

Predation changes the shape of thermal performance curves for population growth rate

Thomas M. LUHRING^{*}, John P. DELONG

School of Biological Sciences, University of Nebraska, Lincoln, Nebraska 68588, USA

*Address correspondence to Thomas M. LUHRING. E-mail: tomluhring@gmail.com

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Abstract

Ectotherms generally demonstrate non-linear changes in performance (e.g., movement speed, individual growth, population growth) as a function of temperature that are characterized by thermal performance curves (TPC). Predation risk elicits phenotypic and behavioral changes that likewise impact performance measures. We tested whether exposure to predation *Orthocyclops modestus* impacts the maximum population growth rate (r_{\max}) TPC of the protist *Paramecium aurelia*. We fit predator and non-predator exposed *P. aurelia* population growth rates to a function previously shown to best describe *Paramecium* population growth rate TPC's (Lactin-2) and compared subsequent parameter estimates between curves. For *Paramecium* exposed to predation risk, maximum population growth increased more rapidly as temperatures rose and decreased more rapidly as temperatures fell compared to the initial temperature. The area under each TPC curve remained approximately the same, consistent with the idea of a trade-off in performance across temperatures. Our results indicate TPCs are flexible given variation in food-web context and that trophic interactions may play an important role in shaping TPCs. Furthermore, this and other studies illustrate the need for a mechanistic model of TPCs with parameters tied to biologically meaningful properties.

Keywords: paramecium, phenotypic plasticity, predation, reaction norm, temperature, thermal performance curve

Ectotherm biology is dominated by the effects of temperature-dependent processes (Brown et al. 2004; Kingsolver 2009). Likewise, predation risk has a strong influence on organism physiology (e.g., Bernard 2004; Handelsman et al. 2013), behavior (Lima and Dill 1990; Sih 1994), and life history (Riessen 1999). Behavioral and phenotypic plasticity enables organisms to optimize fitness by matching traits to environmental pressures such as temperature and predation risk (Lima and Dill 1990; Stearns 1992; Sih 1994; Bernard 2004; Handelsman et al. 2013). How species respond to changing trophic interactions and temperature regimes influence their ability to persist through environmental perturbations such as those caused by invasive species, habitat loss, and a warming and more variable climate (Vasseur et al. 2014; Walsh et al. 2015). Organisms attend to and allocate resources among traits in order to optimize fitness in response to several simultaneously varying environmental pressures (Stearns 1992). However, most studies investigate plasticity in response to a single varying type of environmental pressure (Miner et al. 2015).

Temperature-dependent performance, particularly of population growth rate, is often used to predict the effects of climate change (mainly changing temperature means and variation) on the persistence of species (Deutsch et al. 2008; Bozinovic et al. 2011; Clusella-Trullas et al. 2011; Kingsolver et al. 2013; Vasseur et al. 2014). Temperature-dependent performance can be characterized by thermal performance curves (TPCs) that describe variation in a performance metric across a range of temperatures (Scheiner 2002; Angilletta 2009; Kingsolver 2009). These curves generally adhere to a shape characterized by an exponential increase at cooler temperatures and a peak at an optimal temperature (T_{opt}) where performance is maximized. Above the T_{opt} there is a steep decrease in performance until it reaches some critical upper temperature (CT_{max}) (Scheiner 2002; Angilletta 2009; Kingsolver 2009). While this general shape is repeated across taxa, there is considerable variation in TPC shape within and among species attributable to genetic, latitudinal, or acclimation effects (Izem and Kingsolver 2005; Kingsolver 2009; Angilletta 2009; Clusella-Trullas et al. 2011). Predation risk also shapes the TPCs of individual-level performance metrics (e.g., burst speed, relative growth rate; Culler et al. 2014; Katzenberger et al. 2014), but whether and how TPCs of population-level metrics (e.g., population growth rate) change in response to predation risk is unknown. If population-level TPCs do change shape in response to predation, then TPCs that are measured without predation risk (the majority of those reported to date) will be less useful for predicting the consequences of future temperature change than currently thought.

