

RGui

File Edit View Misc Packages Windows Help

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```
# Data for photograph preferences
age_groups <- c("5", "15", "25", "35", "45")
A <- c(18, 2, 20)
B <- c(22, 28, 10)
C <- c(20, 40, 40)

# Creating a data frame
data <- data.frame(A, B, C)

# (i) Calculate the sample covariance matrix for the preferences
cov_B_C <- cov(data[,2:3])
print(paste("Covariance Matrix between B and C:", cov_B_C))

# (ii) Calculate the sample covariance matrix for all preferences
cov_matrix <- cov(data)
print("Covariance Matrix:")
print(cov_matrix)

# (iii) Calculate the sample correlation matrix for the preferences
cor_B_C <- cor(data[,2:3])
print(paste("Correlation between B and C:", cor_B_C))

# (iv) Calculate the sample correlation matrix for all preferences
cor_matrix <- cor(data)
print("Correlation Matrix:")
print(cor_matrix)
```

R Console

```
> # (ii) Calculate the sample covariance matrix for the preferences
> cov_matrix <- cov(data)
> print("Covariance Matrix:")
[1] "Covariance Matrix:"
> print(cov_matrix)
      A      B      C
A 97.33333 -74 -46.66667
B -74.00000 84 -20.00000
C -46.66667 -20 133.33333

> # (iii) Calculate the sample correlation between B and C
> cor_B_C <- cor(data[,2:3])
> print(paste("Correlation between B and C:", cor_B_C))
[1] "Correlation between B and C: -0.188982236504614"

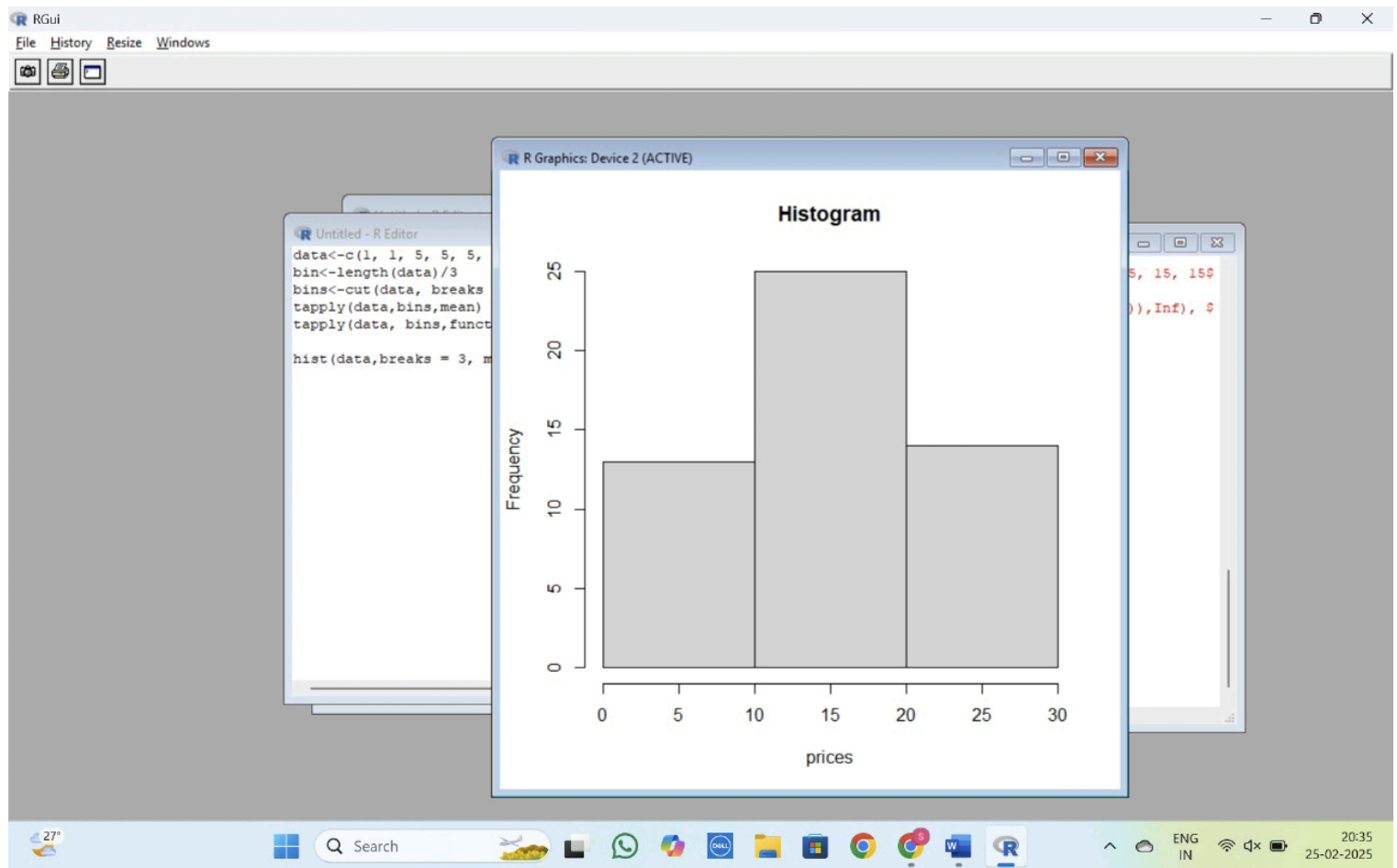
> # (iv) Calculate the sample correlation matrix for the preferences
