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3 **Pollinators show flower colour preferences but flowers with similar colours do not**  
4 **attract similar pollinators.**

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10

11 **Running title:** Flower colour and pollinator composition

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1 **Abstract**

2

3 *Background and aims.* Colour is one of the main floral traits used by pollinators to locate  
4 flowers. Although pollinators show innate colour preferences, the view that the colour of a  
5 flower may be considered an important predictor of its main pollinators is highly  
6 controversial because flower choice is highly context-dependent, and initial innate preferences  
7 may be overridden by subsequent associative learning. Our objective is to establish whether  
8 there is a relationship between flower colour and pollinator composition in natural  
9 communities.

10

11 *Methods.* We measured the flower reflectance spectrum and pollinator composition in four  
12 plant communities (85 plant species represented by 109 populations, and 32305 plant-  
13 pollinator interactions in total). Pollinators were divided into six taxonomic groups: bees,  
14 ants, wasps, coleopterans, dipterans and lepidopterans.

15

16 *Key results.* We found consistent associations between pollinator groups and certain colours.  
17 These associations matched innate preferences experimentally established for several  
18 pollinators and predictions of the pollination syndrome theory. However, flowers with similar  
19 colours did not attract similar pollinator assemblages.

20

21 *Conclusions.* The explanation for this paradoxical result is that most flower species are  
22 pollination generalists. We conclude that although pollinator colour preferences seem to  
23 condition plant-pollinator interactions, the selective force behind these preferences has not  
24 been strong enough to mediate the appearance and maintenance of tight colour-based plant-  
25 pollinator associations.

1

2 *Key words:* colour preferences, floral colour, floral reflectance spectra, generalization, floral  
3 traits, phylogenetic signal, plant-pollinator interactions, pollinator assemblage, pollination  
4 syndromes.

5

## 6 **INTRODUCTION**

7

8 Pollinators use flower traits, such as odour, shape, size and colour, as cues to locate pollen-  
9 nectar sources and discriminate between different flower species (Chittka and Raine 2006).  
10 Although ultimate flower choice undoubtedly depends on a combination of stimuli, various  
11 studies have demonstrated that some pollinators rely strongly on colour to make their foraging  
12 decisions (Dafni et al. 1990; Heiling et al. 2003; Ômura and Honda 2005; Dötterl et al. 2014).

13

14 Diurnal pollinators have well-developed colour vision, which in most cases covers a wider  
15 range of the spectrum than human vision. The vast majority of pollinators in temperate  
16 regions are insects, and different visual models have been developed for different groups  
17 (Chittka 1992; Troje 1993). The best studied species are the western honey bee, *Apis*  
18 *mellifera*, and the bumblebee *Bombus terrestris*, which share a similar trichromatic system.  
19 The visual system of the honey bee, which is usually used as a model for all bees, ranges  
20 between 300 and 700 nm, with three types of photoreceptors peaking in the UV, blue and  
21 green regions of the spectrum (corresponding to 344, 438 and 560 nm, respectively). Most  
22 other insects studied so far also have a trichromatic system, but there are also species with  
23 dichromatic (certain flies and coleopterans) and tetrachromatic (mostly butterflies) systems  
24 (Briscoe and Chittka 2001). The trichromatic state is supposed to be ancestral, and the loss or  
25 gain of photoreceptor types would be secondary (Briscoe and Chittka 2001). Insect colour

1 vision appeared long before the emergence of the angiosperms (Chittka 1997). For this  
2 reason, it is believed that flowers tuned their visual signals to the sensory system of  
3 pollinators, thus becoming as conspicuous and attractive as possible to them. In support of  
4 this view, some pollinator species have been found to show innate preferences for certain  
5 colours (Lunau and Maier 1995; Raine et al. 2006; Raine and Chittka 2007; Willmer 2011).

