1	Diminished warming tolerance and plasticity in low latitude populations of a marine gastropod
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7	Lay Summary
8	The physiology of populations within species can determine sensitivity to climate change. Experiments
9	on marine snails indicate that southern populations are most susceptible to warming because they have
10	minimal physiological flexibility and are most threatened by extreme heat events. Management of this
11	species depends on the specific population in question.
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Abstract

23 Models of species response to climate change often assume that physiological traits are invariant 24 across populations. Neglecting potential intraspecific variation may overlook the possibility that some 25 populations are more resilient or susceptible than others, creating inaccurate predictions of climate 26 impacts. In addition, phenotypic plasticity can contribute to trait variation and may mediate sensitivity to 27 climate. Quantifying such forms of intraspecific variation can improve our understanding of how climate 28 can affect ecologically important species, such as invasive predators. Here, we quantified thermal 29 performance (tolerance, acclimation capacity, developmental traits) across seven populations of the 30 predatory marine snail Urosalpinx cinerea from native Atlantic and non-native Pacific coast populations 31 in the United States. Using common garden experiments, we assessed the effects of source population and developmental acclimation on thermal tolerance and developmental traits of F1 snails. We then estimated 32 33 climate sensitivity by calculating warming tolerance (thermal tolerance – habitat temperature), using field 34 environmental data. We report that low latitude populations had greater thermal tolerance than their high latitude counterparts. However, these same low latitude populations exhibited decreased thermal tolerance 35 when exposed to environmentally realistic higher acclimation temperatures. Low latitude native 36 37 populations had the greatest climate sensitivity (habitat temperatures near thermal limits). In contrast, invasive Pacific snails had the lowest climate sensitivity, suggesting that these populations are likely to 38 persist and drive negative impacts on native biodiversity. Developmental rate significantly increased in 39 40 embryos sourced from populations with greater habitat temperature, but had variable effects on clutch 41 size and hatching success. Thus, warming can produce widely divergent responses within the same 42 species, resulting in enhanced impacts in the non-native range and extirpation in the native range. 43 Broadly, our results highlight how intraspecific variation can alter management decisions, as this may clarify whether management efforts should be focused on many or only a few populations. 44

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Introduction

48 Understanding the sensitivity of species to climate change is a primary aim of global change ecology (Calosi et al., 2008; Williams et al., 2008; Bennett et al., 2019). Ecological forecasts are a suite 49 50 of modeling tools that can aid conservation practitioners in determining species sensitivity to climate 51 change by correlating occupied distribution environments or known physiological limits with predictions of future climate scenarios (Pearson and Dawson, 2003; Helmuth, 2009; Chown et al., 2010; Cacciapaglia 52 53 and van Woesik, 2018). In a conservation and management context, ecological forecasts can be used to 54 identify species at risk and prioritize efforts and management actions on species and ecosystems of 55 concern (Payne et al., 2017; Tulloch et al., 2020). However, these models often use physiological 56 measures from a single population to infer the capacity of a species to respond to environmental change 57 (Pearman et al., 2010; D'Amen et al., 2013; Valladares et al., 2014; Lecocq et al., 2019) and implicitly 58 assume that physiological niches are homogenous across populations within a species (Peterson, 1999, 59 2011; Bennett et al., 2019). However, populations within species often exhibit physiological variation that 60 reflects heterogeneity in environmental conditions and potential local adaptation (Moran *et al.*, 2016; 61 Peterson et al., 2019). Ignoring the potential for such locally-adapted variation greatly risks under- or 62 over-estimating species sensitivity to climate change (Pearman et al., 2010; Valladares et al., 2014; Cacciapaglia and van Woesik, 2018). For example, populations of widely distributed species can differ in 63 64 thermal tolerance by up to 1.5-3.8°C (e.g., Fangue et al., 2006; Pereira et al., 2017). In contrast, thermal 65 tolerance may be invariant across a species range, a pattern that is described as niche conservatism (Lee 66 and Boulding, 2010; Pearman et al., 2010; Gaitán-Espitia et al., 2017). If populations are niche 67 conserved, then modeling a species as a single unit is appropriate. However, the management implications of assuming niche conservatism or local adaptation can be starkly divergent; when modelled as having 68 69 homogenous physiology throughout its range, a Porites coral species was expected to increase its range 70 by 5-6% by 2100, while when modelled as five distinct populations the range was forecasted to decrease 71 by 50% (Cacciapaglia and van Woesik, 2018). Thus, intraspecific variation in thermal performance may

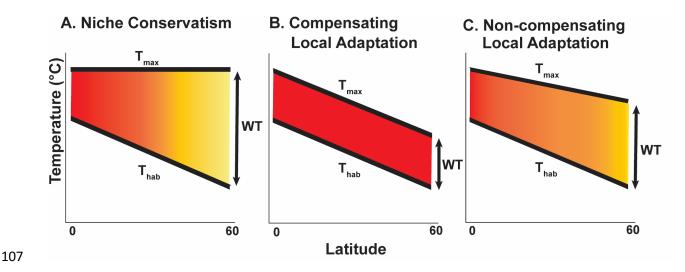
be crucial to understanding species sensitivity to climate change, but our understanding of mechanismsunderlying such variation remains incomplete.

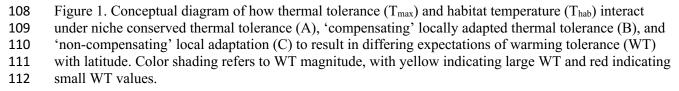
74 Climate sensitivity may also be mediated by phenotypic plasticity. Acclimation is one form of 75 plasticity that is defined as within generational phenotypic change in response to an altered environmental 76 change and may allow an organism to rapidly adjust physiology to changing environmental conditions 77 (Seebacher et al., 2012; Beaman et al., 2016). For example, higher acclimation temperatures tend to 78 increase thermal tolerance, primarily due coordinated molecular adjustments such as increased heat shock 79 protein expression to maintain or regain homeostasis (Hofmann, 1999; Basu et al., 2002; Guy et al., 80 2008). The majority of these studies examine plasticity within a focal life stage (Marshall, 2008; Moore 81 and Martin, 2019). However, organismal life stages do not act as 'firewalls' past which the effects of 82 thermal challenge cannot penetrate (Marshall and Morgan, 2011). The effects of marine climate change 83 will impact all life stages of marine organisms and exposure to thermal stress in one life stage can result 84 in latent or carry-over effects to future life stages (Pechenik, 2006; Hettinger et al., 2012). Developmental 85 acclimation should increase adult thermal tolerance to a point, beyond which we expect a reduction in 86 adult tolerance when acclimation temperature exceeds the thermal optima of organismal performance 87 (Overgaard et al., 2011; Scharf et al., 2015; Truebano et al., 2018). Therefore, identifying the potential 88 effects of developmental acclimation are critical to understanding actual organismal reactions in later life 89 stages.

90 Even though plastic trait expression is often linked to environmental exposure, the extent of 91 plasticity capacity itself can be adapted to local conditions (De Jong, 2005; Valladares et al., 2014). 92 Under the latitudinal variability hypothesis, which predicts how thermal phenotypic plasticity might vary 93 between latitudinally separate populations, high-latitude but non-polar populations should have higher plasticity in response to seasonally variable temperatures (Bozinovic et al., 2011; Gunderson and 94 95 Stillman, 2015; Barria and Bacigalupe, 2017). In contrast, tropical and polar species that experience 96 minimal seasonality are expected to have lower plasticity in response to limited environmental fluctuations (Tewksbury et al., 2008; Overgaard et al., 2011; Peck et al., 2014). It has also been suggested 97

98 that lower plasticity in warm adapted populations may reflect a trade-off between plasticity and greater overall tolerance (trade-off hypothesis; Stillman, 2003; Magozzi and Calosi, 2015; Sasaki and Dam, 99 2019; Heerwaarden and Kellermann, 2020). Plasticity can buffer species' susceptibility to warming 100 101 temperatures, and thus it is important to quantify this trait in order to fully assess warming sensitivity 102 (Palumbi et al., 2014). However, among species variation in plasticity means warming sensitivity will 103 also vary, requiring study on a species-by-species basis to accurately understand warming sensitivity 104 (Seebacher et al., 2012). Considering the role of plasticity, in addition to potential local adaptation, are 105 critical to determining organismal susceptibility to thermal stress (Valladares *et al.*, 2014).

