1	Divergence in female damselfly sensory structures is consistent with a species recognition
2	function but shows no evidence of reproductive character displacement
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11	isolation, damselfly
12	
13	Abstract
14	Males and females exchange signals prior to mating that convey information such as sex,
15	species identity, or individual condition. In some animals, tactile signals relayed during physical
16	contact between males and females before and during mating appear to be important for mate
17	choice and reproductive isolation. This is common among odonates, when a male grasps a
18	female's thorax with his terminal appendages prior to copulation, and the female subsequently
19	controls whether copulation occurs by bending her abdomen to complete intromission. It has
20	been hypothesized that mechanosensory sensilla on the female thoracic plates mediate mating
21	decisions, but is has been difficult to test this idea. Here, we use North American damselflies in
22	the genus Enallagma (Odonata: Coenagrionidae) to test the hypothesis that variation in female

23	sensilla traits is important for species recognition. Enallagma anna and E. carunculatum
24	hybridize in nature, but experience strong reproductive isolation as a consequence of
25	divergence in male terminal appendage morphology. We quantified several mechanosensory
26	sensilla phenotypes on the female thorax among multiple populations of both species and
27	compared divergence in these traits in sympatry versus allopatry. Although these species
28	differed in features of sensilla distribution within the thoracic plates, we found no strong
29	evidence of reproductive character displacement among the sensilla traits we measured in
30	regions of sympatry. Our results suggest that species-specific placement of female
31	mechanoreceptors may be sufficient for species recognition, although other female sensory
32	phenotypes might have diverged in sympatry to reduce interspecific hybridization.

34 Introduction

35 For sexual species, maintenance of species boundaries relies on reproductive isolation 36 (RI) between recently diverged species (Mayr, 1942). Premating reproductive isolating barriers, 37 including behavioral or sexual isolation, often evolve earlier in the speciation process than 38 postmating barriers in a variety of animal taxa (e.g., McMillan et al., 1997; Price and Bouvier, 39 2002; Mendelson and Wallis, 2003; Dopman et al., 2010; Sánchez-Guillén et al., 2012; Williams 40 and Mendelson, 2014; Castillo et al., 2015; Barnard et al., 2017). Behavioral isolation requires 41 that mate recognition signals and/or preferences diverge between populations, which 42 ultimately results in the ability of individuals to discriminate conspecifics from heterospecifics. 43 Species recognition signals may rely on a variety of sensory modalities such as color (Wiernasz and Kingsolver, 1992; Sætre et al., 1997; Jiggins et al., 2001; Boughman et al., 2005; Kronforst 44 45 et al., 2006; Williams and Mendelson, 2014), courtship behavior (Stratton and Uetz, 1986), 46 sound/vibration (Ewing and Bennet-Clark, 1968; Wells and Henry, 1998; Shaw, 2000; Gerhardt 47 and Huber, 2002; Arthur et al., 2013), and volatile chemicals (Coyne et al., 1994; Noor and Coyne, 1996; Trabalon et al., 1997; Rafferty and Boughman, 2006). Often, multiple signals act 48 49 in concert to affect species recognition (e.g., Costanzo and Monteiro, 2007; Girard et al., 2015). 50 Although much is known about the importance of visual, auditory, and chemical signals 51 and responses in sexual communication and species recognition, we know relatively little about 52 other sensory modalities that may have strong effects on individual mating decisions. Tactile 53 signals have been hypothesized to contribute to mating decisions (Mendelson and Shaw, 2012), 54 but it is unclear whether tactile cues could represent a primary species recognition signal, given 55 that visual, auditory, and chemical cues usually act earlier during the mating sequence.

Research on the prevalence of tactile signals in mating decisions is limited (Coleman, 2008)
partly because of the experimental challenge it poses: whereas other sensory modalities
present male signals to a focal female from a distance, studying female preference for tactile
cues requires contact between males and females, which is not always easily achieved or
quantified under controlled conditions.

Despite this challenge, understanding the role of tactile signals along the continuum 61 62 between intraspecific mate choice and interspecific RI is important because it broadens our 63 understanding of the causes and consequences of a common pattern in nature— the rapid 64 divergence of male genital morphology between species. It has been suggested that rapid 65 genital differentiation can cause RI (Dufour 1844), although mechanical incompatibilities 66 between heterospecific male and female genitalia do not appear to be a common cause of RI 67 (Shapiro and Porter, 1989; Masly, 2012; Simmons, 2014). However, observations both within 68 (Eberhard, 1994; Edvardsson and Göran, 2000; Briceño and Eberhard, 2009a; Briceño and 69 Eberhard, 2009b; Frazee and Masly, 2015) and between species (Patterson and Thaeler Jr, 70 1982; Robertson and Paterson, 1982; Eberhard, 1992; Coyne, 1993; Barnard et al., 2017) 71 suggest that male reproductive structures may convey tactile information to females that 72 affects their subsequent behavior and/or physiology. Although female genital structures often 73 appear invariant among closely related species (Shapiro and Porter, 1989), subtle 74 morphological differences (e.g., Kamimura and Mitsumoto, 2011; Yassin and Orgogozo, 2013) 75 could enable females to detect variation among males' morphology. Female variation in 76 detection ability could also occur in signal processing at the level of neurons, neural networks, 77 and/or in the distribution and morphology of sensory structures that receive male tactile

78	signals. These sensory structures may exist not just in the female genitalia or reproductive
79	tract, but in any region of the female that receives contact from male structures.
80	Female sensory structures that reside in body regions that contact species-specific male
81	structures during mating have been documented in several arthropods, including flies
82	(Eberhard, 2001; Ingram et al., 2008) and damselflies (Cordoba-Aguilar, 1999, 2002, 2005;
83	Robertson and Paterson, 1982). Other studies have demonstrated that tactile cues from male
84	grasping organs influence female mating responses, either via experimental manipulation of
85	male structures and desensitization of females (Eberhard, 2002; Briceño et al., 2007; Briceño
86	and Eberhard, 2009a; Eberhard, 2010; Myers et al., 2016), or via comparison of female behavior
87	when females are grasped by males with varying terminal appendage morphologies (Sánchez-
88	Guillén et al., 2012; Sánchez-Guillén et al., 2014; Barnard et al., 2017). Premating tactile
89	isolation may also be important in vision-limited vertebrates. For example, contact cues via the
90	lateral line system may influence female mate choice in a cavefish (Plath et al., 2004; but see
91	Rüschenbaum and Schlupp, 2013).
92	Tactile signals appear to be a significant cause of RI in Zygoptera, the damselfly suborder
93	of Odonata (Krieger and Krieger-Loibl, 1958; Loibl, 1958; Robertson and Paterson, 1982; Corbet,
94	1999). Concentrations of cuticular mechanoreceptors (sensilla) on the female thorax have been
95	described in several coenagrionid damselfly genera. The morphology of these sensilla is
96	consistent with a mechanosensory function and does not indicate that they conduct signals
97	related to olfaction, hygroreception, or temperature reception (McIver, 1975; Robertson and
98	Patterson, 1982). These sensilla reside in areas where males' grasping appendages contact the
99	female thorax before and during mating, which has led to speculation that they allow females

