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Does avian malaria reduce fledging success: An experimental test of the selection hypothesis

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24 **Abstract**

25 Like many parasites, avian haematozoa are often found at lower infection intensities in
26 older birds than young birds. One explanation, known as the “selection” hypothesis, is that
27 infected young birds die before reaching adulthood, thus removing the highest infection
28 intensities from the host population. We tested this hypothesis in the field by experimentally
29 infecting nestling rock pigeons (*Columba livia*) with the malaria parasite *Haemoproteus*
30 *columbae*. We compared the condition and fledging success of infected nestlings to that of
31 uninfected controls. There was no significant difference in the body mass, fledging success, age
32 at fledging, or post-fledging survival of experimental versus control birds. These results were
33 unexpected, given that long-term studies of older pigeons have demonstrated chronic effects of
34 *H. columbae*. We conclude that *H. columbae* has little impact on nestling pigeons, even when
35 they are directly infected with the parasite. Our study provides no support for the selection
36 hypothesis that older birds have lower parasite loads because parasites are removed from the
37 population by infected nestlings dying. To our knowledge, this is the first study to test the
38 impact of avian malaria using experimental inoculations under natural conditions.

39

39 **Introduction**

 40 Parasites influence fundamental aspects of the evolutionary ecology of their hosts, such
 41 as population dynamics (Anderson and May 1978; Anderson 1979) and life history evolution
 42 (Hochberg et al. 1992). The impact of parasites on host fitness depends partly on the age at
 43 which hosts become infected. A common pattern in host-parasite interactions is that younger
 44 individuals have higher parasite loads than adults (Gregory et al. 1992; Hudson and Dobson
 45 1997). Sol et al. (2003) considered three hypotheses to explain this pattern. The “selection”
 46 hypothesis suggests that highly parasitized juvenile hosts die before they reach adulthood,
 47 removing large numbers of parasites from the population. The “immunity” hypothesis suggests
 48 that the developing immune system of juveniles is not yet capable of killing parasites, while
 49 adults are much more effective at reducing parasite intensity. The “vector exposure” hypothesis
 50 suggests that adult behavior reduces their exposure to infected vectors, and thus parasites,
 51 compared to juveniles.

 52 Sol et al. (2003) evaluated these hypotheses using data from a study of feral rock pigeons
 53 (*Columba livia*) infected with malaria parasites (*Haemoproteus columbae*) vectored by pigeon
 54 louse flies (Hippoboscoidea: *Pseudolynchia canariensis*). The authors rejected the vector
 55 exposure hypothesis because they found that adult pigeons (> 6 months old) are not, in fact,
 56 exposed to fewer vectors than juvenile pigeons (Sol et al. 2000). Although the authors reported
 57 higher rates of juvenile mortality (61%) compared to adult mortality (33%), consistent with the
 58 selection hypothesis, selection in their study was not strong enough to explain the lower number
 59 of parasites observed in adult birds. The youngest birds in Sol et al.'s study had already fledged
 60 from the nest; however, the greatest impact of *H. columbae* on pigeons may occur while birds are
 61 still in the nest. We conducted a study to test the impact of *H. columbae* on the condition and

62 fledging success of younger, nestling rock pigeons. We used an experimental approach in which
 63 we compared nestlings injected with *H. columbae* to control birds not injected with the parasite.

64 At least 200 species of *Haemoproteus* are known to infect birds worldwide (Martinsen et
 65 al. 2008). Perez-Tris et al. (2005) classified *Haemoproteus* as an avian malaria parasite because
 66 members of the genus were nested phylogenetically within the genus *Plasmodium*. *H. columbae*
 67 is a parasite of pigeons and doves that uses blood-feeding pigeon flies as vectors (Valkiūnas
 68 2005). The parasite enters a feeding fly and reproduces in its midgut, where *H. columbae*
 69 oocysts attach to the gut wall. Once mature, the oocysts burst and release infective sporozoites
 70 that migrate from the fly's gut into its salivary glands. The fly then injects these sporozoites into
 71 a pigeon when it feeds. *H. columbae* reproduces asexually in the lungs of the pigeon, then
 72 invades and matures in the red blood cells (Ahmed and Mohammed 1978).

73 *Haemoproteus* species can have several negative effects on host fitness. These effects
 74 include reductions in host body condition (Merino et al. 2000), lower reproductive success
 75 (Marzal et al. 2004; Tomas et al. 2007), and even death (Atkinson and Forrester 1988; Sol et al.
 76 2003). Studies of the impact of malaria on juvenile birds have consisted of observational studies
 77 in the field (Sol et al. 2003), and experimental studies using captive birds (Yorinks and Atkinson
 78 2000; Garvin et al. 2003). The goal of our study was to use an experimental approach under
 79 field conditions. We infected nestling birds with malaria parasites to test the impact on body
 80 mass, fledging success, age at fledging, and post-fledging survival of experimental versus control
 81 birds. Studies with captive birds suggest that the most pathogenic phase of the *Haemoproteus*
 82 life cycle occurs when parasites enter red blood cells to mature (Atkinson and Forrester 1988;
 83 Atkinson and van Riper 1991). In the case of *H. columbae* this takes place about 24-37 days
 84 after infection (Ahmed and Mohammed 1978). Since pigeons fledge at about 32 days of age, it

85 is not possible to be sure that fledglings are infected with malaria parasites, short of
 86 experimentally infecting them. Experimental manipulation is the most powerful approach for
 87 testing the impact of parasites on hosts in any case (McCallum and Dobson 1995). To our
 88 knowledge, this is the first study to test the impact of avian malaria parasites using experimental
 89 inoculation under natural conditions.

90

91 **Materials and Methods**

92 We experimentally manipulated *H. columbae* in nestling rock pigeons. The study took
 93 place August-November 2009 under a highway overpass in Draper, Utah, USA (40°31'36" N,
 94 111°53'28" W). We visited the field site every 2-3 days throughout the study period. Nestlings
 95 were weighed at each visit to the nearest 1.0 g with a pesola scale. Our experiment was
 96 restricted to nests with two nestlings, the normal number for rock pigeons. Nests were
 97 sequentially assigned to one of three treatment groups: experimental (n = 12 nests), control (n =
 98 13), or background (n = 12). When nestlings were 4 - 7 days old (50 - 150 g), those at
 99 experimental nests were injected with a suspension of *P. canariensis* flies infected with *H.*
 100 *columbae* (Ahmed and Mohammed 1978). We created the infected fly suspension by feeding
 101 flies (bred from wild stock) on heavily infected captive birds. Following 10-12 days on a bird,
 102 flies were placed in vials and taken to the field site, where batches of ten live flies were
 103 macerated in 1000 μ L of phosphate buffered saline for three minutes. Experimental nestlings
 104 were injected intraperitoneally with 500 μ L of the infected fly suspension using a 0.5 cc syringe.
 105 Control birds were injected with 500 μ L of another suspension made using uninfected flies.
 106 Background birds were handled but not injected.

107 Prior to the field experiment, we conducted a test of the inoculation method using 27 wild
 108 trapped, captive rock pigeons. After blocking by capture date and site, 13 randomly chosen birds
 109 were injected with a suspension of infected flies, as described above. Fourteen control birds
 110 were injected with a suspension of uninfected flies. At 25, 35, and 42 days post injection, blood
 111 samples were taken from all birds and smears were prepared for examination. Each smear was
 112 carefully examined under oil immersion at 1000x for 10 minutes; if parasites were detected, then
 113 the number of parasites was quantified in 25 microscope fields per bird. All 13 experimental
 114 birds were infected with *H. columbae*, while none of the 14 control birds was infected.

115 When nestlings were approximately ten days old they were fitted with a numbered
 116 aluminum band and three plastic color bands. To score fledging success we observed and
 117 identified birds after they left the nest on the basis of their color band combinations. We
 118 conducted a thorough census of all birds at the bridge during each visit to the field site. We also
 119 searched for banded birds at other bridges within 8 km of the study site in order to determine
 120 whether newly fledged birds were dispersing from the natal site.

121 We continued to monitor birds at the bridge for 50 days post injection (ca. 25 days post
 122 fledging) because peak parasitemia can be delayed for this long after injection (extrapolated from
 123 Ahmed and Mohammed (1978)). To confirm experimental infections, we examined the blood of
 124 birds after they fledged. We used walk-in traps to capture pigeons from 30-50 days post
 125 injection. Blood samples were taken and birds immediately released. Blood smears were
 126 prepared and examined back in the lab.

127 Data were analyzed using Prism[®] v.5.0b (GraphPad Software, Inc.). Power analyses
 128 were conducted in G*Power 3 with an error probability set at 0.05 (Buchner et al. 1997). Where
 129 necessary, data were log transformed for normalization. To avoid pseudoreplication (Hurlbert

130 1984) we averaged values for nestlings within each nest. We used one-way ANOVAs to
 131 compare parasite abundance and host age and mass at fledging among treatments. A repeated-
 132 measures ANOVA was used to compare the number of birds per nest at hatching, fledging, and
 133 one, two, and three weeks post-fledging.

135 **Results**

136 Three times as many experimental birds were infected as control or background birds
 137 (Fig. 1a); the three groups also differed in parasite abundance (Fig. 1b; ANOVA $F_{2,17} = 4.25$, $P <$
 138 0.05). Dunnett's post hoc comparisons confirmed that experimental birds had significantly more
 139 parasites than controls ($P < 0.05$), while control and background birds did not differ significantly
 140 ($P > 0.05$).

141 There was no significant difference in the age of birds at fledging, nor body mass prior to
 142 fledging (see Table 1). There was no significant difference in the proportion of nests that
 143 fledged at least one offspring ($\chi^2 = 0.005$, $P = 0.99$). There was also no significant effect of
 144 treatment on the mean number of birds fledged per nest, nor the number of birds observed after
 145 fledging (Fig. 2; repeated measures ANOVA, treatment $F_{2,34} = 0.64$, $P = 0.53$). There was a
 146 significant effect of time (Fig. 2; time, $F_{4,136} = 43.32$, $P < 0.0001$), but no significant interaction
 147 between time and treatment (time*treatment, $F_{8,136} = 0.49$, $P = 0.86$).

148 We reanalyzed the data after excluding naturally infected control and background birds,
 149 as well as experimental birds for which we could not confirm infection. We still found no
 150 significant difference in age at fledging ($F_{2,31} = 0.53$, $P = 0.60$) mass at fledging ($F_{2,31} = 1.01$, $P =$
 151 0.38), or the proportion of nests that fledged at least one offspring ($\chi^2 = 0.01$, $P = 0.99$).

152 Our experiment had considerable power (1.0) to detect the level of juvenile mortality
 153 (61%) reported by Sol et al. (2003); we had power of 0.8 to detect mortality of at least 30%
 154 (effect size of $f = 0.55$).

156 Discussion

157 Our goal was to experimentally test the “selection” hypothesis. This hypothesis,
 158 reviewed by Gregory et al. (1992), states that lower parasite loads of adults, compared to
 159 juveniles, are the result of heavily infected juveniles dying before adulthood, removing parasites
 160 from the population. Previous tests of this hypothesis involving avian malaria have focused on
 161 juvenile (fledged) birds and relied on observational data (Sol et al. 2003; van Oers et al. 2010).
 162 These studies provided some support for the selection hypothesis, but the intensity of selection
 163 measured could not fully explain differences in juvenile and adult parasite loads. It was
 164 conceivable, therefore, that the greatest impact of *H. columbae* on pigeons takes place while they
 165 are still in the nest.

166 Our results provided no support for the selection hypothesis because there was no impact
 167 of malaria on any of the components of host fitness we measured. Specifically, there was no
 168 significant difference in the body mass, fledging success, age at fledging, or post-fledging
 169 survival of experimental versus control birds. We are confident that our measures of post-
 170 fledging survival were accurate because none of the birds from our study were observed at other
 171 bridges (see methods). Young pigeons do not normally disperse until three months of age, in any
 172 case (Johnston and Janiga 1995).

173 The results of our study were unexpected, given that Sol et al.'s longer-term study
 174 demonstrated that *H. columbae* has a significant negative impact on pigeon fitness. The fact that

175 malaria had no detectable impact on fledging success in our study was not due to unusually low
 176 rates of fledging in both experimental and control birds. Fledging success was 73% (Fig. 2),
 177 similar to that in other studies of feral pigeons (reviewed by Johnston and Janiga (1995), Table
 178 18.4 (values adjusted for hatching rates)). Similarly, the fact that malaria had no detectable
 179 impact on fledging was not due to methodological problems with the creation of experimental
 180 infections. The malaria parasite levels in our study were comparable to those observed in other
 181 studies of naturally infected pigeons (Kartman 1949; Klei and DeGuisti 1975; Paperna and
 182 Smallridge 2002). However, *H. columbae* may affect hosts only at levels higher than what we
 183 observed (Earle et al. 1993; Paperna and Smallridge 2002). For example, the *H. columbae* levels
 184 in Sol et al.'s (2003) study were among the highest ever recorded for feral rock pigeons.

185 Another factor that could conceivably contribute to why the birds in our study did not
 186 appear to be affected by *H. columbae*, compared to the reduction in survival shown for older
 187 birds by Sol et al. (2003), is that nestling pigeons could have higher tolerance to parasites than
 188 older birds. Nestlings are fed a rich diet of crop milk by both parents. The milk, which consists
 189 of the sloughed lining of the parents' crop, is very high in fat and protein (Johnston and Janiga
 190 1995). It would be interesting to test the impact of *H. columbae* on nestlings fed a less nutritious
 191 diet.

192 A few control and background birds were naturally infected with *H. columbae*. However,
 193 infection levels were still significantly higher in the experimental groups than the control or
 194 background groups. Even after excluding the naturally infected birds, we did not find that
 195 malaria parasites affected age, mass, or fledging success.

196 Since *H. columbae* had no apparent effect on nestling rock pigeons, our study does not
 197 provide support for the "selection hypothesis". Sol et al. (2003) reported results that were

198 consistent with selection hypothesis; however, selection in their study was not strong enough to
 199 explain the differences in parasitemia they observed between juvenile and adult pigeons.
 200 Because Sol et al. (2000, 2003) reported data ruling out the “vector exposure” hypothesis, they
 201 suggested a combination of the selection and immunity hypotheses may explain the fact that
 202 juvenile birds have higher parasitemia than adult birds. Our data provide no reason to disagree
 203 with this assessment.

204 To our knowledge, this is the first study to test the impact of avian malaria parasites using
 205 experimental inoculation under natural conditions. This approach has several advantages. First,
 206 like many malaria parasites, *H. columbae* takes several weeks to appear in the peripheral blood
 207 after the host is infected. This fact makes early infections difficult to detect without more
 208 invasive methods, such as collection of organ tissues (Valkiūnas 2005; Cosgrove et al. 2006).
 209 Experimental infections get around this problem. Second, inoculating hosts with parasites has
 210 the strong advantage of controlling for factors that could lead to spurious negative correlations
 211 between parasite load and host fitness (Hawlana et al. 2006; Blanchet et al. 2009). The greatest
 212 limitation of our study is that the modest sample sizes limit our ability to detect relatively small
 213 effect of malaria parasites on birds. For example, to detect a 10% reduction in juvenile survival
 214 with a power of 0.8 would require a sample of 93 nests per treatment for a total of 279 nests. A
 215 study of this magnitude may be feasible in the future using feral Rock Pigeons and *H. columbae*.

216
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226

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293

293 Table 1. Age of birds at fledging and body mass prior to fledging. Values are grand means
 294 (\pm SE) of the mean value per nest.

295		Experimental	Control	Background	Test statistic	<i>P</i>
296	Age in days	32.3 \pm 0.5	31.8 \pm 0.6	32.3 \pm 0.6	<i>F</i> = 0.25	0.78
297	(number of nests)	(11)	(12)	(11)		
298	Mass in grams	298 \pm 14.9	311.3 \pm 12.5	313.1 \pm 11.4	<i>F</i> = 0.35	0.71
299	(number of nests)	(11)	(12)	(11)		

316 **Figure legends**

317

318 Fig. 1. Prevalence (a) and mean abundance (+ SE) (b) of malaria parasites 30-50 days after
319 treatment.

320

321 Fig 2. Mean (\pm SE) offspring observed per nest. The mean (\pm SE) number of offspring fledged
322 per nest did not differ significantly among treatments; see text.

Fig 1a

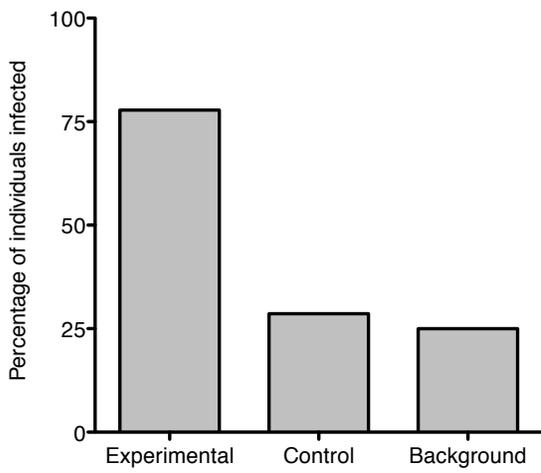


Fig 1b

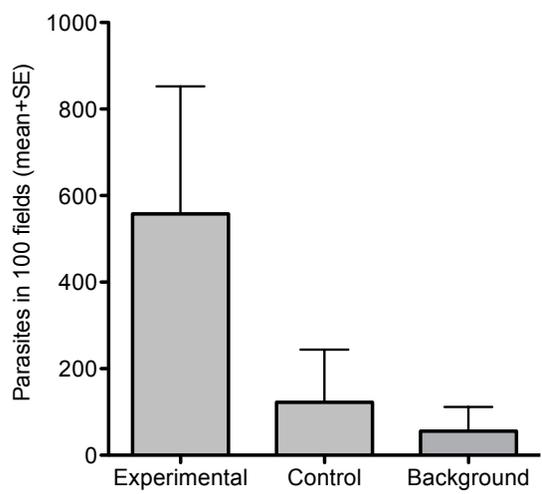


Fig 2

