

1	Title: Using experimental evolution to study adaptations for life within the
2	family
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15	Nicrophorus vespilloides, interacting phenotypes
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Abstract

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20 Parents of many species provision their young and the extent of parental provisioning 21 constitutes a major component of the offspring's social environment. Thus a change 22 in parental provisioning, whether evolved or plastic, can alter the form or strength of 23 selection on offspring and result in the coevolution of parental provisioning and traits 24 expressed in offspring. Although this reasoning is central to our evolutionary 25 understanding of family life, there is little direct evidence that selection by parents 26 causes evolutionary change in their young. Here we use experimental evolution to 27 examine directly how populations of burying beetles, Nicrophorus vespilloides, adapt 28 to a change in post-hatching parental provisioning. We measured the performance of 29 larvae descended from independent lab populations that had been propagated for 30 several generations with and without post-hatching parental care (Full Care and No 31 Care populations respectively). We found that adaptation to the absence of post-32 hatching care led to rapid and consistent changes in larval survival in the absence of care. Specifically, larvae descended from No Care populations had higher survival in 33 34 the absence of care than larvae descended from Full Care populations. Other 35 measures of larval performance, such as the ability of larvae to consume a breeding 36 carcass and larval mass at dispersal did not differ consistently between the Full Care 37 and No Care populations. Nevertheless, our results show that populations can adapt 38 rapidly to a change in the extent of parental care and that experimental evolution can 39 be used to study such adaptation.

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Introduction

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45 All animals provision their young in some form. This provisioning may come 46 as yolk deposited in eggs, nutrients transferred between mothers and embryos via a 47 placenta, or as resources provided to dependent young after birth or hatching. 48 Understanding why animals vary in the mode by which they provision their young, 49 and the amount of resources they provide, has been a major focus of evolutionary and 50 behavioral ecology (Clutton-Brock 1991; Roff 1992, 2002). Much of this work has 51 considered parental provisioning as an adult life-history trait that is likely to 52 experience different strengths or forms of selection in different environments (Smith 53 and Fretwell 1974; Roff 1992, 2002). If levels of parental provisioning are heritable, 54 then selection for different levels of parental provisioning in different populations can 55 result in divergence in traits such as egg size or offspring size at birth (e.g. 56 Schwarzkopf et al. 1999; Czesak and Fox 2003; Heath et al. 2003; Fischer et al. 57 2006). Indeed, several studies have found associations between environmental variables and egg size or size at birth suggesting that parental provisioning is often 58 59 locally adapted (e.g. Reznick and Endler 1982; Johnston and Leggett 2002; Heath et 60 al. 2003: Schrader and Travis 2012). 61 Parental provisioning is not only an important adult life history trait. In 62 animals with parental care it also constitutes an important part of the offspring's 63 social environment (Wolf et al. 1998). Therefore a change in parental provisioning, 64 may alter the form or strength of selection on offspring phenotype resulting in the

65 further evolution of traits expressed in offspring (Kirkpatrick and Lande 1989; Wolf

66 et al. 1998; Lock et al. 2004; Moore et al. 1997). That offspring will adapt to

67 variation in parental provisioning is a central assumption of models of both parent-

68 offspring conflict and coadaptation. For example, parent-offspring conflict is 69 predicted to favor adaptations in offspring that allow them to extract more care from 70 parents than is optimal for parents to provide and models of coadaptation predict that 71 selection on offspring will result in the joint evolution of traits expressed in parents 72 (e.g. supply of resources) and offspring (e.g. demand for resources) that interact to 73 influence offspring fitness (Trivers 1974; Wolf and Brodie 1998; Wolf 2000; Hinde et 74 al. 2010). Support for this assumption comes mainly from quantitative genetic 75 estimates of social epistasis or genetic correlations between traits involved in parental 76 supply of resources and offspring demand for these resources (reviewed in Kölliker et 77 al 2012). The presence of such correlations suggests that selection has favored 78 specific combinations of these traits (i.e. there has been correlational selection on 79 parent-offspring interactions). In most cases however, it is not known how these 80 genetic correlations became established and explanations for the forces that generated 81 them are inferred from the direction of the correlation (Agrawal et al. 2001; Kölliker 82 et al. 2005; Lock et al. 2004; reviewed in Kölliker et al. 2012). For example, Agrawal 83 et al. (2001) found a negative genetic correlation between parental provisioning and 84 offspring begging in Burrower bugs (Sehirus cinctus) and the direction of this 85 correlation is consistent with predictions of parent-offspring coadaptation theory that 86 assume the presence of stabilizing selection on offspring phenotype (Wolf and Brodie 87 1998; Agrawal et al. 2001). In short, previous work suggests that traits involved in 88 parent-offspring interactions should coevolve and some studies have uncovered 89 genetic correlations between traits expressed in parents and offspring consistent with 90 such coevolution. However, no studies have measured whether a change in parental 91 behavior causes an evolved change in traits expressed by the offspring.