> cor_matrix <- cor(data)
> print("Correlation Matrix:")
[1] "Correlation Matrix:"
> print(cor_matrix)
      A      B      C
A 1.0000000 -0.8183918 -0.4096440
B -0.8183918 1.0000000 -0.1889822
C -0.4096440 -0.1889822 1.0000000
> |
```

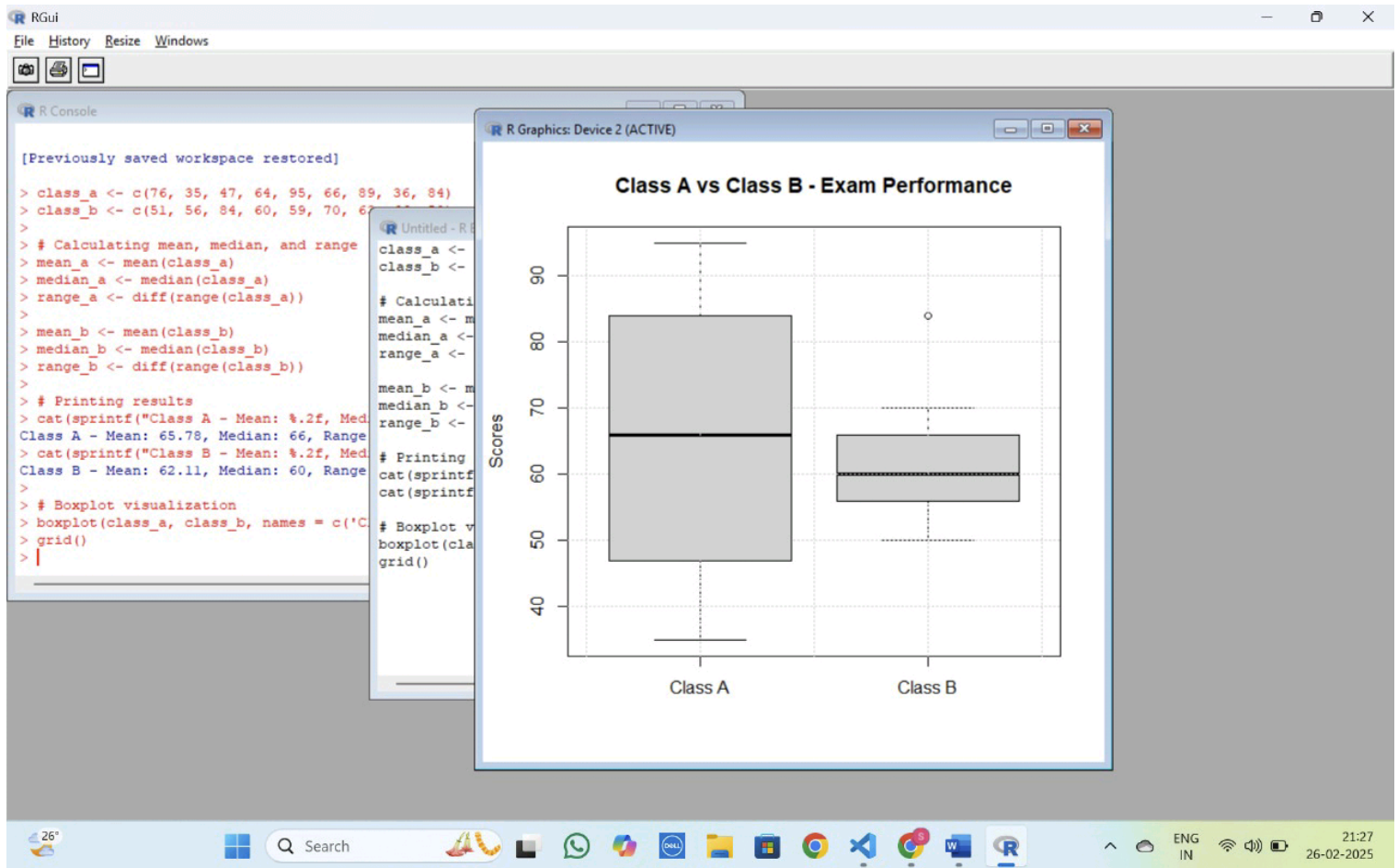
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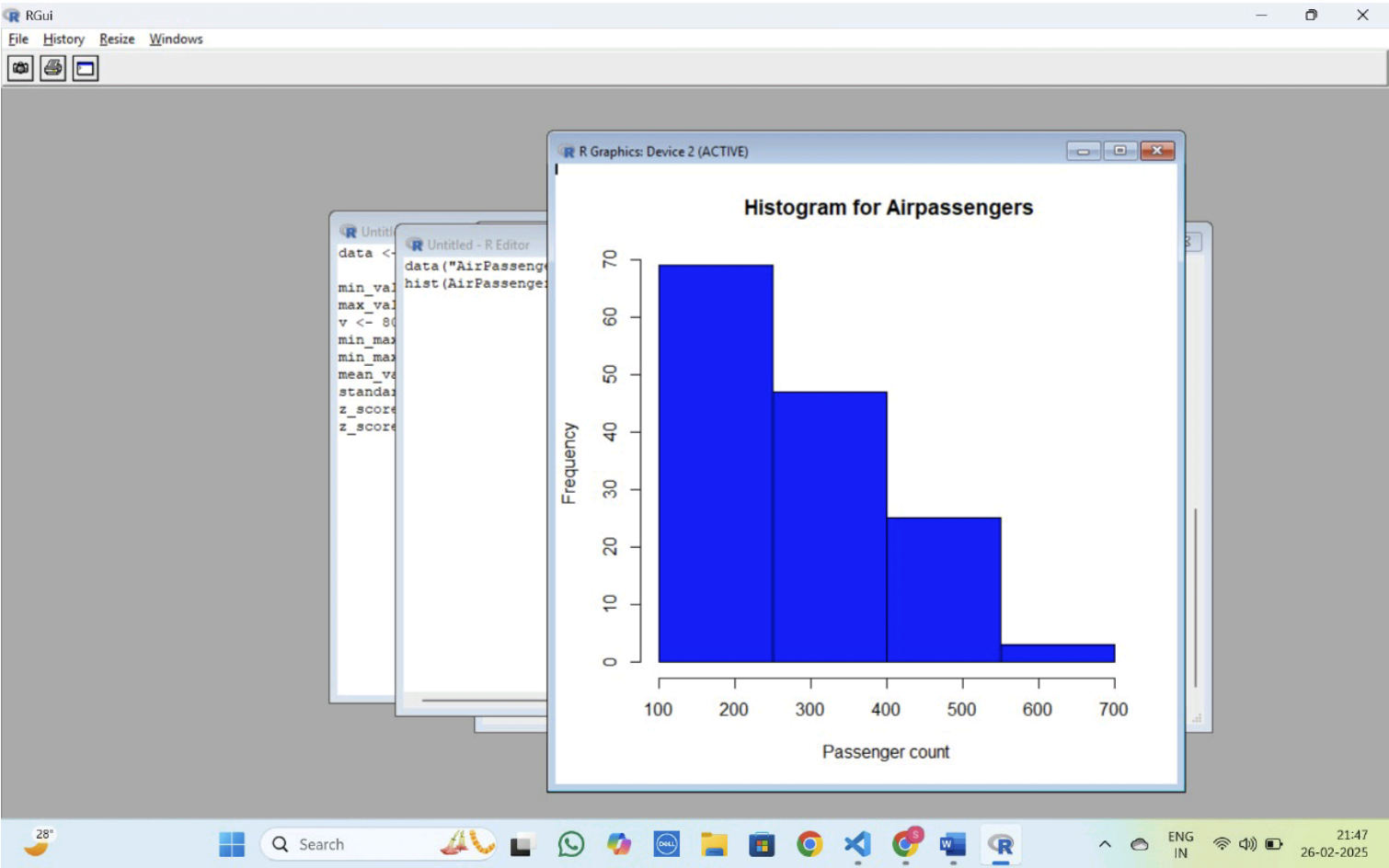
Untitled - R Editor

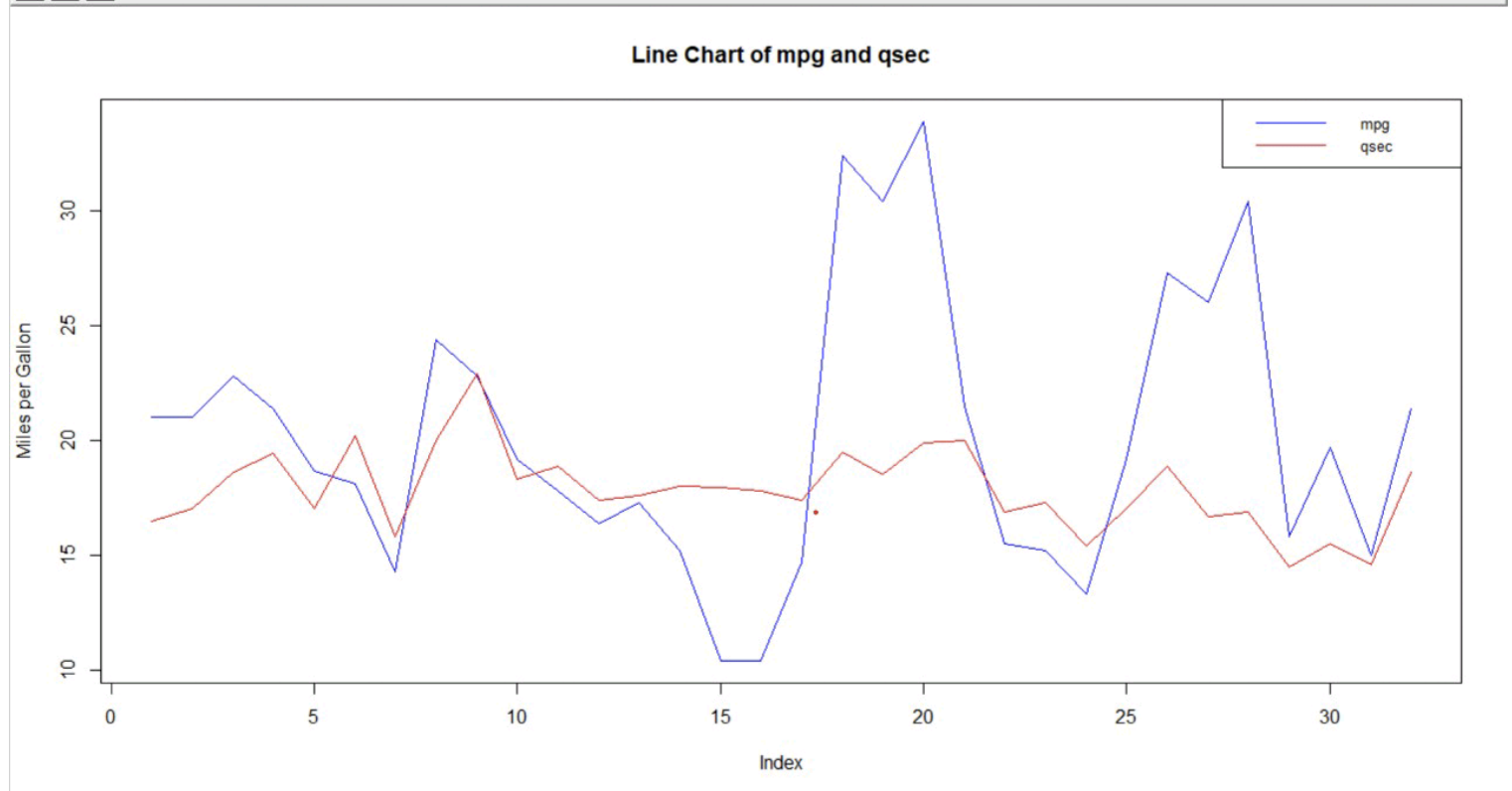
```
data <- c(200, 300, 400, 600, 1000)
min_value <- 50000
max_value <- 100000
v <- 80
min_max_normalized <- (v - min_value) / (max_value - min_value)
min_max_normalized
mean_value <- mean(data)
standard_deviation <- sd(data)
z_score_normalized <- (v - mean_value) / standard_deviation
z_score_normalized
