

1 Arceo-Gómez et al. Patterns and correlates of variation in heterospecific pollen receipt

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3 **Patterns of among- and within-species variation in heterospecific pollen receipt: the**  
4 **importance of ecological generalization<sup>1</sup>**

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- 32 • *Premise of the study:* Co-flowering plants are at risk of receiving pollen from  
33 heterospecifics as well as conspecifics, yet evidence shows wide variation in the degree  
34 that this occurs. Evaluation of patterns and correlates of among- and within-species  
35 variation in heterospecific pollen (HP) receipt is key to understanding its importance for  
36 floral evolution and species coexistence; however, the rarity of deeply-sampled  
37 multispecies comparisons has precluded such an evaluation.
- 38 • *Methods:* We evaluated patterns of among- and within-species variation in HP load size  
39 and diversity in 19 species across three distinct plant communities. We assessed the  
40 importance of phenotypic specialization (floral phenotype), ecological specialization  
41 (contemporary visitor assemblage) and conspecific flower density as determinants of  
42 among-species variation. We present hypotheses for different accrual patterns of HP  
43 within species based on the evenness and quality of floral visitors, and evaluated these by  
44 characterizing the relationship between conspecific pollen (CP) and HP receipt.
- 45 • *Key results:* We found that within-species variation in HP receipt was greater than  
46 among-species and among-communities variation. Among species, ecological  
47 generalization emerged as the strongest driver of variation in HP receipt irrespective of  
48 phenotypic specialization. Within-species variation in HP load size and diversity was  
49 predicted most often from two CP-HP relationships (linear or exponentially decreasing)  
50 suggesting that two distinct types of plant-pollinator interactions prevail.
- 51 • *Conclusions:* Our results give important insights into the potential drivers of among- and  
52 within-species variation in HP receipt. They also highlight the value of explorations of

53 patterns at the intraspecific level which can ultimately shed light on plant-pollinator-  
54 mediated selection in diverse communities.

55 **Keywords:** co-flowering community, diversity, ecological generalization, floral specialization,  
56 heterospecific pollen, floral visitor assemblage, pollinator sharing

57

58           The movement of pollinators between plants in multi-species plant communities can  
59 result in the transfer of pollen from heterospecifics as well as conspecifics (e.g., Feinsinger et al.,  
60 1986; Montgomery and Rathcke, 2012; Fang and Huang, 2013). However, the likelihood of  
61 heterospecific pollen (hereafter, HP) receipt varies widely among species. For instance, a review  
62 of published studies (Ashman and Arceo-Gómez, 2013) revealed variation among species in both  
63 the percentage of flowers that received any HP (2-100%), as well as in the average intensity of  
64 HP receipt (% HP in total stigmatic pollen load: 0.1% to 80%). Furthermore, Fang and Huang  
65 (2013) uncovered variation among species within a single community not only in HP load size  
66 but also in the number and identity of HP donors. Some species received large multispecies loads  
67 while others received small monospecific loads of HP (Fang and Huang, 2013). Variation in HP  
68 receipt among individuals within a single species can also be extensive (e.g., 1-95% of total load  
69 in *Mimulus guttatus*; Arceo-Gómez and Ashman, unpublished data), but variation at this level  
70 has received much less attention. In fact, a systematic comparison of all aspects of HP receipt  
71 (e.g., mean and variance of HP load size and diversity) among and within species has not been  
72 attempted in any community. Characterization of the pattern of variation at inter- and intra-  
73 specific levels as well as identifying the factors that contribute to variation at both levels are  
74 crucial for gaining a complete understanding of the role of HP transfer in floral evolution and  
75 species coexistence in diverse plant communities.

76           Phenotypic specialization is thought to reflect past selection to maximize conspecific  
77 pollen (hereafter, CP) receipt and export while minimizing HP transfer (Rathcke, 1983; Waser,  
78 1983; Armbruster, 1995; Waser et al., 1996; Muchhala and Potts, 2007; Pauw, 2013). Flower  
79 shape (e.g., symmetry), in particular, can restrict visitors by body size, tongue length and/or  
80 learning capabilities leading to predictions of lower HP receipt by zygomorphic flowers than

81 actinomorphic ones (Faegri and van der Pijl, 1979; Feinsinger et al., 1986; Waser, 1986;  
82 McLernon et al., 1996). Irrespective of floral symmetry, however, species with large flowers,  
83 long life times and/or large exerted stigmas are accessible to a wide array of pollinators (see  
84 below) and thus are expected to receive large and possibly diverse loads of HP (Montgomery and  
85 Rathcke, 2012; Ashman and Arceo-Gómez, 2013; Fang and Huang, 2013; Huang and Shi, 2013).  
86 In addition, species with such floral features (i.e., actinomorphic and large, long-lived flowers)  
87 would be predicted to also exhibit less variation among plants, as every flower would receive at  
88 least some HP over their life time. In contrast, species with more restrictive floral phenotypes  
89 and small, short-lived flowers are expected to receive, on average, smaller and less diverse HP  
90 loads, but may have higher variance among conspecifics as HP delivery is predicted to be more  
91 stochastic. These predictions for HP load size among species have been partially borne out  
92 (McLernon et al., 1996; Eaton et al., 2012; Montgomery and Rathcke, 2012; Fang and Huang,  
93 2013; Huang and Shi, 2013), but those for variance among and within species have yet to be  
94 explored.

95         Separate from the evolutionary state of specialization reflected in floral phenotype  
96 addressed above, the ecological state of specialization, that is, the contemporary community of  
97 floral visitors (*sensu* Feinsinger et al., 1986; Waser et al., 1996; Fenster et al., 2004) is expected  
98 to be an important determinant of the among-species variation in amount and diversity of HP  
99 deposited on stigmas. From this perspective, a plant species can range from being highly  
100 specialized, i.e., visited by only a single taxon (Pellmyr et al., 1996; Flemming and Holland,  
101 1998; Weiblen, 2004) to widely generalized, for example, be visited by up to 70 different taxa  
102 (Waser et al., 1996; Olesen and Jordano, 2002; Vazquez and Aizen, 2003; Alarcón et al., 2008;  
103 Bosch et al., 2009). As a consequence, plant species that host a wider array of visitors (that may

104 also visit other plant species) are predicted to receive larger, more diverse loads of HP with little  
105 variance among flowers than species with less diverse pollinator assemblages, all else being  
106 equal (e.g., pollen placement). This relationship between HP receipt and pollinator diversity has  
107 been explored indirectly via the presumed association between ecological specialization and  
108 phenotypic specialization (e.g., generalized pollination-actinomorphic flowers vs. specialized  
109 pollination-zygomorphic flowers; McLernon et al., 1996). A direct test of the effect of ecological  
110 specialization on HP receipt, however, requires measures of the contemporary community of  
111 floral visitors. Simultaneous tests of floral morphology and the degree of ecological  
112 specialization are needed to evaluate their relative importance in determining patterns of HP  
113 receipt across species (see Fang and Huang, 2013). Furthermore, for species visited by generalist  
114 pollinators, the floral neighborhood can also be important in influencing patterns of HP receipt  
115 (Rathcke, 1983; Feinsinger et al., 1986). For instance, the local abundance of conspecifics  
116 relative to heterospecifics may affect HP receipt, as rare species are expected to receive more  
117 (and more diverse) HP than common species (Rathcke, 1983; Vamosi et al., 2006; Alonso et al.,  
118 2010). To date, only a small handful of comparative studies have addressed these factors (see  
119 McLernon et al., 1996; Fang and Huang, 2013) and as of yet no study has incorporated all three.