92 Here we describe an experiment in which we examine directly how 93 populations of burying beetles, *Nicrophorus vespilloides* adapt to a change in parental 94 provisioning. Like all species in the genus, N. vespilloides breeds on the carcasses of 95 small vertebrates. Upon encountering a carcass, parents mate and prepare the carcass 96 for their young to feed upon. Carcass preparation involves shaving the fur or feathers 97 from the carcass, rolling it into a ball, and smearing the surface of the carcass with 98 anal exudates that delay decomposition (Scott 1998). The eggs, which are laid near 99 the carcass, hatch and the larvae migrate to the carcass where they feed. N. 100 vespilloides larvae exhibit begging behaviors and parents respond to these behaviors 101 by regurgitating predigested carrion that larvae consume. Post-hatching parental care 102 is facultative in *N. vespilloides*. Although larvae are able to complete development 103 with no post-hatching care, measures of breeding success and larval performance are 104 typically higher when parents are allowed to provision larvae than when they are not 105 (Eggert et al. 1998). 106 We took advantage of the facultative nature of post-hatching care in this 107 species to ask how populations adapt to the complete removal of post-hatching care. 108 To do this we used experimental evolution. This approach involves establishing 109 replicate experimental populations, exposing these populations to different

110 environments for many generations, and then comparing traits between populations

111 that have evolved in these different environments. Experimental evolution has been

used to study how populations adapt to changes in environmental variables such as

113 predation risk (e.g. Reznick et al. 1997) as well as social aspects of the environment

such as the mating system (e.g. Hollis and Kawecki 2014) but has not yet been used to

study the evolution of parent-offspring interactions. We allowed *N. vespilloides*

116 populations to evolve for several generations with and without post-hatching parental

117	care (Full Care and No Care populations respectively). Then, for each experimental
118	population, we measured offspring performance with and without post-hatching
119	parental care. We predicted that adaptation to the absence of post-hatching parental
120	care would reduce larval dependence upon care such that, in the absence of parental
121	care, larvae from No Care populations would perform better than larvae from Full
122	Care populations.
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124	Methods
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126	Establishment and maintenance of experimental populations
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128	The experimental populations used in this study were created as part of a
129	larger experiment designed to test whether post-hatching parental care influences the
130	response to selection on body size (Jarrett et al. in prep). For the purposes of this
131	study we ignore selection on body size and focus on four experimental populations
132	that differed in the presence of post-hatching parental care. A complete description of
133	the protocols used to establish and maintain these populations as well as data on
134	differences in body size between the populations is provided in the supplemental
135	material.
136	The experimental populations were descended from beetles collected in 2012
137	from two wild populations (Byron's Pool and Wicken Fen) in Cambridgeshire, UK.
138	These populations were interbred over the course of four generations, with 40 pairs
139	breeding each generation. We then interbred this population with a laboratory
140	population for a single generation (breeding 160 pairs) to create a large, genetically
141	diverse stock population. From this stock population, we created two populations that

142 were maintained without post-hatching parental care (No Care 1 and 2) and two

143 populations that were maintained with full parental care (Full Care 1 and 2).

144 Each No Care (NC) population was initiated by breeding 60 pairs of beetles from the stock population. We placed each pair in a box with commercially bought 145 146 compost and a thawed mouse carcass (8-14 g) and then placed these boxes in a dark 147 cupboard to simulate underground conditions. We removed both parents from the 148 breeding box 53 hours after pairing. This is enough time for females to complete egg 149 laying and carcass preparation but is before eggs begin to hatch (Boncoraglio and 150 Kilner 2012, Smiseth et al. 2006). After removing both parents, we returned the box 151 to the cupboard where it remained until larval dispersal (8 days after pairing). On the 152 eighth day we counted and removed all of the larvae from each breeding box, 153 weighed each brood, and then calculated the average mass of larvae in each brood 154 (total brood mass / brood size). We measured brood size and brood mass 8 days after 155 pairing because by this time larvae have nearly always left the carcass (even in cases 156 where some flesh remains) and are wandering in the soil. Furthermore, in our 157 population, leaving the larvae for longer increases that probability that one of the 158 parents will eat some of the wandering larvae.

After measuring brood size and larval mass, we placed the larvae in 25-celled
eclosion boxes (box dimensions, length x width x depth: 10 cm x 10 cm x 1.8 cm)

161 with one larva in each cell (individual cell dimensions: 2 cm x 2 cm x 1.8 cm),

162 covered them with damp peat, and left them to pupate for 17 days. Most individuals

had eclosed by 17 days. However, those that had not were allowed more time to

164 pupate (usually an additional day). After eclosion we sexed and photographed each

165 beetle and then placed individual beetles in plastic boxes (box dimensions, length x

166 width x depth: 12 cm x 8 cm x 2 cm), with damp compost and a small amount of

ground beef. Adult beetles remained in these boxes for two weeks and were fed
ground beef twice per week. Two weeks after eclosion, we bred beetles from each
population as described above. The number of pairs bred in each generation varied.
In No Care 1 (NC1) we bred between 64 and 120 pairs per generation (mean =
110.75) and in No Care 2 we bred between 39 and 120 pairs per generation (mean =
82.7)
The Full Care (FC) populations were each initiated by breeding 40 pairs of

beetles and were treated in exactly the same way as the NC populations except that
we allowed parents to remain with their larvae until larval dispersal. In the Full Care
population (FC1) we bred 80 pairs per generation and in the Full Care 2 (FC2) we
bred between 6 and 80 pairs per generation (mean = 75.7).

