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1 **Transgenerational effects of maternal sexual**
2 **interactions in seed beetles**

3

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18

19

20 **Keywords**

21 Intergenerational effects, cross-generational effects, sexual conflict, multiple mating,

22 polyandry, net fitness, inclusive fitness, indirect costs, indirect benefits, maternal

23 effects, multigenerational economics, non-genetic inheritance, lifespan, reproductive

24 success

25 **Abstract**

26 Mating often bears large costs to females, especially in species with high levels of sexual
27 conflict over mating rates. Given the direct costs to females associated with multiple
28 mating, which include reductions in lifespan and lifetime reproductive success, past
29 research focused on identifying potential indirect benefits (through increases in offspring
30 fitness) that females may accrue. Far less attention has, however, been devoted to
31 understanding how costs of sexual interactions to females may extend across
32 generations. Hence, little is known about the transgenerational implications of variation
33 in mating rates, or the net consequences of maternal sexual activities across generations.
34 Using the seed beetle, *Callosobruchus maculatus*, a model system for the study of sexual
35 conflict, we investigate the effects of mating with multiple males versus a single male, and
36 tease apart effects due to sexual harassment and those due to mating *per se*, over three
37 generations. A multigenerational analysis indicated that females that were exposed to
38 ongoing sexual harassment and who also were permitted to mate with multiple males
39 showed no difference in net fitness compared to females that mated just once without
40 ongoing harassment. Intriguingly, however, females that were continually harassed, but
41 permitted to mate just once, suffered a severe decline in net fitness compared to females
42 that were singly (not harassed) or multiply mated (harassed, but potentially gaining
43 benefits via mating with multiple males). Overall, the enhanced fitness in multiply mated
44 compared to harassed females may indicate that multiple mating confers
45 transgenerational benefits. These benefits may counteract, but do not exceed (i.e. we
46 found no difference between singly and multiply mated females), the large
47 transgenerational costs of harassment. Our study highlights the importance of examining
48 transgenerational effects from an inclusive (looking at both indirect benefits but also

49 costs) perspective, and the need to investigate transgenerational effects across several
50 generations if we are to fully understand the consequences of sexual interactions, sexual
51 conflict evolution, and the interplay of sexual conflict and multi-generational costs and
52 benefits.

53 **Introduction**

54 Sexual interactions usually bear large costs on the participants. Often, investment in
55 current reproduction trades off against future reproduction and lifespan (Reznick, 1985;
56 Williams, 1966). While sexual interactions and mating are necessary to ensure
57 fertilisation in sexually reproducing organisms, and hence are the cornerstone for the
58 production of progeny, associated costs can be substantial. Specifically, females of many
59 species incur large direct costs; elevated mating rates can substantially depress fecundity
60 and longevity for females (Arnqvist and Nilsson, 2000; Blanckenhorn et al., 2002;
61 Crudginton and Siva-Jothy, 2000; Gavrilets et al., 2001). This has been particularly well
62 documented in the fruit fly *Drosophila melanogaster*, where seminal proteins that are
63 transferred with the male ejaculate decrease female longevity (Chapman et al., 1995).
64 However, even co-habitation and harassment without successful copulation have been
65 shown to be detrimental for female fitness in this species (Partridge and Fowler, 1990).

66

67 Mating at a higher frequency than is required to fertilise a complete set of ova can be a
68 result of sexual conflict over mating rates, which is common in the animal kingdom
69 because of strong selection on males to maximise their reproductive success (Arnqvist
70 and Rowe, 2005; Chapman et al., 2003; Parker, 2006). High rates of female sexual
71 interactions may also evolve adaptively if direct benefits, such as mating gifts or paternal
72 care, are gained (Arnqvist and Nilsson, 2000), or if females benefit indirectly (via genetic
73 benefits) by producing fitter offspring as a result of elevated sexual interactions and
74 matings with multiple males (Jennions and Petrie, 2000; Kokko et al., 2003). In many
75 species, there are no apparent direct benefits associated with mating with multiple males
76 (Arnqvist and Kirkpatrick, 2005; Jennions and Petrie, 2000), and for this reason, the

77 majority of research in this field has focussed on whether the direct costs associated with
78 multiple mating can be compensated by the production of fitter offspring (Chapman et
79 al., 2003; Holland and Rice, 1998; Kokko et al., 2003).

80

81 Theory predicts that indirect genetic benefits are unlikely to outweigh the direct costs
82 incurred by females (Cameron et al., 2003). Empirically this has been supported by a
83 range of studies (see Arnqvist and Nilsson, 2000), for example in the common lizard
84 *Lacerta vivipara* (Le Galliard et al., 2008) and in the fruit fly *D. melanogaster* (Brommer
85 et al., 2012; Orteiza et al., 2005; Stewart et al., 2008, 2005). Nonetheless, there are also
86 studies in *D. melanogaster* acknowledging major fitness benefits of mating with multiple
87 males due to genetic benefits (i.e. indirect benefits). For example, more fecund *D.*
88 *melanogaster* daughters compensate for the direct costs of mating incurred by their
89 mothers (Priest et al., 2008b, 2008a). Mating multiple times with different males may
90 allow females to mate with more attractive mates, generating genetic benefits. Some
91 studies show that fitness benefits via attractive sons may indeed outweigh direct costs,
92 such as in the house cricket *Acheta domesticus* (Head et al., 2005), and *D. melanogaster*
93 (Rundle et al., 2007). Similarly, benefits in the form of increased offspring viability cancel
94 out the direct costs (decrease in female longevity) of mating in the Australian field cricket,
95 *Teleogryllus oceanicus* (Garcia-Gonzalez and Simmons, 2010).

