BMI and Glycemia Statistical Analysis

Statisticians World

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Importing the Data

In this section, we load a dataset containing patient BMI and glycemia levels.

```
# Set the working directory (modify this path if necessary)
setwd("C:/Users/0&1/OneDrive/Documents/R-Youtube")

# Load the dataset
my_data <- read.csv("BMI1.csv")

# Preview the data
head(my_data)</pre>
```

```
##
    name gender glycemia BMI
## 1
                     98 24.1
       1
             1
              2
## 2
                    117 25.0
## 3
       3
             1
                     98 27.6
## 4
       4
              1
                     85 23.2
       5
## 5
              2
                     139 28.2
## 6
                     147 26.4
```

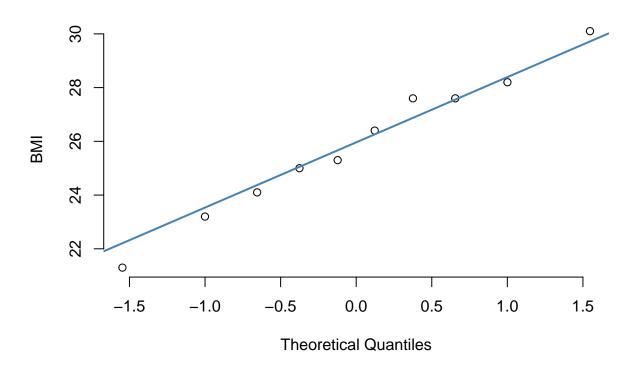
Assessing Normality

We use Q-Q plots to visually check the normality of BMI and glycemia. A straight-line pattern indicates approximate normality.

Q-Q Plot for BMI

```
qqnorm(my_data$BMI, ylab = "BMI", pch = 1, frame = FALSE)
qqline(my_data$BMI, col = "steelblue", lwd = 2)
```

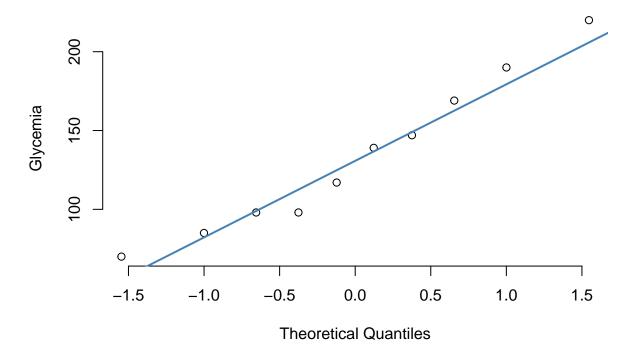
Normal Q-Q Plot



Q-Q Plot for Glycemia

```
qqnorm(my_data$glycemia, ylab = "Glycemia", pch = 1, frame = FALSE)
qqline(my_data$glycemia, col = "steelblue", lwd = 2)
```

Normal Q-Q Plot

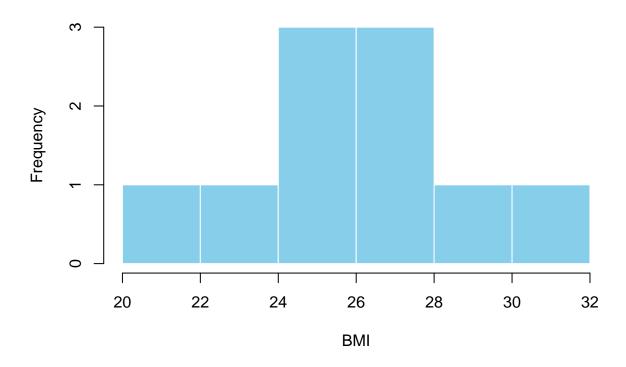


Exploring the BMI Distribution

We use a histogram to examine the shape of the BMI distribution.

```
hist(my_data$BMI,
    main = "Histogram of BMI",
    xlab = "BMI",
    ylab = "Frequency",
    col = "skyblue",
    border = "white")
```



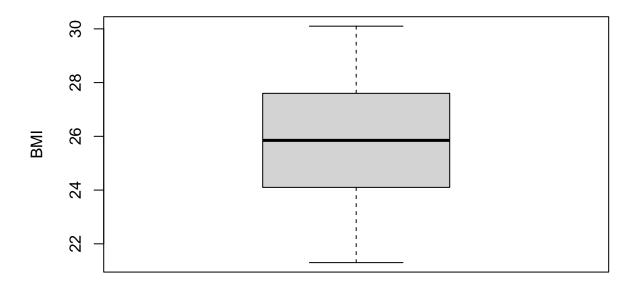


Detecting Outliers

A boxplot is useful for identifying potential outliers in BMI values.

```
boxplot(my_data$BMI,
    ylab = "BMI",
    main = "Boxplot of BMI",
    col = "lightgray",
    border = "black")
```

Boxplot of BMI



One-Sample T-Test

We now perform a one-sample t-test to determine if the mean BMI significantly differs from a known population mean of 28.4.

Hypotheses

Null hypothesis (H): The mean BMI = 28.4

Alternative hypothesis (H): The mean BMI 28.4

Perform the T-Test

Set population mean

```
pop_mean <- 28.4
```

Perform the test

```
result <- t.test(my_data$BMI, mu = pop_mean)</pre>
```

Output result

result

5.3 Interpretation If the p-value is less than 0.05, we reject the null hypothesis, concluding that the mean BMI differs significantly from 28.4.

Summary

We assessed normality of BMI and glycemia using Q-Q plots.

BMI distribution was visualized with a histogram and boxplot.

A one-sample t-test was conducted to compare the sample BMI to a population mean of 28.4.

Statistical results support inference about differences in the BMI distribution.

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