

ESE 326: Final Project

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1 Introduction

The main objective of this project involved R's built-in dataset, Iris. Through graphical exploration and mathematical analysis, the researchers determine whether there are clear rules as to which of the features determine the species of a given specimen.

2 Methods

2.1 Exploratory Analysis

2.2 Confidence Interval Estimate

2.3 Hypothesis Test

$\sigma_1 = \sigma_2$ T-test using S_p :

$$\frac{(\bar{X}_1 - \bar{X}_2) - (\mu_1 - \mu_2)}{\sqrt{S_p^2 \left(\frac{1}{n_1} + \frac{1}{n_2} \right)}} \sim T_{n_1+n_2-2} \quad (1)$$

$$S_p^2 = \frac{(n_1 - 1)S_1^2 + (n_2 - 1)S_2^2}{n_1 + n_2 - 2} \quad (2)$$

$\sigma_1 \neq \sigma_2$ T-test using γ :

$$\frac{(\bar{X}_1 - \bar{X}_2) - (\mu_1 - \mu_2)}{\sqrt{\frac{S_1^2}{n_1} + \frac{S_2^2}{n_2}}} \sim T_\gamma \quad (3)$$

$$\gamma = \frac{\left(\frac{S_1^2}{n_1} + \frac{S_2^2}{n_2} \right)^2}{\frac{\left(\frac{S_1^2}{n_1} \right)^2}{n_1 - 1} + \frac{\left(\frac{S_2^2}{n_2} \right)^2}{n_2 - 1}} \quad (4)$$

3 Results and Observations

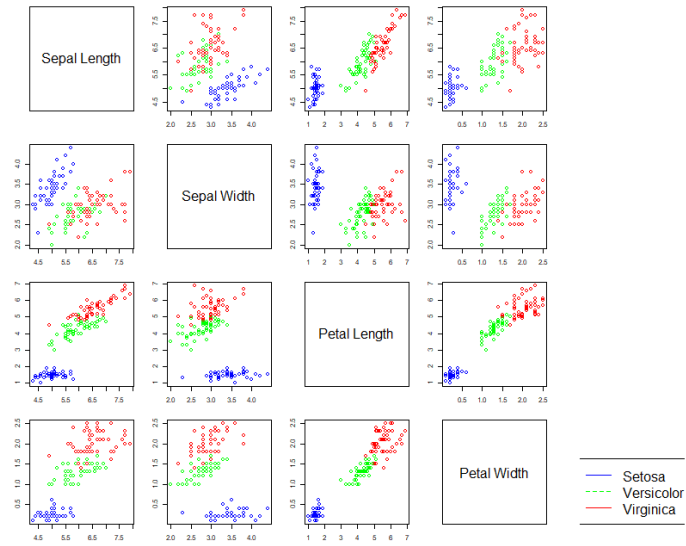


Figure 1: A visualization of the Iris dataset showing scatterplots of each pair of the features.

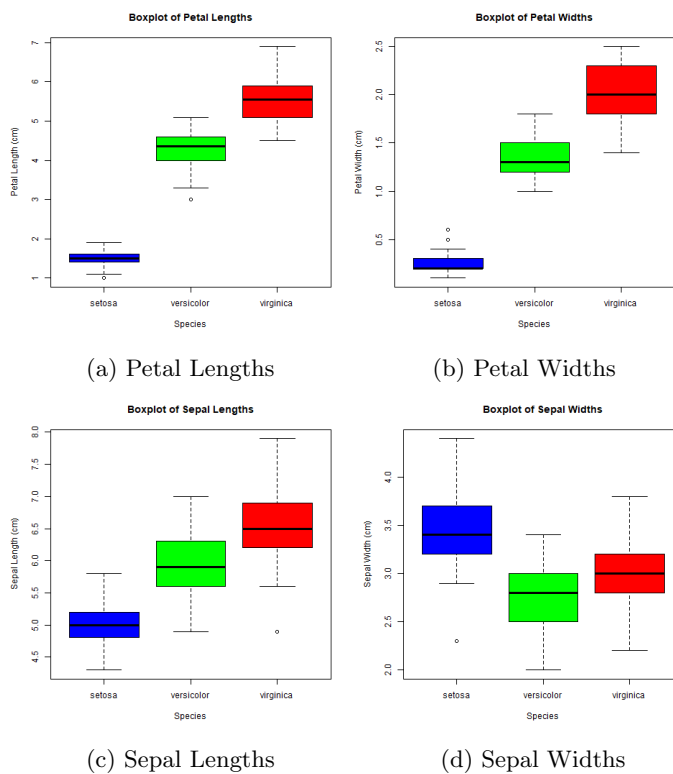


Figure 2: Boxplots for each of the four features.

Figure 2a is petal length

4 Conclusions

5 Appendix

5.1 R-scripts

5.2 Extra Figures and Tables