```

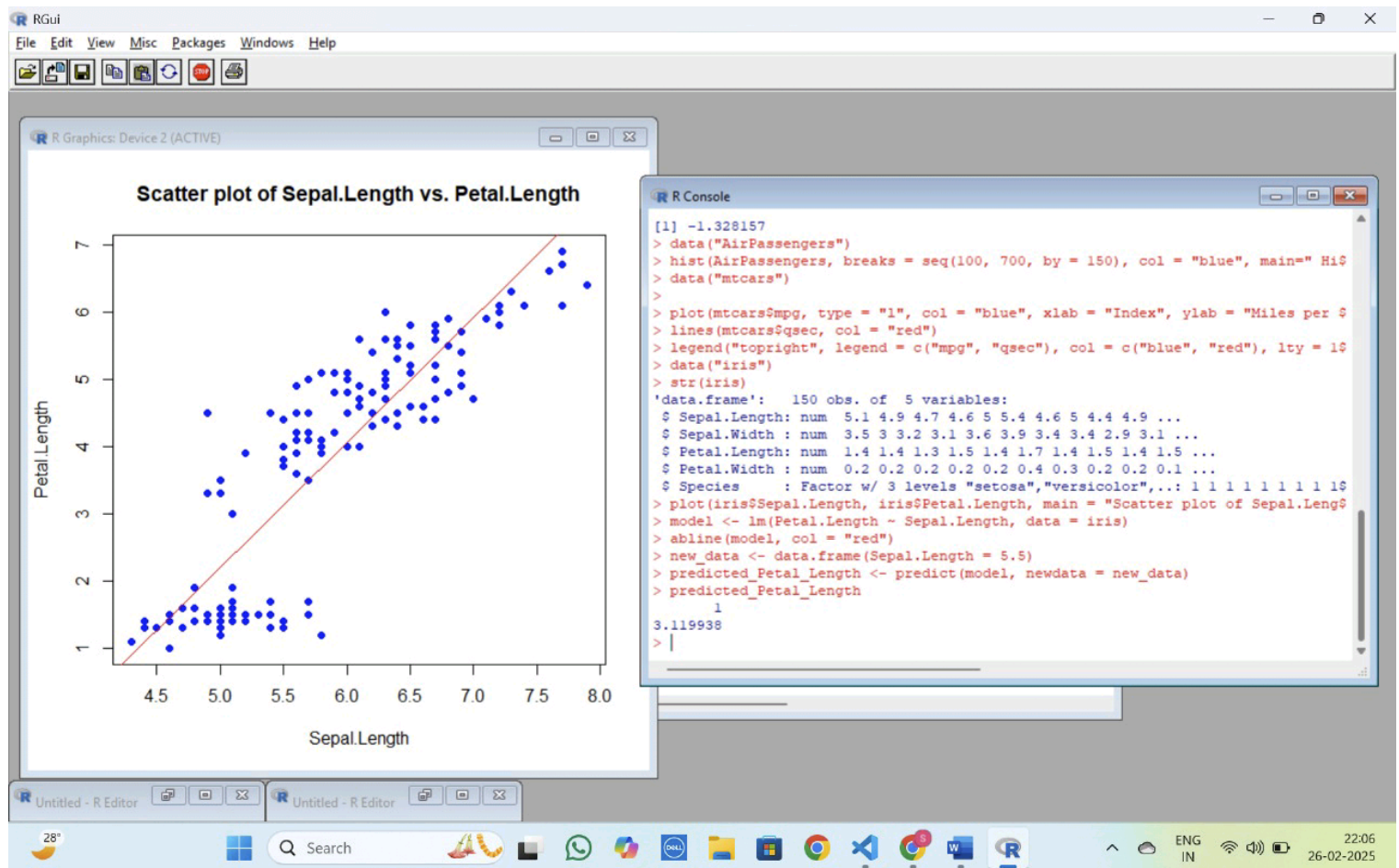
R Console

```
> range_b <- diff(range(class_b))
>
> # Printing results
> cat(sprintf("Class A - Mean: %.2f, Median: %d, Range: %d\n", mean_a, median_a, range_a))
Class A - Mean: 65.78, Median: 66, Range: 60
> cat(sprintf("Class B - Mean: %.2f, Median: %d, Range: %d\n", mean_b, median_b, range_b))
Class B - Mean: 62.11, Median: 60, Range: 34
>
> # Boxplot visualization
> boxplot(class_a, class_b, names = c('Class A', 'Class B'), main = 'Class A vs B')
> grid()
> data <- c(200, 300, 400, 600, 1000)
>
> min_value <- 50000
> max_value <- 100000
> v <- 80
> min_max_normalized <- (v - min_value) / (max_value - min_value)
> min_max_normalized
[1] -0.9984
> mean_value <- mean(data)