At any given temperature, mortality risk (e.g., predation) can alter individual-level behavior or physiology in ways that cause a concomitant change in r_m (Cole 1954; Lima 1998; Riessen 1999). Furthermore, the effects of these pressures on r_m differ with respect to size-dependent selection on body size and death rate. Earlier age at first reproduction reduces the likelihood of dying prior to first reproduction when mortality is size-independent and constant (Brown and Sibly 2006) and size does not confer additional benefits (e.g., Peters 1983; Kingsolver and Pfennig 2004; Luhring and Holdo 2015). This is seen in size-dependent predation risk in *Daphnia* where cues from gape-limited predators induced a larger body size through delayed age of first reproduction (which would decrease r_m), but cues from predators with a preference for larger prey items induced a smaller size at maturity and earlier age of first reproduction (which would increase r_m) (Riessen 1999). Since TPCs connect performance at one temperature to performance at another temperature, predation-induced changes in performance at one temperature may come with a cost to performance at another temperature, generating a mechanism by which overall TPC shape might change under predation risk.

To test whether predation pressure can change the performance of temperature-dependent population-level metrics, we investigated the effects of exposure to predation on r_m TPCs in *Paramecium aurelia*. Cyclopoid copepods are generalist predators that apply substantial predation pressure on protists and other zooplankton in freshwater ponds (Kalinowski and

DeLong 2015). *Orthocyclops modestus* co-occurs with *P. aurelia* in freshwater ponds and forages at a high rate on many similar-sized protists (Novich et al 2014). Because all size classes of *P. aurelia* are susceptible to predation by copepods, we expect that their TPCs will emphasize accelerated growth rates at temperatures closest to their natal temperature. However, it is unclear whether such increases, if detected, would occur to the same degree across all temperatures, or whether increase in growth rate at some temperatures would be accompanied by decreases in growth rate at other temperatures (e.g., a trade-off). Because changing physiology and behavior in response to predation risk are generally governed by trade-offs that incur costs (Lima and Dill 1990; Sih 1994), we predict that although predation will change the shape of r_m TPCs, the area under the TPC curves will be constrained. Here we test directly whether predation risk alters the temperature dependence of growth rates in *P. aurelia* by estimating the growth rate TPCs for populations with and without exposure to predation risk.

Materials and Methods

Study species rearing and maintenance

We collected the ciliate *P. aurelia* and the copepod *O. modestus* (hereafter cyclops) from a freshwater pond at the Spring Creek Prairie Audubon Center in Southeastern Nebraska (Novich et al. 2014). *Paramecium aurelia* was isolated from pond water samples and grown in media made from filtered and autoclaved local pond water and liquid protozoa medium from Carolina Biological Supply (Burlington, NC, USA) at a ratio of 1:9 liquid protozoa medium to water. The media was autoclaved and then bacterized with a mix of unidentified bacteria from the local source by passing local water through a 5- μm syringe filter onto an agar plate, allowing the bacterial community to grow for a few days, and then inoculating the protozoan medium with these bacteria. We kept cyclops in filtered pond water in Petri dishes and fed them with a range of protist prey. We kept both cultures at room temperature, $\sim 24^\circ\text{C}$.

Predation Trials

We initiated six replicate *P. aurelia* populations in 60-mm Petri dishes. Three of these were predation treatments and three were controls without predation. Population densities were ~ 740 (± 98 SD) cells mL^{-1} to start, estimated by counts of three separate 0.2-mL samples from the stock culture. We rinsed cyclops in sterile pond water, added three cyclops to each of the three predation dishes, and allowed them to predate the *P. aurelia* for four days. We added the equivalent amount of rinse water to the no-predation dishes to control for any potential differences generated by rare bacteria or other microorganisms that might have entered the cultures through the rinse water. We allowed the *P. aurelia* in the control dishes to grow without predation risk over this same four-day period. After four days, mean *P. aurelia* densities had doubled in the control dishes ($1,856 \pm 718$ cells mL^{-1}) but remained approximately constant in the predation dishes (790 ± 217 cells mL^{-1}), indicating substantial predation on *P. aurelia* by cyclops.

