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Effects of the pathogenic water mold *Saprolegnia ferax* on survival of amphibian larvae

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ABSTRACT: Infectious diseases are a significant threat to worldwide biodiversity. Amphibian declines, a significant part of current biodiversity losses, are in many cases associated with infectious disease. Water molds are one group of pathogens affecting amphibians on a worldwide basis. Although water molds have been studied extensively for their effects on host embryos, little information is available about how they affect post-embryonic amphibians. We tested the effects of one species of water mold, *Saprolegnia ferax*, in a comparative study of larvae of 4 amphibian species: *Pseudacris regilla* (Pacific treefrog), *Rana cascadae* (Cascades frog), *Ambystoma macrodactylum* (long-toed salamander), and *R. aurora* (red-legged frog). *S. ferax* can kill amphibians at the embryonic and juvenile life history stages, depending on the amphibian species. In the present study, a 1 wk exposure to *S. ferax* killed *P. regilla* larvae and a 2 wk exposure killed *R. aurora* larvae. Larvae of the other host species were unaffected after 1 wk of exposure to *S. ferax*. Our results suggest that *S. ferax* can kill amphibian larvae and further suggest that evaluation of how pathogens affect amphibians at the population level requires investigation at various life stages.

KEY WORDS: Water mold · *Saprolegnia* · Amphibian · *Ambystoma macrodactylum* · *Pseudacris regilla* · *Rana aurora* · *Rana cascadae*

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INTRODUCTION

Biodiversity losses characterized by population declines, range reductions, and species extinctions are occurring at alarming rates (Raven 1987, Myers 1993, Singh 2002). One contributing factor is infectious disease (Daszak et al. 2000, Harvell et al. 2002, Altizer et al. 2003). As part of these worldwide losses in biodiversity, many amphibian populations are declining and disappearing (Houlahan et al. 2000). Although many factors probably contribute to amphibian population declines (Blaustein & Kiesecker 2002), infectious disease may be a major cause (Daszak et al. 1999). For example, the pathogenic fungus *Batrachochytrium dendrobatidis* is associated with many amphibian population declines (e.g. Berger et al. 1998). However, other pathogens, including viruses, bacteria, trematodes, mesomycetozoans,

and water molds, may also contribute to increased mortality and population declines in amphibians (e.g. Blaustein et al. 1994, Green et al. 2002).

Water molds (Stramenopila: Oomycota: Oomycetes: Saprolegniales) are fungus-like protists that inhabit aquatic habitats and moist soils (Dick 1990). Most aquatic water molds are saprobes that grow on dead organic matter (Dick 1990), and some water molds, including aquatic species, are capable of parasitism (Dick 1990, Johnson et al. 2002). Water molds that infect amphibians include *Achlya flagellata*, *Aphanomyces* sp., *Saprolegnia ferax*, *S. parasitica*, and probably a number of other species (Tiffney & Wolf 1939, Blaustein et al. 1994, Berger et al. 2001, Lefcort et al. 1997). *S. ferax* contributes to massive mortality of frog embryos (Blaustein et al. 1994, Kiesecker & Blaustein 1995). In addition, a mixed-species culture containing

multiple species of both *Achlya* and *Saprolegnia* caused mortality and sublethal effects in embryos of *Rana sylvatica* (wood frog) and *Bufo americanus* (American toad) (Touchon et al. 2006, Gomez-Mestre et al. 2006). The effects of water molds on amphibian embryos appear to be influenced by several environmental factors, including temperature, pH, ultraviolet-B radiation, precipitation, plant cover, and oviposition behavior (Kiesecker & Blaustein 1995, Green 1999, Kiesecker et al. 2001a, Ruthig 2006, additional references in Romansic et al. 2006).

