PART OF A SPECIAL ISSUE ON POLLINATOR-DRIVEN SPECIATION

Novel adaptation to hawkmoth pollinators in *Clarkia* reduces efficiency, not attraction of diurnal visitors

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Received: 11 July 2013  Returned for revision: 6 August 2013  Accepted: 21 August 2013

**Background and Aims** Plant populations experiencing divergent pollination environments may be under selection to modify floral traits in ways that increase both attractiveness to and efficiency of novel pollinators. These changes may come at the cost of reducing overall effectiveness of other pollinators. The goal of this study was to examine differences in attractiveness and efficiency between *Clarkia concinna* and *C. breweri*, sister species of annual plants with parapatric distributions.

**Methods** An assessment was made as to whether observed differences in visitors between natural populations are driven by differences in floral traits or differences in the local pollination environment. Differences in floral attractiveness were quantified by setting out arrays of both species in the geographical range of each species and exposing both species to nocturnal hawkmoths (*Hyles lineata*) in flight cages. Differences in visitor efficiency were estimated by measuring stigma–visitor contact frequency and pollen loads for diurnal visitors, and pollen deposition on stigmas for hawkmoths.

**Key Results** The composition of visitors to arrayed plants was similar between plant species at any particular site, but highly divergent among sites, and reflected differences in visitors to natural populations. Diurnal insects visited both species, but were more common at *C. concinna* populations. Hummingbirds and hawkmoths were only observed visiting within the range of *C. breweri*. Despite attracting similar species when artificially presented together, *C. concinna* and *C. breweri* showed large differences in pollinator efficiency. All visitors except hawkmoths pollinated *C. concinna* more efficiently.

**Conclusions** Differences in the available pollinator community may play a larger role than differences in floral traits in determining visitors to natural populations of *C. concinna* and *C. breweri*. However, floral traits mediate differences in pollinator efficiency. Increased effectiveness of the novel hawkmoth pollinator on *C. breweri* comes at relatively little cost in attractiveness to other visitors, but at large cost in their efficiency as pollinators.

**Key words:** *Clarkia concinna*, *Clarkia breweri*, floral traits, *Hyles lineata*, parapatric divergence, pollinator effectiveness, pollinator efficiency, pollinator shift, speciation.

INTRODUCTION

Changes in floral phenotypes to attract novel pollinators have been common and important in the evolutionary history of the angiosperms (Grant, 1949; Baker, 1959; Stebbins, 1970; Kay et al., 2006). Divergence in pollinator use can reduce gene flow among populations, potentially facilitating speciation (Grant, 1994; Schemske and Bradshaw, 1999; Kay and Sargent, 2009). The external pollinator environment is probably paramount in driving adaptation to novel pollinators (Stebbins, 1970; Harder and Barrett, 2007; Thomson and Wilson, 2008; Kay and Sargent, 2009), although there may be an important additional role for certain types of mutations that cause large changes in colour or reward in facilitating or constraining pollinator transitions (e.g. Bradshaw et al., 1998; Bradshaw and Schemske, 2003; Rausher, 2008). Geographical variation in the attributes of pollinator assemblages or the co-occurring plant community may result in range-wide differences in visitation rates of pollinators (Grant and Grant, 1965; Eckert, 2002; Aigner, 2005; Moeller, 2005; Hersch and Roy, 2007). These differences can drive selection for ecotypes that are better matched with the local pollination environment (Johnson, 1997; Johnson and Steiner, 1997; Perez-Barrales et al., 2007; Kay and Sargent, 2009; Anderson et al., 2010; Newman et al., 2012).

Adaptation to novel pollinators may come at the cost of reduced effectiveness of existing pollinators, as a single floral phenotype is unlikely to be optimal for all pollination environments. These pollinator effectiveness trade-offs may play a large role in determining the ease of transition to a novel floral phenotype (Aigner, 2001, 2004; Sargent and Otto, 2006; Muchhala, 2007; Kay and Sargent, 2009; Dell’Olivo and Kuhlemeler, 2013). Altering the floral phenotype to increase the effectiveness of a novel pollinator may result in a large decrease in effectiveness of a former pollinator (Fig. 1). In this so-called ‘con cave’ trade-off, an intermediate phenotype may suffer reduced fitness due to poor effectiveness of both pollinators, making the evolutionary transition difficult. Alternatively, increasing the effectiveness of a novel pollinator may come with little or no cost in the effectiveness of the former pollinator (Fig. 1). If the transition involves a ‘convex’ trade-off or no trade-off at all, then a new pollinator may be more readily acquired (Aigner, 2001; Sargent and Otto, 2006). However, a convex trade-off is less likely to reduce gene flow.
Once changes in attractiveness traits occur, selection can act to increase the efficiency of novel visitors. If these visitors differ substantially in size, morphology or behaviour, a floral phenotype that is efficiently pollinated by both former and novel pollinators may not be possible. Therefore, efficiency trade-offs may often be concave (Muchhala, 2007). If novel visitors have higher visitation rates than former visitors in the local environment, selection for novel efficiency traits may occur despite a concave trade-off. While differences in visitation rates between closely related species or populations are frequently studied, differences in pollinator efficiency are less commonly quantified (reviewed by Kay and Sargent, 2009).