6

7 To this extent, colour constitutes one of the main traits used in pollination syndrome theory  
8 (Faegri and Van der Pijl 1979). According to this theory, unrelated plant species adapted to  
9 the same pollinators should show convergence of floral traits, including colour. For example,  
10 bee-visited flowers are expected to be blue or violet, and beetle-visited flowers are expected  
11 to be white or cream (Faegri and Van der Pijl 1979). Accordingly, some studies have shown  
12 that colour changes among related plant species or between populations within a same species  
13 are sometimes accompanied by changes in pollinator composition (Bradshaw and Schemske  
14 2003; Rodríguez-Gironés and Santamaría 2004; Wolfe and Sowell 2006). Other studies have  
15 shown pollinator colour preferences through experimental manipulations of flower colour  
16 (Campbell et al. 2010). Ultimately, flower colour could be considered an important predictor  
17 of pollinator group (Fenster et al. 2004). However, other studies do not support this view.  
18 Colour preferences predicted by pollination syndromes are not always matched by innate  
19 colour preferences (Lunau and Maier 1995), which in many cases are displayed at a finer  
20 taxonomic level, such as genus or species, rather than functional group (Raine et al. 2006).  
21 Other studies report that changes in flower colour are not always associated with distinct  
22 pollinator assemblages (Cooley et al. 2008).

23

24 Even if pollinators show innate preferences for certain colours, actual flower choice in natural  
25 communities may not reflect these preferences for two reasons. First, in a natural setting,

1 flower choice may be influenced by the presence and abundance of other co-pollinators  
2 potentially competing for flower resources (Waser 1983; Lázaro et al. 2009). Second, colours  
3 may also be used by pollinators as signals of floral rewards, so that initial innate preferences  
4 may be modulated by subsequent associative learning (Gumbert 2000; Goyret et al. 2008).  
5 Individual pollinators show remarkable plasticity and are known to switch plants in response  
6 to changes in pollen or nectar levels (Heinrich 1979). Thus, the role of colour in determining  
7 ultimate flower choice will depend on the interaction between innate and learned preferences.  
8 A handful of studies have analysed the influence of floral colour on pollinator partitioning in  
9 plant communities, and most of them conclude that floral colour is an important cue (McCall  
10 and Primack 1992; Bosch et al. 1997; Hingston and McQuillan 2000; Wolfe and Sowell 2006;  
11 Lázaro et al. 2008). However, colours in these studies were categorized as perceived by  
12 humans. Waser et al. (1996) characterized colour based on wavelength measures that were  
13 incorporated into the visual model of the honeybee to generate colour categories. In their  
14 study, the association between floral colour category and pollinator composition narrowly  
15 failed significance. Clearly, further research on this topic is needed.

16

17 The aim of this study is to establish whether there is a relationship between flower colour and  
18 pollinator visitation in natural communities. To do this, we studied the flower reflectance  
19 spectrum and flower visitor assemblages in four plant communities comprising 85 species and  
20 109 plant populations. The four communities are located in the same geographic region, and  
21 therefore have similar climatic conditions and share the same regional pool of pollinators.  
22 Because the association between flower colour and pollinator composition may be affected by  
23 phylogeny, our analyses account for phylogenetic relatedness. We ask two questions: 1) Do  
24 different pollinator groups show preferences for certain flower colours? 2) Do plant species  
25 with similar colours attract similar pollinator assemblages? Although apparently similar, these

1 two questions address colour-pollinator relationships from two different and complementary  
2 perspectives. The first question addresses plant-pollinator relationships from the pollinators'  
3 point of view, and the second from the plants' perspective. The existence of colour  
4 preferences by different pollinator groups does not necessarily imply that plants with the same  
5 colour attract similar pollinator assemblages. For example, dipterans may preferentially visit  
6 flowers with high levels of reflectance in the yellow region of the spectrum, but different  
7 yellow flowers may be visited by different pollinator assemblages. The distinction is  
8 important because pollination syndrome theory relies on the assumption that plants with  
9 similar traits have evolved to attract similar pollinators.

