

Relationship between floral colour and pollinator composition in four plant communities

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The present study started on February 2014. I was provided with the database of pollinator censuses I needed for the study. My contribution to this work has been: 1) Conducting fieldwork along the flowering season to obtain the flower samples and also performing censuses to expand the database; 2) conducting flower colour measurements with the spectrometer; 3) definition of the objectives and methodology (with supervisors' advice); 4) analysis and interpretation of results (with supervisors' advice); 5) writing of the document (with supervisors' advice).

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Abstract

Pollinators use a variety of floral cues to locate resources, but the relative importance of these different cues on pollinator foraging decisions is unclear. However, floral colour is undoubtedly one of the most important, as evidenced in previous works revealing the importance of floral colour on pollinator choices and determining flower visitor composition. Our purpose is to establish whether there is a relationship between flower colour and pollinator composition in natural communities. We measured the floral reflectance spectrum of 109 plant populations, from 300 to 700 nm, and divided it in four broad bands of 100 nm each. We found a phylogenetic signal on floral colour variables, and considered this in our analyses. Our results show a lack of concordance between colour spectra and pollinator assemblages. Nevertheless, this work proves that colour determines plant-pollinator relationships in some degree. Thus, particular pollinator groups display preferences for certain bands of the colour spectrum. These preferences match those expected from the pollination syndromes theory. Future work on this issue should combine floral colour along with other floral traits, to assess the importance of the combination of the main floral features for determining pollinator attraction and behaviour together.

Keywords: colour preferences, flower colour, phylogenetic signal, pollinator composition, pollination syndromes.

24 Introduction

25

26 Pollinators use floral cues such as odour, shape and colour to locate pollen-nectar sources [1].
27 These same cues allow pollinators to discriminate between different flower species and to maintain
28 flower constancy (or fidelity) within and between foraging bouts [2–4]. Although the relative
29 importance of these different cues on pollinator foraging decisions is unclear, some studies have
30 demonstrated that certain pollinators rely more strongly on colour than fragrance cues [5–7].
31 Pollinators may show innate preferences for certain colours [8,9]. In addition, colours may be used
32 by pollinators as signals of floral rewards, so that initial innate preferences may be modulated by
33 subsequent associative learning [10,11]. Thus, the role of colour in determining ultimate flower
34 choice will depend on the interaction between innate and learned colour preferences.

35

36 Pollinators have well-developed colour vision, which in most cases covers a wider range of the
37 spectrum than human vision. The vast majority of pollinators in temperate regions are insects, and
38 different visual models have been found for different groups [12,13]. The best studied species is the
39 honeybee (*Apis mellifera*), with a trichromatic visual system. Honeybee vision ranges between 300
40 and 700 nm with its three receptors peaking at the UV, blue and green parts of the spectrum
41 (corresponding to 344, 438 and 560 nm respectively). Most other studied insect groups share a
42 trichromatic visual system, but there are also known di-chromatic (certain flies and beetles) and
43 tetrachromatic models (butterflies) [14]. The trichromatic state is supposed to be ancestral, and the
44 loss or gain of photoreceptor types is secondary [14]. Insect colour vision appeared long before the
45 emergence of Angiosperms [15]. Therefore, it is believed that flowers tuned their visual signals to
46 the sensory system of pollinators, thus becoming as conspicuous as possible to them.

47 Traditionally, colour has been considered an essential cue in determining flower choice by
48 pollinators. To this extent, colour constitutes one of the main traits used in pollination syndrome
49 theory [16], according to which flowers have adopted different suites of traits as an adaptation to
50 attract different pollinator functional groups. Such a view implies that unrelated plant species
51 adapted to the same pollinators show convergence of floral traits, including colour. For example,
52 bee-visited flowers are expected to be blue or violet and moth flowers are expected to be light-
53 coloured [16]. In agreement with this view, flower colour is considered by some authors to be an
54 important predictor of pollinator group [17]. However, this view is highly controversial. Although
55 some pollinators are known to present innate floral colour preferences, these preferences do not
56 always match the colour expected from pollination syndromes [8]. In most cases, colour
57 preferences are displayed at a finer taxonomic level, such as genus or species, rather than at the
58 functional group level. A number of studies have analyzed pollinator choices among differently
59 coloured flower species within a genus or between colour morphs within a species. Some of these
60 studies have found evidence of the importance of flower colours in determining flower visitor
61 composition [18–21], but others have not [22,23]).