96

97 Recent studies showing that transgenerational costs may exacerbate the direct costs to
98 females (Dowling et al., 2014; Gasparini et al., 2012), or invoke opposing effects across
99 different generations (Brommer et al., 2012) add further insights into understanding the
100 fitness consequences of sexual interactions.. These transgenerational effects (TGE),
101 which may be inherited non-genetically (e.g., including epigenetic mechanisms), include

102 indirect genetic effects (IGE), and encompass maternal effects, paternal effects and effects
103 via interacting phenotypes (i.e. non-sire influences on offspring life history trajectories;
104 see Garcia-Gonzalez and Dowling, 2015; García-González and Simmons, 2007). TGEs have
105 been identified as important factors influencing the fitness of offspring across
106 generations following sexual interactions. Recent studies have demonstrated
107 transgenerational costs to females that are brought about by sexual interactions: the
108 effects of heightened (in both intensity and frequency) sexual interactions and increased
109 harassment lead not only to longevity costs in female *Drosophila melanogaster*
110 themselves (direct costs), but also to longevity costs in their offspring, adding therefore
111 a transgenerational cost (Dowling et al., 2014). Similarly, higher levels of male sexual
112 harassment in female guppies (*Poecilia reticulata*), led to lower reproductive success for
113 their sons and daughters (Gasparini et al., 2012). Moreover, a study in *D. melanogaster*
114 reported that females that were exposed to mating at different rates, produced sons with
115 increased fitness, but grandsons with decreased fitness (Brommer et al., 2012). Opposing
116 effects in descendants of different sexes may be due to negative genetic correlations for
117 fitness between the sexes, or parents and offspring. Such negative genetic correlations
118 have been reported in *D. melanogaster* (Brommer et al., 2012; Chippindale et al., 2001;
119 Pischedda and Chippindale, 2006), and also in other species such as the southern ground
120 cricket *Allonemobius socius* (Fedorka and Mousseau, 2004) and red deer (*Cervus elaphus*,
121 Foerster et al., 2007). These studies highlight the importance of considering the sex-
122 specific nature of cross-generational costs and benefits, because the benefits of mating
123 with multiple males may disappear due to conflicting effects across generations or due to
124 opposing effects within the sexes.

125

126 Here, we report effects of maternal mating history on female lifetime reproductive
127 success across three generations, and on offspring and grand-offspring longevity, in both
128 sexes, in the seed beetle *Callosobruchus maculatus*. Specifically, after an initial baseline
129 mating, which rendered females non-virgins, we exposed females to one of three
130 maternal mating treatments. These were a treatment of no further male exposure (single
131 mating), a treatment of harassment by multiple emasculated males incapable of
132 insemination, and a treatment of multiple mating with multiple males capable of
133 harassing females and successfully inseminating them. We investigated differences
134 across treatments in offspring production in each generation separately, and also
135 calculated the net consequences of each of the mating treatments by examining female
136 offspring production across all three generations, to gain an understanding of the
137 multigenerational economics of maternal sexual interactions. We discuss how exposure
138 to sexual interactions may influence the evolution of mating systems, and the importance
139 of these interactions and ensuing TGEs for population growth rates. Our study highlights
140 the effects of non-genetic inheritance and the transgenerational consequences of sexual
141 interactions on net fitness and population growth rates.

142 **Methods**

143 We used virgin male and female seed beetles (*Callosobruchus maculatus*) in our
144 experiments. These beetles were sourced from an outbred population (South Indian
145 stock population, SI, obtained from a replicate held at Uppsala University and prior to this
146 kept by C. W. Fox at the University of Kentucky), which exhibits substantial phenotypic
147 and genetic variance for a range of traits and behaviours (see for instance Fox et al., 2003;
148 Berg and Maklakov, 2012; Berger et al., 2014; Bilde et al., 2008). The stock population at

149 Doñana Biological Station was established in 2013 using more than 450 founders and has
150 been cultured since then in non-overlapping generations on organic mung beans (*Vigna*
151 *radiata*) that are frozen prior to use. The stock population is kept across multiple
152 containers, each of which typically generates over a thousand adults per generation.
153 Around 50 non-virgin adults (25 males and 25 females) are randomly selected in each
154 container each generation and allowed to reproduce in a new container with uninfested
155 beans. The effective population size for each replicated population exceeds 75
156 individuals, as the 50 adults are non-virgins collected from containers with
157 approximately 1000 individuals and females mate multiply. The high rates of female
158 multiple mating in these populations mean that our estimate of N_e is likely to be an
159 underestimate. Offspring from the different containers are admixed and redistributed
160 every few generations, and thus the stock population is maintained at large population
161 sizes (in excess of 300 individuals). Beetles are kept in walk-in climate chambers
162 (Fitoclima 10000 EHF, Aralab) at a constant 29°C temperature with 40 % humidity and a
163 12hour/12hour light/dark cycle.

164

165 ***Maternal mating treatment***

166 We individually paired 120 virgin females and males and allowed them to mate once (Day
167 0). Seven pairs were excluded, as they did not mate in the time allocated (30 s). After
168 mating, each female was transferred immediately into an empty 30 ml container. On day
169 1, the F0 females were separated at random into three treatment groups: 1) single mating
170 (monogamous treatment, M) – kept as is, with no further interactions allowed, 2)
171 harassment (H) – four males that had been incapacitated to mate were added to each
172 female's container. Incapacitation was conducted on five day old males, under CO₂

173 anaesthetisation. Relaxation due to anaesthesia led to the eversion of the male aedeagus,
174 which was surgically shortened by approximately 1/3 in length using microscissors,
175 removing the spiny tip of the aedeagus. The efficacy of this procedure was confirmed in
176 preliminary tests: males did not achieve successful copulations but continued to harass
177 females and attempt mating. In the last treatment, 3) multiple mating (polyandrous
178 treatment, P) – each female was placed with four same-age stock males (who were not
179 emasculated but were briefly anesthetized, similarly to males used in the H treatment,
180 prior to their use) that could both harass and successfully mate with females.