Post-predation growth rates

We measured maximum population growth rates by allowing four cells to grow overnight (20–23 hours) in 35-mm Petri dishes with 2-mL of bacterized protozoan medium. No predators were used in these trials – we were measuring the changes in population growth rate induced by ~ 8 generations of growth under predation risk. We collected individual cells with

pipettes, and while watching through a stereoscopic microscope, placed four cells into each dish. We visually confirmed that all four cells made it into each dish alive. By starting these trials with very low densities and high resource levels, we ensured that the observed growth rates were close to maximal. Three replicate growth trials for the predation and no-predation treatments were conducted at each of nine temperatures (12, 16, 20, 24, 27, 30, 33, 36, 40°C), for a total of 54 trials. The following day, counts were made of the total number of cells in each dish. Growth rates r were calculated using the formula for exponential growth: $r_m = \frac{\ln \frac{N_{t+1}}{N_t}}{t}$, where N_t is population size at time t , and t is the actual time elapsed during each trial.

Curve-fitting and statistical analyses

We used the Lactin-2 function (eq. 1) to fit temperature-dependent r across temperature treatments because it allows for negative values at both cold and hot extremes and provides a relatively good fit for r_m TPCs (Lactin et al. 1995; Krenke et al. 2011):

$$\text{Eq 1. } \exp(\rho T) - \exp\left(\rho T_{Max} - \frac{T_{Max} - T}{\Delta T}\right) + \lambda$$

In this function, T_{max} changes the temperature at which the TPC begins to decelerate and approach the x -axis at the higher end, ΔT is a reference temperature, and λ and ρ are constants that determine the steepness of the rising portion of the curve and the height of the overall curve, respectively (see S1 for effect of changes to each fit parameter). Because the shape and x -intercept of the fit is dependent on multiple parameters, T_{max} is not the same as CT_{max} (critical upper temperature where r_m is 0). We fit the Lactin-2 curve to the r_m values across temperatures (constrained to the temperature range used in the experiment) for each treatment (predator or no predator) using the curve fitting tool in MatLab (R2015b) to produce estimates and 95% confidence estimates of the four parameters comprising the curve. Differences in parameter values between treatments were determined by comparing 95% confidence bounds. We calculated the area under the curve for the mean, upper and lower 95% confidence limits of each fitted curve to compare curve areas between treatments.

Results

TPC parameters and area under the curve

Both treatments demonstrated negative r_m at extreme temperatures (12°C and 40°C) with no survival at 40°C (Figure 1). Data from 40°C treatments were dropped from analysis because r_m cannot be calculated when abundances are zero. The Lactin-2 function fit the shape of predator and non-predator treatment TPCs well and captured the negative values at both temperature extremes (non-predator adjusted $R^2=0.90$; predator adjusted $R^2=0.92$). All four parameter estimates differed in either variance (ρ and ΔT) or means (T_{Max} and λ) between the predator and non-predator treatments (Figure 2; Table 1). The areas under the non-predator and predator TPCs were nearly identical for each mean, upper and lower prediction bound (Fig. 3). The parameter means that differed between the treatments were associated with vertical (λ) and horizontal shift (T_{Max}) of the curve. *Paramecium aurelia* exposed to predators demonstrated a subsequently higher r_m across a span of 9.5°C (21.9-

31.4°C) compared to non-predator treatments. However, predator exposed *P. aurelia* had poorer performance at cooler temperatures (< 21.9°C), a 3.6°C narrower thermal breadth (18.7°C vs 22.4°C), and a 2.2°C cooler T_{OPT} (30.7°C) than the non-predator control (32.9°C)(Table 2).

