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13 Introduction

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Materials and Methods

36 Data

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37 Analyses

We used R for all analyses (R Core Team, 2018).

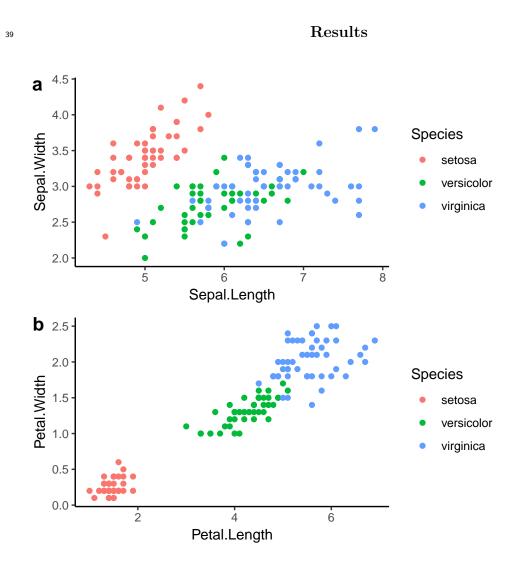


Figure 1. Sepals and petals in three *Iris* species. (a) Relationship between sepal length and width. (b) Relationship between petal length and width.

Table 1

Mean values for floral organ traits in three Iris species.

Species	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
setosa	5.006	3.428	1.462	0.246
versicolor	5.936	2.770	4.260	1.326
virginica	6.588	2.974	5.552	2.026

- Length and width of sepals were not correlated (r = -0.12, p = 0.15, Figure 1a). In contrast, length and width of petals showed a strong positive correlation (r = 0.96, p = 0.00, figure 1b).
- $_{43}$ The mean species values for each trait are shown in table 1

44 Discussion

45 References

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Supplement

2 Software used

71

- We used R (Version 3.5.1; R Core Team, 2018) and the R-packages cowplot (Version
- ⁷⁴ 0.9.3; Wilke, 2018), *dplyr* (Version 0.7.8; Wickham et al., 2018a), *forcats* (Version 0.3.0;
- Wickham, 2018a), ggplot2 (Version 3.1.0; Wickham, 2016), papaja (Version 0.1.0.9842; Aust
- ⁷⁶ & Barth, 2018), purrr (Version 0.2.5; Henry & Wickham, 2018), readr (Version 1.2.1;
- Wickham et al., 2018b), stringr (Version 1.3.1; Wickham, 2018b), tibble (Version 1.4.2;
- Müller & Wickham, 2018), tidyr (Version 0.8.2; Wickham & Henry, 2018), and tidyverse
- 79 (Version 1.2.1; Wickham, 2017) for all our analyses.