120         Additional insight into the dynamics of HP transfer in natural communities can be gained  
121 from more detailed analysis of intraspecific variation. For instance, the evenness of pollinator  
122 quality within the assemblage could affect the pattern and variance in HP receipt among  
123 conspecific individuals. Specifically, because flower-visiting taxa (or functional groups) have  
124 differences in body size, flight distances and floral constancy they can vary in the size and purity  
125 (i.e., CP vs. HP) of the pollen load they deposit ('pollination quality'; Herrera, 1987). For  
126 instance, Hymenoptera deposited more CP than Lepidoptera and Diptera in Mediterranean

127 *Lavandula latifolia* (Herrera, 1987) and bats deposited larger amounts of HP relative to  
128 hummingbirds and hawkmoths in tropical *Aphelandra acanthus* (Muchhala et al., 2009). Thus,  
129 low quality floral visitors are those that deposit little CP but large amounts of HP, whereas high  
130 quality visitors deposit a lot of CP but little or no HP (Herrera, 1987). As a result, within an  
131 assemblage, the evenness of flower-visiting taxa (or functional groups) that differ in their quality  
132 may be an important determinant of the distribution of HP among flowers. This hypothesis can  
133 be assessed for a single plant species by estimating the pollination quality of each visiting group  
134 within its assemblage (e.g., Herrera, 1987; Muchhala et al., 2009), but such an approach is  
135 prohibitive for comparing among multiple plant species. Here we propose that the pattern of CP  
136 and HP receipt among plants within a species will reflect the joint effect of quality and evenness  
137 of their floral visitor assemblage and that these CP-HP relationships can be readily compared  
138 among species.

139         Specifically, we envision three ways HP receipt could covary with CP receipt in flowers  
140 within a species. In the first, HP receipt increases monotonically as CP increases (type 1 in Fig.  
141 1). This relationship would occur when both CP and HP grains are deposited with each pollinator  
142 visit, as expected when an evenly diverse assemblage of pollinators of similar quality visit  
143 flowers. Little variation in HP load size is expected among flowers and HP loads should be  
144 diverse because the diverse visitors are also visiting many other plant species (i.e., generalists;  
145 Neiland and Wilcock, 1999). The second possibility is that the delivery of HP is independent of  
146 delivery of CP, leading to no relationship between CP and HP receipt across conspecifics (type 2  
147 in Fig. 1). This pattern could occur when a species is visited by only one or very few high quality  
148 pollinators that deliver almost pure CP loads (e.g., specialists) and thus HP receipt occurs only as  
149 a result of pollinator ‘mistakes’ (e.g., Wang et al., 2013) which lead to unpredictable patterns of



150 HP deposition. A third alternative is that HP receipt decreases or increases exponentially as CP  
151 receipt increases (type 3A and 3B in Fig. 1). These patterns are expected to occur when a species  
152 is visited by a highly uneven pollinator assemblage that is composed of one or few frequent,  
153 high-quality pollinators and also several less frequent, low-quality pollinators (e.g., Gómez et al.,  
154 2010). We note that these predictions rest on the assumption that high quality pollinators also  
155 tend to visit flowers more frequently than low quality ones-- a pattern often observed in natural  
156 communities (e.g., Motten et al., 1981; Herrera 1987; Sahli and Conner 2006; Gómez 2000;  
157 Gómez et al., 2007; Gómez et al., 2010). Thus in species with exponential patterns of CP-HP,  
158 receipt HP would be deposited mostly during visits by low-quality pollinators that are less  
159 frequent and exhibit low constancy. As a result the HP load varies with CP (either positively or  
160 negatively) depending on the CP load size deposited by the more frequent high-quality  
161 pollinators in the assemblage. For instance, if high-quality pollinators deliver large and pure CP  
162 loads whereas HP is delivered along with few CP grains by low-quality pollinators then HP  
163 would decrease exponentially as CP receipt increases (type 3A in Fig. 1). Alternatively, if  
164 numerous high-quality pollinators deliver small but pure CP loads and infrequent low-quality  
165 pollinators deliver large, mixed pollen loads then HP will increase exponentially with CP (type  
166 3B in Fig. 1). Due to the unevenness of pollinator quality in the assemblage, HP loads will be  
167 unequally distributed across flowers, i.e., HP will be aggregated in only a few flowers, and there  
168 will be high within-species variance in HP load size. Moreover, the diversity of HP loads is  
169 predicted to be lower than that of species with a linear CP-HP relationship (type 1) because it  
170 reflects infrequent and random visitation events, rather than regular, consistent co-transport.  
171 These predictions can be tested by fitting curves to natural variation in CP and HP in flowers  
172 within species and comparing the evenness of pollinator assemblages and diversity of HP  
173 between groups defined by HP-CP relationships.

174           Accordingly, the aim of this study is to evaluate patterns of among- and within-species  
175 variation in HP receipt (load size and diversity) and identify potential drivers of such variation.  
176 First, we assess the relative importance of phenotypic and ecological (i.e., contemporary floral  
177 visitor assemblage) specialization, as well as conspecific flower density as determinants of  
178 among-species differences in mean and variance of HP load size and diversity received by  
179 flowers. Second, we evaluate patterns of within-species variation by characterizing CP-HP  
180 relationships (Fig. 1) and the diversity in the HP load and determine whether differences among  
181 species are related to the evenness of their floral visitor assemblage. We conduct these analyses  
182 with 19 plant species from three geographically distinct co-flowering communities and thereby  
183 assess the generality of these patterns across a wide range of ecological contexts.

184

## MATERIALS AND METHODS

185           *Study systems*- Heterospecific pollen receipt was studied in 19 species sampled across three  
186 geographically and ecologically distinct co-flowering plant communities: serpentine seeps of  
187 California, USA (SS), dolomite outcrops in Andalusia, Spain (DO) and dry scrublands in  
188 Yucatan, Mexico (DS) (Table 1). While each community has a unique evolutionary history, all  
189 three are seasonal flowering, species-rich communities dominated by small woody or herbaceous  
190 perennials and annuals (Estrada-Loera, 1991; Médail and Quezel, 1997, 1999; Safford et al.,  
191 2005; Freestone and Inouye, 2006; Alonso et al., 2013) that support many functional groups of  
192 insect flower visitors (e.g., Herrera et al., 2006; Campos-Navarrete et al., 2013; Koski et al.,  
193 2015). Thus, these communities reflect ecologically and evolutionarily independent replicates of  
194 similarly diverse plant-pollinator communities (Table 1).