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Block One

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181 We maintained NC1 and FC1 for four generations as described above. We 182 then passed both populations through a fifth generation in which larvae were reared 183 with full parental care, following the protocol for the Full Care populations (Figure 1). 184 This was done to minimize differences between lines in transgenerational effects of 185 care. The resulting offspring were used as the parents in the sixth generation. 186 In the sixth generation, we bred beetles from the NC1 and FC1 populations 187 with and without post-hatching parental care (see Figure 1). To do this we randomly 188 paired unrelated adult males and females from within each experimental population (n 189 = 47 pairs from the NC1 population; n = 69 pairs from the FC1 population). In the 190 NC1 population we bred 22 pairs without post-hatching parental care and 25 pairs 191 with post-hatching parental care. In the FC1 population, we bred 41 pairs without

192	post-hatching parental care and 28 pairs with post-hatching parental care. Eight days
193	after pairs were bred, we recorded several measures of larval performance. First we
194	recorded two proxies of larval survival: whether the breeding attempt produced at
195	least one dispersing larva (breeding success, scored as a binary variable), and for pairs
196	that bred successfully, the number of larvae at dispersal (brood size). We also
197	recorded two measures of larval performance: whether the larvae had consumed the
198	entire breeding carcass (carcass consumption) and the average mass of dispersing
199	larvae (larval mass). Carcass consumption was scored as a binary variable. If there
200	was no flesh remaining on the carcass at the time of larval dispersal we considered the
201	carcass to be completely consumed, otherwise, we scored the carcass as unconsumed
202	(as in Rauter and Moore 2002). The average mass of dispersing larvae was calculated
203	as the total brood mass at dispersal divided by the number of larvae in the brood.
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204 205	Block Two
204 205 206	Block Two
204 205 206 207	<i>Block Two</i> We replicated the experiment described above using NC2 and FC2
204 205 206 207 208	<i>Block Two</i> We replicated the experiment described above using NC2 and FC2 populations. These populations had been maintained for seven generations and were
204 205 206 207 208 209	<i>Block Two</i> We replicated the experiment described above using NC2 and FC2 populations. These populations had been maintained for seven generations and were each passed through an eighth generation in which larvae were reared with full
204 205 206 207 208 209 210	<i>Block Two</i> We replicated the experiment described above using NC2 and FC2 populations. These populations had been maintained for seven generations and were each passed through an eighth generation in which larvae were reared with full parental care (Figure 1). The resulting offspring were used as the parents in the ninth
204 205 206 207 208 209 210 211	<i>Block Two</i> We replicated the experiment described above using NC2 and FC2 populations. These populations had been maintained for seven generations and were each passed through an eighth generation in which larvae were reared with full parental care (Figure 1). The resulting offspring were used as the parents in the ninth generation. In the ninth generation, we bred beetles from the NC2 and FC2
204 205 206 207 208 209 210 211 211	<i>Block Two</i> We replicated the experiment described above using NC2 and FC2 populations. These populations had been maintained for seven generations and were each passed through an eighth generation in which larvae were reared with full parental care (Figure 1). The resulting offspring were used as the parents in the ninth generation. In the ninth generation, we bred beetles from the NC2 and FC2 populations with and without post-hatching parental care (Figure 1). Our methods
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204 205 206 207 208 209 210 211 212 213 214	<i>Block Two</i> We replicated the experiment described above using NC2 and FC2 populations. These populations had been maintained for seven generations and were each passed through an eighth generation in which larvae were reared with full parental care (Figure 1). The resulting offspring were used as the parents in the ninth generation. In the ninth generation, we bred beetles from the NC2 and FC2 populations with and without post-hatching parental care (Figure 1). Our methods were identical to those used in block one although sample sizes were different. In the NC2 population we bred 50 pairs without post-hatching parental care and 30 pairs
204 205 206 207 208 209 210 211 212 213 214 215	<i>Block Two</i> We replicated the experiment described above using NC2 and FC2 populations. These populations had been maintained for seven generations and were each passed through an eighth generation in which larvae were reared with full parental care (Figure 1). The resulting offspring were used as the parents in the ninth generation. In the ninth generation, we bred beetles from the NC2 and FC2 populations with and without post-hatching parental care (Figure 1). Our methods were identical to those used in block one although sample sizes were different. In the NC2 population we bred 50 pairs without post-hatching parental care and 30 pairs with post-hatching parental care. In the FC2 population, we bred 50 pairs without

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Analysis

219 220 Our experimental design was a 2 x 2 factorial with two population types (No 221 Care and Full Care) and two levels of environment (No Care and Full Care). The 222 entire experiment was replicated twice using independent No Care populations that 223 had been evolving without parental care for different amounts of time (4 generations 224 in block one and 7 generations in block two). To account for this, we included block 225 as an additional factor in all analyses. When interactions involving block were 226 significant we analyzed the data from each block separately. 227 We examined the effect of population, environment, block, and their

