

RESEARCH ARTICLE

Subtle short-term physiological costs of an experimental augmentation of fleas in wild Columbian ground squirrels

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ABSTRACT

Parasites affect many aspects of host physiology and behavior, and thus are generally thought to negatively impact host fitness. However, changes in form of short-term parasite effects on host physiological markers have generally been overlooked in favor of fitness measures. Here, we studied flea (*Oropsylla idahoensis* and *Oropsylla opisocroistis tuberculata*) parasitism on a natural population of Columbian ground squirrels (*Urocitellus columbianus*) in Sheep River Provincial Park, AB, Canada. Fleas were experimentally added to adult female *U. columbianus* at physiologically demanding times, including birth, lactation and weaning of their young. The body mass of adult females, as well as their oxidative stress and immunity were recorded multiple times over the active season under flea-augmented and control conditions. We also measured the prevalence of an internal parasite (*Trypanosoma otospermophilii*). Doubly labeled water (DLW) was intraperitoneally injected at peak lactation to examine energy expenditure. Effects of parasites on oxidative stress were only observed after offspring were weaned. There was no direct effect of experimentally heightened flea prevalence on energy use. A short-term 24 h mass loss (−17 g) was detected briefly after parasite addition, likely due to *U. columbianus* preferentially allocating time for grooming. Our parasite augmentation did not strongly affect hosts and suggested that short-term physiological effects were unlikely to culminate in long-term fitness consequences. Columbian ground squirrels appear to rapidly manage parasite costs, probably through grooming.

KEY WORDS: Parasitism, Immune cost, *Urocitellus columbianus*, Host–parasite, Energy expenditure, Oxidative stress

INTRODUCTION

The resources that parasites extract from their hosts are often thought to produce negative effects on host fitness (Møller et al., 1994; Delahay et al., 1995; Careau et al., 2010). Parasitism can induce direct costs through sapping resources from their hosts (Nelson et al., 1975) and indirect costs through changes in behavioral activity (Giorgi et al., 2001; Scantlebury et al., 2007), acting as pathogen vectors (Smith et al., 2006), or modifying physiological tradeoffs (Bertrand et al., 2006b; Sorci et al., 2017). These host–parasite links are illustrated by eastern grey kangaroos selectively foraging away from better quality, but fecally contaminated grass

patches (Garnick et al., 2010) or male grey squirrels suffering from higher flea parasitization when upregulating testosterone levels (Scantlebury et al., 2010). As a result, parasites, when numerous, have the potential to generate a high resource toll on their hosts (Khokhlova et al., 2002; Krasnov et al., 2008). For example, when feral rock doves had lice levels experimentally increased, they steadily lost feather and body mass, resulting in compromised integument insulation (increased thermal conductance) and increased metabolic rate (Booth et al., 1993).

It is often difficult to distinguish direct resource loss to the parasite from the costs of anti-parasite defense or environmental effects (Bonneaud et al., 2003), such as effects of temperature extremes (Cohen et al., 2017) on energetics. Additionally, it is hard to discern whether parasites are the cause of poor host health and body condition or the result of it (Boonstra et al., 1980). Thus, it is useful to examine parasite effects through controlled experimental manipulation (Keymer and Read, 1991) to directly address these questions. Such research has been conducted in laboratory studies, but these have often failed to account for natural host–parasite dynamics such as the ‘80:20 rule’, an aggregated negative binomial distribution where a few hosts (20%) harbor the majority of parasites (80%) in a population (Galvani, 2003; Poulin, 2004; Craig et al., 2007). This underlines the parasite preference for hosts in terms of age, sex, condition and season (Dick and Patterson, 2007; Bize et al., 2008; Liberman et al., 2011), which are often overlooked and indicate the value of further field studies.


These factors, coupled with experimental studies that frequently focus on long-term fitness costs, may explain findings of muted parasite effects in wild studies (Khokhlova et al., 2002) compared with laboratory tests. An alternative research design might quantify more subtle short-term physiological modifications while preserving natural features of wild conditions. An emphasis of short-term effects on physiological changes serves two purposes. Firstly, short-term physiological shifts should be more detectable and directly quantifiable than multi-faceted fitness outcomes. Secondly, collection of these chronically sustained short-term effects may allow improved interpretation of potential long-term costs.

In this study, we experimentally tested the effects of ectoparasitic fleas (*Oropsylla idahoensis* and *Oropsylla opisocroistis tuberculata*) on a wild population of adult female Columbian ground squirrels (*Urocitellus columbianus*). We subjected a group of Columbian ground squirrels to an experimental increase in their natural flea loads and compared their physiological responses with those of a group of ground squirrels in which flea loads were left unchanged. Columbian ground squirrels are hibernating sciurid rodents with a 3–4 month annual active season, during which reproduction takes place (Dobson et al., 1992). Parental care is restricted to the highly territorial mothers during the 24 days of gestation, 27 days of lactation and a short period after weaning (Murie and Harris, 1982). This species is naturally parasitized by both ecto- (ticks, mites, botflies) and endo-parasites (helminths,

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coccidia, trypanosomes). The most visible of these are fleas (Raveh et al., 2011, 2015) that seem to follow the aggregated 80:20 distribution. As fleas are often localized to individual hosts and burrows, natural parasite dispersal is low, thus allowing enhanced isolation of parasite effects on hosts. As such, breeding female *U. columbianus* are an ideal model system to reveal parasite costs because of the lack of confounding factors such as parasite transfer or dispersal (Krasnov et al., 2003b; Hawlena et al., 2005).

