

How phylogenetic relatedness and floral traits are involved in heterospecific pollen effect in an artificial co-flowering community

Jose B. Lanuza ^{1,2} true

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

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

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

* corresponding author: barragansljose@gmail.com

Possible journals to publish: New phytologist, journal of ecology, oikos. . .

Keywords: heterospecific pollen, plant reproduction, fitness, competition

INTRODUCTION

Paragraph 1

In natural systems plant species normally coexist and share their floral visitors with other species (Bascompte et al., 2003). This pollinator sharing from the plant perspective can be negative due to competition (refs) or positive due to facilitation (refs). Moreover, once the pollinator has landed on the stigma some other issues for the species fitness may arise, the arrival of foreign pollen and conspecific pollen loss.

Paragraph 2

The effect of heterospecific pollen has been widely studied (refs). Native species are supposed to have greater negative effects than native ones Arceo-Gómez and Ashman (2016)

species-specific. What kind of pollen interaction? Size just mechanical?

Paragraph 3

28 Relatedness. What we know?

29 **Paragraph 4**

30 The great difficulty of working with pollen in a coflowering community make the understanding of
31 heterospecific pollen effect a real challenge. For this reason we have created an artificial co-flowering
32 community in a glasshouse to test the effect with all the possible combinations among them. Where we
33 test the following hypothesis: 1) Does heterospecific pollen reduce seed set, if so, 2) Does heterospecific
34 pollen effect depend on the relatedness of the species, 3) Does heterospecific pollen effect depend on any
35 floral trait?

36 **METHODS**

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

45 **Hand-pollination**

46 Foreign pollen effect was studied through two different treatments, one with 50% conspecific pollen and
47 50% heterospecific pollen and a second one with 100% foreign pollen (N=10). The proxy of effect were
48 seed set and “pollen tubes”. Moreover, hand cross pollination, hand self pollination, apomixis (bagged
49 emasculated flowers) and natural selfing were tested (N=10). Flowers were emasculated the day prior
50 anthesis and hand pollinated next day with a toothpick. Hand-pollination was realized with 3-4 gentle
51 touches on the surface of the stigma. The mixes of pollen were performed on an eppendorf based on the
52 pollen counts made with Neubauer chamber (each anther was counted 4 times for 20 different anthers)

53 per species).

54 **Evolutionary distance**

55 Two types of evolutionary distances were calculated with MEGA7 using kinds of markers: 1) Internal
56 transcribed spacer (ITS) and 2) ribulose-bisphosphate carboxylase (RBCL)

57 **Traits**

58 Several traits of the ten species were measured. Pollen per anther was counted, number of ovules,
59 stigma width and length and stigmatic area, style width and length, ovary width and length. Moreover
60 stigma type was tested. Self-incompatibility was

61 We used the statistical language R (R Core Team 2018) for all our analyses. These were implemented in
62 dynamic markdown documents using `knitr` (Xie 2014, 2015, 2018) and `rmarkdown` (Allaire et al.
63 2018) packages. All the multilevel models were fitted with `lme4` (Bates et al. 2015).

64 **RESULTS**

65 **DISCUSSION**

66 Discuss.

67 CONCLUSIONS

68 ACKNOWLEDGEMENTS

69 REFERENCES

- 70 Allaire, J., Y. Xie, J. McPherson, J. Luraschi, K. Ushey, A. Atkins, H. Wickham, J. Cheng, and W.
71 Chang. 2018. Rmarkdown: Dynamic documents for r.
- 72 Arceo-Gómez, G., and T.-L. Ashman. 2016. Invasion status and phylogenetic relatedness predict cost
73 of heterospecific pollen receipt: Implications for native biodiversity decline. *Journal of Ecology*
74 104:1003–1008.
- 75 Bates, D., M. Mächler, B. Bolker, and S. Walker. 2015. Fitting linear mixed-effects models using lme4.
76 *Journal of Statistical Software* 67:1–48.
- 77 R Core Team. 2018. R: A language and environment for statistical computing. R Foundation for
78 Statistical Computing, Vienna, Austria.
- 79 Xie, Y. 2014. Knitr: A comprehensive tool for reproducible research in R. *in* V. Stodden, F. Leisch,
80 and R. D. Peng, editors. *Implementing reproducible computational research*. Chapman; Hall/CRC.
- 81 Xie, Y. 2015. *Dynamic documents with R and knitr*. 2nd editions. Chapman; Hall/CRC, Boca Raton,
82 Florida.
- 83 Xie, Y. 2018. Knitr: A general-purpose package for dynamic report generation in r.