Although it is difficult to determine the order in which floral traits evolved by studying the current evolutionary endpoints, we can assess the differences in attraction and efficiency of potential pollinators in carefully chosen sister taxa. In this study, we quantify differences in visitor effectiveness between Clarkia concinna and C. breweri (Onagraceae), two closely related sister species with parapatric ranges that occur across a steep environmental gradient of precipitation and habitat types in the coastal ranges of California, USA. Clarkia concinna is probably more similar to the ancestral phenotype (Lewis and Lewis, 1955; Sytsma et al., 1990), and is visited by a variety of diurnal insects (MacSwain et al., 1973; Groom, 1998). Clarkia breweri is unique in the genus in having sweet-scented, night-opening flowers that are predicted to attract hawkmoths (MacSwain et al., 1973; Raguso, 1995; Raguso and Pichersky, 1995). We specifically chose to study parapatric species, as they are more likely to be representative of the early stages of geographical divergence in response to different pollination environments, and floral traits are unlikely to have been shaped by sympatric competition for pollinators (e.g. Armbruster et al., 1994), facilitation of pollination (e.g. Moeller, 2004) or selection against hybridization (i.e. reinforcement; e.g. Kay and Schemske, 2008).

Our study uses C. concinna and C. breweri to investigate the transition to a novel pollinator. We first quantify the between-species differences in floral morphology that may be involved in this transition. Next, we compare the visitor assemblages and the amount of diurnal and nocturnal pollen receipt in natural populations of both species. We expect nocturnal hawkmoths to visit and pollinate C. breweri, and diurnal insects to visit and pollinate C. concinna. We then compare visitation rates of different functional groups between plant species when placed together in each species’ geographical range. We predict that, when provided a choice, hawkmoths would prefer C. breweri, and all other visitors will not show a strong preference. Finally, we compare estimates of pollinator efficiency for each functional group of pollinators between plant species. We predict that diurnal insects will more efficiently pollinate C. concinna and hawkmoths will more efficiently pollinate C. breweri.

MATERIALS AND METHODS

Study system

Clarkia concinna (Fisch. & C.A.Mey.) Greene and C. breweri Greene are sister species of self-compatible, annual herbs
endemic to the Northern and Central California Coast Ranges, USA, with *C. breweri* replacing *C. concinna* at the south-eastern (drier) edge of the latter species’ range (Lewis and Lewis, 1955; Gottlieb and Weeden, 1979; Fig. 2). The more broadly distributed *C. concinna* occurs in chaparral, mixed evergreen forest and mesic oak/pine woodland, whereas *C. breweri* is restricted to xeric, exposed hill-sides in chaparral with a *Pinus sabiniana* overstorey. Sites with *C. breweri* populations average half the precipitation of *C. concinna* sites [42.8 cm (s.d. 9.1 cm) and 83.6 cm (s.d. 26.9 cm), respectively and populations bloom approximately 1 month earlier (April–May versus May–July; T. Miller, unpubl. res.). Both species have four showy, lobed and clawed petals, a long, narrow hypanthium containing a nectar reward, four stamens and an inferior ovary (Fig. 3). Flowers are protandrous with the four-lobed stigma opening and becoming receptive 2–7 d after anthesis. Previous observations of natural populations indicate that *C. concinna* is visited by long-tongued flies, butterflies and bees (MacSwain et al., 1973; Groom, 1998). *Clarkia breweri* is much paler pink in colour than the bright magenta of *C. concinna*. Additionally, the former is more strongly scented, due to the production of sweetly scented linalool and its pyranoid oxide (in addition to novel aromatic volatiles) at rates 250 times higher than *C. concinna* per unit floral mass (Pichersky et al., 1994; Raguso and Pichersky, 1995).

Both species provide nectar as a floral reward. The nectar differs between species in composition (mean sucrose/hexose ratios are 2 : 3 for *C. concinna* \((n = 10)\) and 3 : 2 for *C. breweri* \((n = 10)\), as determined via high-performance liquid chromatography; see Raguso et al., 2003 for methods), but total percentage sugar content is similar [brix, Bellingham & Stanley refractometer; *C. breweri* \((n = 18)\) 33.9 ± 9.9 % per flower; *C. concinna* \((n = 22)\) 35.0 ± 7.5 % per flower, \(t = 0.40, \text{ d.f.} = 38, P = 0.69\). *C. breweri* produces more nectar per flower [nectar from greenhouse-grown plants mean ± SEM of standing crop: *C. breweri* \((n = 18)\) 6.2 ± 2.8 μL per flower; *C. concinna* \((n = 22)\) 2.3 ± 0.8 μL per flower, \(t = 6.32, \text{ d.f.} = 38, P < 0.0001\). The paler colour and fragrance of *C. breweri* (MacSwain et al., 1973), more copious nectar and the tendency of *C. breweri*’s anthers to dehisce in the evening (K. Kay and D. Picklum, unpubl. res., R. Raguso, unpubl. res.) suggest a nocturnal moth pollination syndrome. The current species distributions and novelty of the floral traits of *C. breweri* support a pattern of recent evolution of the xeric-adapted *C. breweri* from a mesic-adapted *C. concinna*-like ancestor (Lewis and Lewis, 1955; Sytsma et al., 1990).