Prior studies on parasite effects in *U. columbianus* have resulted in variable outcomes (little to no effect: Raveh et al., 2011, 2015; negative effect: Neuhaus, 2003). These studies applied experimental reductions of fleas, in a species that naturally has relatively low levels of infestation, to assess fitness consequences on individuals. Such detection of parasite effects may have been limited by the natural parasite distribution when using the approach of parasite removal in lightly infested populations. Parasite costs might only be relevant when present in resource-deficient hosts. Co-evolution of host–parasite interactions might be favored by natural selection when they minimize negative effects of the parasite on the host (Hinnebusch et al., 2017). In these cases, lowering the level of parasites is unlikely to have strong effects on fitness. Adding parasites to wild hosts provides an improvement over previous tests of ectoparasite effects reported in the literature (Booth et al., 1993; Warburton et al., 2016), because treated hosts should be more likely to show parasite consequences as a result of exacerbated costs.

Our approach to understanding host–parasite dynamics thus has two novel features: augmentation of fleas, which is more likely to reveal costs, and physiological measures that can expose such costs. Short-term parasite effects on physiological metrics were assessed during energetically constrained time points, such as lactation (Rogowitz, 1998; Naya et al., 2008), to augment the perceptibility of costs through a higher energy budget (Metcalf and Monaghan, 2013). In particular, we expected that *U. columbianus* would employ behavioral and immune defenses against flea-induced stress. As fleas can serve as a vector for pathogens (Durden and Hinkle, 2019), such as the blood parasite *Trypanosoma otospermophili* (Freedman, 1947; Lizundia et al., 2011), trypanosome levels might also increase in the flea-treated group. We expected higher trypanosome prevalence to lead to stimulation of nitric oxide (NO), which has been shown to rise in response to trypanosome infections (Vespa et al., 1994). By doing so, parasite infestation would be positively correlated with energy use (Kam et al., 2011) and a subsequently enhanced oxidative stress due to a non-specific innate immune response (Plumel et al., 2016; Bertrand et al., 2006a). A difference in the dynamics of mass, oxidative stress, immunity and energetic demand of heavily infested individuals would provide evidence supporting a short-term physiological consequence of parasite infestation in *U. columbianus*.

MATERIALS AND METHODS

Ethics statement

Animal care was carried out in accordance with Auburn University IACUC protocol no. 2018-3227 (with additional approval by the University of Calgary). Authorization for conducting research and collecting samples in Sheep River Provincial Park was obtained from Alberta Environment and Parks (research permit no. 58954) and Alberta Tourism, Parks, and Recreation (research and collection permit no. 18-448).

Population monitoring

Columbian ground squirrels, *U. columbianus* (Ord 1815), were studied at Sheep River Provincial Park, AB, Canada (Meadow B; 50°38′11.3″N, 114°39′56.7″W; 1550 m elevation) from April to

August 2018. The entire *U. columbianus* population at Meadow B has been continuously monitored since 1992 (Wiggett and Boag, 1986; Viblanc et al., 2010; Rubach et al., 2016), from the onset of emergence from hibernation in late April, to the end of offspring weaning in early July. Female *U. columbianus* have a short active season and a single reproductive period each year (Dobson et al., 1992). Each squirrel in this population is permanently identified with unique numbered fingerling ear tags (#1-Monel metal, National Band and Tag Company, Newport, KY, USA), and is given a unique hair dye marking (Clairol, Stamford, CT, USA) at the start of the season so it can be identified from a distance during field observations. We followed all reproductive females ($n=31$) to determine their mating day from behavioral observations and inspection of their genitalia (Murie and Harris, 1982). *Uroditellus columbianus* were trapped using Tomahawk live traps (13×13×40 cm; Tomahawk, Hazelhurst, WI, USA) baited with a small amount of peanut butter. Some of the females ($n=5$) either disappeared during the breeding season or were not re-captured and were thus excluded from analyses.

Experimental manipulation of flea load

We experimentally increased ectoparasite load on 16 females (treatment group) and compared them with 15 control females (control group; see below). At the start of the experiment, we ascertained *U. columbianus* body condition and then randomly assigned females of similar condition and age to control and treatment groups. Body condition was estimated by regressing individual body mass on zygomatic arch breadth (an index of structural size; Dobson, 1992). Fleas were collected from squirrels at a neighboring meadow less than 400 m away from the study site (50°38′19.7″N, 114°39′47.1″W) by brushing individuals with a fine-tooth flea comb (Four Paws, Hauppauge, NY, USA) into an aerated plastic container and transferring the fleas on the same day to experimental subjects. Because of the need for the same-day transfer to new hosts, fleas were not identified to species or sex, and were assumed to belong to one of the two common species found in prior studies (Hubbard, 1947; Hilton and Mahrt, 1971). Prior to flea addition, each squirrel was carefully combed on all sides of the body, including the head, to assess initial natural flea numbers. After counting, all initially present fleas were returned to their host.

An average of 20 fleas were added to each experimental subject at each time point, in addition to their inherent number of parasites (see Results). Fleas were added at three separate time points during the season: gestation (t_1), at lactation onset (t_2) and at peak lactation (t_3) prior to weaning. These time points were chosen because they represent important transitions in the breeding cycle and likely would incur elevated physiological demands. Additionally, they coincided with other manipulations of the long-term study, hence reducing animal handling and stress. We re-captured all non-hibernating females at a 4th time point (t_4 ; roughly a week before the onset of hibernation), but did not re-infest any animals, as a negative control. Fleas were deposited on the ventral side of the restrained animal and rubbed into the fur. We ensured all fleas had entered the animal's pelage before releasing it. The control group had their pelage rubbed in a similar manner to simulate flea addition, but with no change in number of natural fleas.

Trypanosomes

We assayed presence of *T. otospermophili* through collection of 100 μ l whole blood in capillary tubes. After collection, we centrifuged the capillary tubes at 5000 g for 10 min to apply quantitative buffy coat methodology. Centrifugation of whole blood

serves to concentrate trypanosomes in the buffy coat of the solute and enhance parasite detection (Sato et al., 2007); 5 μ l of the buffy coat was spread on a glass slide in a thin smear and Wright–Giesma stained (Shandon Kwik-Diff stain, Thermo Fisher Scientific, Waltham, MA, USA), followed by count estimates of trypanosomes.