10

## 11 **MATERIALS AND METHODS**

12

### 13 **Study area**

14

15 We conducted our study in four Mediterranean communities near Barcelona (NE Spain),  
16 whose geographic coordinates are given in Table 1. Distance between sites ranged from 10 to  
17 66 km. The climate is Mediterranean, with a strong sea influence. Summers are dry and most  
18 precipitation occurs in spring and autumn. Weather conditions are very similar across the four  
19 sites (Table 1). We studied the vast majority of the entomophilous species in each community,  
20 without selecting species based on their flower traits or pollinator visitation patterns (Table 1).  
21 Therefore our plant sample was unbiased and representative of the flora of the region. Some  
22 species were present in more than one community. In these cases, we sampled flower colour  
23 and pollinator composition of each population separately (total: 85 species, 109 populations).

24

### 25 **Pollinator assemblages**

1

2 Pollinator data were obtained throughout the main general flowering period in the study area  
3 (February-July). Very few species bloom in August. Each species was surveyed on several  
4 days covering its entire flowering period. In two communities (CA and CO), pollinator  
5 surveys were carried out by slowly walking along 25- or 50-m-long x 1-m-wide vegetation  
6 transects. This was done several times throughout the day, from early morning to evening. In  
7 the other two communities (GA and PA), selected individuals of each plant species in bloom  
8 were observed during 4- or 5-minute periods every 1-2 h. All insects observed contacting the  
9 reproductive organs of flowers were identified by sight and recorded. Pollinators were  
10 assigned to one of six taxonomic groups: bees, ants, wasps, coleopterans, dipterans and  
11 lepidopterans. From these surveys, we characterized the pollinator assemblage of each plant  
12 species in each of the four communities as the proportion of visits from each pollinator group.

13

#### 14 **Flower colour measurement**

15

16 Flowers of each species were collected and transported with a portable cooler to the  
17 laboratory, where they were temporarily placed in a cold room at 4°C. Flower reflectance  
18 spectra were measured using an USB4000 spectrometer with a USB-DT deuterium tungsten  
19 halogen source (Ocean Optics Inc., Dunedin, FL, USA) connected to a computer running  
20 SpectraSuite (Ocean Optics). The light spectrum analysed ranges from 300 to 700 nm divided  
21 into 0.22 nm intervals, and the spectrometer sensor was fixed at an angle of 45° from the  
22 measuring area. Petals were mounted on an adhesive tape to obtain a flat surface, thus  
23 minimizing reflectance variability due to uneven distances between the petals and the sensor.  
24 For small flowers, we had to use several petals from different flowers to cover the entire  
25 measuring area. Some species had corollas displaying two or more clearly different colours.

1 When these different colours were largely separated (for example, some Fabaceae in which  
2 the keel is clearly differently coloured than the wings and the standard), we took colour  
3 measures of the different parts separately and calculated a weighted mean (according to the  
4 surface occupied by each part in the corolla), thus obtaining a single colour value. In corollas  
5 showing only small colour markings such as nectar guides or small dots, a single measure was  
6 taken.

7

8 Spectrometer readings incorporate a certain degree of noise. We used Avicol (Gomez 2006)  
9 to clean this noise, thus smoothening the reflectance spectra. This correction (triangular  
10 smoothing) is based on a floating mean with weights and a window size of 15 nm. We  
11 measured 4-5 flowers per species except for three species for which we could only obtain one  
12 measure. Different measures were averaged to obtain a single value per species and  
13 community.

14

## 15 **Colour characterization**

16

17 We characterized flower colour using two sets of variables: colour descriptors and “colour  
18 composition”. Colour descriptors (brightness, chroma and hue) are physical properties of  
19 colours extracted directly from the reflectance spectrum (Endler 1990; Smith 2014).  
20 Brightness is a measure of the total intensity of light reflected by a surface; chroma is a  
21 measure of purity or saturation of a colour, and it is a function of how rapidly intensity  
22 changes with wavelength; hue, represents the usual meaning of colour (such as red, pink,  
23 yellow, etc.), and it is a function of the shape of the spectrum. These three colour descriptors  
24 were calculated following Smith (2014). To obtain “colour composition”, we divided the  
25 reflectance spectrum into 4 broad bands of 100 nm each (300-400, 400-500, 500-600, 600-