**Seed sources**

We collected seed for all experiments from two natural populations in 2008 and 2010 near centres of abundance for both species (Fig. 2). Seed of *C. breweri* originated from a site 15 km east of the summit of Mount Hamilton along San Antonio Road, Santa Clara County, California (WGS 1984 coordinate system: 37°35′ N, 121°56′ W, elevation 630 m; MH). Seed of *C. concinna* originated from a site along Knoxville Road, south of Lake Berryessa, Napa County, California (38°49′ N, 121°24′ W, 160 m; KN). Both sites have habitats typical of their respective species. The seeds were stored in dry silica at 4 °C until transport to the field sites. In the winters of 2010 and 2011, we planted seeds in 3.8-cm-diameter containers in a mixture of four parts Pro-Mix HP Mycorrhizae potting soil to one part perlite. Germination occurred in Conviron E-15 growth chambers (15 °C, 14-h light; 10-h 10 °C dark) and we moved plants to a greenhouse when secondary leaves had emerged. Plants were kept in the greenhouse at 13–21 °C until transport to the field sites.

**Floral morphology**

We characterized floral morphology by taking measurements with digital calipers from one flower with a freshly open stigma on 17 greenhouse-grown plants of each species. We measured corolla diameter from the tip of the left petal to the tip of the right petal, and petal width at the widest point of the top petal. We measured style and filament length from the hypanthium opening and hypanthium width just below the opening and just above the ovary. We used a discriminant analysis to compare floral measurements between species. All statistics were performed in JMP version 10.0 (SAS, USA).

**Floral visitors to natural populations**

To quantify differences in pollinator assemblies, we observed insect and hummingbird visitors to natural populations of both species. All observations occurred during peak flowering. Diurnal observations were conducted from mid-morning until late afternoon, corresponding with the peak in diurnal pollinator activity. Nighttime observations were conducted with the aid of a red-filtered flashlight from dusk until complete darkness, from approx. 2000 to 2200 h, with the exact interval dependent on the
time of year. This time of day corresponds with the peak in hawkmoth activity. We observed *C. breweri* in April and May 1990–1994 at the seed collection site, and at an additional site along Del Puerto Canyon Road, near Frank Raines Regional Park, Stanislaus County, California (37°42′4″N, 121°36′9″W, 310 m; FR; Fig. 2). Total observation time was 55 h (36 daytime and 19 nighttime hours over 37 d). We observed *C. concinna* in May 1990 and June 2010 at the seed collection site and five additional nearby sites (within 15 km). Visitor communities were qualitatively similar between years, so the 1990 and 2010 data were combined. Total observation time was 32 h (5 daytime hours over 2 d in 1990, 21 daytime and 6 nighttime hours over 6 d in 2010).

We classified visitors into six functional groups based on size and behaviour (Fenster *et al.*, 2004): long-tongued flies...
(Bombiliidae and Acroceridae), small bees (Andrenidae, Megachilidae, Halictidae and Apidae in the genera Anthophora and Cera- 

tina and Archilochus alexandri). Short-tongued flies (Syrphidae and Calliphoridae) and beetles also visited both the 

natural populations and the floral arrays at low rates, but were not observed contacting reproductive structures of either species and 

thus are excluded from analyses. We observed a visitor for the duration of its foraging bout and counted the total number of visits. As 

the goal was to accumulate a large number of pollinator observations, the total number of flowers was not standardized, and 

varied between several to a few hundred depending on population density and observer position. We calculated the number of visits 

per hour of observation and the relative proportion of the observations for each functional class.

Pollinator exclosures

We compared pollen deposition from nocturnal and diurnal visitors in natural populations of both species by covering 

flowers during the day or night. Clarkia breweri exclosures (n = 66 per treatment) were performed in May 1991 at FR and in 

May 1993 at the seed collection site. Clarkia concinna exclosures (n = 27 per treatment) were performed in June 2013 at the Donald 

and Sylvia McLaughlin Natural reserve (38.869°N, 122.378°W, 510 m). To measure only pollinator-mediated pollen deposition, 

we emasculated flowers with unopened stigmas and covered them with bridal veil. Day exposed flowers were uncovered from 

dawn until dusk and night exposed flowers from dusk until dawn. The treatment continued until corolla dehiscence, 36–72 h after treatment initiation. Then, stigmas were collected and 

pollen grains counted. Clarkia pollen is easily distinguished from other pollen by its large size, pink colour and triangular 

shape, and we examined pollen loads when our species of interest was the only blooming Clarkia in the vicinity. Five-day exposed 

C. concinna stigmas were eaten by beetles and were not included in the analyses. For each species separately, we compared the pro-

portion of stigmas with pollen deposition between treatments (day or night) with chi-squared tests. Stigmas with fewer than five 

grains of pollen were scored as no pollen deposition. We compared the natural logarithm of pollen deposited using an ANOVA with 
treatment (day or night), species and their interaction as factors. A significant interaction indicates a between-species difference 
in the timing of pollination.

