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Parasitized Mates Increase Infection Risk for Partners

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ABSTRACT: Individuals can gain fitness benefits and costs through their mates. However, studies on sexual selection have tended to focus on genetic benefits. A potentially widespread cost of pairing with a parasitized mate is that doing so will increase an individual's parasite abundance. Such a cost has been overlooked in systems in which parasites are indirectly transmitted. We manipulated the abundance of the nematode parasite *Trichostrongylus tenuis*, an indirectly transmitted parasite, within pairs of wild red grouse *Lagopus lagopus scoticus* in spring. Parasite levels were correlated within pairs before the experiment. We removed parasites from males, females, or both members of the pair and evaluated individual parasite uptake over the subsequent breeding period. At the end of the breeding season, an individual's parasite abundance was greater when its mate had not been initially purged of parasites. This cost appeared to be greater for males. We discuss the implications of our results in relation to the costs that parasites may have on sexual selection processes.

Keywords: red grouse, sexual selection, direct benefits, parasitism.

Introduction

The ability to choose a high-quality mate is known to provide individuals with genetic benefits (Fisher 1930; Zahavi 1975, 1977; Kodric-Brown and Brown 1984). Indeed, most of our understanding of the evolution of sexual selection comes from studies that have examined these forms of benefits (Jennions and Petrie 2000; Kotiaho and Puurtinen 2007). However, individuals may also gain direct fitness benefits from partners who invest more in the production of offspring, typically through better parental care, food provisioning, fecundity, or fertility (Andersson 1994; Cornwallis and Uller 2009). Yet these forms of benefits have received scant attention, especially in vertebrates. Parasites may play a key role here, because they can negatively

affect the fitness benefits that individuals receive from their partners (Møller et al. 1999). If an individual is heavily parasitized, this could influence its mate's parasite load, its future reproductive output (Hudson 1986a; Gustafsson et al. 1994), and its survival (Marzal et al. 2008; Martínez-de la Puente et al. 2010).

Because of the deleterious effects of parasites on hosts, it is expected that the parasite abundance of one pair member might influence levels of parasitism in its mate (Able 1996). The infection risk of mating with a parasitized individual may depend on the mode of parasite transmission. Parasites may be transmitted through direct contact (e.g., as occurs in sexually transmitted diseases) or indirectly via the environment. The risks and costs of direct transmission between mates have been well studied in sexually transmitted parasites and diseases (Sheldon 1993; Knell and Webberley 2004; White et al. 2010). However, the costs of indirectly transmitted parasites have been poorly studied, and experimental evidence for costs is lacking. These types of parasites are widespread and are transmitted via the environment, either with or without intermediate vectors. Blood parasites, such as *Trypanosoma* species, and hemsporidians are common in reptiles, birds, and mammals and are transmitted from host to host via the bite of blood-feeding invertebrates (Loye and Zuk 1991). The effects of these parasites can have profound fitness-related consequences on their hosts (Merino et al. 1996, 2000; Potti and Merino 1996; Lachish et al. 2011). Similarly, many intestinal parasites, such as coccidia or nematodes, are indirectly transmitted and can cause a wide range of negative effects on hosts (Hudson et al. 1998; Fox and Hudson 2001; Horak et al. 2004, 2006; Mougeot et al. 2005a, 2010a, 2010b; Martínez-Padilla et al. 2007; Lemus et al. 2010). Indirectly transmitted parasites can have a profound effect within a sexual selection context, because choosing a highly parasitized mate may impose the cost of increasing the partner's risk of infection throughout the breeding period.

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Most vertebrate pairs share their space and time during the breeding period, so their parasites share a common environment, which increases the risk of infection between mates. Two hypotheses consider the risk of infection in these situations. The avoidance hypothesis suggests that individuals will avoid highly parasitized mates (Borgia 1986), whereas the contagion indicator hypothesis (Able 1996) focuses on the risks of infection for both offspring and mates. Both hypotheses suggest the existence of mechanisms and behavioral responses that facilitate the recognition, and therefore the avoidance, of highly parasitized mates in a wide variety of taxa (Borgia 1986; Able 1996; Loehle 1997; Ehman and Scott 2002; Kavaliers et al. 2005). A premise underlying both hypotheses is that risk of parasite transmission between pairs is high. Although highly plausible, empirical evidence for indirectly transmitted parasites is scarce and generally limited to circumstantial support (*sensu* Clayton 1991) or to laboratory or captivity conditions (Hillgarth 1996; Luong et al. 2000; Abbot and Dill 2001).