195           Within each community four to nine insect-pollinated plant species with overlapping  
196 flowering times were selected (Table 1). Plant species represented nine families and varied in  
197 flower color, size, longevity, mating system and type of rewards (Table 2, Appendix 1, Alonso et  
198 al., 2013). Both, zygomorphic and actinomorphic species were included from each community  
199 and were similarly represented in the data set (10 vs. 9; Appendix 1).

#### 200       ***Data collection-***

201   *Pollen receipt*– Within each community an average of 135 wilted flowers (range: 52-222) was  
202 collected from each species along 3-5 transects at bi-weekly intervals during peak flowering  
203 (Table 2). This sampling scheme allowed us to capture both temporal and spatial variation in  
204 pollen receipt for each species. Across the 19 species, styles from a total of 2,566 flowers were  
205 collected and stored in 70 % ethanol until they were stained with decolorized aniline blue and  
206 prepared for scoring. Conspecific pollen and HP on stigmas were visualized with the aid of  
207 fluorescence microscopy (Kearns and Inouye, 1993). Both CP and HP were counted and  
208 summed across all styles within a flower when more than one style was present. A pollen library  
209 was constructed for each community to aid in the identification of pollen to species when  
210 possible or to morphotype when not. Heterospecific pollen was classified based on morphology  
211 (size, shape and exine ornamentation). For analysis HP was pooled into 16 to 32 morphotype  
212 categories per community (Table 1). Each morphotype consisted of 1-6 possible species known  
213 to co-flower (Koski et al., 2015; Alonso, Ashman and Parra-Tabla unpublished data). The mean  
214 and coefficient of variation [CV] in HP load size (number of HP grains per flower) and diversity  
215 (number of HP morphotypes per flower) were calculated for each species (Appendix 1).

216

217 *Floral visitors*– The contemporary assemblage of floral visitors was characterized for each  
218 species by conducting 3-5 min observations of flowers within fixed plots scaled appropriately for  
219 each community (119 2m<sup>2</sup> plots in SS, 112 4m<sup>2</sup> plots in DO, and 60 4m<sup>2</sup> plots in DS). Pollinator  
220 censuses were conducted during the period of highest pollinator activity (i.e., between 8:00 and  
221 16:00h) twice per plot per week for the duration of the flowering season. A total of 252.5, 32.5  
222 and 183.7 hrs of observation was conducted at SS, DO and DS respectively. During each census  
223 the number of open flowers and the number and identity of floral visitors on a given species was  
224 recorded. Only species that received at least 25 visits were analyzed ( $N = 15$  species; Appendix  
225 1).

226 Floral visitors are often categorized into functional groups based on their morphology (e.g.,  
227 body size), energetic requirements and flight ability, which can determine their foraging and  
228 flower-handling behavior, the range of flowers they can visit, as well as their conspecific pollen  
229 transfer abilities and thus the type of selection they generate (Fenster et al., 2004; Moretti et al.,  
230 2009). In this sense, insects within a pollinator functional group are also expected to be  
231 equivalent in their contribution to HP transfer to a given plant species. Members of a functional  
232 group are thus more similar to each other in their quality as pollinators than to members of other  
233 groups (Zamora, 2000; Fenster et al., 2004; Greenleaf et al., 2007; Geslin et al., 2013; Rosas-  
234 Guerrero et al., 2014; Koski et al., 2015) and the presence/absence of a particular group can have  
235 important effects on plant fitness (Gómez et al., 2010; Albrecht et al., 2012; Fründ et al., 2013).  
236 Thus, in this study, floral visitor diversity was calculated based on 11 behaviorally and  
237 functionally defined groups based on their body size, energetic requirements and  
238 foraging/feeding behavior (Appendix 2). These functional groups were present across all three  
239 communities and included: large solitary bees, small solitary bees, large social bees, beetles, bee

240 flies, large syrphid flies, small syrphid flies, other flies, butterflies, walking insects (e.g., ants)  
241 and wasps (Appendix 2). Simpson's reciprocal diversity index ( $1/D$ ) and evenness were  
242 calculated for each study species using data on number of visits by these functional groups.  
243 Simpson's reciprocal index ( $1/D$ ) was chosen because it is an abundance weighted diversity  
244 estimate ( $1/D = 1/\sum[n / N]^2$  where  $n$  = total number of organisms in a particular functional group,  
245 and  $N$  = total number of organisms in all functional groups) that is very robust even with small  
246 sample sizes and thus is a reliable measure of diversity (Magurran, 2004). Rarefaction analyses  
247 demonstrated that our sampling effort was sufficient to adequately characterize the diversity of  
248 pollinator functional groups visiting plant species as we were able to capture between 80-98% of  
249 all possible plant-pollinator interactions (Koski et al., 2015; Alonso et al., unpubl. data).  
250 Functional group diversity was not correlated with the total number of visits observed for a given  
251 species ( $r = 0.09$ ,  $P = 0.7$ ).

252 Simpson's evenness was calculated by dividing Simpson's reciprocal index by the total  
253 number of pollinator functional groups in the sample (Magurran, 2004). Thus, evenness varies  
254 from 0 (completely uneven) to 1 (complete evenness).

255 *Conspecific density*– Conspecific flower density for each focal plant species was determined  
256 from the records of the number of open flowers per plot during each floral visitor census. To  
257 produce a single value per species of local conspecific density (flowers/m<sup>2</sup>) flower densities were  
258 averaged across the season for every plot where that species flowered. This estimate of local  
259 conspecific flower density is correlated with its global floral abundance in the community ( $r =$   
260  $0.99$ ;  $P = 0.001$ ,  $N = 19$ ).

261 *Flower traits*– Flower biomass was used as a proxy of flower size. For each species, we  
262 collected a minimum of 15 flowers and these were dried in bulk at 60°C for 24 h and then  
263 weighed to the nearest 0.0001 g on an AE200 Mettler analytical balance (Mettler-Toledo,  
264 Columbus, Ohio, USA). Total dry weight was then divided by the total number of flowers to  
265 estimate average flower biomass for each species. Flower longevity was estimated by recording  
266 the number of days a flower remained open (from anthesis to senescence) on a minimum of five  
267 individuals (excluded from pollinators) per species during peak flowering in each community.  
268 The average flower size and longevity for each species was used in analyses. Floral shape was  
269 scored as zygomorphic (bilateral symmetry) or actinomorphic (radial symmetry).