Floral arrays

To assess whether observed differences in visitors to natural populations of the two species are driven by floral trait differences or 

by differences in local pollinator assemblages, we placed arrays of greenhouse-grown plants of the two species near two natural 

populations of each species. Plants were transported to the sites in their original containers, watered frequently in the field, and 

kept in tents, vehicles or under bridal veil when not being used in an array. Observations at experimental arrays occurred from 

April to June in 2010 and 2011, while the nearby Clarkia population was blooming. Two of the sites, FR and KN, were locations of 
natural population observations. Additionally, we conducted array observations at two sites on Mount Hamilton, Santa Clara County, California. The species ranges are adjacent at this location, with C. breweri occurring on the south-eastern slope and C. concinna on the north-western slope. One array location was near a 

C. breweri population (37-340°N, 121-633°W; elevation 1120 m; MHS), and the other was near a C. concinna population (37-343°N, 121-649°W; elevation 1090 m; MHN; Fig. 2).

Arrays consisted of 10–30 plants of each species. In 2010, we used an approximately equal number of plants of both species, placed on six trays composed entirely of one species each (three trays per species). Within a tray, plants were evenly spaced at 

approx. 15-cm intervals. Because C. concinna produces more flowers per plant than C. breweri, this arrangement led to an unbal-
nanced number of flowers. Therefore, in 2011, we used approx-

imately the same number of flowers of both species by reducing the proportion of C. concinna plants. We observed arrays for 96 
daytime hours, which were divided into 2-h observation blocks (FR 2010: 4 d, 26 h, FR 2011: 2 d, 10 h; KN 2010: 3 d, 10 h, 

KN 2011, 2 d, 6 h; MHN 2010: 3 d, 18 h, MHN 2011: 1 d, 4 h; 

MHS 2010: 3 d, 18 h, MHS 2011: 1 d, 4 h). We also conducted 40 total hours of nighttime array observations at all four sites using infrared cameras equipped with motion detectors (FR 2010: 6 nights, 9 h; FR 2011: 3 nights, 6 h; KN 2010, 3 nights, 

5 h; KN 2011: 3 nights, 6 h; MHN 2010: 2 nights, 3 h; MHN 

2011: 2 nights, 4 h; MHS 2010: 2 nights, 3 h; MHS 2011: 

2 nights, 4 h). However, we observed no hawkmoth visits and too 

few nocturnal moth visits to effectively analyse. During some ob-

servation nights at FR and KN, we also attempted to determine local hawkmoth activity using a black light to illuminate a white 

sheet. We failed to detect any hawkmoths despite previous suc-

cess using this method at FR in 1991 (Raguso, 1995). Diurnal visitors were classified into functional groups as in the 
natural population observations. We measured visitation rates for each functional group during each observation period as 

visits per flower per hour. We then compared visitation rates of functional groups using a MANOVA with plant species, site, 

year and all interactions as factors. We compared the relative pro-

portion of array visits for each site and species to diurnal natural 

population visits with a hierarchical cluster analysis using 

Ward’s minimum variance method. If visitors to each species 

cluster with the natural population observations of the same 

species regardless of site, then the differences in natural popula-

tion observations are due to divergent floral traits. Alternatively, 

if visitors to both species at a site cluster together with nearby 

natural population observations, then differences are due to differ-

ences in the local pollinator assemblages.

Diurnal floral visitor efficiency

To quantify the efficiency of each functional group of diurnal visitors, we measured stigma–visitor contact rates and visitor 
pollen loads. Although not as accurate as quantifying pollen depos-

ition on emasculated stigmas (or the resulting seed production), 

using these two methods allowed us to obtain large sample sizes 

for all functional classes. Moreover, a visitor carrying pollen and 

contacting the stigma is much more likely to effect pollination 

than a visitor doing only one or neither of these acts. We measured 

contact rates during observations of both natural populations and 

floral arrays as the proportion of visits in which the visitor contacted 

the receptive surface of the stigma. In almost all cases of stigma
contact, the anthers were also contacted, while the converse was not necessarily true. Contact was only assessed when the observer had an unobstructed view of the visitor. We quantified pollen loads on insects collected with a net as they departed from natural populations, placing them in a clean kill jar and then transferring them to individual micro-centrifuge tubes or larger plastic vials. We were not able to quantify hummingbird pollen loads. We dabbed the bodies of collected specimens with basic fuchsin glycerine jelly (Beattie, 1971; Kearns and Inouye, 1993), transferring the jelly to glass microscope slides and melting it with a butane lighter. We counted pollen grains under 80 × magnification. Additionally, we checked the bodies of specimens for any remaining pollen, which was also counted under a dissecting microscope. We compared differences in contact rates between species for each functional group with chi-squared tests. We compared between-species differences in log-transformed pollen loads for each functional group with t-tests.

