Running title: Parasite effects on stickleback

The association of feeding behavior with the resistance and tolerance to parasites in recently diverged sticklebacks

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Abstract

Divergent natural selection regimes can contribute to adaptive population divergence, but can be sensitive to human-mediated environmental change. Nutrient loading of aquatic ecosystems, for example, might modify selection pressures by altering the abundance and distribution of resources and the prevalence and infectivity of parasites. Here, we used a mesocosm experiment to test for interactive effects of nutrient loading and parasitism on host condition and feeding ecology. Specifically, we investigated whether the common fish parasite *Gyrodactylus* sp. differentially affected recently diverged lake and stream ecotypes of three-spined stickleback (*Gasterosteus aculeatus*). We found that the stream ecotype had a higher resistance to *Gyrodactylus* sp. infections than the lake ecotype, and that both ecotypes experienced a cost of parasitism, indicated by negative relationships between parasite load and both stomach fullness and body condition. Overall, our results suggest that in the early stages of adaptive population divergence of hosts, parasites can affect host resistance, body condition, and diet.

Keywords: three-spined stickleback, diet, parasites, environmental change, eutrophication, aquatic ecosystems, metabolic condition, trade-offs, adaptive divergence

Introduction

Contrasting environmental conditions can generate the divergent selection pressures that drive adaptive population divergence (Schluter, 2000; Nosil, 2012), and during this process several traits linked to species interactions can evolve rapidly, including those associated with feeding behaviours and defences against predators and parasites (Kortet et al., 2010; Eizaguirre & Lenz, 2010; Karvonen & Seehausen, 2012; Schluter, 2000). The evolution of such traits might influence the buildup of reproductive isolation (Servedio et al., 2011; Nosil, 2012; Eizaguirre et al., 2009a), particularly if they are also involved in mate choice (Maan & Seehausen, 2011) and local adaptation (Eizaguirre & Lenz, 2010; Eizaguirre et al., 2011; Konijnendijk et al., 2013; Bassar et al., 2010; Arnegard et al., 2014). During the early stages of adaptive population divergence, environmental change can weaken divergent selection regimes, and this might erode ecological and genetic differentiation between populations (Hendry et al., 2009; Seehausen et al., 2014; Vonlanthen et al., 2012). In most cases, however, we lack a mechanistic understanding of how environmental change, be it abiotic or biotic, can influence performance variation among and within diverging populations. This makes it difficult to predict the effect of environmental change on the dynamics of population divergence (Nosil, 2012; MacColl, 2011).

The resistance and tolerance of hosts to parasites are traits that can both evolve rapidly (Råberg, 2014; Siva-Jothy & Thompson, 2002; Murray et al., 1998) and influence host energy acquisition and resource assimilation over a range of environmental conditions (Pianka, 1981; Huey et al., 2001; Howick & Lazzaro, 2014). Resistance is the ability of individuals to limit their parasite load, and is measured as the number of parasites acquired per individual host over a given time period (Råberg et al., 2009). Tolerance, on the other hand, is the ability of individuals to reduce the harmful effects of a given parasite load, and can be estimated (at the population level) as the slope of the relationship between parasite load and proxies of fitness (Schneider & Ayres, 2008; Råberg et al., 2009; Kaufmann et al., 2014). The evolution of host resistance and tolerance strongly depends on how parasites affect trade-offs between immune defense and life history traits (Poulin, 2007; Råberg, 2014; Hamilton & Zuk, 1989). For instance, when hosts lack the genetic material to fight off infections, they may increase energy acquisition to compensate for the costs of parasitism (Howick & Lazzaro, 2014; Ponton et al., 2011; Zuk & Stoehr, 2002). If individual hosts

are in positive energy balance (e.g. full guts, ample fat reserves), they might be able to either better cope with the physiological demands of the infection or to mount a more effective immune response (Zuk & Stoehr, 2002). In general, host responses to parasites tend to be strongly dependent on environmental conditions that determine host energy balance (Wong & Candolin, 2015; Budria & Candolin, 2014).

In aquatic systems, nutrient loading can have strong effects on host-parasite interactions (Johnson et al., 2008, 2007; Budria & Candolin, 2014), by changing the chemical (e.g. phosphorus and oxygen levels), physical (e.g. light, turbidity) and biological properties of aquatic ecosystems (Smith & Schindler, 2009). Nutrient loading has been shown to alter resource availability and competitive interactions (Leach et al., 1977; Talbot & Hole, 1994), predation risk (Cothran et al., 2012; Van de Meutter et al., 2005), mating behaviour (Cothran et al., 2012; Jaervenpaeae & Lindstroem, 2004), and the prevalence and virulence of parasites (Zuk & Stoehr, 2002; Kortet et al., 2010; Johnson et al., 2012; Domenici et al., 2007; Halstead et al., 2014). Previous experiments in aquatic systems have found strong effects of nutrient loading on the morphology and behaviour of vertebrates (Johnson et al., 2010; Budria & Candolin, 2014). However, less is known about how nutrient loading might affect parasite resistance, tolerance, and feeding ecology of hosts in the early stages of adaptive divergence.