#### 270 *Data Analyses-*

271 *Sources of variation in HP receipt*- To evaluate the pattern of variation in size (number of HP  
272 grains) and diversity (number of HP types) of the HP load per flower, we conducted nested  
273 mixed effects ANOVAs (proc mixed; SAS 2010). To account for the potential influence of  
274 shared evolutionary history on species patterns of HP receipt, family and species (nested within  
275 family) were treated as random effects, while community was treated as a fixed effect. We then  
276 used this same model to partition the total variance in HP load size and diversity among families,  
277 species, communities and individuals within species (residual variation; proc varcomp, method =  
278 MIVQUE0; SAS 2010). Load size and diversity of HP types were (square root + 0.5)  
279 transformed in order to meet assumptions of normality of residuals.

280 *Among-species variation in HP receipt*- Means and CV of HP load size and diversity were  
281 calculated for each species and used in among-species regressions to determine the relative  
282 importance of ecological specialization (contemporary floral visitor diversity), conspecific

283 flower density and phenotypic specialization (flower size, longevity and symmetry) in explaining  
284 interspecific variation in HP receipt. Given the minor, but significant, effect of community on  
285 HP load size and diversity (see results), the effect of community was removed using a single  
286 effect linear model (proc glm; SAS 2010) and using the residuals in subsequent regression  
287 analyses across all species regardless of their community (e.g., Jakob et al., 1996; Essenberg,  
288 2013). There was no effect of community on any of the independent variables studied (all  $P >$   
289 0.3) so raw data were used in the regression analyses. Standardized regression coefficients were  
290 calculated for each variable (each observation minus the mean and divided by the variance) to  
291 facilitate comparisons among independent variables. To test for an effect of continuous variation  
292 in phenotypic specialization (flower size and longevity), ecological specialization and  
293 conspecific flower density, multiple regressions were conducted. To test for an effect of  
294 categorical variation in phenotypic specialization (flower symmetry: actinomorphic vs.  
295 zygomorphic) and its potential interaction with floral visitor diversity and conspecific density an  
296 ANCOVA was run (proc glm; SAS 2010). When interactions between flower symmetry and  
297 visitor diversity or conspecific density were non-significant ( $P > 0.5$ ) they were removed from  
298 the models. Heterospecific pollen load size, diversity of HP types (square root + 0.5) and  
299 conspecific flower density (log10) were transformed to improve the distribution of residuals.  
300 Insect visitation and flower size data were not available for four and one species respectively,  
301 (Appendix 1), so the final data set for the analysis of species-specific traits on HP receipt  
302 included a total of 14 species.

303 *Within-species variation in HP receipt-* To characterize within-species relationships between HP  
304 and CP receipt (Fig. 1), linear and non-linear regressions between the amount of CP and HP  
305 received per flower were performed separately for each species. First we tested for the

306 significance of both linear (proc reg; SAS 2010) and non-linear relationships between CP and HP  
307 receipt. The non-linearity of the relationship was tested with the inclusion of a non-parametric  
308 component (spline) using a generalized additive model with a cubic spline and a Poisson error  
309 distribution (proc gam; SAS 2010). If only a significant linear relationship was found then the  
310 species was classified as type 1. If both (linear and non-linear) were found to be non-significant  
311 then the species was categorized as type 2 (amount of HP received independent of the amount of  
312 CP). Only when the non-linear relationship was found significant was an exponential model  
313 ( $HP = a * (\exp[b_1 * CP])$ ) then fit to the data (proc nlin; SAS 2010). Species with a significant  
314 negative  $b$  have an exponentially decreasing CP-HP relationship (type 3A), while those with  
315 positive  $b$  have an exponentially increasing CP-HP relationship (type 3B). If both linear and non-  
316 linear relationships were found to be significant, then we assessed the fit of the models by  
317 comparing adjusted  $R^2$ s and the model with the best fit was selected (larger  $R^2$  values). Pseudo  $R^2$   
318 for non-linear models was estimated as  $1 - (\text{Error sum of squares} / \text{Total sum of squares}$   
319 [corrected]) (Introduction to SAS. UCLA: Statistical Consulting Group [online]. Website  
320 <http://www.ats.ucla.edu/stat/sas/notes2> [accessed 29 May 2015]). Finally, we performed linear  
321 models to evaluate whether species with different patterns of CP-HP receipt (linear vs. nonlinear:  
322 type 1 vs. 3A and 3B; see results) also differed significantly in evenness of flower visiting fauna,  
323 CV of HP load size or total HP diversity received. These models were based upon the residuals  
324 from a one-way ANOVA where community was the predictor of each response variable.

325

## RESULTS

326 **Sources of variation in HP receipt-** We found extensive variation among species in mean HP  
327 load size (mean  $\pm$  SE:  $24.7 \pm 11.4$ ; range: 0.04-191.5 pollen grains) and diversity ( $1.3 \pm 0.25$ ;



328 0.03-3.8 morphotypes) per flower as well as in their respective CVs ( $302.6 \pm 30.5$ ; 102.9-612.1  
329 and  $140.6 \pm 27.1$ ; 42.1-547.6 for HP load size and diversity respectively; Fig 2; Appendix 1).

330 Variation among individuals within a species (residual variation) accounted for the highest  
331 proportion of the total variation in HP load size and diversity (46 and 50% respectively).  
332 Nevertheless, the contribution of species-level variation to total variation for both response  
333 variables was also substantial (28 and 10% respectively). Variation among communities  
334 explained 23 and 7%, and plant family explained 3 and 33% of the total variation in HP load size  
335 and diversity, respectively.

336 In the global analysis, the average HP load size and diversity received per flower was  
337 significantly different among species ( $Z = 1.86$ ,  $P < 0.03$  and  $Z = 2.18$ ,  $P < 0.01$ , respectively).  
338 Heterospecific pollen load size and diversity also varied significantly among communities ( $F_{2,}$   
339  $_{2547} = 5.77$ ,  $P = 0.003$  and  $F_{2, 2547} = 4.28$ ,  $P = 0.01$ , respectively) while family had no effect in  
340 either response variable (both  $Z < 1.4$ ,  $P > 0.05$ ).

341 ***Among-species variation in HP receipt-*** There was wide variation in floral visitor diversity  
342 ( $2.02 \pm 0.25$ ; 1.05-4.09), conspecific flower density ( $19.7 \pm 1.25$ ; 0.6–217.2 flowers/m<sup>2</sup>), flower  
343 longevity ( $3.8 \pm 0.79$ ; 0.5-12 days) and flower biomass ( $0.008 \pm 0.001$ ; 0.0006-0.29g) among  
344 species providing ample discriminatory ability in regressions even with a small number of  
345 species (Appendix 1).