Hawkmoth effectiveness

We used captive hawkmoths (Hyles lineata) to quantify floral preferences and relative efficiency of nocturnal pollinators. This species was the most common hawkmoth observed at C. breweri populations. Hawkmoth eggs and larvae were obtained opportunistically from Colorado Springs, Eldorado County, CO, during the summer of 2009 and were used to establish a laboratory colony maintained in a growth chamber (16 h light, 8 h dark, 24 ± 1 °C, 60 ± 5 % relative humidity) at Cornell University. This species is abundant across the Americas and disperses widely (Raguso et al., 1996), so there was no a priori reason to assume that a colony established from Colorado would not be representative of coastal California moths. Larvae were raised on a corn-meal-based diet as described by Goyret et al. (2009). We sorted approx. 60 pupae by sex into 0.4 × 0.4 × 1-m flight cages in separate growth chambers until they eclosed (14 h light, 10 h dark, 23 ± 1 °C, regular misting to increase relative humidity).

To test for floral preference, we allowed hawkmoths to forage on mixed arrays of C. breweri and C. concinna. Trials were performed in a 1.8 × 1.8 × 1.8-m flight cage in a windowless indoor room with approximately equal numbers of flowers of each species (between-species difference in flower number ≤2, 56–70 total flowers per trial) haphazardly arranged in the cage. Trials consisted of a feeding bout of an individual hawkmoth from the first visit of a flower until it left the array for longer than approx. 1 min. We classified an individual visit to a flower as either an approach (close investigation of flower but no insertion into the hypanthium) or a probe (tongue inserted into the hypanthium of the flower). Trials of five or fewer probes were not included, and hawkmoths were not reused. We compared the number of approaches, probes and total visits between plant species with separate paired t-tests using each trial (n = 10) as a replicate. There was no difference in preference between the sexes. Because we did not test hawkmoth behaviour on single species arrays, we were unable to measure innate differences in latency (time to first visit) between the plant species. Furthermore, because of the relatively small size of our flight cage, our work only addresses hawkmoth preference once they are in close proximity to the flowers, and have committed to foraging for nectar. There may be additional differences between C. breweri and C. concinna in the degree to which they affect feeding motivation, long-distance attraction and latency to visit by H. lineata and other hawkmoths, due to differences in floral scent, corolla shape and reflectivity (see Goyret et al., 2007; Hoballah et al., 2007; Schlumberger and Raguso, 2008).

To estimate hawkmoth efficiency, we conducted single-species pollen deposition trials in 0.4 × 0.4 × 1-m flight cages. Nine separate trials for each species were conducted with 1–4 emasculated flowers with open stigmas and 2–9 unmanipulated flowers with fresh pollen per trial. The ratio of emasculated to perfect flowers did not differ between species [mean of 0.43 (± 0.15 s.e.) for C. breweri and 0.56 (± 0.22 s.e.) for C. concinna, t = 1.42, P = 0.17]. We placed an individual hawkmoth in the cage and allowed it to forage until it had probed all emasculated flowers after first visiting at least one perfect flower. The number of probes per emasculated flower did not differ between species [mean of 2.80 (± 1.71 s.e.) for C. breweri and 2.92 (± 1.34 s.e.) for C. concinna, t = 0.16, P = 0.87]. We collected and squashed stigmas on microscope slides, and counted pollen under 80 × magnification. Efficiency for each trial was calculated as the natural logarithm of the mean number of grains of pollen deposited per probe. Efficiency was compared between plant species with a two-tailed t-test.

RESULTS

Floral morphology

Greenhouse-grown Clarkia breweri and C. concinna differ markedly in floral traits (Table 1). Discriminant analysis correctly classified the species identity of all flowers based on floral morphology (n = 34, log likelihood <0.001). Clarkia breweri flowers have smaller corolla diameters, but wider individual petals than C. concinna. Reproductive parts, especially the anthers, are exserted further and the hypanthium is wider in C. breweri.