> standard_deviation <- sd(data)
> z_score_normalized <- (v - mean_value) / standard_deviation
> z_score_normalized
[1] -1.328157
>
```

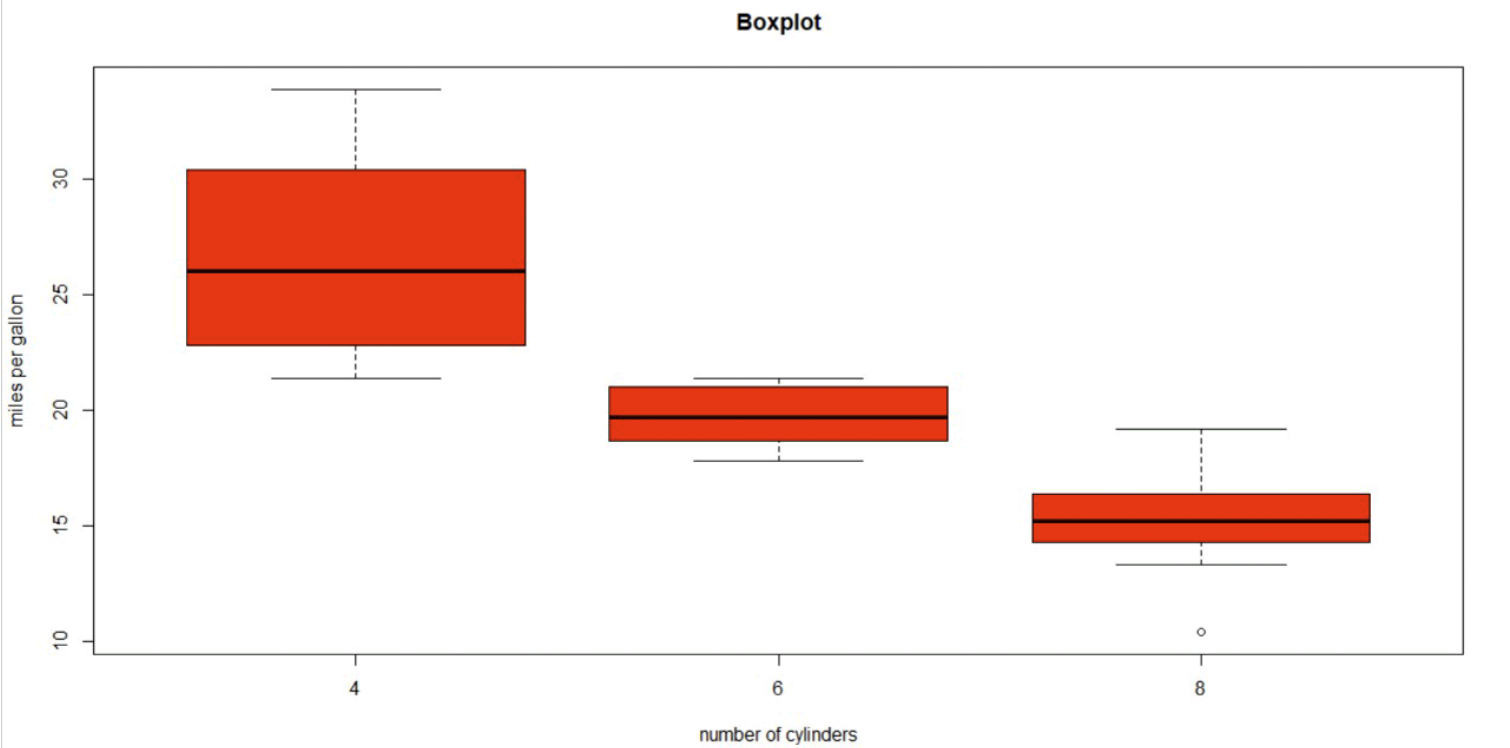
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```
RGui - [R Console]
File Edit View Misc Packages Windows Help

> min_value <- 50000
> max_value <- 100000
> v <- 80
> min_max_normalized <- (v - min_value) / (max_value - min_value)
> min_max_normalized
[1] -0.9984
> mean_value <- mean(data)
> standard_deviation <- sd(data)
> z_score_normalized <- (v - mean_value) / standard_deviation
> z_score_normalized
[1] -1.328157
> data("AirPassengers")
> hist(AirPassengers, breaks = seq(100, 700, by = 150), col = "blue", main="Histogram for Airpassengers", xlab = "Passenger count", ylab = "Frequency")
