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PHYSIOLOGICAL ECOLOGY - ORIGINAL RESEARCH

Interactions between environmental stressors: the influence of salinity on host–parasite interactions between *Daphnia magna* and *Pasteuria ramosa*

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Abstract Interactions between environmental stressors play an important role in shaping the health of an organism. This is particularly true in terms of the prevalence and severity of infectious disease, as stressors in combination will not always act to simply decrease the immune function of a host, but may instead interact to compound or even oppose the influence of parasitism on the health of an organism. Here, we explore the impact of environmental stress on host–parasite interactions using the water flea *Daphnia magna* and its obligate parasite *Pasteuria ramosa*. Utilising an ecologically relevant stressor, we focus on the combined effect of salinity and *P. ramosa* on the fecundity and survival of the host, as well as on patterns of infectivity and the proliferation of the parasite. We show that in the absence of the parasite, host fecundity and survival was highest in the low salinity treatments. Once a parasite was introduced into the environment, however, salinity and parasitism acted antagonistically to influence both host survival and fecundity, and these patterns of disease were unrelated to infection rates or parasite spore loads. By summarising the form of interactions found in the broader *Daphnia* literature, we highlight how the combined effect of stress and parasitism will vary with the type of stressor, the trait used to describe the health of *Daphnia* and the host–parasite combination under observation. Our results highlight how the context-dependent nature of interactions between stress and parasitism inevitably

complicates the link between environmental factors and the prevalence and severity of disease.

Keywords Bacteria · Environmental conditions · Infection · Infectious disease · Salt stress · Stress · Virulence

Introduction

Throughout their lives animals are challenged by a range of environmental stressors that impact on the health and fitness of the individual. By definition, a stressor is any external factor that influences the ability of an organism to maintain its natural regulation or functioning to such an extent that fitness is reduced (Selye 1950; Brett 1958; but see Hoffmann and Parsons 1991); as such, a stressor can relate to an inexhaustible range of abiotic and biotic factors. Variation in environmental characteristics, such as temperature and food availability, together with the presence of predators and parasites have all been shown to impact on the physiological function and life-history of an individual (reviewed in Hofmann and Todgham 2010; Hawlena and Schmitz 2010; Flatt and Heyland 2011; Schmid-Hempel 2011). These naturally occurring stressors, however, are unlikely to act independently (i.e. additively) but instead may interact to compound or even oppose the influences of other stressors on the health of an organism (synergistic vs. antagonistic interactions, respectively). Viewing environmental stressors in isolation, therefore, can lead to a distorted picture of how individual stressors would potentially combine in natural populations to influence the health of an organism (Marcogliese and Pietrock 2011).

Of particular interest for studies of infectious disease is how the impact of parasites and pathogens is linked to

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other environmental sources of stress (Esch et al. 1975; Lafferty and Holt 2003; Marcogliese and Pietrock 2011). In the biomedical literature, it has been recognised that physical or even psychological stress can downregulate various components of the immune response (Khansari et al. 1990; Glaser et al. 1999), making individuals more susceptible to infectious disease. Environmental stressors will also influence the ability of an organism to acquire, process, and store nutrients, reducing the pool of resources that individuals can invest in disease resistance (e.g., condition dependence of immunocompetence, see Zuk and Stoehr 2002; Schmid-Hempel 2003). While the impact of environmental stress is, by definition, negative for the host, a more complex pattern may exist for the parasite. Early in the infection process, stress-induced suppression of the immune system will potentially aid the ability of a parasite to establish infection (Le Moullac et al. 1998; Malham et al. 2003). In contrast, once the parasite has evaded this initial immune response, any further stress will adversely affect host condition, decreasing the resources available for the parasite to grow and multiply within the host (Frost et al. 2008; Seppälä et al. 2008). Accordingly, both the timing and intensity of any environmental stress are important factors influencing how environmental challenges are linked to parasite fitness and the characteristics of infectious disease.