100 to evaluate male morphologies and discriminate conspecific from heterospecific males (Jurzitza, 101 1974, 1975; Tennessen, 1975; Robertson and Paterson, 1982; Battin, 1993a, 1993b). Each 102 mechanoreceptor is associated with a single sensory neuron (McIver, 1975; Kiel, 1997). The 103 thoracic sensilla thus represent a spatial matrix that can transmit signals to the female central 104 nervous system based on the pattern in which the sensilla are stimulated. Greater numbers of 105 these receptors are expected to enhance a female's sensory resolution by increasing the 106 combinatorial complexity of tactile signals that she can perceive. For example, if a female 107 possesses 25 sensilla, and each sensillum has two response states ("on" if contacted and "off" if 108 not contacted), then the number of unique tactile patterns that the female could distinguish is $2^{25} = 3.4 \times 10^7$. A female that possesses just one additional sensillum would be able to 109 distinguish among roughly twice as many tactile patterns $(2^{26} = 6.7 \times 10^7)$. Should individual 110 111 sensilla respond to quantitative variation in touch (rather than a binary response), this would 112 dramatically increase the number of response states and therefore further enhance tactile 113 acuity (e.g., Gaffin and Brayfield, 2017). Female damselfly thoracic sensilla thus present an 114 external, quantifiable phenotype in which to investigate the mechanistic basis of tactile stimuli 115 and female mating decisions.

The North American damselfly genus *Enallagma* includes several recently diverged species that often co-occur in the same habitats (Johnson and Crowley, 1980; McPeek, 1998), and do not engage in premating courtship (Fincke et al., 2007; Barnard et al., 2017) or use chemical cues for mate selection (Rebora et al., 2018). A female's first opportunity to assess a potential mate occurs when the male uses his terminal appendages to grasp the mesostigmal plates on the female thorax to form "tandem", the premating position. The male superior

122 grasping appendages (cerci) have species-specific morphologies, and differences in the 123 morphology of these structures are the primary cause of RI in this genus (Paulson, 1974; 124 Barnard et al., 2017). Two species, *E. anna* and *E. carunculatum*, occasionally hybridize in 125 nature to produce males and females with reproductive structure morphologies that are 126 intermediate to each of the pure species (Miller and Ivie, 1993; Donnelly, 2008; Johnson, 2009; 127 Barnard et al., 2017). Females of both pure species discriminate strongly against both 128 heterospecific and interspecific hybrid males that take them in tandem, which shows that 129 female E. anna and E. carunculatum can detect not only large differences in male cercus 130 morphologies, but also more subtle differences such as those that distinguish conspecific and 131 hybrid males (Barnard et al., 2017).

132 Because it appears that mesostigmal sensilla mediate species recognition, they might be 133 expected to show signs of reproductive character displacement (RCD): increased divergence of 134 traits involved in RI in regions of sympatry between E. anna and E. carunculatum relative to 135 regions of allopatry (Brown and Wilson, 1956; Howard, 1993; Pfennig and Pfennig, 2009). RCD 136 can manifest phenotypically as divergence in either signaling traits or mate preferences, in 137 which sympatric females display stronger discrimination against heterospecific males than do 138 allopatric females of the same species (e.g., Gerhardt, 1994; Gabor and Ryam, 2001; Albert and 139 Schluter, 2004; Wheatcroft and Qvarnstrom, 2017). This strengthening of preference in 140 sympatry may evolve via direct selection on adult prezygotic phenotypes, or via reinforcement, 141 where selection against interspecific hybrids gives rise to selection for enhanced premating 142 isolation between species (Dobzhansky, 1937). Enallagma anna and E. carunculatum can 143 interbreed, but their hybrids experience significantly reduced fitness (Barnard et al., 2017).

144	Female Enallagma experience frequent mating attempts from heterospecific males where both
145	species co-occur (Paulson, 1974; Fincke et al., 2007; Barnard et al., 2017). These findings
146	suggest that in sympatry, females may experience selection for stronger species discrimination
147	ability. Studies of several Enallagma species (not including E. anna or E. carunculatum) have
148	revealed that male cercus shape varies little among populations, even across large geographical
149	regions (McPeek et al., 2011; Siepielski et al., 2018). Enallagma anna and E. carunculatum
150	appear to show similar patterns, at least in the western part of their distributions (A. Barnard,
151	unpublished data). It is possible, however, that females in sympatry with other species are
152	more sensitive to variation among males than are females of the same species in regions of
153	allopatry, and this variation in sensitivity may be reflected in female sensilla traits.
154	Here, we use sensilla number, density, and location as proxies for female preference, to
155	test the hypothesis that variation in female sensilla phenotypes supports a function in species
156	recognition. We tested this hypothesis by quantifying sensilla on the mesostigmal plates of a
157	large set of <i>E. anna</i> and <i>E. carunculatum</i> females from multiple populations across the western
158	United States and comparing phenotypes of each pure species from sympatric and allopatric
159	populations to identify patterns consistent with RCD. We predicted that in sympatric
160	populations, females would possess higher sensilla numbers, higher sensilla density, and/or
161	different spatial distributions of sensilla within their mesostigmal plates when compared to
162	females from allopatric populations.
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163