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113 Understanding geographic variation in thermal performance is central to identifying which

114 populations may be at the greatest risk of extinction caused by climate change. Both across species and

115 populations, evidence suggests that upper thermal tolerances increase with decreasing latitude (e.g.

116 Stillman and Somero, 2000; Sgrò et al., 2010; Zippay and Hofmann, 2010; Sunday et al., 2011; Pereira et

117 *al.*, 2017; Jensen *et al.*, 2019; Sasaki and Dam, 2019). However, quantifying thermal tolerance alone does

118 not reveal climate sensitivity, as it does not factor in the 'environmental distance' between thermal

119 tolerance and the *in situ* temperature regime. It is therefore necessary to integrate habitat temperature with 120 organismal tolerance. An organism's 'warming tolerance' (WT) quantifies this buffer by calculating the difference between thermal tolerance and habitat temperature (e.g., mean annual temperature; Deutsch et 121 122 al., 2008). In the absence of rapid thermal adaptation, populations at greatest risk of warming are those 123 with diminished warming tolerance (Deutsch et al., 2008; Bennett et al., 2019). In populations with 124 invariant thermal limits (niche conservatism), warming tolerance may be greater at high latitudes, as the 125 difference between habitat temperature and the conserved thermal tolerance will be large (Fig. 1A; 126 Bennett et al., 2019). In contrast, low latitude populations would be most sensitive because of the small 127 difference between habitat temperature and thermal tolerance, assuming habitat temperatures decrease 128 more or less linearly from the equator to the poles (Tewksbury et al., 2008; Diamond et al., 2012; Bennett et al., 2019; Pinsky et al., 2019). However, if thermal tolerance varies across populations ('compensating' 129 local adaptation), warming tolerance may actually be similar across populations, suggesting sensitivity 130 131 across the entire species range (if warming tolerance is low) or resilience to changing temperatures (if 132 warming tolerance is high; Fig. 1B; Bennett et al., 2019). Finally, local adaptation in thermal tolerance 133 may exist, but may not track perfectly with habitat temperature ('non-compensating' local adaptation), 134 resulting in greater sensitivity to climate warming in populations with greater thermal exposure (Fig. 1C). 135 Thus, integrating intraspecific measures of physiological performance with environmental data is a 136 promising approach that can clarify population sensitivity to climate change. For conservation 137 stakeholders, this integration can better inform whether management needs to be focused on a few 138 sensitive populations, many populations throughout a species range, or none. 139 In a management and conservation context, knowledge of physiological performance can also 140 clarify our understanding of impacts of invasive species under climate change (Zerebecki and Sorte,

141 2011; Sorte *et al.*, 2013; Lennox *et al.*, 2015). Greater thermal tolerance breadths and plasticity are traits

that can contribute to the success of invasive species, particularly in the face of climate change (Chown *et*

143 *al.*, 2007; Slabber *et al.*, 2007; Sorte *et al.*, 2010; Zerebecki and Sorte, 2011; Seebacher *et al.*, 2012;

144 Kelley, 2014). These adaptations may allow invasive species to survive challenging transport conditions

145 and to rapidly colonize habitats with thermal conditions that differ from their native range (Diez et al., 146 2012). These same traits are also predicted to confer climate resilience to invasive species as habitats experience elevated and increasingly variable temperatures (Dukes and Mooney, 1999; Stachowicz et al., 147 148 2002; Diez et al., 2012; Sorte et al., 2013). Forecasting the impacts of invasive species under climate 149 warming may be informed by knowledge of thermal physiology in both the native and invasive ranges 150 because adaptation in the native range provides the standing genetic material that founds invasive 151 populations. For invasive populations, thermal tolerance and plasticity may be locally adapted to novel 152 environments, even those that are warmer or colder than their native range environment (Beaumont et al., 153 2009; Griffith et al., 2014; Tepolt and Somero, 2014; Wesselmann et al., 2020). Altogether, there exists a 154 range of possible climate sensitivities of invasive populations that may not be accurately described by native range thermal physiology. Neglecting the potential for novel trait performance in invasive 155 populations can decrease the accuracy of ecological forecasts to climate change that are solely based on 156 157 the native range (Broennimann et al., 2007; Fitzpatrick et al., 2007; Loo et al., 2007; Beaumont et al., 158 2009). Thus, evaluating the range of thermal physiology across native and invasive populations of single 159 species can shed light on the range of current adaptations within a species and thus clarify the extent of 160 current sensitivity, as well as the potential for future evolutionary adaptation to climate change (Beaumont et al., 2009; Henkel et al., 2009; Hill et al., 2013; Wesselmann et al., 2020). 161

162 To address the roles of local adaptation and plasticity in determining thermal sensitivities across 163 native and invasive ranges, we quantified intraspecific variation in thermal performance of invasive and 164 native populations of an ecologically important predatory marine snail (Atlantic oyster drill, Urosalpinx 165 *cinerea*). We used split-brood common garden experiments to assess thermal performance of laboratory 166 reared F1 juveniles sourced from native and invasive populations across a latitudinal gradient on the Atlantic $(32.7^{\circ} - 43.1^{\circ} \text{ N})$ and Pacific $(38.1^{\circ} - 40.8^{\circ} \text{ N})$ coasts of the United States, respectively. Our 167 specific objectives were to: 1) determine if variation in thermal tolerance and developmental traits occurs 168 169 among native and invasive populations, 2) quantify plasticity in thermal tolerance and developmental

170 traits by manipulating temperature during embryonic incubation, and 3) estimate climate sensitivity of each population using warming tolerance (Deutsch et al., 2008). We hypothesized that 1) thermal 171 tolerance would increase with environmental temperature, thereby suggesting local adaptation, 2) 172 173 elevated acclimation temperature during development would result in greater juvenile thermal tolerance, 174 and 3) plasticity would be highest in cold origin populations. Because latitude itself is not a perfect 175 predictor of the actual environmental temperatures experienced by populations, particularly across coastal 176 latitudinal gradients, we also evaluated the correlation between a suite of environmental metrics (e.g., 177 maximum and mean temperature) and thermal and warming tolerance (Helmuth, 2009; Kuo and Sanford, 178 2009). Our broader goal was to quantify intraspecific thermal performance across a species' native and 179 invasive ranges to determine what populations are likely most sensitive to climate warming, and therefore identify which populations of *Urosalpinx* are likely to persist in the long term without management 180 181 intervention.