228 interactions on breeding success using two complementary analyses. We first scored 229 each breeding attempt as being either a success or a failure and analyzed the data 230 using a GLM with a binomial error term. For this analysis, we scored pairs that laid 231 eggs but did not produce any dispersing larvae as failures and pairs that had at least 232 one dispersing larva as successes. Second, for pairs that bred successfully, we 233 examined the effect of population, environment, block, and their interactions on the 234 number of dispersing larvae (brood size) using a GLM with a Gaussian error term. 235 Although brood size only takes on integer values, we chose to use a Gaussian error 236 term instead of a Poisson error term because the overall mean brood size was fairly 237 high (14.19) and the residuals from a Gaussian model and a quasipoisson model (the 238 Poisson model was overdispersed) behaved similarly. We note however that using a 239 quasipoisson error term did not qualitatively affect our results. Carcass mass did not 240 influence brood size in our experiments so we did not include it as a covariate in this

analysis [linear regression of carcass mass on brood size in both blocks pooled; slope

242 (±SE) = 0.11 (0.46),
$$R^2$$
 = 0.0004, $F_{1,181}$ = 0.066, P = 0.80).

We restricted our analysis of carcass consumption to the No Care environment since parents also feed upon the carcass making it impossible to attribute carcass consumption to offspring when parents are also present (we note however that carcass consumption was > 90 % in all treatments where parents were allowed to remain with their larvae). We examined the effect of population (Full Care or No care), brood size, block, and their interactions on carcass consumption using a GLM with a binomial error term.

250 We intended to examine the effects of population, environment, and block on 251 larval mass using a GLM with brood size as a covariate. We originally included the 252 interaction between brood size and environment (Care versus No Care) in this 253 analysis. This interaction was highly significant (P < 0.01) in both blocks of the 254 experiment indicating that the homogeneity of slopes assumption of the model was 255 violated. Because the relationship between brood size and mean larval mass differs 256 between the two parental care environments it is inappropriate to proceed with testing 257 the significance of the main effects assuming a common slope. Below we report the 258 results of the analysis of larval mass excluding brood size as a covariate.

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Results

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Breeding success, measured as the proportion of pairs producing at least one dispersing larva, was lower without post-hatching parental care than with posthatching parental care. However, the magnitude of this difference varied between the No Care and Full Care populations (Figure 2, Table 1). In the absence of post-

266 hatching care the No Care populations had higher breeding success than the Full Care 267 populations (50 % versus 34 % in block one, 70 % versus 52 % in block two). In the 268 presence of post-hatching care, breeding success was lower when the beetles were 269 from the No Care populations than when they were from the Full Care populations 270 (80 % versus 93 % in block one, 88 % versus 97 % in block two). 271 In both blocks of the experiment, post-hatching parental care led to an 272 approximately 60 % increase in mean brood size at dispersal (pooling lines within 273 each environment). The effect of parental care on average brood size did not differ 274 between the No Care and Full Care populations (Figure 3, Table 2). 275 In both blocks of the experiment, the proportion of broods that successfully 276 consumed the carcass appeared higher when the larvae were from the No Care 277 populations than when they were from the Full Care populations (Figure 4). In block 278 one carcass consumption was 4.6 times greater when beetles were descended from the 279 No Care line than when they were from the Full Care line and in block two carcass 280 consumption was 1.7 times greater when the beetles were descended from the No 281 Care line than when they were from the Full Care line (Figure 4). However, after 282 accounting for the effect of brood size on carcass consumption, there was no evidence 283 for a difference between populations in carcass consumption (Table 3). 284 The analysis of larval mass was complicated by differences between the Full 285 Care and No Care environments in the relationship between larval mass and brood 286 size. In the Full Care environment there was a negative relationship between average 287 offspring size and brood size while in the No Care environment this relationship was 288 shallower and hump-shaped (Figure S2). We have observed the same relationship in

289 other experiments and discuss its potential causes elsewhere (Schrader et al. in

290 review). However, because the relationship between brood size and mean larval mass

differs between the Full Care and No Care environments we report the results of theanalysis of larval mass excluding brood size as a covariate.

293 Initial analyses of the larval mass revealed a significant treatment by block 294 interaction so we analyzed each block separately. In block one mean larval mass was 295 significantly influenced by carcass mass [linear regression of mean larval mass on carcass mass: slope (\pm SE) = 0.008615 (0.00319), P = 0.0086, R² = 0.086, n = 71] so 296 297 we included carcass mass as a covariate in the analysis. After controlling for 298 variation in carcass mass, we a found a significant effect of environment and 299 marginally significant effects of population and the population by environment 300 interaction on mean larval mass (Table 4). The presence of parental care increased 301 larval mass, however this effect was more pronounced when beetles were from the 302 Full Care population than the No Care population (Table 4, Figure 5). 303 In block two, the relationship between carcass mass and mean larval mass was 304 marginally non-significant [linear regression of mean larval mass on carcass mass; slope (\pm SE) = 0.003386 (0.00182) P = 0.066, R^2 = 0.022, n = 110]. Our results are 305 306 qualitatively similar whether or not we include carcass mass as a covariate and for 307 simplicity we present the results excluding carcass mass. In contrast to block one, 308 there was no evidence that parental care increased mean larval mass, nor was there 309 evidence that the Full Care and No Care populations differed in larval mass or the 310 effects of parental care on larval mass (Table 4, Figure 5). 311