346 Among species, mean HP load size increased significantly with increasing floral visitor  
347 diversity and flower biomass (Table 3A). Floral visitor diversity was the only significant factor  
348 affecting HP load diversity and this was a positive effect (Table 3B). Conspecific flower density  
349 and flower longevity did not significantly affect either aspect of HP receipt (Table 3A, B).

350 Although there was substantial variation among species in the CV of HP load size and diversity  
 351 (see above), none of the factors we measured significantly explained this variation (Table 3C,  
 352 D).

353 When flower symmetry was included in a separate model to account for categorical  
 354 variation in floral phenotype (see data analysis above) the results did not change. That is, the  
 355 strong effect of floral visitor diversity and the non-significant effect of conspecific density, ( $F_{1,9} = 14.5$ ,  $P = 0.004$ ;  $F_{1,11} = 0.4$ ,  $P > 0.5$  respectively) and diversity ( $F_{1,9} = 9.96$ ,  $P = 0.01$ ;  $F_{1,9} =$   
 356  $0.8$ ,  $P > 0.4$ ) on HP load size remained. There were no significant interactions of flower  
 357 symmetry with either continuous variable ( $F_{1,9} = 0.04-1.3$ ; all  $P > 0.3$ ). And even though average  
 358 HP load size and diversity were slightly elevated in actinomorphic flowers compared to  
 359 zygomorphic ones, these were not statistically significant differences (HP load size:  $27.2 \pm 18.3$   
 360 vs.  $22.1 \pm 14.2$  grains;  $F_{1,11} = 0.18$ ,  $P > 0.6$ ; HP diversity:  $1.6 \pm 0.3$  vs.  $0.9 \pm 0.3$ ;  $F_{1,11} = 1.5$ ,  $P =$   
 361  $0.24$ ).

363 ***Within-species variation in HP receipt-*** Of the 19 species evaluated seven exhibited  
 364 significant linear relationships between the amount of CP and HP received (type 1 in Fig. 1; Fig.  
 365 3A for an example), eight showed an exponentially decreasing relationship (type 3A in Fig. 1;  
 366 Fig. 3C) and three showed an exponentially increasing relationship (type 3B in Fig. 1; Fig. 3D;  
 367 Table 2). Only in one species did HP vary independently of CP (type 2 in Fig. 1; Fig. 3B; Table  
 368 2). Furthermore, differences in CP-HP patterns (linear type 1 vs. exponential type 3A and 3B)  
 369 reflected differences in HP load size and diversity and pollinator evenness in the predicted  
 370 directions, although none of these relationships were statistically significant (HP diversity:  $1.8 \pm$   
 371  $0.3$  vs.  $0.9 \pm 0.3$ ;  $F_{1,16} = 3.21$ ,  $P = 0.09$ ; CV in HP load size:  $291 \pm 36$  vs.  $270 \pm 42$ ;  $F_{1,16} = 0.11$ ,

372  $P > 0.7$ ; pollinator evenness:  $0.44 \pm 0.08$  vs.  $0.34 \pm 0.06$ ;  $F_{1, 16} = 1.16$ ,  $P > 0.4$ ). There was also  
373 no association between flower symmetry and CP-HP relationship type ( $\chi^2_{16} = 0.01$ ,  $P > 0.9$ ).

## 374 DISCUSSION

375 Wide variation in mean HP load size and diversity in naturally pollinated flowers was  
376 revealed in this detailed comparison of 19 species across three plant communities. Diversity of  
377 the contemporary flower-visiting community emerged as a strong and consistent driver of  
378 variation in HP receipt among species, more so than evolutionary degree of specialization, as  
379 represented by floral phenotype (symmetry, biomass and longevity). Furthermore, within-species  
380 variation in HP load size and diversity could be associated with patterns of CP receipt and two  
381 dominant CP-HP relationships emerged (linear or exponentially decreasing) indicating the  
382 potential for promising explorations of mechanisms at this level in the future. We discuss these  
383 results and their broader ecological implications in detail below.

384 ***Among-species variation in HP receipt-*** Plant species identity accounted for more variation in  
385 HP load size than family membership suggesting that the processes that influence variation in HP  
386 load size are likely independent of common evolutionary history. However, this was not the case  
387 for HP load diversity where among-family variation was more important. This latter result is  
388 consistent with studies that have shown, across many plant-pollinator communities, that  
389 pollinator guilds tend to visit closely related plant species (e.g., Vamosi et al. 2014) and thus  
390 species within the same plant family are likely to receive similar diversity of HP grains.  
391 Furthermore, HP receipt varied across communities but variation among them explained less  
392 than variation among species and individuals in both HP load size and diversity, suggesting that  
393 underlying processes that influence HP receipt may operate across a wide range of ecological

394 and evolutionary contexts. Interestingly, each community had both species with high mean HP  
395 receipt and species with very low HP receipt (Fig. 2). For example, in the serpentine seeps  
396 *Delphinium uliginosum* and *Triteleia peduncularis* received hundreds of HP grains—ten times  
397 the HP per flower received by *Zigadenus venenosus*, and in the dolomite outcrops *Silene*  
398 *lasiostyla* received twenty times that of *Thymus orospedanus* or *Sideritis incana* (Fig. 2;  
399 Appendix 1). Similar dramatic differences have been observed between species within alpine  
400 (Fang and Huang, 2013), prairie (Montgomery and Rathcke, 2012), and old field communities  
401 (McLernon et al., 1996) suggesting that avoidance of HP may not be the only evolutionary stable  
402 strategy for co-existence. That is, there may be two evolutionary strategies: selection for  
403 avoidance of HP on one hand and selection for tolerance of the effects of HP receipt on the other  
404 that could contribute to species coexistence (Ashman and Arceo-Gómez, 2013). Studies that link  
405 natural patterns of HP receipt to their causes (see below) as well as fitness costs are needed to  
406 test these ideas (Ashman and Arceo-Gómez, 2013).

407         There was also substantial among-species variation in the diversity of HP loads.  
408 However, given our use of morphotypes (which each could represent 1-6 species) these data  
409 most likely represent underestimates of the diversity of interactions. In fact, when species have  
410 been identified up to 15 species of HP can occur on one flower (Bartomeus et al., 2008; Fang and  
411 Huang, 2013). Such extensive HP transfer indicates that pollinator sharing is leading to  
412 multispecies interactions on the style for most plants in the community, the consequences of  
413 which can be greater than just the sum of individual species' effects (Arceo-Gómez and Ashman,  
414 2011). What is clear, however, is that this variation in load size and diversity provides a window  
415 into understanding the ecological drivers of HP receipt, as well as can help pinpoint where  
416 opportunities for natural selection to avoid or tolerate HP receipt lie (see below).