In contrast to the extensive research on the effects of water molds on amphibian embryos, there is relatively little information available about how water molds affect post-embryonic amphibians. There is experimental evidence that *Saprolegnia* influences competitive interactions between larvae of *Rana cascadae* (Cascades frog) and *Pseudacris regilla* (Pacific treefrog), apparently by mediating densities of the competing species through differential effects on embryos (Kiesecker & Blaustein 1999). Water mold infections have been reported in amphibian larvae in nature (Bragg & Bragg 1958, Bragg 1962, Berger et al. 2001, Converse & Green 2005), and occur occasionally in larvae of North American species (Converse & Green 2005). Berger et al. (2001) reported massive mortality of *Bufo marinus* (cane toad) larvae associated with *Aphanomyces* sp. infection, and *Saprolegnia* has been found on dead amphibian larvae after mass mortality events (Bragg & Bragg 1958, Bragg 1962). Romansic et al. (2006) demonstrated that *Saprolegnia* can kill larvae of *R. aurora* (northern red-legged frog), and Romansic et al. (2007) found that *Saprolegnia* can kill newly metamorphosed *R. cascadae*. The *Saprolegnia* isolates used in these studies were recently identified as *S. declina* (used in the Romansic et al. 2006 study of larvae) and *S. ferax* (used in the Romansic et al. 2007 study of metamorphs, see 'Materials and methods' of the present paper for identification methods). As in amphibian embryos, the effects of *Saprolegnia* on amphibian larvae appear to be influenced by environmental stressors (Lefcort et al. 1997, Romansic et al. 2006). In addition, Walls & Jaeger (1987) found that *Saprolegnia* infection in *Ambystoma maculatum* (spotted salamander) larvae was associated with bite wounds and mortality from aggressive *A. talpoideum* (mole salamander) larvae.

Recently published models suggest that mortality of post-embryonic amphibians contributes significantly to population declines (Biek et al. 2002, Vonesh & De la Cruz 2002). Therefore, a full investigation of the potential effects of amphibian pathogens at the population level requires information on how these pathogens affect different life history stages, not just a single stage such as embryos. Therefore, we conducted a comparative study of the effects of *Saprolegnia ferax* in

larvae of 4 amphibian species: *Pseudacris regilla*, *Rana cascadae*, *Ambystoma macrodactylum* (long-toed salamander), and *R. aurora*.

MATERIALS AND METHODS

Collection and maintenance of amphibians. Amphibians were kept in a laboratory maintained at approximately 13 to 15°C in 2002 and 10.5 to 14°C in 2003. All water used in the study, unless otherwise noted, was tap water conditioned with NovAqua® and Amquel® water conditioners. Amphibians were kept under a natural photoperiod; however, *Pseudacris regilla* and *Rana cascadae* (used in Expt 1) were transferred from a natural photoperiod to a 12 h light:12 h dark cycle at the start of Expt 1, as were *Ambystoma macrodactylum* larvae at the start of Expt 2. Prior to experimentation, frog embryos and larvae were kept in 38 l aquaria (length × width × height = 50 × 25 × 31 cm). Frog larvae were fed a mixture (3:1 by volume) of rabbit chow and Tetramin® fish flakes (hereafter: tadpole food) ad libitum. Metamorphosed frogs (≥50% tail resorption) were removed.

Pseudacris regilla larvae (Gosner development stages 25 to 35, Gosner 1960) were collected on 2 September 2002 from the Potholes subalpine meadow, located approximately 0.6 km NW of Todd Lake, Deschutes County, Oregon, USA (~1980 m elevation). Prior to experimentation, *P. regilla* were maintained in aquaria filled with about 35 l of water and not aerated. *P. regilla* were kept at a density of approximately 5 larvae l⁻¹ water for 4 d, then transferred to new tanks and maintained at a density of approximately 2.3 larvae l⁻¹. Larvae were then transferred to new tanks every 7 d.

Rana cascadae larvae were collected from the Potholes meadow on 23 August and 18 September 2002. *R. cascadae* collected on 23 August ranged from Gosner stages 25 to 35, and those collected on 18 September ranged from stages 36 to 40. Prior to experimentation, *R. cascadae* were maintained unaerated in aquaria filled with about 35 l of water. *R. cascadae* were housed at a density of approximately 1.6 larvae l⁻¹ and transferred to new tanks with new water every 6 to 9 d. *R. cascadae* were kept at a lower density than *Pseudacris regilla* because they were larger in size.

