

1 **Male genital titillators and the intensity of postcopulatory sexual**
2 **selection across bushcrickets**

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23

24 **Abstract**

25 Animal genitalia are diverse and a growing body of evidence suggests that they evolve
26 rapidly under post-copulatory sexual selection. This process is predicted to be more intense
27 in polyandrous species, although there have been very few comparative studies of the
28 relationship between the complexity of genital structures in males and measures of the
29 degree of polyandry. In some bushcricket families, males possess sclerotised copulatory
30 structures known as titillators, which are inserted into the female's genital chamber and
31 moved rhythmically. Like other genital structures, bushcricket titillators are widely used as
32 important taxonomic characters and show considerable variation across species in structure,
33 shape and the extent to which they are spined. Here, we examine relationships between the
34 presence/absence of titillators, titillator complexity and both mating frequency and the
35 degree of polyandry in bushcrickets, using phylogenetic comparative analyses. Using
36 published sources combined with original observations, data were obtained for the mean
37 level of polyandry, the duration of the male and female sexual refractory periods and the
38 level of complexity of titillators. To analyse data, we fitted phylogenetic generalised least
39 squares models. No significant relationships were found between titillator presence or
40 complexity and either the level of polyandry, duration of the male's sexual refractory
41 period or the ratio of the female and male sexual refractory periods. The duration of the
42 female's refractory period, however, was positively associated with titillator presence and
43 negatively associated with titillator complexity. The data therefore partially support the
44 hypothesis that post-copulatory sexual selection drives genital evolution in this taxon.

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46

47 Genital structures are extremely diverse and are therefore often used as taxonomic
48 characters to separate closely related species (Eberhard 1985, 2010). Genital traits appear to
49 diverge more rapidly than non-genital traits (Rowe and Arnqvist 2012). The selective
50 pressures that drive such rapid evolutionary change in genitalia have been the subject of
51 considerable debate. These include selection for species isolation (the “Lock and Key”
52 hypothesis, reviewed in Mikkola 2008), neutral evolution (Pleiotropy hypothesis) and
53 sexual selection (reviewed in Eberhard 2010; Simmons 2014). The sexual selection
54 hypothesis potentially involves three closely inter-related processes of post-copulatory
55 sexual selection: sperm competition, sexually antagonistic co-evolution and cryptic female
56 choice (Simmons 2014). Strong support for the role of sexual selection in genital evolution
57 comes from comparative studies that have found a positive relationship between indices of
58 the degree of polyandry (and hence the intensity of sexual selection) and the degree of
59 evolutionary divergence or elaboration of genital structures in mammals (Ramm 2007; Orr
60 and Brennan 2016) and insects (Arnqvist 1998, Rowe and Arnqvist 2012).

61

62 Bushcrickets (Orthoptera: Tettigoniidae) are an excellent model taxon in which to test
63 hypotheses relating to the evolution of copulatory structures. They are a diverse and
64 relatively well studied family of insects, which exhibit varying degrees of polyandry
65 (Gwynne 2001; Vahed 2006, 2007). Two types of sclerotized copulatory structures occur in
66 male bushcrickets: the cerci and titillators. The cerci act as genital claspers (Rentz 1972,
67 Vahed et al. 2014). Comparative evidence suggests that sexual conflict over the duration of
68 ejaculate transfer has affected the form and function of the cerci in bushcrickets (Vahed et
69 al. 2014, Lehmann et al. 2016). Titillators (Figure 1) are concealed structures that are

70 inserted into the female's genital chamber prior to spermatophore transfer and are moved
71 rhythmically with contractions of the male's abdomen and phallic complex (Wulff et al.
72 2015, 2017; Wulff & Lehmann 2016). The tips of the male's titillators contact the sensilla-
73 rich, un-sclerotised, soft tissues of the opening of the female's genital chamber during
74 copulation, prior to spermatophore transfer (Wulff et al. 2015, 2017). Like other genital
75 structures, bushcricket titillators are widely used as important taxonomic characters
76 (Chamorro-Rengifo and Lopes-Andrade 2014) and show considerable variation across
77 species in structure, shape and the extent to which they are spined (Vahed et al. 2011).
78 Comparative evidence indicates that the presence of titillators is associated with longer
79 copulation durations (prior to spermatophore transfer) in bushcrickets (Vahed et al 2011),
80 but the relationship between titillator complexity and polyandry has not previously been
81 examined. In fact, we are not aware of any previous studies that have tested for a
82 relationship across species between direct measures of the degree of polyandry and genital
83 complexity in any animal taxon (for studies that have used in-direct measures of the degree
84 of polyandry or used a binary "monandrous versus polyandrous" classification, see Ramm
85 2007; Arnqvist 1998; Rowe and Arnqvist 2012; Orr and Brennan 2016; Kuntner et al.
86 2016). Bushcrickets are one of the few animal groups in which data on the lifetime degree
87 of polyandry are available for a range of species (Vahed 2006).

