

HOST STARVATION DECREASES PARASITE LOAD AND MEAN HOST SIZE IN EXPERIMENTAL POPULATIONS

KATJA PULKKINEN¹ AND DIETER EBERT²

Zoological Institute, University of Basel, Rheinsprung 9, 4051 Basel, Switzerland

Abstract. While host stress in vertebrate populations has often been linked to outbreaks of epidemics, which are attributed to the immuno-compromise of the stressed hosts, no predictions have been made about the response of invertebrate host populations to stressful conditions. Experiments conducted on individual invertebrate hosts, however, suggest that starved hosts may be a poor resource for parasites and that heavily infected old hosts may be more susceptible to stress, causing parasite populations to decline when their host population faces food shortages. In this epidemiological experiment, we exposed infected and uninfected *Daphnia magna* populations, which had been kept for many generations under a constant high food supply, to reduced food resources. Using the microsporidian gut parasite *Glugoides intestinalis*, which is exclusively horizontally transmitted, we tracked changes in parasite and host population size as well as host body length to elucidate how food shortages for the hosts influence host and parasite population dynamics. In both infected and uninfected populations, food shortage led to an approximately equal reduction in host density and changes in host body length distribution. Large hosts suffered a higher mortality than smaller hosts, which significantly reduced the mean body length in the host populations; however, this change was stronger in the infected populations and went hand-in-hand with a reduction in parasite spore load (a measure of intensity of infection) and prevalence. This effect disappeared after six weeks of food shortage, when the populations reached a new equilibrium. Our results indicate that in this system food stress impairs parasite spread and that host mortality is an important factor in regulating parasite abundance at the population level.

Key words: *Daphnia magna*; experimental epidemiology; *Glugoides intestinalis*; microsporidia; stress-induced immunosuppression; stress-induced mortality.

INTRODUCTION

Conventional wisdom holds that when living conditions deteriorate for hosts, for example during periods of overcrowding or food shortage, infectious diseases spread more rapidly, leading to an increase in parasite population size (Anderson and May 1981, Lloyd 1995). Laboratory experiments have established that malnutrition can lead to higher parasite loads in individual hosts (Wakelin 1989, Lloyd 1995). At the population level, however, this contention is based mostly on observations of vertebrate animals, where overcrowding and malnutrition are often coupled with high burdens of parasites and high mortality of infected animals (Anderson and May 1981, Gulland 1992, 1995). A plausible explanation for this observation is that stress impairs the defense mechanisms of the hosts and leads to an increase in parasite within-host growth and transmission. However, the size of a parasite population is

determined by a balance between the increase of parasites via transmission and within-host growth and the removal of parasites via parasite and host mortality (Anderson 1978, 1979a, Anderson and May 1978). If host stress has a stronger effect on parasite removal than on their growth and transmission, then the net effect of stress may be a decline in prevalence rather than an increase. This can occur when stress-induced mortality increases among infected hosts (Anderson 1979a). The balance of factors leading to an increase or a decrease of the parasite population size is likely to depend on the host–parasite system (Anderson and May 1979).

Research on vertebrates has shown that malnutrition can suppress immunocompetence, resulting in higher parasite loads due to increased within-host growth (see, e.g., reviews by Wakelin [1989] and Lloyd [1995]) and leading, in turn, to increased transmission rates. Invertebrates differ from vertebrates with regard to their immune system (e.g., they lack acquired immunity; Briggs et al. 1995) and their generation times and life spans are much shorter than those of vertebrates, and infections are usually chronic (i.e., there is no recovery from infections). Further, as a considerable proportion of host biomass is often converted into parasite biomass (e.g., parasitoids, baculoviruses), many parasites of invertebrates depend strongly on host biomass and re-

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¹ Present address: Department of Biological and Environmental Sciences, University of Jyväskylä, P.O. Box 35, 40014 University of Jyväskylä, Finland.

² Present address: Ecology and Evolution, Department of Biology, Fribourg University, Chemin du Musée 10, CH-1700 Fribourg, Switzerland.

sources. Therefore, host stress may have a stronger negative effect on parasite within-host growth and survival (via host deaths) and a more important role in determining the size of the parasite population in invertebrates. In the present study, we tested experimentally how resource limitation imposed on invertebrate host populations affects both host and parasite populations. Our prediction was that food stress would induce high mortality in heavily infected host individuals, thus removing a large part of the parasite population and leading to a net decrease in the parasite load in the host population.

As a model system we used the freshwater crustacean *Daphnia magna* and a horizontally transmitted microsporidian parasite that reproduces directly within the host. The *Daphnia*–microparasite system is especially suitable for studying epidemiological questions because it enables one to simultaneously manipulate a large number of replicate populations and to compare the effects of perturbation to the equilibrium conditions. In the absence of predators or parasites, the *Daphnia*'s mean population size is determined mainly by the rate of feeding (Slobodkin 1954). At constant food supply, the population settles to an equilibrium in which the total number of animals, reproductive rates, growth rates, and size-frequency distribution remain largely constant over time (Slobodkin 1954, Perrin et al. 1992). Epidemiological models predict that when a parasite persists in a host population under a certain parameter space, it regulates the host population at or around an equilibrium level (Anderson 1978, 1979b). Since *Daphnia* can be maintained in laboratory conditions such that they reproduce only asexually, it is possible to exclude effects of host genetics on disease dynamics and to concentrate purely on epidemiological questions. This corresponds to the *Daphnia*'s natural situation, as they reproduce mostly clonally during summer; in small water bodies, especially, the whole population may consist of only one or a few clones (Ebert et al. 2002). In many zooplankton communities, scarcity of food is considered to be a major controlling force (Lampert and Muck 1985), and natural populations of *Daphnia* frequently suffer from population crashes after food shortages (Tessier and Goulden 1982, Ghilarov 1985). Microsporidian parasites are the most common parasites of *Daphnia* in natural populations (Stirnadel and Ebert 1997, Ebert et al. 2001). Therefore the question of how nutrition influences parasite epidemiology is also relevant for real *Daphnia* populations.

