
versicolor <- subset(iris, Species == "versicolor")$Sepal.Length</pre>
t_test <- t.test(setosa, versicolor)</pre>
print(t_test)
```

```
Welch Two Sample t-test
```

```
data: setosa and versicolor
t = -10.521, df = 86.538, p-value < 2.2e-16
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
   -1.1057074 -0.7542926
sample estimates:
mean of x mean of y
   5.006   5.936</pre>
```

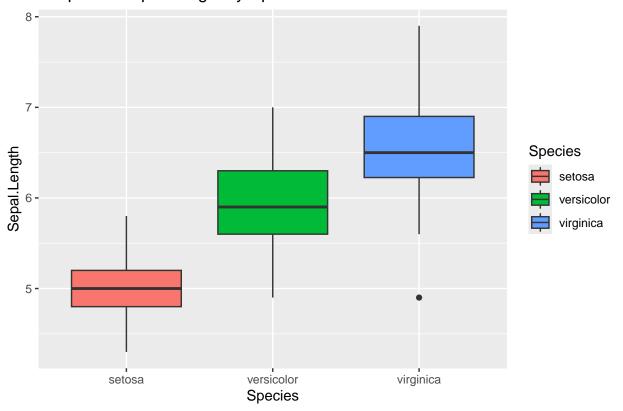
```
# Visualization of Iris Dataset
# Histogram for Sepal.Length
ggplot(iris, aes(x = Sepal.Length)) +
  geom_histogram(binwidth = 0.3, fill = "blue", color = "black") +
  ggtitle("Histogram of Sepal Length in Iris Dataset")
```

Histogram of Sepal Length in Iris Dataset



```
# Boxplot for Sepal.Length across Species
ggplot(iris, aes(x = Species, y = Sepal.Length, fill = Species)) +
  geom_boxplot() +
  ggtitle("Boxplot of Sepal Length by Species in Iris Dataset")
```

Boxplot of Sepal Length by Species in Iris Dataset



```
print("---- Palmer Penguins Dataset Analysis ----")
```

[1] "---- Palmer Penguins Dataset Analysis ----"

```
# Remove rows with missing values
penguins_clean <- na.omit(penguins)

# Mean
penguins_mean <- sapply(penguins_clean[, 3:6], mean, na.rm = TRUE)
print(paste("Mean of Palmer Penguins dataset:", penguins_mean))</pre>
```

- [1] "Mean of Palmer Penguins dataset: 43.9927927927928"
- [2] "Mean of Palmer Penguins dataset: 17.1648648648649"
- [3] "Mean of Palmer Penguins dataset: 200.966966966967"
- [4] "Mean of Palmer Penguins dataset: 4207.05705705706"

Median

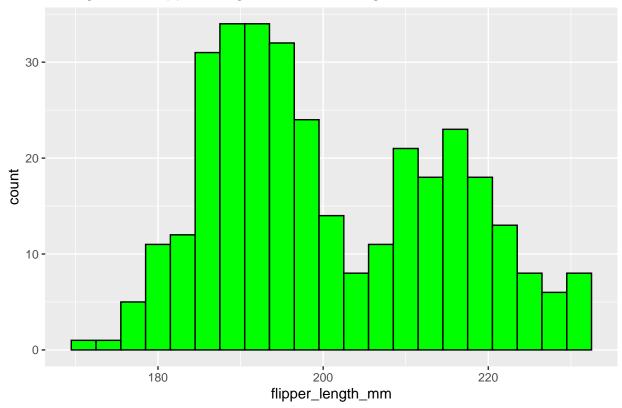
```
penguins_median <- sapply(penguins_clean[, 3:6], median, na.rm = TRUE)
print(paste("Median of Palmer Penguins dataset:", penguins_median))</pre>
```

- [1] "Median of Palmer Penguins dataset: 44.5"
- [2] "Median of Palmer Penguins dataset: 17.3"

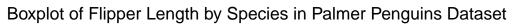
```
[3] "Median of Palmer Penguins dataset: 197"
[4] "Median of Palmer Penguins dataset: 4050"
# Mode
penguins_mode <- sapply(penguins_clean[, 3:6], calc_mode)</pre>
print(paste("Mode of Palmer Penguins dataset:", penguins_mode))
[1] "Mode of Palmer Penguins dataset: 41.1"
[2] "Mode of Palmer Penguins dataset: 17"
[3] "Mode of Palmer Penguins dataset: 190"
[4] "Mode of Palmer Penguins dataset: 3800"
# Variance
penguins_variance <- sapply(penguins_clean[, 3:6], var, na.rm = TRUE)</pre>
print(paste("Variance of Palmer Penguins dataset:", penguins_variance))
[1] "Variance of Palmer Penguins dataset: 29.9063334418756"
[2] "Variance of Palmer Penguins dataset: 3.87788830999674"
[3] "Variance of Palmer Penguins dataset: 196.441676616375"
[4] "Variance of Palmer Penguins dataset: 648372.487698542"
# Standard Deviation
penguins_sd <- sapply(penguins_clean[, 3:6], sd, na.rm = TRUE)</pre>
print(paste("Standard Deviation of Palmer Penguins dataset:", penguins_sd))
[1] "Standard Deviation of Palmer Penguins dataset: 5.46866834264756"
[2] "Standard Deviation of Palmer Penguins dataset: 1.9692354633199"
[3] "Standard Deviation of Palmer Penguins dataset: 14.0157652882879"
[4] "Standard Deviation of Palmer Penguins dataset: 805.215801942897"
# Skewness
penguins_skewness <- sapply(penguins_clean[, 3:6], skewness, na.rm = TRUE)</pre>
print(paste("Skewness of Palmer Penguins dataset:", penguins_skewness))
[1] "Skewness of Palmer Penguins dataset: 0.0451359779776739"
[2] "Skewness of Palmer Penguins dataset: -0.149044996398334"
[3] "Skewness of Palmer Penguins dataset: 0.358523654622741"
[4] "Skewness of Palmer Penguins dataset: 0.470116171418382"
# Kurtosis
penguins_kurtosis <- sapply(penguins_clean[, 3:6], kurtosis, na.rm = TRUE)</pre>
print(paste("Kurtosis of Palmer Penguins dataset:", penguins_kurtosis))
```

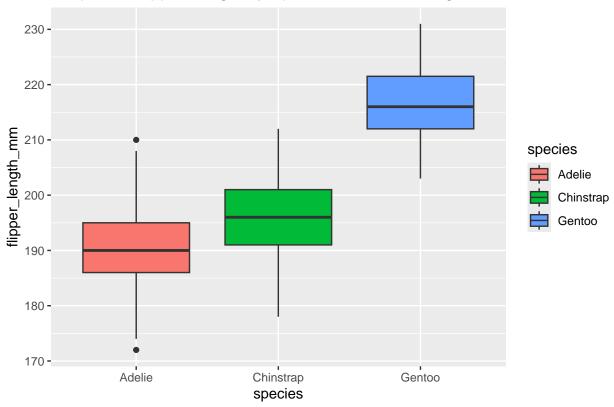
```
[1] "Kurtosis of Palmer Penguins dataset: 2.11182658541194"
[2] "Kurtosis of Palmer Penguins dataset: 2.10341274887238"
[3] "Kurtosis of Palmer Penguins dataset: 2.03516741259049"
[4] "Kurtosis of Palmer Penguins dataset: 2.25951411974012"
# Hypothesis Testing (t-test) between flipper_length_mm of Adelie and Gentoo species
adelie <- subset(penguins_clean, species == "Adelie")$flipper_length_mm</pre>
gentoo <- subset(penguins_clean, species == "Gentoo")$flipper_length_mm</pre>
t_test_penguins <- t.test(adelie, gentoo)</pre>
print(t_test_penguins)
    Welch Two Sample t-test
data: adelie and gentoo
t = -33.506, df = 251.35, p-value < 2.2e-16
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
-28.72740 -25.53771
sample estimates:
mean of x mean of y
 190.1027 217.2353
# Visualization of Palmer Penguins Dataset
# Histogram for flipper_length_mm
ggplot(penguins_clean, aes(x = flipper_length_mm)) +
  geom_histogram(binwidth = 3, fill = "green", color = "black") +
 ggtitle("Histogram of Flipper Length in Palmer Penguins Dataset")
```

Histogram of Flipper Length in Palmer Penguins Dataset



```
# Boxplot for flipper_length_mm across Species
ggplot(penguins_clean, aes(x = species, y = flipper_length_mm, fill = species)) +
   geom_boxplot() +
   ggtitle("Boxplot of Flipper Length by Species in Palmer Penguins Dataset")
```





Data Import, Cleaning, and Export with Titanic Dataset and Adult Income Dataset

Date of Execution - 2025-09-09

Objective: Real World Data Cleaning Processes and emphasis on usage of advanced data wrangling techniques in R.

```
# Load necessary libraries
library(tidyverse)
library(titanic)
library(dplyr)
library(caret)
library(ggcorrplot)
# Load the Titanic dataset
data <- titanic::titanic_train</pre>
# Handle the missing data
# Replace missing values in the 'Age' column with the median age
data$Age[is.na(data$Age)] <- median(data$Age, na.rm = TRUE)</pre>
# Replace missing values in the 'Embarked' column with the mode
mode_embarked <- as.character(names(sort(table(data$Embarked), decreasing = TRUE)[1]))</pre>
data$Embarked[is.na(data$Embarked)] <- mode_embarked</pre>
\# Define the numeric columns for z-score and correlation calculation
numeric_columns <- c("Age", "SibSp", "Parch", "Fare", "Survived", "Pclass")</pre>
# Remove outliers using z-score
z_scores <- as.data.frame(scale(data[, numeric_columns]))</pre>
# Identify the rows that have z_scores greater than 3 or less than -3 (outliers)
outlier_rows <- apply(z_scores, 1, function(row) any(abs(row) > 3))
# Filter out Outliers
data_cleaned <- data[!outlier_rows, ]</pre>
```

```
# Summarize the dataset before and after cleaning
summary_before <- summary(data)
summary_after <- summary(data_cleaned)

# Calculate Correlation Matrix (fixed)
correlation_matrix <- cor(data_cleaned[, numeric_columns], use = "complete.obs")

# Export cleaned data onto a new CSV file
write.csv(data_cleaned, "titanic_cleaned.csv", row.names = FALSE)

# Display Summaries
print("Summary Before Cleaning:")</pre>
```

[1] "Summary Before Cleaning:"

print(summary_before)

Passe	engerId	Surv	ived	Pcla	SS	Name	
Min.	: 1.0	Min.	:0.0000	Min. :	1.000	Length:8	91
1st Qu	1.:223.5	1st Qu.	:0.0000	1st Qu.:	2.000	Class :c	haracter
Mediar	n :446.0	Median	:0.0000	Median :	3.000	Mode :c	haracter
Mean	:446.0	Mean	:0.3838	Mean :	2.309		
3rd Qu	1.:668.5	3rd Qu.	:1.0000	3rd Qu.:	3.000		
Max.	:891.0	Max.	:1.0000	Max. :	3.000		
Se	ex		Age	Si	bSp	Pa	rch
Length	n:891	Min.	: 0.42	Min.	:0.000	Min.	:0.0000
Class	:characte	r 1st	Qu.:22.00	1st Qu	.:0.000	1st Qu	.:0.0000
Mode	:characte	r Medi	an :28.00	Median	:0.000	Median	:0.0000
		Mean	:29.36	Mean	:0.523	Mean	:0.3816
		3rd	Qu.:35.00	3rd Qu	.:1.000	3rd Qu	.:0.0000
		Max.	:80.00	Max.	:8.000	Max.	:6.0000
Tio	cket		Fare	Ca	bin		
Length	n:891	Min.	: 0.0	0 Lengt	h:891		
Class	:characte	r 1st	Qu.: 7.9	1 Class	:chara	cter	
Mode	:characte	r Medi	an : 14.4	5 Mode	:chara	cter	
		Mean	: 32.2	.0			
		3rd	Qu.: 31.0	0			
		Max.	:512.3	3			
Emba	arked						

Length:891

Class :character

Mode :character

print("Summary After Cleaning:")

[1] "Summary After Cleaning:"

print(summary_after)

PassengerId		Surv	ived	Pclass		Name		
Min.	: 1.0	Min.	:0.0000	Min. :1	.000 Le	ength:820		
1st Qu	1.:226.8	1st Qu.	:0.0000	1st Qu.:2	.000 C1	ass :character		
Mediar	:446.5	Median	:0.0000	Median :3	.000 Mc	de :character		
Mean	:445.7	Mean	:0.3902	Mean :2	.311			
3rd Qu	1.:661.2	3rd Qu.	:1.0000	3rd Qu.:3	.000			
Max.	:891.0	Max.	:1.0000	Max. :3	.000			
Se	ex		Age	Sib	Sp	Parch		
Length	1:820	Min.	: 0.42	Min.	:0.0000	Min. :0.0000		
Class	:characte	r 1st	Qu.:23.00	1st Qu.	:0.0000	1st Qu.:0.0000		
Mode	:characte	r Medi	an :28.00	Median	:0.0000	Median :0.0000		
		Mean	:29.44	Mean	:0.3488	Mean :0.2549		
		3rd	Qu.:35.00	3rd Qu.	:1.0000	3rd Qu.:0.0000		
		Max.	:66.00	Max.	:3.0000	Max. :2.0000		
Tic	cket		Fare	Ca	bin			
Length	1:820	Min.	: 0.00	00 Lengt	h:820			
Class	:characte	r 1st	Qu.: 7.89	96 Class	:charact	er		
Mode	:characte	r Medi	an : 13.00	00 Mode	:charact	er		
		Mean	: 25.83	36				
		3rd	Qu.: 27.00	00				
		Max.	:164.86	67				

 ${\tt Embarked}$

Length:820

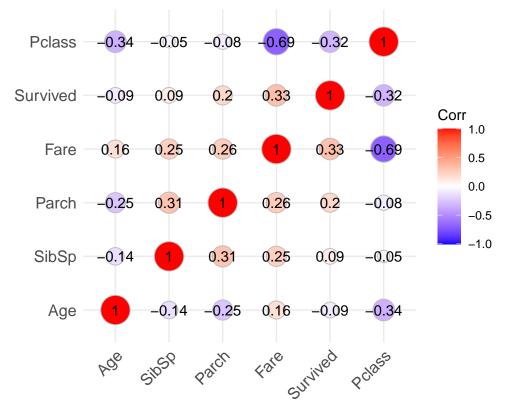
Class :character
Mode :character