Behavior

After flea addition, we released squirrels at the place of initial capture. We then visually observed control and treated squirrel behavior for 15 min to gauge how differentially parasitized squirrels allocated their time-budget to body maintenance. We recorded the number of seconds spent self-grooming.

Oxidative status and innate immunity

Individual oxidative stress levels and innate immunity were estimated during time points t_1 , t_2 , t_3 and t_4 . Blood (0.5 ml) was collected from the saphenous vein using a 27 G needle fitted to a 1 ml heparinized syringe. We kept blood on ice packs in a cooler box while in the field. After centrifugation (5000 g for 10 min), within 1–2 h of collection, plasma was separated and kept frozen at -20°C until the end of the field season, before transportation on dry ice and subsequent frozen storage at -80°C until laboratory analysis.

We assessed female oxidative status in plasma by global measures of oxidative damage (d-ROMs test; 8 μ l of plasma) and antioxidant defenses (OXY-absorbent test; 5 μ l of 1:100 diluted plasma) (Diacron International, Grosseto, Italy) (for details, see Costantini et al., 2011; Costantini et al., 2016; Viblanc et al., 2018). In addition, we measured NO in plasma (diazotization assay; 10 μ l plasma; for details, see Bourgeon et al., 2007) as a reflection of macrophage activation by intracellular pathogens (Playfair and Bancroft, 2004). Reactive oxygen metabolite (ROM) and antioxidant capacity (OXY) sample measurements were run in duplicate and NO was run once per sample. Intra-plate variation was 5.15% for ROMs and 12.1% for OXY. Inter-plate variation based on a standard sample repeated over plates was 2.74% for ROMs, 8.61% for OXY and 1.51% for NO.

Estimation of total daily energy expenditure (DEE)

Field protocol

DEE for treated and control females was determined only during peak lactation (day 25; t_3), when reproductive demands on females were expected to be the highest. DEE was estimated using the doubly labeled water (DLW) technique, as extensively described elsewhere (Kenagy et al., 1990; Rimbach et al., 2018), including in *U. columbianus* (Skibieli et al., 2013). Briefly, females were weighed (to the nearest 5 g using a spring scale; 1 kg, Pesola Ag, Baar, Switzerland) and a first blood draw (100 μ l) was collected from the saphenous vein using a 30 G needle in two 100 μ l non-heparinized capillary tubes to establish background levels of ^{18}O and ^2H . Capillaries were immediately sealed with a micro-jet flame and stored at room temperature until analysis (within 3 months). Squirrels were then injected intra-peritoneally with a premixed 5 g kg $^{-1}$ dose of DLW (10% H_2^{18}O and 99.9% $^2\text{H}_2\text{O}$, Cambridge Isotopic Laboratories, Cambridge, MA, USA). After injection, females ($n=26$) were held in traps in a quiet, shaded location and covered with a dark cotton pillowcase for 75 min to allow for isotope equilibration (Król and Speakman, 1999; Simmen et al., 2010; Skibieli et al., 2013). Following the equilibration period, another blood sample was drawn ($n=26$), fleas were added to the experimental animals, and the subjects were released. As part of the DLW test, subsequent blood samples and weight measurements were taken at 24 h and 72 h post-enrichment ($n=26$) to estimate isotope elimination

rates (Speakman and Racey, 1987). During the DLW experiment, we recorded the average ambient temperature (T_A) experienced by individuals to control for potential thermoregulatory effects on metabolic rate. We used thermo-logging iButtons (DS1921G, Maxim Integrated, San Jose, CA, USA), which recorded T_A with 15 min intervals over the course of the experiment. iButtons were centrally located in the colony, attached to the bases of elevated observation benches, with the iButtons 1 m above ground level.

Isotope analyses

Sealed capillary tubes were vacuum distilled for 10 min and the resulting water distillate analyzed by a continuous flow isotope ratio mass spectrometer (IRMS; IRMS-DELTA V PLUS, Thermo, Bremen, Germany) as described previously (Chery et al., 2015; Mahlerl et al., 2018). Distillates were pyrolyzed at 1400°C into H_2 and CO_2 gases in a glassy carbon tube under pure He flow at 90 ml min $^{-1}$. H_2 and CO_2 were separated at 110°C on a molecular sieve GC column before sequential analysis in IRMS. Results were normalized using the VSMOW2/SLAP2 international scale. In addition, memory-effect and drift-corrections were applied as needed. All analyses were performed in quadruplet and samples were re-analyzed if the standard deviation exceeded 2% for ^2H or 0.2% for ^{18}O in more than three out of the four analyses. We calculated the total body water (TBW) from the ^{18}O dilution space divided by 1.007 to correct for *in vivo* isotopic exchange (Racette et al., 1994). The mean \pm s.d. isotope dilution space ratio was 1.029 ± 0.016 . We calculated the CO_2 production rate from the single pool model as recommended for the body size of *U. columbianus* (Speakman et al., 1993; Speakman and Hambly, 2016). We converted CO_2 production to DEE using a modification of Weir's equation and an assumed food quotient of 0.85 based on the prior literature involving *U. columbianus* and DLW (Skibieli et al., 2013). For 6 animals, we observed either capillary leakage or incomplete DLW equilibration occurring within the standardized equilibration period, thus prompting their removal from subsequent analyses.