Discussion

Our study demonstrated a multi-generational effect of predation risk on maximum population growth rate (r_m) thermal performance curves (TPCs) in *P. aurelia*. The overall shape of *P. aurelia* TPCs changed in response to predation (Figure 1) indicating that the effect of predation on subsequent performance is temperature dependent. Furthermore, the lack of difference in total TPC area between treatments supports the idea that improving performance at one area of the TPC comes at the cost of performance in another (i.e., trade-off) (Kingsolver2009).

Paramecium aurelia appear to increase r_m in response to predation risk from cyclops, as would be consistent with animals reproducing earlier (in this case, dividing at a smaller size) in response to increased size-independent mortality risk. However, the ability to increase r_m was temperature-dependent with predator-exposed *P. aurelia* showing an increased r_m across a 9.5°C range of temperatures close to the rearing temperature and a slightly elevated CT_{max} (Table 2). This enhanced performance at temperatures close to the rearing temperature was accompanied by a 2.2°C cooler T_{OPT} and 3.7°C narrower breadth of temperatures over which r_m was positive in post-predation trials (Table 2). These changes are consistent with a trade-off between performance at a current temperature and ability to adjust to a broader range of temperatures (e.g., generalist-specialist trade-off; Izem and Kingsolver 2005; Kingsolver 2009). Furthermore, as predicted by the “hotter is better” rule (Kingsolver and Huey 2008) the decreased T_{opt} seen in the predator-exposed population, was accompanied by a decreased overall maximum performance P_{max} (Table 2). *Paramecium* experiencing elevated mortality risk from predation can increase their r_m at current temperatures, however this appears to come at the cost of a decreased ability to survive cooler temperatures and a decreased maximum population growth rate (P_{max}).

The shape and bounds of TPCs are important for predicting the effects of climate change on population persistence (Deutsch et al. 2008; Martin and Huey 2008; Vasseur et al. 2014). For example, the location of an organism’s T_{opt} relative to the current environmental temperature, known as the thermal safety margin (TSM; Deutsch et al. 2008), is thought to determine the impact of warming on performance. Animals with larger TSM’s are predicted to benefit more from increased warming than animals with smaller TSM’s because of their closer proximity to the rapid decline in performance above the T_{opt} , assuming that the decline is equally precipitous in both cases. Here, predation pressure reduced T_{opt} by 2.2°C and thus reduced the calculated TSM of our study species when compared to the non-predator control. However, the predator-exposed population TPC showed a less precipitous decline in performance above the T_{opt} compared to that of the non-predator treatment. This predator-induced change in TPC shape beyond the T_{opt} complicates the use TSM’s for predicting vulnerability to climate change.

Projected increases in both climate variability and the occurrence of extreme weather events (Katz et al. 1992; Easterling et al. 2000; Overpeck and Udall 2010) could pose problems for organisms with narrow thermal tolerance breadth. In our study, predation narrowed the breadth of temperatures over which positive population growth occurred by 3.7°C, mostly through an increased vulnerability to colder temperatures (decreased cold tolerance of 4.4°C). Models using TPCs to predict population vulnerability to climate change generally focus on changes to the upper limits of TPCs and in some cases,

do not account for negative population growth at colder temperatures (e.g., Vasseur et al. 2014). In our study, predation risk induced an apparent trade-off between increased performance at current or warmer temperatures and a decreased survivability at colder temperatures. Thermal performance curves derived for *P. aurelia* populations raised without predation pressures would thus potentially overestimate the ability of this species to tolerate colder temperatures in natural food webs.

The temperature-dependent nature of biological processes makes TPCs important and widely used tools for predicting effects of climate change on species persistence (Deutsch et al. 2008; Kingsolver et al. 2013; Vasseur et al. 2014). However, an assumption of using TPCs characterized outside of the context of a functioning food web is that recent trophic interactions do not influence their shape. Our results suggest that such assumptions may be violated in systems where predation pressure may induce trade-offs that reshape TPCs. This effect may be weaker in performance measures such as burst speed in field-collected, long-lived animals, which are not far removed from the context of their environment. In contrast, measuring r_m as a response to temperature generally requires the use of several replicate populations of animals with short generation times raised at different controlled temperatures in a laboratory setting. The controlled nature of the lab may be necessary to get measurements of fitness that relate strongly to population persistence such as r_m , but may also increase the chance of creating a TPC that does not necessarily match that of animals *in situ*.