```
print("Correlation Matrix:")
[1] "Correlation Matrix:"
print(correlation_matrix)
                          SibSp
                                     Parch
                                                 Fare
                                                         Survived
                Age
        1.00000000 -0.14391182 -0.2517719 0.1598100 -0.08602643
Age
        -0.14391182 1.00000000 0.3072105 0.2472157 0.09445934
SibSp
Parch
        -0.25177192  0.30721046  1.0000000  0.2599031  0.20107069
Fare
        0.15981001 0.24721568 0.2599031 1.0000000 0.33043946
Survived -0.08602643 0.09445934 0.2010707 0.3304395 1.00000000
Pclass
        -0.33698055 -0.05231213 -0.0783660 -0.6917198 -0.32230582
             Pclass
        -0.33698055
Age
        -0.05231213
SibSp
Parch
        -0.07836600
Fare
        -0.69171982
Survived -0.32230582
Pclass
         1.00000000
# Visualize Correlation Matrix (fixed)
ggcorrplot(correlation_matrix,
          method = "circle",
```

lab = TRUE) +

ggtitle("Correlation Matrix of Titanic Dataset")

Correlation Matrix of Titanic Dataset



Objective - Data Import, Cleaning, and Export with Adult Income Dataset

```
# Load necessary libraries
library(tidyverse)
library(dplyr)
library(caret)
library(ggcorrplot)
# Load the Adult Income dataset
data <- read.csv("D:/Coding/Coding/Time Series Analysis/Lab 4/adult.data", header = FALSE)</pre>
# Assign column names based on the dataset documentation
colnames(data) <- c('age', 'workclass', 'fnlwgt', 'education', 'education_num',</pre>
                     'marital_status', 'occupation', 'relationship', 'race', 'sex', 'capital_gain',
                     'capital_loss', 'hours_per_week', 'native_country', 'income')
# Handle missing values represented by '?'
data[data == '?'] <- NA</pre>
# Replace categorical missing values with mode
replace_mode <- function(x){</pre>
  mode_val <- as.character(names(sort(table(x), decreasing = TRUE)[1]))</pre>
```

```
x[is.na(x)] <- mode_val
  return(x)
}
data <- data %>%
 mutate_if(is.character, replace_mode)
# Replace numeric missing values with median
data <- data %>%
  mutate_if(is.numeric, ~ifelse(is.na(.), median(., na.rm = TRUE), .))
\# Define the remove_outliers function
remove_outliers <- function(x){</pre>
  z_scores <- scale(x)</pre>
 x[abs(z\_scores) \le 3]
# Remove outliers using z-score
numeric_columns <- sapply(data, is.numeric)</pre>
# Apply z-score outlier removal to numeric columns
data_cleaned <- data %>%
  filter(!apply(as.data.frame(scale(data[, numeric_columns])), 1,
                function(row) any(abs(row) > 3)))
# Summarize before and after cleaning
summary_before <- summary(data)</pre>
summary_after <- summary(data_cleaned)</pre>
# Calculate correlation Matrix
correlation_matrix <- cor(data_cleaned[, numeric_columns], use = "complete.obs")</pre>
# Export as CSV
write.csv(data_cleaned, "cleaned_adult_income_data.csv", row.names = FALSE)
# Display Summaries
print("Summary Before Cleaning:")
```

[1] "Summary Before Cleaning:"

print(summary_before)

age	workclass	fnlwgt
Min. :17.00	Length:32561	Min. : 12285
1st Qu.:28.00	Class :character	1st Qu.: 117827
Median :37.00	Mode :character	Median : 178356
Mean :38.58		Mean : 189778
3rd Qu.:48.00		3rd Qu.: 237051
Max. :90.00		Max. :1484705
education	education_num	marital_status
Length: 32561	Min. : 1.00	Length: 32561
Class :characte	r 1st Qu.: 9.00	Class :character
Mode :characte	r Median :10.00	Mode :character
	Mean :10.08	
	3rd Qu.:12.00	
	Max. :16.00	
occupation	relationship	race
Length: 32561	Length: 32561	Length: 32561
Class :characte	r Class :characte	r Class :character
Mode :characte	r Mode :characte	r Mode :character

sex	capital_gain	capital_loss	hours_per_week	
Length: 32561	Min. : 0	Min. : 0.0	Min. : 1.00	
Class :character	1st Qu.: 0	1st Qu.: 0.0	1st Qu.:40.00	
Mode :character	Median: 0	Median: 0.0	Median :40.00	
	Mean : 1078	Mean : 87.3	Mean :40.44	
	3rd Qu.: 0	3rd Qu.: 0.0	3rd Qu.:45.00	
	Max. :99999	Max. :4356.0	Max. :99.00	

native_country income

Length:32561 Length:32561

Class:character Class:character

Mode:character Mode:character

print("Summary After Cleaning:")

[1] "Summary After Cleaning:"

print(summary_after)

age	workclas	S	fnlwgt				
Min. :17.00	Length:29	828 M:	in. : 12285				
1st Qu.:27.00	Class :ch	aracter 1	st Qu.:117509				
Median :37.00	Mode :ch	aracter Me	edian :177667				
Mean :38.14	1	Me	ean :185193				
3rd Qu.:47.00)	31	rd Qu.:234279				
Max. :79.00)	Ма	ax. :506329				
education	educat	ion_num ma	arital_status				
Length:29828	Min.	: 3.00 Le	ength:29828				
Class :charac	cter 1st Qu	.: 9.00 C	lass :character				
Mode :charac	cter Median	:10.00 Mo	ode :character				
	Mean	:10.08					
	3rd Qu	.:12.00					
	Max.	:16.00					
occupation	relati	onship	race				
Length:29828	Length	:29828	Length:29828				
Class :charac	cter Class	:character	Class :character				
Mode :charac	cter Mode	:character	Mode :character				

sex		capital_gain					capital_loss				
Length:29828		Min.		:	0.	0	Min		:	0.000	
Class	:charact	er	1st	Qu.	:	0.	0	1st	Qu.	:	0.000
Mode	:charact	er	Medi	ian	:	0.	0	Med	ian	:	0.000
			Mear	1	: 5	70.	2	Mea	n	:	1.209
			3rd	Qu.	:	0.	0	3rd	Qu.	. :	0.000
			Max.		:220	40.	0	Max		:12	58.000
hours_per_week nat			cive_country				income				
Min.	: 4.0	Len	gth:2	29828	3		Lei	ngth	:298	328	
1st Qu	1.:40.0	Cla	ss :	chara	acte	r	Cla	ass	:cha	arac	ter
Mediar	ı:40.0	Mod	e :	chara	acte	r	Мо	de	:cha	arac	ter
Mean	:39.9										
3rd Qu	1.:45.0										
Max.	:77.0										

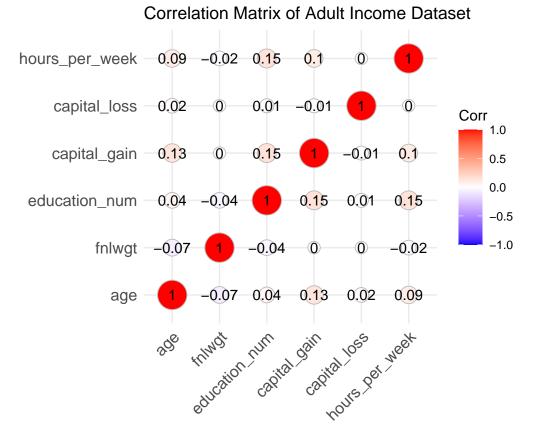
print("Correlation Matrix:")

[1] "Correlation Matrix:"

print(correlation_matrix)

```
fnlwgt education_num capital_gain
                       age
age
                1.00000000 -0.074427786
                                          0.041427102 0.131043981
               -0.07442779   1.000000000   -0.037482414   -0.002378925
fnlwgt
                                         1.000000000 0.154844283
               0.04142710 -0.037482414
education_num
capital_gain
               0.13104398 -0.002378925
                                          0.154844283 1.000000000
                0.02082465 0.002583047
                                         0.009481359 -0.009038231
capital_loss
hours_per_week 0.09219535 -0.015375555
                                         0.150513483 0.097209049
               capital_loss hours_per_week
                0.020824647
                               0.092195352
age
fnlwgt
                0.002583047
                              -0.015375555
education_num
               0.009481359
                             0.150513483
capital_gain
              -0.009038231
                             0.097209049
capital_loss
                1.000000000
                             -0.003089539
hours_per_week -0.003089539
                               1.000000000
```


ggtitle("Correlation Matrix of Adult Income Dataset")



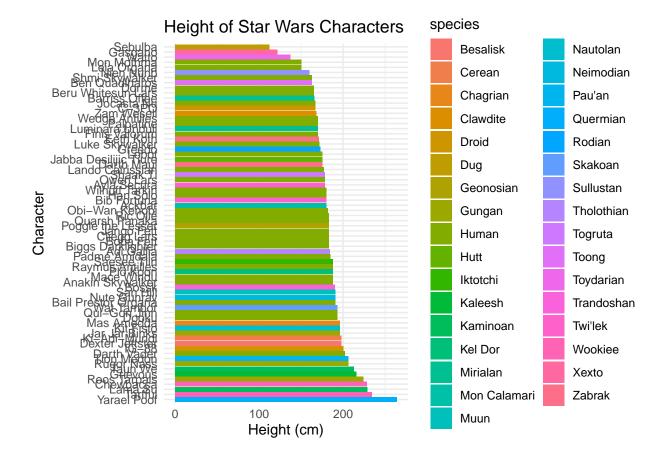
Advanced Data Manipulation with dplyr and Complex Grouping

Date of Execution - 2025-09-16

Objective - The goal of this program is to test advanced data manipulation techniques using the dplyr package.

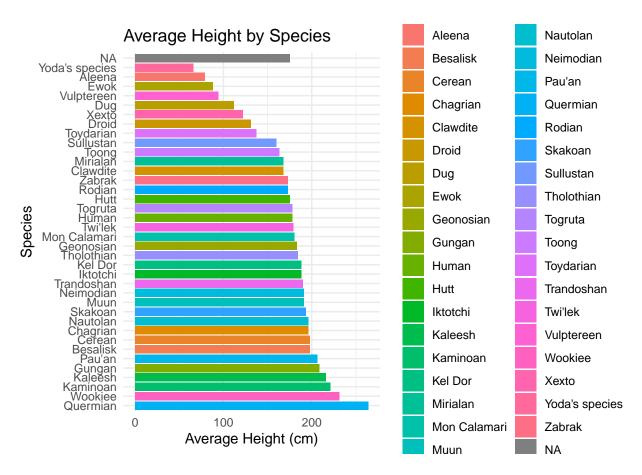
```
## Load necessary libraries
library(dplyr)
library(nycflights13)
library(ggplot2)
library(zoo)
# Preview the Star Wars Dataset
data("starwars")
head(starwars)
# A tibble: 6 x 14
        height mass hair_color skin_color eye_color birth_year sex
 name
  <chr>
         <int> <dbl> <chr>
                                 <chr>
                                            <chr>
                                                           <dbl> <chr>
1 Luke ~
            172
                   77 blond
                                 fair
                                            blue
                                                                 male
2 C-3PO
           167
                  75 <NA>
                                 gold
                                                           112
                                            yellow
                                                                 none
3 R2-D2
           96
                32 <NA>
                                 white, bl~ red
                                                            33
                                                                 none
4 Darth~
            202
                                                            41.9 male
                136 none
                                 white
                                            yellow
5 Leia ~
           150
                 49 brown
                                 light
                                            brown
                                                            19
                                                                 fema~
            178
                 120 brown, gr~ light
                                            blue
                                                                 male
# i 6 more variables: gender <chr>, homeworld <chr>, species <chr>,
    films <list>, vehicles <list>, starships <list>
# Select specific columns (name, species, height, mass),
# filter out rows with missing species or height,
# and arrange by height in descending order
starwars_filtered <- starwars %>%
  select(name, species, height, mass) %>%
  filter(!is.na(species) & !is.na(height) & height > 100) %>%
  arrange(desc(height))
# Display the filtered data
head(starwars_filtered)
```

```
# A tibble: 6 x 4
  name
               species height mass
  <chr>
               <chr>
                          <int> <dbl>
1 Yarael Poof Quermian
                            264
                                   NA
2 Tarfful
               Wookiee
                            234
                                  136
3 Lama Su
               Kaminoan
                            229
                                   88
4 Chewbacca
               Wookiee
                            228
                                  112
                            224
                                   82
5 Roos Tarpals Gungan
6 Grievous
               Kaleesh
                            216
                                  159
```



```
# Grouping by species, calculating average height and mass, and counting observation
species_summary <- starwars %>%
group_by(species) %>%
```