Statistical analysis

All statistics were done in R 3.5.1 (R Core Team 2018; <https://www.R-project.org>). We proceeded in a 3-step analysis. First, we assessed the efficiency of our treatment by comparing the initial and final parasite loads of our control and treated individuals at each time point of infestation. For this, we used either linear mixed models or generalized linear mixed effects models (LMMs and GLMMs; lme4 package in R; Bates et al., 2015) with initial or final (initial+additional fleas) parasite counts entered as the dependent variable, time point (t_1 to t_4), treatment group (control or treatment) and their interaction entered as independent variables, and female ID as a random factor to account for longitudinal data collection. For initial flea levels, we used a Poisson distribution as is appropriate when working with count data and given the distribution of initial flea loads. For final flea loads, the addition of ca. 20 fleas per squirrel in the treated group normalized the distribution of residuals in our model. Second, we investigated changes in body mass, oxidative status (ROM and OXY) and innate immunity (NO and trypanosomes) over the season using a similar procedure. Mass, ROM, OXY, NO or trypanosome levels were entered as dependent variables in separate LMMs and we tested for the fixed effects of time point, treatment and their interaction. In those models, we added female ID and age as random factors to account for repeated observations and potential age effects on physiological variables. The number of observations (n) and corresponding number of

individuals (N) are indicated for each model. Because of repeated observations on individuals, $n > N$. Finally, we analyzed the effects of our treatment on female energy expenditure during peak lactation, the period of highest reproductive demand. We compared treatment and control groups in terms of body mass, DEE, oxidative status and immunity using linear models (LMs). We included female age, litter mass at weaning (reproductive investment) and T_A (average temperature between release of the individual after DLW injection and collection of last blood sample) as covariates in the model to test for their effects on DEE. For all analyses, we visually inspected the distribution of model residuals using $Q-Q$ plots to ensure a reasonable fit to normality. Mean differences between groups (time points or treatment) were assessed using *post hoc* Tukey's HSD test ('glht' package in R; Hothorn et al., 2008). For each model, we calculated effect sizes (Cohen's d) and their 95% confidence intervals ('effsize' package in R; <https://CRAN.R-project.org/package=effsize>). We used benchmarks $d=0.2$, 0.5, 0.8 to indicate small, medium and large effect sizes, respectively (Nakagawa and Cuthill, 2007). The α -level was set at 0.05 for all statistical analyses and results are presented as means \pm 1 s.e.m. unless stated otherwise.

RESULTS

Changes in parasite load and individual condition over the experiment

Fleas

Over the course of the experiment, treated individuals were infested at three different time points with a mean of 19.59 ± 0.71 fleas per animal on top of the originally present 0.93 ± 0.24 fleas. In contrast, control animals had on average 0.61 ± 0.16 fleas per animal at each time point (Fig. 1B). Thus, after treatment, treated individuals averaged 33.63 times the flea load of the control individuals (LMM; $t=22.94$, $P < 2e^{-16}$, $n=90$ observations, $N=31$ individuals). The number of fleas in the treated individuals had returned to a level near the initial level by the next experimental infestation (Fig. 1A).

Mass

We did not observe differential mass changes between groups (Table 1) from t_1 to t_4 . During the DLW experiment, at t_3 , initial mass was not significantly different between control and treatment groups prior to flea application (Table 1): treated *U. columbianus* experienced a short-term (24 h) mass loss compared with control animals but this disparity was negligible at 72 h post-flea application. As a result of incomplete DLW equilibration, some individuals ($n=5$) were removed from mass analyses.

Behavior

After controlling for age, treated *U. columbianus* responded to flea infestation by increasing their self-grooming behavior by 73.31% compared with controls (Fig. 2; t_1-t_3 20.45 ± 3.41 s, control 11.8 ± 2.18 s; LMM; $t=2.22$, $P=0.03$, $n=89$, $N=31$). Grooming decreased for both groups at peak lactation (t_3).

Trypanosomes

Trypanosome prevalence steadily decreased over the season in both treatment and control groups (Fig. 3A). At t_4 , we observed a statistically insignificant increase in average count of trypanosomes (per 5 μ l of buffy coat) in the control group (329.82 ± 272.43 ; LM; $t=1.789$, $P=0.09$, $n=23$, $N=23$) compared with the treatment group (25.46 ± 19.14). Upon removal of an extreme outlier, there was no appreciable difference between treatment and control groups (Fig. 3A; control 62.8 ± 59.72 ; LM; $t=-0.66$, $P=0.52$, $n=22$, $N=22$).

Total DEE at peak lactation (t_3)

During peak lactation, treated individuals did not show significantly higher DEE than controls (Fig. 4; LM; $t=0.31$, $P=0.76$, $n=20$, $N=20$), even when we accounted for differences in fat-free mass. DEE significantly increased with age (Fig. 4; LM; $t=2.48$, $P=0.02$, $n=20$, $N=20$). Older breeders did not have larger litter masses at weaning (LM; $t=0.84$, $P=0.41$, $n=28$, $N=28$), nor did litter mass