88

89 Here, we use phylogenetic comparative analyses to test the prediction that genital
90 complexity in males will be positively related to the intensity of sexual selection in
91 bushcrickets. We examine the relationship between titillator complexity and two related
92 measures of the intensity of sexual selection: the lifetime degree of polyandry and the

93 relative potential reproductive rates (the reciprocal of the sexual refractory period) of males
94 and females. Where the male's potential reproductive rate exceeds that of the female (i.e.
95 where the operational sex ratio is male biased), males are assumed to be under stronger
96 sexual selection (Clutton-Brock & Parker 1992, Ahnesjö et al. 2001: see also Kokko et al.
97 2014 for a critical review).

98

99 It should be noted that our study is not designed to distinguish between the different
100 mechanisms of postcopulatory sexual selection (i.e. sperm competition, cryptic female
101 choice and sexually antagonistic co-evolution). One mechanism of postcopulatory sexual
102 selection, however, may be distinguishable because it potentially makes the opposite
103 prediction: If complex genitalia cause damage to the female's reproductive tract (see
104 Crudginton and Siva-Jothy 2000) and thereby delay the female from re-mating, then more
105 elaborate genitalia could be associated with a longer "time out" from mating (and therefore
106 a lower potential reproductive rate in females and a lower degree of polyandry, see
107 Stockley 2002; Kuntner et al. 2016).

108

109 **Methods**

110 Data on the mean lifetime degree of polyandry were obtained for 29 species of tettigoniid
111 bushcrickets from published sources including our own studies (Supplementary Table S1).
112 For the majority of these species, data on polyandry were obtained from counts of
113 spermatoduses within the spermatheca of tettigoniids sampled at the end of the season in
114 the field (Gwynne 1984, Vahed 2006, Vahed et al. 2011, Gwynne & Lorch 2013, Kaňuch et
115 al. 2015; Gao Yong, pers. comm.). Spermatoduses are capsules of sperm that are formed
116 within the female's sperm storage organ (spermatheca) each time she mates (Vahed 2003,
117 2006, Parker et al. 2017). Microsatellite analysis of sperm from spermatoduses in the
118 bushcricket *Pholidoptera griseoptera*, which had mated under natural field conditions,
119 revealed that in over 80% of females, each spermatoduse within the spermatheca was from
120 a different male (Parker et al. 2017). Re-mating between pairs in bushcrickets is unlikely
121 due to the long sexual refractory periods of both males and females that occur after each
122 mating, which are associated with the transfer of the large spermatophores that are typical
123 of this family (Vahed 2007). Because of this, spermatoduse counts can be used to estimate
124 the degree of polyandry in bushcrickets (Gwynne 1984, Vahed 2006, Vahed et al. 2011,
125 Gwynne & Lorch 2013, Kaňuch et al. 2015). Complementary data were obtained from
126 assessments of caged populations (in one species, *Lluciapomaresius stalii*, Bateman 1998),
127 or marked field populations (in six taxa, Heller and von Helversen 1991, McCartney 2010)
128 and, in two species, molecular analysis of sperm in the spermatheca and offspring of field-
129 mated females (Hockham et al. 2004, Simmons et al. 2007). Caged populations could yield
130 an exaggerated degree of polyandry (females may not be able to evade the males as

131 effectively as they would in the field). It should be noted, however, that in the bushcricket
132 species in this study which was examine in caged populations, *L. stalii*, the males are
133 unable to force the female to copulate and the females play an active role in pair formation
134 and in mounting the male to initiate copulation (Bateman 1998). In addition, the degree of
135 polyandry for this species was at the lower end of the spectrum (mean of 2.1 mates), rather
136 than high. Conversely, studies of marked individuals in field populations could under-
137 estimate the degree of polyandry because matings could be missed. The large
138 spermatophore, which remains attached to the female for several hours after copulation,
139 however, means that matings are unlikely to have been overlooked in these studies (Heller
140 and von Helversen 1991, McCartney 2010).