```
summarise(
    avg_height = mean(height, na.rm = TRUE),
  avg_mass = mean(mass, na.rm = TRUE),
   count = n()
  ) %>%
  arrange(desc(count))
# Display the species summary
head(species_summary)
# A tibble: 6 x 4
 species avg_height avg_mass count
  <chr>
                <dbl>
                       <dbl> <int>
1 Human
                178
                          81.3
                                 35
2 Droid
                          69.8
                131.
                                   6
3 <NA>
                 175
                          81
                                   4
4 Gungan
                 209.
                          74
                                   3
5 Kaminoan
                 221
                          88
                                   2
6 Mirialan
                 168
                          53.1
                                   2
# Plotting the average height
ggplot(species_summary, aes(x = reorder(species, -avg_height), y = avg_height, fill = species)) +
  geom_bar(stat = "identity") +
 coord_flip() +
  labs(title = "Average Height by Species",
      x = "Species",
       y = "Average Height (cm)") +
 theme_minimal()
```



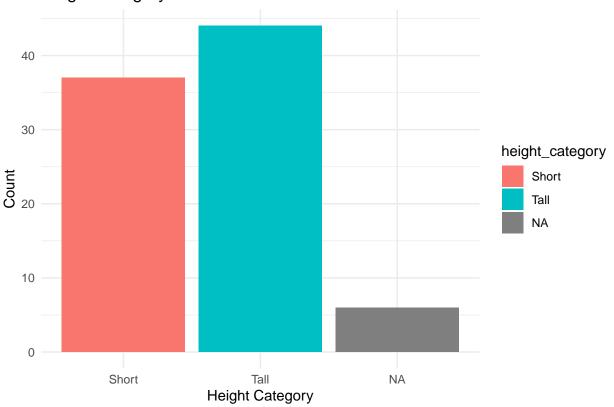
```
# Adding a new column that classifies characters based on height
starwars_classified <- starwars %>%
   mutate(height_category = ifelse(height < 180, "Short", "Tall"))
# Display the classified data
head(starwars_classified)</pre>
```

A tibble: 6 x 15

name height mass hair_color skin_color eye_color birth_year sex <chr> <int> <dbl> <chr> <chr> <chr> <dbl> <chr> 172 77 blond 1 Luke ~ fair blue 19 male 2 C-3PO 75 <NA> gold 167 yellow 112 none 3 R2-D2 32 <NA> white, bl~ red 96 33 none 4 Darth~ 202 136 none white yellow 41.9 male 5 Leia ~ 150 49 brown light brown 19 fema~ 6 Owen ~ 178 120 brown, gr~ light blue 52 male

- # i 7 more variables: gender <chr>, homeworld <chr>, species <chr>,
- # films <list>, vehicles <list>, starships <list>,
- # height_category <chr>

Height Category Distribution



```
# Joining with another dataset (flights dataset from nycflights13)
data("flights")
data("airlines")

# Inner join flights with airlines on the common column "carrier"
flights_inner_join <- flights %>%
    inner_join(airlines, by = "carrier")

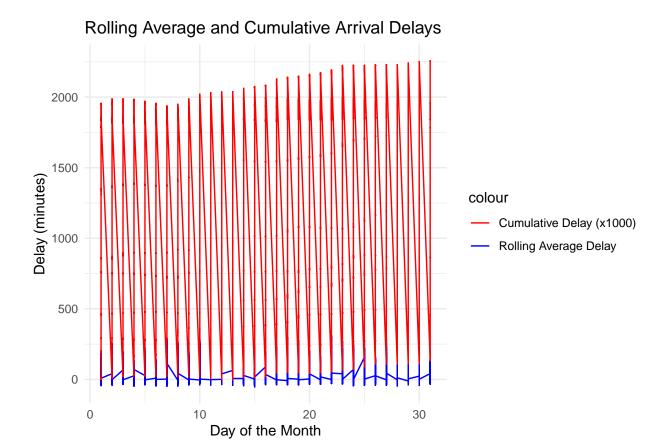
# Outer join flights with airlines on the common column "carrier"
flights_outer_join <- flights %>%
    full_join(airlines, by = "carrier")

# Display the joined data
head(flights_inner_join)
```

```
# A tibble: 6 x 20
  year month
              day dep_time sched_dep_time dep_delay arr_time
  <int> <int> <int>
                       <int>
                                       <int>
                                                 <dbl>
                                                          <int>
1 2013
                                                     2
                  1
                         517
                                         515
                                                            830
2 2013
                                         529
                  1
                         533
                                                     4
                                                            850
 2013
                         542
                                         540
                                                     2
                                                            923
            1
                  1
  2013
                         544
                                         545
                                                    -1
                                                           1004
 2013
5
            1
                  1
                         554
                                         600
                                                    -6
                                                            812
                         554
                                         558
                                                    -4
                                                            740
6 2013
                  1
# i 13 more variables: sched_arr_time <int>, arr_delay <dbl>,
    carrier <chr>, flight <int>, tailnum <chr>, origin <chr>,
    dest <chr>, air_time <dbl>, distance <dbl>, hour <dbl>,
   minute <dbl>, time_hour <dttm>, name <chr>
#
head(flights_outer_join)
# A tibble: 6 x 20
                day dep_time sched_dep_time dep_delay arr_time
  year month
  <int> <int> <int>
                       <int>
                                       <int>
                                                 <dbl>
                                                          <int>
                                                     2
1 2013
                         517
                                         515
                                                            830
2 2013
                         533
                                         529
                                                     4
                                                            850
3 2013
                         542
                                         540
                                                            923
                  1
                                                     2
            1
 2013
                         544
                                         545
                                                    -1
                                                           1004
5 2013
                         554
                                         600
                                                    -6
                                                            812
            1
                  1
                         554
                                         558
                                                            740
 2013
                  1
                                                    -4
# i 13 more variables: sched_arr_time <int>, arr_delay <dbl>,
    carrier <chr>, flight <int>, tailnum <chr>, origin <chr>,
#
    dest <chr>, air_time <dbl>, distance <dbl>, hour <dbl>,
   minute <dbl>, time_hour <dttm>, name <chr>
# Calculating a 5 period rolling average of arrival delays and cumulative sum
flights_rolling <- flights %>%
  arrange(year, month, day) %>%
  mutate(
    arr_delay = ifelse(is.na(arr_delay), 0, arr_delay),
   rolling_avg_delay = zoo::rollmean(arr_delay, 5, fill = NA),
    cumulative_delay = cumsum(arr_delay)
  )
# Display the transformed data
head(flights_rolling)
```

```
# A tibble: 6 x 21
  year month day dep_time sched_dep_time dep_delay arr_time
  <int> <int> <int>
                       <int>
                                      <int>
                                                <dbl>
                                                         <int>
1 2013
                                                    2
                                                           830
            1
                  1
                         517
                                        515
2 2013
                                        529
                                                           850
            1
                  1
                         533
                                                    4
3 2013
                         542
                                        540
                                                    2
                                                           923
            1
                  1
4 2013
                         544
                                        545
                                                   -1
                                                          1004
                  1
5 2013
                                        600
            1
                  1
                         554
                                                   -6
                                                           812
                         554
                                        558
                                                   -4
                                                           740
6 2013
            1
                  1
# i 14 more variables: sched_arr_time <int>, arr_delay <dbl>,
    carrier <chr>, flight <int>, tailnum <chr>, origin <chr>,
   dest <chr>, air_time <dbl>, distance <dbl>, hour <dbl>,
   minute <dbl>, time_hour <dttm>, rolling_avg_delay <dbl>,
#
    cumulative_delay <dbl>
# Plotting the rolling average and cumulative delays
```

Warning: Removed 4 rows containing missing values or values outside the scale range ('geom_line()').



Program - 6

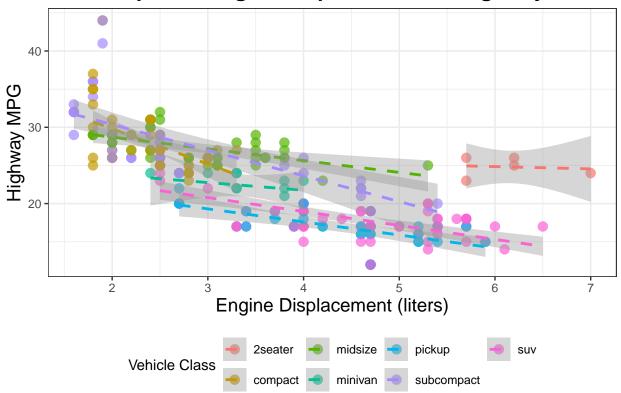
Data Visualisation with ggplot2 and Customisations

Date of Execution - 2025-09-23

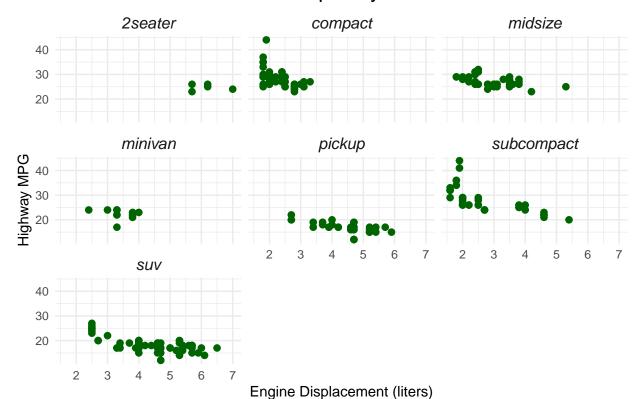
Objective - This program evaluates students' ability to create and customize complex data visualizations using the ggplot2 package.

```
# Load necessary libraries
library(ggplot2)
library(dplyr)
library(reshape2)
# Scatterplot with regression line and confidence intervals
data("mpg")
ggplot(mpg, aes(x = displ, y = hwy, color = class)) +
  geom_point(size = 3, alpha = 0.7) +
  geom_smooth(method = "lm", se = TRUE, linetype = "dashed") +
  labs(title = "Scatterplot of Engine Displacement vs Highway MPG",
       x = "Engine Displacement (liters)",
       y = "Highway MPG",
       color = "Vehicle Class") +
  theme_bw() +
  theme(plot.title = element_text(hjust = 0.5, size = 16, face = "bold"),
        axis.title = element_text(size = 14),
        legend.position = "bottom")
```

Scatterplot of Engine Displacement vs Highway MPG



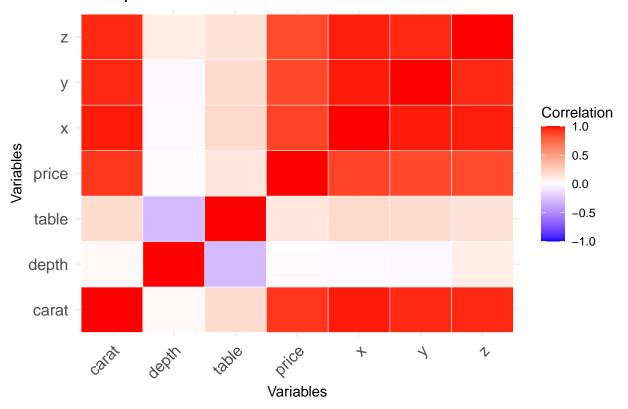
Faceted Scatterplot by Vehicle Class



Heatmap of correlatio matrix data("diamonds") # Calculate correlation matrix for numeric variables cor_matrix <- cor(diamonds[sapply(diamonds, is.numeric)], use = "complete.obs")</pre> #Convert to tidy format cor_melt <- melt(cor_matrix)</pre> # Create heatmap ggplot(cor_melt, aes(Var1, Var2, fill = value)) + geom_tile(color = "white") + scale_fill_gradient2(low = "blue", high = "red", mid = "white", midpoint = 0, limit = c(-1, 1), space = "Lab", name = "Correlation") + labs(title = "Heatmap of Correlation Matrix for Diamonds Dataset", x = "Variables", y = "Variables") + theme minimal() + theme(axis.text.x = element_text(angle = 45, hjust = 1, size = 12), axis.text.y = element_text(size = 12),

```
plot.title = element_text(hjust = 0.5, size = 16))
```

Heatmap of Correlation Matrix for Diamonds Dataset



```
# Enhancing the scatterplot with annotations
ggplot(mpg, aes(x = displ, y = hwy, fill = class)) +

geom_point(size = 3, shape = 21, alpha = 0.8) +

theme_light() +

scale_color_brewer(palette = "Set2") +

labs(title = "Customised Scatter Plot",

x = "Engine Displacement (liters)",

y = "Highway MPG",

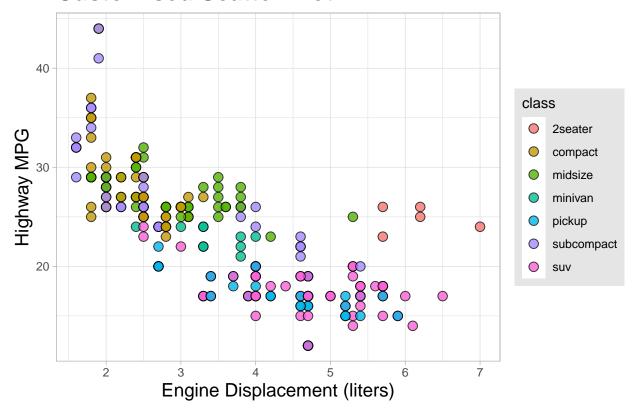
color = "Vehicle Class") +

theme(plot.title = element_text(face = "bold", size = 18),

axis.title = element_text(size = 14),

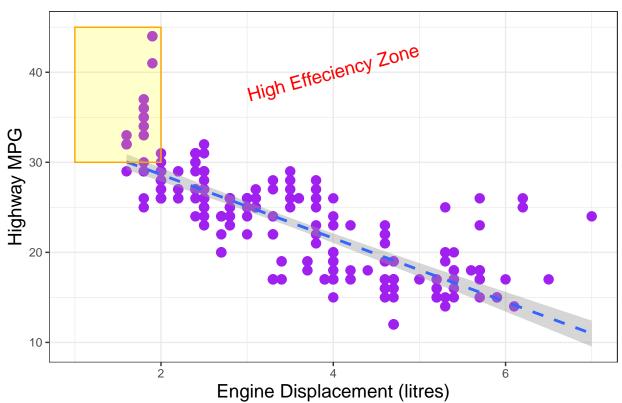
legend.background = element_rect(fill = "gray90"))
```

Customised Scatter Plot



```
# Annotate plots and save as image files
annotated_plot <- ggplot(mpg, aes(x = displ, y = hwy)) +</pre>
  geom_point(size = 3, color = "purple") +
  geom_smooth(method = "lm", se = TRUE, linetype = "dashed") +
  labs(title = "Annotate the Plot",
       x = "Engine Displacement (litres)",
       y = "Highway MPG",
       color = "Vehicle Class") +
  theme_bw() +
  theme(plot.title = element_text(hjust = 0.5, size = 16, face = "bold"),
        axis.title = element_text(size = 14),
        legend.position = "bottom") +
  annotate("text", x = 4, y = 40, label = "High Effeciency Zone",
           color = "red", size = 5, angle = 15) +
  annotate("rect", xmin = 1, xmax = 2, ymin = 30, ymax = 45,
           alpha = 0.2, fill = "yellow", color = "orange")
annotated_plot
```

Annotate the Plot



ggsave("annotated_scatterplot.png", plot = annotated_plot, width = 8, height = 6)

Program - 7

Linear and Multiple Regression Analysis with Interaction Terms

Date of Execution - 2025-10-14

Objective - This program focuses on regression modeling, interaction effects, and model diagnostics.

