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Running Title: Sexual conflict and speciation

**Patterns of reproductive isolation within- and between- species
characterised by a sexual conflict over mating**

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18 ABSTRACT

19

20 Theory suggests that under some circumstances sexual conflict over mating can lead to
21 divergent sexually antagonistic coevolution (SAC) among populations for traits
22 associated with mating, and that this can promote reproductive isolation and hence
23 speciation. However, sexual conflict over mating may also select for traits, such as male
24 willingness to mate, that enhance gene flow between populations, limiting population
25 divergence. Here we compare pre- and post-mating isolation within and between two
26 species characterised by male-female conflict over mating rate. We quantify sexual
27 isolation among five populations of the seed bug *Lygaeus equestris* collected from Italy
28 and Sweden, and two replicates of a population of the sister-species *Lygaeus simulans*,
29 also collected from Italy. We find no evidence of reproductive isolation amongst
30 populations of *L. equestris*, suggesting that sexual conflict over mating has not led to
31 population divergence in relevant mating traits in *L. equestris*. However, there was strong
32 asymmetric pre-mating isolation between *L. equestris* and *L. simulans*: male *L. simulans*
33 were able to mate successfully with female *L. equestris*, but male *L. equestris* were
34 largely unable to mate with female *L. simulans*. We found little evidence for strong post-
35 mating isolation between the two species however, with hybrid F₂ offspring being
36 produced. Our results suggest that sexual conflict over mating has not led to population
37 divergence, and indeed perhaps supports the contrary theoretical prediction that male
38 willingness to mate may retard speciation by promoting gene flow.

39

40

41 Key words:

42 *Lygaeus* – population divergence – sexually antagonistic co-evolution – sexual conflict –
43 sexual isolation – speciation

44

45 INTRODUCTION

46

47 The evolution of reproductive isolation is the key event in speciation, and understanding
48 how and why reproductive isolation arises remains a central topic in evolutionary biology
49 (Coyne & Orr, 2004; Butlin *et al.*, 2012). Over the last decade or so, interest has turned to
50 the role that sexual conflict, and in particular sexual conflict over mating, might play in
51 reproductive isolation (Holland & Rice, 1998; Parker & Partridge, 1998; Gavrillets, 2000;
52 Arnqvist *et al.*, 2000; Martin & Hosken, 2003; Arnqvist & Rowe, 2005). Sexual conflict
53 arises when different evolutionary optima exist for males and females for a given trait
54 (Parker, 1979; Chapman *et al.*, 2003; Arnqvist & Rowe, 2005). In terms of mating, sexual
55 conflict may arise over whether mating between two individuals takes place at all (for
56 instance if male and female mate preferences do not coincide) or over the frequency of
57 mating. Generally speaking, males are typically thought to be selected to mate more often
58 than is optimal for females (Bateman, 1948; Trivers, 1972; Arnqvist & Nilsson, 2000)
59 and to be less discriminating in terms of their mating partners, leading to sexual conflict
60 between males and females over mating rate (reviewed by Arnqvist & Rowe, 2005).

61

62 Opposing selection on males and females may be an important driver of evolutionary
63 change, especially if it results in sexually antagonistic co-evolution (SAC). Within
64 populations, SAC from inter-locus sexual conflict may be rapid, taking the form of
65 irresolvable “arms-races” between the sexes, or cyclical dynamics as males and females
66 constantly co-evolve in response to adaptations and manipulation of the other sex
67 (Parker, 1979; Rice, 2000; Arnqvist & Rowe, 2005; Lessells, 2006). As such, SAC has
68 received increasing attention due to its potential to drive populations along divergent co-
69 evolutionary trajectories, facilitating population divergence, sexual isolation, and finally
70 speciation. Laboratory studies have demonstrated that sexual conflict over mating rate
71 can lead to SAC, with the evolution of increased (or decreased) female resistance to
72 mating under situations of increased (or decreased) conflict over mating (Holland & Rice,
73 1999; Wigby & Chapman, 2004; Stewart *et al.*, 2005; Rice *et al.*, 2006). Some studies
74 have also confirmed that SAC over mating rate can result in population divergence
75 (Hosken *et al.*, 2002; Martin & Hosken, 2003), although divergence has not always

76 resulted (Wigby & Chapman, 2006; Bacigalupe *et al.*, 2007; Gay *et al.*, 2009; Maklakov
77 *et al.*, 2010).

78

79 The role of sexual conflict in terms of reproductive isolation is not therefore clear-cut.
80 First, SAC may be less commonly expressed in the field than under laboratory conditions
81 (Chapman, 2006). Indeed, studies conclusively showing past operation of SAC remain
82 limited in number (Arnqvist & Rowe, 2002a,b; Koene & Schulenburg, 2005; Bergsten &
83 Miller, 2007; Anthes *et al.*, 2008). Fewer studies still demonstrate active SAC within
84 species to be driving current population divergence in the field (but see Hebets &
85 Maddison, 2005; Sugano & Akimoto, 2007; Gagnon & Turgeon, 2011). However,
86 continued cycles of antagonistic co-evolution are not the only outcome of opposing
87 selection, and may not even be the most likely (Parker, 1979; Lessells, 2006). Second,
88 SAC over mating is expected to select for generally persistent, or exploitative, males and
89 concomitant female resistance to male mating attempts (Parker & Partridge, 1998). This
90 means that among-population or even hetero-specific matings may occur *more* readily
91 than conspecific matings, if males are selected to be highly coercive and if females are
92 less able to resist males that they have not co-evolved with (Jennions & Petrie, 1997).
93 Sexual conflict, therefore, may actually retard population divergence rather than
94 promoting it, by promoting or maintaining gene-flow across populations (Holland &
95 Rice, 1998; Markow & Hocutt, 1998; Parker & Partridge, 1998; Gavrilets *et al.*, 2001).
96 Indeed more recent theoretical studies suggest that population divergence from sexual
97 conflict may be much less likely than first thought, even when SAC is apparent, with
98 only two of six possible SAC dynamics resulting in population divergence (Gavrilets &
99 Hayashi, 2005); this may account for the lack of population divergence observed amongst
100 some laboratory evolution studies (see above). Moreover, the renewed appreciation of
101 phenomena such as same-sex matings (Bailey & Zuk, 2009) and reproductive
102 interference (Burdfield-Steel & Shuker, 2011) reminds us that mate discrimination is
103 sometimes far from perfect, both in the laboratory and the field.