In the study reported here, we explored the impact of environmental stress on host–parasite interactions using the water flea *Daphnia magna*. *Daphnia* is increasingly being used as a model system for research into environmental stress (Eads et al. 2008) and host–parasite interactions (Ebert 2005, 2008). Studies have shown how food quality (Pulkkinen and Ebert 2004; Frost et al. 2008), maternal stress (Ben-Ami et al. 2010; Stjernman and Little 2011; Hall and Ebert 2012), temperature (Mitchell et al. 2005; Vale et al. 2008), predation (Yin et al. 2011) and pesticides (Coors et al. 2008; Coors and De Meester 2011) can all impact on host–parasite interactions. Here, we manipulated an additional stressor of *Daphnia* (i.e. salinity) and characterised the impact of the obligate bacterium parasite *Pasteuria ramosa* on host traits. For *D. magna* of coastal rock-pool habitats, salinity is an important environmental stressor. At our study location, peak fluctuations in salinity have been recorded up to 12.5 parts per thousand (approximately 12.5 g L⁻¹), although normal variation between rock pools is much lower [mean ± standard deviation (SD) parts per thousand: 1.97 ± 1.47, as summarised in Ranta 1979]. Such variation affects the overall physiological condition of individuals by influencing a range of key processes, such as feeding, swimming speed, metabolism, growth, reproduction and survival (Arnér and Koivisto 1993; Baillieul et al. 1998). Despite the importance of salinity as an environmental stressor for rock-pool

populations of *Daphnia*, however, the link between salinity and parasitism has not been directly explored, although studies have explored such interactions in other crustaceans (Dunn and Hatcher 1997; Piscart et al. 2007; Studer and Poulin 2012).

By exposing a single clone of *D. magna* to four salt concentrations (0.33–3 g L⁻¹ as commercial sea salt) and manipulating the presence or absence of the parasite *P. ramosa*, we aimed to understand how a natural stressor, salinity, can impact host–parasite interactions. In our study, we focused exclusively on the within-host dynamics of infection. To achieve this we first used a compatible combination of host and parasite clones (Luijckx et al. 2010). Second, we manipulated the salt concentration only after the individuals were exposed to the parasite. In this way we excluded the potential effects of stress on feeding or other behaviours, which may reduce the encounter rates between the parasite and the host, and explored the impact of stress directly on host physiology. We then measured characteristics of both host and parasite fitness, such as survival, offspring production, infectivity and spore production. In doing so, the aim of our study was to address three important questions: (1) how does host performance change along a salinity gradient; (2) how does the interaction between salinity and *P. ramosa* infection compare to the broader patterns of environmental stress and parasitism in the *Daphnia* literature; (3) can any differences in the severity of disease be accounted for by changes in the growth or infectivity of the parasite along a salinity gradient?

Materials and methods

Host and parasite clones

The host *Daphnia magna* Straus is a freshwater cladoceran that reproduces through cyclical parthenogenesis and inhabits eutrophic shallow ponds throughout Eurasia. The parasite *Pasteuria ramosa* Metchnikoff 1888 is a bacterial obligate endoparasite of *Daphnia* that causes severe loss of host fitness via castration and reduced survival (Ebert et al. 2004). Following infection, the parasite grows within the body cavity of its host, producing endospores that give the infected individual a distinctive reddish-brown colouration. Transmission only occurs horizontally with the spores released from the decaying cadaver of a formerly infected host.

For our study, all host individuals were derived from a single *D. magna* clone (SP1-2-3) originating from a rock-pool on the island of Spikarna, Tvärminne archipelago, in south-western Finland. In this small coastal rock-pool habitat, salinity can change within very short periods as the

result of a sudden invasion of Baltic Sea water or as a consequence of strong rain. The parasite clone (C19) originated from single spore infection based on an isolate (P1) of *P. ramosa* collected from Germany (Luijckx et al. 2010). This parasite is naturally infective to *D. magna* clone SP1-2-3 and resembles *P. ramosa* clones from the same rock-pool habitat (Luijckx et al. 2010). Parasite suspensions were produced by grinding up infected *D. magna* and adjusting the spore density to the desired concentration. A placebo suspension was produced with the same amount of uninfected *Daphnia*.