181 Females were kept in their respective treatments until day 4 and the containers were
182 checked daily for dead males, which were replaced immediately. While being kept
183 without beans can suppress both egg laying as well as remating rates in the study species
184 (Eady et al., 2004), continuous exposure to males is expected to lead to highly elevated
185 harassment and remating rates. For instance, Eady et al (2004) found that even under
186 conditions of suppressed oviposition between 20-60 % of females readily remated during
187 a relatively short (<45min) second mating opportunity, which was provided 24 h after an
188 initial mating. Undoubtedly, remating rates in H and P females, each of which were
189 continuously housed with four additional males for several days, would had been much
190 higher. On day 5, males were discarded and females were transferred into single
191 containers filled with approximately 40-70 mung beans for egg laying. Females were
192 placed in containers with 40-70 new (i.e., uninfested) beans 0 hrs (day 5), 24 hrs (day 6)
193 and 72 hrs (day 8) after the end of the mating treatment. They were kept and checked
194 daily for survival in the last container until death. We set up a total of 41 F0 females in
195 the M, 39 in the H, and 34 in the P treatment. Female *C. maculatus* lay one egg per bean
196 when provided with sufficient resources (Messina, 1991), and the provision described

197 above ensured that there was no larval competition (i.e., no more than one egg per bean),
198 as bean provision matched the patterns of egg laying: fecundity is highest during the first
199 day, decreasing quickly during the following days (Credland and Wright, 1989). On
200 average (\pm SE), the females in our experiment produced 27 (\pm 0.66) adult offspring (53%)
201 within the first 24hrs of oviposition, 14 (\pm 0.51) adult offspring (27%) in the subsequent
202 48 hours, and only 10 (\pm 0.35; 20%) adult offspring in the remaining time of their lives.
203 The sum of adult offspring from all containers constitutes our measure of lifetime
204 reproductive success (LRS) for each female, calculated for each generation separately.
205 After allowing one week of larval development, we collected 12 single inoculated beans
206 from the first egg containers that had been provided ("day 5": eggs laid 0-24 hrs after end
207 of mating treatment) and placed them individually in Eppendorf tubes with holes for
208 airflow, where they were kept until virgin adult beetles emerged. Of these, four males and
209 four females randomly selected from each clutch were used as focal F1 individuals. The
210 remaining beans from the first egg container and the remaining containers were kept
211 until all offspring had hatched, and were frozen for later counting.

212 **F1: First offspring generation - sons and daughters**

213 Two days post emergence into adulthood, virgin daughters (up to four from each treated
214 female) ($N_{\text{daughters}} = 405$) were each paired with same-age single virgin males derived from
215 a standardized heterozygous line (cross between two near-isogenic lines that had been
216 generated after following a brother-sister mating protocol for 33 generations). We
217 utilized tester individuals with a standardized genetic background to minimize variance
218 in reproductive success that would be attributable to genetic variance among the tester
219 males. After 24hrs, F1 females were provided clean beans (as described above for F0: 0
220 hrs, 24 hrs and 72 hrs after separation from mate) for egg laying. We acknowledge that

221 F1 and F2 females were younger at time of first bean provisioning, which may contribute
222 to differences in offspring production between the generations. However, as we were
223 especially interested in the variation in LRS across treatments within and across
224 generations, rather than in the within-treatment changes over generations, this is not
225 considered a problem. Females were monitored for lifespan daily. Grand-offspring were
226 sourced from the first egg laying (0hrs) container as before, but 8 instead of 12 inoculated
227 beans were isolated from each female this time. Due to equipment failure beyond our
228 control, approximately 50% of containers in the second instalment (24-72 hrs since start
229 of egg laying) for our assessment of lifetime reproductive success in this generation did
230 not contain viable offspring. As we could not be certain that the containers in which
231 offspring had emerged were unaffected (overall, unusually low numbers were observed),
232 we excluded all containers from this instalment for the calculation of lifetime
233 reproductive success in F1. We hence used only numbers for adult offspring from eggs
234 that that were laid at 0-24 hrs and between 72 hrs until death for F1 LRS.
235 Up to four virgin sons per female were kept in individual Eppendorf tubes and monitored
236 for lifespan, and survival checked once per day ($N_{\text{sons}}=393$).

237 **F2: Second offspring generation - grandsons and granddaughters**

238 We mated two females from each daughter in the same manner as described for the
239 previous generation ($N_{\text{granddaughters}}$ that successfully produced offspring=675). Emerging
240 F3 offspring were frozen and counted. Lifespan was monitored as before by checking
241 survival once per day in these F2 females ($N=647$ instead of 675, due to some females
242 escaping at late-age) and in two additional virgin male offspring per family
243 ($N_{\text{grandsons}}=679$).

244 **Economics across three generations**

245 To calculate the across-generation female productivity for each treatment, in addition to
246 comparisons of LRS in each of the generations separately, we approximated an index of
247 LRS, based on average offspring numbers across individuals and their contribution to the
248 next respective generation (for a hypothetical calculation example please refer to Table
249 S1).

250 We counted total offspring numbers for LRS without distinguishing between the sexes,
251 but assumed that offspring were produced in equal sex ratios (Reece et al., 2005). We
252 only used 50% of the counted LRS numbers from each female in F0 and F1 for our
253 calculation, due to the fact that we here only assayed female reproductive success. For
254 each F0 female that successfully produced descendants through to F3, the index was
255 calculated as:

256

257 $\frac{1}{2}$ (F0 LRS) * $\frac{1}{2}$ (F1 average LRS) * F2 average LRS

258

259 **Statistical analyses**

260 All analyses were carried out in R (version 3.4.0, R Development Core Team, 2012). Mixed
261 model analyses on lifetime reproductive success and on lifespan were conducted using
262 *lme4* (Bates et al., 2015) and p-values extracted using *lmerTest*, (Kuznetsova et al., 2013.
263 See [http://cran.r-project.org/web/packages/lmerTest.](http://cran.r-project.org/web/packages/lmerTest/)), using mating treatment as a
264 fixed factor. In analyses of F1 data, F0 female ID was added as a random variable. In
265 analyses of F2 data, F1 ID nested within F0 ID was included as a random variable.
266 Normality of residuals was visually confirmed. To run survival analyses and compare
267 survival probabilities across the treatments, we used mixed Cox proportional hazard
268 models using the R package *coxme* (Therneau, 2015. See <http://cran.r->

269 project.org/web/packages/coxme.), with female IDs included as a random effect as
270 detailed above. To further investigate potential trade-offs between survival and
271 reproduction, LRS of the respective generation was added as a covariate into the model
272 (see supplemental Table S2). We used Tukey multiple comparisons of means (TukeyHSD)
273 to investigate differences between treatments in the lifetime reproductive success assays,
274 and the *glht* function in package *multcomp* (Version 1.4-7, Hothorn et al., 2017,
275 <https://cran.r-project.org/web/packages/multcomp>) for posthoc tests on lifespan and
276 survival. Visual displays of the results (barplots) are based on means. Additional analyses
277 investigating mother-offspring correlations in reproductive success (Table S3) and
278 lifespan are presented in the supplemental material (Table S4).