104

105 Measuring the degree of sexual isolation among allopatric populations that show
106 variation in sexual conflict over mating is one way to gain insight into the role of sexual

107 conflict in population divergence and speciation in the field (Gay *et al.*, 2009; Gagnon &
108 Turgeon, 2011). Population crosses are an important first step towards identifying current
109 episodes of population divergence, and in exploring the involvement of sexual conflict
110 and SAC in such diversification episodes (e.g. Hebets & Maddison, 2005; Long *et al.*,
111 2006; Panhuis *et al.*, 2006; Sugano & Akimoto, 2007), although interpreting patterns of
112 sexual conflict themselves from population crosses requires caution (Long *et al.*, 2006).

113

114 Here, we explore patterns of pre- and post-mating reproductive isolation among
115 populations of two closely related species of seed bug characterised by sexual conflict
116 over mating, *Lygaeus equestris* (LE) and *Lygaeus simulans* (LS). These species have
117 similar ecologies, including aposematic warning colouration, and both have promiscuous
118 mating systems characterised by sexual conflict such that multiple mating is extremely
119 costly to females in terms of reduced longevity and fecundity (Shuker *et al.*, 2006). Males
120 therefore benefit more from extra copulations than females do. Two LE populations were
121 derived from Sweden, three from Northern Italy, and a LS population was twice sampled
122 from Central Italy. In terms of predictions, given that the two species are presumed sister
123 species, first we expect pre-zygotic isolation to be more developed than the post-zygotic
124 isolation (Coyne & Orr, 2004). Second, if sexual conflict is associated with increased
125 population divergence for mating traits, then we should expect variation in the extent to
126 which populations of LE mate with each other, and perhaps also in the fitness of hybrid
127 offspring. If this isolation is in part associated with ecological factors then we might also
128 expect LE individuals from the Swedish or Italian populations to be more likely to mate
129 with individuals from the same region as themselves. We performed two sets of within-
130 and between-species no-choice mating experiments to test these predictions. The first
131 assayed mating over a short period for reproductively mature individuals, allowing
132 investigation of the latency of individuals to mate among four populations (three LE and
133 one LS). The second experiment expanded the number of populations studied to seven
134 populations (five LE and two LS), and assayed mating over a longer period of adult life,
135 as well as the production of an F₁ and F₂ offspring generation.

136

137 MATERIALS AND METHODS

138

139 *The study species*

140 *Lygaeus equestris* (Linnaeus; Hemiptera: Lygaeidae) is an aposematic seed feeding insect
141 with a wide geographic distribution, ranging from Spain to Russia (Deckert, 1985;
142 Solbreck *et al.*, 1989; Péricart, 1998; for a general review of lygaeid biology see
143 Burdfield-Steel & Shuker 2014a). It is a generalist seed predator of various composites
144 (Solbreck & Kugelberg, 1972; Kugelberg, 1973; Solbreck *et al.*, 1989) however its
145 preferred host plant through much of Europe is *Vincetoxicum hirundinaria* (Gentianales,
146 Asclepiadaceae: Kugelberg, 1974, 1977; Solbreck *et al.*, 1989). It is largely univoltine,
147 producing one offspring generation per year. However, *L. equestris* retains the potential
148 for multi-voltinism across its distribution (Solbreck & Sillén-Tullberg, 1981; Solbreck,
149 1991) which is more likely to occur in southern regions (Solbreck *et al.*, 1989; Shuker *et al.*,
150 2006). Relatively recently, studies concluded the presence of a sister species, *Lygaeus*
151 *simulans* (Deckert, 1985), in central and southern Europe (Deckert, 1985; Péricart, 1998;
152 Tadler *et al.*, 1999). *Lygaeus simulans* differs to *L. equestris* in the morphology of the
153 base of the antennae and male parameres (genital claspers: Deckert, 1985; Péricart,
154 1998), but clear discrimination with the naked eye is difficult (Tadler *et al.*, 1999). Much
155 of the ecology of the two species is thought to be similar, and both display polygamous
156 mating systems characterised by sexual conflict over mating (Sillén-Tullberg, 1981;
157 Deckert, 1985; Tadler, 1999; Tadler *et al.*, 1999; Micholitsch *et al.*, 2000; Shuker *et al.*,
158 2006). The two species are parapatric across parts of their range and to date no hybrids
159 have been found in the field (Maschler, 2002).

