How Phylogeny and traits relate to heterospecific pollen effect

Jose B. Lanuza ^{1,2} true

⁴
⁵
¹US Environmental Protection Agency, Office of Research and Development, National Health and Environmental Effects
6 Research Laboratory, Atlantic Ecology Division, 27 Tarzwell Drive Narragansett, RI, 02882, USA

⁷
⁸
²Big Name University, Department of R, City, BN, 01020, USA

⁹
⁹
¹⁰
³Estacion Biologica de Donana (EBD-CSIC), E-41092 Sevilla, Spain

11
^{*}
^{*}
corresponding author: f.rodriguez.sanc@gmail.com

3 Write your abstract here.

4 Keywords: rmarkdown, reproducible science

INTRODUCTION

16 METHODS

- 17 The study was conducted in a glasshouse at University of New England (Armidale, Australia) from
- November 2017 to March 2018. Rooms were temperature controlled depending on the requirements of
- 19 the species with day and night temperature differences. The species selected (Table 1) belonged to
- three different families, Solanaceae, Brassicaceae and Convolvulaceae. The criteria of species/family
- selection was based on close/distant related species (see phylogenetic tree for relatedness fig 1),
- 22 heterogeneous traits, low structural flower complexity and fast life cycle. For the purpose of the
- 23 experiment all the species where considered as pollen recipient and as pollen donor (see interaction
- 24 matrix, fig 2). Species were watered once or twice per day and fertilized weekly (NPK 23: 3.95: 14).

Hand-pollination

- 26 Foreign pollen effect was studied through two different treatments, one with 50% conspecific pollen and
- 50% heterospecific pollen and a second one with 100% foreign pollen (N=10). Moreover, hand cross

- 28 pollination, hand self pollination, apomixis (bagged emasculated flowers) and natural selfing were
- 29 tested (N=10). Flowers were emasculated the day prior anthesis and hand pollinated next day with a
- 30 toothpick. Had-pollination was realized with 3-4 gentle touches on the surface of the stigma. The mixes
- of pollen were performed based on the pollen counts (Neubaeur chamber) on an eppendorf.
- We used the statistical language R (R Core Team 2018) for all our analyses. These were implemented in
- dynamic rmarkdown documents using knitr (Xie 2014, 2015, 2018) and rmarkdown (Allaire et al.
- ³⁴ 2018) packages. All the multilevel models were fitted with lme4 (Bates et al. 2015).

85 RESULTS

- Trees in forest A grew taller than those in forest B (mean height: 25 versus 13 m). And many more
- 37 cool results that get updated dynamically.

38 DISCUSSION

Discuss.

40 CONCLUSIONS

41 ACKNOWLEDGEMENTS

42 REFERENCES

- ⁴³ Allaire, J., Y. Xie, J. McPherson, J. Luraschi, K. Ushey, A. Atkins, H. Wickham, J. Cheng, and W.
- 44 Chang. 2018. Rmarkdown: Dynamic documents for r.
- ⁴⁵ Bates, D., M. Mächler, B. Bolker, and S. Walker. 2015. Fitting linear mixed-effects models using lme4.

- 46 Journal of Statistical Software 67:1–48.
- 47 R Core Team. 2018. R: A language and environment for statistical computing. R Foundation for
- 48 Statistical Computing, Vienna, Austria.
- ⁴⁹ Xie, Y. 2014. Knitr: A comprehensive tool for reproducible research in R. in V. Stodden, F. Leisch,
- 50 and R. D. Peng, editors. Implementing reproducible computational research. Chapman; Hall/CRC.
- 51 Xie, Y. 2015. Dynamic documents with R and knitr. 2nd editions. Chapman; Hall/CRC, Boca Raton,
- 52 Florida.
- Xie, Y. 2018. Knitr: A general-purpose package for dynamic report generation in r.

List of Tables

55 List of Figures