```
# Load necessary libraries
library(MASS)
library(ggplot2)
library(dplyr)
library(caret)
library(car)
library(pROC)
library(corrplot)
# Load the Boston Housing dataset
data("Boston")
head(Boston)
                                          dis rad tax ptratio
     crim zn indus chas
                         nox
                                rm
                                   age
1 0.00632 18 2.31
                   0 0.538 6.575 65.2 4.0900
                                                1 296
                                                         15.3
2 0.02731 0 7.07 0 0.469 6.421 78.9 4.9671
                                                2 242
                                                         17.8
3 0.02729 0 7.07 0 0.469 7.185 61.1 4.9671
                                                2 242
                                                         17.8
                   0 0.458 6.998 45.8 6.0622
                                                3 222
4 0.03237 0 2.18
                                                         18.7
5 0.06905 0 2.18
                     0 0.458 7.147 54.2 6.0622
                                                3 222
                                                         18.7
                     0 0.458 6.430 58.7 6.0622
6 0.02985 0 2.18
                                                3 222
                                                         18.7
  black 1stat medv
1 396.90 4.98 24.0
2 396.90 9.14 21.6
3 392.83 4.03 34.7
4 394.63 2.94 33.4
5 396.90 5.33 36.2
6 394.12 5.21 28.7
# q1. Preprocessing
# Check for missing values
sum(is.na(Boston))
```

Summary Statistics

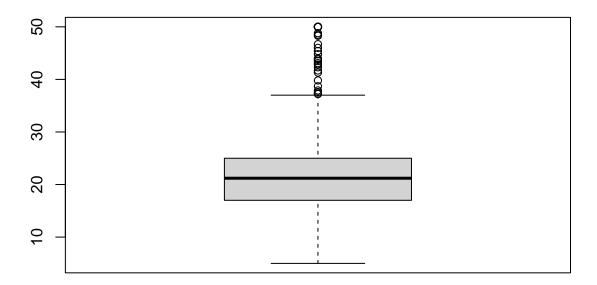
summary(Boston)

```
crim
                      zn
                                    indus
Min. : 0.00632
                 Min. : 0.00
                                 Min. : 0.46
1st Qu.: 0.08205
                 1st Qu.: 0.00
                                 1st Qu.: 5.19
Median : 0.25651
                 Median: 0.00
                                 Median: 9.69
Mean : 3.61352
                 Mean : 11.36
                                 Mean :11.14
3rd Qu.: 3.67708
                 3rd Qu.: 12.50
                                 3rd Qu.:18.10
Max. :88.97620
                 Max. :100.00
                                 Max. :27.74
    chas
                    nox
                                     rm
                                                   age
                                              Min. : 2.90
Min. :0.00000
                Min. :0.3850
                                Min. :3.561
1st Qu.:0.00000
                1st Qu.:0.4490
                                1st Qu.:5.886 1st Qu.: 45.02
Median :0.00000
                Median :0.5380
                                Median:6.208
                                              Median: 77.50
Mean
     :0.06917
                Mean
                     :0.5547
                               Mean :6.285
                                              Mean : 68.57
3rd Qu.:0.00000
                3rd Qu.:0.6240
                                3rd Qu.:6.623
                                              3rd Qu.: 94.08
Max. :1.00000
                Max. :0.8710
                                Max. :8.780
                                              Max. :100.00
    dis
                   rad
                                   tax
                                                ptratio
Min. : 1.130
               Min. : 1.000
                               Min. :187.0
                                             Min. :12.60
1st Qu.: 2.100
               1st Qu.: 4.000
                               1st Qu.:279.0
                                             1st Qu.:17.40
Median : 3.207
               Median : 5.000
                               Median :330.0
                                            Median :19.05
Mean : 3.795
               Mean : 9.549
                               Mean :408.2
                                             Mean :18.46
3rd Qu.: 5.188
               3rd Qu.:24.000
                               3rd Qu.:666.0
                                              3rd Qu.:20.20
Max. :12.127
               Max. :24.000
                               Max. :711.0
                                             Max. :22.00
   black
                   lstat
                                  medv
Min. : 0.32
               Min. : 1.73
                              Min. : 5.00
1st Qu.:375.38
               1st Qu.: 6.95
                              1st Qu.:17.02
Median :391.44
               Median :11.36
                              Median :21.20
Mean :356.67
               Mean :12.65
                              Mean :22.53
3rd Qu.:396.23
               3rd Qu.:16.95
                              3rd Qu.:25.00
Max. :396.90
               Max. :37.97
                              Max. :50.00
```

Check for outliers using boxplots

boxplot(Boston\$medv, main = "Boxplot of Median Value (medv)")

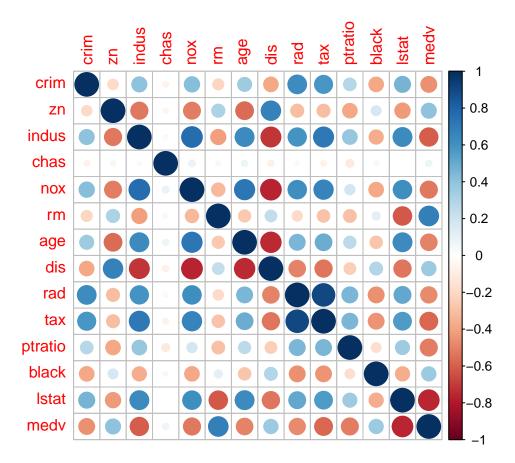
Boxplot of Median Value (medv)



```
# Remove potential outliers (optional, based on domain knowledge)
Boston <- Boston %>% filter(medv < 50)

# 2. Feature Selection

# Calculate correlation matrix
corr_matrix <- cor(Boston)
corrplot(corr_matrix, method = "circle")</pre>
```



```
# High correlation observed between 'medu', 'lstat', and 'rm'
# We will use 'lstat' and 'rm as predictors based on this analysis.

# 3. Simple Linear Regression Model
simple_model <- lm(medv ~ lstat, data = Boston)
summary(simple_model)</pre>
```

Call:

lm(formula = medv ~ lstat, data = Boston)

Residuals:

Min 1Q Median 3Q Max -13.992 -3.313 -0.941 1.914 21.246

Coefficients:

Estimate Std. Error t value Pr(>|t|)

(Intercept) 32.54041 0.48150 67.58 <2e-16 ***

lstat -0.84374 0.03268 -25.82 <2e-16 ***

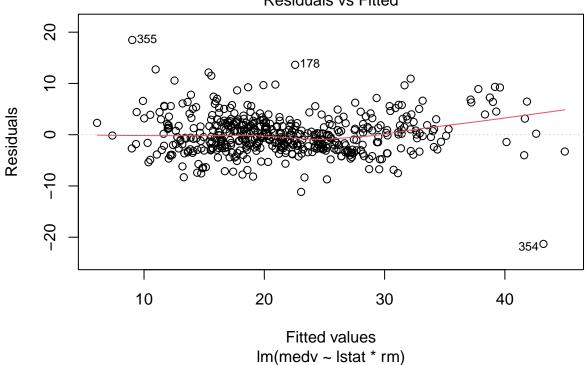
--
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1

```
Multiple R-squared: 0.5774, Adjusted R-squared: 0.5765
F-statistic: 666.6 on 1 and 488 DF, p-value: < 2.2e-16
# Interpretation
# - The negative coeffecient for 'lstat' suggests that higher 'lstat' values
  (higher percentage of lower status population) are associated with lower 'medu'
# (median home value)
# - The p-value (<0.05) indicates that the relationship is statistically significant.
# 4. Multiple Linear Regression
multiple_model <- lm(medv ~ lstat * rm, data = Boston)</pre>
summary(multiple_model)
Call:
lm(formula = medv ~ lstat * rm, data = Boston)
Residuals:
    Min
             1Q Median
                             3Q
                                     Max
-21.3064 -2.4982 -0.3056 1.8635 18.4779
Coefficients:
           Estimate Std. Error t value Pr(>|t|)
(Intercept) -25.99970 3.07045 -8.468 2.98e-16 ***
            lstat
            rm
           lstat:rm
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 3.845 on 486 degrees of freedom
Multiple R-squared: 0.7625, Adjusted R-squared: 0.761
F-statistic: 520 on 3 and 486 DF, p-value: < 2.2e-16
# Interpretation:
# - Significant coeffecients for 'lstat', 'rm', and the interaction term ('lstat:rm')
# - Indicates that the relationship between 'lstat' and 'medv' depends on the value of 'rm'
# - The adjusted R^2 has improved, suggesting better fit when compared to simple model
# 5. Model Performance Evaluation
```

Residual standard error: 5.119 on 488 degrees of freedom

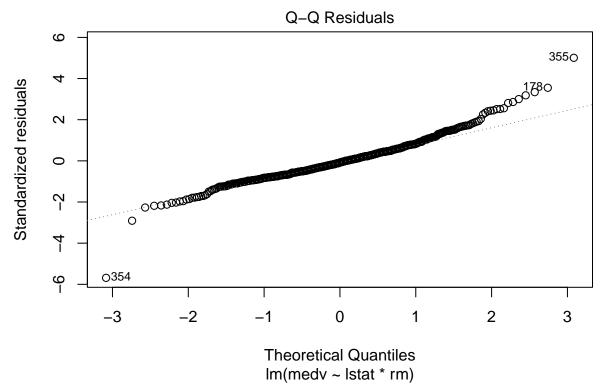
Residuals v/s Fitted Plot

Residuals vs Fitted



```
# Q-Q plot for checking normality of residuals
plot(multiple_model, which = 2, main = "Normal Q-Q Plot")
```

Normal Q-Q Plot



Linear Regression

490 samples
2 predictor

```
Resampling: Cross-Validated (10 fold)
Summary of sample sizes: 441, 441, 442, 441, 441, 440, ...
Resampling results:
 RMSE
           Rsquared
                      MAE
  3.855714 0.7597212 2.868657
Tuning parameter 'intercept' was held constant at a value of TRUE
# Interpretation:
# - Cross Validation RMSE provides a estimate of prediction error
# - Lower RMSE indicates better model performance
# 8. ROC Curve Analysis (Classification Approach)
# Convert 'medv' to a binary classification problem: High (>=25) or Low (<25)
Boston$medv_class <- ifelse(Boston$medv >= 25, 1, 0)
# Fit a logistic regression model for classification
logistic_model <- glm(medv_class ~ lstat*rm, data = Boston, family = "binomial")</pre>
summary(logistic_model)
Call:
glm(formula = medv_class ~ lstat * rm, family = "binomial", data = Boston)
Coefficients:
            Estimate Std. Error z value Pr(>|z|)
(Intercept) -19.80238 7.45022 -2.658 0.00786 **
lstat
           -0.02301 0.75334 -0.031 0.97563
             3.29921 1.12727 2.927 0.00343 **
rm
            -0.03989 0.11492 -0.347 0.72851
lstat:rm
Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
(Dispersion parameter for binomial family taken to be 1)
   Null deviance: 536.34 on 489 degrees of freedom
Residual deviance: 254.70 on 486 degrees of freedom
AIC: 262.7
```

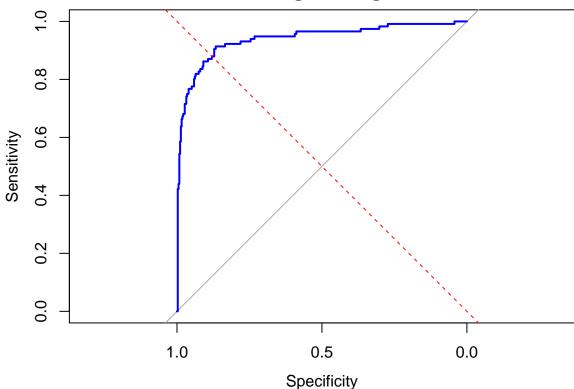
No pre-processing

Number of Fisher Scoring iterations: 8

```
# Predict probabilities and compute ROC Curve
pred_probs <- predict(logistic_model, type = "response")
roc_curve <- roc(Boston$medv_class, pred_probs)

# Plot ROC Curves
plot(roc_curve, main = "ROC Curve for Logistic Regression Model", col = "blue")
abline(a=0, b=1, lty = 2, col = "red")</pre>
```

ROC Curve for Logistic Regression Model



```
cat("AUC: ", auc(roc_curve), "\n")
```

AUC: 0.9392864

Interpretation

- The ROC Curve evaluates the trade-off between sensitivity and specificity

- The Area Under the Curve (AUC) indicates the model's discriminatory ability

(AUC closer to 1 is better)

Program - 8

K-Means Clustering and PCA for Dimensionality Reduction

Date of Execution - 2025-10-28

Objective - This program tests the student's knowledge of clustering techniques and dimensionality reduction through PCA.