Our understanding of sexual selection is based largely on the idea that females choose males and increase their fitness by doing so (Hamilton and Zuk 1982; Folstad and Karter 1992). However, recent studies suggest that males can also gain direct fitness benefits (Thrall et al. 1997), thus promoting the coevolution of preferences toward phenotypic traits that indicate quality (i.e., that reveal the benefits that the bearer can offer) in both sexes. In a variety of species, females display traits that may signal their quality (Potti and Merino 1996; Vergara et al. 2009, 2011; Martínez-Padilla et al. 2011) as well as their reproductive value (Ilmonen et al. 2000). To better understand whether mating with parasitized partners represents a cost for individuals within a sexual selection context, we need to test whether the cost of mating with a parasitized partner could be asymmetrical or sex-dependent (i.e., greater for one sex than for the other).

Here, we experimentally test how parasite abundance in one member of the pair determines parasite abundance in its mate. Our study species is the red grouse *Lagopus lagopus scoticus*, a territorial, monogamous bird species that lives on heather moorlands in the United Kingdom (Watson and Moss 2008). In autumn, males establish their territories and keep them over winter. In spring, successful territory holders are chosen by females, with males that are in better condition and that have bigger ornaments being more likely to obtain a mate (Redpath et al. 2006a). Families stay together until autumn, when young reach independence (Watson and Moss 2008). We studied infection patterns in *Trichostrongylus tenuis*, the main nematode parasite of red grouse, which is transmitted indirectly and has no alternative host or vectors. It reduces fecundity and survival and has a destabilizing effect on red grouse

population dynamics (Hudson 1986a, 1998; Martínez-Padilla et al. 2007; Mougeot et al. 2009, 2010a, 2010b; Redpath et al. 2006b).

We initially explored patterns of parasite abundance within pairs before manipulation and then altered parasite abundance within individuals during pair formation in early spring. Specifically, we removed parasites of either males, females, or both members of the pair and explored subsequent patterns of parasite uptake of the mate throughout the breeding period. We first expected that parasite reinfection rates would be lower for both males and females if their mate was purged of parasites, compared with a naturally infected mate. Second, if both males and females gain equivalent direct fitness benefits from reduced parasite transmission risk, then treatment should have similar effects on male and female parasite uptake rates.

Material and Methods

Catching and Tagging

In spring 2009, we caught 77 pairs in 4 populations, including 3 populations in England (Moorhouse, Geltsdale, and Catterick) and 1 population in Scotland (Invermark; see Martínez-Padilla et al. 2011 for further details). We caught wild red grouse pairs by dazzling (rendered immobile by bright spotlight) and netting them at night. Each pair member was individually ringed and fitted with a radio collar (TW3-necklace tag, Biotrack), which allowed subsequent location and recaptures. From autumn until courtship in early spring, male red grouse are highly territorial; they have to keep a territory to survive overwinter and to attract a female in spring (Watson and Moss 2008). Before spring, females associate with territorial males. Once paired, males guard their mates intensively before egg laying, so pairs stay close to each other. After hatching, pairs are no longer territorial, but males and females stay together during brood rearing until the beginning of autumn, when families break up (Watson and Moss 2008).

Experiment

At first capture (C1; March 7, ± 3 weeks; see fig. A1 for further details), a subsample of 45 of the 77 captured pairs were randomly assigned to one of two treatments: (1) focal individual dosed and mate not dosed (T1; 23 pairs) or (2) both pair members dosed (T2; 22 pairs). The sex of treated birds was randomly chosen. Capture time did not differ between treatment groups for either males or females (both $P > .463$). Dosed individuals were treated orally with 1 mL of anthelmintic (Levamisole hydrochloride 3%, Nilverm Gold, Schering-Plough Animal Health, Welwyn Gar-

den City, UK). This Levamisole dose has been shown to be highly effective at purging red grouse from their adult *Trichostrongylus tenuis* nematodes (Mougeot et al. 2009; Martínez-Padilla et al. 2011). When only 1 individual in a pair was dosed, its mate was given 1 mL of water as a control.