The stickleback-*Gyrodactylus* model system is useful for investigating the interactive effects of parasitism and environmental change (i.e. nutrient loading) on the evolution of host resistance during adaptive population divergence. Lake and stream three-spined stickleback (hereafter, sticklebacks) populations that are genetically and phenotypically differentiated (i.e. ecotypes) have evolved multiple times in the Northern hemisphere since the last glaciation ($\simeq 12$ 000 years) (Hendry et al., 2009; Feulner et al., 2015). For our study, we chose a pair from the Lake Constance region in central Europe that has diverged very recently (<150 years) (Lucek et al., 2010), concomitant with dramatic changes in nutrient levels in Lake Constance. Lake Constance has experienced a prolonged phase of eutrophication (1950-1980: total Phosphorus, TP increased from $\simeq 10$ to 80 $\mu g/L$) and re-oligotrophication (1980-2010: TP decreased from $\simeq 80$ to 10 $\mu g/L$) (Jochimsen et al., 2013), but the ecological and evolutionary effects of these nutrient dynamics on the resident stickleback populations are unknown. The specific lake-stream pair that we used is phenotypically divergent in body size, growth rate (Lucek et al., 2012), and

morphology (Berner et al., 2011; Lucek et al., 2013), and is genetically differentiated at several putatively adaptive loci (Marques et al., 2016). For a parasite, we chose the monogean (Platy-helminthes) flatworm, *Gyrodactylus spp.* (hereafter, *Gyrodactylus*), because it is a common ec-toparasite of sticklebacks and other fish (Bakke et al., 2007; Rahn et al., 2015; Kalbe et al., 2002; Raeymaekers et al., 2011; Stephenson et al., 2015), and it is known to drive local adaptation in other stickleback populations (Eizaguirre & Lenz, 2010; Konijnendijk et al., 2013). *Gyrodactylus* is a flatworm that reproduces on the skin and gills of fish (Bakke et al., 2007), and can increase host mortality (Bakke et al., 2007), reduce body condition (Eizaguirre et al., 2011), and reduce life-time reproductive success (Eizaguirre et al., 2009b). It has no intermediate host, and it is generally more common in stream than in lake environments (Bakke et al., 2007; Eizaguirre et al., 2011; Kalbe et al., 2002).

We performed a 7-week mesocosm experiment to investigate how closely related lake and stream ecotypes of stickleback differ in their resistance and tolerance to the monogenean ectopar-asite *Gyrodactylus* in aquatic ecosystems with low and high levels of nutrient loading. In this study, we tested the following two predictions. First, we predicted that lake and stream ecotypes would differ in their resistance and tolerance to *Gyrodactylus* infections. This prediction is based on previous work showing rapid parasite-mediated evolution of stickleback immune systems in other populations (Eizaguirre et al., 2012a,b). Second, we predicted that nutrient loading might differ-entially affect the resistance and tolerance of both ecotypes to parasite exposure, partly because of the potential impacts of nutrient loading on the energy balance (e.g. stomach fullness), diet composition (i.e. gut contents), and body condition (a fitness proxy) of sticklebacks (Schlotz et al., 2013; Budria & Candolin, 2014; Howick & Lazzaro, 2014).

Materials and methods

Mesocosms Experiment

Our experimental setup consisted of 40 outdoor mesocosms, and was a complete randomized block design, with factorial combinations of fish ecotype (lake and stream), nutrient level (high and low nutrients, HN and LN), and *Gyrodactylus* exposure (exposed and unexposed, G+ and G-,

Fig. 1). The experiment lasted 7 weeks, from May 3 to June 21, 2013. Each mesocosm was filled with 1000 L of filtered water from Lake Lucerne (Switzerland) and seeded with a mixture of sand, gravel, benthic substrate, and zooplankton. We amended nutrient levels by adding 20 ml of NaNO$_3$ and HNa$_2$PO$_4$ stock solution to achieve the following target concentrations: 640 μg N/L and 40 μg P/L in HN tanks, and 80 μg N/L and 5 μg P/L in LN tanks, which represents approximately the median and the minimum, respectively, of the total phosphorus concentration in Lake Constance over the past 50 years (Jochimsen et al., 2013). Each mesocosm received either 6 or 7 stickleback so as to standardize biomass among mesocosms (Mean-lake= 24.55 g $\pm 0.621 SE$, Mean-stream= 23.15 g $\pm 0.292 SE$)