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Discussion

Changes in parental provisioning are predicted to drive evolutionary changesin offspring, however few studies have directly examined how offspring adapt to a

316 change in a parental effect. In this experiment we used experimental evolution to 317 investigate how populations of *N. vespilloides* evolve in response to the elimination of 318 post-hatching parental care. We found that breeding success in the absence of post-319 hatching care was higher when the beetles had evolved in the No Care selection 320 regime than when they evolved in the Full Care selection regime. This observation is 321 consistent with rapid adaptation to the absence of post-hatching parental care. 322 However, other components of larval performance related to resource use did not 323 differ consistently between populations evolving with and without care.

324 Our results suggest that breeding success in the No Care populations has 325 become less dependent upon post-hatching parental care, and can envisage two routes 326 through which this may have evolved. The first possibility involves the evolution of a 327 trait expressed in the larvae. Eggert et al. (1998) found that the presence of parents 328 during the first 12 hours after larval hatching greatly increased larval survival and 329 growth and suggested that this effect was due to parents chewing a hole in the carcass, 330 thereby making it more accessible for the larvae. In our experiment, it may be that 331 larvae from the No Care lines are better able to chew an opening in the breeding 332 carcass themselves, whereas larvae from the Full Care lines are still dependent on 333 their parents for this task. This change in the larvae could occur through either a 334 behavioral (e.g. increased self feeding) or morphological (e.g. jaw structure) 335 adaptation. We are currently examining these possibilities. 336 The second route involves evolutionary change in traits expressed in the 337 parents. It is well-known from earlier studies that parental investment in burying 338 beetles involves activities before and after hatching (Scott 1998). Furthermore, there 339 is individual variation in the relative magnitude of the pre- and post-hatching 340 components of parental care (Lock et al. 2004; Lock et al. 2007; Steiger 2013). By

341 eliminating post-hatching parental care in one of our experimental evolution 342 treatments, it is possible that we selected for increased pre-hatching care. In other 343 words, larvae from the No Care populations may be less dependent on post-hatching 344 care because their parents have invested more heavily in pre-hatching care. 345 There are two general ways this might happen. First, parents from the No 346 Care populations may have invested more in carcass preparation than parents from the 347 Full Care populations, possibly making it easier for larvae to penetrate and use the 348 breeding carcass. Around the time of hatching, N. vespilloides parents create a 349 feeding cavity in the carcass that larvae recruit to after hatching. Adaptation to the No 350 Care selection regime could theoretically have selected parents that create this 351 depression earlier than usual (prior to parental removal at 53 hours after pairing). 352 However, we think such a shift is unlikely to explain our results because we have 353 never seen a feeding cavity (or the beginning of one) as early as 53 hours after 354 pairing. Another possibility is that selection in the No Care environment has favored 355 parents that create small holes in the carcass (as opposed to the rather large feeding 356 cavity) prior to parental removal. This could happen as a byproduct of parents 357 feeding on the carcass and neglecting to reseal the holes they create. We did not 358 inspect the carcasses in this experiment for small holes. Nevertheless, some support 359 for this idea comes from another experiment (Schrader et al. in revision). In that 360 experiment, we removed parents at 53 hours post-pairing and manipulated brood sizes 361 and inspected carcasses at larval hatching (70 hours post-pairing). At 70 hours post-362 pairing, we found that 13.75 % of prepared carcasses (11/80) did indeed have small 363 holes visible. Although we cannot say with certainty that these holes were created by 364 the parents, it is likely that at least some of them were. Nevertheless, this same 365 experiment also found that larvae are able to penetrate the breeding carcass, even if

there is no preexisting hole. Determining whether adaptation to the No Care selection
regime has involved a subtle change in carcass preparation will require more careful
examination of carcasses.

369 A second possibility is that selection in the No Care populations has favored 370 increased maternal investment into individual eggs. In many organisms, egg size or 371 size at birth is positively correlated offspring performance (e.g. Sinervo 1990; 372 Takakura 2004; Bashey 2006; Boivin and Gauvin 2009; Monteith et al. 2012; 373 reviewed in Fox and Czesak 2000). In N. vespilloides, the relationship between egg 374 size and larval performance depends upon the presence of post-hatching care 375 (Monteith et al. 2012). When parents are allowed to provision larvae, there is no 376 relationship between egg size and larval growth, but when post-hatching parental care 377 is removed this relationship is positive. These results suggest that an increase in egg 378 size can partially compensate for the absence of post-hatching parental care. It is 379 unclear however, whether the benefits of increased egg size observed by Monteith et 380 al (2012) would be realized in our experiment. For example, Monteith et al. (2012) 381 removed parents 72 hours after pairing (around the time of larval hatching). By this 382 time parents have usually created the feeding cavity in the carcass. In our experiment, 383 parents were removed well before they begin creating this cavity (as described 384 above). In addition, Monteith et al.'s (2012) No Care treatment involved cutting a 385 hole in the carcass with a razor blade immediately after removing the parents. This 386 was done to facilitate larval entrance into the breeding carcass. Our No Care treatment did not involve cutting a hole in the carcass. Thus, larvae had to chew their 387 388 way in on their own. Whether offspring from larger eggs are better at chewing their 389 way into the carcass remains unknown.