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METHODS

184 Species Selection

185 We used the snail U. cinerea (hereafter Urosalpinx) as our focal species because of its limited dispersal 186 that drives a high potential for local adaptation, its wide range across latitude and thermal regimes, and its tractability in the egg and juvenile life stages (Cheng et al., 2017). Urosalpinx undergoes direct 187 188 development, laying benthic egg cases that each contain 4-16 embryos that develop for 26-56 days after 189 which they emerge as hatchlings (Carriker, 1955). Because of this direct development, dispersal and gene 190 flow are likely limited among populations, suggesting a high potential for local adaptation (Kawecki and 191 Ebert, 2004). Further, we sampled populations from both the invaded and native ranges of Urosalpinx 192 with the goal of understanding if trait performance differs between invaded and native populations under 193 different thermal regimes (Zerebecki and Sorte, 2011). Urosalpinx is native on the Atlantic coast of North America from south Florida to Massachusetts and cryptogenic (of unknown origin) north to Nova Scotia 194 (Fofonoff et al., 2020). In the late 1800s, Urosalpinx was introduced to multiple locations on the Pacific 195

196 coast of North America, ranging from San Francisco Bay north to Puget Sound, via importation of

- 197 Eastern oysters (*Crassostrea virginica*; (Carriker, 1955; Fofonoff *et al.*, 2020). The high biomass (1.7
- 198 million kg) and diverse origins of oysters transported to these Pacific sites (Hoos *et al.*, 2010) indicates
- initial *Urosalpinx* populations were likely large, suggesting limited founder effects. In the invasive range,
- 200 Urosalpinx can virtually eliminate native oysters and other native species via predation (Carriker, 1955;
- 201 Kimbro et al., 2009; Cheng and Grosholz, 2016).

202 Broodstock Collection

203 We examined physiological performance of F1 offspring in order to ensure a common garden 204 environment for the entire embryonic and juvenile life phases. This approach does not fully account for 205 the possibility of maternal or transgenerational effects but is a reasonable starting point for assessing 206 intraspecific patterns of thermal performance. To produce F1 offspring for experimentation, we collected broodstock adult Urosalpinx from seven sites, five from the Atlantic and two from the Pacific that 207 encompassed a wide range of their latitudinal distribution (Fig. 2). All collections were conducted from 208 15 March - 9 June, 2019. We chose collection sites to be within 15 km of in situ environmental data 209 210 loggers (e.g. National Data Buoy Center, National Estuarine Reserve System, NOAA Ocean Observing System, Table S1). At each site, we hand-collected at least 30 adult male and female ovster drills in the 211 212 extreme low intertidal and subtidal zones from both natural and artificial substrate, including oyster reefs, 213 pier pilings, and boulders, within a 30-meter radius. We then transported snails in aerated coolers of 214 seawater from collection sites, kept cool with ice packs and separated by population. Water conditions 215 within the coolers were monitored to maintain 100% dissolved oxygen saturation and temperature within 4° C of collection temperature. Samples from Humboldt Bay and Tomales Bay (Pacific populations) were 216 collected in a similar fashion except that they were overnight mailed in plastic bags with saltwater-217 218 moistened paper towels but without seawater. Snails were kept cool with ice packs and upon arrival were 219 immediately placed in holding tanks separated by population at the University of Massachusetts, 220 Amherst. No mortalities occurred as a result of collection or shipping.

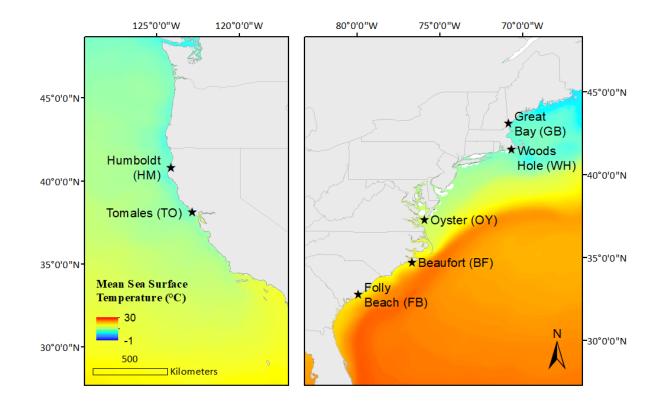


Figure 2. Urosalpinx cinerea collection sites on the Atlantic and Pacific seaboards of the United States.
 Mean SST is an annual composite of 2018 5km data (data source: NOAA/NESDIS Geo-Polar (Maturi et al., 2017); annual SST composite data from NOAA Coral Reef Watch 2018 v3.1).

226	We maintained Urosalpinx populations in a recirculating seawater system at 12 °C (salinity 30
227	PSU) until they were needed for experimentation and as other populations were collected. Populations
228	were kept separate in plastic aquaria with aeration. We fed broodstock Urosalpinx with blue mussels
229	(Mytilus edulis), acorn barnacles (Semibalanus balanoides), and eastern oyster flesh (Crassostrea
230	virginica) ad libitum. To initiate egg case laying, we raised the system water temperature by 1°C/day until
231	20°C was reached and then moved all broodstock to an identical recirculating seawater system at the
232	Gloucester Marine Station (UMass Amherst). We performed daily water changes on the broodstock
233	recirculating system using ambient coastal seawater maintained at 20°C. We also monitored ammonia
234	levels (API Mars Fishcare, Inc., Chalfont, PA) to ensure levels stayed below 0.25 ppm. Ammonia varied

between 0-0.25 ppm with one spike to 0.5 ppm caused by overfeeding, remedied with daily waterchanges.

237 Egg case collection and developmental acclimation

238 Our primary goal was to quantify thermal tolerance and plasticity (measured as developmental 239 acclimation capacity at 20 and 24 °C) across populations. We selected 20°C to enable comparison with prior work on Urosalpinx (Cheng et al. 2017) and chose 24°C to represent a warmer temperature that 240 *Urosalpinx* likely already experiences during summer and is below a previously recorded juvenile thermal 241 optima (26.5°C; Cheng et al., 2017). Thus, we hypothesized that an increase in acclimation temperature 242 from 20 to 24 °C would result in an increase in thermal tolerance. We performed daily inspections for egg 243 cases from July 5th - 31st, 2019, and collected a total of 122 egg cases. Mothers typically laid eggs in 244 245 clusters of 5-8 cases. In cases where a mother was discovered laying the egg cluster, we affixed a plastic numbered tag to the mother with cyanoacrylate glue to track the identity of egg laying mothers. We 246 247 tracked mother identity, as well as unique clutches (group of egg cases found together), to differentiate 248 egg cases laid by different individual mothers. Because some eggs were laid by unidentified mothers 249 (n=21), we used unique clutches to control for maternal effects.

250 We collected eggs the day they were laid and incubated them using two methods to facilitate 251 collection of different data types. For development time, we placed single eggs into plastic tea strainers 252 (Tops Permabrew, Darien, CT) that were divided in half with nylon fabric. Each tea strainer therefore 253 held two eggs from a single egg cluster and allowed us to track time to hatching of individual egg cases. For thermal tolerance, the remaining eggs were housed in undivided tea strainers separated by population 254 255 (20 - 30 egg cases per strainer) until hatchling emergence. Both types of strainers were submerged in seawater maintained at 20 or 24 °C (salinity = 30 PSU), which served as our developmental acclimation 256 257 for the egg stage. In each aquarium, we monitored temperature at least twice daily; temperature within the 258 aquaria never varied by more than ± 0.4 °C for the duration of egg development.

259 Immediately after hatching, we combined F1 snails from different mothers and of the same 260 population and acclimation temperature into strainers and fed F1 snails C. virginica oyster spat ad libitum (3 mm shell diameter; Muscongus Bay Aquaculture, Bremen, Maine). F1 snails were housed in strainers 261 262 between 8 and 16 days submerged within a tank (39 L) maintained at 20°C or 24°C before they were placed in the thermal tolerance experiment, and thus acclimation extended post-hatch. The mass of 263 264 juvenile snails as recorded immediately before the thermal tolerance experiment was not significantly different between both acclimation temperature treatments (GLM, $F_{1.649} = 2.90$, P = 0.0892). The egg 265 266 cases acclimated at 24°C from Great Bay, NH and Woods Hole, MA, did not produce enough juveniles to 267 enter in a heat bar trial, and so were not included in our analysis.