417           After accounting for community differences, HP load size and diversity reflected species-  
418 specific floral visitor diversity, more than other measures of ecological context (conspecific  
419 density) or phenotypic specialization (floral symmetry, longevity and biomass). The only other  
420 study of HP patterns that directly characterized contemporary floral visitor assemblages also  
421 found that more generalized plants tended to receive larger and more diverse loads of HP than  
422 specialized ones (Fang and Huang, 2013). Moreover, in our study the increase in HP load size  
423 and diversity with increasing floral visitor diversity occurred irrespective of flower symmetry, a  
424 finding that supports the notion that phenotypic specialization may not always reflect ecological  
425 specialization (Ollerton, 1996). Such a disconnect may also explain patterns of HP receipt  
426 contrary to the presumed level of specialization based on symmetry, i.e., higher HP receipt in  
427 zygomorphic than actinomorphic flowers found in other studies (McLernon et al., 1996).

428           Floral traits that defined specialization or restrictiveness, while being quite varied in our  
429 data set (Appendix 1), were not strong determinants of variation in any aspect of HP receipt.  
430 Only flower biomass affected HP load size among-species, with species with heavier (larger)  
431 flowers receiving larger and more diverse HP loads than lighter (smaller) ones. Because we  
432 accounted for visitor diversity, the effect of flower biomass is not through differences in  
433 generalization level but rather may be mediated through enhanced visitation rates (e.g., Galen  
434 and Newport, 1987; Conner and Rush, 1996) or larger/more exposed stigmas (Montgomery and  
435 Rathcke, 2012; Fang and Huang, 2013). Interestingly, flower symmetry only had a minor effect  
436 on mean HP load size, and HP receipt appeared to scale similarly with floral visitor diversity in  
437 both actinomorphic and zygomorphic flowers. Furthermore, we did not find any evidence to  
438 support the prediction that sparse (rare) species are more likely to receive HP than abundant ones  
439 (Rathcke, 1983; McLernon et al., 1996), contrasting with the results of Jakobsson et al. (2009).

440 Despite the observation of a wide range in CVs of HP load size and diversity across  
441 species (Appendix 1), none of the predictors we tested could account for this variation (Table 3).  
442 The strongest non-significant indicator of this variation was flower longevity, possibly reflecting  
443 increased visitation rate per flower, and/or increased likelihood of stochastic visitation events by  
444 inconstant visitors. But more study is needed to pinpoint the sources of variation in this  
445 important aspect of HP receipt.

446 ***Within-species variation in HP receipt-*** Within species, HP load size and diversity varied 6-  
447 10 fold in magnitude (Appendix 1) and variance at this level was greater than the combined  
448 contributions of family and species identity (> 46% of the total). Our exploration of within-  
449 species variation in HP load size could be predicted from two dominant CP-HP relationships  
450 (type 1 vs. 3A in Fig. 1; Table 2) and suggests that two distinct types of plant-pollinator  
451 interactions prevail. We found approximately half of the species studied showed significant  
452 linear increases in HP with CP (and also tended to have higher diversity of HP types) while most  
453 of the remaining species showed exponentially decreasing relationships (and fewer HP types).  
454 And while the difference in pollinator evenness between these two groups was in the predicted  
455 direction (greater for species with type 1), it was not significant. Nevertheless, the evaluation of  
456 CP-HP relationships among flowers provides insight into the genesis of variation in HP receipt  
457 that cannot be discerned from means or variances alone and our comparative data provide first  
458 insight into the possible factors responsible for within-species variation in HP receipt. For  
459 instance, the fact that evenness alone did not capture the within-species variance in pollen  
460 transfer dynamics may suggest that variation in pollinator quality is more important than  
461 variation in abundance in influencing deposition dynamics, or that variation not captured by our  
462 floral visitor functional groups is important. Pollinator quality, in particular, is likely a complex

463 trait that results from not only inconstancy but also pollinator grooming (Thomson, 1986;  
464 Holmquist et al., 2012), pollen carryover (Thomson and Plowright 1980) and pollen loss during  
465 transport (Wilcock and Neiland, 2002), and thus, it may be difficult to capture with only  
466 information on the visitor functional groups defined herein.

467       Nevertheless, more species–level studies are needed to tease apart the contribution of  
468 pollinator evenness from pollinator quality as well as to take into account fine-scale  
469 neighborhood variation in flower diversity and abundance. Such work would provide insight into  
470 the potential for ongoing evolution of either avoidance (e.g., pollen placement [Armbruster et al.,  
471 1994], site of stigma contact [Armbruster et al., 2014], stigma size [Montgomery and Rathcke,  
472 2012]) or tolerance strategies (e.g., stigma secretions [Ashman and Arceo-Gómez, 2013]) in  
473 contemporary communities. While it is a daunting task to perform these studies across many  
474 species within communities, such studies are needed to understand how traits evolve in multi-  
475 species contexts.

476       **Conclusions-** This study illustrates how widespread the phenomenon of HP receipt is both  
477 within and across diverse plant communities. In doing so it highlights ecological generalization  
478 as an important determinant of interspecific variation in both the size and diversity of HP load.  
479 Indicating that contemporary changes in pollinating fauna (loss of species, shifts in flowering  
480 time) will likely translate into changes in the plant-plant interactions on the style (e.g., Brosi and  
481 Briggs 2013) We further confirm that HP and CP receipt are not independent of each other at  
482 intraspecific level for some species and propose that the shape of such relationship reflects past  
483 evolution and may provide substrate for future evolution to shape avoidance of HP transfer.

484

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**Table 1.** Location, community type (code), flowering period, number of focal species, total co-flowering species richness, flower visitor functional groups observed and number of pollen morphotypes identified at each location.

<b>Location</b>	<b>Community</b>	<b>Flowering period sampled</b>	<b>Number of focal species</b>	<b>Number of insect pollinated co-flowering species</b>	<b>Number of flower visitor functional groups</b>	<b>Number of pollen morphotypes</b>
California	Serpentine seeps (SS)	June-July 2010	6	50	11	20
Andalusia	Dolomite outcrops (DO)	May-June 2010	9	67	11	32
Yucatan	Dry scrublands (DS)	July-September 2011	4	38	10	16

**Table 2.** Nineteen study species (species code used in Fig 2; for complete taxonomic names see Appendix 1), plant family (family codes), community type and sample size ( $N$ ). For each, the evenness of flower visiting fauna, within-species variance (CV) in HP load size (pollen grains), and mean number of HP morphotypes receive is given, as well as the type of CP-HP relationship (described in Fig. 1) observed. For each CP-HP regression the coefficient ( $b$ ) and its significance are given. ‘.’ denotes data not available.