Floral visitors to natural populations

We observed 2913 visits over 32 h to C. concinna and 1695 visits over 55 h to C. breweri (Fig. 4). The majority of visits to C. concinna at all six observed populations were from long-tongued flies and large bees, with additional visits by diurnal lepidopterans and small bees. No nocturnal visitors or hummingbirds were observed. The majority of diurnal visits to C. breweri at both populations were from hummingbirds, a group that did not visit

<table>
<thead>
<tr>
<th>Trait</th>
<th>C. breweri</th>
<th>C. concinna</th>
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</thead>
<tbody>
<tr>
<td>Corolla diameter</td>
<td>48.4 (1.2)</td>
<td>56.9 (1.3)</td>
</tr>
<tr>
<td>Petal width</td>
<td>19.4 (0.7)</td>
<td>12.9 (0.4)</td>
</tr>
<tr>
<td>Style length</td>
<td>23.1 (0.7)</td>
<td>14.6 (0.4)</td>
</tr>
<tr>
<td>Stamens length</td>
<td>13.9 (0.7)</td>
<td>11.2 (0.3)</td>
</tr>
<tr>
<td>Top hypanthium width</td>
<td>2.05 (0.07)</td>
<td>1.51 (0.07)</td>
</tr>
<tr>
<td>Bottom hypanthium width</td>
<td>1.24 (0.04)</td>
<td>0.83 (0.02)</td>
</tr>
<tr>
<td>Floral colour</td>
<td>Pale pink</td>
<td>Bright magenta</td>
</tr>
<tr>
<td>Floral scent</td>
<td>Strong, sweet</td>
<td>Weak</td>
</tr>
</tbody>
</table>
Clarkia breweri also received a small number of visits from all other diurnal functional groups, as well as hawkmoths and other nocturnal lepidopterans.

Pollinator exclosures

Clarkia breweri flowers were primarily pollinated during the day (95% of day-exposed stigmas had pollen deposition, 19% of night-exposed stigmas; $\chi^2 = 28.8$, $P < 0.001$), and Clarkia concinna flowers during the night (26% day-exposed, 59% night-exposed; $\chi^2 = 15.0$, $P < 0.001$). More pollen was deposited during the day for Clarkia concinna, and during the night for Clarkia breweri (d.f. = 3, 177; treatment $f$ = 1.85, $P = 0.07$; species $f$ = 0.98, $P = 0.32$, treatment by species $f$ = 7.09, $P < 0.001$; Fig. 5).

Floral arrays

The identity of the plant species had no effect on the composition of visitors to the array flowers, nor was there an effect of species by year or by location interactions. Location, year and the location by year interaction all significantly affected visitor composition to the arrays (Table 2, Fig. 6). Therefore, visitation rates were determined by where and when the array was observed rather than the species identity of the flower. The three-way interaction of species, location and year was also significant. The effect of year may be attributed to between-year differences in the pollination environment or the changes in our experimental design. No functional class showed any between-species difference in visitation rates (long-tongued flies: $t = 0.47$, d.f. = 92.0, $P = 0.64$; large bees: $t = 0.18$, d.f. = 73.9, $P = 0.85$; small bees: $t = 1.04$, d.f. = 77.6, $P = 0.29$; diurnal lepidopterans: $t = 1.55$, d.f. = 66.9, $P = 0.12$; hummingbirds: $t = 0.88$, d.f. = 84.5, $P = 0.37$). The composition of visitors to flowers of both species at FR clustered with Clarkia breweri natural populations, and the composition of visitors to both species at KN clustered with Clarkia concinna natural populations (Fig. 7).

Diurnal floral visitor efficiency

Diurnal visitors of all five functional groups were potentially efficient pollinators of Clarkia concinna, but not of Clarkia breweri (Fig. 8). All functional groups contacted the stigmas of Clarkia concinna flowers more frequently than those of Clarkia breweri (Fig. 8A). Contact rates ranged from 35 to 90% in Clarkia concinna and 4 to 12% in Clarkia breweri (long-tongued flies $\chi^2 = 33.26$; large bees $\chi^2 = 33.26$; small bees $\chi^2 = 33.26$; diurnal lepidopterans $\chi^2 = 33.26$; hummingbirds $\chi^2 = 33.26$).
134.61; small bees $\chi^2 = 13.12$; lepidopterans $\chi^2 = 19.98$; hummingbirds $\chi^2 = 152.17$; $P < 0.001$ for all tests). Long-tongued flies and large bees both had larger pollen loads of *C. concinna* than *C. breweri* (Fig. 8B; long-tongued flies: $t = 9.80$, d.f. = 20.82, $P < 0.001$; large bees: $t = 11.01$, d.f. = 11.60, $P < 0.001$). Small bees carried a substantial amount of pollen of both species, although amounts were marginally higher for *C. concinna* ($t = 1.77$, d.f. = 12.98, $P = 0.10$). Diurnal lepidopterans carried little pollen of either species ($t = 0.94$, d.f. = 18.80, $P = 0.35$).

**TABLE 2.** MANOVA of the response of floral visitation rates (visits per flower per hour) of five functional groups of visitors to species (*C. concinna* or *C. breweri*), location (FR, KN, MHN, MHS), year (2010 or 2011) and all interactions

<table>
<thead>
<tr>
<th>Effect</th>
<th>Wilks' $\lambda$</th>
<th>$F$-value</th>
<th>No. of variables</th>
<th>d.f.</th>
<th>$P$</th>
</tr>
</thead>
<tbody>
<tr>
<td>Species</td>
<td>0.11</td>
<td>2.10</td>
<td>2</td>
<td>477</td>
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<td>11.66</td>
<td>4</td>
<td>12, 204</td>
<td>&lt; 0.0001</td>
</tr>
<tr>
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<td>0.30</td>
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<td>16</td>
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Significant differences ($P < 0.05$) are highlighted in bold.