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Table 1. Curve fit parameters for temperature-dependent growth rate (r_m) in *P. aurelia* raised with or without predators. Confidence bounds (CB) are presented from fitting a Lactin-2 function to the data for predator and non-predator treatments. Adjusted R^2 for non-predator (0.90) and predator (0.92) treatments indicated a good fit for both curves.

Parameter	Non-Predator ($\mu \pm 95\%$ CB)	Predator ($\mu \pm 95\%$ CB)
ρ	0.048 (0.006)	0.084 (0.092)
ΔT	1.22 (1.15)	6.37 (8.57)
T_{Max}	36.61 (0.61)	39.25 (1.39)
λ	-1.93 (0.40)	-3.59 (0.46)

Table 2. TPC parameters for temperature-dependent growth rate (r_m) in *P. aurelia* raised with or without predators. Parameters were solved from the equation of the TPC fitted to the predator and non-predator treatments. Lower critical temperature - CT_{min} ; upper critical temperature - CT_{max} ; optimal temperature - T_{opt} ; maximum r_m at T_{opt} - P_{max} ; Thermal Performance Breadth – TPB; Thermal Safety Margin - TSM. Effect of predation shown as difference between non-predator and predator treatments.

Parameter	Non-Predator	Predator	Effect of Predation
CT_{min}	13.7	18.1	+4.4
CT_{max}	36.1	36.8	+0.7
T_{opt}	32.9	30.7	-2.2
P_{max}	2.6	2.5	-0.1
TPB	22.4	18.7	-3.7
TSM	8.9	6.7	-2.2