390 We have suggested ways in which removing post-hatching care may have led 391 to the evolution of traits in either offspring or parents. However, we wish to stress 392 that these hypotheses are not mutually exclusive. Adaptation to the absence of post-393 hatching parental care may have involved the joint evolution of traits expressed in 394 offspring and parents (i.e. coadaptation). For example, selection in the No Care 395 populations may have favored parents that make small holes in the carcass and larvae 396 that are able to exploit these holes effectively. Whatever traits may be involved, the 397 rapid evolution of reduced offspring dependence on post-hatching care in the No Care 398 lines suggests that there is extensive genetic variation for traits affecting larval 399 survival in the absence of post-hatching care. It is possible that such variation persists 400 cryptically in N. vespilloides because post-hatching parental care typically shields it 401 from natural selection.

402 Similar to a previous study (Eggert et al. 1998), we found that post-hatching 403 parental care positively affected the number of dispersing larvae. This effect did not 404 differ between the Full Care and No Care lines however. This suggests that 405 adaptation to the No Care selection regime has not involved traits that regulate brood 406 size after larvae become established on the carcass, either because selection at this 407 stage has been weak or because there is little heritable variation for the larval 408 performance at this stage. Our experiment does not allow us to distinguish between 409 these two possibilities. Although the effect of post-hatching parental care on brood 410 size did not differ between the Full Care and No Care lines, we note that in the 411 absence of care, the average brood size at dispersal in the No Care lines was 34-65 % 412 higher than the average brood size in the Full Care lines (Figure 3). It is possible that 413 further adaptation to the No Care selection regime will lead to greater divergence in brood size. 414

415 Although adaptation to the No Care selection regime reduced offspring 416 dependence on parental care it did not lead to consistent differences in the sensitivity 417 of larval mass to post-hatching care. In one block of the experiment post-hatching 418 care increased larval mass, but only when the beetles were descended from the Full 419 Care population. In the other block of the experiment, there was no effect of care on 420 larval mass in beetles descended from either population. The absence of a consistent 421 effect of post-hatching care on larval mass is somewhat surprising as previous studies 422 of *Nicrophorus* beetles have found that post-hatching care increases larval mass 423 (Rauter and Moore 1997; Eggert et al. 1998, Monteith et al. 2012). However, these 424 previous studies examined the effect of parental care under different environmental 425 conditions than we did. For example, Rauter and Moore (1997) and Eggert et al. 426 (1998) measured the impact of post-hatching care in broods whose size had been 427 standardized to minimize variation in larval mass caused by variation in brood size. 428 We did not manipulate brood size and were unable to control for variation in brood 429 size statistically because the presence or absence of post-hatching care substantially 430 changed the relationship between brood size and larval mass (see above). Second, 431 although Monteith et al. (2012) measured the effect of care on larval growth without 432 manipulating brood size, they used larger carcasses than we did and their No Care 433 treatment involved cutting a hole in the breeding carcass. Finally, all previous studies 434 examining the impact of parental care on larval mass have removed care at larval 435 hatching or very soon after. In contrast, our experiment involved removing parents 436 well before larvae hatch.

437 Despite these methodological issue, the lack of divergence between lines in
438 larval mass isn't entirely surprising for two reasons. First, the No Care and Full Care
439 selection regimes probably did not involve consistent differential selection on larval

440 mass. There was no selection on larval mass in the Full Care lines (mating was 441 random with respect to adult size, which is positively correlated with larval mass). In 442 the No Care lines, there was a history of selection for either large or small adults but 443 any history of this selection was likely erased by the protocol we used to create the 444 experimental populations (see supplemental methods). Second, we know from 445 previous studies that the total heritability of adult body size in N. vespilloides (which 446 is strongly positively correlated with larval mass at dispersal) is very low suggesting 447 that a response to even very strong selection on body size is unlikely to result in 448 evolutionary change over the short-term (Head et al. 2012; Schrader et al. unpublished 449 data).