In *Daphnia*, resistance to starvation has been shown to depend on the energy reserves and thus on size of the individual (Tessier and Goulden 1982, 1987, Tessier et al. 1983, Tessier and Consolatti 1989, Perrin et al. 1992). We therefore predicted that a food shortage in uninfected populations would increase the mortality of young, small *Daphnia*. However, the effect of parasites on host survival correlates positively with the intensity

of infection (Anderson 1979a, Anderson and May 1979, Ebert et al. 2000b). In infected *Daphnia* populations the largest animals have the highest parasite load due to their higher filtering rate and accumulation of parasites through time (Ebert 1994, Mangin et al. 1995, Stirnadel and Ebert 1997). Therefore, our second prediction was that, as opposed to uninfected populations, the large animals in infected populations would be among the first to die in a food shortage. The high mortality of large animals would lead to a decrease in the mean body length of the *Daphnia* and to a removal of a large part of the parasite population. Furthermore, it has been shown that the level of host nutrition affects parasite growth in *Daphnia*, so that parasites grow more slowly in hosts kept on low diet (Ebert et al. 2000b, Bittner et al. 2002). This would further inhibit parasite population from increasing in a starved host population. However, we predicted that when food deprivation continued for some time, the populations would reach a new equilibrium at the lower food level, with the population numbers remaining low, but the body length distribution returning to approximately prestarvation levels. In infected populations, this could either lead to an increase in the mean parasite load or, because of the decrease in the density of the host population, it could act in an opposite direction (Anderson and May 1979). Thus, the final outcome was not easily predictable. In this paper we present the results of an epidemiological experiment in which well-fed uninfected or infected *Daphnia* populations were subjected to a food shortage. The aim of the study was to understand the relative role of the forces that influence host and parasite population size and host body length structure and to use these population-level findings to expand our knowledge of host–parasite interactions as observed on the individual level.

METHODS

The host–parasite system

The model system we used in the experiment consisted of a freshwater crustacean, *Daphnia magna* Straus (Crustacea: Cladocera), and its obligatory parasitic microsporidian, *Glugoides intestinalis* (Microspora: Glugeidae; Larsson et al. 1996). *D. magna* is a cyclic parthenogen that can be maintained in the laboratory in clonally reproducing populations. Males and fertilized resting eggs are produced when conditions deteriorate (e.g., due to crowding). Under laboratory conditions, resting eggs usually do not hatch without dormancy, and thus the genetic composition of a monoclonal *Daphnia* population does not change during experimentation.

Due to the small size of *D. magna* (maximum length 4 mm), large populations can be maintained in small vessels in the laboratory, enabling us to maintain many populations under controlled conditions. The vessels are small enough to exclude spatial heterogeneity in

the populations; still the population numbers are large enough for disease dynamics to occur. With a constant food supply and without parasites, *Daphnia* populations reach an equilibrium population size, representing a carrying capacity determined by the food quantity (Slobodkin 1954, Perrin et al. 1992, Ebert et al. 2000a). At 20°C, the generation time of *D. magna* is between 8 and 16 days (depending on resources available).

G. intestinalis (Larsson et al. 1996) is an intracellular parasite in the gut epithelium of *Daphnia* that infects its host through waterborne spores. It reproduces directly, and transmission is horizontal between hosts (Ebert 1995). It best fits the definition of a microparasite used in epidemiological models (May and Anderson 1979). Hosts do not recover from infection (Ebert et al. 2000a). The parasite is transmitted from live hosts and dead hosts (Ebert 1995), but transmission through dead hosts is less likely (D. Ebert, *personal observation*). Reinfection of the same host with spores released with the feces due to ingestion during filter feeding is common (Ebert 1995). The number of parasite spores found inside the host is directly correlated with the parasite's transmission success (Ebert 1994). Exposure to larger doses (higher concentration of the waterborne spores) leads to higher parasite loads (Ebert and Mangin 1997).

The *Daphnia* clone used in the experiment originated from a pond in North Germany near Gaarzerfeld. It was collected in 1997 and maintained in the laboratory since. The *G. intestinalis* strain used in the experiment was isolated from this host clone and propagated in the same host genotype in the laboratory.

Population dynamics experiment

We cultured *Daphnia* populations in glass jars filled with 1 L of artificial freshwater medium (Klüttgen et al. 1994) that we modified by using only one-twentieth of SeO₂ concentration and adding one-fifth water from a local well. We fed the animals daily with suspension of the green algae *Scenedesmus gracilis* grown in chemostats. We kept the cultures in an air-conditioned room at 20°C with a 16:8 light:dark period and randomized the position of the jars every week.

We bred uninfected *Daphnia* in the laboratory and began the experiment by distributing them evenly into 40 populations of 250 animals each so that the body length distribution was as similar as possible among all populations. We randomly chose 20 populations to be infected with *G. intestinalis*. To each of these populations we added 20 infected adult females within a net enclosure for 1 wk. The mesh size of the net (250 µm) kept the infected and uninfected *Daphnia* from mixing, but allowed algae and parasite spores to pass through. The nets were shaken every day to maximize the transport of spores into the culture. To the 20 uninfected populations, we added uninfected *Daphnia* in the net enclosures.