```
# Load required libraries
library(rattle)
                    # For Wine dataset
library(ggplot2)
                    # For visualization
library(cluster)
                   # For silhouette scores
library(factoextra) # For PCA and clustering visualization
library(dplyr)
                    # Often useful for data manipulation
# Normalize function (Min-Max Scaling)
normalize <- function(data) {</pre>
  return((data - min(data)) / (max(data) - min(data)))
}
# Analysis for WINE Dataset
# Step 1: Load Wine dataset and normalize
data(wine)
wine_data <- wine[, -1] # Remove the class label
wine_norm <- as.data.frame(lapply(wine_data, normalize))</pre>
# Step 2: Apply PCA
wine_pca <- prcomp(wine_norm, scale. = TRUE)</pre>
summary(wine_pca)
```

Importance of components:

```
PC1 PC2 PC3 PC4 PC5 PC6
Standard deviation 2.169 1.5802 1.2025 0.95863 0.92370 0.80103
Proportion of Variance 0.362 0.1921 0.1112 0.07069 0.06563 0.04936
Cumulative Proportion 0.362 0.5541 0.6653 0.73599 0.80162 0.85098
PC7 PC8 PC9 PC10 PC11 PC12
```

```
      Standard deviation
      0.74231
      0.59034
      0.53748
      0.5009
      0.47517
      0.41082

      Proportion of Variance
      0.04239
      0.02681
      0.02222
      0.0193
      0.01737
      0.01298

      Cumulative Proportion
      0.89337
      0.92018
      0.94240
      0.9617
      0.97907
      0.99205

      PC13

      Standard deviation
      0.32152

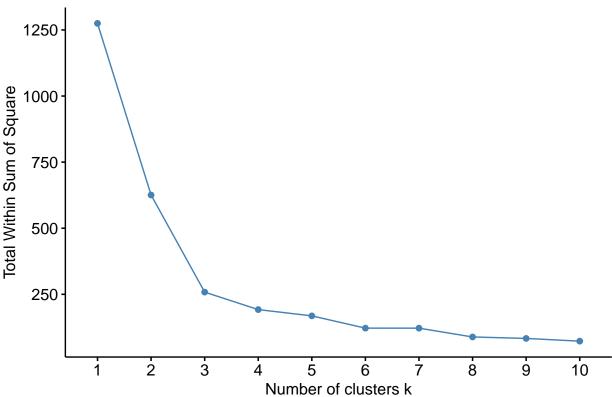
      Proportion of Variance
      0.00795

      Cumulative Proportion
      1.00000
```

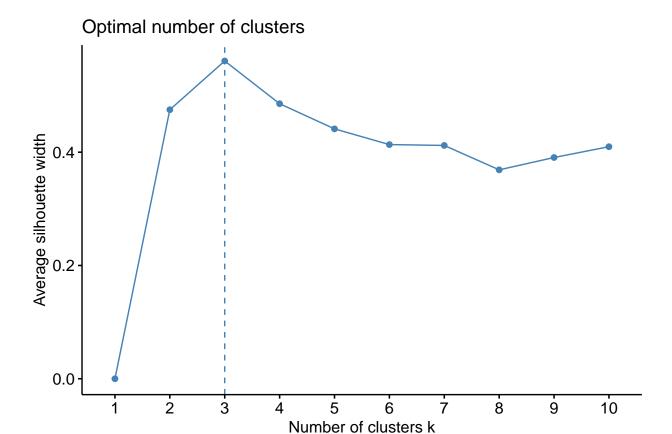
```
# Reduce to top 2 principal components
wine_pca_data <- as.data.frame(wine_pca$x[, 1:2])

# Step 3: Determine the optimal number of clusters (Elbow method)
elbow_wine <- fviz_nbclust(wine_pca_data, kmeans, method = "wss")
print(elbow_wine)</pre>
```





```
# Step 4: Silhouette analysis
silhouette_wine <- fviz_nbclust(wine_pca_data, kmeans, method = "silhouette")
print(silhouette_wine)</pre>
```

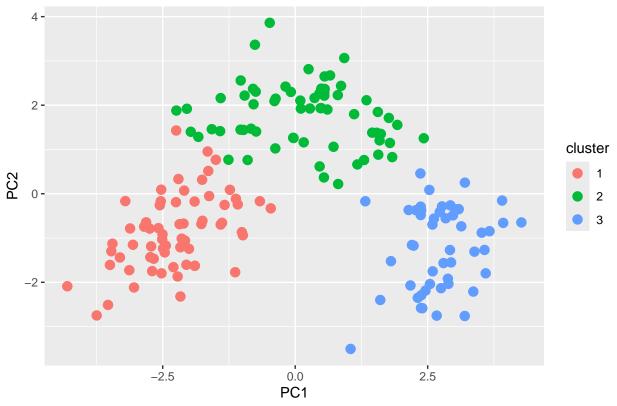


```
# Step 5: Apply K-means clustering (using centers=3 based on known structure/analysis)
set.seed(123)
wine_kmeans <- kmeans(wine_pca_data, centers = 3, nstart = 25)

# Step 6: Visualize clusters
wine_pca_data$cluster <- as.factor(wine_kmeans$cluster)

p1 <- ggplot(wine_pca_data, aes(x = PC1, y = PC2, color = cluster)) +
    geom_point(size = 3) +
    labs(title = "K-Means Clustering on Wine Dataset")
print(p1)</pre>
```





```
# Step 7: Interpret results
cat("Wine Dataset Clustering Results:\n")
```

Wine Dataset Clustering Results:

summary(bc_pca)

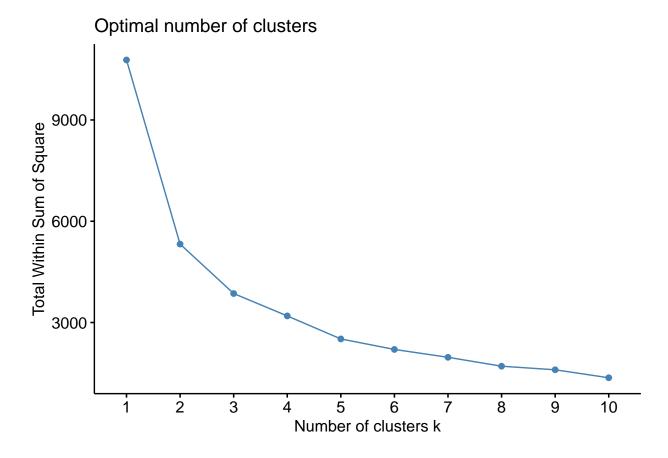
```
Importance of components:
```

```
PC1
                                 PC2
                                         PC3
                                                 PC4
                                                         PC5
                                                                 PC6
Standard deviation
                       3.6444 2.3857 1.67867 1.40735 1.28403 1.09880
Proportion of Variance 0.4427 0.1897 0.09393 0.06602 0.05496 0.04025
Cumulative Proportion 0.4427 0.6324 0.72636 0.79239 0.84734 0.88759
                           PC7
                                   PC8
                                          PC9
                                                 PC10
                                                        PC11
                                                                PC12
Standard deviation
                       0.82172 0.69037 0.6457 0.59219 0.5421 0.51104
Proportion of Variance 0.02251 0.01589 0.0139 0.01169 0.0098 0.00871
Cumulative Proportion 0.91010 0.92598 0.9399 0.95157 0.9614 0.97007
                          PC13
                                  PC14
                                          PC15
                                                  PC16
                                                          PC17
                       0.49128 0.39624 0.30681 0.28260 0.24372
Standard deviation
Proportion of Variance 0.00805 0.00523 0.00314 0.00266 0.00198
Cumulative Proportion 0.97812 0.98335 0.98649 0.98915 0.99113
                          PC18
                                  PC19
                                          PC20
                                                 PC21
                                                         PC22
                                                                 PC23
Standard deviation
                       0.22939 0.22244 0.17652 0.1731 0.16565 0.15602
Proportion of Variance 0.00175 0.00165 0.00104 0.0010 0.00091 0.00081
Cumulative Proportion 0.99288 0.99453 0.99557 0.9966 0.99749 0.99830
                         PC24
                                 PC25
                                         PC26
                                                 PC27
                                                         PC28
                                                                 PC29
Standard deviation
                       0.1344\ 0.12442\ 0.09043\ 0.08307\ 0.03987\ 0.02736
Proportion of Variance 0.0006 0.00052 0.00027 0.00023 0.00005 0.00002
Cumulative Proportion 0.9989 0.99942 0.99969 0.99992 0.99997 1.00000
                          PC30
Standard deviation
                       0.01153
Proportion of Variance 0.00000
Cumulative Proportion 1.00000
# Reduce to top 2 principal components
bc_pca_data <- as.data.frame(bc_pca$x[, 1:2])</pre>
```

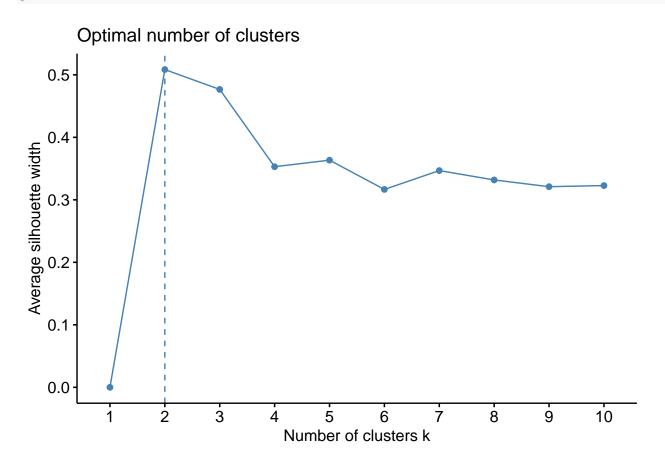
Optimal number of clusters (Elbow method)

print(elbow_bc)

elbow_bc <- fviz_nbclust(bc_pca_data, kmeans, method = "wss")</pre>



```
# Optimal number of clusters (Silhouette analysis)
silhouette_bc <- fviz_nbclust(bc_pca_data, kmeans, method = "silhouette")
print(silhouette_bc)</pre>
```

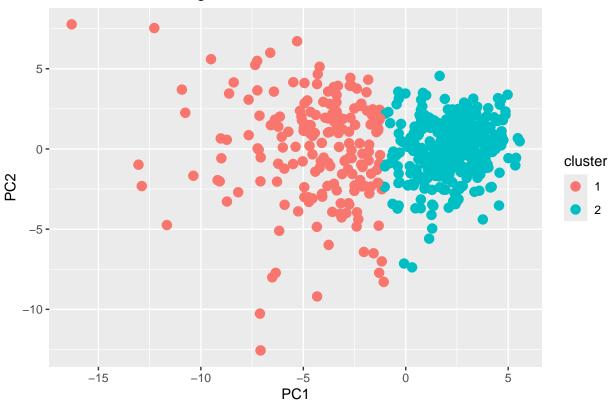


```
# Apply K-means clustering (using centers=2 based on binary classification/analysis)
set.seed(123)
bc_kmeans <- kmeans(bc_pca_data, centers = 2, nstart = 25)

# Add cluster assignments to PCA data
bc_pca_data$cluster <- as.factor(bc_kmeans$cluster)

# Visualize clusters
p2 <- ggplot(bc_pca_data, aes(x = PC1, y = PC2, color = cluster)) +
    geom_point(size = 3) +
    labs(title = "K-Means Clustering on Breast Cancer Dataset")
print(p2)</pre>
```

K-Means Clustering on Breast Cancer Dataset



```
# Interpret results
cat("Breast Cancer Dataset Clustering Results:\n")
```

Breast Cancer Dataset Clustering Results:

```
cat("Cluster Sizes: ", bc_kmeans$size, "\n")
```

Cluster Sizes: 191 378

Program - 9

Time Series Analysis using ARIMA and Seasonal Decomposition

Date of Execution - 2025-10-28

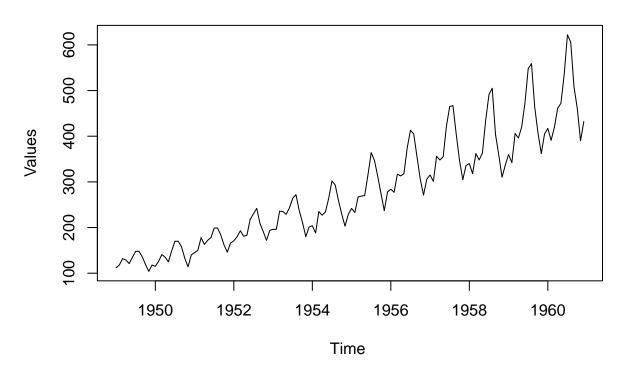
Objective - This program evaluates students' skills in time series analysis, model fitting, and forecasting.