1 700 nm; Chittka *et al.* 1994). Breakpoints delimiting bands were chosen assuming that the  
2 majority of the inflection points in floral reflectance spectra are located near 400, 500 and 600  
3 nm (Chittka and Menzel 1992). Roughly, the first band corresponds to the UV part of the  
4 spectrum, the second to the blue-violet, the third to the green-yellow and the fourth to the  
5 orange-red (henceforth U, B, Y and R bands). We calculated the proportion of the reflectance  
6 spectrum attributable to each band, obtained by dividing the brightness of each band by the  
7 total brightness of the sample. By using the proportion, instead of the raw values of brightness  
8 of each band, we avoid differences between chromatically equivalent spectra, i.e., spectra  
9 with the same shape but different brightness (Endler 1990). It is important to note that flower  
10 colours result from the combination of reflectance levels across the spectrum. For example,  
11 white flowers reflect from 400 to 700 nm and yellow flowers from 500 to 700 nm. **Lilac-pink**  
12 **flowers reflect in the blue and red regions with a variable proportion in the yellow region, and**  
13 **purple flowers reflect in the UV, blue and red regions of the spectrum** (see Table S1  
14 **[Supplementary Information]).**

15

## 16 **Data analysis**

17

### 18 *Phylogenetic signal of colour variables*

19

20 To establish whether colour was phylogenetically constrained, we built a phylogenetic tree of  
21 the 85 species using Phylocom (Webb et al. 2008), with family names following the  
22 Angiosperm Phylogeny Group classification (The Angiosperm Phylogeny Group 2009). We  
23 used the “bladj” function in Phylocom to achieve an ultrametric rooted tree. Polytomies  
24 generated by the program were hand-resolved. All distances between families (assessed as  
25 millions of years of divergence) and some distances between genera were obtained from the

1 database [www.timetree.org](http://www.timetree.org) (Hedges et al. 2006). Additional distances between genera and  
2 distances between species were extracted from the literature [Allan and Porter, 2000; Lavin *et*  
3 *al.*, 2005 (Fabaceae); Barres *et al.*, 2013 (Cardueae); Bremer, 2009 (Ericales); Guzmán and  
4 Vargas, 2009 (Cistaceae); Mansion *et al.*, 2009 (Boraginaceae); Couvreur *et al.*, 2010  
5 (Brassicaceae); Bendiksby *et al.*, 2011; Drew and Sytsma, 2012 (Lamiaceae); Koopman and  
6 Baum, 2008 (Malvaceae); Ruiz-Sanchez *et al.*, 2012 (Papaveraceae); Riina et al. 2013  
7 (Euphorbiaceae); Banasiak et al. 2013 (Apiaceae)]. The resulting tree is shown in Figure S1  
8 **[Supplementary Information]**.

9  
10 We tested for the presence of phylogenetic signal in brightness, chroma and hue with the  
11 Blomberg's K test (Blomberg et al. 2003) using the “phylosig” function of the R package  
12 “phytools” (Revell 2012). This was done for each community separately and for the 85  
13 species together. We also tested for the presence of phylogenetic signal in flower colour  
14 composition. To do this, we divided the colour spectrum into 40 bands of 10 nm each and  
15 obtained the mean reflectance value of each band. We used 40 bands instead of the 4 bands  
16 used to define colour composition to increase the resolution of this analysis. We then used the  
17 generalized K statistic described by Adams (2014),  $K_{mult}$ , specifically developed for high-  
18 dimensional multivariate data. This analysis was performed with the function “physignal” of  
19 the package “geomorph” for R (Adams and Otarola-Castillo 2013). Again, we conducted this  
20 analysis for each community separately and for the 85 species together.

21

## 22 *Association between pollinator groups and regions of the colour spectrum*

23

24 We explored possible preferences of the different pollinators groups for certain regions of the  
25 colour spectrum by means of Canonical Correspondence Analyses (CCA) including the

1 pollinator assemblage of each population and relative brightness of the four above-mentioned  
2 colour bands (U, B, Y and R). This was again done for the four communities separately and  
3 for all species together. CCAs were performed using the function “CCorA” in the R package  
4 “vegan” (Oksanen et al. 2015).