268 Thermal Tolerance

269 We quantified thermal tolerance and developmental acclimation across populations using LT_{50} 270 methodology with an aluminum heat bar (Kuo and Sanford, 2009; Cheng et al., 2017). The heat bar is drilled to accommodate 5 ml centrifuge tubes that can house individual snails that are then exposed to a 271 272 gradient of temperatures along the length of the heat bar. This heat bar was constructed with a solid 273 aluminum block similar to Kuo and Sanford (2009), but heat was applied with a silicone heating element (Omega SRFGA-406/2-P 60 watt, Omega Engineering, Norwalk, CT, USA) and adjusted with a 274 proportional integral derivative (PID) controller (ITC-100, Inkbird, Shenzhen, PRC). Cooling was 275 maintained by circulating 3-5 °C water through the opposing end of the heat bar. Although Urosalpinx 276 277 experiences aerial and aquatic thermal stress, this species is commonly found in both subtidal and low-278 intertidal habitats with limited aerial exposure (Carriker, 1955; Cheng and Grosholz, 2016; Cheng et al., 279 2017). Thus, we chose to quantify thermal tolerance in water to avoid the confounding effect of aerial 280 desiccation (Stillman and Somero, 2000).

In heat bar trials, individual snails were placed in 5 ml centrifuge tubes filled with 5 ml of aerated
seawater at the same acclimation temperature the snail experienced during development. We inserted a 2
x 2 cm 200 µm nitex mesh square into the tube using a plastic collar so that approximately 0.5 ml of the

284 tube's water was above the mesh. This prevented the snail from crawling out of the water, ensured free exchange of oxygen with the water in the tube, and enabled us to record water temperature without 285 disturbing the snail. We randomly assigned one of the three possible row positions along the heat bar, so 286 287 that each population was represented in a column but in a random row. Thus, we tested up to three 288 different population-acclimation treatments(each of which was defined as a "trial") at a time on the heat 289 bar array (Fig. S1). Each heat bar "run" was defined as a ramping of three trials in the heat bar with 18-30 290 snails from three populations and a single acclimation temperature. We quantified wet weight of each live 291 snail (Ohaus Pioneer PX Scale, Ohaus Corporation, Parsippany, NJ) prior to the run to account for age 292 and size effects, as age and age-linked size can affect thermal tolerance (Nyamukondiwa and Terblanche, 293 2009; Truebano et al., 2018). However, there was little evidence that age (as measured by body mass) predicted thermal tolerance (Table S2). Therefore, we removed body mass as a predictor from our 294 295 models. The shell length of these juvenile snails ranged approximately from 1-2 mm.

296 We used the PID controller to control the temperature ramp along the heat bar, increasing the controller setpoint by 5 °C every 30 minutes in steps from 25 °C to 60 °C for a total period of four hours. 297 In the final hour, we held the heat bar at 60 °C, so each snail was exposed to a heat ramp lasting five 298 299 hours (Table S3, Fig. S2). We measured the temperature in each column every hour using a 300 thermocouple. After the heat ramp, we removed the centrifuge tubes from the heat bar and allowed them 301 to recover in aerated seawater at the appropriate acclimation temperature (20 or 24°C) overnight. After 302 the recovery period, we evaluated snails for mortality using a stereomicroscope and a probe classifying 303 snails that did not retract their foot upon stimulus as dead and those that reacted as alive (Cheng et al., 2017). In total, we conducted 22 independent heat bar trials (20°C n = 14, 24°C n = 8) for seven 304 305 populations using a total of 652 juvenile snails (Table S4). Individual snail sample sizes between acclimations were uneven, with n = 418 at 20°C and n = 234 at 24°C, due to egg case availability. 306

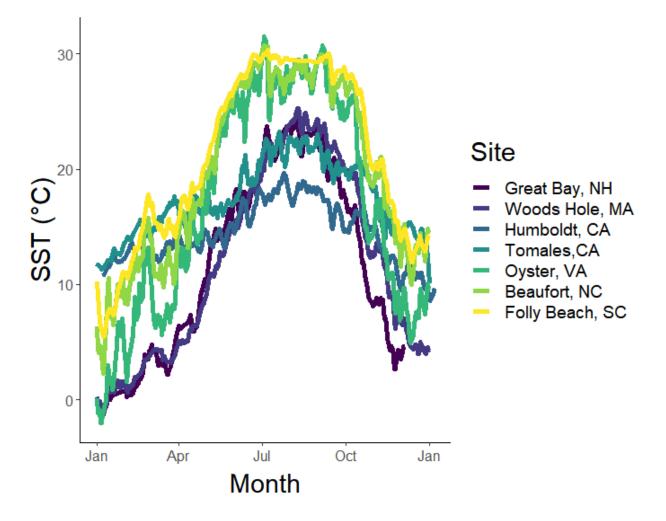
307 Developmental Metrics

308 In addition to thermal tolerance, we quantified the effects of temperature on development across 309 populations by measuring: 1) hatching success, 2) clutch size, and 3) developmental rate. To assess embryo hatching success over the incubation period, we counted the number of successfully hatched 310 311 snails and compared this to the number of unsuccessful embryos using a microscope (Leica S9i, Leica 312 Microsystems, Inc, Buffalo Grove, IL. USA). We also counted the number of initial embryos per egg case 313 to evaluate clutch size. To measure developmental rate of embryos within egg cases, we noted the lay 314 date of each case within two days of laying and checked egg cases daily for hatching. We classified an 315 entire egg case as hatched when the first hatchling snail emerged from the opening at the top of each egg 316 case, allowing hatchlings to crawl freely out of the egg case.

317 Environmental Metrics

318 While latitude is a commonly used metric of the types of environmental conditions experienced by a 319 population (e.g. Sunday et al., 2014), we chose to evaluate multiple site level environmental temperature 320 metrics as potential predictors of thermal tolerance and developmental traits because latitude may not be 321 an accurate predictor of local scale temperatures experienced by organisms (Kuo and Sanford, 2009). 322 Moreover, while latitude can be a useful predictor that is correlated with environmental conditions, habitat temperatures can differ at the same latitude based on ocean (Pacific vs. Atlantic) and local (inner 323 324 estuary vs. outer estuary) conditions, and is thus another potential direct driver of environmentally 325 adapted traits (Kuo and Sanford, 2009; Baumann and Conover, 2011; Sunday et al., 2011). Thus, we 326 extracted a series of environmental temperature predictors with the goal of understanding what aspect of 327 habitat temperature (e.g. mean vs. maximum temperature) best predicted patterns in thermal tolerance. 328 From these temperature data, we calculated five environmental predictors: 1) mean annual temperature, 2) 329 summer mean temperature, 3) upper 25th percentile of the summer period, 4) the upper 10th percentile of 330 the summer period, and 5) the maximum summer temperature (Table S5). We used each environmental predictor by itself in each model to evaluate which predictor best explained trait performance patterns 331 332 using model selection, including a null model. We selected site temperature data based on the

completeness of the record in 2018, the proximity of the temperature data to the collection site (no more
than 15 km; Table S1), and from locations representative of collection sites (e.g. environmental data was
collected from buoys in tidal creeks if the collection site was in a tidal creek). When available, we
selected only continuous 2018 temperature records, but the two data sources from the Pacific only had
continuous data from 2015 (Table S1, Fig. 3). Summer was classified as between June 1 and September
30.



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Figure 3. Sea surface temperature (SST) from sources near broodstock collection sites. Each time series represents one year of data from January 1 to December 31, 2018 (except for Pacific sites, where data

ranged between January 1 to December 31, 2015) for comparison of thermal regime across populations.

343 Lines represent the daily mean temperature at each site. Sites are presented in order of annual mean

temperature. See Table S1 for source list and sampling dates.