```
# Load required libraries for time series analysis and modeling
library(forecast)
library(ggplot2)
library(TSA)
library(tseries)
# Function to perform Exploratory Data Analysis (EDA) on the time series data
perform_eda <- function(ts_data, dataset_name) {</pre>
  cat(" Exploratory Data Analysis for ", dataset_name, "\n")
  print(summary(ts_data)) # Print summary of the dataset
  plot(ts_data, main = paste(dataset_name, " Time Series "), ylab = " Values ", xlab = "Time ")
  cat("ACF and PACF plots :\n")
  acf(ts_data, main = paste("ACF of", dataset_name)) # Autocorrelation plot
 pacf(ts_data, main = paste(" PACF of", dataset_name)) # Partial autocorrelation plot
}
# Function to decompose the time series into trend , seasonal , and residual components
decompose_ts <- function(ts_data, dataset_name) {</pre>
  cat(" Decomposing the time series for ", dataset_name, "\n")
  decomposition <- decompose(ts_data) # Decompose the time series
 plot(decomposition) # Plot the decomposition
  return(decomposition) # Return the decomposition result
# Function to fit an ARIMA model to the time series data
fit_arima <- function(ts_data, dataset_name) {</pre>
  cat(" Fitting ARIMA model for ", dataset_name, "\n")
  adf_test <- adf.test(ts_data, alternative = "stationary") # ADF test for stationarity
  cat("ADF Test p- value :", adf_test$p.value, "\n")
  # If p- value > 0.05 , data is non - stationary , so we difference the data
  if (adf_test$p.value > 0.05) {
    ts_data <- diff(ts_data) # Difference the data to make it stationary
```

```
plot(ts_data, main = paste(dataset_name, " Differenced Time Series "))
  }
  auto_model <- auto.arima(ts_data, seasonal = FALSE) # Fit ARIMA model (non - seasonal)
  print(summary(auto_model)) # Print ARIMA model summary
  forecast_result <- forecast(auto_model, h = 12) # Forecast next 12 periods
 plot(forecast_result, main = paste(dataset_name, " ARIMA Forecast ")) # Plot ARIMA forecast
  return(auto_model) # Return the fitted ARIMA model
}
# Function to fit a Seasonal ARIMA ( SARIMA ) model to the time series data
fit_sarima <- function(ts_data, dataset_name) {</pre>
  cat(" Fitting SARIMA model for ", dataset_name, "\n")
  auto_sarima <- auto.arima(ts_data, seasonal = TRUE) # Fit SARIMA model ( seasonal )</pre>
  print(summary(auto_sarima)) # Print SARIMA model summary
  sarima_forecast <- forecast(auto_sarima, h = 12) # Forecast next 12 periods</pre>
 plot(sarima_forecast, main = paste(dataset_name, " SARIMA Forecast ")) # Plot SARIMA forecast
  return(auto_sarima) # Return the fitted SARIMA model
}
# Function to compare ARIMA and SARIMA models by evaluating forecast accuracy
compare_models <- function(arima_model, sarima_model, ts_data) {</pre>
  cat(" Comparing ARIMA and SARIMA models :\n")
  h <- min(12, length(ts_data)) # Forecast horizon of 12 or adjusted based on dataset length
  arima_forecast <- forecast(arima_model, h = h) # ARIMA forecast</pre>
  sarima_forecast <- forecast(sarima_model, h = h) # SARIMA forecast</pre>
  actual_values <- ts_data[(length(ts_data) - h + 1): length(ts_data)] # Comparison
  # Calculate accuracy of both models
  arima_accuracy <- accuracy(arima_forecast$mean, actual_values)</pre>
  sarima_accuracy <- accuracy(sarima_forecast$mean, actual_values)</pre>
  cat(" ARIMA Forecast Accuracy :\n", arima_accuracy) # Print ARIMA accuracy
  cat(" SARIMA Forecast Accuracy :\n", sarima_accuracy) # Print SARIMA accuracy
}
# Function to visualize the comparison of ARIMA and SARIMA forecast performance
plot_forecast_comparison <- function(actual_values, arima_forecast, sarima_forecast, time_points) {</pre>
  arima_rmse <- sqrt(mean((arima_forecast - actual_values)^2)) # Calculate RMSE for ARIMA
  sarima_rmse <- sqrt(mean((sarima_forecast - actual_values)^2)) # Calculate RMSE for SARIMA</pre>
  # Color coding for better and worse RMSE
```

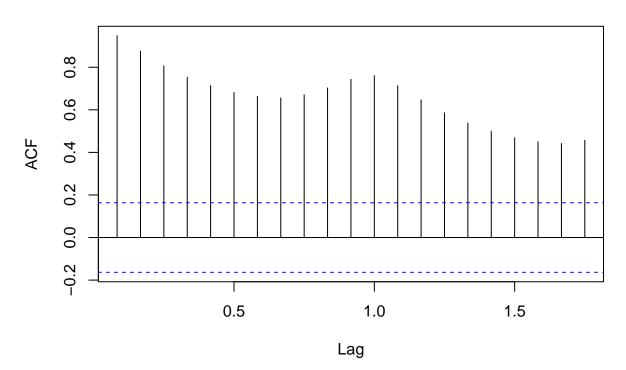
```
better_color <- ifelse(arima_rmse < sarima_rmse, "green", "red")</pre>
  worse_color <- ifelse(arima_rmse < sarima_rmse, "red", "green")</pre>
  # Plot actual values and forecasts
  plot(time_points, actual_values, type = "o", col = "blue", pch = 16, lty = 1, xlab = " Time ",
       ylab = " Values ", main = " Forecast Comparison ")
  lines(time_points, arima_forecast, col = better_color, lty = 2, lwd = 2) # ARIMA forecast line
  lines(time_points, sarima_forecast, col = worse_color, lty = 3, lwd = 2) # SARIMA forecast line
  # Add a legend to the plot
  legend("topright", legend = c("Actual Values", paste("ARIMA (RMSE =", round(arima_rmse, 2), ")"),
                                paste("SARIMA (RMSE =", round(sarima_rmse, 2), ")")),
         col = c("blue", better_color, worse_color), lty = c(1, 2, 3), lwd = c(1, 2, 2),
        pch = c(16, NA, NA))
}
# AirPassengers Dataset Analysis
data("AirPassengers")
air_data <- AirPassengers</pre>
cat("\n- - - AirPassengers Dataset - - -\n")
- - - AirPassengers Dataset - - -
perform_eda(air_data, "AirPassengers")
 Exploratory Data Analysis for AirPassengers
  Min. 1st Qu. Median Mean 3rd Qu.
                                          Max.
  104.0 180.0
                  265.5 280.3 360.5
                                          622.0
```

AirPassengers Time Series

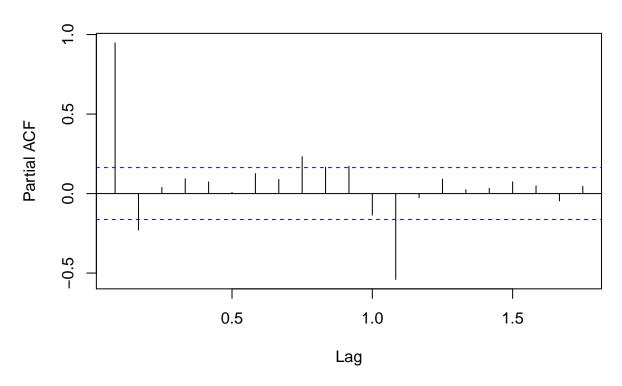


ACF and PACF plots :

ACF of AirPassengers



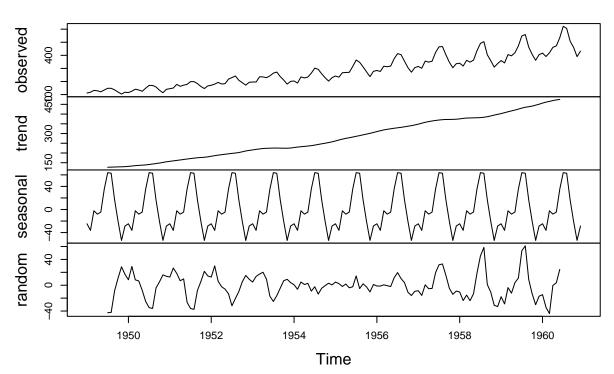
PACF of AirPassengers



decompose_ts(air_data, "AirPassengers")

Decomposing the time series for AirPassengers

Decomposition of additive time series



\$x

 Jan
 Feb
 Mar
 Apr
 May
 Jun
 Jul
 Aug
 Sep
 Oct
 Nov
 Dec

 1949
 112
 118
 132
 129
 121
 135
 148
 148
 136
 119
 104
 118

 1950
 115
 126
 141
 135
 125
 149
 170
 170
 158
 133
 114
 140

 1951
 145
 150
 178
 163
 172
 178
 199
 199
 184
 162
 146
 166

 1952
 171
 180
 193
 181
 183
 218
 230
 242
 209
 191
 172
 194

 1953
 196
 196
 236
 235
 229
 243
 264
 272
 237
 211
 180
 201

 1954
 204
 188
 235
 227
 234
 264
 302
 293
 259
 229
 203
 229

\$seasonal

	Jan	Feb	Mar	Apr	May
1949	-24.748737	-36.188131	-2.241162	-8.036616	-4.506313
1950	-24.748737	-36.188131	-2.241162	-8.036616	-4.506313
1951	-24.748737	-36.188131	-2.241162	-8.036616	-4.506313
1952	-24.748737	-36.188131	-2.241162	-8.036616	-4.506313
1953	-24.748737	-36.188131	-2.241162	-8.036616	-4.506313
1954	-24.748737	-36.188131	-2.241162	-8.036616	-4.506313
1955	-24.748737	-36.188131	-2.241162	-8.036616	-4.506313
1956	-24.748737	-36.188131	-2.241162	-8.036616	-4.506313
1957	-24.748737	-36.188131	-2.241162	-8.036616	-4.506313
1958	-24.748737	-36.188131	-2.241162	-8.036616	-4.506313
1959	-24.748737	-36.188131	-2.241162	-8.036616	-4.506313
1960	-24.748737	-36.188131	-2.241162	-8.036616	-4.506313
	Jun	Jul	Aug	Sep	Oct
1949	35.402778	63.830808	62.823232	16.520202	-20.642677
1950	35.402778	63.830808	62.823232	16.520202	-20.642677
1951	35.402778	63.830808	62.823232	16.520202	-20.642677
1952	35.402778	63.830808	62.823232	16.520202	-20.642677
1953	35.402778	63.830808	62.823232	16.520202	-20.642677
1954	35.402778	63.830808	62.823232	16.520202	-20.642677
1955	35.402778	63.830808	62.823232	16.520202	-20.642677
1956	35.402778	63.830808	62.823232	16.520202	-20.642677
1957	35.402778	63.830808	62.823232	16.520202	-20.642677
1958	35.402778	63.830808	62.823232	16.520202	-20.642677
1959	35.402778	63.830808	62.823232	16.520202	-20.642677

1960 35.402778 63.830808 62.823232 16.520202 -20.642677

Nov Dec

1949 -53.593434 -28.619949

1950 -53.593434 -28.619949

1951 -53.593434 -28.619949

1952 -53.593434 -28.619949

1953 -53.593434 -28.619949

1954 -53.593434 -28.619949

1955 -53.593434 -28.619949

1956 -53.593434 -28.619949

1957 -53.593434 -28.619949

1958 -53.593434 -28.619949

1959 -53.593434 -28.619949

1960 -53.593434 -28.619949

\$trend

	Jan	Feb	Mar	Apr	May	Jun	Jul
1949	NA	NA	NA	NA	NA	NA	126.7917
1950	131.2500	133.0833	134.9167	136.4167	137.4167	138.7500	140.9167
1951	157.1250	159.5417	161.8333	164.1250	166.6667	169.0833	171.2500
1952	183.1250	186.2083	189.0417	191.2917	193.5833	195.8333	198.0417
1953	215.8333	218.5000	220.9167	222.9167	224.0833	224.7083	225.3333
1954	228.0000	230.4583	232.2500	233.9167	235.6250	237.7500	240.5000
1955	261.8333	266.6667	271.1250	275.2083	278.5000	281.9583	285.7500
1956	309.9583	314.4167	318.6250	321.7500	324.5000	327.0833	329.5417
1957	348.2500	353.0000	357.6250	361.3750	364.5000	367.1667	369.4583
1958	375.2500	377.9167	379.5000	380.0000	380.7083	380.9583	381.8333
1959	402.5417	407.1667	411.8750	416.3333	420.5000	425.5000	430.7083
1960	456.3333	461.3750	465.2083	469.3333	472.7500	475.0417	NA
	Aug	Sep	Oct	Nov	Dec		
1949	127.2500	127.9583	128.5833	129.0000	129.7500		
1950	143.1667	145.7083	148.4167	151.5417	154.7083		
1951	173.5833	175.4583	176.8333	178.0417	180.1667		
1952	199.7500	202.2083	206.2500	210.4167	213.3750		
1953	225.3333	224.9583	224.5833	224.4583	225.5417		
1954	243.9583	247.1667	250.2500	253.5000	257.1250		
1955	289.3333	293.2500	297.1667	301.0000	305.4583		
1956	331.8333	334.4583	337.5417	340.5417	344.0833		
1957	371.2083	372.1667	372.4167	372.7500	373.6250		
1958	383.6667	386.5000	390.3333	394.7083	398.6250		
1959	435.1250	437.7083	440.9583	445.8333	450.6250		

1960 NA NA NA NA

\$random

	Jan	Feb	Mar	Apr	May
1949	NA	NA	NA	NA	NA
1950	8.4987374	29.1047980	8.3244949	6.6199495	-7.9103535
1951	12.6237374	26.6464646	18.4078283	6.9116162	9.8396465
1952	12.6237374	29.9797980	6.1994949	-2.2550505	-6.0770202
1953	4.9154040	13.6881313	17.3244949	20.1199495	9.4229798
1954	0.7487374	-6.2702020	4.9911616	1.1199495	2.8813131
1955	4.9154040	2.5214646	-1.8838384	1.8282828	-3.9936869
1956	-1.2095960	-1.2285354	0.6161616	-0.7133838	-1.9936869
1957	-8.5012626	-15.8118687	0.6161616	-5.3383838	-4.9936869
1958	-10.5012626	-23.7285354	-15.2588384	-23.9633838	-13.2020202
1959	-17.7929293	-28.9785354	-3.6338384	-12.2967172	4.0063131
1960	-14.5845960	-34.1868687	-43.9671717	-0.2967172	3.7563131
	Jun	Jul	Aug	Sep	Oct
1949	NA	-42.6224747	-42.0732323	-8.4785354	11.0593434
1950	-25.1527778	-34.7474747	-35.9898990	-4.2285354	5.2260101
1951	-26.4861111	-36.0808081	-37.4065657	-7.9785354	5.8093434
1952	-13.2361111	-31.8724747	-20.5732323	-9.7285354	5.3926768
1953	-17.1111111	-25.1641414	-16.1565657	-4.4785354	7.0593434
1954	-9.1527778	-2.3308081	-13.7815657	-4.6868687	-0.6073232
1955	-2.3611111	14.4191919	-5.1565657	2.2297980	-2.5239899
1956	11.5138889	19.6275253	10.3434343	4.0214646	-10.8989899
1957	19.4305556	31.7108586	32.9684343	15.3131313	-4.7739899
1958	18.6388889	45.3358586	58.5101010	0.9797980	-10.6906566
1959	11.0972222	53.4608586	61.0517677	8.7714646	-13.3156566
1960	24.555556	NA	NA	NA	NA
	Nov	Dec			
1949	28.5934343	16.8699495			
1950	16.0517677	13.9116162			
1951	21.5517677	14.4532828			
1952	15.1767677	9.2449495			
1953	9.1351010	4.0782828			
1954	3.0934343	0.4949495			
1955	-10.4065657	1.1616162			
1956	-15.9482323	-9.4633838			
1957	-14.1565657	-9.0050505			
1958	-31.1148990	-33.0050505			
1959	-30.2398990	-17.0050505			

1960 NA NA

\$figure

[1] -24.748737 -36.188131 -2.241162 -8.036616 -4.506313 35.402778

[7] 63.830808 62.823232 16.520202 -20.642677 -53.593434 -28.619949

\$type

[1] "additive"

attr(,"class")

[1] "decomposed.ts"

arima_air <- fit_arima(air_data, "AirPassengers")</pre>

Fitting ARIMA model for AirPassengers

Warning in adf.test(ts_data, alternative = "stationary"): p-value smaller than printed p-value

ADF Test p- value : 0.01

Series: ts_data

ARIMA(4,1,2) with drift

Coefficients:

ar1 ar2 ar3 ar4 ma1 ma2 drift
0.2243 0.3689 -0.2567 -0.2391 -0.0971 -0.8519 2.6809
s.e. 0.1047 0.1147 0.0985 0.0919 0.0866 0.0877 0.1711

 $sigma^2 = 706.3$: log likelihood = -670.07

AIC=1356.15 AICc=1357.22 BIC=1379.85

Training set error measures:

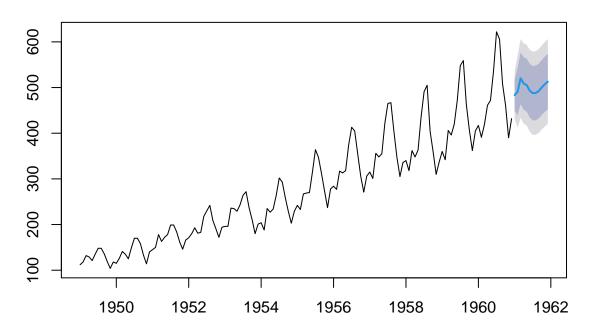
ME RMSE MAE MPE MAPE MASE

Training set -1.228696 25.82793 20.59211 -1.665245 7.476447 0.6428946

ACF1

Training set 0.0009861078

AirPassengers ARIMA Forecast



sarima_air <- fit_sarima(air_data, "AirPassengers")</pre>

Fitting SARIMA model for AirPassengers

Series: ts_data

ARIMA(2,1,1)(0,1,0)[12]

Coefficients:

ar1 ar2 ma1

0.5960 0.2143 -0.9819 s.e. 0.0888 0.0880 0.0292

 $sigma^2 = 132.3$: log likelihood = -504.92

AIC=1017.85 AICc=1018.17 BIC=1029.35

Training set error measures:

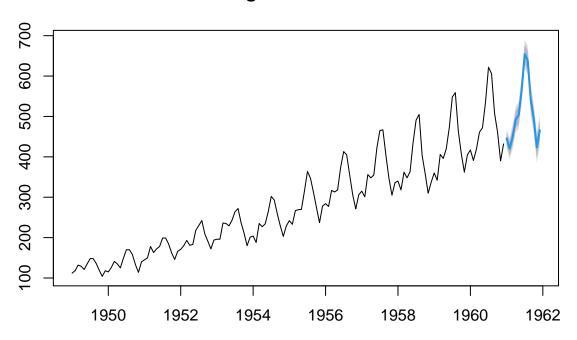
ME RMSE MAE MPE MAPE MASE

Training set 1.3423 10.84619 7.86754 0.420698 2.800458 0.245628

ACF1

Training set -0.00124847

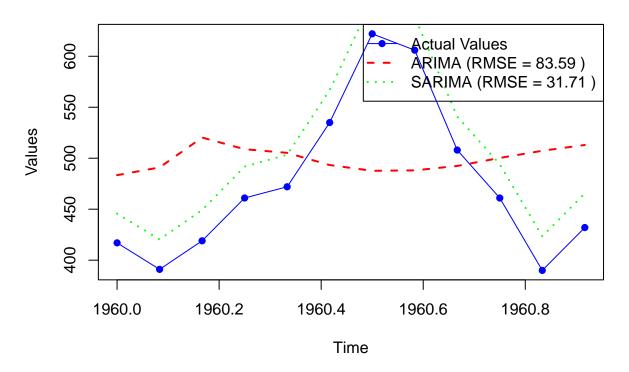
AirPassengers SARIMA Forecast



compare_models(arima_air, sarima_air, air_data)