One month later, we recaptured (second capture; C2) at least 1 member of the pair (T1: 21 pairs, 14 males, 16 females; T2: 19 pairs, 10 males, 8 females). Time from C1 to C2 did not differ between treatment groups for males or females (both $P > .577$). It was not always possible to catch both pair members. However, at night, pair members roost no more than 5–10 m apart, so we could always confirm by radio tracking whether the individual that we caught was still paired with the same mate. After C2, we continued monitoring pairs to ensure that mates stayed together during the whole reproductive period. During laying and at hatching (late April through May), we also relocated all individuals during the day to confirm that they were with the same mates. In August, although red grouse families remain together (Watson and Moss 2008), we located the pairs again during the day for the same purpose. When one member of the pair died (10 males and 4 females died), the remaining individual was subsequently excluded from the analyses. We also excluded from analyses those pairs for which we were unsure whether they had been together during the whole study period (9 pairs). In early September, we recaptured all the individuals alive (third capture; C3). We caught both birds, or at least one member of the pair (T1: 25 individuals, 13 males, 12 females; T2: 24 individuals, 14 males, 10 females). Capture time did not differ between treatment groups for males or females (both $P > .420$).

T. tenuis Life Cycle and Parasite Abundance Estimates

Most red grouse older than 2 months are parasitized by *T. tenuis* (Hudson 1986b), which has a life cycle with no alternative hosts (fig. A2). Prevalence is close to 100%, and abundance may reach approximately 30,000 worms per bird (Hudson 1986b). At low worm intensity, the effect of worms on the life-history traits of individuals are not apparent (Hudson 1986a). Adult worms inhabit the birds' caeca, and their eggs are voided in caecal droppings on the moor. After approximately two weeks, depending on weather conditions (Moss et al. 1993), they develop into infective larvae that migrate out of caecal droppings onto terminal heather shoots, ready to reinfect new grouse. Once ingested, larvae migrate to the caecum, where they develop into adult worms (Watson and Moss 2008). Depending on environmental conditions, infective larvae can arrest their development for periods ranging from weeks to months. After ingestion, mature female worms can

quickly produce eggs that have to be excreted to develop into new infective larvae (Shaw and Moss 1989a, 1989b; Shaw et al. 1989).

We estimated *T. tenuis* worm abundance by determining parasite egg concentration in individual fecal samples, using a previously validated technique (Seivwright et al. 2004, 2005). At each capture, birds were kept overnight in individual pens to collect fecal samples and were released in the early morning. Not all individuals produced fecal samples overnight, which led to reduced sample size for some capture events. All samples were analyzed within 3 weeks after collection to reliably estimate *T. tenuis* abundance (Seivwright et al. 2004).

Statistical Analyses

We used SAS, version 9.2. Parasite abundance data were fitted to models using a negative binomial distribution of errors and a log link function. To analyze the association between male and female worm burdens within pairs, we used generalized linear models (GENMOD procedure) with female worm burden as the dependent variable, male worm burden as a covariate, and site included as a factor. To analyze the effectiveness of parasite dosing and the experiment effects on parasite abundance, we used generalized linear mixed models (GLIMMIX procedure) with recapture (C1 and C2) and sex as factors to explore variation in male and female parasite levels over time. Individual was included as a random factor to avoid pseudoreplication. To test the effect of our treatments in autumn (C3), estimated worm abundances in males and females were considered as dependent variables (GENMOD procedure). Site and treatment were included as fixed effects. We also tested the effect of sex and pair treatment on final parasite levels. In this model, sex, treatment, and their interaction, together with site, were included as factors. Post hoc comparisons were used to show the interactions between groups (T1 and T2) and sex. Data are given as arithmetic means (\pm SEs).

Results

Association of Parasites between Males and Females within Pairs

Parasite abundance differed between sites ($\chi^2 = 10.39$, $P = .016$). After controlling for site effects, worm burdens of females were positively associated with those of their mates ($\chi^2 = 5.38$, $P = .020$; fig. 1), and this relationship was consistent across sites (site \times male parasite burden interaction: $\chi^2 = 2.19$, $P = .534$).