We removed one-fourth of the medium together with the *Daphnia* from each population every week after mixing it in a Folsom plankton sample divider, and we replaced the removed water with fresh medium. We counted the animals removed every week and measured them for body length (from the top of the head to the base of the tail spine) every other week. All animals used to determine parasite spore load and for the experiment for time until death under complete starvation were from this removed portion. We kept all populations well fed (110×10^6 algae cells·jar⁻¹·d⁻¹). When approximate population size equilibrium was reached, we took a subsample of the animals from the weekly removed portion to determine spore load in the populations. These samples, and all later spore samples, were frozen for later counting. From this point onwards, we randomly assigned 10 of the 20 uninfected and 10 of the 20 infected populations to the food reduction treatment. Starved populations received one-eighth of the standard food ration (13.8×10^6 algae cells·jar⁻¹·d⁻¹). The other populations were fed as before. One week after the food reduction treatment began, we took a second sample to ascertain spore load. We continued the food reduction treatment until those populations settled to equilibrium at the lower food level. We then sampled the infected populations to determine the spore load for the third time.

Once per month we checked populations for the presence (infected populations) or absence (uninfected populations) of infection by dissecting five large females from each population and checking for spore clusters in the guts. All infected populations remained infected until the end of the experiment, although one was lost accidentally. We also discovered one of the uninfected populations to be infected on week 21 and therefore excluded it from all analyses.

Spore load dynamics

To determine mean parasite load, we took 10 animals per sampling from each population. The animals were taken in proportion to the size distribution of the animals in each population. The animals were measured for body length and placed singly in 50 µL of distilled water in micro test tubes and frozen for later spore counting. Spore counting was done by grinding up the whole *Daphnia* in the 50 µL of the water it was frozen in and counting the spores in a bacterial counting chamber (Thoma ruling). Because this method has a detection threshold of ~200 spores per animal, it is possible that some early infections in animals with only a few spore clusters were missed, perhaps leading to a slight underestimation of prevalence. However, since the heavily infected animals had more than 100 000 spores, spore load counts were not affected. As large animals contribute more volume to the total volume in which we counted the spores, we corrected the sample volume for this body length effect. We calculated the volume of each animal by converting their body length ac-

ording to a power equation that we obtained by weighing 96 *Daphnia magna* with a size range from 0.9 to 3.7 mm and assuming that the density of the *Daphnia* equals that of water. The power equation used was volume = $0.2418 \times \text{length}^{2.593}$.

Mortality during food reduction in the population experiment

In addition to population censuses, we tracked mortality in the populations directly by collecting the dead animals from all the populations daily during days 3 to 7 after the food reduction treatment was begun. We measured the length of the dead animals to see if the size distribution of dying animals differed according to treatments. Since the size frequency distribution was bimodal, we divided the animals into two size classes, 0.9–2.0 mm (small) and 2.1–3.7 mm (large), which corresponds approximately to nonreproducing and reproducing *Daphnia* size classes. In each population, we calculated the probability of death within both size classes for each day and calculated means separately for all treatments. We calculated the number alive each day by subtracting the cumulative number of dead animals from the number alive before food reduction. Since we calculated the probability of death starting from those alive on day zero, the values we obtained for each day are a slight underestimation of the real values, but the proportions of the probabilities between days are not affected.

Experiment for time until death under complete starvation

To test for size-specific mortality in our populations, we measured the time until death under complete starvation for different sizes of individual animals. We took five animals from each of the 40 populations on week 14 and placed them singly in 100-mL jars in artificial medium. The medium was filtered through a 0.2- μm pore size filter to remove any bacterial growth that might serve as food for the *Daphnia*. The medium was replaced every day. At the beginning of this starvation experiment, we measured the body length of the animals. We recorded the survival of the animals daily and ran the experiment until the death of the last experimental animal. We could not verify the infection status of the animals taken from infected populations, since in most cases they degraded too quickly after their death. However, from other parasite checks in our experimental populations we know that in an infected population most hosts carry the parasite. We calculated the mean time to death for both uninfected and infected animals in the two size classes (small and large).

Data analysis

The effect of food reduction treatment and infection on population counts and mean body lengths of *Daphnia* were analyzed with GLM repeated-measures ANOVA using log-transformed data. Week was included

as a within-subject factor and infection status and food reduction treatment as between-subject factors. In this analysis, a statistically significant food reduction effect in itself is not an indication of the effect of food reduction treatment; instead this is indicated only by a statistically significant interaction between food reduction and week after the food reduction treatment began in week 21. In these analyses, we included only measurements from week 15 onwards, because the marked fluctuations in population counts before this time were likely to be caused by the starting conditions. Since the circularity assumptions of the variance-covariance matrix (von Ende 1993) were not met, we used Huynh-Feldt corrected significance levels (Potvin and Lechowicz 1990). Among the four treatments, the skewness and kurtosis values of the *Daphnia* populations after food reduction treatment were compared with ANOVA on weeks 22 and 27 separately for each week. We used two parameters to measure the spore load in infected populations: mean spore load and prevalence. Of these two, prevalence is most closely related to transmission success in microparasites. We calculated the mean spore load of a population as the mean of the spore load of the 10 animals subsampled, including uninfected animals and prevalence as a proportion of infected animals in the subsample. The effect of the food reduction treatment on mean parasite load and prevalence of infection were tested with *t* tests between well-fed and starved populations separately for each sampling point. The probabilities of infection at a certain body length for individual *Daphnia* were compared between the starved and well-fed groups separately for each of the three sampling points with a logistic regression. The data on the probabilities of death during the 3–7 d after food reduction were not normally distributed, so we analyzed them using non-parametric Kruskal-Wallis tests. We compared the probability of death between well-fed and starved populations in the two size classes and between the size classes among well-fed and starved populations. We also compared uninfected and infected animals within each of the four groups. The mean time to death of the individual animals under complete starvation was tested between small and large animals separately for uninfected and infected animals with Mann-Whitney *U* tests, because the data were not normally distributed. We used Bonferroni correction to adjust the levels of statistical significance for multiple tests. All statistical analyses were done with SPSS 10.0 (SPSS 1999). In the figures included in this paper, the well-fed animals are always depicted with open symbols and the starved animals with filled symbols.