```
Comparing ARIMA and SARIMA models :
 ARIMA Forecast Accuracy :
 -23.06901 83.58979 74.65149 -7.376947 16.02863 SARIMA Forecast Accuracy :
 -31.66945 31.70666 31.66945 -6.784721 6.784721
# Forecasting and plot comparison for AirPassengers dataset
h_air <- 12 # Define forecast horizon for AirPassengers dataset (12 months ahead)
# Extract the actual values for the last 12 months of the AirPassengers data
air_actual_values <- air_data[(length(air_data) - h_air + 1): length(air_data)]
# Generate ARIMA forecast for the next 12 months
arima_air_forecast <- forecast(arima_air, h = h_air)$mean</pre>
\# Generate SARIMA forecast for the next 12 months
sarima_air_forecast <- forecast(sarima_air, h = h_air)$mean</pre>
# Extract the time points for the last 12 months
time_points_air <- time(air_data)[(length(air_data) - h_air + 1): length(air_data)]</pre>
# Plot and compare the forecasts from ARIMA and SARIMA models against the actual values
plot_forecast_comparison(air_actual_values, arima_air_forecast, sarima_air_forecast,
                         time_points_air)
```

Forecast Comparison



```
# Monthly Milk Production Dataset Analysis
data(milk) # Load the Monthly Milk Production dataset
milk_data <- milk # Assign the dataset to a variable
cat("\n- - - Monthly Milk Production Dataset - - -\n")</pre>
```

- - - Monthly Milk Production Dataset - - -

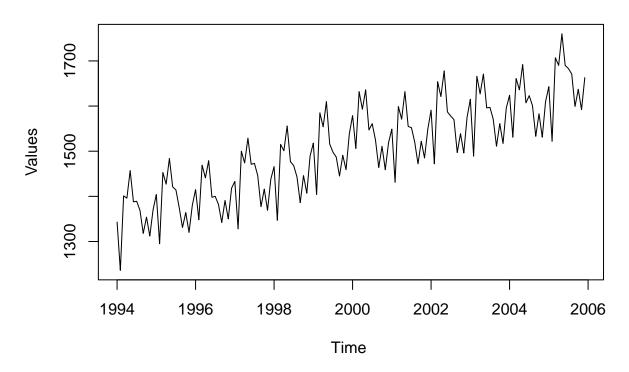
Perform Exploratory Data Analysis (EDA) for the Milk Production dataset
perform_eda(milk_data, "Monthly Milk Production")

Exploratory Data Analysis for $\mbox{Monthly Milk Production}$

milk

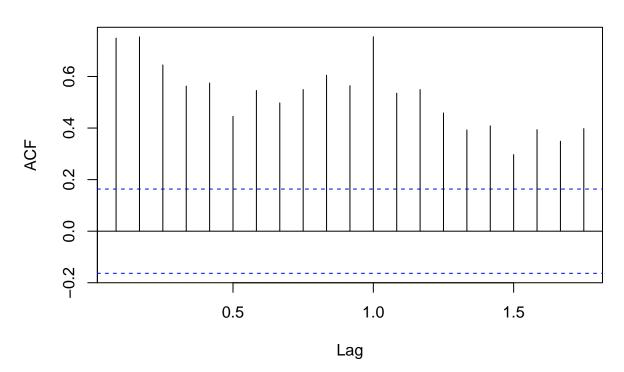
Min. :1236 1st Qu.:1420 Median :1504 Mean :1504 3rd Qu.:1588 Max. :1760

Monthly Milk Production Time Series

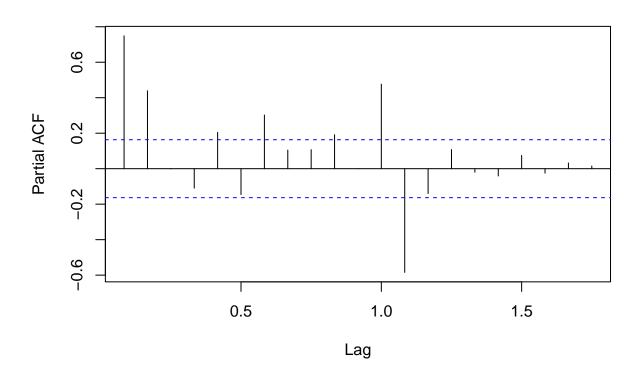


ACF and PACF plots :

ACF of Monthly Milk Production



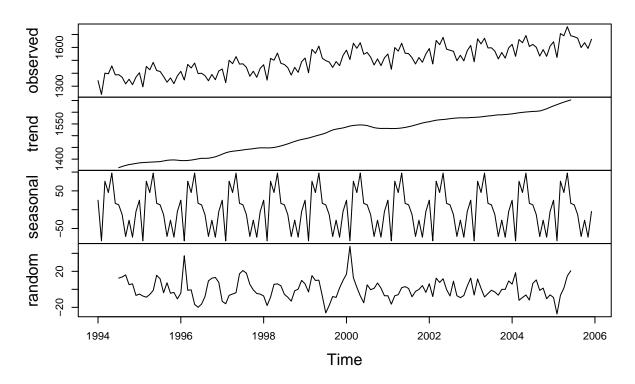
PACF of Monthly Milk Production



Decompose the Milk Production time series into trend , seasonal , and residual components
decompose_ts(milk_data, "Monthly Milk Production")

Decomposing the time series for Monthly Milk Production

Decomposition of additive time series



\$x

 Jan
 Feb
 Mar
 Apr
 May
 Jun
 Jul
 Aug
 Sep
 Oct
 Nov
 Dec

 1994
 1343
 1236
 1401
 1396
 1457
 1388
 1389
 1369
 1318
 1354
 1312
 1370

 1995
 1404
 1295
 1453
 1427
 1484
 1421
 1414
 1375
 1331
 1364
 1320
 1380

 1996
 1415
 1348
 1469
 1441
 1479
 1398
 1400
 1382
 1342
 1391
 1350
 1418

 1997
 1433
 1328
 1500
 1474
 1529
 1471
 1473
 1446
 1377
 1416
 1369
 1438

 1998
 1466
 1347
 1515
 1501
 1556
 1477
 1468
 1443
 1346
 1446
 1407
 1489

 1999
 1518
 1404
 1585
 1554
 1610
 1516
 1498
 1487
 1445
 1451</t

\$seasonal

	Jan	Feb	Mar	Apr	May	Jun
1994	25.16919	-82.90657	75.61237	45.65783	97.34343	16.80934
1995	25.16919	-82.90657	75.61237	45.65783	97.34343	16.80934
1996	25.16919	-82.90657	75.61237	45.65783	97.34343	16.80934
1997	25.16919	-82.90657	75.61237	45.65783	97.34343	16.80934
1998	25.16919	-82.90657	75.61237	45.65783	97.34343	16.80934

25.16919 -82.90657 75.61237 45.65783 97.34343 16.80934 1999 2000 25.16919 -82.90657 75.61237 45.65783 97.34343 16.80934 2001 25.16919 -82.90657 75.61237 45.65783 97.34343 16.80934 2002 25.16919 -82.90657 75.61237 45.65783 97.34343 16.80934 2003 25.16919 -82.90657 75.61237 45.65783 97.34343 16.80934 2004 25.16919 -82.90657 75.61237 45.65783 97.34343 16.80934 2005 25.16919 -82.90657 75.61237 45.65783 97.34343 16.80934 Jul Oct Nov Dec Aug Sep 1994 12.89646 -13.32323 -71.29293 -27.92929 -73.19066 -4.84596 1995 12.89646 -13.32323 -71.29293 -27.92929 -73.19066 -4.84596 1996 12.89646 -13.32323 -71.29293 -27.92929 -73.19066 -4.84596 1997 12.89646 -13.32323 -71.29293 -27.92929 -73.19066 -4.84596 1998 12.89646 -13.32323 -71.29293 -27.92929 -73.19066 -4.84596 1999 12.89646 -13.32323 -71.29293 -27.92929 -73.19066 -4.84596 2000 12.89646 -13.32323 -71.29293 -27.92929 -73.19066 -4.84596 12.89646 -13.32323 -71.29293 -27.92929 -73.19066 2001 -4.84596 2002 12.89646 -13.32323 -71.29293 -27.92929 -73.19066 -4.84596 2003 12.89646 -13.32323 -71.29293 -27.92929 -73.19066 -4.84596 2004 12.89646 -13.32323 -71.29293 -27.92929 -73.19066 -4.84596 2005 12.89646 -13.32323 -71.29293 -27.92929 -73.19066 -4.84596

\$trend

Jan Feb May Jul Mar Apr Jun. 1994 NA 1363.625 NANA NANA NA 1995 1384.042 1385.333 1386.125 1387.083 1387.833 1388.583 1389.458 1996 1393.917 1393.625 1394.375 1395.958 1398.333 1401.167 1403.500 1997 1421.208 1426.917 1431.042 1433.542 1435.375 1437.000 1439.208 1998 1448.208 1447.875 1448.125 1449.750 1452.583 1456.292 1460.583 1999 1486.750 1489.833 1494.125 1498.458 1502.500 1506.708 1511.292 2000 1536.875 1541.083 1543.458 1545.083 1545.917 1545.125 1543.083 2001 1530.958 1530.375 1530.500 1531.292 1532.833 1535.167 1538.167 2002 1559.667 1562.833 1565.958 1567.708 1568.875 1570.417 1572.500 2003 1577.375 1578.208 1578.833 1580.333 1582.125 1583.875 1585.125 2004 1593.083 1595.417 1597.583 1599.417 1600.917 1602.083 1603.458 2005 1626.917 1632.333 1638.000 1643.000 1647.792 1652.542 Aug Sep Oct. Nov Dec 1994 1368.625 1373.250 1376.708 1379.125 1381.625 1995 1392.125 1395.000 1396.250 1396.625 1395.458 1996 1403.417 1403.875 1406.542 1410.000 1415.125 1997 1441.375 1442.792 1444.542 1446.792 1448.167 1998 1465.125 1470.417 1475.542 1480.000 1483.875

1999 1518.083 1524.292 1527.875 1530.583 1532.958
2000 1538.708 1534.208 1531.917 1530.833 1531.000
2001 1541.625 1545.625 1550.000 1554.000 1557.250
2002 1574.208 1575.417 1576.167 1576.125 1576.208
2003 1587.250 1588.792 1588.958 1590.208 1591.542
2004 1603.875 1605.417 1609.583 1614.667 1620.958
2005 NA NA NA NA NA NA

\$random

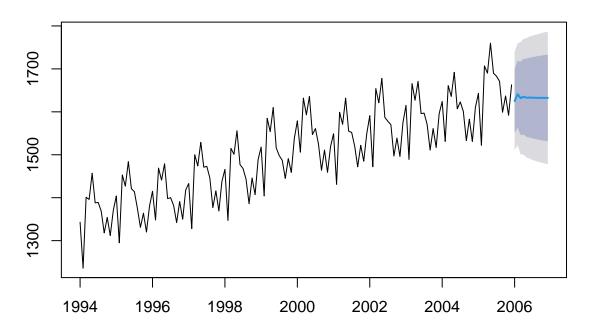
	Jan	Feb	Mar	Apr	May
1994	NA	NA	NA	NA	NA
1995	-5.21085859	-7.42676768	-8.73737374	-5.74116162	-1.17676768
1996	-4.08585859	37.28156566	-0.98737374	-0.61616162	-16.67676768
1997	-13.37752525	-16.01010101	-6.65404040	-5.19949495	-3.71843434
1998	-7.37752525	-17.96843434	-8.73737374	5.59217172	6.07323232
1999	6.08080808	-2.92676768	15.26262626	9.88383838	10.15656566
2000	16.95580808	47.82323232	12.92929293	2.25883838	-7.26010101
2001	-7.12752525	-16.46843434	-7.11237374	-5.94949495	1.82323232
2002	6.16414141	-7.92676768	12.42929293	7.63383838	11.78156566
2003	12.45580808	-6.30176768	11.55429293	1.00883838	-8.46843434
2004	5.74747475	18.48989899	-12.19570707	-9.07449495	-6.26010101
2005	-9.08585859	-27.42676768	-6.61237374	1.34217172	14.86489899
	Jun	Jul	Aug	Sep	Oct
1994	NA	12.47853535	13.69823232	16.04292929	5.22095960
1995	15.60732323	11.64520202	-3.80176768	7.29292929	-4.32070707
1996	-19.97601010	-16.39646465	-8.09343434	9.41792929	12.38762626
1997	17.19065657	20.89520202	17.94823232	5.50126263	-0.61237374
1998	3.89898990	-5.47979798	-8.80176768	-13.12373737	-1.61237374
1999	-7.51767677	-26.18813131	-17.76010101	-7.99873737	-8.94570707
2000	-14.93434343	5.02020202	-0.38510101	1.08459596	7.01262626
2001	3.02398990	0.93686869	-8.30176768	-2.33207071	-0.07070707
2002	-0.22601010	-7.39646465	9.11489899	-7.12373737	-9.23737374
2003	-4.68434343	-1.02146465	-2.92676768	-6.49873737	-0.02904040
2004	-11.89267677	6.64520202	10.44823232	-1.12373737	1.34595960
2005	20.64898990	NA	NA	NA	NA
	Nov	Dec			
1994	6.06565657	-6.77904040			
1995	-3.43434343	-10.61237374			
1996	13.19065657	7.72095960			
1997	-4.60101010	-5.32070707			

1998 0.19065657 9.97095960