346 Statistical Analysis

347 Thermal Tolerance

348 To evaluate thermal tolerance across populations we used a two-step approach. First, we extracted LT_{50} estimates for each heat bar trial ("trial" = 18-30 snails from a population-acclimation treatment in the heat 349 350 bar) using Firth's bias-reduced logistic regression (Heinze and Schemper, 2002) due to complete 351 separation of the survival data. Complete separation occurs when a predictor perfectly discriminates 352 between binomial states. In our case, survival in each trial was consistent up to a certain temperature threshold after which all individuals died, and thus was completely separated (Fig. 4; Cheng et al. 2017). 353 354 This lack of variation is problematic for traditional model estimation, thus necessitating the alternate 355 approach. For these analyses we used the *brglm2* package in R (Kosmidis, 2021) to model the effect of final heat bar temperature on survival for each population and acclimation temperature treatment. Thus, 356 357 each "run" of the heat bar produced three LT₅₀ measurements for the three "trials" of populationacclimation treatments. 358

359 Second, as opposed to modeling LT_{50} as a function of population (e.g. using ANOVA), we used a 360 regression-based approach using five environmental variables from each population to understand drivers of thermal tolerance over an environmental cline (Table S5). Once we extracted the LT_{50} from each trial, 361 362 we then tested for geographic patterns in thermal tolerance by pairing each population's environmental 363 data (Table S5) with their extracted LT_{50} estimates. These environmental data were then used as a suite of predictors, in addition to the acclimation temperature of each trial, in a model-selection framework. We 364 constructed generalized linear models with gaussian error distributions using this set of environmental 365 and acclimation predictors, and used small sample adjusted Akaike's Information Criterion (AICc) to 366 select models which had the greatest support against a null model. We chose our cut-off of well-supported 367 368 models for model selection throughout as $\Delta AICc < 2$ (Burnham and Anderson, 2002).

369 We further examined the difference between calculated thermal tolerances (LT_{50}) and the habitat 370 temperature of each population (hereafter referred to as warming tolerance; Deutsch et al., 2008). We calculated warming tolerance as $WT = LT_{50} - T_{hab}$, with T_{hab} as the maximum summer temperature. This 371 372 method accounts for maximum water temperatures an organism could experience, which is likely to be a 373 selective force across populations (Kingsolver et al., 2013; Sunday et al., 2014). We calculated separate 374 warming tolerance estimates using LT_{50} values from the 20 and 24°C acclimation temperatures to assess how thermal history may influence thermal sensitivity estimates. While we included the two Pacific sites 375 376 in the data, we did not model an effect of invasion status because there was no overlap in T_{hab} values 377 between oceans and due to limited population replication in the Pacific. 378 **Developmental Traits** 379 We used generalized linear mixed models to assess the fixed effects of acclimation temperature and 380 environmental predictors, and their interaction on developmental traits (hatching success, clutch size, development time). We included clutch as a random effect. For clutch size, we used a Conway-Maxwell 381 382 Poisson error distribution because of initial overdispersion in the data (Chanialidis et al., 2018). For 383 hatching success of snails, we used a binomial error distribution with logit link function. For development time, we used a gaussian distribution. For all development analyses, we used environmental predictors as 384 defined in Table S5. For these analyses we used the glmmTMB package (Brooks et al., 2017). We 385 386 performed all thermal tolerance and developmental trait statistical analyses in R (v. 3.5.1, R Core Team, 387 2018).

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RESULTS

390 Thermal Tolerance

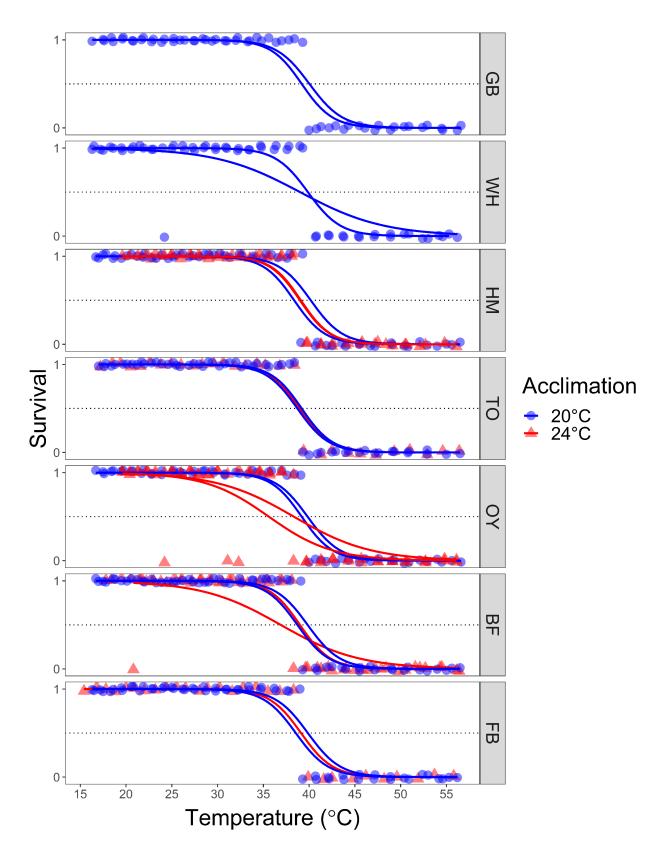


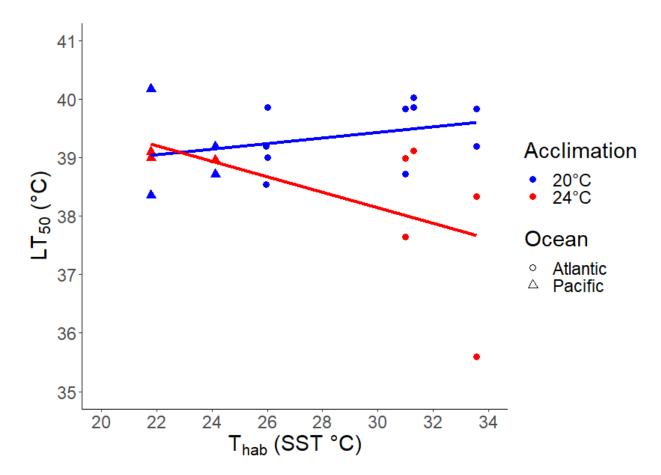


Figure 4. Survivorship of *Urosalpinx* hatchlings (survival = 1, mortality = 0) as a function of final temperature within the heat bar array, separated by acclimation temperature. Model estimates represent independent heat bar trials. Dotted line represents the threshold for calculating LT_{50} . Populations are ordered by ascending mean temperature within the native and invasive (HM and TO) range. Site codes are defined as in Fig. 1. Points jittered for visual clarity.

398

399	The most supported model describing spatial patterns of thermal tolerance in Urosalpinx
400	contained habitat temperature (T_{hab}) as measured by the maximum summer habitat temperature at each
401	site with an interactive effect with acclimation temperature (T_{acc}). When acclimated at 20°C, thermal
402	tolerance increased with habitat temperature significantly but with high variability (GLM, $F_{3,18} = 4.51$, P
403	= 0.0417; Fig. 5, Table 1). When acclimated at 24° C, thermal tolerance decreased significantly with
404	habitat temperature (GLM, $F_{3,18} = 4.51$, $P = 0.0352$; Fig. 5, Table 1). Urosalpinx acclimated at 20°C and
405	24°C had a cross-population mean thermal tolerance of 39.3 ± 0.61 °C ($n = 14$) and 38.3 ± 1.22 °C ($n = 14$)

406 8; mean \pm SD), respectively.



408 Figure 5. LT_{50} estimates of *Urosalpinx* hatchlings over their habitat maximum summer temperature and 409 two experimental acclimation temperatures. T_{hab} is the maximum summer temperature.