Ambystoma macrodactylum larvae were collected on 28 August and 2 September 2002 from the Potholes meadow. To prevent cannibalism, which sometimes occurs among *A. macrodactylum* larvae (Walls et al. 1993), larvae were maintained individually in plastic cups (bottom diameter = 8 cm, top diameter = 11 cm, height = 11 cm) filled with 0.6 l of water. Prior to their use in Expt 2, *A. macrodactylum* were transferred to new cups with new water every 1 to 9 d and fed *Tubifex* sp. worms.

Six *Rana aurora* egg masses (approximately 290 to 350 embryos mass⁻¹, Gosner stages 18 to 21) were collected from Coast Pond (~8 km south of Waldport, Lincoln County, Oregon, USA) on 16 January 2003. Embryos were placed in aerated aquaria filled with about 30 l of water. *R. aurora* were transferred to new aquaria with new water every 7 to 8 d prior to being tested in Expt 3. Hatchlings were separated from unhatched individuals 15 d after hatching began and used in Expt 3. During the time between separation of hatchlings and Expt 3, *R. aurora* larvae ranged in density from 2.5 to 9.5 larvae l⁻¹.

Collection, isolation, and culture of *Saprolegnia ferax*. *S. ferax* was isolated from a water sample collected on 10 September 2002 at the shore of Lost Lake in the Oregon Cascade Range (Linn County, 1220 m elevation), a site where *S. ferax* has contributed to massive mortality of *Bufo boreas* embryos (Blaustein et al. 1994, Kiesecker & Blaustein 1995, Kiesecker et al. 2001a). This isolate can kill juvenile frogs (Romansic et al. 2007). *S. ferax* was isolated using hemp seeds and YpG (yeast-glucose) agar media (Fuller & Jaworski 1987) and identified as *S. ferax* using available keys and standard methods (Seymour 1970, Johnson et al. 2002). Identification was confirmed using a DNA barcoding procedure and comparison to available DNA sequences in GenBank (J. E. Johnson et al. unpubl. data). To obtain *S. ferax* for use in experiments, sterile hemp seeds were added to Petri dishes containing *S. ferax* cultures on YpG agar media to allow seeds to become inoculated with *S. ferax*. Seeds were then removed and added to standardized Petri dishes (diameter = 85 mm, height = 12 mm, 50 seeds per dish) filled approximately half full with ultrapure water and incubated for 7 d at approximately 20 to 23° C. *S. ferax* hyphae grew between seeds in dishes, producing clumps of seeds connected by a mycelium of *S. ferax* containing hyphae and zoosporangia. These clumps of seeds were used to apply *S. ferax* treatments in experiments.

Experimental setup. To minimize handling stress, larvae were not measured at the start of experiments. Instead, within 1 d of the start of each experiment, leftover larvae not used in the experiment were measured for Gosner stage (frogs only) and total length in mm. These measurements serve as estimates of these parameters for the exposed and control larvae at the start of the experiments. All experiments were checked once per day. Larvae were examined visually for hyphae consistent with water mold infection in amphibian larvae (Converse & Green 2005). Dead larvae were removed and preserved in 70% ethanol. Larvae surviving to the end of their experiment were euthanized using MS-222 and preserved in 70% ethanol.

Expt 1. Mid-late stage *Pseudacris regilla* and *Rana cascadae* larvae: *P. regilla* and *R. cascadae* larvae were exposed to *Saprolegnia ferax* in a 2 × 2 factorial experiment that manipulated frog species and *S. ferax*. Experimental units consisted of plastic boxes (dimensions: 31 × 18 × 8 cm) filled with 2 l of water to a depth of 4 cm and stocked with 6 larvae. Larvae were added to units haphazardly, except that each unit received either *P. regilla* or *R. cascadae*. On the day following the start of the experiment, 10 larvae of each species were haphazardly selected from the unused larvae. Selected *P. regilla* ranged in Gosner stage from 31 to 39 and in total length from 24.5 to 33.2 mm, while selected *R. cascadae* ranged from stage 33 to 41 and in total length from 31.5 to 53.0 mm. There were 2 pathogen treatments: *S. ferax* and control. There were 5 replicates of each treatment–frog species combination, for a total of 20 experimental units. Treatment–frog species combinations were assigned to units randomly. The *S. ferax* treatment consisted of a clump of 50 hemp seeds overgrown with *S. ferax* hyphae and zoosporangia. The control treatment consisted of 50 sterile hemp seeds. A cage consisting of a small plastic box (length × width × height = 9.5 × 9.5 × 6.5 cm) with sides of 1 mm fiberglass mesh and no top was placed in the center of each unit. Seeds were placed inside the cage to prevent larvae from feeding on *S. ferax* and thereby reducing the production of zoospores. However, the mesh sides of the cage allowed passage of *S. ferax* zoospores. The experiment began on 30 September 2002 and lasted for 7 d. During this experiment, larvae were fed tadpole food ad libitum.