```
1999
     1.60732323 9.88762626
2000 1.35732323 -7.15404040
2001 4.19065657 -3.40404040
2002 -6.93434343 3.63762626
2003 -0.01767677 9.30429293
2004 -10.47601010 -6.11237374
2005
              NA
$figure
 [1] 25.16919 -82.90657 75.61237 45.65783 97.34343 16.80934
 [7] 12.89646 -13.32323 -71.29293 -27.92929 -73.19066 -4.84596
$type
[1] "additive"
attr(,"class")
[1] "decomposed.ts"
# Fit ARIMA model for the Milk Production dataset
arima_milk <- fit_arima(milk_data, "Monthly Milk Production")</pre>
Fitting ARIMA model for Monthly Milk Production
Warning in adf.test(ts_data, alternative = "stationary"): p-value
smaller than printed p-value
ADF Test p- value : 0.01
Series: ts_data
ARIMA(2,1,1)
Coefficients:
        ar1 ar2
                        ma1
     0.2066 0.3330 -0.9109
s.e. 0.0879 0.0869 0.0336
sigma^2 = 3373: log likelihood = -782.65
AIC=1573.29 AICc=1573.58 BIC=1585.14
Training set error measures:
                  ΜE
                        RMSE MAE MPE
                                                   MAPE
```

Training set 10.57461 57.26184 47.71957 0.5838515 3.166587 1.536337

Monthly Milk Production ARIMA Forecast



Fit SARIMA model for the Milk Production dataset

sarima_milk <- fit_sarima(milk_data, "Monthly Milk Production")</pre>

Fitting SARIMA model for $\,$ Monthly Milk Production

Series: ts_data

ARIMA(1,0,0)(2,1,2)[12] with drift

Coefficients:

ar1 sar1 sar2 sma1 sma2 drift
0.8638 0.0607 -0.4074 -1.0121 0.4831 2.1882
s.e. 0.0475 0.1862 0.1173 0.1994 0.1881 0.2174

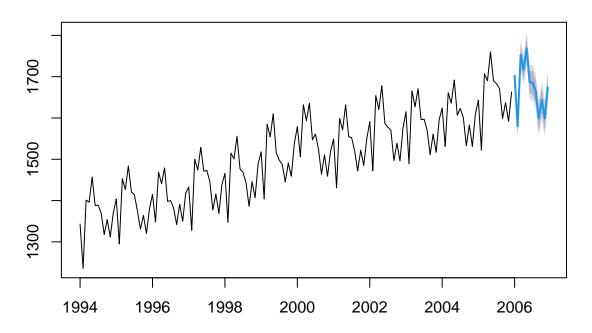
sigma^2 = 137.9: log likelihood = -518.84 AIC=1051.67 AICc=1052.57 BIC=1071.85

Training set error measures:

ME RMSE MAE MPE MAPE
Training set -0.1211196 10.98512 8.342375 -0.01115387 0.5520753

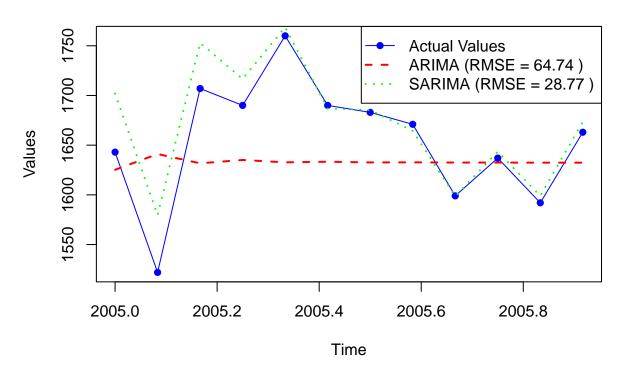
MASE ACF1

Monthly Milk Production SARIMA Forecast



```
# Compare ARIMA and SARIMA models based on their forecast accuracy
compare_models(arima_milk, sarima_milk, milk_data)
 Comparing ARIMA and SARIMA models :
 ARIMA Forecast Accuracy :
 21.88452 64.7399 54.04997 1.188661 3.264899 SARIMA Forecast Accuracy :
 -17.81697 28.77242 19.52245 -1.093511 1.195402
# Forecasting and plot comparison for Milk Production dataset
h_milk <- 12 # Define forecast horizon for Milk Production dataset (12 months ahead )
# Extract the actual values for the last 12 months of the Milk Production data
milk_actual_values <- milk_data[(length(milk_data) - h_milk + 1): length(milk_data)]
# Generate ARIMA forecast for the next 12 months
arima_milk_forecast <- forecast(arima_milk, h = h_milk)$mean</pre>
# Generate SARIMA forecast for the next 12 months
sarima_milk_forecast <- forecast(sarima_milk, h = h_milk)$mean</pre>
# Extract the time points for the last 12 months
time_points_milk <- time(milk_data)[(length(milk_data) - h_milk + 1): length(milk_data)]</pre>
# Plot and compare the forecasts from ARIMA and SARIMA models against the actual values
plot_forecast_comparison(milk_actual_values, arima_milk_forecast, sarima_milk_forecast,
                         time_points_milk)
```

Forecast Comparison



Program - 10

Interactive Visualization with plotly and Dynamic Reports with RMark-down

Date of Execution - 2025-10-28

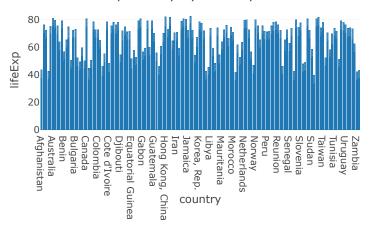
Objective - This program tests students' abilities to create interactive visualizations and generate dynamic reports using plotly and RMarkdown.

```
# Load necessary libraries
library(plotly)
library(gapminder)
library(dplyr)
# Load Gapminder dataset
data("gapminder")
# Scatter plot with plotly
# Scatter plot of GDP vs Life Expectancy by Continent
scatter_plot <- gapminder %>%
  plot_ly(x = ~gdpPercap, y = ~lifeExp, color = ~continent, size = ~pop,
          hoverinfo = 'text', text = ~paste("Country:", country, "<br/>br>GDP per Capita:", gdpPercap),
          type = 'scatter', mode = 'markers') %>%
  layout(title = 'GDP vs Life Expectancy by Continent',
         margin = list(1 = 20, r = 20, b = 20, t = 30)
  )
# Display the scatter plot
scatter_plot
Warning: 'line.width' does not currently support multiple values.
```

GDP vs Life Expectancy by Continent Africa 80 Americas Asia 70 Europe Oceania 60 lifeExp 50 40 30 20 50k 100k gdpPercap

```
# Bar chart with plotly
# Filter for year 2007 and create a bar chart of life expectancy by country
bar_chart <- gapminder %>%
filter(year == 2007) %>%
plot_ly(x = ~country, y = ~lifeExp, type = 'bar',
```

Life Expectancy by Country in 2007

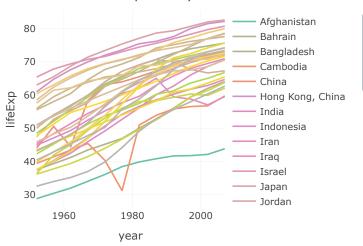


```
# Line chart with plotly
# Filter data for Asia and create a line chart showing life expectancy trends over time
line_chart <- gapminder %>%
filter(continent == 'Asia') %>%
plot_ly(x = ~year, y = ~lifeExp, color = ~country, type = 'scatter', mode = 'lines') %>%
```

Warning in RColorBrewer::brewer.pal(max(N, 3L), "Set2"): n too large, allowed maximum for palette Set2 Returning the palette you asked for with that many colors

Warning in RColorBrewer::brewer.pal(max(N, 3L), "Set2"): n too large, allowed maximum for palette Set2 Returning the palette you asked for with that many colors

Life Expectancy Trend in Asia



```
# Combine the Plots
# Combine the scatter, bar, and line charts into one interactive layout
dashboard <- subplot(scatter_plot, bar_chart, line_chart, nrows = 1) %>%
    layout(title = 'Gapminder Data Visualization',
        margin = list(1 = 20, r = 20, b = 20, t = 30)
```

)

Warning: 'line.width' does not currently support multiple values.

Warning in RColorBrewer::brewer.pal(max(N, 3L), "Set2"): n too large, allowed maximum for palette Set2

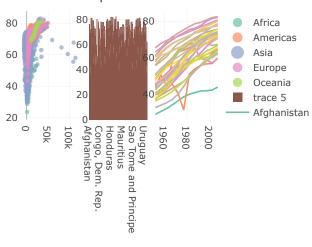
Returning the palette you asked for with that many colors

Warning in RColorBrewer::brewer.pal(max(N, 3L), "Set2"): n too large, allowed maximum for palette Set2 Returning the palette you asked for with that many colors

Display the dashboard

dashboard

Gapminder Data Visualization





B.M.S. College of Engineering

Dept. of CSE (Data Science)

Basavanagudi, Bangalore-19