Fish Collection and Parasite Exposure

We collected stream sticklebacks from two streams (Aubach: $47^o19'37.45''$ N $9^o34'12.82''$ E; and Zapfenbachkrummensee Kanal: $47^o21'20.65''$ N $9^o36'11.94''$ E) and lake sticklebacks from the shore of an inlet delta of Lake Constance (Staad: $47^o29'8.30''$ N $9^o32'38.25''$ E) in early spring 2013. Twenty lake and stream sticklebacks were dissected to estimate the abundance of *Gyrodactylus* parasites in these natural populations. For the experiment, we initially removed *Gyrodactylus* by treating wild-caught fish with a 1:4000 diluted solution of formalin (Raeymaekers et al., 2011; Buchmann & Kristensson, 2003). A visual inspection of all fish under stereomicroscope revealed no living parasites after disinfection. After one week, we then performed a standardized infection of the fish that we used for the mesocosm experiment. Fish were anesthetized ($N = 278$) with 0.1% MS222 (Gilderhus & Marking, 1987) and placed on a petri dish filled with saline solution ($6.4g/L$ NaCl). With a fine brush, we manually added exactly four individual parasites onto each individual that was part of the infection treatment (G+). We used two parasites that originated from either naturally infected lake fish or stream fish, so as to account for possible differences in host-parasite coevolution (Eizaguirre et al., 2011; Konijnendijk et al., 2013). Unexposed fish (G-) used in the experiment were handled in the same way but received no parasites. After this procedure, the fish were kept in $5L$ aquaria for a week prior to their introduction into the mesocosms.

Parasite levels, fish condition and diet

At the end of the experiment, we caught the surviving fish ($N = 199$ fish out of $N = 278$), euthanized them using an overdose of MS222, and counted the number of *Gyrodactylus* per fish (i.e. parasite load). Three tanks were excluded from the experiment because of low survival (see Fig-S1 and Table-S3). Additionally, one randomly chosen fish from each mesocosm was excluded from all analyses ($N = 40$), because it was needed for another experiment (Brunner et al *submitted*). For the remaining 159 fish used in the current study, we measured standard length (mm, $\pm$ 1mm), body weight (g, $\pm$ 0.01 g), liver weight (mg, $\pm$ 0.001 mg), and wet gut mass (mg, $\pm$ 0.001 mg) (see Table-S3 for descriptive statistics). Fish guts were stored in saline solution (6.4 g/L NaCl) and frozen at -20 $°C$ in separate vials for later dietary analyses.

The hepatosomatic index (HSI), was used as a proxy of individual variation in body condition, where $HSI = [LM/(BM - W_{FS})]x100$, and LM is the wet liver mass (g) and BM wet body mass (g) (Chellappa et al., 1995; Hammerschmidt & Kurtz, 2005; Kurtz et al., 2004). We estimated energy balance based on the relative fullness index of the fish guts (FI), where $FI = [W_{SC}/BM]x100$, and W_{SC} is the difference between the mass of the full stomach and empty stomach (Magnusson et al., 2003). To quantify diet composition, the whole content of each gut was flushed into petri dishes and the prey items were identified to the lowest taxonomic level possible (Streble & Krauter, 2006; Tachet et al., 2000). Individual prey were counted on a 2.5 x 2.5 mm grid under a stereomicroscope, and in our analyses we focused on the six most abundant prey items, namely Collembola, Nymphs, Chironomidae, Ostracoda, Chydoridae, and Cyclopoida (Table-S1).

Statistical Analyses

The effects on fish mortality were tested using generalized linear mixed effect models (GLMM), with a binomial family (logit-link); fish ecotype, nutrient levels, parasite exposure and their interactions were fixed effects and block was used as random effect. To analyse variation in parasite load (number of *Gyrodactylus* per fish), we used a GLMM with the number of parasites as the response variable, body length as an offset, a negative binomial error family (square-root link), and tanks nested within blocks as random effects. Parameters were estimated by Penalized Quasi Maximum

Likelihood (PQL) with the R package *MASS*, which produces a better dispersion of the fitted and predicted values compared to other R packages (e.g., *lme4* and *glmmADMB*, Zuur et al., 2009; Bolker et al., 2009).

We tested for the effects of our treatments on body condition (HSI) and the fullness index (FI) with linear mixed effect models (LMM) with tank nested within block as random effects and parasite load as a continuous variable. To test if nutrient loading altered the resistance and tolerance of the ecotypes to *Gyrodactylus*, we performed mixed effects ANCOVA with either body condition (HSI) or diet (FI) as responses variables, with the interaction of ecotype x nutrient x parasite load as explanatory variables (Råberg et al., 2009), and with tanks nested within block as a random effect. Differences between slopes were tested using likelihood ratio tests and the X^2 test statistic following (Hayward et al., 2014). To test if slopes were significantly different from 0 (i.e. tolerant or intolerant) we used a $t-test$ with the R package *lmertest*.