5

### 6 *Relationship between flower colour and pollinator assemblages*

7

8 To test whether flowers with similar colours had similar pollinator assemblages, we  
9 conducted a partial Mantel test between distance matrices of colour composition (combination  
10 of the proportion of the four colour bands) and pollinator assemblages, including the  
11 phylogenetic distance matrix as covariable. We then followed the same approach to test the  
12 association between pollinator assemblages and the other colour variables (brightness,  
13 chroma, hue). We used Bray-Curtis distances between pairs of species for pollinator  
14 composition and colour composition, and Euclidean distances between species for brightness,  
15 chroma and hue. These analyses were done for each community separately and for the four  
16 communities lumped together. In the latter case, because the same species may attract  
17 different pollinators in different communities, we maintained all 109 populations. For plant  
18 species present in more than one community, we assigned a very low value of divergence  
19 (1000 years) between conspecific populations. These analyses were performed with the  
20 function “mantel.partial” in the R package “vegan”.

21

## 22 **RESULTS**

23

24 We recorded 32,315 plant-pollinator interactions. The number of interactions recorded in each  
25 community ranged from 3505 (CA) to 13673 (CO) (Table S2 [**Supplementary**

1 **Information**]). Mean number of interactions per population was 206 in CA (range: 43 to  
2 1454); 297 in CO (range: 42 to 1911); 364 in GA (range: 26 to 1730) and 307 in PA (range:  
3 31 to 1359). Overall, 88.5 % of the populations surveyed had more than 50 recorded  
4 interactions. Most of the pollinators recorded were bees, accounting for 48.9% of the flower  
5 visits. The second most frequent group was coleopterans (21.5% of the interactions), followed  
6 by ants (14.7%), dipterans (8.8%), lepidopterans (3.5%) and wasps (2.5%). Bees and  
7 coleopterans were the two most abundant groups in all four communities, except CA, which  
8 was largely dominated by ants (Table S2 [**Supplementary Information**]).

9  
10 The four communities showed a high degree of similarity in flower colours. The most  
11 common floral colour in the four communities was lilac-pink (30-50% of the species),  
12 followed by white (16-29%) (Table S1 [**Supplementary Information**]). UV-yellow flowers  
13 were also well represented (14-24%), although they were lacking in GA. Yellow (12- 15%),  
14 purple (4- 9%) and green (4- 6%) flowers were less frequent. The association between colour  
15 categories and pollinator composition is shown in Table S3 [**Supplementary Information**].

16

### 17 *Phylogenetic signal of colour variables*

18

19 All colour descriptors considered showed significant phylogenetic signal when the four  
20 communities were pooled together (Table 2). In most cases, however, significance was lost  
21 when the communities were analysed separately, possibly due to small sample sizes. In all  
22 cases,  $K$  and  $K_{mult}$  values were lower than 1, indicating that related species were less similar  
23 than expected under the Brownian motion evolution model.

24

### 25 *Association between pollinator groups and regions of the colour spectrum*

1

2 CCAs revealed clear associations between certain pollinator groups and certain colours (Figs.  
3 1 and 2). Visual inspection of the resulting biplots revealed that some of these patterns were  
4 relatively consistent across the four communities (Fig. 1, Table 4).

5

6 Overall, bees were associated with purple flowers, and ants with UV-Yellow and green  
7 flowers. Wasps and dipterans were mostly associated with UV-yellow flowers. Coleopterans  
8 were associated with white and yellow flowers, and lepidopterans with pink flowers (Table 4,  
9 Fig. 2).

10

#### 11 *Relationship between flower colour and pollinator assemblages*

12

13 Results from the partial Mantel test showed no significant association between flower colour  
14 and pollinator assemblages (Table 3). Plants with similar colour descriptors, including colour  
15 composition, did not attract similar pollinator assemblages in any of the communities, and  
16 similar results were obtained when data from the four communities were lumped together.