164 Materials and methods

165 **Population sampling**

166	We measured the sensilla traits of 29 <i>E. anna</i> females across 13 populations, and 74 <i>E.</i>
167	carunculatum females across 19 populations (Fig. 1, Table 1). We classified each population as
168	allopatric, locally allopatric, or sympatric. Sympatric populations are those where E. anna and
169	E. carunculatum co-occur temporally as well as spatially. Because E. anna's geographic range
170	falls completely within E. carunculatum's range, only E. carunculatum has completely allopatric
171	populations. We designated populations as locally allopatric at sites within the area of range
172	overlap, but where only one species is known to occur based on occurrence data from
173	OdonataCentral (Abbott, 2016). Although some specimens were collected as early as 1945, the
174	majority of samples (82 of 103) we studied were collected between 2012 and 2016.
175	
176	Trait imaging and quantification
177	We photographed each damselfly using a Nikon D5100 camera (16.2 MP; Nikon
178	Corporation, Tokyo, Japan). We dissected the ventral thoracic cuticle from each female using
178 179	Corporation, Tokyo, Japan). We dissected the ventral thoracic cuticle from each female using forceps and imaged the mesostigmal plates using scanning electron microscopy. Specimens
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179 180	forceps and imaged the mesostigmal plates using scanning electron microscopy. Specimens were mounted on aluminum stubs with carbon tape, sputter-coated with gold-palladium, and
179 180 181	forceps and imaged the mesostigmal plates using scanning electron microscopy. Specimens were mounted on aluminum stubs with carbon tape, sputter-coated with gold-palladium, and imaged at, 200X magnification and 3kV using a Zeiss NEON scanning electron microscope.
179 180 181 182	forceps and imaged the mesostigmal plates using scanning electron microscopy. Specimens were mounted on aluminum stubs with carbon tape, sputter-coated with gold-palladium, and imaged at, 200X magnification and 3kV using a Zeiss NEON scanning electron microscope. To avoid any potential bias during measurements, we blind-coded image files before
179 180 181 182 183	forceps and imaged the mesostigmal plates using scanning electron microscopy. Specimens were mounted on aluminum stubs with carbon tape, sputter-coated with gold-palladium, and imaged at, 200X magnification and 3kV using a Zeiss NEON scanning electron microscope. To avoid any potential bias during measurements, we blind-coded image files before measuring all traits. We measured abdomen length (abdominal segments 1-10, excluding
179 180 181 182 183 184	forceps and imaged the mesostigmal plates using scanning electron microscopy. Specimens were mounted on aluminum stubs with carbon tape, sputter-coated with gold-palladium, and imaged at, 200X magnification and 3kV using a Zeiss NEON scanning electron microscope. To avoid any potential bias during measurements, we blind-coded image files before measuring all traits. We measured abdomen length (abdominal segments 1-10, excluding terminal appendages) on the full-body photos as an estimate for body size using the segmented

188 showed that left plate and right plate sensilla counts were highly correlated within individual 189 females (r = 0.85). In cases where we quantified the left plate, we flipped the image 190 horizontally, so it was in the same orientation as a right plate. We standardized the position of 191 the mesostigmal plate in each image by cropping and rotating the image so that the lower 192 medial corner of the plate was in line with the lower left corner of each image. We counted 193 sensilla and obtained their x and y coordinates in ImageJ using the multi-point selection tool. 194 We traced an outline around the plate image, excluding the lateral carina (Fig. S1), using a 195 Wacom Cintig 12WX tablet and stylus (Wacom, Saitama, Japan) and the freehand selection tool 196 in ImageJ. This procedure produced x and y coordinates that described the plate outline. We 197 performed all measurements twice for each specimen. Measurements across the two technical 198 replicates were highly correlated ($r_{abdomen} = 0.95$, n = 78; $r_{count} = 0.95$, n = 103; $r_{plate area} = 0.99$, n 199 = 86), so we used the mean trait values of the two replicates in subsequent analyses. 200 Seventeen samples were imaged at angles that allowed counting of the sensilla, but distorted 201 the plate shape or distances between the sensilla. Those samples are included in analyses of 202 sensilla number, but were not included in the analyses of sensilla density or distribution. 203

204 Sensilla trait analyses

205 We conducted all morphometric and statistical analyses using R v. 3.4.1 (R Core Team, 206 2015). We used the mesostigmal plate outline coordinates to calculate each plate's two-207 dimensional area. To calculate the area of the sensilla-covered region of each plate, we 208 generated a polygon connecting the coordinates of the outermost sensilla and calculated the 209 area within this outline. We determined the proportion of each plate that was covered by

210 sensilla by dividing the sensilla area by total plate area. We calculated sensilla density in two 211 ways. First, we divided sensilla number by the area of the sensilla-covered region. This 212 measures the number of sensilla that occur in a particular area, but does not capture the 213 relative arrangement of sensilla within that area. Second, we computed the nearest neighbor 214 distances among all sensilla within each plate based on their x and y coordinates and then 215 calculated the mean and median nearest neighbor distances between the sensilla for each 216 female. Nearest neighbor mean and median distances were highly correlated ($r_{E, carunculatum} =$ 0.83; $r_{E,anna} = 0.81$), so we used the mean values for these measures in our analyses. 217 218 To determine whether larger females possess more sensilla, we regressed sensilla 219 number against abdomen length. We found no significant relationship between these traits in either species (*E. anna*: R_{adi}^2 = -0.007, $F_{1.25}$ = 0.82, *P* = 0.737; *E. carunculatum*: R_{adi}^2 = 0.01, $F_{1.48}$ = 220 221 0.52, P = 0.47). We thus present the results that compare sensilla counts without correcting for differences in body size. 222

223

224 Sensilla spatial analyses

To quantify sensilla distributions within each plate, we generated kernel density estimates (KDEs) for populations with at least four sampled individuals (two *E. anna* and six *E. carunculatum* populations) using the R package ks (Duong, 2016). First, we randomly selected one of the two replicate sets of sensilla and plate outline coordinates for each female. To prepare the coordinate data for KDE analyses, we concatenated the sensilla and plate coordinates for each female and adjusted all plate outlines to have an area of one. This standardized each set of sensilla coordinates for size, while maintaining their relative positions

232	within each plate. Next, we translated each set of coordinates to place the origin of the
233	coordinate system at the plate outline's centroid. We concatenated sensilla coordinates for all
234	females sampled within each population to compute a representative KDE for each population.
235	Within each species, we conducted pairwise tests to compare each population's KDE
236	against every other population using the function kde.test with the default settings. This test
237	returns a <i>P</i> -value that reflects the probability of generating the two respective KDEs from the
238	same distribution of points. Because we performed multiple pairwise tests among <i>E</i> .
239	carunculatum populations, we adjusted the resulting <i>P</i> -values using the false discovery rate
240	(Benjamini & Hochberg, 1995).
241	We generated an average plate outline for each population on which to visualize the
242	KDEs. The total number of coordinates that describe each plate outline varied among females,
243	ranging from 647-1078 for <i>E. anna</i> and 688-1028 for <i>E. carunculatum</i> . We standardized the
244	number of coordinates representing each plate by retaining the points for each of the upper
245	and lower medial corners and randomly sampled 198 points in between. We then treated each
246	of these, 200 points as landmarks (the corners represented fixed landmarks and the remaining
247	points were designated as sliding semilandmarks) and used the R package geomorph (Adams
248	and Otarola-Castillo, 2013) to perform general Procrustes analysis (Rohlf, 1999) and obtain an
249	average two-dimensional plate shape for each population.
250	