160

161 *Experiment 1: Four populations*

162 In terms of general husbandry, both species were maintained in stock cages in an
163 incubator at 29°C with a 22:2 hour light: dark cycle in order to prevent the initiation of
164 reproductive diapause. Stock cages (plastic sandwich boxes measuring 30x15x15cm with
165 fine mesh over part of the lids) were provisioned with organic sunflower seeds (Goodness
166 Direct, UK) to a depth of about 3cm (Shuker *et al.*, 2006). A piece of cotton wool was
167 added to the cages for bugs to cling on to and hide in. Oviposition was either amongst the
168 sunflower seeds or on this piece of cotton wool. Each week two universal tubes were

169 filled with distilled water and stoppered with cotton wool were provided as a water
170 source. Populations comprised two to three stock cages and were maintained in
171 continuous culture with overlapping generations, with stock cages being replaced
172 approximately every 6-8 weeks. A sample of around 60 individuals of all age classes was
173 haphazardly removed from the population cages to initiate a new one. Egg-to-adult
174 development under these conditions is broadly similar for both species, taking
175 approximately 23-28 days.

176

177 To ensure a continual supply of bugs for the experiments, we transferred late larval
178 instars periodically (every 4-5 days) from continuous culture cages into smaller nymph
179 development cages for adult eclosion. In our first experiment, mating interactions were
180 examined for sexually mature individuals from four populations, in a 4x4 reciprocal
181 design (i.e. 16 replicated combinations). The populations were two Swedish LE
182 populations (from Morga and Geta), a LE population from the Dolomites region of
183 northern Italy, and a population of LS sampled from the Tuscany region of central Italy.

184

185 From each population, we removed newly eclosed adults from nymph cages every two
186 days, and placed them in same sex pots (within their respective populations) with others
187 of the same age to develop, ensuring virginity. Densities were restricted to six bugs per
188 pot (measuring 8 x 8 x 5.5cm, transparent with perforated lids). Each pot contained a
189 30mm diameter Petri dish lid and base containing water soaked cotton wool and organic
190 sunflower seeds respectively. Seeds and water were replaced every three days ensuring
191 an *ad libitum* supply of both. All bugs were retained in these conditions for at least seven
192 days prior to experimentation ensuring sexual maturity.

193

194 Adult bugs aged between seven and 12 days post-adult eclosion were then used in ‘no
195 choice’ mating trials. We randomly assigned these sexually mature males and females to
196 partners of the opposite sex from each of the four populations (LE: Morga, Geta,
197 Dolomites; LS: Tuscany) in a fully-factorial reciprocal design. Pairs were placed in
198 transparent pots, without any seeds or water, and were randomly distributed in trays,
199 before returning to the incubator. We scored the pairs every 30 minutes, for eight hours,

200 for copulation (stable end to end position: Sillén-Tullberg, 1981; Tadler *et al.*, 1999).
201 Each hour, pots were rotated in their position within the incubator to minimise potential
202 position effects. Pairing treatments were performed in blocks, with at least two replicates
203 of each combination attained per block. Pairs where one or both individuals died during
204 the observational period were discarded, and not included in the analysis. From the eight
205 hour observation period, we recorded whether mating occurred and, for each pair that
206 mated, the time taken for mating to occur. We obtained 25 replicates for each reciprocal
207 cross (total $N = 400$ trials).

208

209 *Experiment 2: Seven populations*

210 In this experiment, we recorded the incidence of mating over a prolonged period of adult
211 development (up to approximately three weeks). In addition, we used individuals from
212 this second experiment to further explore aspects of pre- and post-mating sexual
213 isolation. We reared any F_1 generation offspring produced, and allowed them to
214 reproduce, thus enabling assessment of hybrid F_1 fertility.

215

216 We included two further LE populations from Northern Italy (Ledro and Predazzo), and a
217 replicate LS population from the same area in Tuscany, thus creating a 7×7 reciprocal
218 population design (i.e. 49 replicated combinations). All populations were maintained as
219 before. For this experiment, males and females were paired from 0-2 days post adult
220 eclosion and retained in small pots (8 x 8 x 5.5cm) containing a layer of organic, de-
221 husked, sunflower seeds, and a 7ml plastic tube containing carbon filtered water capped
222 with a cotton wool bung, for a maximum of 20 days. Water tubes were replaced every 10
223 days, or when necessary, to ensure a constant supply of water. From days 3 to 12 of being
224 paired (bugs aged ~ 5 to 14 days post eclosion, allowing for the necessary development
225 time required for sexual maturity), we scored pairs twice daily for mating. Males that
226 died during this time, without mating being observed, were replaced by another virgin
227 male of similar age from the appropriate population. Where males died and mating was
228 observed the female was isolated in a similar pot and left to oviposit. If a female died
229 without any observed matings or eggs the males were likewise re-used with new females.
230 However if a female died after producing eggs, the eggs (within the pot) were retained

231 and left to develop, and the male discarded. After day 12, adults were left to continue
232 mating (without scanning for mating) and females left to oviposit until day 20,
233 whereupon both adults were removed and discarded.

234

235 Following removal of the adults on day 20, eggs were left to develop in their respective
236 pots without their parents. After a further 7 days (when all viable eggs have hatched), we
237 scored pots for the presence and number of offspring (hatched eggs). We then transferred
238 up to 25 nymphs to a fresh pot with seeds and water (as above) and allowed them to
239 develop to adulthood. Pots were then checked for adult eclosion every 5 days and a
240 maximum density of 5 males and 5 females were retained together in fresh pots upon
241 eclosion (with a minimum of at least 3 F₁ males and 3 F₁ females per pot). When either
242 no males or no females eclosed (i.e. one sex absent), virgin bugs from other replicated
243 pots (of the appropriate cross) were used to allow for the mating and fertility of this F₁
244 generation to be assayed. Once again, we replaced the water every 10 days or earlier to
245 allow for a constant supply of water. We retained all pots for up to 20 days and scored
246 them for the presence/absence of F₂ offspring (hatched eggs).