279 **Results**

280 **The maternal generation (F0)**

281 We found no effect of the mating treatment on female lifetime reproductive success
282 ($F_{2,111}=0.625$, $p= 0.5372$, Fig 1A). Furthermore, we detected no effects of mating
283 treatment on lifespan ($F_{2,111}=0.592$, $p= 0.555$, see also Table S2 for no evidence for
284 lifespan-LRS trade-offs) or survival probability ($\chi^2= 0.838$, $df= 2$, $p= 0.658$).
285 Furthermore, we find little evidence for cross-generation correlations of LRS (Table S3).

286 **Offspring (F1 and F2)**

287 **1) Lifetime reproductive success (LRS)**

288 **a) F1: Daughters**

289 The maternal mating treatment conferred strong effects on the lifetime reproductive
290 success of females in the F1 generation ($F_{2,95}= 101.53$, $p< 0.0001$). Daughters from singly

291 mated F0 females produced the largest number of offspring (mean \pm SE, 46.2 \pm 1.01),
292 followed by daughters of multiply mated F0 females (40.8 \pm 1.32), and finally daughters
293 of harassed F0 females produced the lowest number of offspring (22.4 \pm 0.98; all
294 treatments significantly different from each other, Tukey's HSD test: M-H: $p < 0.0001$, M-
295 P: $p = 0.0035$, H-P: $p < 0.0001$, see Fig 1B).

296 ***b) F2: Granddaughters***

297 Grand-maternal mating treatment also affected the lifetime reproductive success of the
298 granddaughters ($F_{2,87}=6.220$, $p < 0.003$), but the pattern was reversed compared to the
299 previous generation. Granddaughters from singly mated F0 females exhibited
300 significantly lower levels of reproductive output (52.7 \pm 1.22) than granddaughters from
301 harassed (60.9 \pm 1.79), but not multiply mated (55.9 \pm 1.65) F0 females (Tukey's HSD test:
302 M -H = -8.171, $p = 0.0004$; M -P = 3.19, $p = 0.3077$; H - P = -4.981, $p = 0.0693$, see Fig 1C).
303 Overall, F2 females appear to show a classical offspring number / lifespan trade-off
304 (Table S2), in contrast to females in the other generations.

305 ***2) Lifespan***

306 ***a) F1: Sons & Daughters***

307 The mean lifespan ($F_{2,104}=4.001$, $p = 0.0212$) and survival probabilities ($\chi^2 = 44.79$, $df=2$,
308 $p < 0.0001$) of F1 daughters differed significantly according to the maternal mating
309 treatment. Specifically, singly mated (M) F0 mothers produced shorter-lived daughters
310 than harassed (H) and multiply mated (P) mothers (see Fig 2 A,C). In sons, neither
311 lifespan ($F_{2,101}=1.80$, $p = 0.2112$, Fig 2B) nor survival ($\chi^2 = 3.61$, $df=2$, $p = 0.165$, Fig 2D)
312 differed with maternal mating treatment. We did not detect any significant mother-
313 offspring correlations in lifespan (see Table S4 A).

314 ***b) F2: Granddaughters & Grandsons***

315 Granddaughters from all three maternal mating treatments differed significantly from
316 each other, both in average lifespan ($F_{2,71}=16.352, p < 0.0001$, Fig 3A) and in survival
317 probability ($\chi^2= 34.17, df=2, < 0.0001$, Fig 3C). Specifically, descendants from the
318 harassment treatment (H) lived longest, M granddaughters were intermediate in lifespan
319 (6% shorter lifespan than H), and P granddaughters lived shortest (12% shorter lifespan
320 than H; Fig 3A,C; Tukey's HSD test: M -H: $z= -3.333, p = 0.0026$; M -P: $z= -2.952, p < 0.009$;
321 H - P: $z= -5.801, p < 0.001$). Interestingly, granddaughter's lifespan was significantly
322 correlated with their grandmothers' (F0) but not their mothers (F1) lifespan (see Table
323 S4 B).

324 The effect was similar for grandsons, with H descendants living the longest (14% longer
325 lifespan compared to M, 12% longer than P descendants; $F_{2,95}=25.984, p < 0.0001$; Tukey's
326 HSD test: M -H: $z= -6.881, p < 0.0001$; M -P: $z= 1.278, p = 0.408$; H - P: $z= -5.324, p <$
327 0.0001 , Fig 3b) and having the highest survival probability ($\chi^2= 108.86, df=2, < 0.0001$,
328 Tukey's HSD test: M - H: $z=-6.980, p < 0.0001$; M-P: $z= 1.316, p = 0.386$; P- H: $z= -5.374,$
329 $p < 0.0001$, Fig 3D).

330 ***Economics***

331 We find that maternal mating treatment had a highly significant effect on net fitness
332 across the three generations examined ($F_{2,106}=6.82, p = 0.0016$, Fig 4), with H females
333 generating less than two thirds of descendants compared to the other treatments
334 (posthoc Tukey comparisons: M -H: $p=0.0014$, M - P: 0.6336 , P -H: 0.0337).