141

142 Data on refractory periods for males and females came from individuals assessed for the
143 time to re-mating in the laboratory (Supplementary Table S1). Individual pairs were
144 assigned to observation cages at a time of day corresponding to the peak activity period for
145 that species (see Vahed 2007 for further details). Data from two Australian species were
146 from mating observations in field cages (Lehmann 2007, Lehmann and Lehmann 2007).
147 These refractory periods correspond with ‘time-outs from matings’ and are therefore tightly
148 linked with the concept of potential reproductive rate ($= 1/\text{time-out}$, Clutton-Brock &
149 Parker 1992, Ahnesjö et al. 2001).

150

151 **Titillator structure**

152 Data on titillator morphology (Figure 1) were obtained from taxonomic sources, chiefly
153 Harz (1969) for the majority of European species (Supplementary Table S1). We developed

154 a ranked classification system to reflect titillator complexity, with emphasis on the
155 complexity (e.g. presence/absence of visible spines; clumping of spines at the tip; single or
156 double pair of projections) of the apical part of the titillators (median projection) which
157 makes contact with the female during copulation (Wulff et al. 2015, 2017) (see Figure 1,
158 Table 1). Titillators were scored blind (by both the first and the last authors) for a previous
159 study (Vahed et al. 2011), i.e. without knowledge of polyandry or duration of male or
160 female refractory periods.

161

162 **Analysis**

163 We split “titillator complexity” into two components analysed in parallel: presence/absence
164 of titillators, analysed as a binary variable (henceforth titillator presence), and complexity
165 of titillators where present (henceforth titillator complexity), analysed as a continuous
166 variable. This is because it would make little sense to treat species without titillators
167 effectively as bearing “titillators of zero complexity”.

168

169 All analyses were conducted in R 3.2.0 (R Core Team 2015). To account for non-
170 independence due to phylogenetic relatedness, we used comparative-phylogenetic statistical
171 methods that account for this non-independence by modelling errors in the model residuals
172 as a function of the phylogenetic distance between species and the underlying model of
173 how traits evolve along branches. We analysed continuous data (refractory periods) using
174 phylogenetic generalized least squares models (PGLS; Grafen, 1989, Martins and Hansen,
175 1997) using the `ppls()` function in the *caper* package (Orme et al. 2013), simultaneously
176 estimating Pagel's λ , i.e. how closely the tree structure can predict covariance among the

177 model's residuals (Pagel 1999, Freckleton et al. 2002). For analysis of binary or Poisson
178 data (titillator presence/absence and polyandry, respectively) we fitted Generalised
179 Estimating Equation (GEE) models (Paradis & Claude 2002), which are more appropriate
180 for non-normally distributed data (Paradis 2006), using the `compar.gee()` function in the
181 *ape* package (Paradis et al 2004). We used standard residual plots to assess model fit by
182 eye.

183

184 We initially tested whether, within our dataset, polyandry was associated with the duration
185 of the female or male refractory periods and whether female and male refractory periods
186 were associated, as would be predicted (Vahed 2006, 2007). We first fitted a PGLS model
187 of polyandry with female or male refractory period as a predictor variable, then a PGLS
188 model with female refractory period as a response and male refractory period as a predictor.
189 Data on polyandry were log-transformed before analysis to improve model fit. Then, for
190 each analysis of titillator structure (binary GEE models of presence; continuous PGLS
191 models of complexity) we fitted models against predictor variables including polyandry,
192 female refractory period and male refractory period. We performed analyses of potential
193 reproductive rate directly on male and female reproductive rate, as this improved model fit.
194 We additionally ran analyses including the reciprocal of the ratio of the male and female
195 refractory period as a predictor variable. This represents the relative reproductive rates of
196 males and females (a measure of the operational sex ratio).

197

198 To avoid confounds due to variable measuring techniques, we re-ran all analyses excluding
199 species where polyandry was assessed using molecular rather than field-based methods (*R.*

200 *verticalis*, *E. ephippiger*) or where field rather than lab-based methods were used to assess
201 the male refractory period (*P. affinis*, *P. v. veluchianus*) or the female refractory period (*P.*
202 *affinis*, *P. v. veluchianus*, *P. v. minor*). See Supplementary Table S1 for details.