247

248 We observed between 8 and 14 replicate pairs for each of the 49 treatment combinations
249 (median $N = 10$). Out of the overall 491 replicates, 12 replicates produced an F₁
250 generation without mating being directly observed between the two daily scans. All
251 combinations of crosses that produced F₁ offspring were observed mating for at least a
252 subset of the replicates however, indicating that twice daily observational scans were
253 adequate for assessing mating (as is likely given the prolonged copulation durations
254 exhibited within these species: Shuker *et al.*, 2006). Of the 223 replicates that
255 successfully produced F₁ offspring, 22 had insufficient numbers of adults emerging (i.e.
256 below our threshold of three males and three females) to assay for the production of F₂
257 offspring within these respective replicate families.

258

259 *Analysis*

260

261 *Experiment 1*

262

263 We analysed the incidence of mating using binary logistic regression with a logit link
264 function. First, we compared the within-population trials to examine if there were
265 population differences in their baseline mating frequencies. Second, male and female
266 identities (their population origin) were used as factors, along with the interaction term
267 between them to test for differences in likelihood of mating across all population
268 combinations. Sexual isolation is indicated by significant interaction effects, and was
269 assessed in each case using likelihood ratio (LR) tests. These analyses were performed
270 using R (R version 2.11.1).

271

272 We further analysed sexual isolation among the populations and species using an overall
273 sexual isolation index (IPSI, Rolan-Alvarez & Caballero, 2000; Perez-Figueroa *et al.*,
274 2005; see the latter paper for a detailed comparison of different indices). This was
275 performed as a global analysis ($I_{PSI_{total}}$) as well as for each pairwise comparison ($I_{PSI_{a,b}}$),
276 and estimates of asymmetry among these crosses ($I_{APSI_{a,b}}$) were also tested (Carvajal-
277 Rodriguez & Rolan-Alvarez, 2006). “Asymmetry” describes the extent to which
278 reproductive isolation in one direction of cross differs from the reciprocal cross.
279 Significance of these sexual isolation indices was tested using bootstrap resampling with
280 10,000 iterations. Where no mating was observed among pairs, zeros were replaced with
281 0.5 to allow for bootstrap resampling. Tests of sexual isolation were performed using the
282 programme JMATING (Carvajal-Rodriguez & Rolan-Alvarez, 2006).

283

284 In terms of the latency to mating, we analysed the time taken to mate using a generalised
285 linear model with a quasi-poisson error distribution and log link to account for
286 overdispersion of the data.

287

288 *Experiment 2*

289

290 Mating propensity and sexual isolation among the populations and species was assessed
291 as above. We also tested F₁ progeny production (from those pairs that were observed *in*
292 *copula*), and the number of progeny produced (of those that produced offspring) by each

293 mating pair. We analysed the incidence of F₁ progeny production among pairs with
294 logistic regression, firstly for conspecific pairs (within populations), and subsequently
295 using all crosses to examine the effects of male and female identity and their interaction
296 on mating success respectively. This was then repeated using only the subset of pairs that
297 mated to investigate potential post mating reproductive barriers, firstly for all pairs and,
298 secondly, among *L. equestris* populations only. We analysed variation in the number of
299 progeny produced (by pairs that produced offspring) again firstly for conspecific pairs
300 (within populations), and subsequently using all crosses to examine effects of male and
301 female identity and their interaction on fecundity respectively, using ANOVA. Lastly, *L.*
302 *equestris* populations were further analysed to test for differential mating success in terms
303 of the number of hatched offspring produced by reproductive pairs. Where the interaction
304 term was not significant ($p > 0.05$), it was removed and the model refitted using only the
305 main effects. Again, these analyses were performed using R (R version 2.11.1). The data
306 presented in Figures 1-3 are also given as raw proportions in the Supplementary
307 Information file.

308

309 RESULTS

310

311 *Experiment 1*

312

313 Mating frequencies for within-population mate trials were similar across all the four
314 populations (mean proportion mating = 0.89, $N = 100$; Likelihood ratio test: LR = 3.00, df
315 = 3, $p = 0.39$; Figure 1). Across all 16 combinations however there was significant
316 behavioural reproductive isolation, generally associated with the inter-specific pairings of
317 *L. equestris* and *L. simulans* (Figure 1). As such, there was a highly significant interaction
318 effect between male and female population on the incidence of mating (interaction effect;
319 LR = 231.12, $df = 9$, $p < 0.0001$) as well as differences among the populations in overall
320 male willingness to mate (LR = 40.55, $df = 3$, $p < 0.0001$) and female willingness to mate
321 (LR = 71.22, $df = 3$, $p < 0.0001$). Within just LE, Morga males mated less overall than
322 males from other populations (LR = 9.17, $df = 2$, $p = 0.01$; see Figure 1), but for LE
323 females there were no population differences in the incidence of mating (LR = 0.35, $df =$

324 2, $p = 0.84$) and no interaction between male identity and female identity on the incidence
325 of mating (LR = 4.73, $df = 4$, $p = 0.32$).