RESULTS

Population dynamics experiment

Population densities fluctuated around 200 individuals (Fig. 1A). At week 21, we considered the popu-

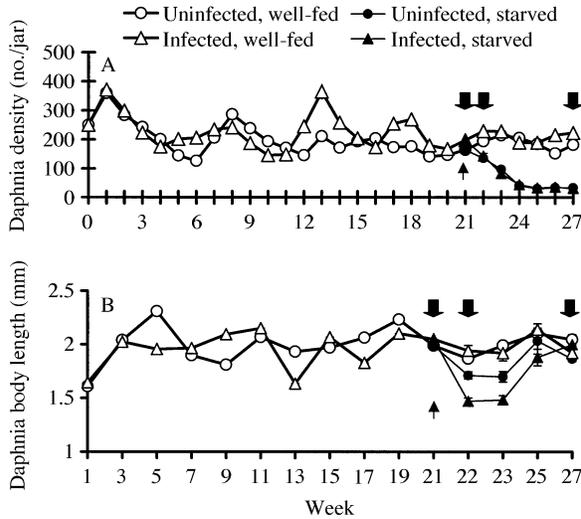


FIG. 1. (A) Population dynamics and (B) body lengths of *Daphnia magna* in uninfected populations and populations infected with *Glugoides intestinalis* (means \pm 1 SE) before (until week 21) and after food reduction treatment. The start of the food reduction treatment is indicated with an arrow from below, and the sampling for spore counts is indicated with block arrows. Note that the scales for week (x-axis) differ slightly.

lations to have reached equilibrium and took the first samples for spore load from the infected populations. Immediately after this, we assigned half of both uninfected and infected populations to the food reduction treatment (one-eighth of the normal food ration) and took a second sample 1 wk later (week 22). In the

populations that were continuously well fed, the densities kept fluctuating around 200 animals per jar, while the starved populations settled to around 20 animals. We took a third sample for determining spore load at week 27 (Fig. 1A). The interaction between food reduction and week was highly significant (repeated-measures ANOVA, within-subjects factors, week \times food reduction, $F = 112.9$, $df = 9.6$, 327.8 , $P < 0.001$), and contrasting between the levels of within-subjects factor (week) revealed that this interaction became significant only after the onset of the food reduction treatment at week 21. The effect of parasitism on the population size was also significant (between-subjects effects, $F = 20.0$, $df = 1, 34$, $P < 0.001$), but the infected populations contained slightly more animals.

During the two weeks after the food reduction treatment was begun (weeks 22 and 23), mean body length of *Daphnia* decreased in the starved populations as compared to the well-fed populations (Fig. 1B, Table 1, within-subjects contrasts). However, when the food reduction had continued for 4 wk (approximately two host generations; week 25), the mean body length returned to its prestarvation level (Fig. 1B, Table 1, within-subjects contrasts). The mean body length of *Daphnia* was also shorter in infected than in uninfected populations (Table 1, between-subjects effects). After food reduction was begun, during week 22, the decrease in mean body length was greater for infected than for uninfected populations (Fig. 1B) as indicated by the contrasting of the three-way interaction between week, food reduction, and infection (Table 1, within-subject contrasts).

TABLE 1. Repeated-measures ANOVA for the mean length of *Daphnia magna*.

Source	df	MS	F	P
Between subjects				
Infection (I)	1	0.035	18.662	<0.001
Food reduction (R)	1	0.124	65.624	<0.001
I \times R	1	0.002	0.927	0.342
Error	34	0.002		
Within subjects†				
Week (W)	4.780	1.077	27.173	<0.001
W \times I	4.780	0.167	4.203	0.002
W \times R	4.780	0.320	8.078	<0.001
W \times I \times R	4.780	0.117	2.943	0.016
Error	162.522	0.040		
Source				
Within-subjects contrasts	W	W \times I	W \times R	W \times I \times R
Week 17 vs. week 15	0.064	<0.001	0.293	0.627
Week 19 vs. previous	<0.001	0.189	0.415	0.732
Week 21 vs. previous	0.481	0.050	0.665	0.529
Week 22 vs. previous	<0.001	0.432	<0.001	<0.001
Week 23 vs. previous	<0.001	0.095	<0.001	0.239
Week 25 vs. previous	0.039	0.864	0.581	0.406
Week 27 vs. previous	0.967	0.139	0.161	0.002

Notes: The *P* values for the within-subjects contrasts are given for the factors having statistically significant within-subjects effects. The contrast is "difference," which compares each level to the mean of all previous levels.

† Huynh-Feldt corrected.