62

63 Even if pollinators show preferences for certain colours, and even if colours reflect reward levels,
64 the role of colour as a driver of ultimate flower choice in natural communities may be difficult to
65 establish for several reasons. First, most plants are pollinator generalists, attracting a wide array of
66 pollinators from different functional groups [24]. Second, in a natural setting, flower choice may be
67 influenced by the presence and abundance of other pollinators potentially competing for flower
68 resources [25]. Pollinators show plasticity at the individual level, and are known to switch plants in
69 response to changes in pollen-nectar levels irrespective of colour [26]. In sum, ultimate flower
70 choice may be highly context-dependent [27]. A handful of studies have analysed the influence of

71 floral colour on pollinator partitioning at the community level [20,28–31], and most of them
72 conclude that floral colour is an important cue. However, colours in these studies were categorized
73 from a human point of view. Waser et al. [24] analyzed pollinator partitioning in a community
74 context and characterized colour based on wavelength measures. In their analysis, the association
75 between floral colour and pollinator group composition narrowly failed significance. Clearly, further
76 research on this topic is badly needed.

77

78 The aim of this study is to establish whether there is a relationship between flower colour and
79 pollinator composition in natural communities. To do this, we studied four plant communities (85
80 species, 109 populations). The four communities were close to one another, and were exposed to
81 similar climatic conditions. Therefore, they shared the same regional pool of pollinator species and
82 pollinator groups. For each flower population, we measured the floral reflectance spectrum and
83 conducted pollinator surveys. The association between flower colour and pollinator composition
84 may be affected by phylogenetic constraints. Closely-related plant species may have similar colours
85 based on shared evolutionary history (e.g., due to similar floral pigments). Therefore, our analyses
86 account for phylogenetic relatedness. We ask the following questions: 1) Is flower colour in our
87 sample of flower species phylogenetically constrained? 2) Are plant species with similar colours
88 visited by similar arrays of pollinators? 3) Do different pollinator functional groups show
89 preferences for certain colours?

90

91

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93

94 **Material and Methods**

95

96 **Study area**

97

98 We conducted our study in four Mediterranean communities near Barcelona (NE Spain), whose
99 coordinates are given in Table 1. The first community (CA) was a herbaceous grassland, dominated
100 by *Hyparrhenia hirta* and *Brachypodium retusum*, located in Canet de Mar. The second community
101 (CO) was a grassland dominated by herbaceous (*Hyparrhenia hirta*, *Foeniculum vulgare*) along with
102 some shrubby species (*Cistus monspeliensis*, *Spartium junceum*), located in Collserola Natural Park.
103 The third community (GA) was a Mediterranean scrubland dominated by *Quercus coccifera*,
104 *Rosmarinus officinalis* and *Pistacia lentiscus*, located in the Garraf Natural Park. The last community
105 (PA) was another Mediterranean scrubland dominated by *Quercus coccifera*, *Ulex parviflorus* and
106 *Pistacea lentiscus*, located in Roques Blanques near El Papiol. Mean distance between the four sites
107 is 36 km (range: 10-66 km). The climate of the area is Mediterranean, with a strong sea influence.
108 Summers are dry and most precipitation occurs in spring and autumn. Weather conditions are very
109 similar across the four sites (Table 1).

110

111 **Pollinator assemblages**

112

113 Pollinator data were obtained throughout the main general flowering period of the area (February-
114 July). Very few species are in bloom during the summer drought. Each species was surveyed on
115 several days covering its entire flowering period. In CA and CO, pollinator surveys were carried out
116 by slowly walking through 25- to 50-m-long x 1-m-wide vegetation transects. This was done several
117 times, from early morning to late evening on every sampling day. In GA and PA, selected individuals

118 of each plant species in bloom were observed during 4-5-minute periods every 1-2 h. All insects
119 observed foraging on the flowers were identified by sight and recorded. Pollinators were assigned
120 to one of six taxonomic groups: bees, ants, wasps, coleopterans, dipterans and lepidopterans. From
121 these surveys, we characterized the pollinator assemblage (proportion of visits from each group) of
122 17 plant species in CA, 46 in CO, 25 in GA and 21 in PA (total: 109 populations, 85 species; some
123 species were present in more than one community).

124

125 **Flower colour measurement**

126

127 Flowers of each population were collected and transported with a portable cooler to the
128 laboratory, where they were temporarily placed in a cold room at 4°C. Flower reflectance spectra
129 were obtained using an USB4000 spectrometer with a USB-DT deuterium tungsten halogen source
130 (Ocean Optics Inc., Dunedin, FL, USA) connected to a computer running SpectraSuite (Ocean
131 Optics). The light spectrum analyzed ranged from 300 to 700 nm divided into 0.22 nm intervals, and
132 the spectrometer sensor was fixed at an angle of 45° from the measuring area. Petals were
133 mounted on an adhesive tape to obtain a relatively flat surface, thus minimizing reflectance
134 variability due to uneven distances between the petals and the sensor. For small flowers, we had to
135 use several petals from different flowers to cover the entire measuring area. Some flowers had
136 corollas with parts displaying two or more clearly different colours. In these cases, we took colour
137 measures of the different parts separately. To obtain a single colour measure for these flowers, we
138 then calculated a weighted mean (according to the surface occupied by each part in the corolla).
139 Most measures were taken on the day of flower collection, but a few were taken the day after.

140

141 Spectrometer outputs showed a certain amount of noise. To minimize noise and obtain smoother
142 reflectance spectra we used the software Avicol [32] to clean negative values and correct by
143 triangular smoothing. The latter correction is based on a floating mean with weights, with a window
144 size of 15 nm. Sample size was 4-5 measures per population, except for three populations for which
145 we could only obtain one measure. Different measures of a population were averaged to obtain a
146 single measure per population.

147

148 **Colour characterization**

149

150 Different pollinator groups are known to have different visual systems (different numbers of
151 photoreceptors and peak sensitivities). Using the reflectance spectrum we avoid biasing our results
152 towards any visual system. We characterized flower colour based on three different sets of
153 variables (I) First, we used the nine colour categories in Chittka et al.[33], representing the main
154 types of floral reflectance spectra in a natural reserve in Germany. Although they did not provide a
155 name for each category, we named floral colours as perceived by humans with the addition of UV
156 where necessary (Appendix 1). (II) Second, we obtained three commonly used [34] colour
157 properties: brightness, chroma and hue. Brightness, calculated as the sum of the reflectance values
158 over the entire spectrum, represents an achromatic value of reflection of the sample. Chroma,
159 calculated as the difference between the maximum and the minimum values of reflectance
160 between the average reflectance of the spectrum, is a measure of spectral purity. Hue, defined as
161 the wavelength with the highest value of reflectance, represents the degree to which a stimulus can
162 be described as similar to, or different from, stimuli that are described as red, green, blue, or
163 yellow. (III) Third, since the majority of the inflection points in floral reflectance spectra are located
164 near 400, 500 and 600 nm [35], we divided the floral reflection spectra in 4 broad bands of 100 nm

each (300-400, 400-500, 500-600, 600-700 nm) following Chittka et al. [33]. We then calculated the proportion of the reflectance spectra that was contained within each band (henceforth colour composition), obtained by dividing the brightness of each of the four bands by the total brightness of the sample. By using the proportion, instead of the raw values of brightness, of each band we avoid differences between chromatically equivalent spectra, i.e., spectra with the same shape, but differences in brightness. Each flower colour can then be characterized by the proportion of each of the four bands (henceforth colour band variables %350, %450, %550 and %650). Roughly, the first band corresponds to the UV part of the spectrum, the second to the blue-violet, the third to the green-yellow and the last one to the orange-red. It is important to bear in mind that some colours as perceived by humans may result from the combination of two or more bands. For example, white flowers, reflect from 400 to 700 nm; yellow flowers, reflect from 500 to 700 nm; and pink flowers, usually reflect in the blue and the red parts of the spectrum with a variable proportion in the yellow part (Appendix 1).

178

179 **Data analysis**

180

181 *Phylogenetic signal of colour variables*

182

183 To know whether colour was phylogenetically constrained, we constructed a phylogenetic tree of
184 the 85 species with Phylocom [36] with family names following the Angiosperm Phylogeny Group
185 classification [37]. We used the “bladj” function in Phylocom to achieve an ultrametric rooted tree.
186 Polytomies generated by the program were hand-resolved. All distances between families (assessed
187 as millions of years of divergence) and some distances between genera were obtained from the
188 database www.timetree.org [38]. Additional distances between genera and distances between

189 species were extracted from the literature [36,37(Fabaceae), 38(Cardueae), 39(Ericales)]. The
190 resulting tree (henceforth regional tree) is shown in Appendix 2.

191

192 We tested for the presence of phylogenetic signal in all colour variables (brightness, chroma, hue,
193 %350, %450, %550 and %650) with the Blomberg's K test [43]. This test compares the actual
194 distribution of traits on the obtained phylogenetic tree with those on trees generated under a
195 Brownian-motion model, and calculates significance through permutations. Blomberg's K performs
196 better than other indices due to its sensitivity to small changes in the distribution of the analyzed
197 traits [44]. K values lower than 1 suggest lack of phylogenetic signal of the trait, while values around
198 1 suggest an evolution of the trait fitting the Brownian motion model and thus occurrence of
199 phylogenetic signal. This test was done for each community separately and for the 85 species
200 lumped together. These analyses were performed with the "phylosig" function of the package
201 "phytools" developed by Liam Revell [45] for the R Statistical Software [46].

202

203 *Relationship between flower colour descriptors and pollinator composition*

204

205 To determine whether plants with similar colour properties attract similar pollinator assemblages,
206 we conducted partial Mantel tests between distance matrices of colour traits and of pollinator
207 assemblages, with a phylogenetic distance matrix as covariable. We conducted four partial Mantel
208 tests, one with colour composition (proportions of the four spectrum bands), one with brightness,
209 one with chroma, and one with hue. We used Bray-Curtis distances between pairs of species for
210 pollinator composition and colour composition, and Euclidean distances between species for
211 brightness, chroma, and hue. Tests were performed with the function "mantel.partial" in the
212 package "vegan" for R [47]. This analysis was done for each community separately and for the 109

213 populations lumped together. In the latter case, we included all populations in the regional tree and
214 assigned a low value of divergence (1000 years) to populations of the same species.

215

216 *Relationship between colour bands and pollinator groups*

217

218 Irrespective of a potential association between flower colour composition and pollinator
219 composition, certain pollinator groups could show a preference for certain colour bands. This would
220 not be detected in the Mantel test described above, which uses composite colour and pollinator
221 variables. To explore potential pollinator group – colour band associations, we performed Canonical
222 Correspondence Analyses (CCA) with the pollinator assemblage of each population and the four
223 colour bands. CCA is a multivariate analysis technique that detects the patterns of variation in a
224 dataset that can be best explained by combinations of the explanatory variables [48]. Again, this
225 was done for the four communities separately and together. CCAs were performed using the
226 function “CCorA” in the package “vegan”.

227

228 **Results**

229

230 The total amount of plant-pollinator interactions recorded was 32,315. Most of the pollinators
231 recorded were bees, accounting for nearly half of the overall interactions. The second most
232 frequent group were coleopterans, with 21% of the interactions, followed by ants (15%) and
233 dipterans (9%). The least important groups in terms of number of interactions were lepidopterans
234 and wasps (3.5% and 2.5%, respectively).

235

236 The four communities presented roughly similar patterns of floral colours (Appendix 1). The most
237 common floral colour in the four communities was pink (30 to 50% of the species), followed by
238 white (16-29%). UV-yellow flowers were also well represented (14-24%), although they were
239 lacking in GA. Yellow (12- 15%), purple (4- 9%) and green (4- 6%) flowers were less frequent.

240

241 *Phylogenetic signal in colour variables*

242

243 Blomberg's tests revealed significant phylogenetic signal in many of the colour traits tested (Table
244 2), that is, phylogeny appears to impose certain constraints on flower colour. All communities
245 showed phylogenetic signal in two or more colour variables. Variables showing phylogenetic signal
246 were similar in CO, GA, PA (Hue, %350, %450), but, surprisingly, CA showed a completely opposite
247 pattern. When analyzing the four communities together, all colour variables showed a strong
248 phylogenetic signal.

249

250 *Relationship between flower colour descriptors and pollinator composition*

251

252 Plants with similar flower colour characteristics did not attract similar pollinator assemblages in
253 three of the four communities or overall (Table 3). Only in one of the four communities, CA, there
254 was a significant association between pollinator composition and two colour properties (brightness
255 and hue).

256

257

258

259

Relationship between colour bands and pollinator groups

CCAs revealed clear associations between certain pollinator groups and certain colour bands (Fig. 1). Visual inspection of the resulting biplots revealed that some of these patterns were relatively consistent across the four communities (Table 4). Bees were positively linked to the UV-blue bands (350-450) of the spectrum, except in CA where they were not clearly related to any band. Ants, wasps and dipterans showed similar colour preferences. In general, they were positively associated with the yellow band (550). In some communities these groups were also associated to the UV-blue bands (350-450), but this relationship was sometimes positive and sometimes negative. Coleopterans and lepidopterans were associated to various colour bands, but these associations were not consistent across communities. When analyzing the four communities together, colour axes were well-defined and the four colour bands were distinctly separated (Fig. 1). The first axis was defined by yellow (550, negative values) and blue (450, positive values). The second axis was defined by the two extreme bands in the spectrum, UV (350) and red (650). Bees were positively associated with the blue and UV bands, ants, wasps and dipterans with the yellow band, coleopterans with the red band, and lepidopterans with the blue and red bands (Fig. 1; Table 4).

Discussion

Our results contrast with other studies analysing phylogenetical constraints on floral colour. Most of previous studies have found floral colour to be a highly labile trait, with low phylogenetic signal. Smith et al. (2008) analyzed variation in brightness, chroma and hue within a plant clade and found no phylogenetic signal. Other studies have analysed phylogenetic clustering on floral colour at the community level and have also found lack of phylogenetic signal [50–52]. In our study, phylogenetic

284 signal at the community level was weak but consistent in three of the four communities studied.
285 Phylogenetic constraints became clearly apparent in all the colour properties considered when
286 plants of all four communities were lumped together. These results indicate that colour changes fit
287 a Brownian-motion evolution model, implying that expected phenotypic differences between
288 species increase with phylogenetic distance. Some studies show that the capacity to produce
289 particular flower pigments has been lost in entire lineages [53], thus providing a biochemical
290 mechanism for the phylogenetic signal found in our study.

291

292 Notwithstanding phylogenetic constraints, the evolution of floral colour may be driven by selective
293 forces exerted by various factors. Traditionally, pollinators have been considered to be the main
294 drivers of floral colour evolution. This belief gained momentum when it became clear that insect
295 pollinator colour vision predated angiosperm radiation, implying that flowers evolved flower
296 colours so as to become as conspicuous as possible to pre-existing pollinators [14]. Later studies
297 suggest that many flower lineages have converged to display floral colours in areas of the spectrum
298 maximizing discrimination by Hymenoptera, both in the northern [33,35] and the southern
299 hemispheres [54]. Surprisingly, however, there is still very little evidence on the potential selective
300 pressures exerted by pollinators on floral colour [55].

301

302 In addition to pollinators, flower predators may also be important drivers of colour evolution. Then,
303 plants would face a trade-off between becoming more conspicuous to pollinators, while reducing
304 conspicuousness to herbivores and parasites, which usually are insects and therefore have similar
305 visual capabilities to pollinators [56]. In such a scenario, flower colour might be under opposing
306 selective pressures. A situation in which opposing selective forces have seemingly resulted in a
307 neutral balance has been described in *Raphanus* [57]. White and yellow *Raphanus* morphs were the

308 most visited by pollinators and were found to be under positive selection. However, these same
309 morphs were under negative selection by herbivores because they were the morphs with lowest
310 concentrations of chemical defences, thus counterbalancing pollinator selective pressure. Floral
311 colour may also evolve in response to abiotic factors, such as solar radiation and drought. Light
312 colours may be advantageous in dry environments, where dark colours may absorb radiation in
313 excess and raise petal temperatures, thereby increasing water loss [58].

314

315 If pollinators are an important selective force behind the evolution of floral colour, we would
316 expect a close association between flower colour properties and pollinators. Some studies have
317 shown that achromatic colour properties are not perceived by many insect groups [59]. That is,
318 insects can easily discriminate objects differing only slightly in reflectance spectrum, but cannot
319 discriminate between objects with similar reflectance spectrum differing only in brightness [59].
320 Therefore, we would expect the association between pollinator composition and colour to be
321 weaker for achromatic than chromatic variables. In agreement to these expectations, our results
322 indicate lack of correspondence between pollinator composition and brightness (achromatic
323 variable). As for chromatic variables, we obtained different results when we analysed colour
324 composition (lack of correspondence when considering all colour bands together) versus each
325 colour band separately (consistent relationships).

326

327 Several factors may contribute to explain the lack of correspondence between pollinator
328 composition and colour composition in a field situation. First, most of the plant species in our study
329 communities are pollinator generalists, and therefore are visited by various pollinator groups
330 [29,60]. This seems to be a general trend in temperate plant-pollinator communities [24,61].
331 Second, in a natural setting, pollinator innate colour preferences may be overridden by learned

332 preferences. It is believed that innate preferences may initially drive flower choice, but pollinators
333 do not restrict foraging to a unique colour pattern and may subsequently switch to other flower
334 colours if these are associated to greater floral rewards [26,62]. In other words, pollinator foraging
335 behaviour is flexible and preferences acquired through associative learning may override innate
336 preferences. This is not to say that innate preferences are not important. Pollinators may still have
337 a tendency to explore flowers with colour properties that can be easily detected by their visual
338 system. Of especial importance is the ability to detect flowers against the background (usually
339 green). Some studies have shown that pollinators forage more efficiently (are able to visit more
340 flowers per unit time) when they forage on flowers of certain colours [63].

341

342 Due to the lack of specialisation by both plants and pollinators, establishing pollinator-colour
343 associations may be easier when considering pollinator groups separately (instead of pollinator
344 assemblages) and specific colour bands (rather than the entire spectrum). This possibility was
345 explored with Canonical Correspondence Analysis (CCA), which yielded some notable coincidences
346 among communities. The overall CCA plot generated four quadrants clearly defined by the four
347 colour bands. The upper-left quadrant included UV-yellow flowers. Ants, dipterans and wasps were
348 linked to the yellow band, with dipterans also showing a slight affinity to the UV band. The upper-
349 right quadrant included purple flowers reflecting in the UV. Bees were strongly linked to this
350 quadrant. Interestingly, closer exploration of the bee data revealed differences between small (< 12
351 mm) and large (> 12 mm) bees (results not shown here). Small bees showed a stronger affinity to
352 the UV band, while large bees were mostly related to the blue band. The lower-right quadrant
353 included blue-red reflecting flowers (pink to human vision). Lepidopterans were tightly related to
354 this colour pattern. Finally, the lower-left quadrant included yellow flowers not reflecting in the UV
355 part of the spectrum. White flowers are related to negative values of the second axis of the biplot,

halfway between the yellow and the blue part of the spectrum along with reflection in the red part. Coleopterans were linked to white, slightly tending to the yellow quadrant. Our ability to detect a correspondence between certain pollinator groups and certain colour bands but not between pollinator composition and overall colour composition is congruent with results found on colour preferences by specific pollinators. Working with *Eristalis tenax*, Lunau [64] found that inexperienced flies land only on human-yellow stimuli, in the small range of wavelengths from 520 to 600 nm.

Interestingly, the above-mentioned pollinator-colour associations are quite consistent with colour preferences described in the pollination syndromes, according to which bees favour blue flowers, flies yellow and white, lepidopterans pink and red, beetles white and cream and wasps favour brown and yellow [9,16]. Other studies have also found similar consistencies. Waser *et al.* [24] explored the association between pollinator groups and bands of the colour spectrum in a natural community in Germany. Their results narrowly failed significance, but the observed trends were consistent with pollination syndromes. Other community studies have also found associations consistent with pollination syndromes [20,28,31]. The pollination syndrome has been highly contested and their ability to predict plant-pollinator associations has been strongly questioned [24,61,65]. However, a recent meta-analysis provides support for the existence of pollinator groups being associated to suites of floral traits [66], and pollination syndromes are still considered by many authors to be a valid framework for the evolution on plant-pollinator interactions [17,67].

Our study provides some evidence of pollinator-colour associations in natural communities, and highlights the fact that these associations are based on particular colour bands, rather than on entire colour patterns. However, flowers are not only coloured spots against a dull background. To

380 fully understand mutualistic relationships between plants and pollinators, flowers have to be
381 assessed as a whole. Future work should include other floral traits potentially influencing pollinator
382 attraction and behaviour (fragrance, shape, size, corolla depth), as well as floral rewards (pollen
383 and nectar). Such an integrative approach may reveal association between floral traits and certain
384 pollinator groups, and provide a more complete answer to the long-asked question of how do
385 pollinators choose flowers.

386

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388

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396

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398

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Table 1. Community descriptors including location, elevation, mean annual precipitation, and mean annual temperature.

Community	Location	Elevation (m.a.s.l.)	Mean annual precipitation (mm)	Mean annual temperature (°C)
CA	41° 35' N, 2° 34' E	50	590	16.1
CO	41° 24' N, 2° 6' E	280	630	14.8
GA	41° 16' N, 1° 55' E	330	580	15.7
PA	41° 27' N, 2° 0' E	150	628	15.4

Table 2. Results of Blomberg's K tests for the presence of phylogenetic signal in colour descriptors in the four study communities and overall (data of the four communities lumped together). Significant results ($p < 0.05$) in bold. Marginally significant results ($0.05 < p < 0.1$) in italics.

Community	Brightness		Chroma		Hue		UV band (350%)		Blue band (450%)		Yellow band (550%)		Red band (650%)	
	K	P	K	P	K	P	K	P	K	P	K	P	K	P
CA	0.73	<i>0.07</i>	0.96	0.01	0.66	0.20	0.44	0.50	0.51	0.31	0.77	0.05	1.14	0.00
CO	0.35	0.62	0.43	0.37	0.71	0.03	0.45	0.26	0.58	0.02	0.39	0.45	0.43	0.26
GA	0.26	0.33	0.21	0.53	0.74	<i>0.06</i>	0.90	0.02	0.60	0.04	0.20	0.45	0.17	0.65
PA	0.34	0.22	0.33	0.27	0.88	0.01	0.73	0.03	0.73	0.02	0.28	0.31	0.19	0.59
CA+CO+GA+PA	0.79	0.00	0.57	0.00	0.81	0.00	0.65	0.00	0.63	0.00	0.89	0.00	0.72	0.00

Table 3. P-values of phylogenetically-controlled Mantel tests between colour descriptors and pollinator composition in the four communities and overall (data of the four communities lumped together). Significant results ($p < 0.05$) in bold.

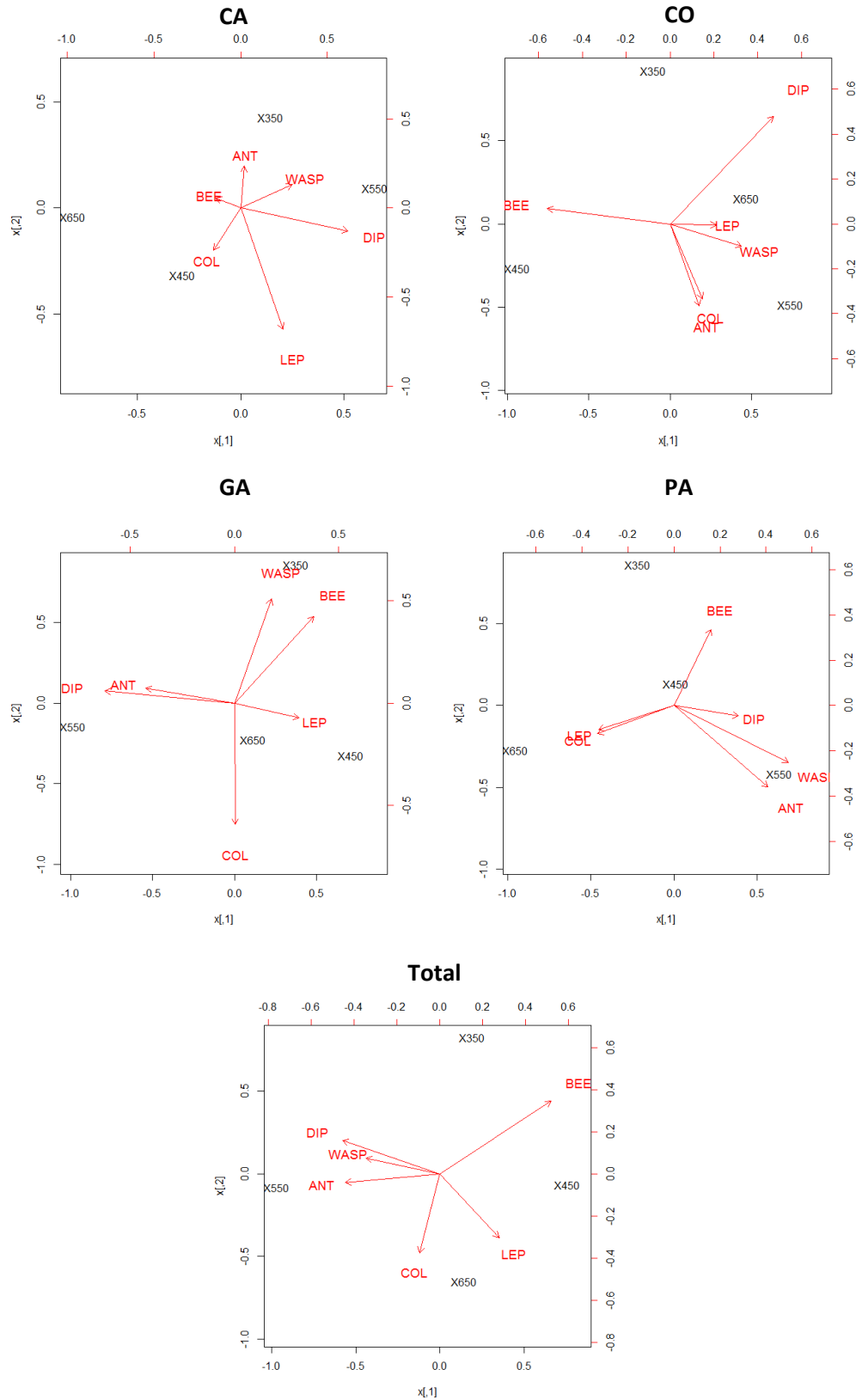
Community	Brightness	Chroma	Hue	Colour composition*
CA	0.016	0.130	0.038	0.303
CO	0.558	0.952	0.188	0.378
GA	0.605	0.655	0.560	0.152
PA	0.186	0.537	0.751	0.200
CA+CO+GA+PA	0.113	0.896	0.553	0.308

* Proportion of UV, blue, yellow, and red bands.

Table 4. Positive and negative relationships between pollinator groups and the four bands of the colour spectrum (350, 450, 550, 650 nm; UV, blue, yellow, and red, respectively) in the four study communities and overall (data of the four communities lumped together), estimated visually from the CCA biplots of Fig. 1.

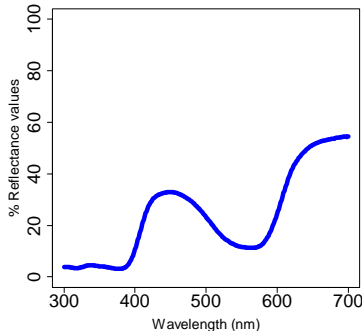

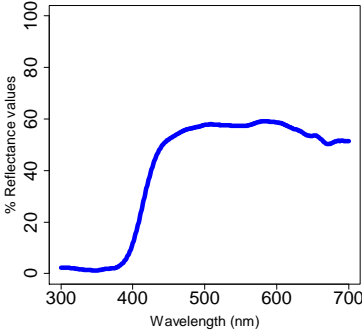

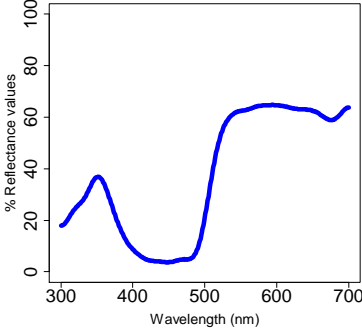

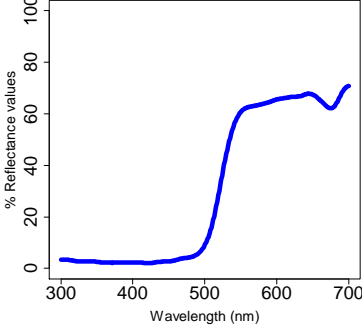

	Type of relationship	CA	CO	GA	PA	CA+CO+GA+PA
BEES	(+)		350, 450	350, 450	350	350, 450
	(-)		500			
ANTS	(+)	350	550	550	550	550
	(-)		350	450	350	450
WASPS	(+)	550	550	350	550	550
	(-)		450			450
DIPTERANS	(+)	550	350, 550	550	550	550
	(-)			450		450
COLEOPTERANS	(+)	450	550		650	650
	(-)		350	350		350
LEPIDOPTERANS	(+)	450, 550		450	650	450, 650
	(-)	350	450	550	550	550

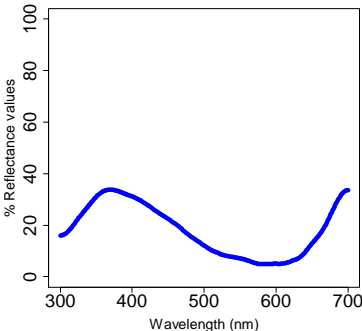

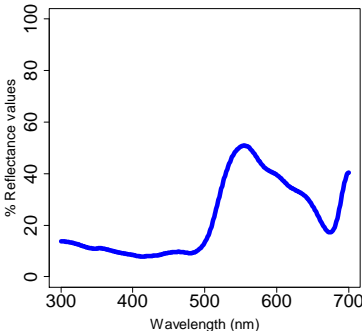

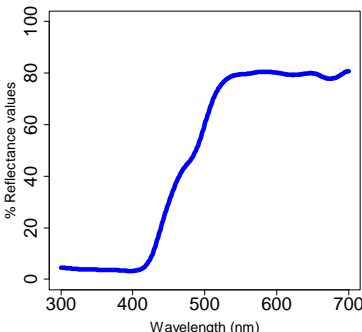

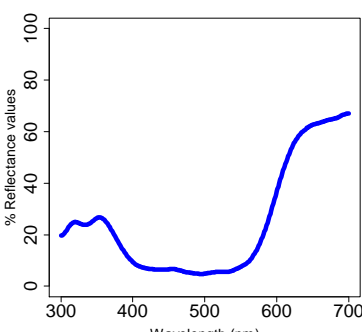

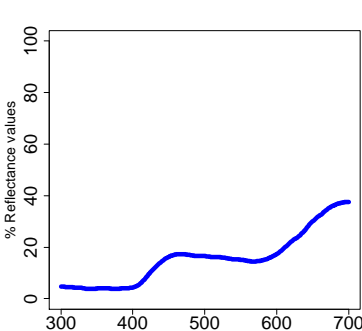

Figure 1. CCA biplots between pollinator groups and four bands of the colour spectrum (X350, X450, X550, X650; corresponding to UV, blue, yellow, and red, respectively) in each of the four communities separately (CA, CO, GA, PA) and for the four communities together (Total). Bee = bees, Ant = ants, Wasp = wasps, Dip = Dipterans, Col = coleopterans, Lep = lepidopterans.



Appendix 1. Flower colour categories

Table S1. Flower colour categories (following Chittka et al. 1994), with an example species of each model and the proportion of each category in each of the four communities (CA, CO, GA, PA).

Colour category	Reflectance spectra	Species	CA	CO	GA	PA
Pink		<i>Cistus albidus</i> L.	29%	30%	52%	38%
						
White		<i>Dorycnium pentaphyllum</i> L.	18%	22%	16%	29%
						
UV-yellow		<i>Sonchus tenerrimus</i> L.	24%	17%	0%	14%
						
Yellow		<i>Ranunculus gramineus</i> L.	12%	15%	12%	14%
						

Purple		<i>Anagallis arvensis</i> L.			
					
		6%	9%	4%	5%
Green		<i>Euphorbia flavicoma</i> DC.			
					
		6%	4%	4%	0%
White-yellow		<i>Biscutella laevigata</i> L.			
					
		0%	2%	8%	0%
UV-red		<i>Papaver rhoeas</i> L.			
					
		6%	0%	0%	0%
Brown		<i>Orobanche latisquama</i> F.W. Schultz			
					
		0%	0%	4%	0%

Appendix 2. Phylogeny

Figure S1. Phylogenetic tree of the 85 species surveyed.