**Hawkmoth effectiveness**

Hawkmoths were more effective pollinators of *C. breweri* than of *C. concinna* (Fig. 9). When presented with both species, hawkmoths visited them at approximately equal rates ($t = 0.86$, d.f. = 9, $P = 0.41$). However, more visits to *C. breweri* resulted in probes ($t = 2.22$, d.f. = 9, $P = 0.052$), whereas more visits to *C. concinna* ended in approaches without a successful probe ($t = 3.14$, d.f. = 9, $P = 0.012$). Single species efficiency trials showed the deposition of ten times more pollen per visit on the...
The large differences in diurnal visitation to natural populations of *Clarkia concinna* and *C. breweri* are largely the result of differences in available pollinator community, not differences in floral attractiveness (Figs 4 and 6). All diurnal visitors are attracted to both species equally when given a choice. Nocturnal hawkmoths, however, show a preference for probing *C. breweri* (Fig. 9A). Assuming that the floral phenotype of *C. breweri* represents a derived condition, the addition of novel traits known to attract hawkmoths (strong floral scent, paler floral colour) do not deter former pollinators, indicating the absence of an attractiveness trade-off. Pollinator efficiency follows a different pattern, apparently due to the increased floral dimensions of *C. breweri*. All diurnal visitors show the potential to consistently pollinate *C. concinna*, but not *C. breweri* (Fig. 8). In contrast, hawkmoths were much more efficient at depositing pollen on *C. breweri* than *C. concinna* (Fig. 9B). The addition of hawkmoths as an effective pollinator for *C. breweri* comes at a large cost in reducing the efficiency of former pollinators.

Natural populations

Our observations of visitors to natural populations (Fig. 4) match previous work showing *C. concinna* is visited by a diverse range of insects (MacSwain et al., 1973; Groom, 1998) and predictions that *C. breweri* is visited by hawkmoths (MacSwain et al., 1973). Diurnal insects do visit *C. breweri*, although at low rates. Surprisingly, hummingbirds are the most common visitors to *C. breweri* populations, despite this flower not fitting a typical hummingbird syndrome. *C. breweri* provides a nectar reward, and local hummingbirds have learned to exploit this resource. Hummingbirds, although present throughout the range of both species, did not visit natural *C. concinna* populations in this experiment, and have not been observed visiting at other *C. concinna* populations (Groom, 1998; T. Miller, unpubl. res.). Community context is a potential explanation for the lack of hummingbird visitation at natural *C. concinna* populations. Hummingbirds may ignore *Clarkia* if more rewarding plant species are abundant.

Pollinator preference

Our lack of a significant plant species effect on functional group visitation at the arrays suggests that there is essentially no trade-off involved in attracting novel hawkmoth visitors (see Aigner, 2001; Sargent and Otto, 2006). The lack of preference by all diurnal visitors contrasts with previous work on bee to hummingbird transitions (Schemske and Bradshaw, 1999; Castellanos et al., 2004), and hummingbird to hawkmoth transitions (Streisfeld and...
were carrying marginally more C. concinna and C. breweri for each visitor functional group to C. concinna, large bees were carrying significantly more pollen, and small bees were carrying marginally more C. concinna pollen. Numbers above columns are sample sizes. All functional groups had significantly higher contact rates to C. concinna. Observations are from both natural populations and floral arrays. Numbers above columns are sample sizes. All functional groups had significantly higher contact rates to C. breweri. Hummingbird pollen load was not measured. Long-tongued flies and large bees were carrying significantly more C. breweri pollen, they may serve as co-pollinators, especially in years when hawkmoth abundance is low. Supplemental pollination by alternative visitors in supposedly specialized plants is becoming increasingly well documented (Kessler and Hodges, 1999; Hirota et al., 2011). However, hawkmoths visited (approaches plus probes) both species at similar rates. Because we presented hawkmoths with Clarkia species in a small flight cage, our study only measured short-distance attraction once hawkmoths had committed to feeding. Although there is some evidence that hawkmoths can use scent to choose between flowers at close distances (Klahre et al., 2011), the scent of C. breweri may primarily be a long-range attractant for hawkmoths, with visual cues guiding floral probing at shorter distances (Goyret et al., 2007). Scent may also induce hawkmoths to visit flowers in general (Raguso and Willis, 2002, 2005), and once latency is overcome, they may visit indiscriminately. This ‘releasing’ function of scent can explain hawkmoth approaches to C. concinna and similarly unscented flowers in other field array experiments (Fulton and Hodges, 1999; Hirota et al., 2012). If scent is important in overcoming latency, the presence of scented plants in the surrounding community may be crucial in the initial stages of acquiring hawkmoth pollinators.