410

411Table 1. Parameter estimates for thermal tolerance, warming tolerance, and developmental rate models.412Bold text denotes significance levels of P < 0.05. Multiple and adjusted R-squared values are presented413for model-averaged and single-model GLMs. For mixed-effect models (developmental rate), the marginal414and conditional R-squared values are given, which estimate model explanatory power between fixed415effects and fixed and random effects combined (Nakagawa and Schielzeth, 2013). Thab as determined via416AICc model selection, are maximum summer temperature for both thermal tolerance and warming

417 tolerance, and mean annual temperature for developmental rate.

Parameter	Estimate	SE	t/z	Р
Thermal Tolerance R ² T _{hab} *Acc (multiple/adjusted): 0.429/0.334				
Acc ₂₀ (Intercept)	17.2	12.4	1.39	0.182
Acc ₂₄	1.04	0.568	1.83	0.0838
T_{hab} * Acc ₂₀	0.956	0.436	2.19	0.0417
T _{hab} * Acc ₂₄	-0.0454	0.0199	-2.28	0.0352
Warming Tolerance				

$R^2 T_{hab}$ *Acc (multiple/adjusted): 0.975/0.971				
Acc ₂₀ (Intercept)	38.0	1.49	25.5	<0.001
Acc ₂₄	4.16	2.27	1.83	0.0838
T _{hab} * Acc ₂₀	-0.951	0.0533	-17.9	<0.001
T _{hab} * Acc ₂₄	-0.182	0.0797	-2.28	0.0352
Developmental Rate R ² _{GLMM} (marginal/conditional): 0.906/0.935				
Acc ₂₀ (Intercept)	46.143	1.756	26.283	<0.001
Acc_{24}	-15.430	1.844	-8.368	<0.001
T _{hab} * Acc ₂₀	-0.463	0.105	-4.397	<0.001
$T_{hab} * Acc_{24}$	0.286	0.111	2.577	0.00996

419 Warming Tolerance

420	We found a strong pattern of decreasing warming tolerance (thermal tolerance – maximum habitat
421	temperature) with increasing summer maximum site temperature for both acclimation temperatures (GLM
422	$F_{3,18} = 11.4, 20^{\circ}C: P < 0.001, 24^{\circ}C: P = 0.0352)$, but that warming tolerance was not significantly
423	different between acclimations (GLM $F_{3,18} = 11.4$, $P = 0.0838$; Fig. 6, Table 1). Invasive pacific
424	populations appeared to have the highest warming tolerance values, although we note that we did not
425	explicitly model invasion status because of the low number of invasive population replicates $(n = 2)$. The
426	minimum calculated warming tolerance occurred in the Virginia population ("Oyster") at 24°C
427	acclimation (2.03 °C), while the largest warming tolerance occurred in the California ("Humboldt")
428	population at 20 °C acclimation (18.4 °C).

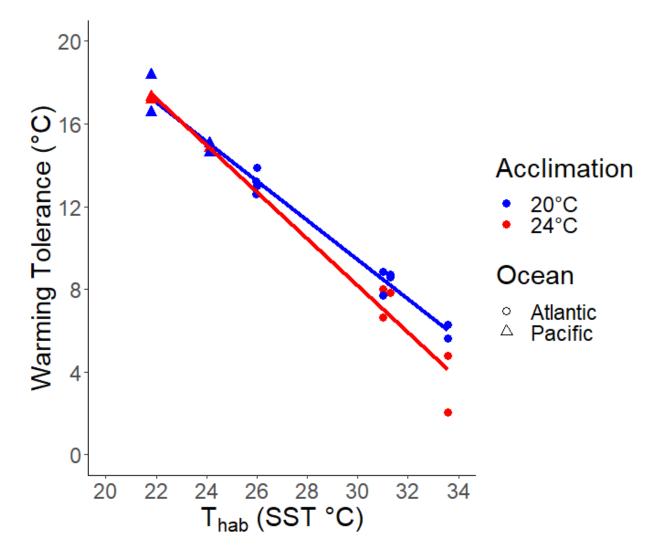




Figure 6. Latitudinal and oceanic trends in warming tolerance $(LT_{50} - T_{hab})$, with T_{hab} being the maximum site summer temperatures. Trendline depicts the significant relationship between warming tolerance and T_{hab} at the 20°C and 24°C acclimations. Note that we include Pacific site data, but omitted invasion status as a predictor from analysis because of low sample size. T_{hab} is the maximum summer temperature.

The hatching time of *Urosalpinx* egg cases decreased with greater mean annual habitat temperature of the source population (GLMM n = 39, P < 0.001) for egg cases reared at a common temperature of 20°C (Fig. 7, Table 1). At 20°C acclimation, the shortest developmental time occurred in egg cases from the southernmost Atlantic site (Folly Beach, 36.5 ± 3.53 days (SD)), while the greatest

⁴³⁵ Developmental Traits

development time occurred in the northernmost Atlantic site (Great Bay, 41.8 ± 2.59 days). When 440 acclimated at the higher temperature of 24 °C, hatching time decreased across all sites (GLMM n = 46, P441 = 0.00996; Fig. 7, Table 1). The shortest development time at 24°C occurred in North Carolina (Beaufort; 442 26.8 ± 1.28 days), and despite the significant negative trend between habitat temperature and time to 443 444 hatching, the slowest development rates occurred at both the northernmost and southernmost Atlantic 445 sites (South Carolina; 29.3 ± 0.577 days and New Hampshire; 29.2 ± 1.47 days). Random effects of clutch gave intercept variance of 0.955 ± 0.977 (SD), and little difference between marginal (0.906; fixed 446 effects only) and conditional (0.935; fixed and random effects) R²_{GLMM} (Nakagawa and Schielzeth, 2013). 447 Multicollinearity was low (VIF ≤ 2.5) for all well-supported developmental trait models. Both clutch size 448 449 and hatching success metrics had multiple well-supported models, so we model averaged top models of 450 clutch size and hatching success. None of the best-supported models were overdispersed (deviance < 451 degrees of freedom).

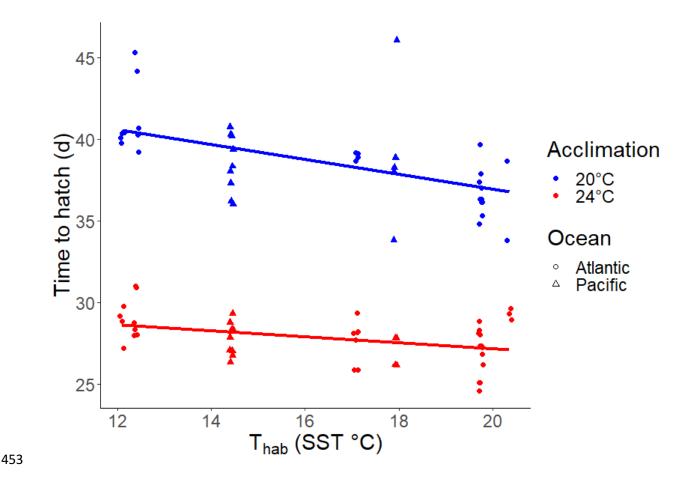
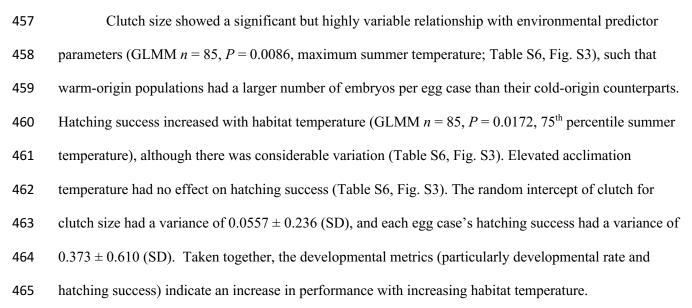


Figure 7. Developmental rate of *Urosalpinx* egg cases when acclimated at 20°C and 24°C. T_{hab} is the mean annual temperature. Points jittered for visual clarity.