251 Statistical analyses

252 Some populations were well-sampled whereas others were represented by a single 253 female (Table 1). To avoid psedoreplication, for each population with N > 1, our analyses of

254	sensilla number, density, and area of each mesostigmal plate covered by sensilla used the
255	population mean of each trait value, so that each population was represented by a single
256	measurement. We arcsin transformed proportion data prior to analysis. To compare traits
257	between <i>E. anna</i> and <i>E. carunculatum</i> , we used Welch's <i>t</i> -tests. We compared traits among
258	sympatric, locally allopatric, and fully allopatric E. carunculatum populations using Kruskal-
259	Wallis tests, and between sympatric and locally allopatric <i>E. anna</i> populations using Welch's t-
260	tests. To understand the relationships between sensilla number, sensilla density, and the area
261	of the plate occupied by sensilla, we performed linear regressions between each pair of traits.
262	Due to the limitation of single samples from some populations, we analyzed E. carunculatum
263	populations in two ways: we first included all samples, then conducted a separate analysis that
264	excluded populations with N < 4. Both analyses yield similar results; we report the results for
265	the analysis using all samples in the main text and provide results from the subset of samples
266	with $N \ge 4$ in the supplemental material (Table S1).
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268 Results

269 Enallagma anna and E. carunculatum females possess distinct sensilla traits

Enallagma anna females possessed significantly more sensilla per plate ($\bar{x} = 49 \pm 2$) than E. carunculatum females ($\bar{x} = 28 \pm 1$, $t_{35.1} = 11.13$, $P = 4.6 \times 10^{-13}$; Fig. 2A). Enallagma anna females also possessed sensilla distributed over a larger proportion of each plate ($t_{39.7} = 11.1$, P $= 8.6 \times 10^{-14}$; Fig. 2B), and larger mean distances between sensilla ($t_{54} = 6.7$, $P = 1.3 \times 10^{-8}$; Fig. 2C). This ultimately results in a lower density of sensilla per unit area in *E. anna* compared to *E.* carunculatum ($t_{99.6} = -12.96$, $P = 2.2 \times 10^{-16}$; Fig. 2D). The sensilla also occurred in different

276	locations on the mesostigma	l plates of each sr	pecies: they were	more medially located in F.
1,0	locations on the mesoslighta	i places of each sp	secres, they were	more meaning located in E.

277 *anna* and more laterally located in *E. carunculatum* (Figs. 3, 4).

278 Both species showed a strong positive relationship between sensilla number and the absolute area of the plate occupied by sensilla (*E. anna*: $R^2_{adi} = 0.33$, $F_{1.27} = 14.71$, P = 0.0007; *E.* 279 carunculatum: $R_{adj}^2 = 0.33$, $F_{1,72} = 37.68$, $P = 4.1 \times 10^{-8}$). Consistent with this result, linear 280 281 regressions also revealed that females with more sensilla also had a larger proportion of the plate occupied by sensilla (*E. anna*: $R^{2}_{adj} = 0.26$, $F_{1, 27} = 10.65$, P = 0.003; *E. carunculatum*: $R^{2}_{adj} = 0.26$ 282 0.20, $F_{1.65} = 18.93$, $P = 4.4 \times 10^{-5}$). Females with more sensilla had smaller mean distances 283 between neighboring sensilla (*E. anna*: $R^2_{adi} = 0.11$, $F_{1,27} = 4.34$, P = 0.046; *E. carunculatum*: R^2_{adi} 284 = 0.09, $F_{1.72}$ = 3.80, P = 0.01). Overall, these results show that a greater number of sensilla was 285 286 more strongly associated with a sensilla distribution that covers a larger area of the 287 mesostigmal plate rather than a greater concentration sensilla within in a smaller area. 288

289 *E. carunculatum* sensilla traits do not show a strong pattern of reproductive character

290 displacement

We made several non-mutually exclusive predictions expected under RCD for the sensilla traits we measured in sympatric populations relative to allopatric populations. In particular, we predicted to observe at least one of the following phenotypic differences in sympatric females relative to allopatric females: (1) more numerous sensilla, (2) denser sensilla, (3) sensilla concentrated in different regions of the mesostigmal plates. We did not find significant differences in any of these traits between sympatric and locally allopatric *E. anna* females (Table 2). However, because our *E. anna* samples included only four females from

298 three locally allopatric populations, we could not perform a robust comparison of *E. anna* 299 sensilla traits between populations that do, or do not encounter *E. carunculatum*. We thus 300 focus our analysis on comparisons between sympatric and allopatric E. carunculatum 301 populations, for which we had larger sample sizes. 302 Sympatric, locally allopatric, and fully allopatric *E. carunculatum* populations did not differ significantly from one another in sensilla number (Kruskal-Wallis $\chi^2_2 = 0.69$, P = 0.71), 303 proportion of the mesostigmal plate covered by sensilla (Kruskal-Wallis χ^2_2 = 2.16, P = 0.34), or 304 sensilla density (overall density: Kruskal-Wallis χ^2_2 = 0.12, *P* = 0.94; mean distance between 305 sensilla: Kruskal-Wallis χ^2_2 = 3.53, P = 0.17). In addition to divergence of mean trait values, RCD 306 307 can also result in reduced trait variance in sympatry without affecting the mean (Pfennig and 308 Pfennig 2009). Sympatric *E. carunculatum* populations displayed less interpopulation variance 309 than allopatric populations in both mean sensilla number (Figure 2A) and mean proportion of the plate covered by sensilla (Figure 2B). However, these trends were not statistically 310 significant (sensilla number: Bartlett's $K_1^2 = 0.83$, P = 0.36; proportion of plate covered by 311 sensilla: Bartlett's $K_{1}^{2} = 1.86 P = 0.17$). 312

Interestingly, although mean trait values did not differ significantly between sympatric
and allopatric populations, sensilla traits displayed considerable variation within the
populations we sampled. For example, within a single population, a particular female might
have twice as many sensilla than another female (Fig. 3). This pattern was also observed in the *E. anna* populations we studied.