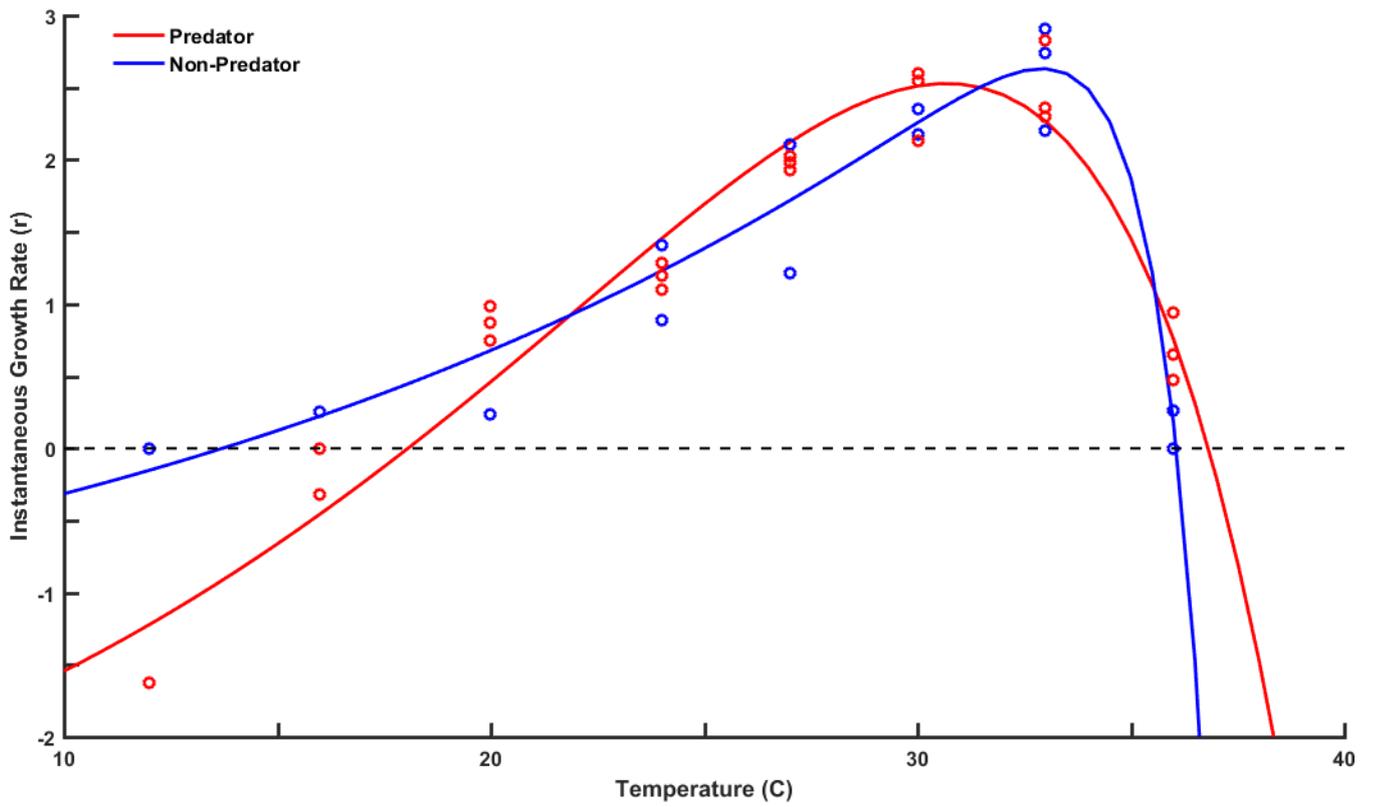


Figure 1 *Paramecium aurelia* TPC's for instantaneous growth rate (r_{max}). *P. aurelia* were raised at 24°C in the absence (blue) or presence of a predator (red) before being raised at a series of temperatures. Curves were fitted to a Lactin-2 function.