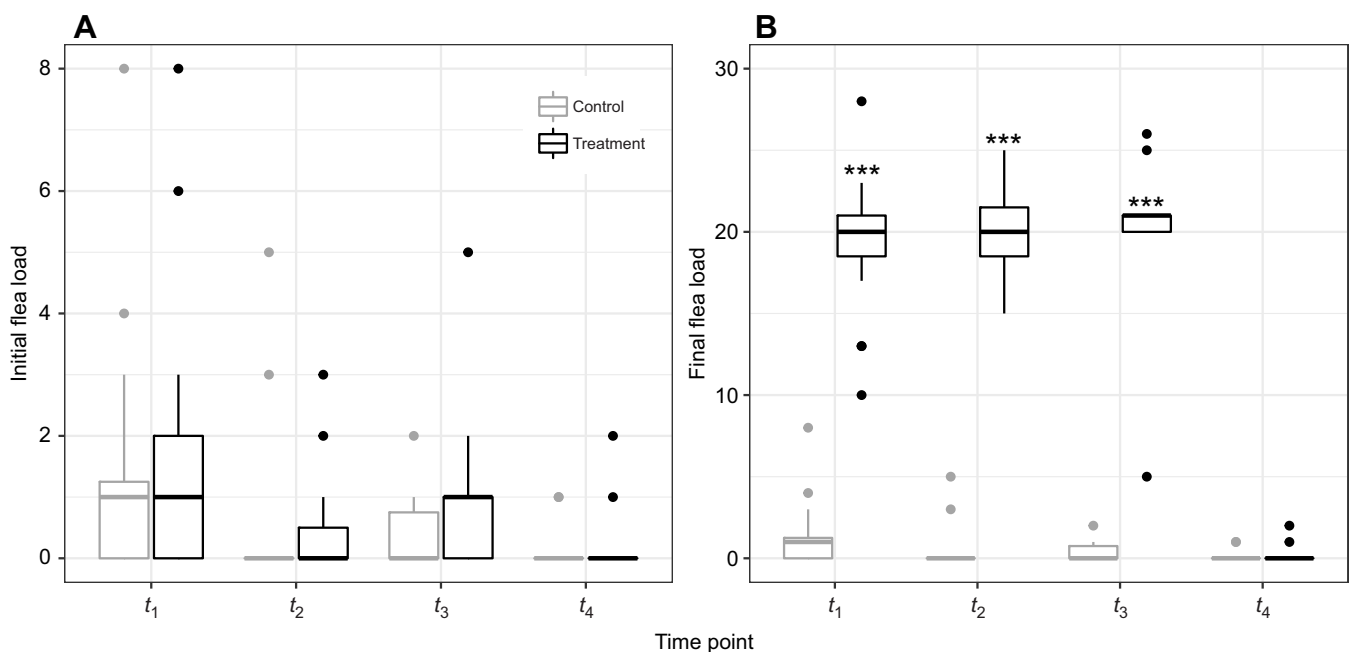


Fig. 1. Flea infestation in Columbian ground squirrels between April and August 2018. (A) Initial and (B) final flea numbers. Time points: t_1 , gestation (16 control, 15 treatment); t_2 , early lactation (14 control, 14 treatment); t_3 , peak lactation (14 control, 15 treatment); t_4 , prior to hibernation immergence (11 control, 13 treatment). Data are medians \pm s.e.m.; asterisks indicate significant differences between groups ($***P \leq 0.001$).

Table 1. Seasonal data

Variable	Control	Treatment	Cohen's <i>d</i>
Age (years)	4.07±0.40	4.13±0.43	0.03 (−0.35, 0.4)
Litter size	1.714±0.37	1.64±0.33	−0.04 (−0.41, 0.34)
Litter mass	179.21±36.99	195.87±40.96	−0.13 (−0.5, 0.24)
Temperature (°C)	13.94±0.29	13.67±0.28	0.27 (−0.54, 1.08)
<i>t</i> ₃			
Mass change 24 h (g)	4.5±6.6	−17.00±6.67	1.02 (0.02, 2.02)
Mass change 72 h (g)	3.5±8.5	−2.22±8.38	0.22 (−0.75, 1.19)
Mass			
<i>t</i> ₁ (g)	552.81±11.36	563.33±8.36	−0.27 (−1.00, 0.47)
<i>t</i> ₂ (g)	538.21±15.61	540.00±16.28	−0.03 (−0.81, 0.75)
<i>t</i> ₃ (g)	529.64±14.21	546.13±19.23	−0.28 (−1.05, 0.48)
<i>t</i> ₄ (g)	545.45±19.93	550.38±19.61	−0.07 (−0.92, 0.78)

*t*₁, gestation (16 control, 15 treatment); *t*₂, early lactation (14 control, 14 treatment); *t*₃, peak lactation (14 control, 15 treatment); *t*₄, prior to hibernation emergence (11 control, 13 treatment). Data are means±95% confidence interval. Cohen's *d* is given (with 95% confidence intervals), with 0.2 being a small effect size, 0.5 a medium effect size and 0.8 a large effect size. Confidence intervals including 0 indicate statistical non-significance.

differ substantially between treated and control individuals (Table 1; LM; $t=0.3$, $P=0.77$, $n=28$, $N=28$). We did not observe a relationship between DEE and ROM (Fig. 5; LM; $t=0.64$, $P=0.54$, $n=20$, $N=20$). A similar analysis on antioxidant defenses (OXY) revealed a relationship of decreased OXY levels (LM; $t=-2.26$, $P=0.04$, $n=20$, $N=20$) with increased DEE in treatment group females. Upon removal of a single outlier, this relationship disappeared (LM; $t=-0.61$, $P=0.55$, $n=19$, $N=19$).

Changes in innate immunity and oxidative status

Innate immunity (NO levels)

The level of inflammation, assayed through NO concentration, was similar in the control and treatment groups and over time (Fig. 3B; LMM with Tukey's *post hoc* test; $t=0.25$, $P=0.63$, $n=69$, $N=29$). NO concentration was not associated with ROM (LMM; $t=0.07$,

$P=0.95$, $n=69$, $N=29$), OXY (LMM; $t=1.17$, $P=0.25$) or trypanosome (LMM; $t=0.893$, $P=0.38$) levels.

Oxidative stress

There was a significant effect of time period on ROM and OXY levels (Fig. 6; ROM $z=-5.08$, $P<0.001$, $n=107$, $N=31$; OXY $z=-3.26$, $P<0.001$, $n=108$, $N=31$). ROM levels decreased from *t*₁ to *t*₃, but slightly rebounded at *t*₄ in the treated group (LMM; Tukey's *post hoc* test; $t=-2.5$, $P<0.02$, $n=54$, $N=15$) compared with the control group. OXY levels steadily decreased in both groups as the season progressed. As for other metrics, ROM (LM; $t=-0.42$, $P=0.68$, $n=69$, $n=24$, $N=24$) and OXY (LM; $t=-0.36$, $P=0.72$) values were not significantly different between the treatment and control groups at peak lactation (*t*₃).

DISCUSSION

Our experimental transformation of an aggregated parasite distribution to a bimodal distribution by adding fleas to some ground squirrels was an attempt to discern short-term parasite effects. However, like other *U. columbianus* studies that removed parasites, our results indicated that these fleas did not significantly impact their hosts over the short term in regards to any of our physiological measures.