318 KDE comparisons did not reveal significant differences in sensilla distributions between 319 sympatric and allopatric *E. carunculatum* populations (Table 3). However, the analysis revealed

significant differences in sensilla distributions between several pairs of allopatric *E*.
 carunculatum populations (Fig. 4E), which indicates that populations isolated from *E. anna* vary
 more among themselves than do populations sympatric with *E. anna*, which share similar
 sensilla patterns. This result is consistent with those described above that indicated higher
 variance in sensilla traits among allopatric populations compared to sympatric populations.

325

326 Discussion

Enallagma anna and E. carunculatum females possess different numbers of sensilla in 327 328 species-specific distributions on their mesostigmal plates. This result supports the idea that 329 receptors that receive male stimuli will occur in patterns that correspond to the male organs 330 during contact (Eberhard, 2010). An association between male morphology and female sensilla 331 has been described for African Enallagma species (Robertson and Paterson, 1982), and our 332 results show a similar pattern for two North American species. *Enallagma anna* male cerci are 333 considerably larger than E. carunculatum cerci, and the observation that E. anna females had a 334 larger number of sensilla compared to E. carunculatum females is consistent with the likelihood 335 that *E. anna* male cerci make greater spatial contact with the mesostigmal plates. 336 When species make secondary contact after initial divergence in allopatry, the possible 337 outcomes are increased species divergence (e.g., Sætre et al., 1997; Noor, 2000; Naisbit et al., 338 2001; Yukilevich, 2012; Dyer et al., 2014), decreased species divergence (e.g., Ritchie et al.,

1989; Shurtliff et al., 2013; Yang et al., 2016), local extinction of one species due to

340 reproductive exclusion (Hochkirch et al., 2007, Groning and Hochkirch, 2008), or no change in

341 either direction (Abbott et al., 2013). Because *E. anna* and *E. carunculatum* produce

reproductively disadvantaged hybrids (Barnard et al., 2017), selection is expected to favor increased premating isolation when the species are sympatric. Within each species, we predicted that female sensilla traits in sympatric populations would diverge from those of allopatric populations indicative of a shift in female preferences to avoid mating with heterospecifics. Contrary to this prediction, sympatric and allopatric *E. carunculatum* populations were not significantly different in mean sensilla trait values (Fig. 2) or sensilla density distributions (Fig. 4E).

349 Although we observed a trend toward more sensilla in sympatric *E. anna* populations 350 relative to allopatric populations (Figs. 2A, 3A), it is difficult to conduct a robust comparison for 351 this species because E. anna's entire geographic range overlaps with E. carunculatum's range 352 and E. anna are often relatively rare (Acorn, 2004; A. Barnard, personal obs.). It was therefore 353 difficult to collect sufficient *E. anna* samples from populations that do not co-occur with *E.* 354 carunculatum. We might, however, expect a stronger pattern of RCD in sympatric E. anna 355 females because E. carunculatum males can take them in tandem relatively easily, whereas E. 356 anna males are typically unsuccessful at taking E. carunculatum females in tandem (Barnard et 357 al., 2017). This means that *E. anna* females may have more opportunities for mating mistakes 358 than *E. carunculatum* females, which can result in stronger asymmetric RCD (Lemmon, 2009; 359 Pfennig and Pfennig, 2009).

There are at least three potential explanations for the absence of RCD in the form of significant differences in the sensilla traits we measured between sympatric and allopatric populations of *E. carunculatum*. First, species-specific sensilla distributions may be sufficiently different to allow females to recognize when they are taken in tandem by heterospecific or

364 conspecific males. If this is true, small degrees of variation within the overall species pattern 365 among females might not affect females' species-recognition abilities. Indeed, a recent study 366 found that intraspecific variation in male cercus morphology appears too minor for *Enallagma* 367 females to show strong discrimination among conspecific males that grasp them (Siepielski et 368 al., 2018). Although RCD is most easily facilitated when the trait under selection already differs 369 between species (Pfennig and Pfennig, 2009), these sensilla traits may have already diverged sufficiently enough to preclude strong selection on further divergence. 370 371 Second, it is possible that the external sensilla phenotypes we measured are not 372 representative of proximate female sensory traits, and the variation that directs mating 373 decisions occurs within the female nervous system. For example, individual sensilla might differ in response rate or ability to distinguish different levels of pressure applied by the cerci and 374 375 grasping pressure might differ between males of each species. The direction of mechanosensor 376 deflection is also important for stimulus detection (Keil, 1997), and different species' cercus 377 morphologies may contact sensilla from different angles. Female mate preferences may also be influenced by the relative frequencies with which females encounter heterospecific and 378 379 conspecific males and female sexual experience (e.g., Svensson et al., 2014). 380 Finally, although we did not detect a statistically significant difference between group 381 means, the small differences we observed may still have biological relevance. If gaining just 382 one additional mechanosensor can (at least) double a female's tactile discriminatory power 383 (Gaffin and Brayfield, 2017), then females in a population with a seemingly minor upward shift

in sensilla number could gain a substantial increase in their ability to detect and avoid mating

with heterospecifics. Similarly, it is difficult to determine the features of sensilla density

distributions that may influence female preference solely by conducting statistical tests
 between KDEs. Small spatial differences within largely similar patterns may not contribute a
 signal large enough to be captured in a statistical test, but still reflect salient variation in the
 way females receive tactile stimuli. This might include three dimensional spatial differences
 that we were unable to measure here.

391 These possible explanations highlight the interesting avenues that female damselfly 392 sensilla provide for investigating the mechanisms underlying how females evaluate male tactile 393 signals to make mating decisions. The ability to quantify the number and locations of female 394 mechanoreceptors in a region contacted by male reproductive structures complements our 395 understanding of patterns of variation in male morphologies (McPeek et al., 2008; McPeek et 396 al., 2009; McPeek et al., 2011; Barnard et al., 2017). Females of both species display substantial 397 intrapopulation variation in sensilla traits (Fig. 3) and this variation may play a role in sexual 398 selection and female preferences within species. Behavioral studies will be crucial to link 399 mechanoreceptor phenotypes to female mating decisions and clarify how sensilla traits 400 influence both species recognition and sexual selection. For example, do females with more 401 sensilla make fewer mating mistakes than females with fewer sensilla (Lemmon, 2009)? 402 Another outstanding question of this system is how the cerci stimulate individual sensilla during 403 tandem. This might be determined by flash-freezing male-female tandem pairs and using 404 micro-CT scanning to understand how the male and female structures interact, similar to a 405 recent approach used in seed beetles (Dougherty and Simmons, 2017). Once we understand 406 how cerci contact the sensilla, functional tests of sensilla electrophysiology could reveal how

407 individual sensilla respond to stimulation and indicate whether certain sensilla make greater408 contributions to reproductive decision-making than others.