Community type	Family (code)	Species code	$N$	Pollinator evenness	CV of HP load size	Mean number of HP types	HP-CP pattern	$b$	$P$
SS	Plantaginaceae (Pl)	Anc	150	0.51	220.9	1.2	Increasing linearly	0.04	0.0007
SS	Ranunculaceae (Ra)	Deu	150	0.68	102.9	2.7	Increasing linearly	0.32	0.0001
SS	Phrymaceae (Ph)	Mig	151	0.19	237.2	2.09	Decreasing exponential	-0.0015	0.0001
SS	Phrymaceae (Ph)	Min	151	0.35	318.8	1.3	Decreasing exponential	-0.002	0.0004
SS	Liliaceae (Li)	Trp	115	0.53	137.4	3.8	Increasing linearly	0.99	0.0001
SS	Liliaceae (Li)	Ziv	151	0.52	156.5	1.8	Increasing linearly	0.032	0.03
DO	Cistaceae (Ci)	Fub	113	.	187.9	1.2	Decreasing exponential	-0.0009	0.0001
DO	Cistaceae (Ci)	Fup	63	0.6	260.4	1.5	Decreasing exponential	-0.004	0.008
DO	Cistaceae (Ci)	Hcr	90	0.23	395.8	1.4	Decreasing exponential	-0.006	0.003
DO	Cistaceae (Ci)	Hci	129	.	612.1	0.27	Decreasing exponential	-0.005	0.001
DO	Lamiaceae (La)	Sii	123	0.37	580.6	0.03	No pattern	0.0001	0.86
DO	Caryophyllaceae (Ca)	Sco	133	.	349.2	2.3	Increasing exponential	0.005	0.002
DO	Caryophyllaceae (Ca)	Spl	200	.	338.8	3.1	Increasing linearly	0.09	0.01
DO	Lamiaceae (La)	Tpo	222	0.31	285.7	0.29	Decreasing exponential	-0.004	0.0001
DO	Lamiaceae (La)	Tho	118	0.23	349.7	0.14	Decreasing exponential	-3.6	0.0001
DS	Plantaginaceae (Pl)	Ang	52	0.31	413.4	0.28	Increasing linearly	0.17	0.0004
DS	Malvaceae (Ma)	Cie	212	0.19	300.4	0.77	Increasing linearly	0.05	0.0001
DS	Lythraceae (Ly)	Cup	97	0.4	256.2	0.27	Increasing exponential	0.01	0.002
DS	Malvaceae (Ma)	Sid	148	0.43	246.3	0.25	Increasing exponential	0.01	0.001

**Table 3.** Multiple regressions for the effects of: flower biomass, longevity, floral visitor diversity and conspecific flower density on mean and variance (CV) of HP load size (number of grains) and diversity (number of morphotypes) ( $N=14$  species). The coefficient of determination ( $R^2$ ) and significance is given for each model along with the standardized regression coefficients ( $\beta$ ) for each independent variable. Significant models and regression coefficients ( $P < 0.05$ ) are denoted in bold face.

Variables		Model			
Dependent	Independent	$\beta$	$P$	$R^2$	$P$
A. Mean HP load size	Flower biomass	0.44	<b>0.05</b>	0.78	<b>0.005</b>
	Flower longevity	0.01	>0.9		
	Flower visitor diversity	0.54	<b>0.02</b>		
	Conspecific density	-0.01	>0.9		
B. Mean HP diversity	Flower biomass	0.29	0.12	0.84	<b>0.001</b>
	Flower longevity	-0.23	0.14		
	Flower visitor diversity	0.53	<b>0.01</b>		
	Conspecific density	-0.23	0.2		
C. CV of HP load size	Flower biomass	0.11	>0.7	0.42	0.24
	Flower longevity	0.24	>0.4		
	Flower visitor diversity	-0.64	0.07		
	Conspecific density	0.02	>0.9		
D. CV of HP diversity	Flower biomass	-0.14	>0.6	0.57	0.07
	Flower longevity	0.56	0.04		
	Flower visitor diversity	-0.16	>0.5		
	Conspecific density	0.1	>0.4		

**Appendix 1** Description of study plant species in each community (com.) serpentine seeps (SS), dry scrubland (DS) and dolomite outcrops (DO). Family (Fam.) and species (Sp.) codes. Information on floral characters (mean flower size [biomass], flower shape (actinomorphic [A], zygomorphic [Z]), mean flower longevity[days]) and ecological characters (mean conspecific flower density [flowers/m<sup>2</sup>], floral visitor diversity [1/D]) and heterospecific pollen (HP) receipt (mean and CV of load size and diversity per flower).

‘.’ denotes data not available.

Com.	Fam. code	Species	Sp. code	Floral characters			Ecological characters		HP load size		HP load diversity	
				Flower biomass (g)	Floral shape	Flower longevity	Flower density	Floral visitor diversity	Mean	CV	Mean	CV
DO	Ca	<i>Sideritis incana</i>	Sii	0.00418	Z	12	25.9	1.1	0.04	580.6	0.03	547.6
DO	Ca	<i>Silene lasiostyla</i>	Spl	0.00966	A	4.5	0.6	.	18.5	338.8	3.13	75.2
DO	Ci	<i>Fumana baetica</i>	Fub	0.00887	A	0.5	1.7	.	9.1	187.9	1.2	94.7
DO	Ci	<i>Fumana paradoxa</i>	Fup	0.00924	A	0.5	3.4	2.4	9.4	260.4	1.5	61.9
DO	Ci	<i>Helianthemum cinereum</i>	Hcr	0.0026	A	2.3	1.9	.	2	612.1	0.27	182.8
DO	Ci	<i>Helianthemum appeninum</i>	Hci	0.01614	A	1	7.6	1.8	14.3	395.8	1.4	129.4
DO	La	<i>Silene colorata</i>	Sco	0.00788	A	.	1.3	.	11.6	349.2	2.3	88.9
DO	La	<i>Thymus orospedanus</i>	Tpo	0.00069	Z	4	217.2	1.8	0.3	349.7	0.14	279
DO	La	<i>Teucrium polium</i>	Tho	0.00253	Z	10	19.5	1.3	0.75	285.7	0.29	197.7
SS	Li	<i>Triteleia peduncularis</i>	Trp	0.0149	A	4.4	3.6	4.2	191.5	137.4	3.8	42.1
SS	Li	<i>Zigadenus venenosus</i>	Ziv	0.004	A	2.8	5.8	2.5	10.6	156.5	1.8	71.3
DS	Ly	<i>Cuphea gaumeri</i>	Cup	0.002	Z	5	12	2.4	0.59	256.2	0.27	161.8
DS	Ma	<i>Cienfuegosia yucatanensis</i>	Cie	0.02	A	1	2.9	1.3	4.09	300.4	0.77	102.3
DS	Ma	<i>Sida acuta</i>	Sid	0.003	A	0.5	3.5	2.1	0.81	246.3	0.25	187.9
SS	Ph	<i>Mimulus guttatus</i>	Mig	0.00638	Z	3.1	14.5	1.6	22.1	237.2	2.09	74.7
SS	Ph	<i>Mimulus nudatus</i>	Min	0.0022	Z	2.2	11.8	1.1	17.1	318.8	1.3	82.9

SS	Pl	<i>Antirrhinum cornutum</i>	Anc	0.0033	Z	6.2	6	1.5	16.09	220.9	1.2	45.4
DS	Pl	<i>Angelonia angustifolia</i>	Ang	.	Z	1	41.4	1.2	8.5	413.4	0.28	198.2
SS	Ra	<i>Delphinium uliginosum</i>	Deu	0.02975	Z	8.2	10.3	4.1	133.2	102.9	2.7	48.5

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**Appendix 2** Description of floral visitor functional groups (adapted from Koski et al., 2015) observed visiting plants in the three studied communities, serpentine seeps (California), dolomite outcrops (Andalusia) and dry scrublands (Yucatan).