335 **Discussion**

336 We demonstrate large transgenerational effects of maternal sexual interactions spanning
337 several generations. Both lifetime reproductive success and lifespan in offspring and
338 grand-offspring were influenced by the maternal mating treatment imposed on F0

339 females. Remarkably, we did not find sizeable costs or benefits in the F0 generation to the
340 mothers themselves. In *C. maculatus*, it has been previously shown that multiply-mated
341 females live shorter than singly mated females (Crudgington and Siva-Jothy, 2000; but
342 see Fox, 1993a; Arnqvist et al., 2004), which is thought to be caused, at least in part, by
343 the sharp male genital spines that puncture the connective tissue within the female
344 reproductive tract during mating (Dougherty et al., 2017; Dougherty and Simmons, 2017;
345 Rönn et al., 2007). Harassment of females by emasculated males (to remove potential
346 effects associated with mating itself) has also been previously reported to lower
347 reproductive success and longevity in this species (den Hollander and Gwynne, 2009).
348 High mating rates have, however, been shown to have beneficial effects on offspring
349 production in this species (Arnqvist et al., 2004), potentially due to effects of large
350 ejaculates on female hydration or nutritional status (Fox, 1993a). Interestingly we find no
351 costs of repeated mating or harassment on lifespan, and no effects of mating regimes on
352 offspring production, in the maternal generation. However, our experiment reveals
353 substantial effects of maternal mating treatment on subsequent generations.

354

355 The evaluation of fitness consequences beyond the parental generation is necessary if we
356 want a comprehensive picture of the consequences of sexual interactions, and to
357 understand the role of all effects, including the effects of the social environment, on the
358 evolution of fitness-related traits. The importance of non-genetic inheritance has been
359 highlighted in this special issue, and here we show that effects attributable to variation in
360 levels of sexual interactions experienced by females in one generation can permeate
361 across several generations, influencing reproductive success and survival patterns of
362 future generations. While maternal effects arising from variations in social conditions
363 have been studied extensively especially within the framework of maternal care

364 (Champagne, 2008), including their large role influencing offspring gene expression, only
365 a few studies to date have investigated the role of transgenerational maternal sexual
366 interactions. Recent studies in *D. melanogaster* found that daughters produce more
367 offspring when their mothers had experienced higher levels of maternal sexual
368 interactions (Garcia-Gonzalez and Dowling, 2015; Priest et al., 2008a), whereas longevity
369 and survival of offspring are negatively affected when produced by mothers that had
370 experienced heightened sexual interactions (Dowling et al., 2014). Such
371 transgenerational effects have also been reported in guppies (*Poecilia reticulata*),
372 whereby increased male presence and harassment led to lower reproductive success in
373 offspring (Gasparini et al., 2012). Our results reflect this pattern found in guppies,
374 because daughters from the harassment treatment had significantly lower levels of
375 lifetime reproductive success than daughters from other treatment groups. However, this
376 pattern was reversed in granddaughters, where descendants from harassed mothers had
377 the highest lifetime reproductive success. Such a reversal of offspring fitness across
378 different generations has also been found in a study in *D. melanogaster*, where sons were
379 found to have increased fitness, but grandsons decreased fitness with increasing
380 maternal sexual interactions (Brommer et al., 2012). In combination, these results
381 highlight the importance of investigating the magnitude and direction of effects across
382 multiple generations to understand the net transgenerational consequences of sexual
383 interactions. Even in the absence of immediate costs and benefits due to sexual
384 interactions potential reversing effects that occur in subsequent generations need to be
385 taken into account to unveil the long-term consequences for the evolution of different
386 mating strategies and mating systems.

387

388 While not as closely reflective of Darwinian fitness as the production of adult offspring
389 over a lifetime, the transgenerational effects we observed on survival are also
390 noteworthy. Daughters from singly mated F0 females produced the largest numbers of
391 offspring, and were found to have the shortest lifespan, reflecting a classic life-history
392 trade-off. Similarly, this trade-off was evident in H daughters, in the other direction, as
393 this group displayed significantly longer survival but lowest reproductive outputs.
394 Granddaughters from harassed F0 females (which had the highest levels of reproductive
395 success), however, lived longest. The survival patterns were similar between the sexes,
396 although male lifespan in the F1 sons was not influenced by their mothers mating
397 treatment, in contrast to daughters. Grandsons from harassed F0 females lived
398 significantly longer than those from singly or multiply mated F0 females, reflecting the
399 pattern observed in granddaughters. This pattern of overall survival benefits to
400 descendants from the harassment treatment is surprising, but highlights the importance
401 of investigating transgenerational effects across a variety of life-history traits, since they
402 may reveal unexpected patterns of trait correlation both within and across generations.
403 We did not find evidence for benefits of mating with multiple males, despite the fact that
404 it has been shown in this species that the receipt of multiple ejaculates can confer direct
405 benefits, in the form of additional hydration (Edvardsson, 2007). It has been suggested
406 that benefits of multiple mating may only be evident under nutrient-limited conditions
407 (Fox, 1993a), which may indicate that our environment was too benign to reveal
408 differences. Lower levels of ambient humidity may be necessary to induce observable
409 effects. However, it is also possible that the negative effects of harassment or harm in the
410 multiply mated group masked the potential benefits that could be conferred directly to
411 females. Negative effects of increasing copulations on female fecundity have been
412 reported previously, while multiple mating conferred the overall benefit of increased

413 fertilisation assurance (Wilson and Tomkins, 2015), and has been associated with
414 increasing egg size (Fox, 1993b).