409	Female preference can drive sexual selection, promote trait divergence, and cause RI
410	between species (Ritchie, 1996). A longstanding presumption in the literature on genital
411	evolution and speciation has been that female reproductive morphologies are less variant or
412	species-specific than male genitalia (reviewed in Shapiro and Porter, 1989). However, recent
413	studies of variation in female reproductive structures suggest that variation does exist among
414	individuals and species (Ah-King et al., 2014), and our data highlight the importance of looking
415	beyond the easily-quantified external morphologies. When male reproductive structure
416	morphologies are obviously divergent, but female morphologies are not, females may possess
417	important variation at neurophysiological levels that affects how they evaluate male tactile
410	at a set of the transfer of the set
418	signals, similar to the way females evaluate signals in other sensory modalities.
418 419	signals, similar to the way females evaluate signals in other sensory modalities.
	Signals, similar to the way females evaluate signals in other sensory modalities.
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419 420	Acknowledgements
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419 420 421 422	Acknowledgements We are grateful to P. Larson for SEM imaging help. We thank O. Fincke, D. Paulson, and B. Mauffray for generously donating damselfly specimens, and D. Gaffin, O. Fincke, R. Knapp, G.
 419 420 421 422 423 	Acknowledgements We are grateful to P. Larson for SEM imaging help. We thank O. Fincke, D. Paulson, and B. Mauffray for generously donating damselfly specimens, and D. Gaffin, O. Fincke, R. Knapp, G. Wellborn, and S. Westrop for helpful discussion during the course of this work. This work was

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664 Figure legends

Figure 1. Sampling sites and species ranges. Enallagma anna's geographic range (red) occurs
within E. carunculatum's geographic range (orange). Names of sites associated with each
number are described in Table 1. Symbol color indicates the species sampled and symbol shape
indicates the population type. (Species ranges are adapted from Johnson, 2009; Paulson, 2009,,
2011).

670

671 Figure 2. Enallagma anna and E. carunculatum sensilla traits by population type. (A) The

number of sensilla on one mesostigmal plate. **(B)** Proportion of the plate that contains sensilla.

673 (C) Mean nearest neighbor distances between sensilla. (D) Sensilla density in the region of the

674 plate that contains sensilla. Within each panel, each open circle represents the mean of one

675 population. Boxplots show the interquartile range. The line within the box shows the median

and whiskers extend to the most extreme observation within 1.5 times the interquartile range.

677

678 Figure 3. Individual trait values for sensilla number, sensilla density, and proportion of plate

679 **containing sensilla.** Each symbol represents a single female, separated by population along the

680 y-axis. Horizontal lines indicate the mean value for each population type (completely allopatric,

locally allopatric, or sympatric), calculated from population means. Populations are described

682 in Table 1.

683

Figure 4. Sensilla locations. (A) White box indicates the location of right mesostigmal plate on
 the thorax. (B) Ultrastructural details of individual sensilla. Scale bar represents 10 μm. (C, D)

686	Scanning electron	micrographs sho	ow the locations of	of sensilla (yellov	v) on the mesostigmal

- 687 plates of *E. anna* (C) and *E. carunculatum* (D). Scale bars represent 100 μm. (E, F) Population
- 688 kernel density estimates for *E. carunculatum* (E) and *E. anna* (F) sensilla. The shading indicates
- 689 different regions of sensilla density: red represents the 75-99th percentile of sensilla density,
- 690 orange represents the 50-74th percentile, and yellow represents the 25th-49th percentile. Each
- 691 outline represents the average mesostigmal plate shape for the population. Asterisks indicate
- 692 *E. carunculatum* populations whose KDEs are significantly different (* *P* < 0.05, *** *P* < 0.001).

Туре	Site (site number*)	Species	Latitude	Longitude	Year	N†	Source [‡]
					collected		
Sympatric	Big Spring, UT (1)	Ea	40.7407	-112.6472	2016	10	AB
		Ec				4	
	Big Sandy Creek, MT	Ec	48.4519	-109.9199	2015	1	AB
	(2)						
	Creston, MT (3)	Ea	48.2437	-114.1406	1972	1	BM
	Dry Sheep Creek, NE	Ea	41.9999	-103.9706	2012	1	BM
	(4)						
	Fish Springs Run, CA	Ea	37.0794	-118.2539	1998	2	BM
	(5)						
	Grace Coolidge	Ea	43.8072	-103.4502	1969	1	BM
	Creek, SD (6)						
	Horseshoe Springs,	Ea	40.6203	-112.7099	2016	1	AB
	UT (7)	Ec				1	
	Long Valley Creek, CA	Ea	39.7315	-120.0434	1973	5	DP
	(8)						
	Murray Creek, NV (9)	Ea	39.2669	-114.8687	2001	1	
	Malad River, UT (10)	Ec	41.8652	-112.1692	1983	2	BM
	Niwot Ditch, CO (11)	Ea	40.1632	-105.1544	2015	2	AB
		Ec				1	

Table 1. Sampling sites for *E. anna* and *E. carunculatum* populations.