326

327 Using the isolation statistics, we confirmed strong pre-mating sexual isolation between
328 LE (Morga, Geta, and Dolomites) and LS (Total $I_{psi} = 0.573$, $p < 0.001$; Table 1), but no
329 sexual isolation effects among LE populations (see Table 1). Furthermore, the sexual
330 isolation effect between LE and LS was asymmetric for the Dolomites*Tuscany cross
331 ($I_{Apsi} = 4.065$, $p = 0.005$) with LS males able to mate with LE females whilst the opposite
332 was not found (Figure 1 and Table 1).

333

334 Overall, when mating did occur there was little difference either within or between
335 species in latency to mating. The time to mating did not differ among the populations or
336 species when paired with conspecific partners (mean = 1.43 hours, SE = 0.15, $F_{3,85} =$
337 0.16, $p = 0.92$). Additionally, including heterospecific pairings revealed no interaction
338 effect between male and female identity on the time to mating ($F_{5,219} = 1.42$, $p = 0.22$),
339 and no overall difference among any of the populations or species in the time taken for
340 males or females to mate (male population: $F_{3,224} = 1.48$, $p = 0.22$; female population:
341 $F_{3,224} = 0.25$, $p = 0.86$). Analysing only LE populations gave the same qualitative results
342 of no differences in time to mating (data not shown).

343

344 *Experiment 2*

345

346 As with the first experiment, there was no difference between the populations, or species,
347 in their mating propensities when paired with conspecific partners (mean proportion
348 mating = 0.813, $N = 64$; LR = 7.32, $df = 6$, $p = 0.29$). However, there was again
349 evidence of inter-specific mating isolation. Including hetero-specific pairings revealed
350 significant heterogeneity among the populations in the likelihood of mating associated
351 with female population and the interaction between male and female population (female
352 population: LR = 126.44, $df = 6$, $p < 0.0001$; interaction effect: LR = 225.11, $df = 36$, p
353 < 0.0001 ; Figure 2), but not associated with the main effect of male population (LR =
354 1.85, $df = 6$, $p = 0.93$). These effects are associated with the hetero-specific trials, as

355 within LE populations there was no significant interaction between male and female
356 population ($N = 255$, interaction effect: $LR = 22.94$, $d.f. = 16$, $p = 0.13$). The LE
357 populations did differ in the overall mating propensity of females ($LR = 11.175$, $d.f. = 4$,
358 $p = 0.025$), suggesting that the LE populations differ in their baseline level of female
359 receptivity to mating (Figure 2), but there were no among-population differences for the
360 overall mating propensity of males ($LR = 3.14$, $d.f. = 4$, $p = 0.54$).

361

362 Again, closer examination using *Ipsi* statistics revealed significant pre-mating sexual
363 isolation between the two species, but not within species (Total *Ipsi* = 0.255, $p = 0.006$;
364 Table 1). Moreover, as with the four population experiment, sexual isolation between the
365 species was significantly asymmetric when LE from Ledro, Morga, and Predazzo, were
366 paired with LS (Table 1 and Figure 2).

367

368 *F₁ offspring production*

369 There was a significant interaction effect between male and female identity on the
370 incidence of hatched offspring ($LR = 167.74$, $d.f. = 36$, $p \ll 0.0001$, male population; LR
371 $= 16.32$, $d.f. = 6$, $p = 0.012$, female population; $LR = 94.30$, $d.f. = 6$, $p \ll 0.001$),
372 indicating strong heterogeneity amongst the populations in the likelihood of successfully
373 interbreeding (Figure 3). As expected, the incidence of hatched offspring production did
374 not differ between populations when individuals were paired with conspecifics (mean =
375 0.64, $N = 64$; $LR = 5.82$, $d.f. = 6$, $p = 0.44$; it is worth highlighting that this means that
376 there was an overall “mating failure” rate of 36%). However, reducing the data set to
377 contain only those pairs that did mate, the likelihood of a male from a particular
378 population producing hatched offspring with a female did not depend on the identity of
379 the female partner and vice versa ($N = 305$, interaction effect; $LR = 36.46$, $d.f. = 27$, $p =$
380 0.11) suggesting limited post-mating reproductive isolation between the species. Refitting
381 the model (using mated pairs) without this interaction effect still revealed population
382 differences for both males and females in the likelihood of producing hatched offspring
383 following successful mating (male population; $LR = 34.35$, $d.f. = 6$, $p < 0.001$, female
384 population; $LR = 25.99$, $d.f. = 6$, $p < 0.001$). Contrary to the result of no population
385 differences in the likelihood of producing hatched offspring in conspecific pairings, this

386 suggests that there is some variability among the populations in the likelihood of
387 producing hatched offspring (Figure 3). Indeed analysis of LE populations only showed a
388 similar story to mating propensity, with a significant effect of female population identity
389 on the incidence of hatched offspring among mated individuals, but no effect of male
390 identity and no interaction between the two ($N = 218$: female identity LR = 17.13, *d.f.* =
391 4, $p = 0.002$; male identity LR = 6.02, *d.f.* = 4, $p = 0.20$; interaction term LR = 17.97, *d.f.*
392 = 16, $p = 0.37$).

393

394 There was no difference among the populations in the number of hatched offspring
395 produced (by reproductive pairs) when paired with conspecific partners (mean = 59.88, se
396 = 5.67, $F_{6,35} = 1.067$, $p = 0.40$). Analysing all population crosses, there were highly
397 significant population differences for males and females in the number of hatched
398 offspring produced overall, (male population: $F_{6,183} = 9.29$, $p < 0.0001$; female
399 population: $F_{6,183} = 7.49$, $p < 0.0001$), which is driven largely by the low numbers of
400 offspring produced among LE by LS population crosses (Figure 3). However, the
401 interaction was found to be non-significant ($F_{25,183} = 0.85$, $p = 0.67$), suggesting that there
402 was little difference in the numbers of hybrid versus pure-bred offspring, although this
403 might be in part due to low statistical power from low sample sizes.