> data("mtcars")
>
> plot(mtcars$mpg, type = "l", col = "blue", xlab = "Index", ylab = "Miles per Gallon", main = "Line Chart of mpg and qsec")
> lines(mtcars$qsec, col = "red")
> legend("topright", legend = c("mpg", "qsec"), col = c("blue", "red"), lty = 1, cex = 0.8)
> data("iris")
> str(iris)
'data.frame': 150 obs. of 5 variables:
 $ Sepal.Length: num  5.1 4.9 4.7 4.6 5 5.4 4.6 5 4.4 4.9 ...
 $ Sepal.Width : num  3.5 3 3.2 3.1 3.6 3.9 3.4 3.4 2.9 3.1 ...
 $ Petal.Length: num  1.4 1.4 1.3 1.5 1.4 1.7 1.4 1.5 1.4 1.5 ...
 $ Petal.Width : num  0.2 0.2 0.2 0.2 0.2 0.4 0.3 0.2 0.2 0.1 ...
 $ Species      : Factor w/ 3 levels "setosa","versicolor",...: 1 1 1 1 1 1 1 1 1 1 ...
> plot(iris$Sepal.Length, iris$Petal.Length, main = "Scatter plot of Sepal.Length vs. Petal.Length", xlab = "Sepal.Length", ylab = "Petal.Length", col = "blue")
> model <- lm(Petal.Length ~ Sepal.Length, data = iris)
> abline(model, col = "red")
> new_data <- data.frame(Sepal.Length = 5.5)
> predicted_Petal_Length <- predict(model, newdata = new_data)
> predicted_Petal_Length
1
3.119938
> data("mtcars")
> boxplot(mpg ~ cyl, data = mtcars, main = "Boxplot", xlab = "number of cylinders", ylab = "miles per gallon", col = "red")
> score <- c(20, 25, 30, 32, 35, 38, 40, 45, 50, 52, 55, 56, 58, 59, 60, 62, 65, 70, 75, 80, 85)
>
> boxplot(score, col = "lightblue", main = "Box Plot of Points Scored by Tennis Players", ylab = "Points Scored")
> |
```