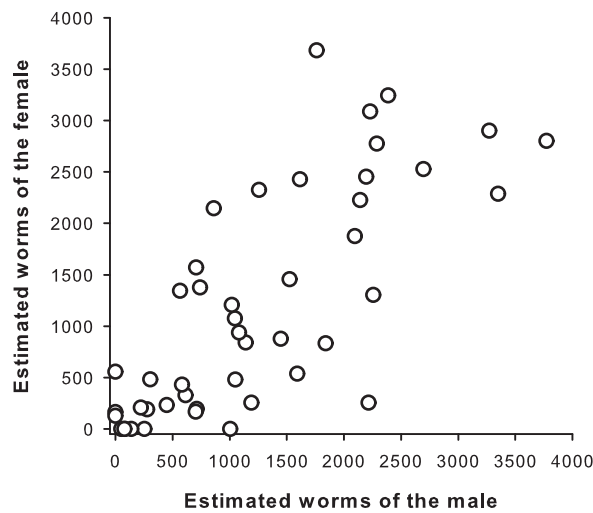


Figure 1: Relationship between the *Trichostrongylus tenuis* parasite abundance of males and females within red grouse *Lagopus lagopus scoticus* pairs in spring.

Effectiveness of the Initial Parasite Purging

Before dosing (C1), initial parasite levels did not differ between treatment groups ($\chi^2 = 0.28$, $P = .594$; fig. 2) or between sexes (sex: $\chi^2 = 1.49$, $P = .222$; treatment \times sex interaction: $\chi^2 = 0.69$, $P = .405$; fig. 2). From C1 to C2, parasite abundance decreased more in dosed individuals than in controls (treatment \times recapture interaction: $\chi^2 = 24.16$, $P < .001$; fig. 2), and a similar decrease occurred in both sexes (treatment \times sex \times recapture interaction: $\chi^2 = 1.49$, $P = .223$). After 1 month, parasite purging reduced *Trichostrongylus tenuis* abundance from 881 ± 144 (C1) to 27 ± 27 (C2) worms per host in males and from $1,320 \pm 283$ (C1) to 106 ± 84 (C2) worms per host in females. In untreated birds, after 1 month, *T. tenuis* abundance changed from $1,312 \pm 321$ (C1) to $1,420 \pm 577$ (C2) worms per host in males and from $1,068 \pm 252$ to $1,243 \pm 407$ worms per host in females.

Sex Mediated Effects of Treatments on Parasite Abundance at the End of the Experiment

In autumn (C3), parasite abundance differed between pair treatment groups (treatment: $\chi^2 = 11.93$, $P < .001$; site: $\chi^2 = 10.14$, $P = .017$). Birds whose mates did not receive a dose had more parasites than did birds whose mates received a dose (T1: $1,329 \pm 164$; T2: 590 ± 59 ; fig. 2). Males and females also differed with respect to their in-

fection levels in autumn, with males having more parasites than females ($\chi^2 = 4.03$, $P = .044$; fig. 3). More importantly, we found that sex mediated the effect of treatments on parasite levels (treatment \times sex interaction: $\chi^2 = 4.00$, $P = .045$). Males that were initially dosed ended up with more parasites if their mate was not dosed, compared with those whose mate was also dosed ($\chi^2 = 5.37$, $P = .020$; T1: $1,952 \pm 175$; T2: 563 ± 94 ; fig. 3a). In contrast, females' worm abundance in autumn (C3) did not differ between the two experimental groups ($\chi^2 = 0.69$, $P = .709$; T1: 794 ± 156 ; T2: 626 ± 68 ; fig. 3b).

Discussion

We found a positive correlation between male and female nematode parasite burdens at the time of pairing in early spring. Moreover, our experiment showed that birds cleared of parasites had greater reinfection rates if their mates had natural levels of parasites than if their mates were free of parasites. This was particularly noticeable in males but was less noticeable in females. To the best of our knowledge, this is the first experimental evidence in wild animals showing that an individual that pairs with a parasitized mate will develop an increased parasite load. Because these parasites reduce the survival and reproductive output of their hosts, our results are consistent with the idea that mating with a parasitized mate may impose a direct cost for individuals.