203

204 The phylogeny used for all analyses was based primarily on the molecular phylogeny by
205 Mugleston et al. (2013) supplemented by the molecular phylogenetic information derived
206 from barcoding of Central European species (Hawlitschek et al. 2016), combined with the
207 morphological phylogeny developed by Naskrecki (2000) (this phylogeny did not use
208 titillators as a character). For the sub-family Tettigoniinae, we used the morphological
209 phylogeny provided by Rentz and Coless (1990) (majority consensus tree of 50 equally
210 short cladograms) because many of the genera were neither included in Mugleston et al.
211 (2013) nor in Naskrecki (2000). For phylogenetic relationships within the genus
212 *Anonconotus* (Tettigoniinae), we used an unpublished molecular phylogeny based on
213 mtDNA (R. Szabo, G. Carron, K. Vahed & M. Ritchie). For the genus *Poecilimon*
214 (Phaneropterinae), we used the molecular phylogeny given in Ullrich et al. (2010), for the
215 *Poecilimon propinquus*-group the mtDNA tree of Lehmann (1998). Branch lengths were
216 not available and so we ran all analyses twice, arbitrarily scaling branches according to
217 node depth, following Grafen (1989), or setting all branch lengths to 1. We conducted our
218 analyses on the datasets for which relevant data were available (n=48). For analyses
219 involving subsets of the full dataset, the full tree was pruned to the appropriate set of taxa
220 only after arbitrary branch lengths had been assigned, in order to preserve the node depths
221 in the full tree. The datasets supporting this article have been uploaded as Supplementary
222 Table S1.

223

224 **Results**

225 Our analyses returned similar results regardless of whether we excluded minority methods,
226 or how we assigned branch lengths (arbitrarily to 1 or scaling according to node depth).
227 Thus, we present only results using all available data regardless of the method used to
228 obtain them, and using Grafen's (1989) node-depth scaling to assign arbitrary branch
229 lengths to the tree.

230

231 *Polyandry*

232 There was a high variation in polyandry in bushcricket females, ranging from as low as a
233 mean of 1.5 matings up to a mean of 25 to 28 matings per female in *Anonconotus* species
234 (Figure 2). The data were largely biased towards species with relatively low polyandry, i.e.
235 between 1.5 and 3.1 numbers of matings over the female lifespan. Polyandry was
236 significantly negatively associated with the duration of the female refractory period (GEE,
237 $df=1$, $P=0.013$, $n=12$) and the male refractory period (GEE, $df=1$, $P=0.047$, $n=14$). Female
238 and male refractory periods were positively associated with each other ($F_{1,25}=18.23$,
239 $p<0.001$, $n=27$); in this model Pagel's λ had wide confidence intervals (0 to 0.906),
240 indicating low certainty in the estimate of phylogenetic signal.

241

242 In the full dataset, titillators were absent in 15 species and present in 32; titillator
243 complexity in species with titillators was distributed as shown in Figure 2. Polyandry was
244 related neither to titillator presence (PGLS: $F_{1, 27}=0.0142$, $P=0.91$, $n=29$; Figure 3a) nor to

245 titillator complexity (PGLS, $F_{1, 20}=2.56$, $P=0.13$, $n=22$; Figure 3b) and vice versa:
246 polyandry was a good predictor neither of titillator presence (GEE, $df=1$, $p=0.98$, $n=29$) nor
247 complexity (PGLS, $F_{1, 20}=2.56$, $P=0.13$, $n=22$). Estimates of Pagel's λ ranged from 0.94 to
248 1.00 [95% CIs $\sim 0.7 - 1$] for these models, indicating a high degree of phylogenetic signal,
249 i.e. covariance in model residuals could be predicted by phylogenetic distance.

250

251 *Male refractory period*

252 The male refractory period can span over several days, the most extreme mean data coming
253 from *Antaxius hispanicus* with 7 days, and *Poecilimon thessalicus* with 6 days. In contrast,
254 several species are able to re-mate within an hour down to a few minutes. Titillator
255 presence did not reliably depend on the male refractory period (PGLS: $F_{1, 30} = 0.51$, $P =$
256 0.41 , $n=32$, Figure 3c) but titillator complexity showed a marginal trend suggesting that
257 species with longer male refractory periods have more complex titillators (PGLS:
258 $F_{1,18}=3.71$, $P=0.069$, $n=20$, Figure 3d). The phylogenetic signal estimate had wide
259 confidence intervals ($\lambda\sim 0.81$ [CI 0.28, 0.96]). The male refractory period did not predict
260 titillator presence (GEE: $df=1$, $P=0.49$, $n=32$) but again showed a marginally positive
261 association with titillator complexity (PGLS: $F_{1, 18}=3.66$, $P=0.072$, $n=20$, $\lambda=0.77$ [CI 0.33,
262 0.95]).