Parasite and salinity challenges

We used a split-brood design where individuals from a single mother were split across the two parasite treatments (exposed and unexposed) and the four salinity treatments (0.33–3 g L⁻¹), resulting in eight treatment combinations. Each treatment comprised 35 individual replicates (280 individuals in total), although nine individuals, distributed evenly across the treatments, were subsequently excluded as they died before the salinity manipulation could be applied. The saline concentrations were obtained by mixing commercial sea salt (hW-Meersalz; Wiegandt GmbH, Krefeld, Germany) in 5 L of artificial medium (ADaM; Klüttgen et al. 1994; modified after Ebert et al. 1998). Each time we created four salinity concentrations from 0.33 g L⁻¹ (the normal salinity of ADaM) and then 1, 2 or 3 g L⁻¹. Beginning with 3-day-old *Daphnia* placed individually in 100-mL jars filled with 20 mL of normal ADaM (0.33 g L⁻¹), we first exposed individuals to the parasite and then applied the salinity challenge. At ages 5 and 6 days, animals received 5,000 spores of the C19 *P. ramosa* clone (10,000 in total) or the equivalent volume of the placebo suspension as the control. At age 7 days, we moved individuals to new jars and replaced the standard (low salinity) medium with 100 mL of the appropriate experimental saline concentrations.

Thereafter, every 3 days (i.e. ages 10, 13, 16 days, etc.) we moved individuals to new jars containing fresh medium of the appropriate salinity treatment and counted the number of offspring produced. Individuals were monitored daily for survival until 30 days post-exposure to the parasite, at which point all *Daphnia* were individually frozen (–20 °C) for the later assessment of infection status and spore production. We counted the number of mature transmission stage spores by crushing the *Daphnia* cadavers and counting two independent samples of this suspension using a Neubauer improved counting chamber (depth 0.1 mm). Throughout the experiment we maintained individuals in a single climate-controlled incubator (light/dark: 16/8 h; 20 ± 0.5 °C) and regularly shuffled the location of the individuals to equalise any positional

effects. To account for the growing demands of the animals, food levels were increased gradually, beginning on day 2 with 0.5 × 10⁶ algae cells of *Scenedesmus obliquus* per *Daphnia* per day, and increasing to 1 × 10⁶, 2 × 10⁶, 2.5 × 10⁶, 3 × 10⁶ and 8 × 10⁶ algal cells per animal per day on days 4, 7, 9, 11 and 13, respectively.

Statistical analysis

We recorded survival and fecundity for 136 control individuals (33–35 individuals per treatment) and survival, fecundity, infection status and spore loads for 135 individuals exposed to *P. ramosa* (32–35 per treatment). In the subsequent analysis, all individuals were used for the estimates of host survival (control and exposed individuals) and infectivity (exposed individuals only). However, to avoid confounding other estimates of host and parasite performance with survival, only individuals that survived until the end of the experiment (day 30 post-exposure to *P. ramosa*) were used for estimates of host offspring production and parasite spore count. Specifically, we included 104 control individuals (23–31 individuals per treatment) and 75 infected individuals (15–21 individuals per treatment). We thus refer to the parasitism treatment as “parasite exposure” for host survival (as individuals may or may not have been infected when they died) and “parasite infection” for host offspring production (as individuals could be reliably identified as infected at day 30 post-exposure).

All statistical analyses were performed in R (ver. 2.12.2; R Development Core Team, available at: www.R-project.org). For host survival and host offspring production, we assessed the effect of both parasite exposure/infection and salinity using a full-factorial model where salinity, parasite exposure/infection and their interaction were included as fixed effects. For the parasite traits, infection status and spore counts, only salinity was included as a fixed effect. We analysed binomial data, such as host survival and infection status, using a generalised linear model with a binomial error link function. Before analyses, we tested for the assumptions of the linear (normality, homogeneity of variances) and generalised linear (over-dispersion) models. In the case of the offspring production data, there was significant heterogeneity in variances between the treatments groups (Levene’s test, $P < 0.001$). Accordingly, we used a white-adjusted analysis of variance to correct for unequal variances as implemented using the *car* package of R (Fox and Weisberg 2011). We also transformed spore load using a natural log transformation, although the figures here are presented on the original scale. Finally, treatment effects were explored post hoc using pairwise comparisons, based on Student’s *t* tests and chi-square tests for the linear and generalised linear models, respectively.

Results

We first examined the impact of salinity alone and salinity in combination with parasite exposure on the survival of *D. magna* over the period of experiment (Fig. 1). All individuals from the treatment groups were included in this analysis, irrespective of the infection status of exposed individuals. There was a significant effect of parasite exposure ($\chi^2_1 = 9.072$, $P = 0.003$) and an interaction between salt concentration and parasite exposure ($\chi^2_3 = 10.111$, $P = 0.018$) on host survival. There was no independent effect of salt concentration alone ($\chi^2_3 = 1.819$, $P = 0.611$). Figure 1 reveals how the interaction between salinity and parasite exposure influences the patterns of host survival. Only at the lowest salt concentration (0.33 g L⁻¹) was there a significant difference between exposed and non-exposed individuals, with the survival for both treatment groups converging as the salt concentrations increase.