404

405 Analysing LE populations in isolation, again there was no significant interaction effect
406 between male and female identity on the number of offspring produced ($F_{16,151} = 1.167$, p
407 = 0.30), however, male and female identity were significant as main effects (male: $F_{4,167}$
408 = 3.12, $p = 0.017$; female: $F_{4,167} = 2.73$, $p = 0.031$), suggesting there is variance in
409 fertility across populations.

410

411 *F₂ offspring production*

412 Intra-specific crosses showed no restrictions in the ability of populations to interbreed
413 and produce fertile hybrids as F₂ generations were produced in each case (Figure 4).
414 However, there was some evidence for partial post-mating reproductive isolation between
415 LE and LS as crosses between LE males and LS females did not produce any F₁ offspring
416 (perhaps not surprising as only one of these pairings mated in the first instance: Figure 2).

417 However, there were more matings in the reciprocal direction (i.e. between male LS and
418 female LE; Figures 1 and 3) and they did produce both an F₁ generation (Figure 3) and a
419 subsequent F₂ generation in some cases (Figure 4). Although low numbers of F₁ offspring
420 were produced from these crosses, sexual isolation between LE and LS would appear to
421 be largely in terms of pre-mating isolation rather than post-mating isolation therefore, and
422 largely in one direction.

423

424 DISCUSSION

425

426 The importance of sexual conflict in facilitating reproductive divergence among
427 populations remains unclear. Here we have investigated the extent of reproductive
428 isolation within and between two sister species of seed bug (*Lygaeus equestris* and *L.*
429 *simulans*), species that are characterised by sexual conflict (Shuker *et al.*, 2006). No
430 sexual isolation was apparent among different LE populations, as might be expected if
431 the sexual conflict over mating had selected for diverging reproductive traits. Instead, our
432 two experiments revealed pre-mating isolation between LE and LS, although this
433 isolation was asymmetric: male LS were able to mate with female LE, but male LE were
434 largely unable to mate with female LS. The results were similar whether we allowed pairs
435 to interact for a few hours or for a few days, and were largely consistent across multiple
436 LE populations and two replicate LS populations. In our first experiment, we found no
437 differences in time to mating for those crosses where mating occurred, again suggesting
438 little divergence in behaviour or key mating signals. We found little evidence for strong
439 post-mating isolation between the two species either however, with hybrid F₁ and F₂
440 offspring sometimes being produced if matings occurred.

441

442 Pre-zygotic sexual isolation may act over many systems involving behavioural,
443 physiological and/or morphological characters (Coyne & Orr 1989, 1997). Although no
444 overt courtship occurs in the two species studied here (Solbreck, 1972; Sillén-Tullberg,
445 1981; Tadler *et al.*, 1999; Dougherty & Shuker, 2014), morphological differences
446 between the species could explain the pre-mating isolation observed to some extent.
447 Indeed, one of the few characters that distinguish the species is the morphology of the

448 male parameres (genital claspers; Deckert, 1985; Péricart, 1998), which are used by
449 males to secure position during mating, facilitating successful copulation (Tadler, 1999;
450 Tadler *et al.*, 1999). We are currently exploring the role that claspers, and also the
451 internal male genitalia, may play in reproductive isolation.

452

453 In addition to morphological differences, we have recently identified differences in
454 cuticular hydrocarbon (CHC) profiles between five species of Lygaeidae, including *L.*
455 *equestris* and *L. simulans* (Burdfield-Steel *et al.*, unpublished data). Our data suggest
456 differences in CHC profiles between these latter two species, including sex-specific and
457 ontogenetic differences. These differences will provide a starting point for experimental
458 tests of species-discrimination mechanisms. Hetero-specific mating attempts are not
459 infrequent across four species of lygaeid more generally however (*L. equestris*, *L.*
460 *creticus*, *Spilostethus pandurus*, and *Oncopeltus fasciatus*: Shuker *et al.*, 2015; see also
461 McLain & Shure, 1987; McLain & Pratt, 1999 for another example from the Lygaeidae),
462 and such hetero-specific mating interactions can be as costly as intra-specific mating
463 harassment for female LE (i.e. there is reproductive interference: Shuker *et al.*, 2015).
464 Given the highly polygamous mating systems of these species, and the occurrence of
465 reproductive interference, our current inference is that despite the availability of cues,
466 such as CHCs, to facilitate species discrimination, males have been selected to be highly
467 opportunistic when it comes to possible mating partners, and this selection has weakened
468 selection for strong species discrimination (Parker & Partridge, 1998; Burdfield-Steel &
469 Shuker 2014b; Shuker *et al.* 2015).

470

471 This means that sexual conflict over mating, although possible in the field, may not be
472 driving divergence and speciation among LE populations. Indeed, our interpretation
473 supports the contrary theoretical proposition, that male willingness to mate may impede
474 speciation through maintaining gene flow across populations (Parker & Partridge, 1998;
475 Gavrillets & Hayashi, 2005), and would concur with other studies. For example, Gagnon
476 & Turgeon (2011) found that despite significant correlations between morphological
477 traits of males and females associated with sexual conflict over mating in populations of
478 *Gerris gillettei*, there were no mating asymmetries among allopatric populations,

479 suggesting that sexual conflict was not driving population divergence. And as highlighted
480 above, even in laboratory evolution studies, support for SAC over mating promoting
481 allopatric population divergence is mixed.