To analyse variation in diet composition, we focused on changes in diet community composition among treatments within experimental blocks. To this end, we used a distance-based redundancy analysis (db-RDA) framework with the function *capscale ()* in the R-package *vegan* in which we: (i) Hellinger transformed the abundance diet matrix by block (Borcard et al., 2011), (ii) averaged the Hellinger transformed matrix by tank and used it as our response matrix in the db-RDA, and (iii) tested for the significance (RDA's F statistic) of our experimental treatments on the transformed matrix by permuting all treatments within a block 999 times. We also tested for the effects of the experimental treatments on each individual prey item from the Hellinger-transformed species matrix (159 x 6 matrix) in a binomial GLMM model, with tank nested within block as a random effect (Table-S3). For all LMM and GLMM, we estimated X^2 and P-values in an $anova$ Type II analysis with the function *Anova ()* from the *car* R-package. All statistical analyses were performed in R-version 3.2.3 (R Core Team, 2015) and the data is available online on Dryad http://dx.doi.org/10.5061/dryad.b0846.

Results

In this experiment, 79 out of 278 fish died during the seven weeks of the experiment. Overall, lake sticklebacks died more than stream sticklebacks (Ecotype: X^2= 4.164, P= 0.041, Fig-S1), and

mortality was highest for lake sticklebacks that were exposed to *Gyrodactylus* (G+) (Ecotype by Exposure interaction: X^2= 6.912, P= 0.008, Fig-S1). We did not find a significant main effect of nutrients (Nutrients: X^2= 0.033, P= 0.854) on mortality.

Prediction 1 (ecotype differences in host resistance and tolerance)- We found that parasite load was highest for lake fish, both in the wild (Lake-wild= 30.4 ±5.23 SE, Stream-wild= 4.68± 1.75 SE, $X^2_{1,40}$= 30.22, P<0.001) and in the experiment (Lake-experiment= 36.27 ± 13.33 SE, Stream-experiment= 3.01 ± 5.65 SE; $X^2_{1,159}$= 20.262, P < 0.001, Fig. 2A). In the experiment, parasite load was negatively related to the body condition ($X^2_{1,159}$= 12.136 P < 0.001), and there was no effect of ecotype on the slope of this relationship (Table-S2), implying that both ecotypes were equally intolerant to parasite load (Fig. 3A). We also found that stomach fullness (FI) was negatively related to parasite load ($X^2_{1,159}$= 4.838, P= 0.028) and that there was no effect of ecotype on the slope of this relationship (Table-S2). However, when we split this analysis by ecotype, we found that the relationship between parasite load and gut fullness was significantly negative for lake fish ($X^2_{1,72}$= 6.197, P<0.05) but not for stream fish ($X^2_{1,87}$= 0.264, P> 0.1, Figure 3D).

Prediction 2 (interactive effects of nutrient)- Overall, fish of both ecotypes were in better condition in HN tanks ($X^2_{1,159}$= 7.406, P < 0.01, Fig. 2B, Table 1). While we found no effects of nutrients on the parasite load of stream fish (Stream-HN= 2.7 ± 0.777 SE, Stream-LN= 3.28 ± 0.919 SE, $X^2_{1,87}$= 0.011, P= 0.916), the parasite load of lake stickleback was lower in HN compared to LN tanks (Lake-HN= 29.60 ±18.9 SE, Lake-LN= 43.74 ±18.94 SE, $X^2_{1,72}$= 7.47, P< 0.01). In addition, we found that the lake fish in the LN tanks had higher parasite loads and there were significant negative relationships between parasite load and body condition (Lake-HN: slope= -0.027 ± 0.024 SE, t-test$_{60.23}$= -1.158, P= 0.251; Lake-LN: slope= -0.063 ± 0.021 SE, t-test$_{61.03}$= -2.942, P < 0.01, Fig. 3B) and stomach fullness (Lake-HN: slope= -0.051 ± 0.039 SE, t-test$_{58.71}$= -1.198, P= 0.199; Lake-LN: slope= -0.084 ± 0.036 SE, t-test$_{63.21}$= -2.307, P = 0.024, see Fig. 3E).

Ecotypes did not differ in their relative gut fullness (Table 1 and Fig. 2C) or in their diet composition (Table 2, Fig. 4). However, initial parasite exposure significantly altered diet composition (F= 2.776, P < 0.05; Table 2 and Fig. 4), such that parasite-exposed fish ate more copepods ($X^2_{1,159}$= 4.152, P= 0.042) and fewer nymphs ($X^2_{1,159}$= 4.703, P= 0.03) than non-exposed fish (see Fig-S2 and Table-S1 and S3).