We then examined the combined effect of salinity and parasite infection on the number of offspring produced by individuals that survived until the end of the experiment (30 days post-exposure). We found that offspring production was influenced by both parasite infection ($F_{1,170} = 1,943.188$, $P < 0.001$) and an interaction between salt concentration and parasite infection ($F_{3,170} = 9.567$,

$P < 0.001$). There was no independent effect of salt concentration alone on offspring production above all other factors ($F_{3, 170} = 1.060$, $P = 0.368$). As shown in Fig. 2, individuals that were exposed to the parasite and become infected produced significantly fewer offspring than those unexposed individuals who received only the salinity manipulation, with the difference between the two groups being greatest in the low salinity treatments (0.33–1.00 g L⁻¹).

Finally, we assessed the impact of salinity on parasite fitness based on estimates of both infectivity and the proliferation of the parasite within the host. Infection rates were unaffected by the salinity treatment ($\chi^2_3 = 0.772$, $P = 0.856$) and appeared to remain relatively constant across the salt gradients (Fig. 3a). The spore load of infected individuals, however, was influenced by the salt concentration ($F_{3,71} = 4.063$, $P = 0.010$). The number of mature spores present in individuals that survived until the end of the experiment (30 days post-exposure) was lowest for intermediate salt concentrations (Fig. 3b).

Discussion

Interactions between parasites and other environmental stressors have increasingly been identified as playing an

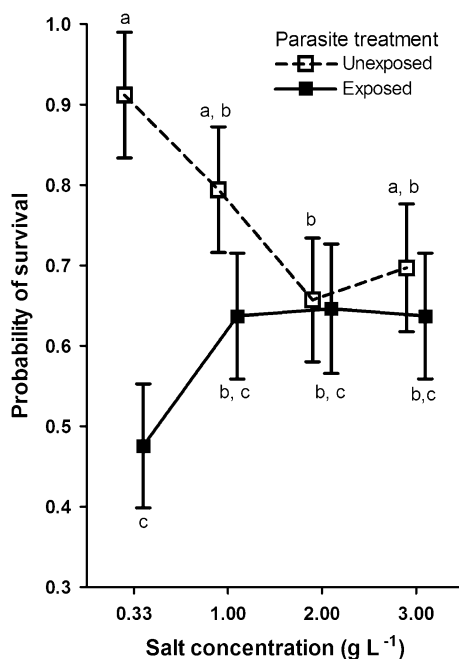


Fig. 1 The impact of salinity and parasite exposure on host survival [mean \pm standard error (SE)] over the duration of the experiment. Estimates of host survival for the parasite exposure treatment included all individuals irrespective of their infection status. Lowercase letters indicate significant groupings by post hoc comparisons using the chi-square test ($P < 0.05$)

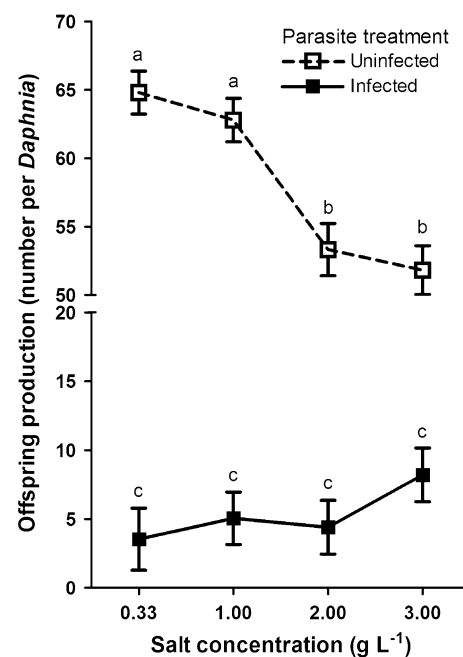


Fig. 2 The impact of salinity and parasite infection on host offspring production (mean \pm SE). Confounding of fecundity and survival estimates was avoided by only including those individuals that survived until the end of the experiment in the final estimates of offspring production. Lowercase letters indicate significant groupings by post hoc comparisons using Student's *t* test ($P < 0.05$)

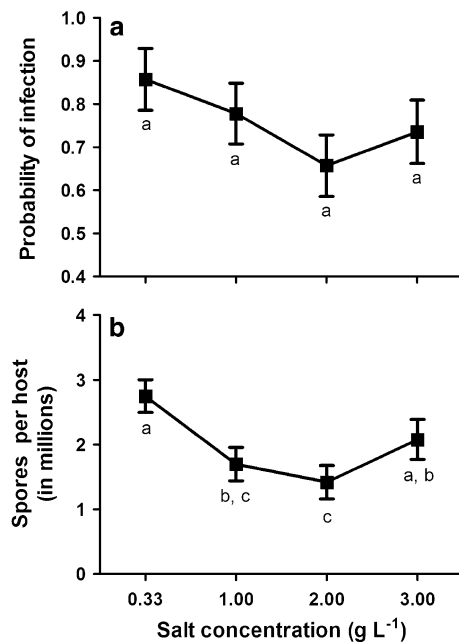


Fig. 3 The impact of salinity on infection rates (a) and spore loads (b) of *P. ramosa*. Only infected individuals that survived until the end of the experiment were included in the final estimates of spore production. For each panel, lowercase letters indicate significant groupings by post hoc comparisons using the chi-square test ($P < 0.05$) and Student's *t* test ($P < 0.05$) for infection rates and spore loads, respectively. Data are presented as the mean \pm SE

important role in shaping the health of an organism (reviewed in Marcogliese and Pietrock 2011). In our study, we focussed on the link between stress and infectious disease by manipulating the presence of the parasite *P. ramosa* and then exposing individuals of *D. magna* to a salinity gradient. Consistent with a range of other studies (Gonçalves et al. 2007; Martínez-Jerónimo and Martínez-Jerónimo 2007; Heine-Fuster et al. 2010), we found a decrease in host fitness at higher salt concentrations. In the absence of the parasite, host fitness was highest at the lower end of the salinity gradient, both in terms of survival (Fig. 1) and reproductive output (Fig. 2). Once a parasite was introduced into the environment, however, our results show that the correlation between salinity and host fitness changed considerably.

For the *Daphnia* and *P. ramosa* host–parasite system, the impact of parasitism is expected to reduce host survival and fecundity (Ebert et al. 2004). In our study, exposure to *P. ramosa* did not simply increase the average impact of salinity on host reproduction and survival (i.e. acted additively), rather it interacted antagonistically with salinity to modify the patterns of host fitness. In terms of host survival, the significant interaction between salinity and parasite exposure reflects that survival between unexposed and exposed individuals differed only at the lower end of the salinity gradient (Fig. 1). The interaction term for host

reproduction, in contrast, arises because the reproductive output of infected individuals remained constant across the salinity gradient (Fig. 2). Such antagonistic interactions have also been found in the amphipod *Gammarus roeseli*, where infection by an acanthocephalan parasite acted to increase the salinity tolerance of the host (Piscart et al. 2007). Antagonism between salinity and parasitism, however, is not necessarily universal, as other studies have found no clear evidence for an interaction between salinity and parasitism on traits such as clutch size (Dunn and Hatcher 1997) and survival (Sousa and Gleason 1989; Reisser and Forward 1991).

Indeed, context-dependent interactions between stressors appear to be a common feature of host–parasite interactions. To highlight this point, a summary of multiple studies across different *Daphnia*–parasite combinations is presented in Table 1 (this study; Bittner et al. 2002; Lass and Bittner 2002; Ebert et al. 2004; Coors and De Meester 2008; Frost et al. 2008; Vale et al. 2008; Coors et al. 2008; Yin et al. 2011). For each host trait and host–parasite combination, we describe the general impact of the environmental manipulation on host traits in the presence and absence of the parasite and classify the impact of multiple stressors as either additive, synergistic or antagonistic. From this summary, it is clear that all possible forms of interactions can be found in studies investigating stress and parasitism in *Daphnia* and that certain host–parasite combinations may also respond differently to a given stressor. In Table 1, for example, the interaction between pesticide exposure and parasitism was found to be additive for the timing and size of the first clutch but synergistic for host survival. Conversely, in Table 1 the effect of predator kairomones and the parasite *Metschnikowia* on the clutch size of *D. longispina* was additive, whereas for *P. ramosa* and the clutch size of *D. magna* it was antagonistic. Thus, the combined effect of stress and parasitism is neither specific to the host–parasite combination, the host trait of interest, nor to the type of environmental stress.

Underlying these different interactions between stressors may be some fundamental aspects of the biology of host–parasite interactions. Synergistic interactions could arise if the environment interfered directly with the ability of the host to cope with infection, whereas antagonistic interactions would occur if the environment interferes with the ability of the parasite to exploit the host. Interactions between stressors may also reflect the priorities placed on a given trait by both hosts and parasites. Under stressful conditions a host may maintain its investment in fecundity at the expense of survival, resulting in a synergistic impact of the parasite on survival under poor conditions. In our study, the antagonistic interactions between salinity and parasitism suggest that the physiological condition of a host is related to the overall severity of disease (i.e.

Table 1 Classifying the combined impact of multiple environmental stressors and parasitism on the survival and reproduction of the host *Daphnia*

Life-history trait	Environmental stress	Impact on host trait when the parasite is absent	Impact on host trait when the parasite is present	Evidence for an interaction between stress and parasitism	Combined effect of stress and parasitism	Host–parasite combination
Survival	Salinity increase	Decrease	Marginal increase	Yes	Antagonistic	<i>Daphnia magna</i> and <i>Pasteuria ramosa</i> ^a
	Food stress	Marginal increase	Decrease	Yes, but not formally tested	Antagonistic	<i>D. galeata</i> and <i>Chilomastix mesnili</i> ^b
	Food stress	Marginal decrease	Increase	Yes, but not formally tested	Antagonistic	<i>D. magna</i> and <i>P. ramosa</i> ^c
	Temperature increase	No change/ marginal decrease	Decrease	Yes	Synergistic	<i>D. magna</i> and <i>P. ramosa</i> ^d
	Predator kairomone	No change	No change	No	No change	<i>D. magna</i> and <i>P. ramosa</i> ^e
	Pesticide exposure	Marginal decrease	Decrease	Yes	Synergistic	<i>D. magna</i> and <i>P. ramosa</i> ^f
	Pesticide exposure	No change	Decrease	Yes	Synergistic	<i>D. magna</i> and <i>P. ramosa</i> ^f
	Pesticide exposure	No change	Decrease	Yes, but not formally tested	Synergistic	<i>D. magna</i> and <i>F. magnivora</i> ^f
	Salinity increase	Decrease	No change	Yes	Antagonistic	<i>D. magna</i> and <i>P. ramosa</i> ^g
	Food stress	Decrease	Reduced decrease	Yes	Antagonistic	<i>D. magna</i> and <i>P. ramosa</i> ^g
Overall fecundity	Temperature increase	Increase	No change	Yes	Antagonistic	<i>D. magna</i> and <i>P. ramosa</i> ^d
	Phosphorus limitation	Decrease	Decrease	No	Antagonistic	<i>D. magna</i> and <i>P. ramosa</i> ^d
	Pesticide exposure	Decrease	Decrease	No	Additive	<i>D. magna</i> and <i>P. ramosa</i> ^g
	Food stress	Decrease	Increase	No	Additive	<i>D. magna</i> and <i>F. magnivora</i> ^f
	Predator kairomone	No change	Decrease	Yes	Antagonistic	<i>D. galeata</i> and <i>C. mesnili</i> ^b
	Predator kairomone	Increase	Increase	Yes	Antagonistic	<i>D. magna</i> and <i>P. ramosa</i> ^f
	Pesticide exposure	Decrease	Decrease	No	Additive	<i>D. longispina</i> complex and <i>Metschnikowia sp.</i> ^h
	Food stress	No change	No change	No, but not formally tested	Additive	<i>D. magna</i> and <i>P. ramosa</i> ^f
	Predator kairomone	Decrease	No change	Yes	No change	<i>D. magna</i> and <i>P. ramosa</i> ^f
	Predator kairomone	Increase	Increase	No	Antagonistic	<i>D. longispina</i> complex and <i>Metschnikowia sp.</i> ^h
Age at first clutch	Predator kairomone	Decrease	Decrease	No	Additive	<i>D. magna</i> and <i>P. ramosa</i> ^f
	Pesticide exposure	Decrease	Decrease	No	Additive	<i>D. magna</i> and <i>P. ramosa</i> ^f
	Food stress	No change	No change	No, but not formally tested	No change	<i>D. magna</i> and <i>P. ramosa</i> ^f
	Predator kairomone	Increase	Increase	Yes	Antagonistic	<i>D. longispina</i> complex and <i>Metschnikowia sp.</i> ^h
First clutch size	Predator kairomone	Decrease	Decrease	No	Additive	<i>D. magna</i> and <i>P. ramosa</i> ^f
	Pesticide exposure	Decrease	Decrease	No	Additive	<i>D. magna</i> and <i>P. ramosa</i> ^f
	Food stress	No change	Increase	Yes	Additive	<i>D. magna</i> and <i>P. ramosa</i> ^f
	Predator kairomone	Increase	Increase	Yes	Additive	<i>D. magna</i> and <i>P. ramosa</i> ^f

