

R Project

2024-03-12

Task 1: Vector based functions.

```
# Set seed for reproducibility
set.seed(123)

# Generate vector vec4
vec4 <- sample(1:100, 100, replace = TRUE)

# Calculate the minimum value of vec4
min_value <- min(vec4)
cat("Minimum value:", min_value, "\n")
```

```
## Minimum value: 4
```

```
# Calculate the maximum value of vec4
max_value <- max(vec4)
cat("Maximum value:", max_value, "\n")
```

```
## Maximum value: 99
```

```
# Calculate the mean value of vec4
mean_value <- mean(vec4)
cat("Mean value:", mean_value, "\n")
```

```
## Mean value: 52.15
```

```
# Calculate the median value of vec4
median_value <- median(vec4)
cat("Median value:", median_value, "\n")
```

```
## Median value: 50
```

```
# Calculate the 25th quantile (1st quartile) of vec4
quantile_25 <- quantile(vec4, 0.25)
cat("25th quantile:", quantile_25, "\n")
```

```
## 25th quantile: 26.75
```

```
# Calculate the standard deviation of vec4
sd_value <- sd(vec4)
cat("Standard deviation:", sd_value, "\n")
```

```
## Standard deviation: 29.57592
```

```
# Calculate the variance of vec4
variance_value <- var(vec4)
cat("Variance:", variance_value, "\n")
```

```
## Variance: 874.7348
```

```
# Calculate the Inter Quartile Range (IQR) of vec4
iqr_value <- IQR(vec4)
cat("Inter Quartile Range (IQR):", iqr_value, "\n")
```

```
## Inter Quartile Range (IQR): 52.75
```

```
# Calculate the total range of vec4
total_range <- diff(range(vec4))
cat("Total range:", total_range, "\n")
```

```
## Total range: 95
```

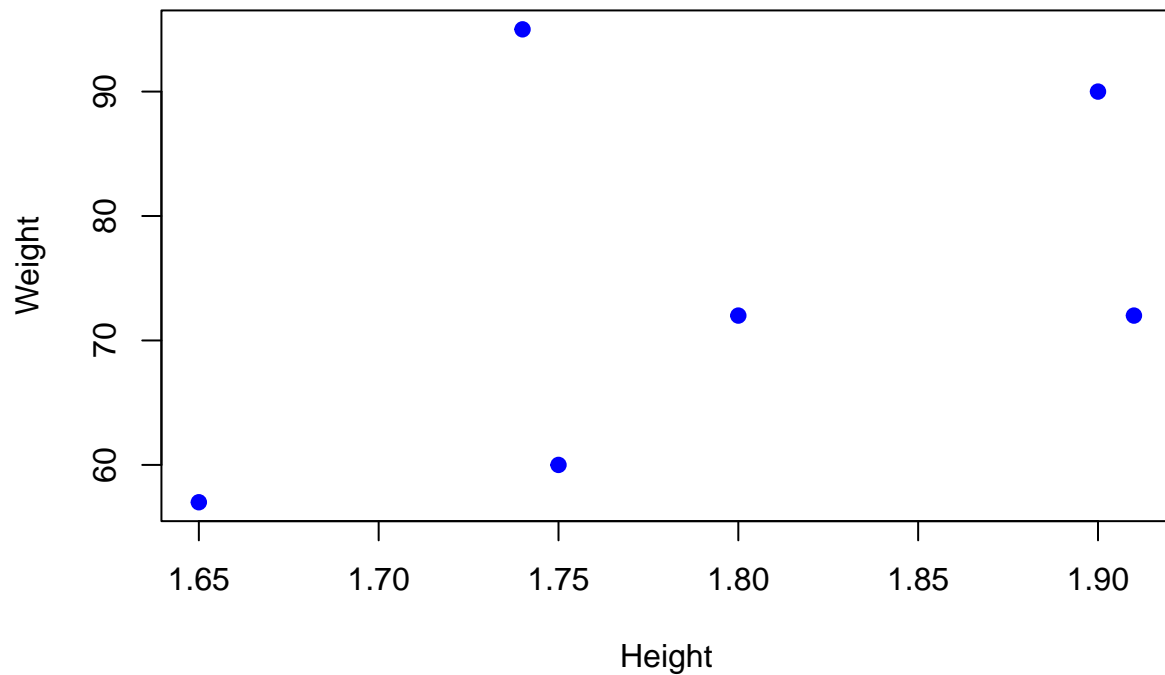
Task 2: Scatter plot

```
# Define the vectors
weight <- c(60, 72, 57, 90, 95, 72)
height <- c(1.75, 1.8, 1.65, 1.9, 1.74, 1.91)
gender <- c("m", "f", "m", "f", "f", "m")

# Convert gender to factor
gender <- factor(gender)

# Plot the scatterplot
plot(height, weight, col = "blue", pch = 19,
      xlab = "Height", ylab = "Weight",
      main = "Scatterplot of Height vs. Weight")
```