263

264 *Female refractory period*

265 The female refractory period in most species was typically longer than that of the male,
266 ranging from 1.07 h to 19 days. Unlike the male refractory period, the female refractory

267 period was positively associated with titillator presence (PGLS: $F_{1,26}=9.04$, $P=0.005$, $n=28$;
268 $\lambda=0.62$, Figure 3e) and negatively associated with titillator complexity (PGLS:
269 $F_{1,13}=18.052$, $n=15$, $P<0.001$; $\lambda=0$, Figure 3f). Analysing the inverse relationships, the
270 female refractory period did not predict titillator presence (GEE, $df=1$, $P=0.15$, $n=28$) but
271 was a good predictor of titillator complexity ($F_{1,13}=18.052$, $P<0.001$, $n=15$; $\lambda=0$).

272

273 *Male vs female potential reproductive rates*

274 The Operational Sex Ratio (measured as the reciprocal of the ratio of female and male
275 refractory periods) was not associated with titillator presence (PGLS: $F_{1,25}=1.164$, $P=0.16$,
276 $n=27$, $\lambda=0.85$; Figure 3g) nor with titillator complexity (PGLS: $F_{1,13}=0.94$, $P=0.35$, $n=15$;
277 $\lambda=0.60$; Figure 3h). The inverse relationships were similarly nonsignificant, as the
278 Operational Sex Ratio predicted neither titillator presence (GEE: $df=1$, $P=0.73$, $n=27$) nor
279 complexity (PGLS: $F_{1,13}=1.00$, $P=0.34$, $n=15$; $\lambda=0.66$).

280

281

282 **Discussion**

283 No significant relationships were found between titillator complexity and any of the degree
284 of polyandry in bushcrickets, the duration of the sexual refractory period in males, or the
285 relative potential reproductive rates of males and females. The duration of the female's
286 sexual refractory period, however, was associated with both titillator presence and
287 complexity. The results, therefore, provide partial support for the hypothesis that post-
288 copulatory sexual selection is a driving force in titillator evolution. It should be noted,
289 however, that while there was a high degree of variation between species in the degree of
290 polyandry (ranging from a mean of 1.5 to a mean of 28 matings per lifetime), there were no
291 entirely monandrous species in this data set. Any effect of post-copulatory sexual selection
292 on genital evolution should be most detectable by comparing monandrous with polyandrous
293 species (Arnqvist 1998). On the other hand, previous studies have found evidence for a
294 positive relationship between indicators of the degree of polyandry, measured on a
295 continuous scale, and the complexity or degree of development of the males' genitalia in
296 both insects and mammals (Ramm 2007, Rowe and Arnqvist 2012, Orr and Brennan 2016).

297

298 The duration of the female's sexual refractory period was significantly longer in species in
299 which titillators were present. On the face of it, this appears to be opposite to the prediction
300 of the sexual selection hypothesis, which predicts that titillators should be more complex in
301 species in which females are more polyandrous (and therefore have *shorter* sexual
302 refractory periods, since there was a significant negative relationship between the duration
303 of the female's sexual refractory period and the lifetime degree of polyandry in our present

304 study). The results are, however, consistent with the hypothesis that complex genitalia
305 delay or deter the female from remating, which can be driven by various mechanisms of
306 post-copulatory sexual selection. Stockley (2002), for example, found that in primates,
307 relatively high penile spinosity was associated with lower potential reproductive rates in
308 females and interpreted this in terms of internal damage caused to the female by the spines.
309 Kuntner et al. (2016) similarly found that in nephilid spiders, the male's genitalia were
310 more complex in polyandrous species than in monandrous species. Titillators in some
311 bushcrickets have spines (see Figure 1) that contact the soft, un-sclerotised lining of the
312 female's bursa copulatrix (Wulff et al. 2015, 2017). However, we found no evidence for
313 any damage by the titillators to the female's reproductive tract (Wulff et al. 2015, 2017;
314 Wulff and Lehmann 2016), in contrast to the action of penile spines in *Callosobruchus*
315 beetles (Hotzy and Arnqvist 2009). Comparative evidence suggests that genital damage
316 selects for females to evolve a thicker wall of the bursa copulatrix to minimise damage by
317 the male's spines in seed beetles (Coleoptera: Bruchidae) (Rönn et al. 2007). The thickness
318 of the parts of the female contacted by the male's genitalia was not measured in the present
319 study. There is strong evidence from other comparative studies of other insect taxa that co-
320 evolution between male and female genital structures does occur (Rönn et al. 2007, Yassin
321 and Orgogozo 2013, reviewed in Simmons 2014) but for several reasons female genitalia
322 are still understudied (Ah-King et al. 2014, Brennan and Prum, 2015) even in bushcrickets.