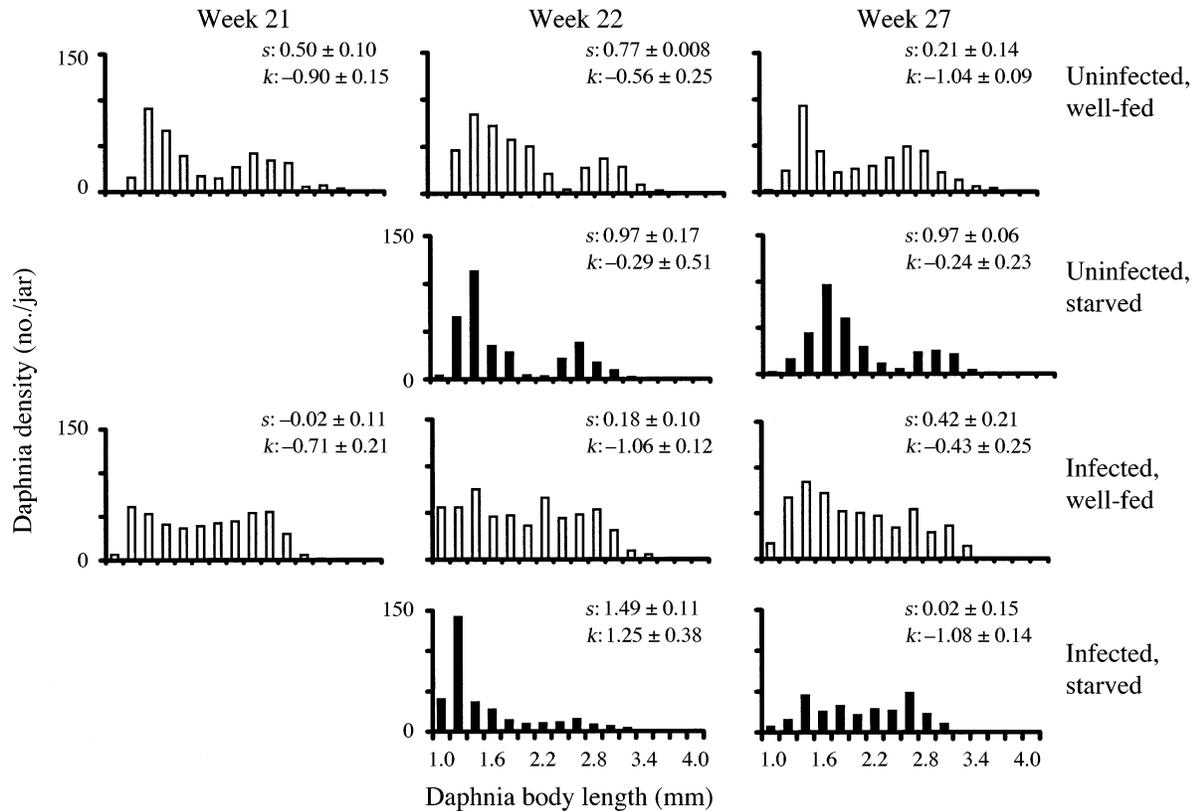


Fig. 2. Length distributions of uninfected *Daphnia magna* populations and populations infected with *Glugoides intestinalis* (sums of replicates) before food reduction treatment (week 21) and 1 and 6 wk (weeks 22 and 27, respectively) after the beginning of food reduction treatment. The numbers within figures show skewness (s) and kurtosis (k) values of the *Daphnia* populations at each time point (means \pm 1 SE). Positive skewness values indicate that populations are skewed to the right, i.e., most of the animals are smaller than the mean of the population. Negative kurtosis values indicate that the distribution is flatter than the normal distribution, and positive values show that the distribution is more peaked than the normal distribution.

In addition to a decrease in mean body length, the shape of the body length distribution of *Daphnias* changed under food reduction (Fig. 2). After 1 wk, the distributions in the starved populations became more positively skewed as compared to the well-fed populations, i.e., there were more small animals than large animals in the populations (Fig. 2; ANOVA, $F = 36.7$, $df = 1, 34$, $P < 0.005$, level of significance Bonferroni adjusted for two tests, $\alpha = 0.001/2$). The increase in skewness was significantly higher for the infected starved populations than for the uninfected starved populations (ANOVA, food reduction treatment \times infection status interaction, $F = 19.6$, $df = 1, 34$, $P < 0.005$). Kurtosis also increased in the starved populations after 1 wk as compared to the well-fed populations (Fig. 2; ANOVA, $F = 13.5$, $df = 1, 34$, $P < 0.005$). The change was much more marked in the infected starved populations (ANOVA, food reduction treatment \times infection status interaction, $F = 8.5$, $df = 1, 34$, $P < 0.025$). After 6 wk, the starved populations returned to the body length distribution before treatment (Fig. 2), showing in week 27 no difference in skewness or kurtosis from the well-fed populations

(ANOVA, $F = 1.4$, $df = 1, 34$, $P = 0.251$ for skewness and $F = 0.1$, $df = 1, 34$, $P = 0.721$ for kurtosis).

Spore load dynamics

Before food reduction (week 21) neither the mean spore load nor the prevalence (proportion of infected hosts) differed significantly in the infected populations later allocated to the food reduction treatment or not (Fig. 3; t test, $t = 1.6$, $df = 17$, $P = 0.139$ for mean spore load and $t = 1.9$, $df = 17$, $P = 0.072$ for mean prevalence). After 1 wk of food reduction (week 22), both the mean spore load and the prevalence decreased significantly in the starved populations compared to the well-fed populations (Fig. 3; t test, $t = 3.9$, $df = 11.2$, $P < 0.003$ for mean spore load and $t = 3.9$, $df = 17$, $P < 0.003$ for prevalence; levels of significance Bonferroni adjusted for three tests, $\alpha = 0.01/3$). In week 27, there was neither a significant difference in the mean spore load (t test, $t = 1.5$, $df = 17$, $P = 0.144$) nor in the prevalence ($t = 0.7$, $df = 17$, $P = 0.516$) between the starved and well-fed populations (Fig. 3).

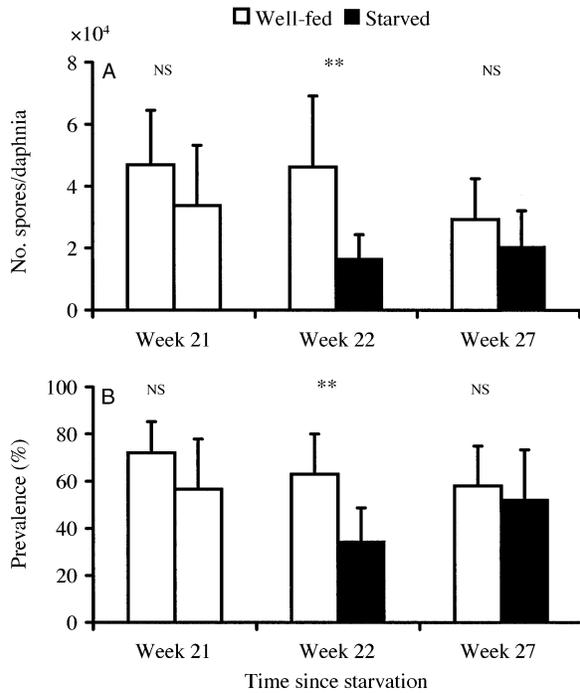


FIG. 3. (A) Spore load and (B) prevalence in well-fed and starved *Daphnia* populations before food reduction treatment (week 21) and 1 and 6 wk (weeks 22 and 27, respectively) after the food reduction treatment was begun (means + 1 SE). ** $P < 0.0033$ (levels of significance Bonferroni adjusted for three comparisons); NS, not statistically significant.