Expt 2. *Ambystoma macrodactylum* larvae: *A. macrodactylum* larvae were exposed to *Saprolegnia ferax* and control treatments. Experimental units consisted of plastic boxes and cages filled with water (as above). To prevent cannibalism, each unit was stocked with only 1 larva. Larvae were assigned to units randomly. The 4 remaining larvae not used in the experiment ranged from 40.8 to 58.2 mm in total length (mean ± SE = 51.6 ± 3.9). There were 2 treatments: *S. ferax* and control. Treatments were assigned to units randomly. There were 17 replicates of each treatment, for a total of 34 units. The *S. ferax* treatment consisted of a clump of 50 hemp seeds overgrown with *S. ferax* hyphae and zoosporangia. The control treatment consisted of 50 sterile hemp seeds. Six days after the experiment began, *A. macrodactylum* were fed *Tubifex* sp. The experiment lasted for 7 d.

Expt 3. Stage 25 *Rana aurora* larvae: *R. aurora* larvae were exposed to *Saprolegnia ferax* and control treatments. Experimental units consisted of plastic boxes and cages filled with water (as above) and stocked with 5 *R. aurora* larvae. Five of the 6 egg masses (Masses 1 through 5) were chosen at random and each unit received one haphazardly selected Gosner stage 25 larva (2 to 3 wk post-hatching) from each

of these masses. Units received larvae in random sequence. At the start of the experiment, 2 Gosner stage 25 *R. aurora* not used in the experiment were haphazardly selected from each of Masses 1 through 5. These larvae ranged in total length from 15.7 to 25.1 mm (mean \pm SE = 20.1 \pm 0.80).

There were 2 treatments: *Saprolegnia ferax* and control. Treatments were assigned to units randomly. There were 5 replicates of each treatment, for a total of 10 units. The *S. ferax* treatment consisted of a clump of 50 hemp seeds overgrown with *S. ferax* hyphae and zoosporangia. The control treatment consisted of 50 sterile hemp seeds. After 7 d, larvae were transferred to new boxes with new water and new cages and treatments were re-applied. Each unit received 0.03 g of tadpole food at the start of the experiment and immediately after transfer to new boxes. The experiment began on 7 February 2003 and lasted for 14 d. Because the lack of *S. ferax*-induced mortality in some species in Expts 1 and 2 may have been due to short exposure times, the duration of Expt 3 was doubled to increase our ability to detect a possible effect of *S. ferax* on survival.

Densities of zoospores and zoospore cysts. For every application of a *Saprolegnia ferax* treatment, the combined number of zoospores and zoospore cysts of *S. ferax* were estimated using a cytometer for seed clumps randomly selected from the unused clumps in the batch used in the application (Table 1). These estimates were extrapolated to an estimated density of zoospores and cysts in the *S. ferax* treatment at application using the volume of water in units (Table 1).

Statistical analyses. In experiments where mortality occurred, survival data had heteroscedasticity that could not be removed by transformation. Therefore, for these experiments, we analyzed survival using non-parametric methods.