415

416 The transgenerational effects in the grand-offspring generation alone indicate positive
417 effects of grand-maternal harassment, both in lifespan and reproductive success. Taking
418 overall fitness gains via female reproduction into account, however, our estimation of
419 population growth via multiplicative fitness indicates that harassment overall bears very
420 large negative fitness consequences, at least when measured across three generations.
421 This indicates that transgenerational effects and non-genetic inheritance of sexual
422 interactions can have important impacts on the evolution of sexual interactions. While
423 the interpretation of opposing patterns in different generations is complex, the strong
424 effects observed in F1 indicate that indirect genetic effects, such as maternal effects, play
425 an important role. Specifically, harassment may induce low levels of fecundity in the
426 offspring generation, for example via elevated stress levels in the F0 females. However,
427 the increase in offspring numbers in the F2 generation may be due to an increase in
428 offspring investment in the F1 mothers following very low levels of investment of the
429 preceding generation. While we did not examine egg size, a possibility is that daughters
430 from harassed mothers invested into fewer but larger eggs, which in turn could have
431 developed into high performing offspring, both in terms of fecundity and longevity.
432 Opposing patterns observed in the different generations may be caused by differences in
433 how females allocate resources toward individual egg size relative to the number of eggs
434 produced. Life history theory predicts that the number and size of offspring should trade-
435 off (Smith and Fretwell, 1974; Stearns, 1989), and how females resolve this trade-off may
436 vary depending on factors like female condition (Wilson et al., 2009), the phenotype of
437 her mate (Kindsvater and Alonzo, 2014; Qvarnström et al., 2000) or the conditions into

438 which her offspring will be born (Fox and Czesak, 2000; Parker and Begon, 1986). The
439 transgenerational fluctuations in LRS seen here for instance could result if harassed F0
440 females invest fewer resources in their eggs leading to the production of low condition
441 daughters (F1), who then themselves go on to produce just a few high quality daughters
442 (F2), who themselves go on to produce high numbers of offspring. Such plastic maternal
443 effects are often linked to limited resources available to devote to offspring production,
444 and are commonly observed across the animal kingdom (Brommer et al., 2012; Brown
445 and Shine, 2009; Lasne et al., 2017; Savalli and Fox, 2002), including this study species
446 (Fox, 1993b; Fox et al., 1999, 1997). In particular, the role of fluctuating maternal effects
447 in mediating the transgenerational fitness consequences of sexual interactions and sexual
448 conflict is an avenue of research that deserves greater attention.

449 **Conclusion**

450 Our results indicate that harassment, when experienced without the apparent benefits of
451 receiving ejaculates from multiple males, leads to suboptimal fitness consequences for
452 daughters. However, different outcomes across generations indicate that it is necessary
453 to include multiple generations when the net consequences of sexual interactions are
454 being investigated. Sex-specific effects of maternal mating history on lifespan indicate the
455 importance of investigating fitness traits in males and females separately. Our finding
456 that costs and benefits may alternate between generations may indicate that socially
457 mediated context-dependent effects may be important drivers of the evolutionary
458 dynamics of sexual interactions.

459

460 **Data archiving**

461 All data presented in this work can be accessed on Dryad. Accession number tbd.

462

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470

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665

666 **Figure legends**

667

668 Figure 1: Lifetime reproductive success in females at the F0, F1 and F2 generation
669 respectively. Light grey: single mating in maternal generation (Mono, M), grey: single
670 mating + harassment (H), dark grey: multiple mating with multiple males (Poly, P). A:
671 maternal generation, B: daughters, C: granddaughters

672

673 Figure 2: Average lifespan and survival curves for male (virgin) and female (mated for
674 24 hrs) offspring (F1). Maternal treatment: Light grey: single mating (Mono, M), grey:
675 single mating + harassment (H), dark grey: multiple mating with multiple males, Poly,
676 P). A: daughters average lifespan, B: sons average lifespan, C: survival curves for
677 daughters (C) and sons (D).

678

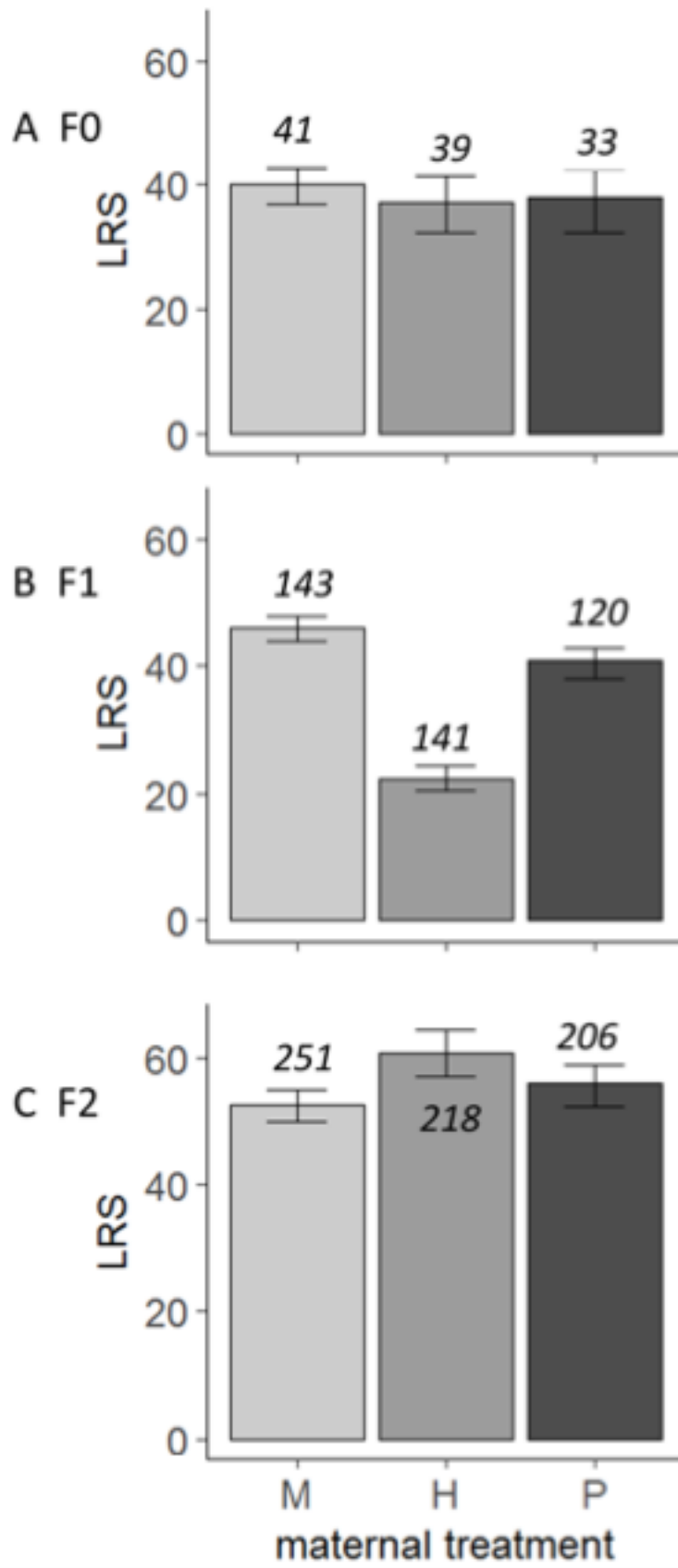
679 Figure 3: Average lifespan and survival curves for male (virgin) and female (mated for
680 24 hrs) grand-offspring (F2). Maternal treatment: Light grey: single mating (Mono, M),
681 grey: single mating + harassment (H), dark grey: multiple mating with multiple males,
682 Poly, P). A: granddaughters average lifespan, B: grandsons average lifespan, C: survival
683 curves for granddaughters (C) and grandsons (D).