Our correlative data suggested that there may have been assortative mating in relation to parasite abundance before breeding in red grouse. We can only speculate about the mechanisms behind this pattern. It has been suggested that parasites decrease the expression of red grouse combs in both males and females (Mougeot et al. 2005a; Martínez-Padilla et al. 2011), which may provide a behavioral explanation for our correlative results. Also, genetic quality and parasite resistance genes may also suggest a potential mechanism. Genes involved in the major histocompatibility complex (MHC) play a key role in immune defense (Klein 1986), and multiple studies have described relationships between specific MHC alleles or heterozygosity with nematode infections (Oliver et al. 2009). In fact, experimental *Trichostrongylus tenuis* infection in red grouse identified genes of the MHC (Webster et al. 2011) as being involved in immune function. Mate choice is often evoked as a means of maintaining diversity of the MHC in vertebrates (Penn and Potts 1999; Tregenza and Wedell 2000). The maintenance of such diversity can be explained by preference for heterozygotes or avoidance of partners with similar MHC alleles (Tregenza and Wedell 2000; Richardson et al. 2005; Bos et al. 2009). Both scenarios may explain the assortative mating that we found regarding parasite abundance in pairs of red grouse before breeding, but

additional research is needed to confirm this finding and to explore the underlying mechanisms.

Our experimental results confirm that red grouse pay a cost by increasing own parasitism risk when mating with a parasitized individual. Our focus was not to identify the mechanism of such parasite intake, but solely to quantify parasite uptake and to test whether parasites represent a cost for breeding partners. These patterns provide empirical support for one of the assumptions that underpins both the contagion and avoidance models of parasite-mediated sexual selection (Able 1996). Both hypotheses suggest that individuals that avoid parasitized mates can have direct benefits through reducing their or their offspring's risk of becoming parasitized. Our results show that parasite infection risk exists and that choosing a highly parasitized mate increases such risk. Therefore, choosy individuals may pay the cost of wrong mate choice by increasing their risk of becoming parasitized.

Several mechanisms may explain our results. Because direct transmission of *T. tenuis* is not possible, indirect infection could be attributable to either variation in susceptibility or exposure to parasites. Aggressiveness may explain the susceptibility of males to parasites in red grouse. Testosterone-dependent aggressiveness may impose the physiological cost of immunosuppression, which may increase susceptibility to parasite infestation (Folstad and Karter 1992; Mougeot et al. 2005b). However, we cannot rule out other potential mechanisms, such as different reproductive costs for parasitized and unparasitized mates. Birds paired with parasitized mates may increase their level of investment in reproduction, thereby increasing their own susceptibility to parasites (Hudson 1986a).

In red grouse, differences in exposure can greatly contribute to explaining differences in parasite abundance levels (Hudson 1986b). A plausible explanation for the infection patterns that we found within pairs is shared territory and, therefore, shared parasite exposure. A close proximity between mates throughout the relatively long breeding season (March–September) means that pair members are exposed to the parasite larvae voided onto the moor by their mate, which implies a high risk of parasite uptake within pairs. Territory sharing may also explain why there is a positive association between parasite burdens in pair members in spring, although our experiment cannot tease apart the cause or consequence of mate choice. We now need to disentangle the relative effect of exposure and susceptibility to understand the consequences of parasitism to mate choice decisions in particular and to sexual selection in general.

Irrespective of the mechanism, our experiment showed that there is a significant risk of parasitism between members of a pair. Our results can be applied to a wide range of organisms, because sharing territories (space) during