Discussion

Understanding how environmental changes affect species evolution is an ongoing challenge. Here we predicted that lake and stream stickleback ecotypes would differ in their resistance and tolerance to *Gyrodactylus* and that nutrient loading would differentially affect the host-parasite interactions for each ecotype. Overall, our results provide partial support for both predictions. First, we found that lake sticklebacks had higher mortality and lower resistance to infection than stream sticklebacks when exposed to *Gyrodactylus* (Fig. 2). However, among the surviving fish, both lake and stream ecotypes were equally intolerant to parasite load (i.e. similar slopes in Fig. 3A). Second, we found that nutrient loading increased the resistance of lake but not stream stickleback, and, exclusively for lake stickleback, we found some evidence that nutrient loading reduced the cost of parasite load (Fig. 3B).

In the wild, lake and stream stickleback ecotypes inhabit different foraging habitats during most of their life cycle, which expose them to contrasting parasite communities (Kalbe et al., 2002; Karvonen et al., 2015; Feulner et al., 2015). Around the Lake Constance basin, stream sticklebacks reside year round in streams, feed primarily on benthic prey items, and are exposed to a lower diversity of parasites than lake fish (Lucek et al., 2013; Moser et al., 2012, 2015b; Karvonen et al., 2015). Lake sticklebacks, on the other hand, live most of their life in the open water of the lake and migrate to nearshore environments and stream channels to breed (Lucek et al., 2013; Moser et al., 2012). They primarily feed on planktonic prey and are exposed to a higher diversity of parasites than stream fish (Moser et al., 2012; Karvonen et al., 2015; Moser et al., 2015b; Lucek et al., 2013). Previous work on other stickleback populations has shown that contrasting foraging habitats and parasite communities are important drivers of phenotypic and genetic differentiation (Feulner et al., 2015; Karvonen et al., 2015), and that rapid evolution of host resistance might be common during the formation of ecotypes (Eizaguirre et al., 2012b,a, 2009a, 2011; Stutz et al., 2014; Oke et al., 2016; Feulner et al., 2015).

Rapid evolution of resistance to contrasting parasite communities is a common outcome of host-parasite interactions (Schmid-Hempel, 2011). The higher level of resistance of stream stickleback could result from coevolution of stream fish with *Gyrodactylus*, one of the most common parasites in stream environments (Kalbe et al., 2002; Eizaguirre et al., 2011). In other stickleback

populations, adaptation to contrasting parasite communities in lakes and streams have resulted in divergent immune responses between ecotypes (Eizaguirre & Lenz, 2010; Feulner et al., 2015; Karvonen & Seehausen, 2012). Mainly because contrasting parasite communities are strongly associated with differences in the frequency of alleles of the major histocompatibility complex (MHC) between populations (Eizaguirre et al., 2012a, 2009a; Matthews et al., 2010). Therefore, if host-parasite coevolutionary dynamics in Lake Constance are similar to other systems around Europe, it is possible that resistance alleles, of either the MHC or other relevant genes, have recently increased in frequency in the stream but not in the lake populations (Eizaguirre & Lenz, 2010; Eizaguirre et al., 2012b,a, 2011). While there is recent evidence for adaptive genomic differentiation between the lake and stream ecotypes from in Lake Constance (Karvonen et al., 2015; Moser et al., 2012, 2015a; Roesti et al., 2015; Lucek et al., 2013; Marques et al., 2016), nothing is known about differentiation of MHC genes for this lake-stream pair and this should be addressed in further studies.

While the evolution of host resistance can influence diversification and maintenance of genetic diversity between and within populations (Buckling & Rainey, 2002; Summers et al., 2003; Eizaguirre et al., 2009b), the evolution of tolerance may favour the spread of parasites (Best et al., 2008; Kaufmann et al., 2014), break down Red-Queen dynamics, and generate multiple stable-states (Best et al., 2014; Råberg, 2014). This is because tolerance can mitigate the negative effects of parasitism, but does not contribute to decreasing parasite load (Best et al., 2014; Råberg, 2014). In our experiment, we did not find significant differences in the degree of tolerance between ecotypes (e.g. slopes in Fig 3A are not different). However, we did find lower parasite loads in the lake ecotype at high nutrient levels (Table 1 and S1), and this suggests that lake fish might be able to mitigate the negative effects of parasite load on body condition if they can increase energy intake in more productive environments (Fig 2B-D and 3E). Such compensatory effects could have significant impacts on host-parasite dynamics if they help maintain susceptible host genotypes in the population (Brunner & Eizaguirre, 2016) or if they allow virulent parasite strains to increase in frequency (Kause & Ødegård, 2012; Best et al., 2014; Råberg, 2014).