DISCUSSION

467 Thermal performance has often historically been assumed to be homogeneous within species, an assumption that can generate inaccurate forecasts of species response to climate change if there is 468 469 adaptive differentiation across populations. There is increasing recognition that intraspecific variation 470 may be common in the ocean (Kuo and Sanford, 2009; Zippay and Hofmann, 2010; Kelly et al., 2012; 471 Hong and Shurin, 2015; Pereira et al., 2017; Sasaki and Dam, 2019). However, observations supporting 472 this view are generally limited, particularly across populations of a species' native and invasive ranges (but see: Henkel et al., 2009; Yu et al., 2012; Hill et al., 2013; Tepolt and Somero, 2014; Wesselmann et 473 474 al., 2020). Here, we found evidence for greater thermal tolerance in southern populations of oyster drills 475 that experience higher habitat temperatures, in support of our hypothesis of local adaptation. However, 476 when developmental acclimation temperature was increased, thermal tolerance decreased in southern 477 populations (2.1-6.4% decrease), contrary to expectations of greater thermal tolerance with higher 478 acclimation. Further, we found diminished warming tolerances of low latitude Atlantic populations as 479 compared to high latitude Atlantic (native) and Pacific (invasive) populations, consistent with the non-480 compensating local adaptation model of warming tolerance (Fig. 1C). This follows our conceptual 481 framework of how variation in thermal tolerance may be inverse to variation in warming tolerance (Fig. 1); thermal tolerance that compensates well with changes in the environment results in low variance in 482 483 warming tolerance. In this study, we show that because thermal tolerance does not scale strongly with 484 environmental temperature, variation in calculated warming tolerance is high between populations. These 485 results suggest a striking contrast. Low latitude native populations appear to exhibit high climate sensitivity and may become extirpated if warming continues ultimately resulting in range contraction. In 486 487 contrast, the two Pacific (invasive) populations of Urosalpinx studied may be more likely to persist in a 488 warming future because of a large buffer between current habitat temperatures and their thermal 489 tolerance. Urosalpinx has well-documented impacts on native, foundational species such as Olympia 490 ovsters, and therefore will likely continue to drive cascading negative effects on native biodiversity into 491 the future (Kimbro et al., 2009; Cheng and Grosholz, 2016).

492 We found interactive effects of source population environment and acclimation temperature on 493 thermal tolerance (Fig. 5). Populations reared at 20°C displayed a positive relationship between thermal tolerance and habitat temperature, consistent with other studies on marine invertebrates (Zippay and 494 495 Hofmann, 2010; Sunday et al., 2011; Kelly et al., 2012; Pereira et al., 2017; Sasaki and Dam, 2019). 496 Although the differentiation in thermal tolerances appears small (~ 1.0 °C across the range tested), this 497 effect size is similar to other studies testing intraspecific variation in thermal tolerance (Fangue *et al.*, 2006; Kuo and Sanford, 2009; Jensen et al., 2019). Interestingly, higher developmental acclimation 498 499 temperature (24 °C) resulted in a negative relationship between habitat temperature and thermal tolerance, 500 or what we define as 'negative plasticity'. At first glance, these results are counterintuitive given the 501 tendency of higher acclimation to result in elevated thermal tolerances (Angilletta Jr., 2009; Pereira et al., 2017; Sasaki and Dam, 2019). However, evidence of a negative response to higher acclimation 502 temperature has been demonstrated in nudibranchs (Armstrong et al., 2019) and salmonids (Blair and 503 504 Glover, 2019; Del Rio et al., 2019) in both developmental and within stage acclimations, albeit not 505 between multiple populations. This negative plasticity in thermal tolerance from southern populations is 506 suggestive of a tradeoff between elevated thermal tolerance and plasticity (Stillman, 2003; Armstrong et 507 al., 2019; van Heerwaarden and Kellermann, 2020). Southern populations have evolved elevated thermal 508 tolerance in response to warm environmental conditions but have done so at the cost of plasticity extent. 509 Because northern/invasive populations have lower evolved thermal tolerances, they do not exhibit such 510 trade-offs with plasticity. It should be noted that our scope of inference is limited here because we were 511 not able to quantify thermal tolerance of northern Atlantic sites at 24°C (Great Bay and Woods Hole) 512 because we were not able to obtain enough juveniles from each treatment to run a heat bar trial. 513 Furthermore, an acclimation of 24°C itself may be stressful for embryonic and newly hatched Urosalpinx. We originally chose 24°C as the higher acclimation temperature because it is below the measured 514 515 thermal optima of juvenile, invasive range Urosalpinx (26.5°C, Cheng et al., 2017), and hatchling 516 survivorship, while invariant with acclimation temperatures in our study, has previously been shown to peak at 20°C and decrease at 25°C (Ganaros, 1958). However, because early life stages are often the 517

518 most vulnerable to thermal stress, it is further possible that physiological stress is incurred in embryos and hatchlings at 24°C (Truebano et al., 2018; Dahlke et al., 2020; McKenzie et al., 2020). Thermal stress can 519 accumulate over time with heightened sublethal temperatures, resulting in reduced survivorship in what 520 521 has been described as a tolerance landscape (Rezende *et al.*, 2020). These developmental acclimation 522 effects are tested less often, but are important because ocean warming is occurring across seasonal cycles 523 and can impact early development when many organisms are the most sensitive (Pechenik, 2006; Marshall and Morgan, 2011; Dahlke et al., 2020). Our results point to the importance of carefully 524 525 considering how seasonality of environmental exposure and ontogeny may affect thermal sensitivity 526 across life stages. This is a critical consideration when designing experimentation tracking local 527 adaptation across generations, especially with complex life stage organisms from environments with strong seasonal thermal fluctuations. Models that predict population persistence using adult thermal 528 529 optima or tolerance may overpredict potential ranges by not considering heightened sensitivity of early 530 life stages and the carry-over effects of warming during development.

531 Among environmental correlates, maximum habitat temperature best explained variation in thermal tolerance. Most studies use mean annual temperature in predicting variation in thermal tolerance, 532 533 perhaps because these data are readily available and explain some variation in tolerance (e.g. Deutsch et al., 2008; Hughes et al., 2018). However, maximum habitat temperature is expected to be the main driver 534 of thermal tolerance both within and across species (Hoffmann, 2010; Kelley, 2014; Pinsky et al., 2019). 535 536 Maximum temperatures should act as a 'filtering' agent such that a locally adapted population will have 537 thermal tolerances selected for from standing genetic variation that allow it to persist in that environment 538 (Bennett et al., 2019; Pinsky et al., 2019). Local thermal heterogeneity, driven by processes such as 539 upwelling, tides, and currents also mean that environmental metrics like latitude or mean temperature are 540 not necessarily correlated with maximum habitat temperature (Baumann and Doherty, 2013). We found that maximum habitat temperature consistently drove variation in thermal tolerance spanning native and 541 542 invasive ranges (Fig. 4). Our temperature records, obtained from buoys within 15 km and of similar

543 habitat type of collection sites, offer a general view of the environmental conditions experienced by 544 populations. However, given organismal body temperature itself may vary as a function of microhabitat and behavior (Helmuth et al., 2010), the exact maximum temperature each Urosalpinx population 545 546 experiences may differ from those obtained via buoy data. As a result, we suggest future work consider 547 testing relationships between upper thermal tolerance and maximum habitat temperatures along with 548 mean temperature and/or latitude, as well as deploying collocated temperature loggers to refine these 549 environmental parameters. By not directly correlating thermal tolerance with a major selective 550 environmental force (i.e. maximum habitat temperature), patterns of local adaptation may be ignored or 551 overstated, potentially wasting resources by managing populations that are not actually sensitive to 552 climate change.