<b>Flower visitor functional group</b>	<b>Description</b>
<b>Large solitary bees</b>	Bees with large bodies ( $\geq 10$ mm), solitary, forage on pollen and nectar, pollen carried on hind legs or the underside of abdomen, includes members of Andrenidae and Megachilidae
<b>Small solitary bees</b>	Bees of very small body size ( $\leq 5$ mm), ectothermic, narrow temperature range of activity, fast flyer, solitary, pollen carried on hind legs, includes members of Halictidae and Apidae
<b>Large social bees</b>	Bees with large bodies ( $\sim 15$ mm), social, efficient forager of pollen and nectar, efficient recruiter, pollen carried on hind legs, includes members of Apidae
<b>Beetles</b>	Pollen eaters, inefficient, mostly ectothermic, narrow daily activity range, short flying range, pollen not carried in specific location on body, includes members of Cleridae and Buprestidae (Coleoptera)
<b>Bee flies</b>	Flies in the family Bombyliidae, fast feeder, narrow thermal range, territorial, narrow spectrum of flowers, long mouth parts, pollen not carried on specific location on body
<b>Large syrphid flies</b>	Large flies in the family Syrphidae ( $\geq 10$ mm), hoverflies, forage on pollen and nectar, fast flying, specialized digestive system for processing pollen, long foraging time, pollen not carried on specific location on body
<b>Small syrphid flies</b>	Small flies in the family Syrphidae ( $\leq 10$ mm), hoverflies, forage on pollen and nectar, fast flying, specialized digestive system for processing pollen, long foraging season, pollen not carried on specific location on body
<b>Other flies</b>	Other flies different from bombyiid and syrphid flies, incidental visitors, small, inefficient, pollen not carried on specific location on body, includes members of Muscidae and Callophoridae
<b>Butterflies</b>	Nectar foragers, long foraging times, visit few flowers per plant, fly long distances, narrow thermal requirements, forage on low concentrated nectar, pollen not carried on specific location on body, includes members of Lepidoptera
<b>Wasp</b>	Carnivorous, secondary flower visitors, some groups visit for nectar, pollen not carried on specific location on body, includes members of Vespidae
<b>Walking insects</b>	Walking insects, pollen not carried on specific location on body, includes members of Formicidae

**Fig 1.** Predicted intraspecific relationships between conspecific (CP) and heterospecific pollen (HP) receipt per flower derived from variation in the evenness and quality of the floral visitor assemblage (see text). Four relationships are hypothesized: 1) HP receipt increases linearly with CP receipt (solid line); 2) HP varies independently of CP receipt and thus no relationship exists between HP and CP receipt (dotted line); 3A) HP decreases or 3B) increases exponentially with increasing CP receipt (dashed lines). Note that curve 2 is shown below curves 3A and 3B for visual clarity and not because HP is predicted to be lower.

**Fig. 2** Mean ( $\pm$  SE) (A and B) and coefficient of variation (CV) (C and D) for heterospecific pollen (HP) load size (number of grains per flower) and HP diversity (number of HP morphotypes per flower) for each of the 19 studied plant species. Species were sampled in three plant communities: serpentine seeps (open bars), dry scrubland (solid bars) and dolomite outcrops (dashed bars; Table 1). Species codes follow names in table 2. Note the broken axis in pane A.

Fig. 3 Examples of the four types of relationships between the amount of conspecific (CP) and heterospecific pollen (HP) received on stigmas (described in Figure 1). A) type 1: HP linearly increases with CP in *Delphinium uliginosum*, B) type 2: HP varies independently with CP in *Silene colorata*; C) type 3A: HP exponentially decreases with CP in *Mimulus guttatus*; D) type 3B: HP exponentially increases with CP in *Sida acuta*. Note the broken axis in pane C.

Figure

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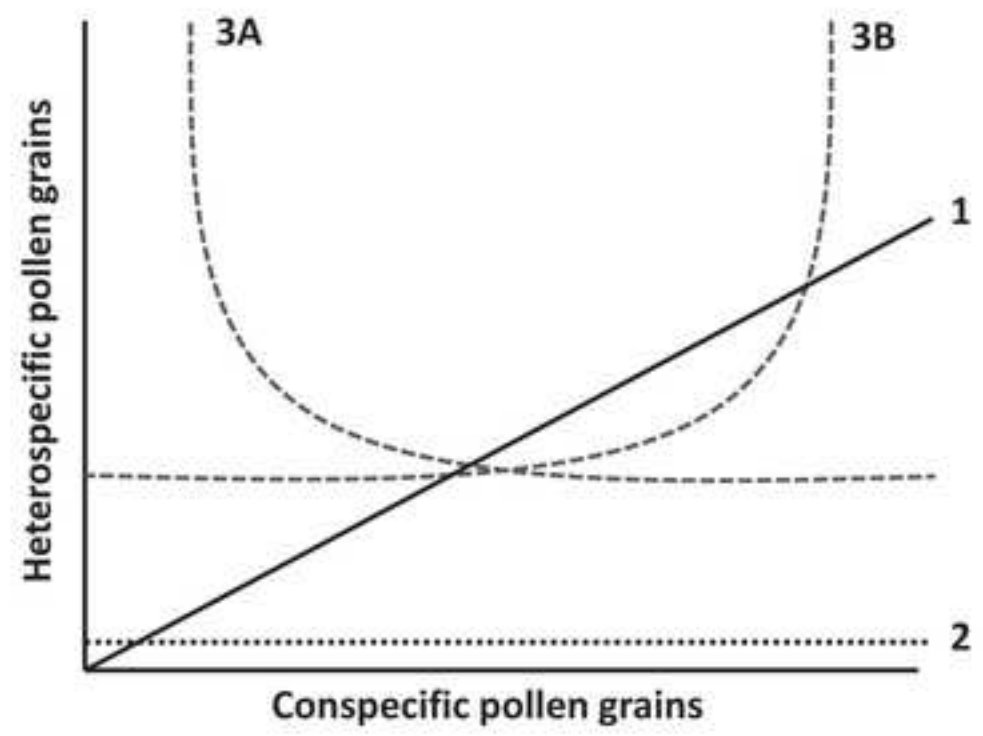




Figure  
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