482

483 In terms of the reproductive isolation between LE and LS, there was a clear pattern of
484 asymmetric pre-mating isolation. Asymmetric sexual isolation is very common, both in
485 terms of pre-mating, behavioural isolation (e.g. Kaneshiro, 1980; Arnold *et al.*, 1996) and
486 also in terms of post-mating isolation (e.g. Turelli & Moyle, 2007). Two such examples
487 of the former include isolation between populations of the grasshopper, *Podisma*
488 *sapporensis* (Sugano & Akimoto, 2007), and between species of Sonoran desert
489 *Drosophila* (Markow & Hocutt, 1998). In both of these cases, divergent sexual
490 antagonistic co-evolution among the populations or species (with differential levels of
491 male vigour and female resistance) was attributed as a likely explanation for the patterns
492 observed, but our failure to link between-species isolation with any among-population,
493 within-species isolation suggests that SAC has not generated important divergence in
494 behaviour between LE and LS. We note here that our analysis of pre-mating isolation
495 was based on a no-choice experimental paradigm. A recent meta-analysis has confirmed
496 that the strength of mate preferences can vary with experimental paradigm, i.e. between
497 choice and no-choice paradigms (Dougherty & Shuker 2015; but see Dougherty &
498 Shuker 2014 for no effect of experimental design on intra-specific mate preferences in *L.*
499 *equestris*). Mate preferences are generally stronger when measured via choice
500 experiments (typically the presentation of two options), although the meta-analysis
501 suggested that experimental paradigm had no effect on the strength of preference for con-
502 over hetero-specifics, as studied here. However, the number of studies available for that
503 comparison was small (Dougherty & Shuker 2015).

504

505 In terms of post-mating isolation, no LE populations (from Sweden or Italy) showed any
506 restriction in their ability to interbreed, producing F₂ offspring and thus demonstrating
507 fertile F₁ hybrids. No breeding restrictions were found between the two *L. simulans*
508 replicate populations either. That is not to say that all matings involve successful sperm
509 transfer, as “mating failures” appear to be rather common, even among con-specifics.

510 Mating failure is probably more common than often realised (Eberhard, 1996; García-
511 González, 2004) and interest in this phenomenon is growing (Rhainds, 2010; Greenway
512 *et al.*, 2015).

513

514 In terms of the inter-specific crosses, only a subset of LE female and LS male replicate
515 crosses produced sufficient numbers of F₁ generation individuals that developed to adults
516 (and could reliably test for an F₂ generation), thus we could not determine with certainty
517 whether all types of crosses are equally (in)compatible, but there appears to be variation
518 in the successful production of F₁ and F₂ offspring. As such, the data we have suggest
519 there might be segregating variation in the genetic basis of any post-zygotic
520 incompatibilities, an increasingly common finding (e.g. Shuker *et al.*, 2005; Cutter,
521 2012). Nonetheless, the production of some hybrid F₂ offspring clearly demonstrates that,
522 given the no-choice mating experiments performed here, the two species can interbreed
523 and produce viable offspring under laboratory conditions.

524

525 There remain a number of outstanding questions. For instance, more needs to be done
526 unravelling how and under what circumstances males and females discriminate between
527 species (e.g. Burdfield-Steel & Shuker, 2014b; Shuker *et al.*, 2015), including further
528 exploration of the role of CHCs (and variation in CHC profiles within and among
529 populations). Second, the work presented here only begins to touch upon the nature and
530 extent of post-mating isolation. Our data show that a proportion of F₁ hybrids are viable
531 and also fertile. However, the extent of variation in hybrid viability or fertility remains to
532 be fully explored, including the extent of any reproductive incompatibilities that fit the
533 pattern of Haldane's rule (Coyne & Orr, 2004). *Lygaeus* have an XO chromosomal sex
534 determination system, therefore we would predict that if failure to produce offspring (or
535 variation in the failure to produce offspring) is associated with only one sex, it should be
536 associated with the heterogametic males. Such experiments are currently underway.

537

538 Finally, our results suggest that this species-pair may be a useful system for exploring the
539 genetic basis of reproductive isolation, as we have behavioural variation across the two
540 species, an asymmetry in the direction of reproductive isolation, and the ability to form

541 viable F2 hybrids, which would facilitate genetic mapping. As the study of reproductive
542 isolation in insects is still somewhat dominated by species of *Drosophila* and a number of
543 Orthoptera (e.g. Coyne & Orr, 2004), this system may provide a welcome new study
544 organism for exploring the evolutionary genetics of reproductive isolation in the wild.

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549

550 ACKNOWLEDGMENTS

551 We are very grateful to NERC for funding (a Studentship to GMVE and an Advanced
552 Fellowship to DMS). Liam Dougherty and Ginny Greenway provided valuable comments
553 on an earlier version of the manuscript, and we are also extremely grateful for the
554 thoughtful and constructive comments of three anonymous referees, whose comments
555 helped us greatly improve the manuscript..

556

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Figure legends

Figure 1. Proportion of pairs mating within and between populations and species ($N = 25$ each, total $N = 400$) in the first, four population experiment. Grey Bars denote pairings within and between populations of *Lygaeus equestris*: Morga, Geta, and Dolomites. White Bars denote pairings involving the sister species *L. simulans*: Tuscany. The data are also presented in tabular form in Supplementary Information.