Our experiment also revealed a number of links between individual variation in parasite load, body condition, and diet (i.e stomach fullness and composition). First, the positive relationship between body condition (HSI) and stomach fullness (FI) is consistent with an expected positive re-

lationship between food acquisition and overall metabolic state. Second, the negative relationship between stomach fullness and parasite number in the lake ecotype suggests potentially higher costs of parasitism for lake fish. This is also consistent with the higher mortality of lake fish when exposed to parasites. In high nutrient environments, lake fish might be able to improve their capacity to cope with *Gyrodactylus* by increasing feeding rates, or changing their diet (e.g. Fig 3E) so as to include more profitable prey (Huey et al., 2001; Arrington et al., 2002; Pianka, 1981). Second, despite known ecotype differences in feeding traits (Lucek et al., 2013; Berner et al., 2011; Marques et al., 2016), in a common foraging environment we found no diet differentiation between ecotypes, but found that stickleback exposed to *Gyrodactylus* ate more cyclopoids and fewer nymphs than non-exposed individuals (Fig. 4, Fig-S1 and Table-S2).

More work on the food quality of different prey items and the effect of these parasites on the functional response of sticklebacks would be necessary to determine if the diet changes associated with parasite exposure were indicative of a compensatory mechanism or not. While it is well known that trophically transmitted parasites can affect host feeding behaviour (e.g. *Schistocephalus* and *Diplostomum* parasites on stickleback feeding rate Lefèvre et al., 2009; Aeschlimann et al., 2000; Jakobsen et al., 1988; Milinski, 1993), our results suggest that directly transmitted parasites can also affect feeding behaviour (e.g. diet preference) and activity (e.g. gut fullness) of their hosts, possibly via effects on host condition.

The stickleback populations used in this experiment were introduced in the Lake Constance region approximately 150 years ago (Lucek et al., 2010; Marques et al., 2016), and over this short time period (< 100 generations) they have evolved divergent life history traits such as growth rate, lifespan, and time of first reproduction (Lucek et al., 2013, 2012; Moser et al., 2012). Building on these previous studies, we found that parasitism can influence both ecological interactions between stickleback and their prey (e.g effects of parasite exposure on diet and stomach fullness) and potentially influence selection gradients experienced by hosts (i.e. relationships between parasite load and body condition). However, more work is needed to understand the mechanisms by which parasitism can affect trade-offs between immunity and feeding behaviour, particularly in natural populations.

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Acknowledgments

Table 1: Response of sticklebacks for parasite load, metabolic condition (HSI), and stomach fullness (FI) to fish ecotype, nutrient levels, parasite exposure, their interactions and parasite load. Significant P values (P < 0.05) are highlighted in bold.

	Parasite load						Metabolic Condition (HSI)						Stomach fullness (FI)					
	Overall		Lake		Stream		Overall		Lake		Stream		Overall		Lake		Stream	
	X^2	P	X^2	P	X^2	P	X^2	P	X^2	P	X^2	P	X^2	P	X^2	P	X^2	P
Gyrodactylus N							12.136	**<0.001**	8.295	**0.004**	3.668	0.055	4.838	**0.028**	6.197	**0.013**	0.264	0.608
Ecotype	20.262	**<0.001**					1.855	0.173					2.981	0.084				
Nutrient	0.023	0.879	7.47	**0.006**	0.011	0.916	7.406	**0.006**	0.365	0.546	5.546	**0.019**	4.463	**0.035**	1.063	0.302	3.281	**0.070**
Exposure	0.008	0.929	6.276	**0.012**	1.19	0.275	1.778	0.182	1.225	0.268	0.445	0.505	0.407	0.524	0.144	0.704	0.162	0.687
Ecotype:Nutrient	0.015	0.903					1.093	0.296					0.195	0.659				
Ecotype:Exposure	5.626	**0.018**					0.053	0.819					0.002	0.968				
Nutrient:Exposure	1.918	0.166	0.354	0.552	2.08	0.149	0.226	0.635	0.348	0.555	0.056	0.813	0.165	0.685	0.001	0.974	0.017	0.897
Ecotype:Nutrient:Exposure	1.229	0.268					0.358	0.55					0.007	0.935				
N	159		72		87		159		72		87		159		72		87	

Table 2: Effects on sticklebacks prey composition at the end of the experiment. R^2 values in the db-RDA correspond to partial R^2 = explained sum of squares (SS) /Total SS Significant P values (P < 0.05) are highlighted in bold.