	Pondera Coulee, MT	Ea	48.1892	-111.3268	2015	1	AB
	(12)	Ec				1	
Locally	Beaver Creek, WY	Ea	42.6417	-108.3475	2015	1	AB
allopatric	(13)						
	Indian Road Camp,	Ec	46.3336	-111.5254	2015	4	AB
	MT (14)						
	Jackson, WY (15)	Ea	43.5363	-110.7629	1971	2	BM
	Muddy Creek, MT	Ea	47.9796	-112.1565	2015	1	AB
	(16)						
	Strawberry River, UT	Ec	40.1692	-110.4229	2016	1	AB
	(17)						
	West Greenbelt, CO	Ec	39.7742	-105.1350	2014	9	AB
	(18)						
Allopatric	Bull Lake, MT (19)	Ec	48.2262	-115.8404	2015	1	AB
	Crab Creek, WA (20)	Ec	46.8317	-119.8431	2016	20	DP
	Clear Lake, IN (21)	Ec	41.7360	-84.8397	1945	1	BM
	Columbia River, WA	Ec	45.83	-122.77	1952	2	BM
	(22)						
	Douglas Lake, MI (23)	Ec	45.5606	-84.6741	2016	17	OF
	Flathead River, MT	Ec	47.3678	-114.5776	2015	4	AB
	(24)						
	Home Lake, CO (25)	Ec	37.5756	-106.0937	2015	1	AB

Little Lake, CA (26)	Ec	35.9490	-117.9023	1967	1	DP
Drumond Island, MI	Ec	46.00	-83.66	2002	1	BM
(27)						
Snake River, ID (28)	Ec	43.7231	-112.0865	1983	2	BM

⁶⁹⁴ *Site number corresponds to the locations numbered in Figure 1

⁶⁹⁵ ⁺ N: number of females that were imaged and measured for this study.

⁴Sources: A. Barnard (AB), Ola Fincke (OF), Bill Mauffray (BM), and Dennis Paulson (DP).

697 **Table 2.** Statistical comparison of sensilla traits in locally allopatric and sympatric *E. anna*

698 populations.

Mean	<u>+</u> s.e.m.
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	Trait	Local allopatry	t	d.f.	P					
	ITali	Local allopatry	Sympatry	L	u.i.	Ρ				
		(N* = 3)	(N = 10)							
	Sensilla number	39.8 <u>+</u> 3.8	48.5 <u>+</u> 2.3	-1.93	3.6	0.13				
	Proportion plate containing	0.67 <u>+</u> 0.27	0.69 <u>+</u> 0.15	-0.25	2.8	0.82				
	sensilla									
	Sensilla density (sensilla	1.5 <u>+</u> 0.2	1.6 <u>+</u> 0.1	-0.43	2.9	0.70				
	mm⁻¹)									
	Mean distance (µm)	, 20.0 <u>+</u> 1.0	, 19.3 <u>+</u> 0.5	0.62	3.13	0.58				
	between sensilla pairs									
699	* N: number of populations analyzed.									
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709 **Table 3**. Results of pairwise comparisons of sensilla kernel density estimates for *E*.

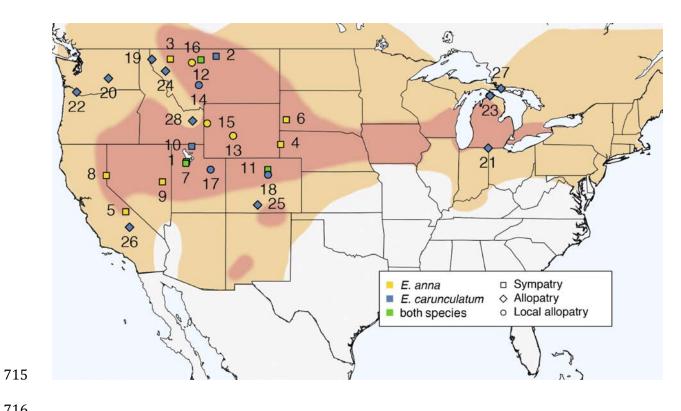
	Big	Crab	Douglas	Flathead	West		
	Springs,	Creek,	Lake, MI	River,	Green-		Population
Population	UT	WA		MT	belt, CO	N*	type
Big Springs, UT	1					4	sympatric
Crab Creek, WA		1				20	allopatric
Douglas Lake, MI	0.263	2.53e ⁻¹⁰	1			17	allopatric
Flathead River, MT	1	0.0103	0.263	1		4	allopatric
							Locally
West Greenbelt, CO	1	0.0625	0.3835	0.502	1	4	allopatric
							Locally
Indian River, MT	1	1	0.0103	0.0625	0.3115	4	allopatric

710 *carunculatum* populations. False discovery rate-adjusted *P*-values are reported[†].

* N: number of females whose sensilla coordinates were used to calculate KDEs.

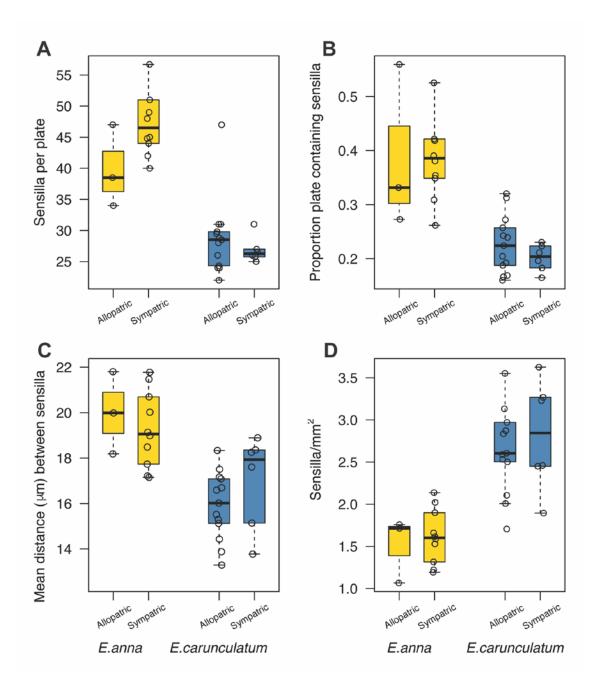
⁷¹² ⁺Bold values indicate *P* < 0.05.

Figure 1.