Figure 2. Proportion of pairs mating within and between populations and species (total $N = 491$) in the second, seven population experiment. Grey Bars denote pairings within and between populations of *Lygaeus equestris*: Morga, Geta, Dolomites, Ledro and Predazzo. White Bars denote pairings involving the sister species *L. simulans*: Tuscany 1 and Tuscany 2. The data are also presented in tabular form in Supplementary Information.

Figure 3. Proportion of pairs producing offspring within and between populations and species (total $N = 491$) in the second, seven population experiment. Grey Bars again denote pairings within and between populations of *Lygaeus equestris*: Morga, Geta, Dolomites, Ledro and Predazzo. White Bars again denote pairings involving the sister species *L. simulans*: Tuscany 1 and Tuscany 2. The data are also presented in tabular form in Supplementary Information.

Figure 4. F2 progeny production for each reciprocal combination. Bugs denote successful production of F2 progeny. Dark coloured cells represent those combinations where no F1 generation was produced. Light shaded cells represent those combinations where F1 progeny were produced but insufficient numbers survived to test reliably for F2 offspring production. Only one combination, denoted as “*”, produced no F2 generation from sufficient numbers of F1 offspring to test this (3 males and 3 females).

Table 1. Global analysis of sexual isolation (*Ipsi*), and estimates of asymmetry (*IApsi*) in mating among populations using F_0 crosses for the four and seven population experiments respectively. SD is the standard deviation and p is the two tail probability of rejecting the null hypothesis being true (isolation/asymmetry = 0) in the bootstrap resampling distribution (derived from 10,000 iterations). Crosses displaying significant pre-mating isolation are shown in bold. Morga, Geta, Dolomites, Ledro and Predazzo are *Lygaeus equestris*, and the Tuscany populations are *L. simulans* (see text for details)..

| Population Cross | <i>Ipsi</i> | SD | p | <i>IApsi</i> | SD | p |
|-------------------------|-------------|-------|-----------------|--------------|-------|--------------|
| <i>Experiment 1</i> | | | | | | |
| Morga * Geta | 0.001 | 0.110 | 0.9922 | 1.003 | 0.044 | 0.935 |
| Morga * Dolomites | 0.024 | 0.109 | 0.8368 | 1.009 | 0.049 | 0.972 |
| Morga * Tuscany | 0.902 | 0.060 | < 0.0001 | 1.505 | 1.006 | 0.535 |
| Geta * Dolomites | -0.044 | 0.107 | 0.6642 | 0.998 | 0.034 | 0.869 |
| Geta * Tuscany | 0.958 | 0.042 | < 0.0001 | 0.438 | 0.620 | 0.543 |
| Dolomites * Tuscany | 0.753 | 0.070 | < 0.0001 | 4.065 | 1.235 | 0.005 |
| Total | 0.573 | 0.049 | < 0.0001 | | | |
| <i>Experiment 2</i> | | | | | | |
| Dolomites * Geta | -0.137 | 0.197 | 0.456 | 1.019 | 0.121 | 0.977 |
| Dolomites * Ledro | -0.030 | 0.195 | 0.857 | 1.013 | 0.164 | 0.834 |
| Dolomites * Morga | -0.009 | 0.185 | 0.948 | 1.011 | 0.131 | 0.888 |
| Dolomites * Predazzo | -0.120 | 0.183 | 0.492 | 0.996 | 0.116 | 0.795 |
| Dolomites * Tuscany(1) | 0.440 | 0.217 | 0.079 | 1.621 | 0.525 | 0.146 |
| Dolomites * Tuscany(2) | 0.844 | 0.131 | 0.001 | 0.739 | 0.658 | 0.947 |
| Geta * Ledro | -0.208 | 0.177 | 0.235 | 0.990 | 0.120 | 0.775 |
| Geta * Morga | 0.140 | 0.209 | 0.513 | 1.091 | 0.293 | 0.892 |
| Geta * Predazzo | -0.069 | 0.199 | 0.702 | 1.014 | 0.132 | 0.993 |
| Geta * Tuscany(1) | 0.640 | 0.192 | 0.021 | 1.423 | 0.533 | 0.268 |
| Geta * Tuscany(2) | 0.835 | 0.142 | 0.005 | 0.725 | 0.642 | 0.941 |
| Ledro * Morga | 0.011 | 0.178 | 0.964 | 0.988 | 0.138 | 0.895 |
| Ledro * Predazzo | -0.086 | 0.180 | 0.600 | 1.001 | 0.107 | 0.840 |
| Ledro * Tuscany(1) | 0.539 | 0.176 | 0.022 | 1.851 | 0.523 | 0.045 |
| Ledro * Tuscany(2) | 0.541 | 0.144 | 0.008 | 2.202 | 0.619 | 0.016 |
| Morga * Predazzo | -0.002 | 0.179 | 0.974 | 1.016 | 0.142 | 0.906 |
| Morga * Tuscany(1) | 0.647 | 0.162 | 0.010 | 1.763 | 0.584 | 0.123 |
| Morga * Tuscany(2) | 0.589 | 0.134 | 0.001 | 2.354 | 0.642 | 0.013 |
| Predazzo * Tuscany(1) | 0.539 | 0.176 | 0.027 | 1.853 | 0.533 | 0.047 |
| Predazzo * Tuscany(2) | 0.627 | 0.145 | 0.004 | 2.016 | 0.611 | 0.056 |
| Tuscany(1) * Tuscany(2) | -0.101 | 0.187 | 0.560 | 0.994 | 0.134 | 0.756 |
| Total | 0.255 | 0.081 | 0.005 | | | |