450 Perhaps more importantly than the individual results described above, our 451 study suggests that experimental evolution can be applied to the study of parental 452 care. Much of what we know about the evolution of parent-offspring interactions 453 comes from quantitative genetic estimates of correlations between parental and 454 offspring traits (reviewed in Kölliker et al. 2012). While these studies have been tremendously valuable, they can only be used inferentially to understand how 455 456 selection has shaped parent-offspring interactions. We suggest that combining 457 quantitative genetic approaches with experimental evolution will provide more direct 458 insights into how parent-offspring interactions coevolve in response to different 459 selection regimes. For example, by manipulating the presence of post-hatching 460 parental care over many generations it might be possible to study not only how 461 offspring adapt to a change in parental care but also whether such adaptation changes 462 the direction or magnitude of genetic correlations between parental and offspring 463 behaviors. Similarly, it might be possible to manipulate the extent of parent-offspring 464 conflict in experimental populations (e.g. through manipulating the mating system)

465	and ask whether correlations between parental and offspring traits evolve in response
466	to antagonistic selection. Insects with flexible patterns of parental care and rapid
467	generation times, such as <i>N. vespilloides</i> , are ideal species for such experiments.
468	
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470	
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575

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Tables

579580 Table 1. Results of a general linear model of the effects of population (No Care or

Full Care), treatment (No Care or Full Care), the population by treatment interaction,
and experimental block on breeding success. See Figure 2 for sample sizes in each

583 treatment.

584

578

Factor	X^2	Р
Population	2.9	0.08
Treatment	44.5	< 0.00001
Population x Treatment	6.50	0.011
Block	7.61	0.0075

585

Table 2. Results of analyses of variance of the effects of population, treatment, the

587 population by treatment interaction, and experimental block on average brood size.

588

Factor	F	df	Р
Population	0.016	1, 177	0.89
Treatment	32.90	1,177	< 0.00001
Population x Treatment	2.35	1,177	0.13
Block	8.80	1,177	0.0034

589

Table 3. Results of a general linear model of the effects of population, brood size,

and block on carcass consumption. See Figure 4 for sample sizes in each treatment.

592

Factor	X^2	Р
Population	1.97	0.16
Brood Size	57.80	< 0.0001
Block	1.51	0.22

593

Table 4. Results of analyses of variance of the effects of population, treatment, and

their interaction on average larval mass for both blocks of the experiment considered

596 separately. Carcass mass was included as a covariate in block one of the experiment 597 but was not included in the analysis of block two.

598

Block one			
Factor	F	df	Р
Population	3.018	1,66	0.087
Treatment	8.39	1,66	0.0051
Population X Treatment	3.71	1,66	0.058
Carcass mass	10.77	1,66	0.0017
Block two			
Factor	F	df	Р
Population	2.19	1,107	0.14
Treatment	0.45	1,107	0.51
Population x Treatment	0.14	1,107	0.71

599

601 602	Figure Legends
602 603 604 605 606 607 608 609 610 611 612 613 614 615 616	Figure 1. A simplified schematic of the experimental populations and design. Dashed lines and solid lines represent populations that had been propagated without and with post-hatching parental care respectively. Numbers to the left of each arrow are the number of generations that each population had been maintained without or with post-hatching parental care. For example, NC1 had been propagated for 4 generations with no post-hatching parental care. All populations were passed through a single generation with full parental care to minimize differences between lines in transgenerational effects (indicated by solid lines enclosed in gray rectangles). The resulting offspring were then bred with or without post-hatching parental care (solid and dashed arrows respectively). For each pair we measured breeding success, brood size at dispersal, mean larval mass, and carcass consumption. A complete description of the establishment and maintenance of the experimental populations can be found in the supplemental material.
617 618 619 620 621 622 623 624	Figure 2. Breeding success of beetles descended from No Care (gray squares and dashed line) and Full Care (black squares and solid line) populations in the absence or presence of post-hatching care (x axis). Breeding success is the proportion of breeding attempts producing at least one dispersing larva. Numbers above each symbol are sample sizes. Data in the top panel are from block one and data in the bottom panel are from block two.
625 626 627 628 629	Figure 3. Mean brood size (± 1 S. E. M.) of beetle pairs descended from No Care (gray squares and dashed lines) and Full Care populations (black squares and solid lines) in the absence or presence of post-hatching care (x axis). Data in the top panel are from block one and data in the bottom panel are from block two.
630 631 632 633	Figure 4. The proportion of successful breeding bouts without parental care in which the breeding carcass was completely consumed when beetle pairs were descended from No Care (gray bars) and Full Care populations (black bars) in each experimental block. The sample size is indicated over each bar.
634 635 636 637 638 639	Figure 5. Mean larval mass (± 1 S. E. M.) of beetles descended from the No Care (gray squares and dashed lines) and Full Care (black squares and solid lines) populations in the absence or presence of post-hatching care. Data in the top panel are from block one and data in the bottom panel are from block two.