323

324 Amongst species with titillators, shorter sexual refractory periods were associated with
325 more complex titillators. These results appear to be in contrast to the analysis based on the
326 presence/absence of titillators (see above). Shorter sexual refractory periods were

327 associated with a higher degree of polyandry (this study), so may be used as an indicator of
328 the degree of polyandry. Comparative studies of water-striders (Heteroptera: Gerridae)
329 have similarly found relationships between indices of the degree of polyandry and genital
330 complexity (Rowe and Arnqvist 2012). The results of the present study should be
331 interpreted with caution, however. The relationship between the duration of the female's
332 sexual refractory period and titillator complexity appear to be driven by the two species in
333 the sub-family Zaprochilinae in our dataset, which have unusually long female sexual
334 refractory periods (Simmons and Gwynne 1991; Lehmann and Lehmann 2007) and simple
335 titillators (Rentz, 1993). This subfamily is phylogenetically distinct from the majority of
336 other Tettigoniid families. It should also be noted that we did not find a significant
337 relationship between the actual degree of polyandry itself (rather than the female's sexual
338 refractory period) and titillator complexity in the present study (the two analyses were
339 based on slightly different subsets of species, see Figure 2).

340

341 It is possible that the classification scheme of titillators used in the present study does not
342 reflect adequately the characters of the titillators that are subject to sexual selection. An
343 alternative approach would be to use more complex morphometric analyses to quantify the
344 shape and/or to measure the relative sizes of parts of the titillators (see, for example Rowe
345 and Arnqvist 2012). In addition, denser sampling of species within selected titillator-
346 possessing genera would allow for a more fine-scale examination of the relationship
347 between titillator morphology and polyandry. In the present study, most titillator-possessing
348 genera were represented by only three species or fewer.

349

350 Although the present study provided mixed support for sexual selection hypotheses for
351 genital evolution, experimental manipulations, in which either one or both titillators were
352 removed, have provided some support for the role of cryptic female choice (Wulff et al.
353 2015, Vahed 2015, Wulff and Lehmann 2016). The results of these manipulations indicated
354 that correct stimulation by titillators is necessary for the proper insertion of the
355 spermatophore and to avoid resistance by females during copulation (Wulff et al. 2015,
356 2017; Wulff and Lehmann 2016). Such results, however, are also consistent with the “Lock
357 and Key” hypothesis, a hypothesis that was not tested in the present study. Simmons (2014)
358 pointed out that female choice that enforces species isolation and female choice that targets
359 variation in male quality within populations may be seen as part of the same continuum.

360 To conclude, the present comparative study provided only partial support for the hypothesis
361 that post-copulatory sexual selection has driven the evolution of titillator complexity in
362 bushcrickets. The inclusion of monandrous species in the sample, examination of the parts
363 of the female’s reproductive tract that are contacted by the titillators and denser sampling of
364 selected genera, however, would be necessary to test the sexual selection hypotheses more
365 fully.

366

367

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371

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544
545

546 **Figure legends**

547 **Figure 1:** Left hand side: Electron micrograph of Titillator tips with spines in the tettigoniid
548 bushcricket *Metrioptera saussuriana*; Right hand side: Variation in titillator morphology in
549 the Tettigoniidae. The numbers represent the system of titillator classification used in the
550 present study (see Table 1). Images adapted from Rentz and Birchim (1968) and Rentz
551 (1993). Note that titillator types 2 and 3 are shown together because they are similar in
552 overall structure (although the apical part that contacts the female, the median projection, is
553 much less strongly projecting in type 2 than in type 3).