684

685 Figure 4: Estimation for multiplicative fitness for females within the different mating
686 treatments after 3 generations. Light grey: single mating (Mono, M), grey: single mating
687 + harassment (H), dark grey: multiple mating with multiple males, Poly, P).

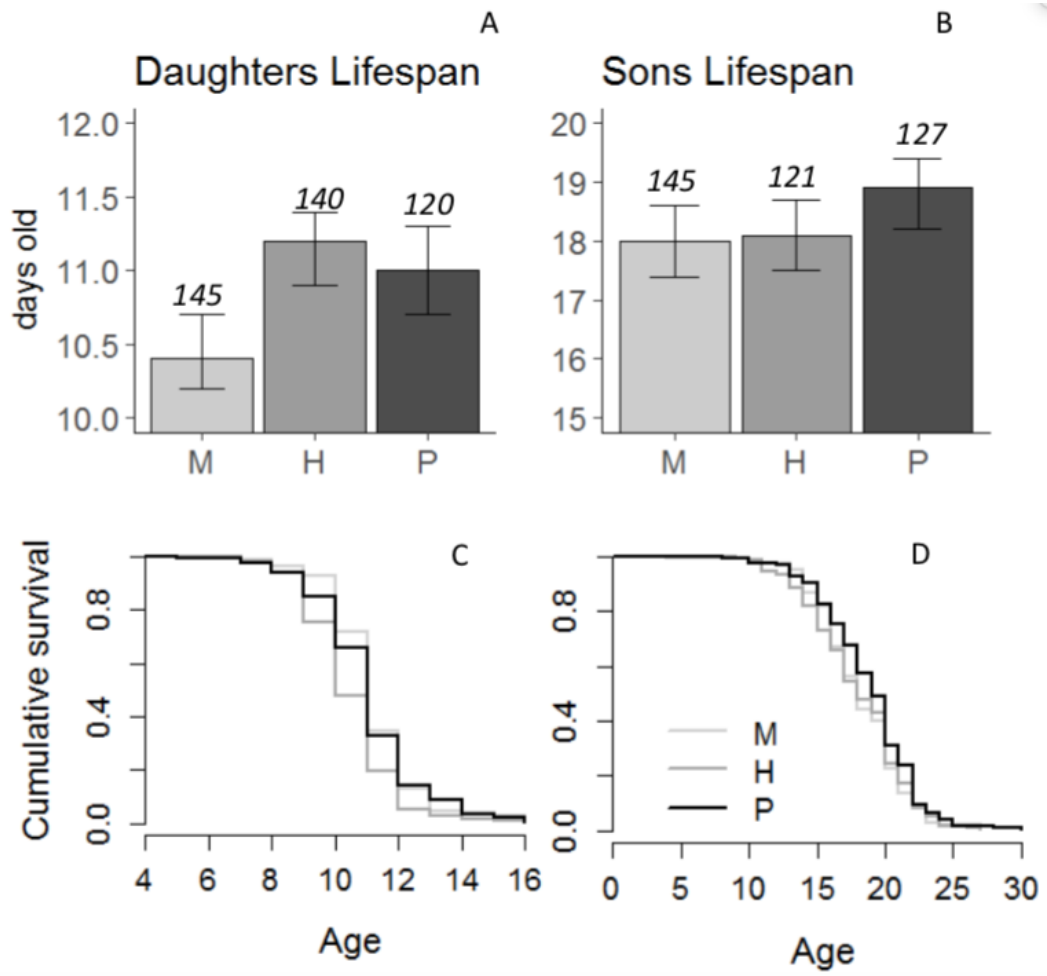
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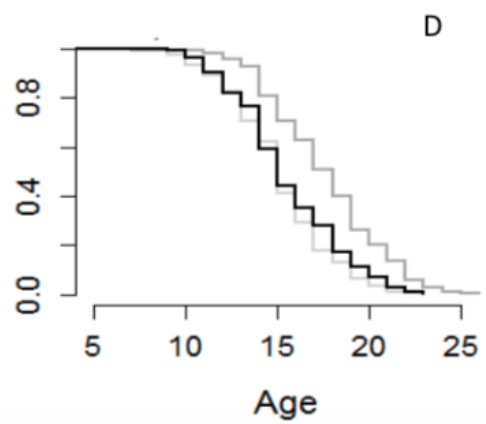