640 Figure 1.



Block two













1	Supplemental Methods
2	
3	The populations used in our experiment were created as part of an experiment
4	in which we artificially selected on adult body size (pronotum width) in populations
5	maintained with either full parental care (Full Care) or no post-hatching parental care
6	(No Care). The goal of this experiment was to test whether the presence of parental
7	care affects the response to selection (Jarrett et al. in prep). Below we describe the
8	selection protocol for the different experimental lines and how these lines were
9	combined to create the experimental populations.
10	
11	Selection on adult size
12	
13	We selected for large body size in two independent Full Care populations [Full
14	Care Large 1 (FCL1) and Full Care Large 2 (FCL2)] and two independent No Care
15	populations [No Care Large 1 (NCL1) and No Care Large 2 (NCL2)]. In these lines
16	we paired unrelated individuals in the largest 1/3 of the population to use as breeding
17	stock each generation. We selected for small body size in two independent Full Care
18	populations [Full Care Small 1 (FCS1) and Full Care Small 2 (FCS2)] and two
19	independent No Care populations [No Care Small 1 (NCS1) and No Care Small 2
20	(NCS2)]. In these lines we paired unrelated individuals in the smallest1/3 of the
21	population to use as breeding stock each generation. We also maintained two
22	replicate Full Care populations without selection on body size as controls [Full Care
23	Control 1 (FCC1) and Full Care Control 2 (FCC2)]. In the control lines, we paired
24	unrelated individuals randomly with respect to body size to use as breeding stock each
25	generation. Our experiment utilized beetles descended from the four No Care

26	populations (NCL1, NCL2, NCS1, NCS2) and the two Full Care Control populations
27	(FCC1 and FCC2). The Full Care Large / Full Care Small populations were not
28	included in the experiment because the lines were staggered temporally making it
29	impossible to breed beetles of similar ages to establish the experimental lines required
30	for the experiment. For the purposes of this study, we ignore differences between
31	lines in selection on adult size but note that there was no significant difference
32	between large and small selected lines used in the experiment (see below).
33	
34	Block 1
35	
36	Four generations after establishing the experimental populations we combined
37	the NCL1 and NCS1 populations into a single population [No Care 1 (NC1)] (see
38	Figure S1A). This was necessary to avoid inbreeding in one of the populations. At
39	this time there was no significant difference between NCL1 and NCS1 in pronotum
40	width (mean pronotum width in generation four: NCL1 = 4.52, mm, n = 239; NCS1 =
41	4.56 mm, n = 150; t $_{365.24}$ = -0.772, P = 0.447). To create the NC1 population, we
42	conducted reciprocal hybrid crosses between NCL1 and NCS1 populations ($n = 5$ in
43	each direction) and within population crosses involving unrelated individuals (NCL1
44	x NCL1, $n = 3$ and NCS1 x NCS1, $n = 7$). At the same time, we created a
45	corresponding full care population (FC1) by conducting reciprocal hybrid crosses
46	between the FCC1 and FCC2 populations ($n = 5$ in each direction) and mating
47	unrelated individuals from within each subpopulation (FCC1 x FCC1 and FCC2 x
48	FCC2, $n = 5$ each). These populations were crossed to be consistent with the protocol
49	described above for the NC1 population. The individuals crossed to create the NC1
50	and FC1 populations were chosen randomly with respect to pronotum width and the

51	crosses were conducted with full parental care to minimize differences between lines
52	in transgenerational effects of care. The offspring from these crosses were used as the
53	parents in the experimental generation.
54	
55	Block 2
56	
57	We replicated the experiment described above using a second pair of
58	experimental laboratory populations that differed with respect to the presence of post-
59	hatching parental care in the previous seven generations. The first population, No
60	Care 2 (NC2), was created by combining the NCL2 and NCS2 subpopulations 7
61	generations after establishing the experimental populations (Figure S1B). At this
62	time, there was no difference between the NCL2 and NCS2 populations in pronotum
63	width (mean pronotum width in generation seven: $NCL1 = 4.45$, mm, n = 417; NCS1
64	= 4.49 mm, n = 188; t $_{442.165}$ = -1.10, P = 0.271). To combine these populations we
65	conducted reciprocal hybrid crosses between the two lines ($n = 10$ in each direction)
66	and within subpopulation crosses involving unrelated individuals (NCL2 x NCL2, $n =$
67	10 and NCS2 x NCS2, $n = 10$). At the same time, we created a corresponding full
68	care population, Full Care 2 (FC2). This population was created by combining the
69	FCC1 and FCC2 subpopulations 7 generations after establishing the experimental
70	populations from the stock. To combine these populations we conducted reciprocal
71	hybrid crosses between the two lines ($n = 10$ in each direction) and within
72	subpopulation crosses involving unrelated individuals (FCC1 x FCC1, n = 10 and
73	FCC2 x FCC2, $n = 10$). The individuals crossed to create the NC2 and FC2
74	populations were chosen randomly with respect to pronotum width and the crosses
75	were conducted with full parental care to minimize differences between lines in

- transgenerational effects of care. The offspring from these crosses were used as the
- 77 parents in the experimental generation.

Supplemental Figures

Figure S1. A detailed schematic showing the creation of the experimental lines used in block one (A) and block two (B) of the experiment. Dashed lines indicate the No Care environment and solid liens indicate the Full Care environment. Abbreviations for each line follow those in the text above. All populations were passed through a single generation with full parental care to minimize differences between lines in transgenerational effects (indicated by solid lines enclosed in gray rectangles). The resulting offspring were then bred with or without post-hatching parental care (solid and dashed arrows respectively). For each pair we measured breeding success, brood size at dispersal, mean larval mass, and carcass consumption.



Figure S2. The relationship between mean larval mass and brood size in the presence
(left) or absence of post-hatching parental care. Data from block 1 are in the top row
and data from block 2 are in the bottom row.