554

555 **Figure 2:** Variation of male refractory period (MRP), female sexual refractory period
556 (FRP), their difference, and lifetime degree of polyandry, across the bushcricket phylogeny.
557 Data have been scaled such that open and closed circles represent the minimum and
558 maximum in the dataset, respectively. For raw data, see Supplementary Table S1.

559

560 **Figure 3:** a) Lifetime polyandry in bushcricket species lacking titillators (“No”) and with
561 titillators (“Yes”); b) Polyandry against titillator complexity (ordinal ranked scale, see
562 Table 1); c) The duration of the male sexual refractory period in bushcricket species lacking
563 titillators (“No”) and with titillators (“Yes”); d) The duration of the male sexual refractory
564 period against titillator complexity; e) The duration of the female sexual refractory period
565 in bushcricket species lacking titillators (“No”) and with titillators (“Yes”); f) The duration
566 of the female sexual refractory period against titillator complexity; g) Operational sex ratio
567 (measured as the reciprocal of the ratio of the female and male refractory periods) in

568 bushcricket species lacking titillators (“No”) and with titillators (“Yes”); h) Operational sex
569 ratio against titillator complexity.

570

571

572 **Table 1:** The titillator classification scheme used in this study (adapted from Vahed et al.
 573 2011).
 574

Numerical classification	Explanation	Examples
0	Titillators absent	<i>Poecilimon</i>
1	No sclerotised titillators, but a densely covered field of small tubercles	<i>Kawanaphila</i>
2	One pair of sclerotised titillators, apical part (median projection) conical and not strongly projecting, may have minute teeth	<i>Ruspolia, Yersinella</i>
3	One pair of sclerotised titillators: apical part strongly projecting, with no teeth (the tip however can be hooked)	<i>Metrioptera roeselii</i>
4	One pair of sclerotised titillators: apical part strongly projecting with clearly visible teeth	<i>Anonconotus, Decticus</i>
5	One pair of sclerotised titillators: apical part strongly projecting with teeth concentrated on the club shaped tip	<i>Metrioptera saussuriana</i>
6	Two pairs of sclerotised titillators: apical part strongly projecting with teeth on one or both pairs.	<i>Gampsocleis, Antaxius</i>

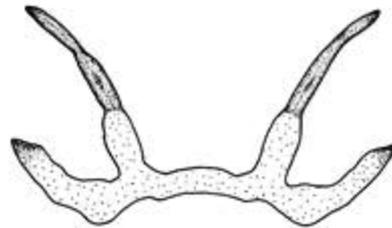
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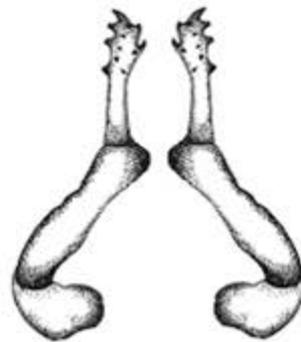
577 **Supplementary Table S1:** Data base for levels of titillator complexity, polyandry and male
 578 and female refractory periods.



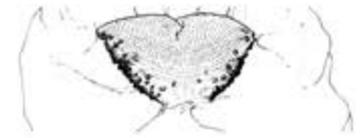
Titillators absent (0)



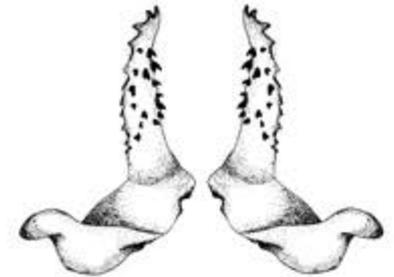
One pair, no teeth (2-3)



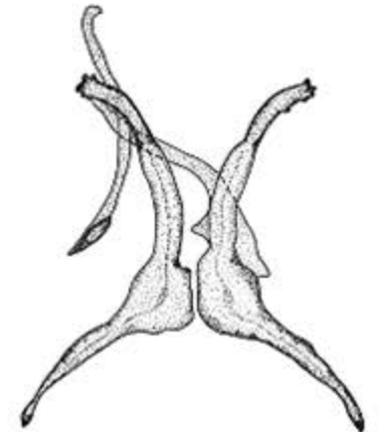
One pair, spines on club shaped tip (5)



Small tubercles (1)



One pair, with teeth (4)



Double pairs of titillators (6)

- Most complex titillators
- Longest MRP
- Longest FRP
- Longest F–M difference
- Greatest polyandry