Scatterplot of Height vs. Weight



Task 3: Creating a Dataframe and plot

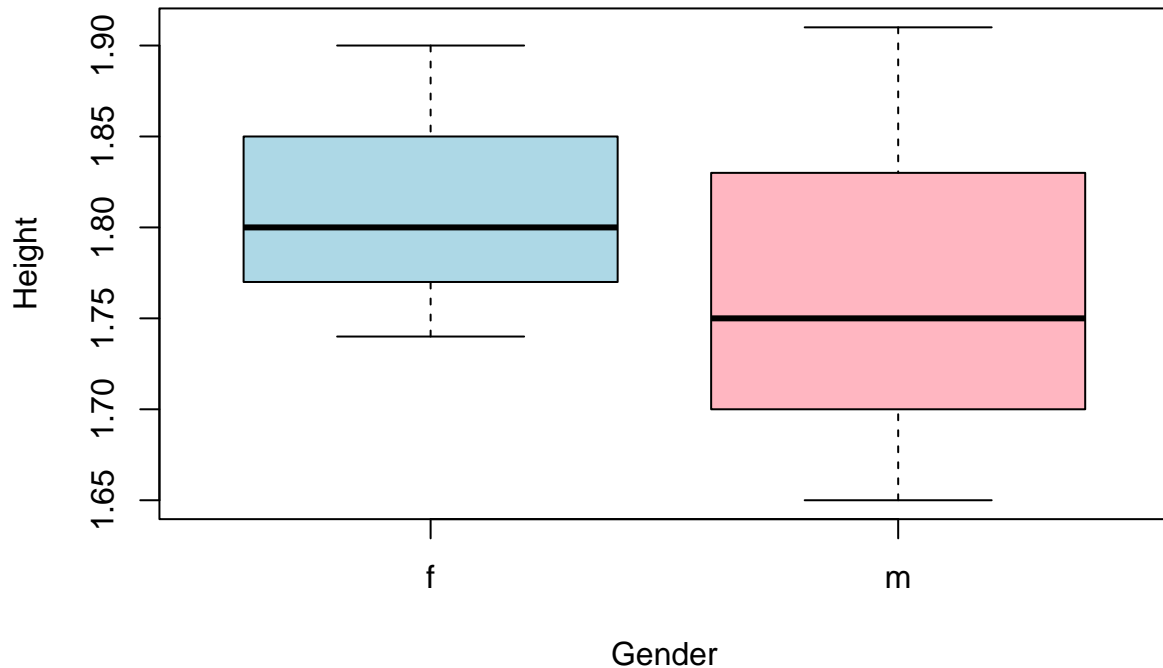
```
# Define the vectors
weight <- c(60, 72, 57, 90, 95, 72)
height <- c(1.75, 1.8, 1.65, 1.9, 1.74, 1.91)
gender <- c("m", "f", "m", "f", "f", "m")

# Convert gender to factor
gender <- factor(gender)

# Create a dataframe
ghw <- data.frame(gender, height, weight)

# Plot a boxplot with height on the y-axis and gender on the x-axis
boxplot(height ~ gender, data = ghw,
        col = c("lightblue", "lightpink"),
        main = "Boxplot of Height by Gender",
        xlab = "Gender", ylab = "Height")
```