553 Diminished warming tolerance at warm-origin sites indicates that southern populations are closer 554 to their thermal limit than their northern counterparts (Fig. 6), and that population origin has a stronger 555 effect on climate sensitivity than does acclimation temperature. Interestingly, this result sets up a third 556 potential pattern of thermal tolerance, habitat temperature, and warming tolerance (see Fig. 1). Despite thermal tolerance being locally adapted, warming tolerance was not constant across populations, 557 558 indicating that a third model of warming tolerance (what we call here 'non-compensating' local 559 adaptation, Fig. 1C) between niche conserved (Fig. 1A) and locally adapted populations (Fig. 1B) are 560 possible. This is likely a result of thermal tolerance not being 1:1 correlated with decreasing habitat 561 temperature. This decreasing relationship between warming tolerance and habitat temperature is 562 consistent with studies that have examined intraspecific sensitivity to climate in crabs, nudibranchs, and 563 leaf miner moths (Gaitán-Espitia et al., 2014; Pincebourde and Casas, 2015; Armstrong et al., 2019), as 564 well as studies of interspecific climate sensitivity (Deutsch et al., 2008; Sunday et al., 2011; Allen et al., 565 2012; Diamond et al., 2012; Vinagre et al., 2016; Comte and Olden, 2017; Janion-Scheepers et al., 2018). Taken together, this evidence supports the view that low latitude populations appear to have high climate 566 567 sensitivity (Tewksbury et al., 2008; Pinsky et al., 2019). In contrast, temperate populations have greater

568 warming tolerance despite reduced thermal tolerance, perhaps because of exposure to lower

environmental temperatures (Deutsch *et al.*, 2008; Janion-Scheepers *et al.*, 2018). Reduced warming
tolerance at the warm edge of a population's range also highlights the potential role of thermal tolerance
in driving extirpation and range contractions at the trailing edge (Sunday *et al.*, 2012; Cahill *et al.*, 2014;
Hardy *et al.*, 2014). Depending on the management goal for a species exhibiting this pattern of warming
tolerance (control for *Urosalpinx*, conservation for others), this potential for local extinction and species
range contraction at the warm trailing edge is of critical interest and may call for resource reallocation
away from warm, trailing-edge populations.

576 We found strong evidence for faster developmental rates for populations sourced from warm habitats, and higher developmental acclimation at 24°C resulting in overall faster growth than at 20°C 577 (Fig. 7). Warm, southern populations developed the fastest at all acclimation temperatures, as expected by 578 579 biogeographic theory of embryonic development rate in marine ectotherms (Lonsdale and Levinton, 580 1985; Collin, 2003; Weydmann et al., 2015). Increased development rate at lower latitudes may result 581 from simple increases in metabolic rate with habitat temperature (Lonsdale and Levinton, 1985), or potentially because of selection arising from heightened risk of predation in tropical low latitude systems 582 583 (Schemske et al., 2009). Interestingly, the fastest development rate occurs at the acclimation temperature 584 (24°C) and populations (low latitude Atlantic) that had the lowest thermal tolerance, suggesting potential 585 trade-offs across life stages (Stillman, 2003). While both were highly variable, hatching success increased 586 with habitat temperature, such that warm populations develop faster and have higher survivorship, and 587 clutch size decreases with higher habitat temperature. Therefore, warm-origin populations spawn smaller 588 egg case clutches, which develop quicker, and have a greater chance of developing successfully. As 589 juveniles, these warm-origin populations show higher thermal tolerance (Fig. 5), but only at a lower 590 acclimation temperature. Additionally, the reduced number of embryos per egg case (low "embryo packing") in warm populations may be a product of a tradeoff between embryo density and oxygen 591 592 availability within each egg case in warm waters (Lee and Strathmann, 1998; Fernández et al., 2007).

593 These results indicate the potential for rapid embryonic development to result in trait performance trade-594 offs in later life stages as a result from increased metabolic demand during embryonic growth (Van Der Have, 2002; Pörtner et al., 2006; Del Rio et al., 2019). If development rate is maximized at each 595 596 acclimation temperature, then enzymatic activity may itself be maximally efficient at these temperatures, 597 and thermal tolerance is reduced due to inefficient enzymatic reactions at elevated temperatures (Somero, 598 1995; Van Der Have, 2002). One caveat of our findings is that we quantified embryonic performance in 599 July in order to synchronize experimental treatments. It is possible that the variation in hatching success 600 and clutch size may reflect phenological shifts in spawning seasons (Carriker 1955). Future efforts should 601 quantify these aspects of spawning and development across the spawning seasons in order to fully resolve 602 the potential range of intraspecific variation and plasticity in Urosalpinx. Our results point to the mechanistic importance of early life stage experiences on trait performance and tradeoffs in subsequent 603 604 life stages, and the need for future research to characterize trait performance and optima across life stages 605 (Pechenik, 2006; Slotsbo et al., 2016).

606 We found invasive and high latitude native *Urosalpinx* populations to be the least sensitive to climate impacts based on their large warming tolerance values, suggesting that these populations will 607 608 persist in their environments. We acknowledge that sampling in the invasive region was limited to two 609 populations and that current data is unable to determine whether greater warming tolerance of these 610 populations is due to population genetics (e.g., founder effect or population bottleneck from introduction) 611 or due to the environment alone (i.e., a large buffer from current habitat temperatures and thermal 612 tolerance). Current efforts are underway to resolve genetic differences among populations in the native 613 and invasive range. Nonetheless, the high warming tolerance observed at the California sites is a concern 614 for native biodiversity because near term warming is likely to increase the predatory impact of *Urosalpinx* 615 on native species, including consumption of Olympia oysters (Ostrea lurida) that are the focus of conservation and restoration efforts (Cheng et al., 2017). Further, heightened development rate at greater 616 acclimation suggests that embryos will develop faster with potentially higher metabolic rates, increasing 617

618 the consumption of newly-hatched juveniles on oysters. From a community ecology perspective, these 619 differing climate sensitivities between Urosalpinx throughout native and invasive ranges demonstrates the potential for indirect impacts of climate change on native biodiversity. Interactions between Urosalpinx, 620 climate, and humans highlights "trophic skew", the reorganization of biological communities with species 621 622 loss from extinction and species gain from invasion (Grosholz, 2002; Duffy, 2003; Byrnes et al., 2007). 623 As marine environments warm, native species will experience both abiotic pressure from warming as well 624 as pressure from the persistence and proliferation of invasive, warm-origin predators like Urosalpinx 625 (Cheng and Grosholz, 2016). Early eradication and control of these resilient invasive predators may assist 626 native species by removing a biotic pressure as natives adapt or migrate in the face of climate change, 627 thereby potentially reducing of trophic skewness (Byrnes et al., 2007; Grosholz and Ruiz, 2009; Cheng et al., 2017). 628

629 In conclusion, our work demonstrates the importance of taking an intraspecific approach to 630 examining thermal performance and sensitivity to climate. Such variation can have large implications for 631 forecasts of species responses to climate change that often assume homogeneity across populations, thereby missing the possibility of more resilient populations under climate change. We found largely 632 633 negative effects of developmental acclimation on thermal tolerance, a crucial consideration given that climate change occurs across temporal scales (e.g. seasons) and will result in biological effects both 634 635 within and across life stages. We also show that integrating environmental data can provide a more 636 complete picture of population-level sensitivity that may drive geographic range contractions. Taken 637 together, this approach can be useful for developing an understanding of climate impacts on populations 638 across their native and invasive ranges. Such a perspective is useful for clarifying potential interactions 639 between climate and biological invasions that can erode native biodiversity. The variation in thermal 640 physiology we demonstrate here supports the necessity of using data from multiple populations when 641 making ecological forecasts of climate change.

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