Survival in Expt 3 was analyzed using a rank-sum test. For Expt 1, we tested for an overall difference between the two treatments and a difference between treatments for each frog species using a multiple comparisons procedure. We used a Bonferroni adjustment (Ramsey & Schafer 1997) to keep $\alpha = 0.05$ for this pro-

cedure. There were 3 pairwise comparisons of interest, so the p-value for rejection of null hypotheses in this procedure was set at 0.0167. Pairwise comparisons were made using rank-sum tests. To investigate whether susceptibility to mortality from *Saprolegnia ferax* differed between *Pseudacris regilla* and *Rana cascadae*, we tested whether the effect of *S. ferax* on survival was different between the 2 species using a rank-based nonparametric contrast (Zar 1999).

RESULTS

Expt 1. Mid-late stage *Pseudacris regilla* and *Rana cascadae* larvae

Median survival was lower in the *Saprolegnia ferax* treatment compared to the control treatment (rank-sum test, $Z = -2.4918$, $p = 0.0127$), indicating a main effect of *S. ferax* treatment (Fig. 1). For *Pseudacris regilla*, median survival was lower in the *S. ferax* treatment compared to the control treatment ($Z = -2.8347$, $p = 0.0046$), while for *R. cascadae*, survival was 100% in both the *S. ferax* and control treatments. However, the effect of the *S. ferax* treatment was not different in *P. regilla* compared to the *R. cascadae* treatment (non-parametric contrast, $S = -4.8941$, $0.10 < p < 0.25$).

In the *Saprolegnia ferax* treatment, there were 13 observations of hyphae consistent with *S. ferax* infection on live *Pseudacris regilla* (Table 2). All larvae that died were in the *P. regilla*-*S. ferax* treatment combination and were partially or completely eaten by conspecifics before inspection for hyphae. Hyphae were not observed on any of the 3 partially eaten carcasses found. Hyphae were not observed on *P. regilla* in the control treatment or on *Rana cascadae* in either treatment.

Expt 2. *Ambystoma macrodactylum* larvae

Survival was 100% in both the control treatment and the *Saprolegnia ferax* treatment. Hyphae were not observed on any *Ambystoma macrodactylum* (Table 2).

Expt 3. Stage 25 *Rana aurora* larvae

Median survival was lower in the *Saprolegnia ferax* treatment (40.0%, 95% CI: 0.0–40.0%) compared to the control treatment (100%, $Z = 2.8247$, $p = 0.0047$). In the *S. ferax* treatment, hyphae consistent with water mold infection were observed on live indi-

Table 1. *Saprolegnia ferax*. Mean estimates of the number and density of the combined number of zoospores and zoospore cysts at each *S. ferax* treatment application

Expt	Application	Zoospores and zoospore cysts		
		Mean no. (± 1 SE)	n	Mean density (l ⁻¹)
1	Sole application	1.3×10^8	1	6.6×10^7
2	Sole application	$1.3 \times 10^8 \pm 6.5 \times 10^6$	3	6.5×10^7
3	Initial application	$4.4 \times 10^7 \pm 7.8 \times 10^6$	2	2.2×10^7
	Re-application	3.3×10^7	1	1.6×10^7

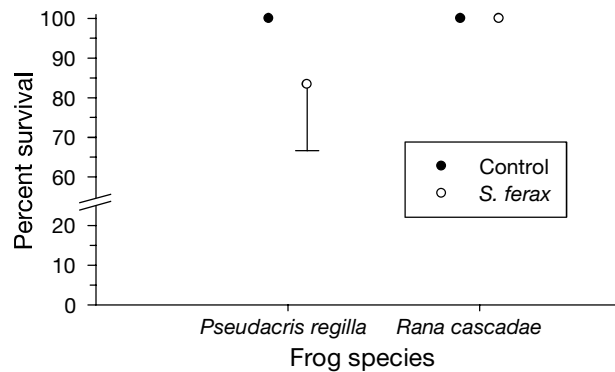


Fig. 1. *Pseudacris regilla* and *Rana cascadae*. Median survival of larvae exposed to *Saprolegnia ferax* and control treatments in Expt 1. The sole error bar is 95% CI

viduals 12 times and on 16 of 17 dead individuals (Table 2). One individual in the *S. ferax* treatment was completely eaten before inspection. No hyphae were observed in the control treatment.