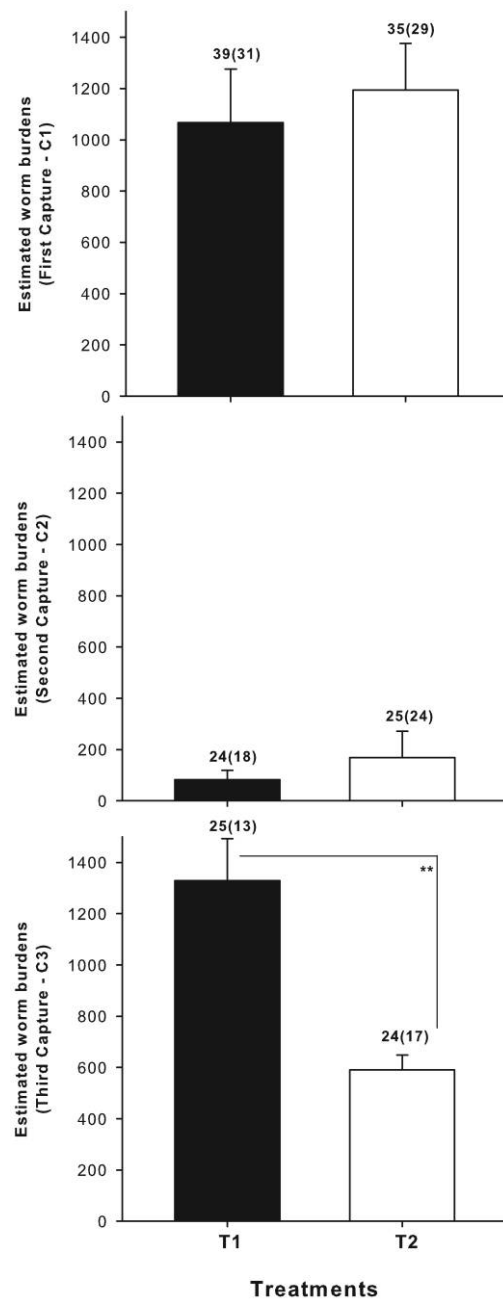


Figure 2: Mean (\pm SE) *Trichostrongylus tenuis* parasite abundance (worms per host) in spring (C1), a month later (C2), and 6 months later (autumn; C3) in relation to the parasite pair treatments. T1 = focal individual dosed + mate not dosed (black bars). T2 = focal individual dosed + mate dosed (white bars). Lines show only the significant differences between treatment groups (two asterisks = P value between .01 and .001). Numbers above bars denote the number of birds captured and, in the parentheses, number of birds from which we collected fecal samples.

reproduction (time) is perhaps the most common social interaction in the animal kingdom. The pattern that we found should encourage new studies aimed at exploring the mechanism that drives it. Because *T. tenuis* is a parasite that clearly reduces reproduction and survival (Hudson 1986a), together with inflicting a plethora of different physiological effects (Hudson 1986b; Redpath 2004; Martínez-Padilla et al. 2007; Mougeot et al. 2009, 2010a, 2010b), we suggest that mating with a parasitized partner may imply a cost that should be taken into account for a better understanding of the selection processes that leads to the avoidance of parasitized mates. Within an evolutionary perspective, our results suggest that mate choice should favor selection of unparasitized mates.

In our experiment, we also explored possible asymmetries in the directionality of parasite uptake within pairs. Since Darwin (1871), it has been traditionally assumed that female mate preference may be the selective force explaining extravagant ornamentation in males. Because of the biased directionality in mate choice studies toward evidence for choosy females (Hamilton and Zuk 1982; Folstad and Karter 1992), it might be expected that females are likely to pay a greater cost for choosing a wrong mate. However, we found that parasite-free males suffered an increased risk of parasite infection when mated with a naturally infected female, but the opposite pattern was not apparent, although sample sizes were small. Our results suggest that the costs of parasitism exist for males and appear greater for them than for females. This is consistent with the idea that mate choice in particular and sexual selection in general might not always be as female biased as has been suggested and might instead, at least in some species, be more reciprocal than has been suspected (Thrall et al. 1997; Prudic et al. 2011). This makes room for sexual selection to operate on male mate choice and the evolution of phenotypic traits of females that indicate their quality (Edward and Chapman 2011). The key factors in the evolution of male mate choice are paternal care, male investment in mating effort, and variation in female quality (Edward and Chapman 2011). Male red grouse provide paternal care from hatching to offspring dispersal, and their help is necessary for breeding successfully (Watson and Moss 2008). This paternal care can even be extended beyond the breeding period, because males also facilitate the territory establishment of their sons (MacColl et al. 2008; Watson and Moss 2008). The expression of testosterone-dependent ornaments in male red grouse (e.g., comb size) can represent an index of mating investment. Males with bigger combs increase their chances to mate and are more likely to mate with more than one female (Redpath et al. 2006a). Also, sperm limitation can be viewed as a form of mating investment and might also occur in red grouse, because mating status seems to con-