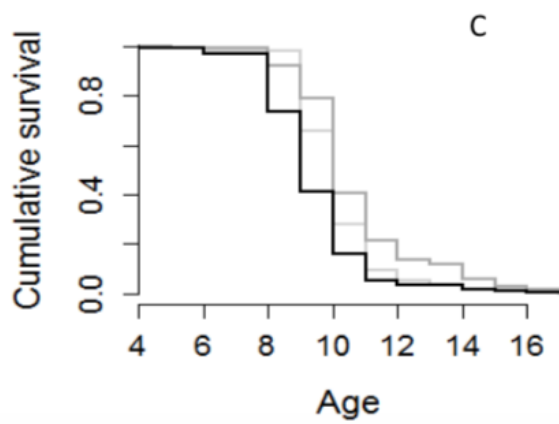
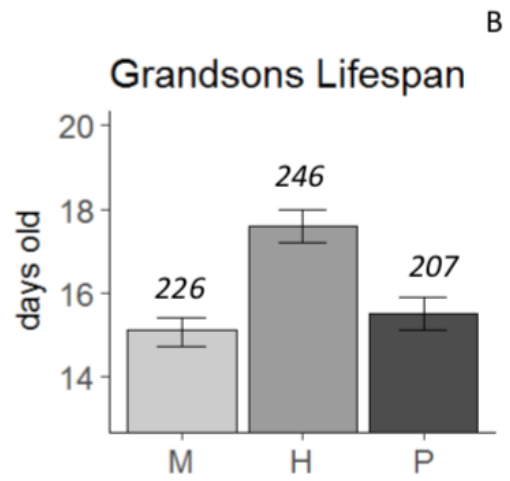
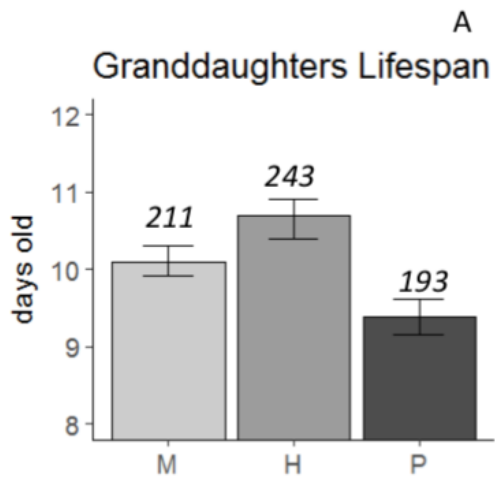
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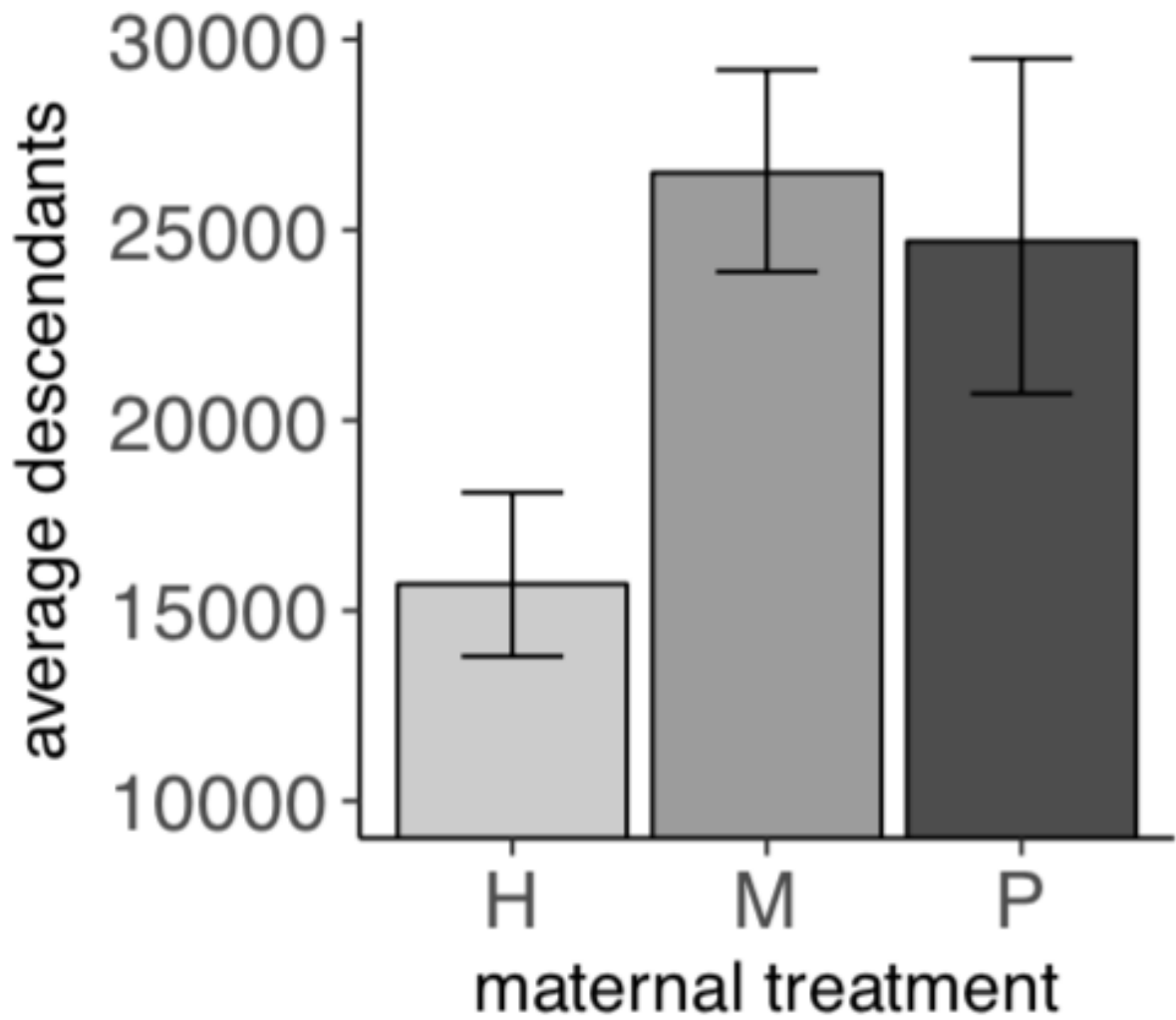


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Multiplicative Fitness



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