DISCUSSION

Expt 1 suggests that 1 wk of exposure to *Saprolegnia ferax* killed mid-late stage *Pseudacris regilla* larvae. In contrast, 1 wk of exposure to *S. ferax* did not kill mid-late stage *Rana cascadae*, although there was insufficient statistical evidence to demonstrate that *P. regilla* was more susceptible to mortality from *S. ferax* than *R. cascadae*. In Expt 2, 1 wk of exposure to *S. ferax* did not kill *Ambystoma macrodactylum* larvae. Expt 3 suggests that 2 wk of exposure to *S. ferax* killed

early stage *R. aurora* larvae. In addition, hyphae consistent with *S. ferax* infection were observed on live *P. regilla* and live and dead *R. aurora* exposed to *S. ferax* treatments. Taken together, the results of the present study demonstrate that *S. ferax* can kill amphibian larvae and suggest that effects on larvae contribute to the overall effects of *Saprolegnia* on populations of its amphibian hosts.

Many of the pathogenic water molds have a wide host range and can persist and reproduce as saprobes on dead organic matter (Tiffney 1936, Dick 1990, Johnson et al. 2002). These characteristics give water molds considerable potential to affect amphibian populations. Water molds are likely to persist at high densities even if the density of one or more amphibian host species declines. Thus, if mortality from a pathogenic water mold contributes to a decline in an amphibian population, such mortality could continue as the population dwindles, potentially driving the host population to extinction. Persistence and reproduction as a saprobe and ability to cause mortality at low host densities are attributes that may aid the fungal pathogen *Batrachochytrium dendrobatidis* in driving to amphibian populations to extinction (Daszak et al. 1999).

Our results, together with previous research on amphibians and water molds (e.g. Kiesecker & Blaustein 1995, Berger et al. 2001, Kiesecker et al. 2001b, Romansic et al. 2006, 2007) suggest that multiple life history stages of amphibians are susceptible to these pathogens. However, the life stage most susceptible to water molds may differ between amphibian species. Additionally, patterns of mortality may depend on the species of water molds present. Our results highlight the need to investigate possible effects on multiple life stages of the host species when evaluating the potential effects of pathogens on host populations. If only one life history stage of the host species is considered, the effect of a pathogen on the host population may be underestimated.

Although infectious diseases are a significant threat to global biodiversity (Daszak et al. 2000, Harvell et al. 2002, Altizer et al. 2003), the degree to which infectious diseases have contributed to the current biodiversity crisis is highly uncertain (Smith et al. 2006). Thus, there is an urgent need for better understanding of disease ecology so that the risk of disease-related biodiversity losses may be more accurately measured, and species and communities at risk of disease-related losses

Table 2. *Pseudacris regilla*, *Rana cascadae*, *Ambystoma macrodactylum* and *R. aurora*. Results of examinations for hyphae. Larvae that were completely eaten before inspection are not included. na: no carcasses

Species	Treatment	No./total no. (%) dead ind. with hyphae	Obs. of hyphae on a live individual
Expt 1			
<i>P. regilla</i>	Control	na	0
	<i>S. ferax</i>	0/3 ^a (0)	13 ^b
<i>R. cascadae</i>	Control	na	0
	<i>S. ferax</i>	na	0
Expt 2			
<i>A. macrodactylum</i>	Control	na	0
	<i>S. ferax</i>	na	0
Expt 3			
<i>R. aurora</i>	Control	na	0
	<i>S. ferax</i>	16/17 ^c (94)	12 ^d

^aAll 3 carcasses were partially eaten
^bAt the end of the experiment, one live individual had hyphae
^cOne carcass with hyphae was partially eaten
^dAt the end of the experiment, hyphae were not found on any live larvae

may be identified (Smith et al. 2006). There is evidence that anthropogenic environmental changes, pathogen pollution, and introduction of non-native pathogens by humans have already increased the impact of infectious disease in some species (e.g. Anagnostakis 1987, Conrad et al. 2005, Johnson et al. 2007). This evidence, combined with the prediction that climate change will cause an overall increase in the impact of infectious diseases on natural communities (Harvell et al. 2002), underscores the importance of making meaningful progress in our understanding of the ecology of infectious disease.

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