**Pollinator efficiency**

In contrast to visitation rates, we found large between-species differences in pollinator efficiency for all functional groups.
(Figs 7 and 8B). The more exserted anthers and stigma, wider hypanthium, and greater herkogamy (spatial separation of anthers and stigma) of *C. breweri* probably evolved through selection for improved hawkmoth efficiency. These morphological changes came at the cost of a substantial reduction in efficiency for all other visitors, including hummingbirds. While few studies have directly quantified trade-offs in pollinator efficiency, we believe they will usually be concave between different pollinator functional groups. It is unlikely that one floral morphology will effect equal pollen transfer across pollinators with divergent sizes, shapes and behaviours. For instance, Muchhala (2007) showed that intermediate phenotypes between bat- and bird-pollinated species of *Burmeistera* have the lowest amounts of pollen transferred due to a poor fit with either type of visitor. In *Polemonium brandegei*, hummingbirds select for exserted sex organs, while hawkmoths select for recessed sex organs, due to differences in pollen removal and placement. Year-to-year variation in pollinator abundance maintains within-population variation in these traits (Kulaba and Worley, 2008).

The foraging behaviour of visitors and floral morphology are informative in explaining potential differences in efficiency. Most hawkmoth probes of *C. breweri* resulted in strong contact with the anthers, leaving pollen hanging from the legs and lower abdomen of the insect (*Clarkia* pollen grains are typically connected with viscin threads). The long style of *C. breweri* puts the receptive surface of the stigma in better alignment with the dangling pollen than the shorter *C. concinna* style (Fig. 3E, F). Additionally, the wider hypanthium tube of *C. breweri* may better accommodate the insertion of the hawkmoth proboscis. We observed hawkmoths failing to probe *C. concinna* flowers due to difficulties in aligning their proboscises with the narrow hypanthium opening. Large bees, lepidopterans and long-tongued flies all foraged for nectar either by inserting their tongue into the hypanthium or by lapping nectar exuded at the hypanthium opening. The large distance between the anthers and the nectar reward in *C. breweri*, compared with *C. concinna*, greatly reduced anther contact, and thus the pollen load, carried by these groups (Fig. 3C, D). Small bees were the only functional group that commonly foraged on both nectar and pollen of both species. When foraging for nectar, small bees rarely contacted the stigma of either species. Individual small bees that were actively collecting pollen carried large pollen loads of both species and frequently contacted *C. concinna* stigmas; however, the greater herkogamy in *C. breweri* decreased the stigma contact rate (Fig. 3A, B). Although hummingbirds are a novel visitor of *C. breweri* natural populations, they contact *C. concinna* stigmas much more frequently due to their upright foraging position (Fig. 3G, H). An additional experiment that quantified hummingbird efficiency on *C. breweri* through florescent dye transfer found similarly low efficiencies (Raguso, 1995). However, hummingbirds can provide at least some supplemental pollination to hawkmoth-specialized plants, such as *C. breweri*, when hawkmoths are rare (Aigner and Scott, 2002).

**Future directions**

Although our results reveal clear differences in between-species patterns of attraction and efficiency, we make the important caveat that we are using current end points to infer past processes. The generality of our results can only be understood through carefully controlled comparisons of both visitation rates and efficiency across multiple types of pollinator transitions. A comparison of attractiveness and efficiency trade-offs between allopatric/parapatric and sympatric sister species is especially warranted. Our prediction from such a comparison is that trade-offs in visitation rates between species that do not co-occur are more likely to be convex or absent than in sympatric species, while efficiency trade-offs are frequently concave irrespective of geographical distributions. Another important avenue of future work is to create intermediate phenotypes to directly measure visitation rates and efficiency of possible ancestral states. Testing explicit predictions about processes involved in adopting novel pollinators will allow us to better understand how pollinator transitions contribute to plant speciation and the evolution of floral phenotypes.

**ACKNOWLEDGEMENTS**

We thank B. Bowman, V. Ford, L. Gottlieb and B. LeNeve for assistance locating populations; M. Minor, J. Bahr, A. Philbin, R. Price, K. Cnop and especially D. Picklum for assistance with field and flight cage experiments; A. Shapiro and R. Thorp for help with insect identification; K. Eberwein and M. Ramos for permission to work at FR; J. Velzy and D. Polk for expert plant care; and J. Yost and two anonymous reviewers for comments on an earlier draft. This work was partially funded by a Committee on
Research Grant to K.M.K. from UCSC. R.A.R. was supported by a Sigma Xi Grant-in-Aid of Research, the California Native Plant Society, the Theodore Roosevelt Fund of the American Museum of Natural History and by NSF grant MCB-9218989 to Eran Pichtersky, University of Michigan. Data presented here include part of a dissertation submitted in partial fulfillment of the requirements for the degree of doctor of philosophy, University of Michigan, 1995. We dedicate this paper to the memory of L. D. Gottlieb for his inspiring work on the genus Clarkia and for supporting and encouraging our work on these fascinating species.

LITERATURE CITED


Miller et al. — Novel pollinator adaptation in Clarkia