Flea augmentation and grooming

By experimentally adding fleas to *U. columbianus*, we expected a multitude of downstream physiological responses. This was partially fulfilled, at least in terms of the flea treatment temporarily enforcing a large short-term increase of parasites. This level of parasites was at the extreme high end of what is normally seen in adult female *U. columbianus* at this specific field site, but not outside the natural range of variation. *Urociellus columbianus*, especially males and juveniles, have the capacity to harbor and maintain numbers of fleas equivalent to or in excess of the treatment level under natural conditions (Raveh et al., 2015;

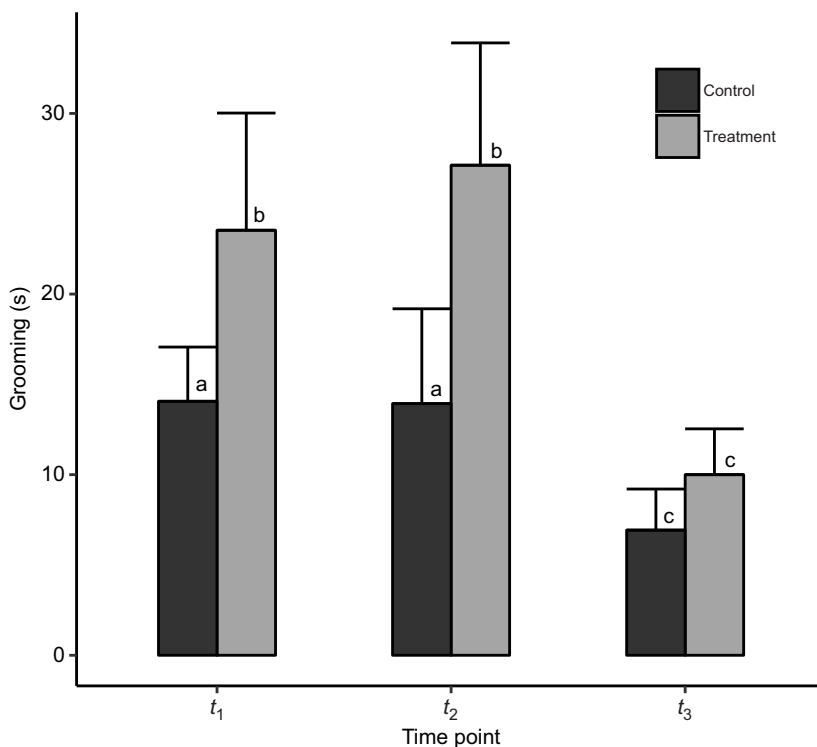


Fig. 2. Time spent grooming in Columbian ground squirrels between April and August 2018. *t*₁, gestation (16 control, 15 treatment); *t*₂, early lactation (14 control, 14 treatment); *t*₃, peak lactation (14 control, 15 treatment). Data are means±s.e.m. Different letters above boxes indicate significant differences between time points (Tukey HSD).

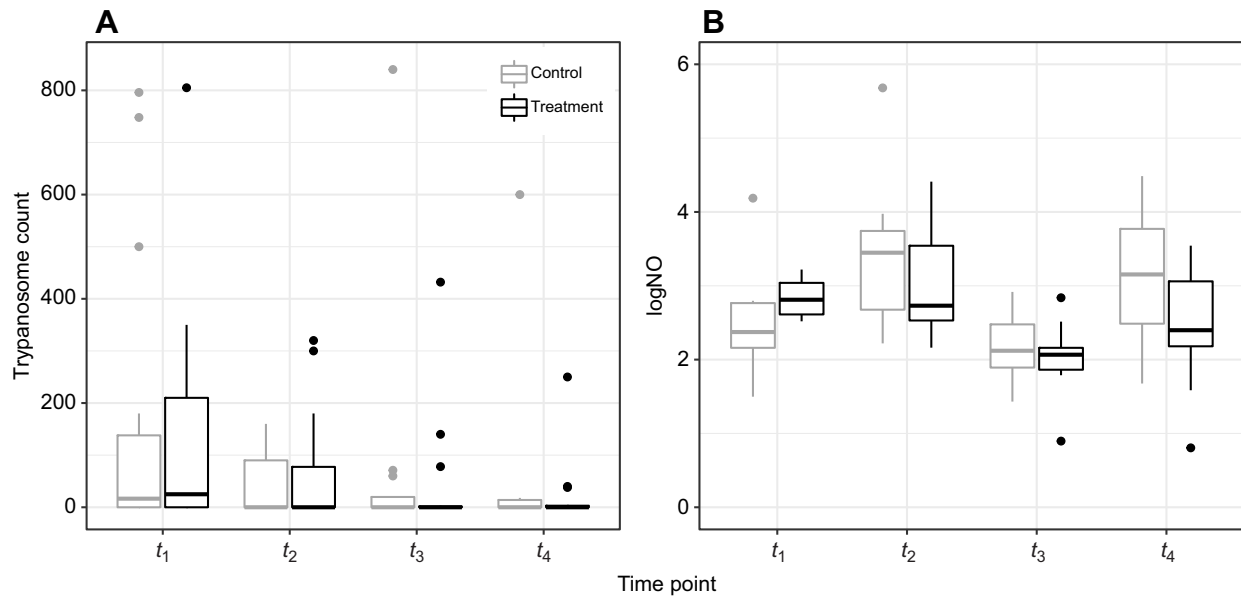


Fig. 3. Innate immunity of Columbian ground squirrels between April and August 2018. (A) Trypanosome count (per 5 μl of buffy coat) and (B) log nitric oxide concentration (NO , $\mu\text{mol l}^{-1}$). t_1 , gestation (16 control, 15 treatment); t_2 , early lactation (14 control, 14 treatment); t_3 , peak lactation (14 control, 15 treatment); t_4 , prior to hibernation emergence (11 control, 13 treatment). Data are medians \pm s.e.m.