		Prey composition (db-RDA)											
		Overall				Lake				Stream			
	DF	SS	F	P	R2	SS	F	P	R2	SS	F	P	R2
Ecotype	1	0.188	1	0.4	0.04								
Nutrients	1	0.283	1.508	0.181	0.06	0.311	1.285	0.277	0.143	0.171	1.227	0.31	0.102
Exposure	1	0.464	2.469	**0.045**	0.099	0.307	1.272	0.302	0.141	0.219	1.574	0.202	0.131
Ecotype:Nutrients	1	0.229	1.217	0.295	0.049								
Ecotype:Exposure	1	0.071	0.379	0.837	0.015								
Nutrients:Exposure	1	0.086	0.458	0.769	0.018	0.046	0.188	0.98	0.021	0.075	0.535	0.711	0.045
Ecotype:Nutrients:Exposure	1	0.079	0.42	0.81	0.017								
Total SS$_{df}$		4.7_{25}				2.176_9				1.671_{12}			

List of figures

1. Experimental design. We crossed in a fully factorial design, lake and stream stickleback (A), exposure to *Gyrodactylus* (B), and nutrient loading in mesocosms environments (C).

2. Effects of the experimental treatments on (A) the individual number of parasites, (B) fish condition measured as the hepatosomatic index (HSI), (C) stomach fullness measured as the fullness index (FI), and (D) the relationship between HSI and FI. Symbols show mean values, shapes and colours represent ecotypes (circles = lake and squares = stream) and *Gyrodactylus* exposure treatments (red = G+ and blue = G-). X axis show the nutrient effects (HN = high and LN = low). Bars in panels A to C represent $\pm$ SE).

3. Association between the number of parasites with fish condition (A-C), and stomach fullness (D-F). Panels A and D show the effects of parasite load on body condition (HIS) and relative stomach fullness (FI) for each ecotype. Panels B,C, E, and D show the effects of nutrient levels on the relationship between parasites and conditions on lake (A an E) and stream (C and F) separately. Symbols' shapes and colours represent ecotypes (circles = lake and squares = stream) and nutrient treatments (dark green = HN and cyan = LN). The lengths of the regression lines represent the rage of infection (resistance) and slope (tolerance) the degree of damage produced by increasing parasite infections. Slopes significantly different from 0 are marked with (*).

4. Effects on diet composition shown in a scaling 3 RDA biplot. Symbols' shapes and colours represent ecotypes (circles = lake and squares = stream) and *Gyrodactylus* exposure treatments (red = G+ and blue = G-). Arrows show the magnitude and direction of change of prey in the constrained multidimensional space (RDA axes). Centroids and bars represent the mean effect of the treatments and the standard errors. RDA explained variances are shown in brackets. $^{*}P < 0.05;^{**}P < 0.01;^{***}P < 0.001$ show significant axes and significant effects on prey items (see supplementary material)