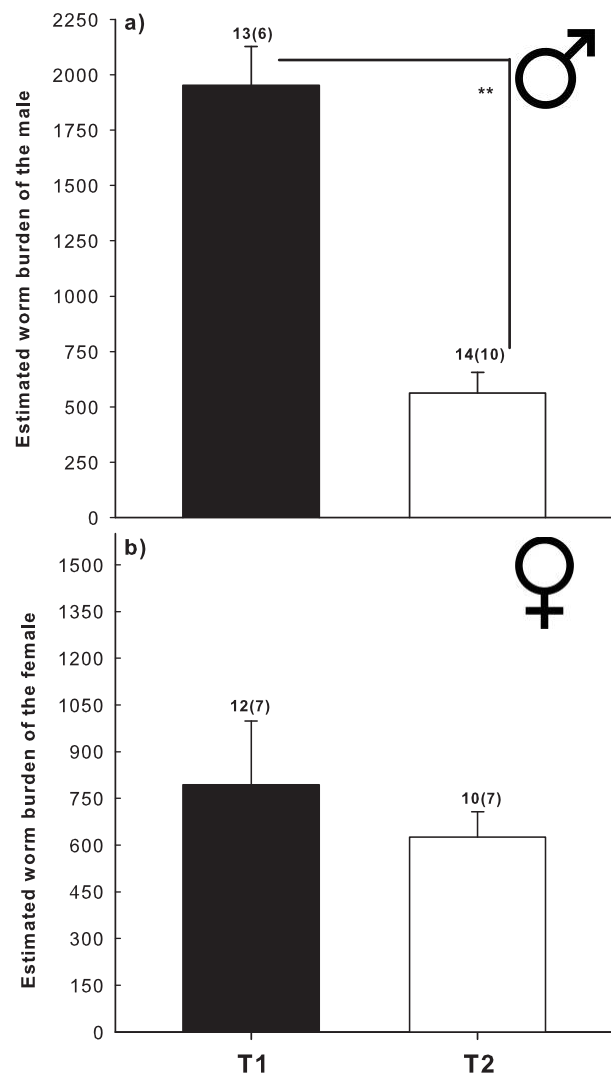


Figure 3: Mean (\pm SE) *Trichostrongylus tenuis* parasite abundance (worms per host) at the end of the experiment (C3) for males (a) and females (b) in relation to the parasite pair treatments. T1 = focal individual dosed + mate not dosed (black bars). T2 = focal individual dosed + mate dosed (white bars). Lines show only the significant differences between treatment groups (two asterisks = P value between .01 and .001). Numbers above bars denote the number of birds captured and, in the parentheses, number of birds from which we collected fecal samples.

strain sperm volume (J. Martínez-Padilla, P. Vergara, F. Mougeot, and S. M. Redpath, unpublished data). Finally, there is high variation in female quality, as has been reported previously (Martínez-Padilla et al. 2011; Vergara et al. 2011). Therefore, male mate choice may operate in red grouse (Martínez-Padilla et al. 2011), and there is some empirical evidence that female ornaments (e.g., comb size) advertise parasite load and body condition (Martínez-

Padilla et al. 2011; Vergara et al. 2011), which provides cues that males could use to evaluate the parasite infection level of a potential female mate.

The directionality of parasitism risk may also have implications for the immunocompetence handicap hypothesis (Folstad and Karter 1992), which was initially developed for males. This hypothesis predicts that testosterone and parasites act in opposite directions, such that testosterone increases the susceptibility to parasitism, and parasites constrain testosterone-dependent behavior. If males mate with highly parasitized females, this extra cost may alter the resolution of the trade-off between investing in testosterone-dependent behavior and resisting parasites in a wide variety of species. Considering parasite levels of female mates might influence how males may resolve the trade-off between aggressiveness and the risk of parasitism. Thus, parasite levels of female mates should be considered for a better comprehension of the effects of the immunocompetence handicap principle.

Our results have implications for our understanding of what causes the cyclic fluctuation seen in red grouse abundance. Two main hypotheses explain these population cycles. The “parasite hypothesis” invokes a destabilizing effect of *T. tenuis* parasites on female fecundity and recruitment (Hudson 1986a; Hudson et al. 1998; see also Redpath et al. 2006b). The “territorial behavior” hypothesis invokes a destabilizing effect of male aggressiveness on territory establishment of young males (Mougeot et al. 2003). Lower female fecundity and increased male aggressiveness caused by parasites and testosterone respectively were suggested as the mechanisms of these two independent hypotheses. From an intrasexual view point, we know that there are also important interactions between the two hypothesis in males; parasite levels influence aggressive behavior and vice versa (Fox and Hudson 2001; Mougeot et al. 2005a; Seivwright et al. 2005). Here, we have provided an important missing link between the two hypotheses, because they are connected from an intersexual point of view. We now know that pairing with a mate that has more parasites increases an individual’s parasite abundance. This appears to be particularly the case for

parasite transmission from females to males. The role of the sexes in driving cyclic dynamics has been little explored, yet our experiment suggests that additional research is needed from two different, but complementary, perspectives. On one hand, males should be considered in the parasite hypothesis, because their parasite abundance can be directly affected by the parasite abundance of their mates. On the other hand, males that mate with highly parasitized females may amplify parasite abundance in the population and alter male aggressive behavior. In both cases, our results highlight the need to consider the role of females and the risk of parasitism within mated pairs in grouse population dynamics.

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APPENDIX A

Chronogram of the Experiment and Parasite Life Cycle

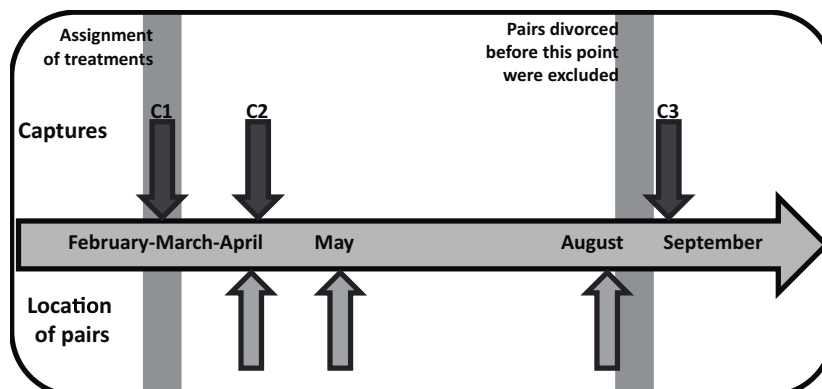


Figure A1: Chronogram of the experiment.

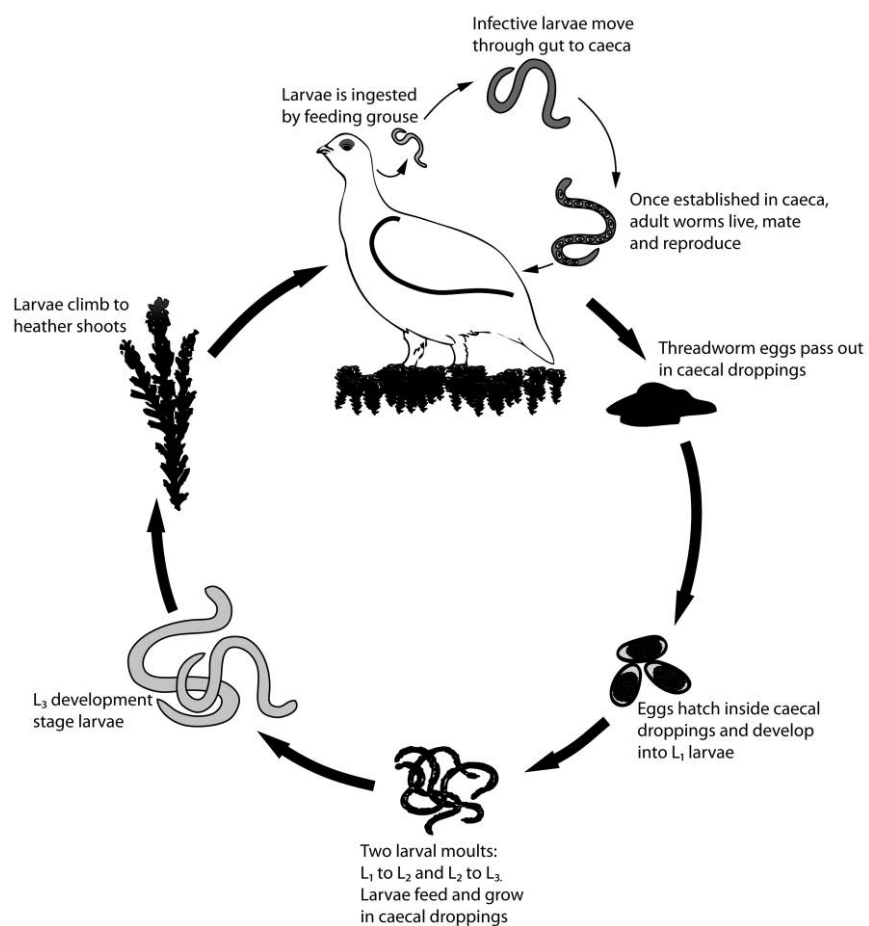


Figure A2: Life cycle of the cecal threadworm *Trichostrongylus tenuis*.

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