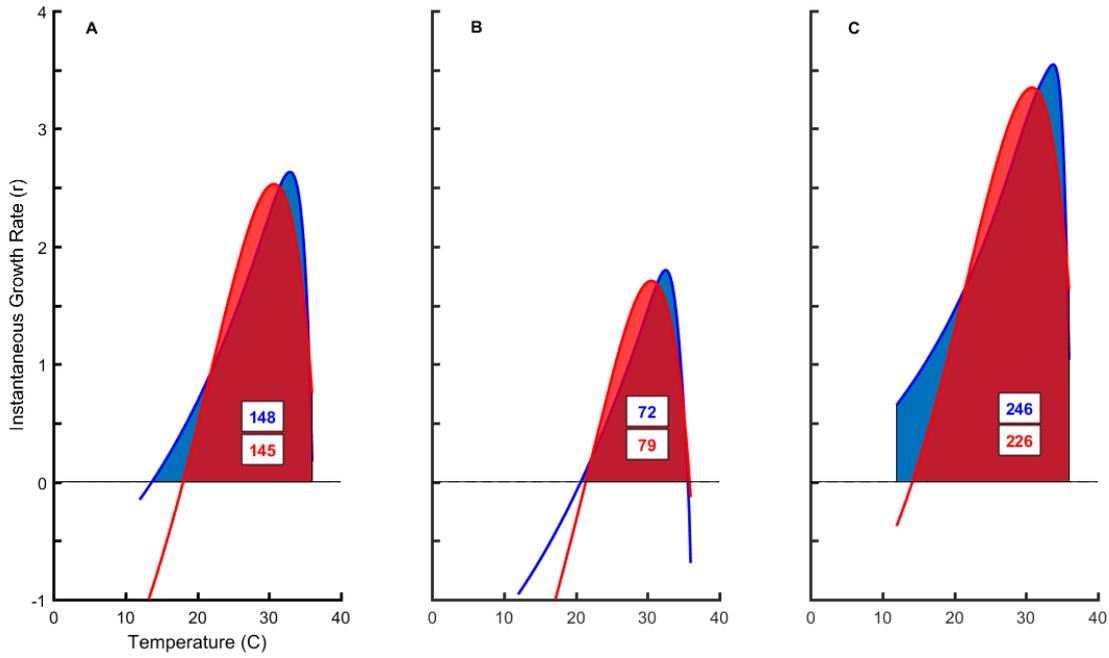
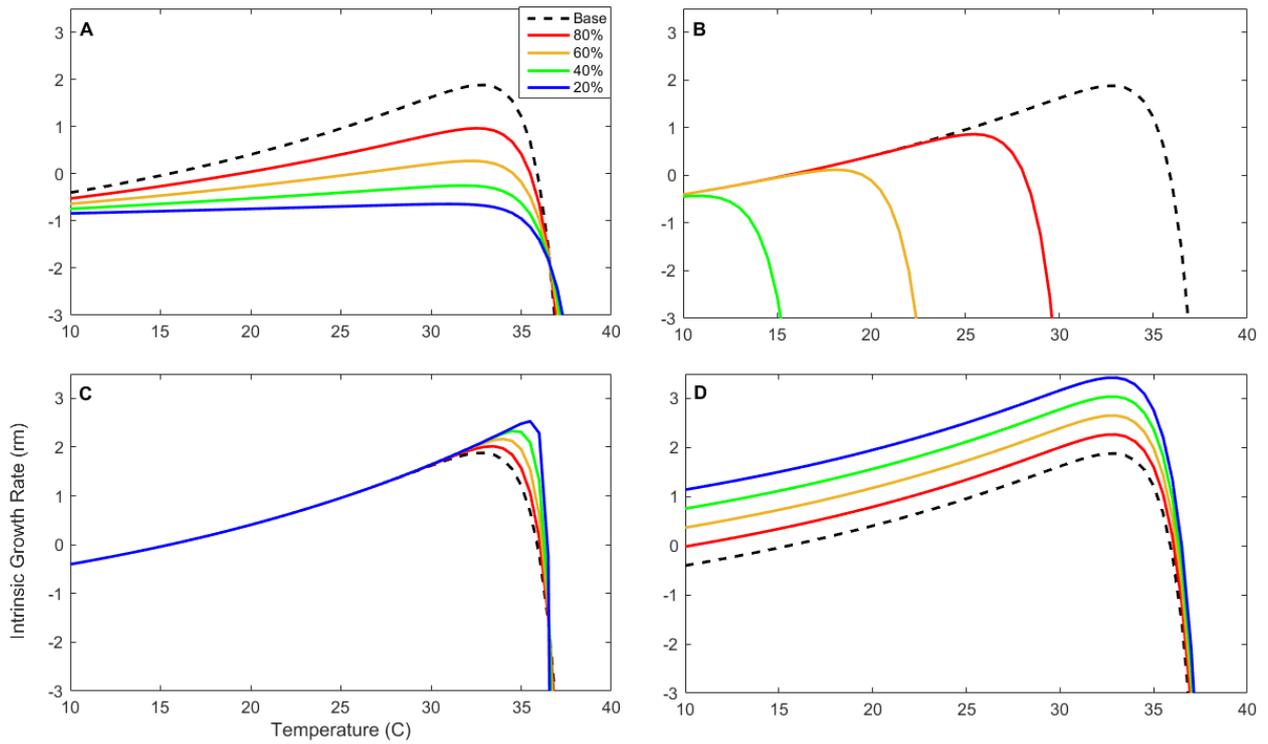


Figure 2 Areas under mean, lower and upper 95% prediction limits of predator and non-predator treatment TPCs. Lactin-2 fitted TPC curves for *P. aurelia* TPC's after being raised in the absence (blue) and presence of a predator (red). Areas under the curve are denoted (within boxes) for fitted model means (A), lower 95% (B), and upper 95% prediction bounds (C).

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Supplementary Figure 1 Lactin-2 parameter effects on TPCs

Lactin-2 TPCs for changes in each parameter: ρ (A), T_{Max} (B), ΔT (C), λ (D), compared to the non-predator treatment baseline (dashed line).