17

## 18 **DISCUSSION**

19

20 We traced the phylogenetic signal of floral colour under a neutral evolution model (Brownian  
21 motion). In this model, changes occur slowly and gradually along phylogenetic branches, and  
22 character similarity between species is related to phylogenetic proximity (Blomberg et al.  
23 2003). Phylogenetic signal for flower colour was mostly lacking when we analysed the four  
24 communities separately, but when all species were lumped together all colour properties  
25 measured showed significant phylogenetic signal (related species were more similar in colour

1 than expected from a random association between species and colours). Notwithstanding this  
2 influence of phylogeny on flower colour, the fact that all K values were lower than 1 indicates  
3 that similarity among related species tends to be lower than expected under Brownian motion.  
4 Most previous studies have found lack of phylogenetic signal in flower colour (Smith et al.  
5 2008; Arnold et al. 2009; McEwen and Vamosi 2010). However, one recent study on four  
6 Himalayan plant communities (Shrestha et al. 2014) obtained results similar to ours  
7 (significant phylogenetic signal with K values  $< 1$ ). A corollary of our phylogenetic analysis  
8 is that the different colour categories considered are widely spread across the phylogenetic  
9 tree. This outcome is in agreement with reports showing important changes in flower colour  
10 as a result of relatively simple genetic changes (Rausher 2008).

11

12 We analysed the colour preferences of the pollinator groups present in our communities.  
13 Results of the CCAs showed notable coincidences among communities and provided  
14 compelling evidence for the association between pollinator groups and certain colours. These  
15 associations were less clear in the communities with lower number of plant species, but were  
16 very apparent when the results of all 109 populations were lumped together (Fig. 2).  
17 Importantly, these associations are notably congruent with innate colour preferences found for  
18 specific pollinators (Lunau and Maier 1995; Willmer 2011). For example, inexperienced  
19 *Eristalis tenax* hover flies were found to land only on artificial flowers with yellow stimuli in  
20 the range of 520 to 600 nm (Lunau and Maier 1995). Naive honey bees showed preference for  
21 stimuli reflecting between 410 and 530 nm (Giurfa et al. 1995), and bumblebees preferred  
22 artificial flowers reflecting in one or two of the three colour bands corresponding to their  
23 photoreceptor types (ultraviolet, blue, green) (Lunau 1992). *Papilio demoleus* butterflies  
24 selected blue, green, and red colours while neglecting yellow (Ilse and Vaidya 1956) and  
25 *Pieris brassicae* preferred artificial flowers reflecting at 450 and 600 nm (Scherer and Kolb

1 1987). The hawkmoth *Macroglossum stellatarum* was shown to have a strong preference for  
2 wavelengths around 440 nm and a weaker preference for wavelengths of 540 nm (Kelber  
3 1997; Telles et al. 2014).

4

5 Importantly, the above-mentioned pollinator-colour associations found in our study are  
6 consistent with colour preferences described in pollination syndromes, according to which  
7 bees favour blue, flies yellow and white, lepidopterans pink and red, coleopterans white and  
8 cream and wasps favour brown and yellow flowers (Faegri and Van der Pijl 1979; Willmer  
9 2011). Waser *et al.* (1996) explored the association between pollinator groups and bands of  
10 the colour spectrum in a natural community in Germany. Their results narrowly failed  
11 significance, but the observed trends were also consistent with associations predicted by  
12 pollination syndromes. Other community studies using more subjective (human-perceived)  
13 colour measures also found associations that were more or less consistent with pollination  
14 syndromes (McCall and Primack 1992; Wolfe and Sowell 2006; Lázaro et al. 2008).

15

16 However, our results cannot be interpreted as supporting pollination syndrome theory because  
17 flowers with similar colours did not attract similar pollinators (Table 3). That is, pollinator  
18 preferences are a necessary but not sufficient condition for the establishment of colour-based  
19 pollination syndromes. The lack of relationship between flower colour and pollinator  
20 composition can be explained by the influence of other traits, or trait combinations, on flower  
21 choices. Some studies have found phenology (Herrera 1988), pollen/nectar rewards (Bosch et  
22 al. 1997), or corolla depth (Stang et al. 2007) to be the main drivers of pollinator partitioning  
23 in flower communities. A second factor contributing to the lack of relationship between  
24 flower colour and pollinator composition is pollinator generalisation. As in most temperate  
25 systems (Herrera 1996; Waser et al. 1996), the majority of plant species in our study are

1 pollinator generalists (with two or more pollinator groups each accounting for more than 10%  
2 of the observed visits), and therefore cannot be readily assigned to a given pollination  
3 syndrome. Even then, we could have found a correspondence between certain flower colours  
4 and pollinator assemblages (rather than single pollinator groups), but this was clearly not the  
5 case.

6

7 Our results are in agreement with the findings of Rezende et al. (2007), who found that plant-  
8 pollinator networks are more strongly associated to animal than to plant phylogeny. That is,  
9 related animal species are more likely to share host plants than related plant species are to  
10 share pollinator visitors. This result is attributed to pollinator mobility (Bascompte and  
11 Jordano 2007), which allows pollinators to play a more active role in the definition of  
12 interaction identity.

13

14 Recently, Renoult et al. (2014) have used a novel approach to analyze the association between  
15 flower colour and generalization in pollinators. They argue that pollinators should  
16 preferentially visit flowers that are most conspicuous to them, and that colour perception  
17 should be analyzed as a measure of colour conspicuousness between emitters (flowers) and  
18 receivers (insects). They measure flower and background colour, and use evidence from  
19 various sources to assign one of eleven described visual systems to each pollinator species.  
20 Then they calculate 'colour matching' for each plant-pollinator pair as a measure of how a  
21 floral colour stands out from the environment for each visual system. They find that colour  
22 matching influences generalization in one of three regions studied. However, they also find  
23 that other factors (spatiotemporal overlap and co-abundance between plants and pollinators)  
24 play a greater role as determinants of plant generalization. The results of Renoult et al. (2014)  
25 in general, and the strong influence of these other factors in particular, support our conclusion



1 that even though pollinator colour preferences seem to be conditioning plant-pollinator  
2 interactions, the selective force behind these preferences may not been strong enough to  
3 mediate the appearance and maintenance of tight colour-based plant-pollinator associations.

4

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14

15

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1

2

1 **Figure 1.** CCA biplots of pollinator groups and bands of the colour spectrum (colour squares)  
2 corresponding to UV, blue, yellow, and red, respectively, in each of the four communities  
3 (CA, CO, GA, PA). Each dot represents a plant population and dot colour corresponds to  
4 flower colour categories as shown in the legend (see Table S1 [**Supplementary Information**]  
5 for example spectra of each category). Bee = bees, Ant = ants, Was = wasps, Dip = dipterans,  
6 Col = coleopterans, Lep = lepidopterans.

7

1 **Figure 2.** CCA biplot of pollinator groups and bands of the colour spectrum (colour squares)  
2 corresponding to UV, blue, yellow, and red, respectively (data from the four communities  
3 lumped together). Each dot represents a plant population and dot colour corresponds to the  
4 flower colour categories as shown in the legend (see Table S1 [**Supplementary Information**]  
5 for example spectra of each category). Bee = bees, Ant = ants, Was = wasps, Dip = dipterans,  
6 Col = coleopterans, Lep = lepidopterans.



1 **Table 1.** Descriptors of the four communities studied, including vegetation type, dominant species, location, elevation, mean annual  
 2 precipitation, mean annual temperature, and number of species sampled.

3

<b>Community</b>	<b>Vegetation</b>	<b>Dominant species</b>	<b>Location</b>	<b>Elevation (m.a.s.l.)</b>	<b>Mean annual precipitation (mm)</b>	<b>Mean annual temperature (°C)</b>	<b>Plant species studied</b>
<b>CA</b>	Grassland	<i>Hyparrhenia hirta</i> , <i>Brachypodium retusum</i>	Canet de Mar 41° 35' N, 2° 34' E	50	590	16.1	17
<b>CO</b>	Open scrubland	<i>Hyparrhenia hirta</i> , <i>Foeniculum vulgare</i> , <i>Cistus monspeliensis</i>	Collserola 41° 24' N, 2° 6' E	280	630	14.8	46
<b>GA</b>	Scrubland	<i>Quercus coccifera</i> , <i>Rosmarinus officinalis</i> , <i>Pistacia lentiscus</i>	Garraf 41° 16' N, 1° 55' E	330	580	15.7	25
<b>PA</b>	Scrubland	<i>Quercus coccifera</i> , <i>Ulex parviflorus</i> , <i>Pistacia lentiscus</i>	El Papiol 41° 27' N, 2° 0' E	150	628	15.4	21

4

1 **Table 2.** Analyses of phylogenetic signal for colour descriptors brightness, chroma, hue  
 2 (Blomberg's K values) and colour composition ( $K_{mult}$  values) in the four study communities  
 3 separately and lumped together. Significant results ( $p < 0.05$ ) in bold.

4

<b>Community</b>	<b>n</b>	<b>Brightness</b>	<b>Chroma</b>	<b>Hue</b>	<b>Colour</b>
	<b>species</b>				<b>composition</b>
<b>CA</b>	17	0.73	<b>0.78</b>	0.85	0.48
<b>CO</b>	46	0.35	<b>0.57</b>	<b>0.71</b>	0.33
<b>GA</b>	25	0.26	0.61	0.44	0.25
<b>PA</b>	21	0.34	<b>0.52</b>	0.76	0.26
<b>CA+CO+GA+PA</b>	85	<b>0.57</b>	<b>0.60</b>	<b>0.71</b>	<b>0.38</b>

5

1 **Table 3.** R values of phylogenetically-controlled partial Mantel tests between colour  
2 descriptors and pollinator composition in the four communities and overall (data of the four  
3 communities lumped together). All results are non-significant.

<b>Community</b>	<b>Brightness</b>	<b>Chroma</b>	<b>Hue</b>	<b>Colour composition</b>
<b>CA</b>	-0.089	0.096	0.199	0.158
<b>CO</b>	-0.002	-0.031	0.015	-0.080
<b>GA</b>	-0.025	0.115	-0.082	-0.038
<b>PA</b>	-0.103	-0.063	-0.035	0.051
<b>Overall</b>	0.021	-0.023	0.019	0.006

4

5

1 **Table 4.** Relationships between pollinator groups and the different floral colours in the four  
 2 communities and overall (data of the four communities lumped together) estimated visually  
 3 from the CCA biplots of Fig. 1 for the four communities and Fig. 2 for the overall tendencies.

4

<b>Community</b>	<b>Bees</b>	<b>Ants</b>	<b>Wasps</b>	<b>Coleopterans</b>	<b>Dipterans</b>	<b>Lepidopterans</b>
<b>CA</b>	Purple UV-yellow	Purple UV-yellow	UV-yellow	White Yellow Pink	White UV-yellow	White Pink
<b>CO</b>	Purple	White Yellow	Yellow	Yellow Green White	Purple UV-yellow	Yellow UV-yellow
<b>GA</b>	Purple Pink	Yellow Green	Purple	White Pink	Yellow Green	Pink
<b>PA</b>	Purple Pink	Yellow Green White	White	Pink	White Pink	Pink
<b>Overall</b>	Purple	UV-yellow Green	UV-yellow	White Yellow	UV-yellow	Pink

5

6

1 **Supplementary Information**

2 **Figure S1.** Phylogenetic tree of the 85 species surveyed. Colour dots represent the colour  
3 category of each plant species (see colour legend in Fig. 1).

4

1 **Table S1.** Percentage of species in each of the four communities (CA, CO, GA, PA)  
2 corresponding to the flower colour categories defined by Chittka *et al.* (1994) with an  
3 example species of each category.

4

1 **Table S2.** Number (and percentage) of pollinators of each pollinator group surveyed in each  
2 of the four communities (CA, CO, GA, PA).

3

1 **Table S3.** Mean percentage of visits of each pollinator group in each colour category (data of  
2 the four communities lumped together).

3

4

5