Boxplot of Height by Gender



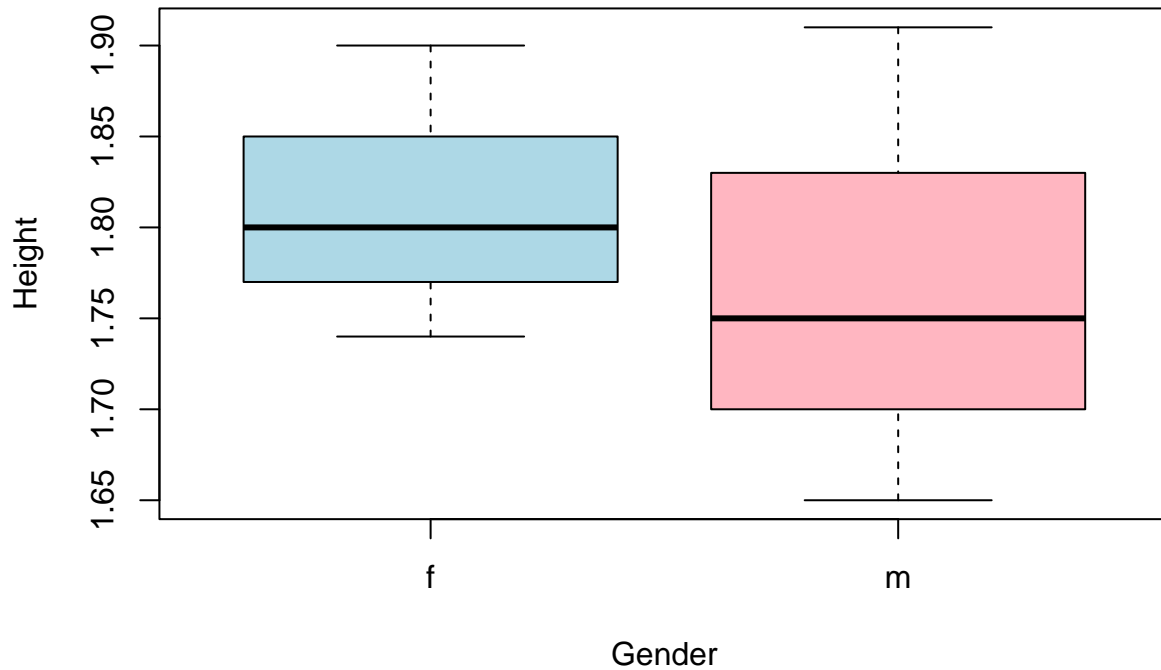
Task 4: Create a dataframe and a plot

```
# Create a dataframe
ghw <- data.frame(
  gender = c("m", "f", "m", "f", "f", "m"),
  height = c(1.75, 1.80, 1.65, 1.90, 1.74, 1.91),
  weight = c(60, 72, 57, 90, 95, 72)
)

# Convert gender to a factor
ghw$gender <- factor(ghw$gender)

# Plot a boxplot with height on the y-axis and gender on the x-axis
boxplot(height ~ gender, data = ghw,
  col = c("lightblue", "lightpink"),
  main = "Boxplot of Height by Gender",
  xlab = "Gender", ylab = "Height")
```

Boxplot of Height by Gender



Task 5: Calculate Corpulence Index(CI)

```
#task 5
# Calculate the Corpulence Index (CI)
ghw$ci <- ghw$weight / (ghw$height)^3

# Display the updated dataframe
ghw
```

```
##   gender height weight      ci
## 1      m   1.75     60 11.19534
## 2      f   1.80     72 12.34568
## 3      m   1.65     57 12.68887
## 4      f   1.90     90 13.12145
## 5      f   1.74     95 18.03333
## 6      m   1.91     72 10.33314
```

Task 6: Calculate the quantiles and plot.

```
# task 6

# g2
g2 <- c(77, 75, 78, 41, 51, 20, 61, 73, 76, 38)

# Calculate quantiles for g2
```

```
quantile_g2 <- quantile(g2)
print(quantile_g2)
```

```
##      0%   25%   50%   75%  100%
## 20.00 43.50 67.00 75.75 78.00
```

```
# Plot a boxplot for g2
boxplot(g2, main = "Boxplot of g2", ylab = "Values")
```