For individual animals, the probability of being infected increased with body length at all sampling dates (Fig. 4; change in the log-likelihood if term removed 168.4, $df = 1$, $P < 0.003$, odds ratio 1539.7 for week 21; 162.1, $df = 1$, $P < 0.003$, odds ratio 256.5 for week 22 and 172.8, $df = 1$, $P < 0.003$, odds ratio 829.9 for week 27; levels of significance Bonferroni adjusted for three tests, $\alpha = 0.01/3$). In weeks 21 and 22, the food reduction treatment did not influence the probability of infection, only body length did. In week 27, the effect of food reduction was also highly significant (change in the log-likelihood if term removed 12.6, $df = 1$, $P < 0.003$, odds ratio 0.1), indicating that individuals in the starved populations had a lower probability of being infected than those in well-fed populations.

Mortality during food reduction in the population experiment

The probability of death was significantly higher for the starved than for the well-fed populations in both size classes (Fig. 5, connected with horizontal braces; Kruskal-Wallis $\chi^2 = 19.6$, $df = 1$, $P < 0.00125$ for small animals; $\chi^2 = 24.3$, $df = 1$, $P < 0.00125$ for large animals; levels of significance Bonferroni adjusted for eight tests, $\alpha = 0.01/8$). In the well-fed populations the probability of death did not differ among size classes (Fig. 5, connected with vertical braces; χ^2

$= 0.4$, $df = 1$, $P = 0.545$), while it did in the starved populations (higher probability of death for the larger hosts; $\chi^2 = 12.0$, $df = 1$, $P < 0.00625$; $\alpha = 0.05/8$). As the larger animals carry the highest parasite load, this result is central to our study and can explain the selective removal of parasites under food reduction (compare Fig. 3).

For the small animals, the probability of death did not differ between uninfected and infected animals either in well-fed populations or in starved populations (Fig. 5A, B; $\chi^2 = 0.9$, $df = 1$, $P = 0.343$ for well-fed; $\chi^2 = 0.0$, $df = 1$, $P = 1.000$ for starved). However, large infected animals in both treatments had a significantly higher probability of death than the uninfected (Fig. 5C, well fed, $\chi^2 = 11.9$, $df = 1$, $P < 0.00625$; Fig. 5D, starved, $\chi^2 = 12.9$, $df = 1$, $P < 0.00125$).

Experiment of time until death under complete starvation

The mean time to death in complete starvation was 150.1 ± 58.4 h (mean \pm SD) for the uninfected *Daphnia*

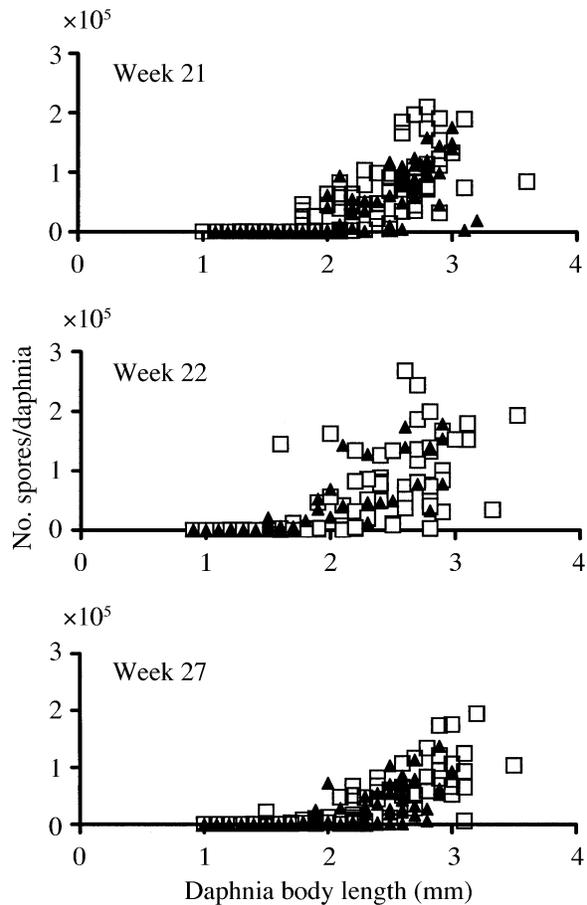


FIG. 4. The relationship between body length and spore load in individual *Daphnia* in well-fed populations (open squares) and in starved populations (closed triangles) before food reduction treatment (week 21) and 1 and 6 wk after (weeks 22 and 27, respectively) the food reduction treatment was begun.