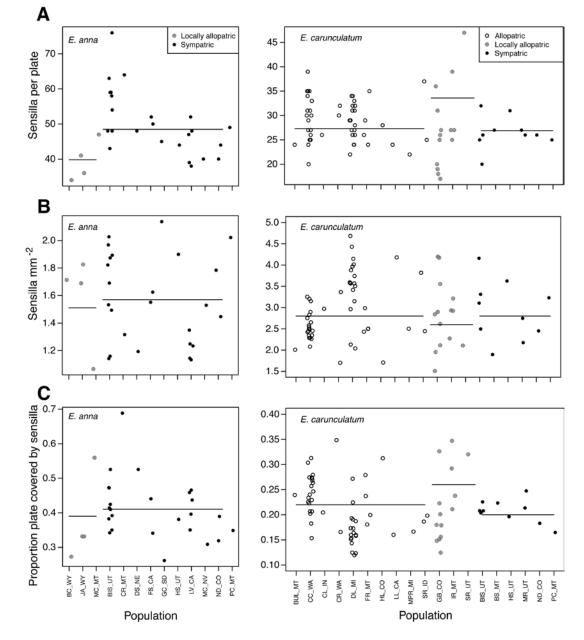


718 Figure 2.

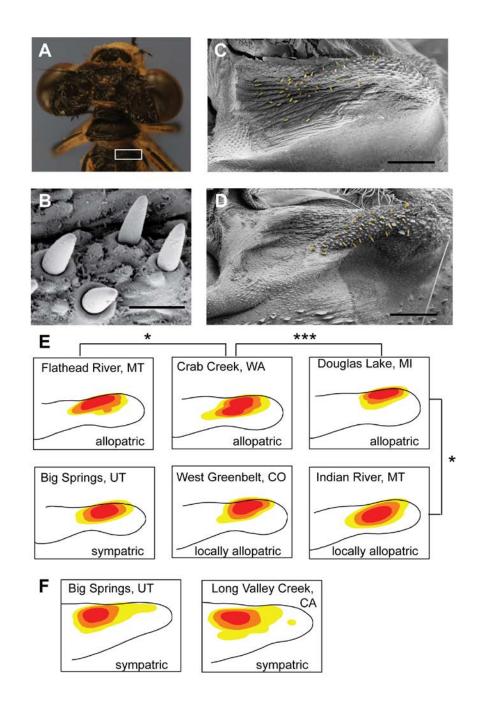
719

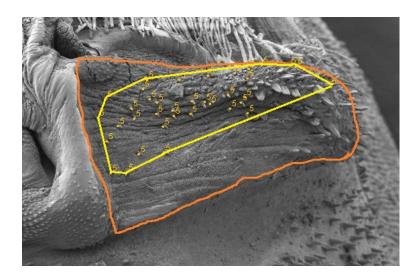


721 Figure 3.



725 Figure 4.



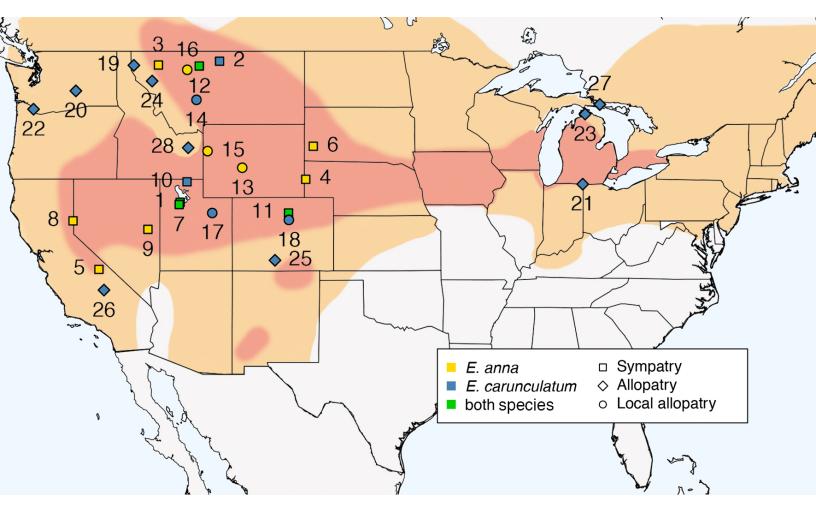


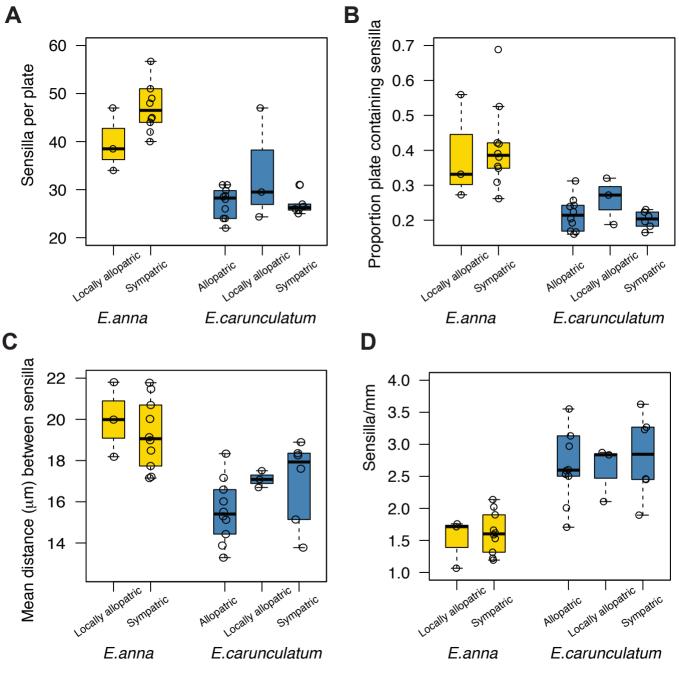


730 **Figure S1.** Method to obtain (x, y) coordinates of mesostigmal plate outline and individual 731 sensilla from scanning electron microscope images. The orange line shows the outline that 732 represents the boundaries of the mesostigmal plate. Yellow dots indicate individual sensilla. 733 The yellow line around the sensilla shows the polygon generated by connecting the outermost 734 sensilla.

- 736 **Table S1.** Statistical comparison of sensilla traits among sympatric, locally allopatric, and fully
- 737 allopatric *E. carunculatum* populations.

			Mean <u>+</u> s.e.m.							
	Trait	Sympatry S	Local allopatry	Allopatry	$\chi^2_2^{\dagger}$	Р				
		(N = 1)	(N = 2)	(N = 3)						
	Sensilla number	25.8	26.9 <u>+</u> 2.6	29.0 <u>+</u> 0.4	2.79	0.25				
	Proportion plate	0.21	0.23 <u>+</u> 0.30	0.21 <u>+</u> 0.23	0.04	0.98				
	containing sensilla									
	Sensilla density (sensilla	15.1	17.3 <u>+</u> 0.2	15.9 <u>+</u> 0.8	2.14	0.34				
	mm ⁻²)									
	Mean distance (µm)	3.27	2.85 <u>+</u> 0.02	2.91 <u>+</u> 0.32	0.86	0.65				
	between sensilla pairs									
738	* N refers to the number of populations analyzed.									
739	$^{\dagger}\chi^{2}_{2}$ refers to the Kruskal-Wallis chi-squared value with 2 degrees of freedom.									
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