J.D.R., F.S.D. and V.A.V., unpublished observations). Post-flea addition, treated *U. columbianus* allocated almost double the time of their non-parasitized counterparts to grooming. More importantly, they allocated energy that would normally be devoted to acquiring resources to maintenance of low flea levels at energetically and nutritionally demanding times. Within 24 h (J.D.R. unpublished observations of recaptured animals), almost all of the added fleas were removed, as evidenced by the equilibration of initial flea levels at each time point. This grooming time frame coincides with the 24 h mass shifts seen only in the treated group at the lactation peak time point (t_3). However, in the context of an acute high infestation, this statistically significant mass loss is hardly biologically costly given the return to normal mass within 72 h. As this population

behaviorally enforces low parasite levels, a situation of prolonged high parasitization is improbable and thus is not of high consequence for most host individuals.

Surprisingly, even during peak lactation, a critical period of the year where female energy demands are typically highest in mammals (Oftedal, 1984; Speakman and McQueenie, 1996), we did not observe an effect of our treatment on female energy expenditure. This result indicates that these parasites do not impose a high energetic cost on their host or that the cost of managing

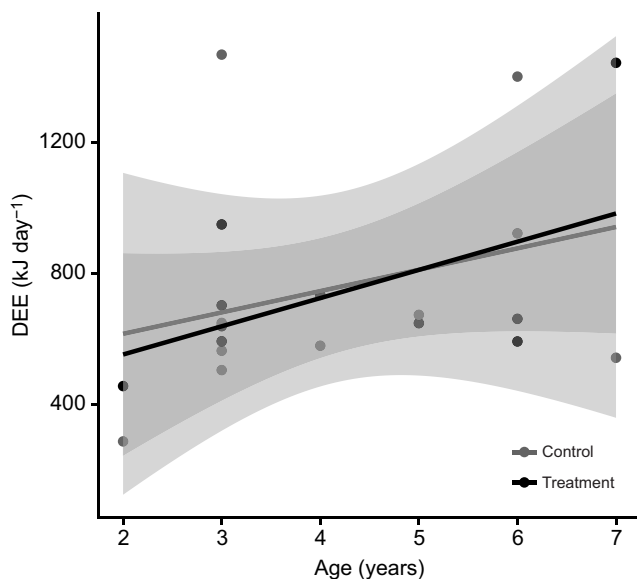


Fig. 4. Regression of age on daily energy expenditure (DEE). Mean data are shown with the 95% confidence interval (grey shading).

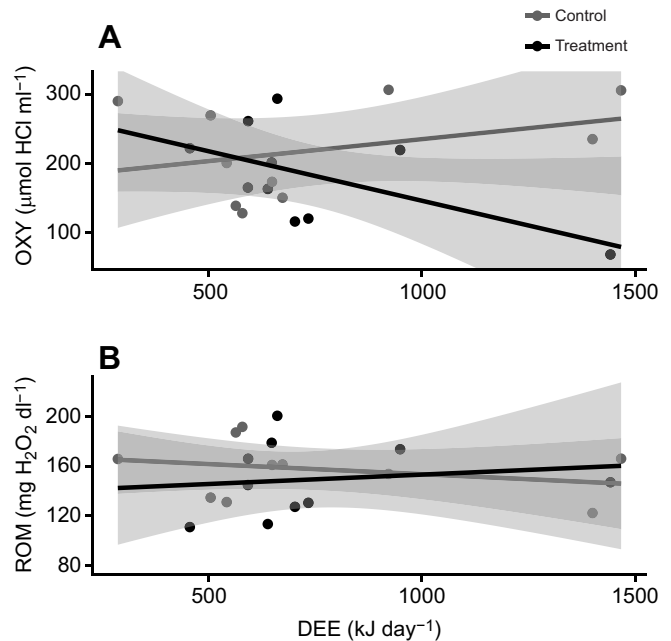


Fig. 5. Impact of DEE on oxidative status in Columbian ground squirrels. (A) Antioxidant capacity (OXY) and (B) reactive oxygen metabolite (ROM) concentration in plasma. Mean data are shown with the 95% confidence interval (grey shading).

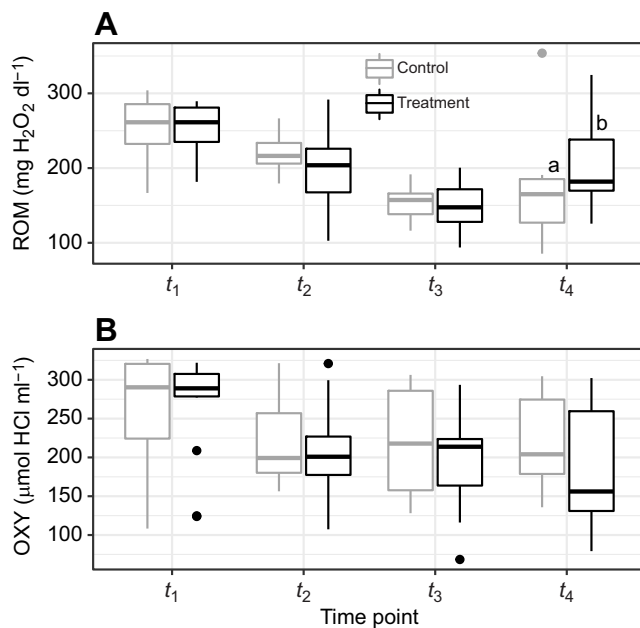


Fig. 6. Oxidative status in Columbian ground squirrels between April and August 2018. (A) ROM concentration and (B) OXY concentration. *t*₁, gestation (16 control, 15 treatment); *t*₂, early lactation (14 control, 14 treatment); *t*₃, peak lactation (14 control, 15 treatment); *t*₄, prior to hibernation emergence (11 control, 13 treatment). Data are medians ± s.e.m. Different letters above boxes indicate significant differences between time points (Tukey HSD).

parasites (i.e. grooming) is compensated through other pathways. Indeed, the time invested in flea removal likely accounts for the loss in body mass through changes in potential energy intake. With both species of fleas being ground squirrel specialists (Hubbard, 1947; Hilton and Mahrt, 1971) and a lack of alternative hosts in the area, it makes sense that the fleas have muted effects on *U. columbianus*. If parasites have co-evolved to specific hosts, they are more likely to deliver less irritable bites and introduce saliva that does not elicit an immune or behavioral response from the host (Dick and Patterson, 2007). Unfortunately, our observations do not account for a sex-biased effect whereupon fleas of a particular sex differentially induce stress (Hawlena et al., 2005; Krasnov et al., 2008). As fleas were collected and assigned randomly to treated individuals, we can assume that any sex bias in the parasites averaged out when the treated and control groups were compared. However, whether potential sex ratio bias in flea populations may differently affect individuals remains to be tested in the future. Given that female fleas consume larger blood meals (Krasnov et al., 2003a), one might expect female-biased populations to have a larger impact on hosts than male-biased populations.

A multitude of studies in other systems have demonstrated that the complexity of parasite and host energetics obscures the quantification of parasite effects (Hicks et al., 2018; Careau et al., 2010). For example, cape ground squirrel (*Xerus inauris*) DEE was similarly unaffected when parasite levels were manipulated (Scantlebury et al., 2007). Rather than parasites, increasing age stimulated slightly higher DEE. Many attributes associated with body composition such as larger litters or heavier young (Adams, 2005) would logically result in increased maternal investment and thus higher DEE. For example, older female *U. columbianus* appear to undergo reproductive senescence and may require extra energy to succeed at breeding (Broussard et al., 2003; 2005). In addition, younger breeding females may exhibit reproductive inefficiencies

when breeding for the first time (Broussard et al., 2008; Rubach et al., 2016), perhaps resulting in younger animals having lower reproduction-associated DEE than older animals (e.g. through reduced milk production). As such, instead of parasites, age-related body composition and activity levels largely influence DEE (Klausen et al., 1997).

Oxidative stress and immunity

Given their relationship, it is natural that a lack of parasite effects on energy use culminated in a similar lack of treatment consequences on immunity and oxidative stress. However, given that NO is a key immune factor involved in the oxidative killing machinery of macrophages (Playfair and Bancroft, 2004) and has been implicated in fighting trypanosome infections (Magez et al., 2006; Gobert et al., 2000), it is somewhat surprising that plasma NO levels did not mirror trypanosome levels in our study. Trypanosome levels fluctuated equally in both groups over the season, but NO concentrations declined in both groups over the season, and NO concentrations of the lack of treatment effect is that this species of flea is simply an inefficient trypanosome vector (Eisen et al., 2009). This putative inefficiency, in addition to the rapid grooming response, affords only a short transmission window and a subsequent lack of NO response to trypanosomes. This absence indicates that, at least in *U. columbianus*, trypanosomes are of little consequence or are at least not managed by NO in macrophages.

In contrast, oxidative stress patterns were more responsive to change over time in that they mirrored prior studies (Viblanco et al., 2018), likely because of the establishment of an oxidative shield early on during lactation (Blount et al., 2016; Vitikainen et al., 2016) that allowed costs to be offset. The oxidative shielding model proposes that mothers increase antioxidant defenses early in reproduction to prevent the transfer of damaged molecules to their offspring, which may occur as maternal oxidative stress increases through gestation and lactation (i.e. the oxidative cost of breeding). In our study, effects of parasites appeared to manifest in the post-shielding period (*t*₄), with treated individuals experiencing larger increases in ROM levels. This may reflect a poorer capacity of parasitized females to buffer reproduction-associated oxidative increases or a potential delayed effect of parasites. Generally, similar flea infestation studies have largely found no results of parasites on oxidative stress (Devevey et al., 2008; Maronde et al., 2018; Wegmann et al., 2015). That said, our finding of an interaction between parasitism and oxidative stress where others have found no relationship is not surprising given the multifaceted and non-linear relationship between the two (Costantini and Møller, 2009).

Conclusion

We attempted to quantify the previously variable costs of parasitism in *U. columbianus* by discriminating between short- and long-term effects (Asghar et al., 2015). However, it became clear that the tendency of *U. columbianus* to prioritize immediate grooming of parasites was likely the reason for the initial low level of fleas (Raveh et al., 2011, 2015) and the lack of seasonal effects of our experimental parasite manipulation. This strong grooming response coupled with oxidative shielding likely resulted in the dampening of any physiologically detectable parasite effects, even during the energetically demanding reproductive period. Some subtle short-term effects do manifest but are unlikely to culminate in detrimental long-term fitness consequences unless parasitemia is chronically sustained. As such, fleas, even when experimentally augmented in number to increase their impact, do not strongly affect these hosts.

This result suggests that while hosts in poor condition may exhibit high flea loads, flea infestation by itself is unlikely to debilitate hosts. Seasonal variance in parasite levels over many species coupled with larger processes (i.e. oxidative shielding t_1 – t_3) temporarily masking costs may result in studies overlooking parasite effects as a result of a short detection time frame. Given the current direction of climate change, it is eminently possible for parasite prevalence to increase (Cohen et al., 2017), and with it, the manifestation of these subtle costs. As such, short-term physiological measurements may be a better approach than long-term fitness estimates to detect parasite costs in wild populations.

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Competing interests

The authors declare no competing or financial interests.

Author contributions

Conceptualization: J.D.R., F.S.D., V.A.V.; Methodology: J.D.R., A.B., V.A.V.; Software: J.D.R.; Validation: J.D.R.; Formal analysis: J.D.R., A.Z., V.A.V.; Investigation: J.D.R., F.S.D., F.C., P.U., V.A.V.; Resources: J.D.R., F.S.D., F.C., A.Z., A.B., V.A.V.; Data curation: J.D.R.; Writing - original draft: J.D.R.; Writing - review & editing: J.D.R., F.S.D., F.C., A.B., V.A.V.; Visualization: J.D.R.; Supervision: J.D.R., V.A.V., F.S.D.; Project administration: J.D.R.; Funding acquisition: J.D.R., F.S.D., V.A.V.

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