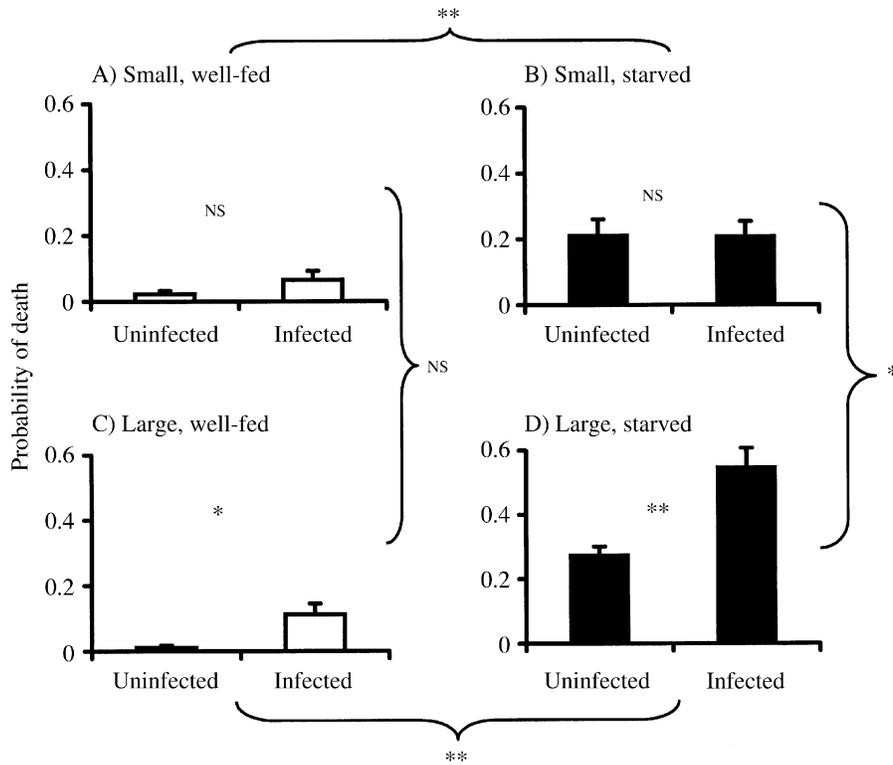


FIG. 5. Probability of death for uninfected and infected *Daphnia magna* 3–7 d after the beginning of the food reduction treatment in two size classes (small, 0.9–2.0 mm; large, 2.1–3.7 mm) for well-fed and starved populations (means + 1 SE). * $P < 0.00625$; ** $P < 0.00125$ (levels of significance Bonferroni adjusted for eight comparisons); NS, not statistically significant.

and 83.6 ± 34.3 h for the infected *Daphnia*. Small animals survived in complete starvation significantly longer both when uninfected (Fig. 6, Mann-Whitney U test $U = 460.5$, $P < 0.005$) and infected ($U = 698.0$, $P < 0.005$; levels of significance Bonferroni adjusted for two tests, $\alpha = 0.01/2$). We could not verify the infection status or spore load of the animals in this

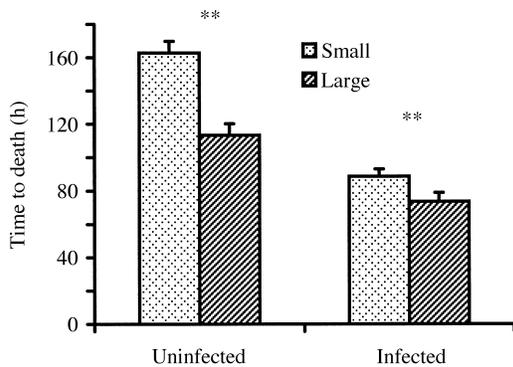


FIG. 6. Starvation times under complete starvation in relation to body length (small, 0.9–2.0 mm; large, 2.1–3.7 mm) for *Daphnia* taken from uninfected and infected populations (means + 1 SE). ** $P < 0.005$ (levels of significance Bonferroni adjusted for two comparisons).

experiment, but because *G. intestinalis* has high transmission rates, even the very young animals in the infection treatment were probably already infected (Ebert 1995, Ebert and Mangin 1997). The results of this starvation experiment are largely consistent with the results of the death rate analysis (Fig. 5), in that infected hosts tended to die at a higher rate than uninfected hosts and large animals with a higher rate than smaller animals.

DISCUSSION

Our study shows that food stress in infected *Daphnia* populations did not facilitate parasite spread. In accordance with our prediction, food shortage caused a disproportionately higher mortality of large animals in the infected populations, which reduced parasite spore load and prevalence (Fig. 3). However, after a few weeks of food reduction, the host population body length distribution regained a shape similar to its pre-treatment shape, and parasite spore load increased to approximately the same level as in the well-fed populations. Thus, our results do not support the conventional wisdom that stress leads to an increase in parasitism at the population level.

As mentioned in the introduction, parasite dynamics are influenced by factors that lead to an increase of population size as well as those that lead to a decrease.

Only their combined effects will reveal the net effect of a treatment. Our results indicate that food stress did not alter the rate at which parasites multiply. If resistance against infection had decreased in individual *Daphnia* in the populations during food reduction, one would expect spore load to increase for given host size or that the first spores would be detectable in smaller hosts (Fig. 4). We did not find this. On the other hand, if reduced resource level in *Daphnia* decreased spore production, as has been shown for two other *Daphnia*-microparasite systems (Ebert et al. 2000b, Bittner et al. 2002), this would result in a decrease in spore loads. We did not find this either. The relationship between body length and spore load in individual *Daphnia* did not change after the food reduction. In summary, our results suggest that food stress does not alter parasite multiplication rates (within-host growth and transmission), but does lead to a selective removal of strongly infected hosts from the stressed host population. This results in a net (temporary) reduction of parasite population size after the onset of food deprivation.

Lafferty and Holt (2003) make a distinction between effects of stress on the growth rate of the disease (R_0) and impact of disease. They stress that the distinction is important because although R_0 describes the parasite population growth, it does not represent the impact of the disease on host population growth. The effects of stress on host vital rates may interact with effects of stress on host susceptibility and lead to either a negative, positive, convex, or concave relationship between stress and disease (Lafferty and Holt 2003). In our experiment, the impact of food reduction on the infected host populations was about a 40% reduction in population size after 1 wk of food deprivation (week 22), while in the populations that remained well-fed, there was a 5% increase in population size. There was a marked decrease in R_0 for both well-fed and starved populations from week 21 to week 22, but in the starved populations this reduction was clearly larger than in the well-fed populations (Figs. 1A and 3). Thus in our experiment the effects of food stress seem to fall to the left upper corner of Fig. 2 in Lafferty and Holt (2003), i.e., food stress decreased R_0 and increased the impact of disease on the host population.

Previous theories about the role of stress in infected populations were based on combined results from studies on individual hosts (e.g., higher pathogenicity when hosts are under stress) and observations from disease outbreaks in natural populations (e.g., epidemic outbreaks of parasites may happen in high-density populations [Anderson and May 1981]). Our experiment is the first to test experimentally for stress effects on the population level. Thus, we are able to link our results to the treatments and exclude confounding factors. However, given the nature of these types of epidemiological experiments in which the treatments define the starting conditions, we cannot draw direct conclusions on cause and effect relationships. For example,

the reduction in food rations lead to a decrease in host density. Density may have an effect on the parasites (e.g., reducing transmission), perhaps influencing (even counteracting) other effects of starvation. Likewise, starvation may lead to a change in host behavior that could influence host and parasite dynamics. Behavior of *Daphnia* has been shown to play an important role in transmission dynamics (Ebert et al. 1997, DeCaestecker et al. 2002). Thus, while experimental epidemiology can include secondary order effects on the population level that may not be visible at the individual level, it cannot fully elucidate mechanistic cause and effect relationships. The results of our experiment must be seen within the complex interactions of two antagonistic populations as they occur. Only a combination of population- and individual-level experiments could lead to a complete understanding of host-parasite epidemiology.

Conventional wisdom was largely based on the demonstration that malnutrition can reduce immunocompetence, leading to higher parasite multiplication rates (Wakelin 1989, Lloyd 1995). For some invertebrates the opposite has been suggested, based on the phenotypically plastic response of the hosts to stress factors (Reeson et al. 2000). For example, crowding may lead to an increase in disease resistance, at least in some social insects (Traniello et al. 2002) and outbreak pest species (Wilson et al. 2002). Food limitation may also lead to changes in transmission via altered behavior of the hosts (Washburn et al. 1991). A more detailed inspection of single cause and effect relationships may reveal many other specific interactions between hosts and parasites that are modified by starvation or other stress factors. Thus, the distinction between vertebrates and invertebrates made here is clearly an oversimplification, as certainly differences among individual systems may disguise such a classification.

The results of this paper show that host starvation may have a significant effect on the epidemiology of microparasites in zooplankton populations. In our experiment, food deprivation induced a disproportionately high mortality among the most heavily infected individuals in the populations and thus decreased the parasite load in the host populations significantly. Periods of severe starvation are rather common for plankton populations (e.g., the clear water stage observed in many lakes) and may reduce host density by several orders of magnitude (Wetzel 1994). This is likely to slow down the parasites' transmission temporarily. In our experimental populations, the parasite load returned to prestarvation level in a few weeks, by which time the starved populations had reached a new equilibrium at the new carrying capacity. With more severe starvation, this may take much longer. The increased mortality rate due to food shortage may, however, also lead to an increase in transmission rates. *G. intestinalis* is known to spread from decaying host cadavers (Ebert 1995), and in our experimental containers, the dead

cadavers were readily reached by *Daphnia*. Spores released from decaying hosts can survive in the sediments for extended periods, and in the natural populations of shallow ponds and lakes, animals browsing for food particles on the sediment surface may become infected (Ebert et al. 1997). As this browsing behavior is intensified during starvation periods, it may lead to the start of epidemics in *Daphnia* inhabiting shallow water bodies (Ebert et al. 1997).

G. intestinalis is a microsporidian parasite of the *Daphnia* gut epithelium. The harm it inflicts on its host is rather minimal compared to other *Daphnia* parasites (Ebert et al. 2000a). Consistent with this, it hardly influenced host density (Fig. 1; see also Ebert et al. [2000a]). Its low virulence allows it to be maintained without intervention for long periods in relatively small laboratory cultures. It is very common in natural populations (Larsson et al. 1996, Stirnadel and Ebert 1997, Decaestecker 2002) and often reaches prevalence near 100% among the adult hosts. Our study shows that the apparent low virulence of this parasite can nevertheless have a profound impact on the size distribution of the host population, which reflects on the patterns of abundance of the parasite itself. There are many more parasites and epibionts of plankton organisms (Green 1974, Threlkeld et al. 1993) that appear harmless but that may show similar effects once properly investigated.

Effect of food stress on survival

Large infected and uninfected animals died sooner under starvation than small ones both in our population experiments and when individual animals were exposed to complete starvation. This was an unexpected result, as earlier laboratory studies found that small (uninfected) *Daphnia* survive less under starvation than larger animals (Tessier and Goulden 1982, 1987, Tessier et al. 1983, Tessier and Consolatti 1989, Perrin et al. 1992). This result has been regarded as universally valid among cladocerans and used to predict outcome of competition (Matveev and Gabriel 1994). However, some field studies have recently questioned this contention and indicated adult mortality is a major component in competitive exclusion (Matveev and Gabriel 1994) and midsummer decline in *Daphnias* (Hülsmann and Weiler 2000). In contrast to our study, individually grown animals in previous laboratory studies were acclimated to high food concentrations prior to starvation. In our experiment, *Daphnia* were kept in populations around the carrying capacity, i.e., at rather low food levels, before the starvation treatments, so they may have been preadapted to low resource levels. Our results therefore conform to the field data (Matveev and Gabriel 1994, Hülsmann and Weiler 2000), rather than to earlier laboratory studies (Tessier and Goulden 1982).

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