We defined the combined effects of stress and parasitism depending on whether or not each stressor acted independently (i.e. additively) or interacted to compound (synergistic) or oppose (antagonistic) the influences of other stressors

^a This study

^b Bittner et al. (2002)

^c Ebert et al. (2004)

^d Vale et al. (2008)

^e Coors and De Meester (2008)

^f Coors et al. (2008)

^g Frost et al. (2008)

^h Yin et al. (2011)

ⁱ Lass and Bittner (2002)

condition-dependence of virulence; Krist et al. 2004; Restif and Kaltz 2006). For both survival and fecundity, differences between the control/uninfected and exposed/infected treatments were greatest in the low salinity treatment (i.e. 0.33 g L^{-1}). Therefore, the virulence of *P. ramosa* appears to depend on host condition, with the greatest reduction in host fitness occurring in the region of the salinity gradient where unexposed individuals performed best (Figs. 1, 2).

The effect of salinity on both host condition and the severity of disease, however, was not directly correlated with the prevalence of disease or the growth of the parasite (Fig. 3). Infection rates remained constant across the salinity gradient, a result that was unexpected given that host susceptibility to *P. ramosa* has been shown to depend on a range of environmental stressors (Frost et al. 2008; Vale et al. 2008; Allen and Little 2010; Hall and Ebert 2012). While studies in other host–parasite systems have also documented the lack of a correlation between salinity and host susceptibility (e.g. the amphipod *Pagurus novizealandiae*; Studer and Poulin 2012), contributing to our findings may have been the timing of the infection process at which the salinity treatment was applied. By using a compatible host–parasite combination and exposing *Daphnia* to the parasite before manipulating the salinity, we may have missed the window in which stress can impact on the ability of a host to clear a parasite infection. As such, we predict that the timing of the environmental stress to impact on the ability of a *P. ramosa* to colonise *Daphnia* (as opposed to influencing parasite growth) may only be a matter of hours.

Nonetheless, the timing of the salinity application would not explain the non-linear effect of salinity on parasite spore loads. The proliferation of the parasite within the host was associated with a small, but significant decrease in parasite spore loads associated with intermediate salinity concentrations. In the *Daphnia*–*Pasteuria* system, other non-linear relationships have previously been reported for temperature and parasite growth (Vale et al. 2008; Vale and Little 2009). In this case, the authors suggest that higher parasite growth at intermediate temperatures is driven by the relationship between *Daphnia* feeding rates and temperature (and presumably nutrient acquisition). Without a physiological understanding of how salinity impacts on both *Daphnia* and *Pasteuria*, however, it is difficult to explain why *P. ramosa* performs better at the extremes of a salinity gradient. Instead, we can point to the growing evidence that interactions between salinity and parasitism may be inherently complex. Short- versus long-term changes in salinity (hours compared to weeks), for example, can differentially influence the production of parasite transmission stages (Koprivnikar and Poulin 2009; Lei and Poulin 2011), and parasites may even be able to manipulate the osmoregulation ability of their host (Brooks and Mills 2011).

In summary, based on our results and a growing number of studies in both *Daphnia* (see Table 1) and other organisms (reviewed in Marcogliese and Pietrock 2011), it is clear that the presence of a parasite can modify the impact of additional stressors that may be present in the environment. Such trait-specific interactions are not only limited to the traits of the host, but they are also reflected in other characteristics of disease, such as the ability of a parasite to colonise and grow within the host. In particular, the timing of the stress and whether the immune system/physiological condition of the host is affected will differentially shape the characteristics of disease. The context-dependent nature of stress and host–parasite interactions, therefore, inevitably complicates how the environment is understood to influence the evolution of virulence, as well as the predictability for which environmental factors can be linked to the prevalence and severity of disease.

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