A

Lake   Stream
Fish ecotype

✖

B

G+   G-
Gyrodactylus **exposure**
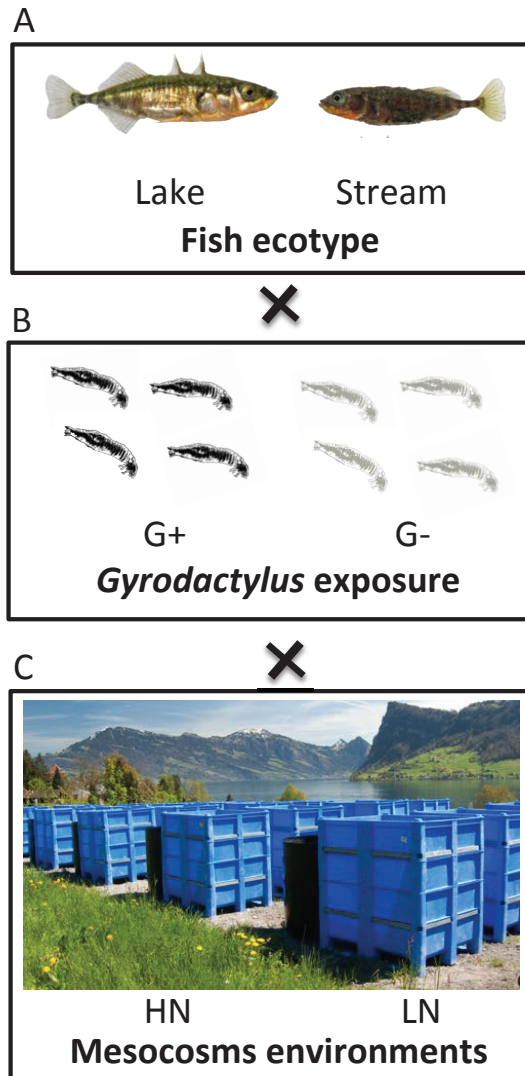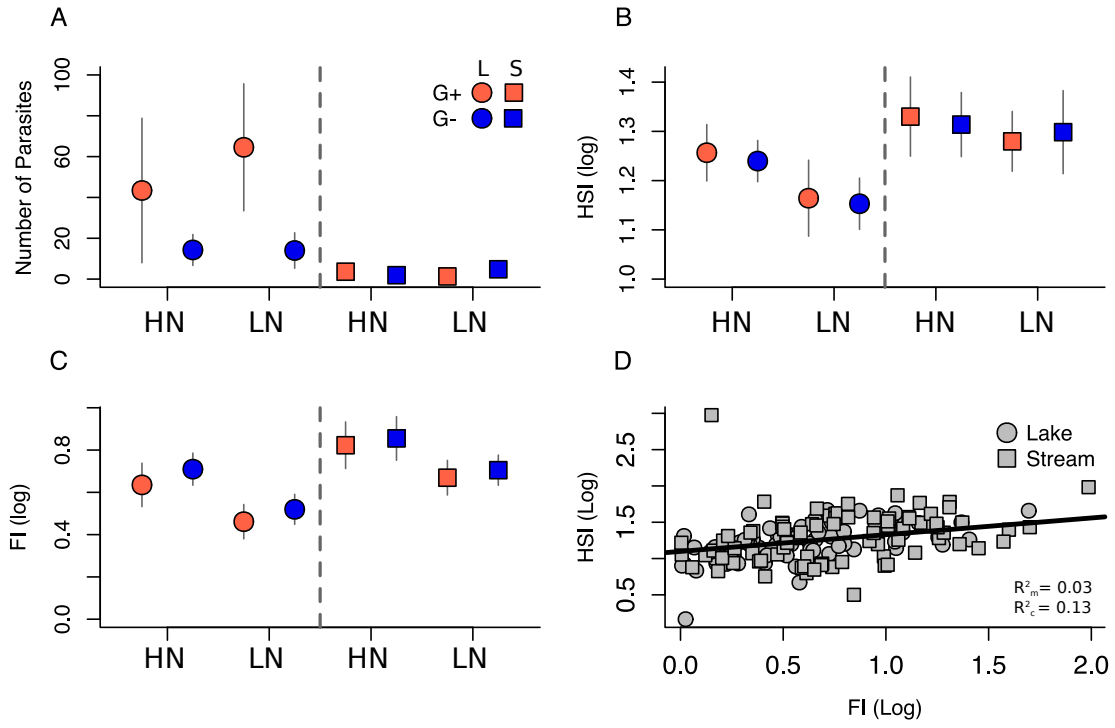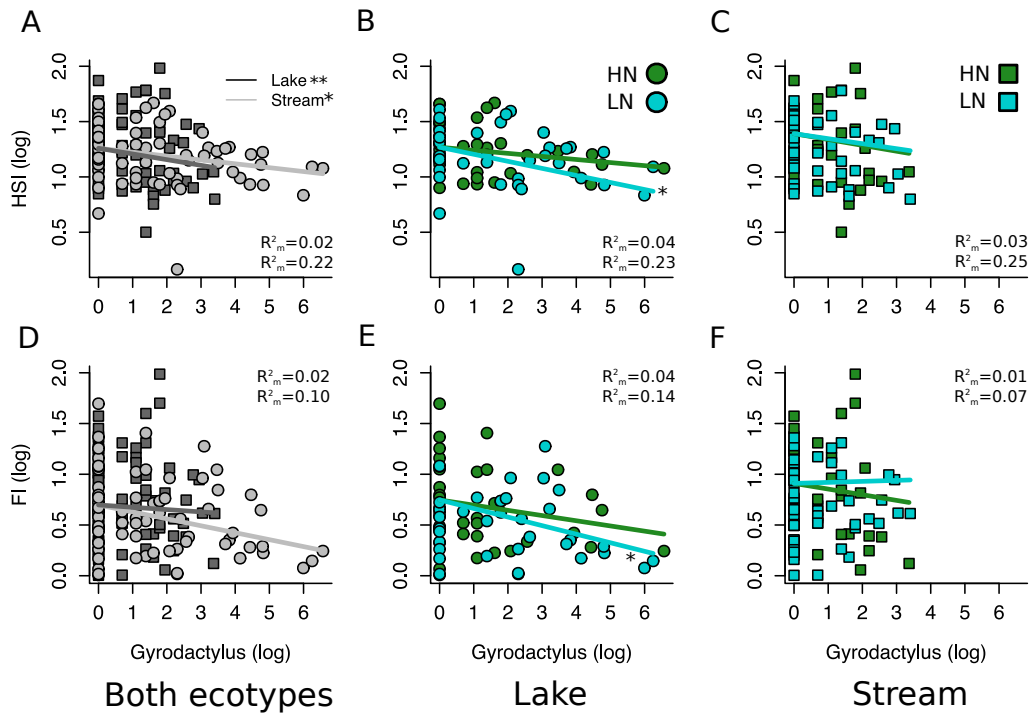
✖

C

HN   LN
Mesocosms environments

Figure 1:

Figure 2:

Figure 3:

Figure 4: