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3 **Urban environments promote adaptation to multiple stressors**
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51 EB: Conceptualization. EB, LL, SGK, AS, CMR, AG, GS, NT, JWED, JCH and RNC:

52 Investigation. EB, PJH, IS, AK and RNC: Formal analysis and visualisation. EB and RNC:

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55

56 **DATA AVAILABILITY STATEMENT**

57 Raw data supporting the findings of this study is available at Drayad:

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60

61 **CONFLICT OF INTEREST**

62 The authors declare no conflicts of interest.

63

64 **ETHICS APPROVAL**

65 Ethical approval was not required for the nature of this work.

66

67

68 **ABSTRACT (150 words)**

69 Anthropogenic activities have drastically changed environmental conditions worldwide,
70 negatively impacting biodiversity and ecosystem services. At the same time, the majority of the
71 human population lives in urban areas that are greatly altered from natural habitats. Nevertheless,
72 many species thrive in urban environments. To improve our knowledge of evolution and
73 adaptation in these anthropogenically impacted habitats, we conducted the widest series of stress
74 experiments to date with three marine taxa: one mussel and two gammarid species. We
75 compared intraspecific populations from protected and human-altered habitats to determine their
76 tolerance to salinity, temperature and partial pressure of CO₂ in water (pCO₂) regimes.
77 Populations from impacted habitats typically outperformed protected habitat populations, with
78 individuals from the most impacted habitat being the most robust. We propose that urban
79 populations are adapting to life in disturbed environments — this adaptation concurrently
80 promotes more resilient rescue populations but potentially confers increased invasion risk from
81 non-native species.

82

83 **INTRODUCTION**

84 The Earth and its environments are being rapidly changed by increasing human populations and
85 economic development (Hawkins, 2012). Anthropogenic activities, including burning of fossil
86 fuels and deforestation, among others, have triggered climate change, pollution, soil erosion, and
87 habitat destruction (Hawkins, 2012; IPCC, 2014). The introduction of non-native species (i.e.,
88 species moved by human activity outside their native region) is also prominent due to the
89 increasing global trade and transport of materials (Hulme, 2009; Sardain *et al.*, 2019).
90 Subsequently, these changes have dramatically affected numerous ecosystems worldwide,
91 leading to biotic homogenization, decreases in biodiversity, and occasionally, to losses of
92 ecosystem functioning and services (Olden *et al.*, 2004; Hawkins, 2012; IPCC, 2014; Linders *et*
93 *al.*, 2019; Pyšek *et al.*, 2020).

94 More than half of the global human population presently lives in urban areas, with these
95 numbers expected to be double by 2050 (Johnson & Munshi-South, 2017; United Nations
96 Statistics Division, 2017). This urbanized world consists of dense settlements made of buildings,
97 roads, and other supporting infrastructures, with urban habitats significantly altered from natural
98 ones (Johnson & Munshi-South, 2017). As a consequence of rising urbanization, environments
99 in cities exhibit altered microclimates, which trend towards higher temperatures, alongside
100 elevated air, water, noise, and light pollution (LLUR, 2001; Grimm *et al.*, 2008; Nikulina *et al.*,
101 2008). They also experience increased habitat fragmentation (Dobbs *et al.*, 2017), lower diversity
102 and abundance of native species, high numbers of non-native species (Aronston *et al.*, 2014;
103 McKinney *et al.*, 2008), and lower phylogenetic diversity within communities (Knapp *et al.*,
104 2012; Sol *et al.*, 2017). Still, urban environments are successfully exploited by many non-human

105 species (Johnson & Munshi-South, 2017; Perry & Göttert, 2024). Pigeons, peregrine falcons,
106 ducks, carps, and goldfish are just some of the examples of those that thrive in urban habitats.

107 While rates of evolution had been deemed to be too slow to have relevance to
108 urbanization, recent research across a wide variety of taxa has demonstrated that organisms can
109 indeed evolve rapidly in response to urban environments (Alberti *et al.*, 2017; Johnson &
110 Munshi-South, 2017; Caizergues *et al.*, 2018; Santangelo *et al.*, 2018; Thompson *et al.*, 2018;
111 Borden & Flory, 2021; Perry & Göttert, 2024). Evolutionary changes have been observed in as
112 little as two generations (Kinnison & Hendry, 2001), and can include behavioral, morphological,
113 and physiological adaptations, or even speciation, in response to altered environmental
114 conditions, novel resources and different physical habitats, among others (Giraudeau *et al.*, 2014;
115 Winchell *et al.*, 2016; Martin *et al.*, 2019; Borden & Flory, 2021; Ålund *et al.*, 2020). These
116 changes may be particularly rapid in urban environments, where significant alterations to
117 numerous abiotic and biotic factors are simultaneous and often predictable. Therefore, studies of
118 adaptation in urban environments may be ideally placed for testing wider evolutionary theory
119 across taxa (McKinney, 2002; Grimm *et al.*, 2008), particularly in the context of multiple
120 stressors (Todgham *et al.* 2005). In turn, understanding the adaptation potential of diverse taxa,
121 as well as ecosystem resilience to anthropogenic stressors, urbanization, and global change, are
122 of paramount importance to inform conservation actions, even in natural habitats.

123 In the last two decades, whereas research into the effects of global change on ecosystems
124 has grown substantially, eco-evolutionary dynamics in urban environments have been neglected
125 outside of the terrestrial realm (Johnson & Munshi-South, 2017; Santangelo *et al.*, 2018;
126 Thompson *et al.*, 2018; Alter *et al.*, 2020; Borden & Flory, 2021; Perry & Göttert, 2024). Marine
127 ecosystems have been particularly overlooked in terms of urban evolution, but provide

128 exceptional model systems considering the presence of multiple stressors and clear disturbance
129 gradients (e.g., harbours versus protected areas). To improve our knowledge on evolution and
130 adaptation potential of taxa in urban and anthropogenically impacted habitats, we conducted a
131 series of laboratory experiments among populations of one mussel and two gammarid species
132 collected at protected and human-altered habitats in the Baltic Sea, to determine their tolerance
133 (i.e., plasticity) to diverse levels of salinity, temperature, and pCO₂ (i.e., partial pressure of CO₂
134 in water). Taking into account the concurrent evolution of adaptation to heterologous stressors
135 (Todgham *et al.* 2005), the overall aim of this study was to test whether populations inhabiting
136 human-altered habitats are better adapted to anthropogenic stressors, including climate change,
137 than populations from protected ones. We tested the null hypothesis that there is no difference in
138 stress resistance between populations inhabiting human-altered and protected habitats.

139

140 **MATERIALS AND METHODS**

141 **Experimental organisms and collection sites**

142 Individuals of populations of one mussel (i.e., *Mytilus* sp.) and two gammarid species (i.e.,
143 *Gammarus locusta* and *G. salinus*) were collected in protected and human-altered (i.e., impacted)
144 habitats in the Baltic Sea in Germany (Figure 1). Only hybrids of *M. edulis* and *M. trossulus* are
145 present at our sampling locations (Knöbel *et al.*, 2021); therefore, we refer to the mussel species
146 used as *Mytilus* sp. In the case of gammarids, each individual was morphologically identified
147 according to Köhn & Gosselck (1989) and Zettler & Zettler (2017). All three taxa used in the
148 experiments are native to the Baltic Sea (Springer & Crespi, 2007; Gaitán-Espitia *et al.*, 2016;
149 Cuthbert *et al.*, 2020; Briski *et al.*, 2024). While both gammarid species (alongside ~99% of
150 global biodiversity) do not yet have an invasion record (Cuthbert *et al.*, 2020; Briski *et al.*, 2024),

151 the mussel *M. edulis* is non-native in Southern Europe, Asia, and North America (Springer &
152 Crespi, 2007; Gaitán-Espitia et al., 2016; Briski et al., 2024); *Mytilus trossulus* is non-native in
153 the Black Sea (Briski et al., 2024). Though, we emphasize that *Mytilus* spp. taxonomy and
154 distribution are complex due to their cryptic morphology and the frequent occurrence of
155 hybridization between species (Springer & Crespi, 2007; Gaitán-Espitia et al., 2016). The
156 protected habitat, Schleimünde (Maasholm), has been a nature reserve since 1972 (Verein
157 Jordsand zum Schutze der Seevögel und der Natur e.V., 2019). According to its ‘EU Water
158 Framework Directive’ status, the Schlei system is in a good chemical condition (LLUR, 2001;
159 Alpert et al., 2015). Though, we emphasize there were no chemical concentrations reported. The
160 human-altered habitat was the Kiel fjord, with three locations sampled, depending on the species:
161 i) downtown Kiel; ii) Kiel canal; and iii) Falckenstein beach (Figure 1). Downtown Kiel is a
162 completely artificial habitat with concrete structures, while the entire Kiel fjord, including
163 downtown Kiel, is heavily impacted by the shipping industry and tourism, and exposed to
164 extensive pollution including heavy metals such as copper, zinc, tin, lead, as well as tributyltin
165 (LLUR, 2001; Nikulina et al., 2008). The levels of copper, zinc, tin, and lead were higher deeper
166 in the Fjord, with values increasing from 40, 80, 4 and 40 mg/kg at Falckenstein beach to 100,
167 240, 16 and 100 mg/kg at downtown Kiel, respectively (Nikulina et al., 2008).

168 Individuals of at least two populations of each species were collected: one population in
169 the protected habitat and one or more in a human-altered habitat. In the case of *G. salinus*, three
170 populations were sampled in human-altered habitats alongside a protected one. In this case, we
171 aimed to determine if there was a difference in performance among populations adapted to
172 various human-altered habitats along a perceived gradient of anthropogenic influence. Species
173 were collected from May 2016 until January 2021, depending on when a particular

174 species/population experiment was conducted as well as our sampling logistics. In the majority
175 of cases, experimentation was in the same year and season for population comparisons from an
176 individual species (Table S1). After collection, the animals were transported to the laboratories at
177 GEOMAR Helmholtz-Zentrum für Ozeanforschung Kiel in their ambient water and acclimatized
178 for at least two weeks before the experiments commenced. During the acclimatization period,
179 animals were kept in their ambient water in 56 L glass aquaria in the laboratory environment
180 (separately per species/population; Table S1). Each aquarium was aerated through a spongy
181 filtration system to maintain water quality. The light regime was 12:12h light and dark. Mussels
182 were fed every three days with commercial live marine phytoplankton (Premium Reef Blend,
183 Sustainable Aquatics), while gammarids were fed weekly with a mixture of commercial
184 crustacean food (Tetra Mix, Tetra Crusta, and Dr. Shrimp Healthy).

185

186 **Experimental design**

187 Two different types of experiments were conducted for each population of each species, with
188 two crossed stressors: i) temperature – pCO₂ (i.e., partial pressure of CO₂ in water); and ii)
189 temperature – salinity (Table 1). In the case of the temperature – pCO₂ experiments, a fully
190 crossed factorial design consisting of two temperature levels and four pCO₂ levels was
191 performed (Table 1). The temperature – salinity experiments consisted of three temperature and
192 two salinity levels (Table 1). The temperature and pCO₂ values were based on current and future
193 predicted values for the Baltic Sea, as well as occasional measurements in Kiel fjord due to heat
194 waves and seasonal upwelling events (Thompsen *et al.*, 2012; Pansch *et al.*, 2018; Meier *et al.*
195 2022). The lower salinity value was based on the regularly measured salinity of the sampled
196 locations, with the higher salinity constituting a rare but ecologically relevant level measured

197 during winter months at the impacted habitats (Kazanavičiūtė et al., 2024; Briski, personal
198 observations).

199 In both types of these multiple-stressor experiments with factorial stressor arrangements,
200 following the acclimatization period to the laboratory environment, the stress tests were
201 conducted using ten mean-size adult individuals per replicate; except in two experiments when
202 eight and nine individuals were used due to a lack of individuals (i.e., in a temperature – pCO₂
203 experiment with *Mytilus* sp. from Schleimünde and a temperature – salinity experiment with *G.*
204 *locusta* from Schleimünde, respectively). The experimental individuals were transferred from the
205 acclimation conditions to the experimental conditions without gradual adjustments (see Tables 1
206 and S1). Three replicates were conducted for each treatment. Mussel experiments were
207 performed in 5 L aquaria, whereas those for gammarids were in 2 L aquaria containing artificial
208 habitat structures to provide complexity of habitat. One aquarium represented one replicate per
209 treatment. The experiments lasted for 30 days, with mortality checked daily. At that time, dead
210 individuals, molted exoskeletons (in the case of gammarids), and newly emerged juveniles (in
211 the case of gammarids) were removed. Light and feeding regimes were kept the same as during
212 the acclimation period. The experimental conditions and treatments differed among
213 species/experiments, but analyses were performed within species (i.e., comparisons were made
214 among intraspecific populations per experiment), and therefore our approach was not
215 confounded by these disparities.

216 In the first type of experiment, i.e., temperature – pCO₂ experiments, the aquaria were
217 placed in water baths at the desired temperature, with pCO₂ levels being randomly assigned
218 within each water bath, and water continuously infused with its determined pCO₂ level using an
219 air stone. The water used to fill the tanks was filtered from the surrounding area of the institute

220 (filtered through 5 µm filter). The aquaria were sealed using a plastic cover to stabilize the pCO₂
221 levels. Throughout the experiments, the salinity was maintained at the ambient condition for
222 each species/population to reduce any additional stress. Temperature, pH (WTW pH 3110 with a
223 SenTix 81 pH electrode), and salinity (WTW Cond 3110 with a Tetracon 325 probe) were
224 recorded daily. Every ten days, 50% of the water was exchanged with previously prepared water
225 to ensure the desired temperature and pCO₂ conditions were maintained. In the second type of
226 experiment, i.e., temperature – salinity, the aquaria were placed in water baths at the desired
227 temperature, with salinity levels being randomly assigned within each water bath. The aquaria
228 were constantly aerated. Salinity and temperature were checked daily, with a 50% water change
229 done every ten days using previously prepared water to ensure the desired salinity and
230 temperature conditions.

231

232 **Statistical analyses and visualization**

233 Mortality curves were constructed for each population for each treatment, described by the
234 equation (Briski et al. 2008, 2011):

$$235 \quad y=100/[1+e^{-Z(t-Q)}] \quad (1)$$

236 where Z is the slope of the mortality rate, t is time and Q is the onset of mortality. All curves
237 were constructed using S-Plus 6.1 (S-Plus[®] 6.1, 2002; Insightful Corp., Seattle, WA, USA).

238 Raw data supporting the findings of this study is available at Dryad under doi:

239 <https://doi.org/10.5061/dryad.3tx95x6qx>.

240 For each species and experiment separately, generalized linear models with binomial
241 error distributions and log links were used to examine survival rates among populations,
242 temperatures, and either salinities or pCO₂ levels at day-15 (mid-point) and day-30 (end-point).

243 Accordingly, twelve models were built in total (3 species \times 2 experimental contexts [temperature
244 + pCO₂/salinity] \times 2 time points). Non-significant terms were removed backward and stepwise
245 to obtain the most parsimonious models. In cases of complete separation (i.e., total survival or
246 mortality in an experimental group), bias reductions were employed to fit the model to the data
247 (Kosmidis, 2020). Residuals were checked for overdispersion via comparisons of simulated and
248 observed residual distributions (Hartig, 2020). Analysis of deviance with type-3 sums of squares
249 was used to compute coefficients where there were more than two populations in a given model
250 (i.e., models considering *G. salinus*). Tukey comparisons were used to examine significant
251 effects and interactions pairwise post-hoc (Lenth, 2020). Significance was inferred at $p < 0.05$,
252 and generalized linear models were fit in R (R Core Development Team, 2022).

253

254 **RESULTS**

255 *Mytilus* sp.

256 In response to temperature and pCO₂ gradients, significant differences for *Mytilus* sp. were
257 observed between populations from human-altered and protected habitats (Table S2), across both
258 time points (Figure 2). Mortality in the population from human-altered habitat tended to peak at
259 the highest pCO₂ level, while mortality in the population from protected habitat peaked at the
260 lowest pCO₂ levels, driving a significant two-way interaction between population and pCO₂
261 (Table S2). Significant population-level differences were also observed between groups from
262 human-altered and protected habitats in response to the experimental temperature and salinity
263 gradients (Table S2), with the protected site population exhibiting greater mortality in all groups
264 at both time points — except at the highest temperature and salinity (Figure 2). In that treatment,
265 there was a much more rapid increase in mortality in the population from the human-altered

266 habitat, where it overtook the population from the protected habitat, as evidenced by a significant
267 three-way interaction term (Table S2).

268

269 *Gammarus locusta*

270 In response to temperature and pCO₂ gradients, population-level responses by *G. locusta* were
271 significantly influenced by pCO₂ at both time points (Table S3), with the population from the
272 protected habitat tending to exhibit higher mortality (Figure 3). Mortality rates of the population
273 from human-altered habitat tended to decrease with increasing pCO₂, whereas the opposite was
274 observed for the population from protected habitat (Figure 3). Therefore, population-level
275 mortalities were most divergent at the highest pCO₂ levels. In contrast, no significant inter-
276 population differences were observed between those from protected and human-altered habitats
277 across temperature and salinity groups at either time point (Table S3). For both populations,
278 mortality significantly increased with temperature and with falling salinity (Figure 3).

279

280 *Gammarus salinus*

281 In the temperature and pCO₂ experiments, *G. salinus* exhibited significant differences according
282 to temperature and population in interaction at day-15 independently of pCO₂ (Table S4).
283 Mortality rates at day-15 in the population from the most impacted human-altered habitat (i.e.,
284 downtown Kiel) decreased with warming, while the other two populations (i.e., Kiel canal and
285 Falckenstein beach) exhibited higher mortality with greater temperature (Figure 4). At day-30, a
286 significant three-way interaction emerged, whereby differences between populations from the
287 most impacted human-altered habitat (i.e., downtown Kiel) and less impacted (i.e., Kiel canal)
288 and protected habitats (i.e., Schleimünde) were greatest at the lower temperature and higher

289 pCO₂ levels (Table S4; Figure 4). In the salinity and temperature experiments, inter-population
290 responses were influenced by temperature and salinity in interaction at day-15, but not at day-30
291 (Table S4). Mortality rates of all populations except the one from the protected habitat (i.e.,
292 Schleimünde) increased with greater temperature and at the lower salinity at day-15. These
293 differences were less obvious by day-30, with mortality rates for human-altered and protected
294 habitat population differences more consistent (Figure 4). The population from the most
295 impacted human-altered habitat (i.e., downtown Kiel) and that from one of the less impacted
296 (i.e., Falckenstein beach) had significantly lower mortality, with differences most pronounced
297 between populations from those sites and populations from the other less impacted (i.e., Kiel
298 canal) and protected habitats (i.e., Schleimünde) at the highest temperature and lower salinity.
299 The population from the most impacted human-altered habitat had significantly lower mortality
300 than all other populations overall.

301

302 **DISCUSSION**

303 Stresses associated with urban areas are pervasive, however, empirical testing of hypotheses
304 linked to adaptation to anthropogenically modified habitats has remained scarce. Our study
305 tested the performance of different populations of the same species of mussels and gammarids
306 inhabiting impacted and protected habitats, and demonstrated that populations from impacted
307 habitats typically performed better under a broad range of environmental conditions than those
308 from the protected habitat. The populations from protected habitat were consistently less robust,
309 or plastic, to multiple anthropogenic stressors. Based on stress performance and phenotypic
310 plasticity, our results indicate that populations in urban, impacted habitats are adapting to life in
311 anthropogenic environments.

312 Our study determined distinct differences between populations from impacted and
313 protected habitats, with impacted environments producing populations more tolerant to multiple
314 abiotic stressors. However, there was also an apparent stress response gradient within
315 populations from urban areas of differing degrees of habitat alteration. This suggests that the
316 development of resistance along anthropogenic gradients is continuous and not binary. Indeed, in
317 the case of both gammarid species, the outer locations in the Kiel fjord at both the Falckenstein
318 beach and Kiel canal sites produced populations less robust to stressed environments, with *G.*
319 *salinus* exhibiting the highest tolerance at the most impacted and innermost downtown Kiel site.
320 Falckenstein beach is sandy, lacking concrete and areas of altered habitat, however, the location
321 is still exposed to frequent shipping transits throughout the year and the tourist industry in
322 summer months. Kiel canal is located deeper in the Kiel fjord and contains a large amount of
323 concrete as well as ship traffic. The most impacted site, downtown Kiel, is a primarily concrete
324 artificial habitat, exposed to diverse anthropogenic impacts from the shipping industry, high
325 heavy metal concentrations, as well as light and noise pollution and tourism (LLUR, 2001;
326 Nikulina *et al.*, 2008). Consequently, while specific environmental conditions driving adaptation
327 require elucidation at these sites, it seems that the protected nature of a habitat does not play the
328 main role in shaping the robustness of taxa. Rather, resistance is determined by the intensity of
329 stressors present in a habitat, likely linked to the distance of the habitat from our cities and
330 industries, and possibly the potential for taxa to move to surrounding refugia. While we made
331 these inferences based on co-occurring temperature, pCO₂, and salinity stressors, which are
332 pervasive in the Baltic Sea context, we acknowledge that other stressors linked to different forms
333 of pollution (e.g., noise, light, heavy metals) require examination. Nevertheless, cross-tolerance
334 among heterologous stressors is possible (Todgham *et al.*, 2005), with the ability of one stressor

335 to increase tolerance to a second heterologous stressor, and therefore our stressor selection could
336 approximate adaptive responses to other stressors.

337 As we did not test the mussel population from Falckenstein beach or the Kiel canal, but
338 only from downtown Kiel, our study cannot show if mussels from these ‘intermediate’ impacted
339 sites demonstrate the same pattern as gammarids. Nevertheless, mussels are sessile filter feeders
340 and once their larvae are settled on a substrate, it is almost impossible for individuals to move
341 further to more hospitable areas and seek refugia (Ruppert *et al.*, 2003). In contrast, gammarids
342 are mobile detritivores that can actively retreat to less hostile environments when needed (Ruppert
343 *et al.*, 2003; Gerhardt *et al.*, 2011). Therefore, we speculate that the mussel population from
344 Falckenstein beach might have been more robust to stressors than the population from the
345 protected habitat. We would speculate the same for other sessile organisms, such as macroalgae
346 and seagrass. We encourage future studies to test our assumptions and assess taxonomic
347 differences in stress resistance and adaptation among taxonomic groups across varying
348 impacted habitats.

349 Occurrences of anthropogenically impacted populations underperforming individuals
350 from the protected site were rare. However, in the highest temperature – salinity treatment, the
351 population of *Mytilus* sp. from the impacted habitat exhibited a significantly faster mortality rate
352 than the population from the protected habitat. As all replicates demonstrated similarly high
353 mortality rates, we believe that there was no laboratory effect causing this rapid mortality of the
354 population, but it was probably due to the combination of stressors being too impactful and
355 exceeding the threshold of the population’s stress tolerance. Similarly, in the case of *G. locusta*
356 in the combination of the higher temperature (*i.e.*, 24 °C) with all pCO₂ levels, the mortality of
357 both populations was remarkably fast, with no survival displayed after two weeks. These

358 examples further support population-level nuances while also highlighting that even populations
359 from anthropogenically-stressed habitats are susceptible to abiotic limits. Hostile environments
360 place extra pressure on organisms, as modifications from optimal environments require
361 additional energetic costs because of, for example, a higher rate of osmoregulation in suboptimal
362 salinity regimes and an increase in metabolic rate with elevated temperatures, leading to a
363 decrease in organisms' fitness (Gillooly *et al.*, 2001; Evens, 2008; Pörtner, 2010; Bruno *et al.*,
364 2015; Rivera-Ingraham & Lignot, 2017). In the case of highly hostile environments where the
365 effects of stressors go beyond the threshold of the population's stress tolerance, those additional
366 energetic costs are disproportionately high, and eventually lead to mortality (Gillooly *et al.*,
367 2001; Evens, 2008; Pörtner, 2010; Bruno *et al.*, 2015; Rivera-Ingraham & Lignot, 2017). Future
368 research should elucidate whether gradual acclimatization to these stressors can mediate levels of
369 mortality within the laboratory through comparison with acute stress exposures. The resilience to
370 the degree of variability in environmental stressors over time should also be examined among
371 populations (Morón Lugo *et al.*, 2020).

372 Furthermore, the 'anthropogenically induced adaptation to invade' (AIAI) hypothesis and
373 urban evolution concept propose that contemporary adaptation of populations to human-altered
374 habitats (e.g., cities, shipping ports, fragmented urban habitats) within the native range can
375 promote establishment success and increase levels of impact in non-native regions (Hufbauer *et al.*
376 *et al.*, 2012; Borden & Flory, 2021; Perry & Göttert, 2024). Indeed, human-altered habitats have
377 relatively similar environmental conditions globally and are increasingly prevalent with human
378 population growth, especially in cities (Grimm *et al.*, 2008). Moreover, most long-distance
379 transport happens between two urban habitats (e.g., between two shipping ports; Hulme, 2009;
380 Kaluza *et al.*, 2010; Sardain *et al.*, 2019), and thus, the presence of adapted populations in the

381 human-altered habitats can experience increased transport probability, higher chances of
382 surviving the harsh transport conditions, and a reduced need for further adaptation in the non-
383 native regions following introduction (Hufbauer *et al.*, 2012; Lockwood *et al.*, 2013; Briski *et al.*,
384 2013, 2014, 2018). Finally, due to broader phenotypic plasticity of those populations, they would
385 also have better chances for further adaptation and evolution in new habitats, particularly if new
386 habitats are more variable and different from their native ones (Buczowski, 2010; Dlugosch *et*
387 *al.*, 2015; Cadotte *et al.*, 2017; van Kleunen *et al.*, 2018; Westneat *et al.*, 2019; Haubrock *et al.*,
388 2021). Therefore, adaptations to anthropogenic and urban stressors could contribute substantially
389 to invasion success of non-native species across all stages of the invasion process – uptake in a
390 native region, transport, introduction, establishment, and spread/impact.

391 These altered ecosystems can also act as experimental ‘time machines’ for studying
392 adaptation potential to predicted future global changes, as these habitats, for example, are often
393 warmer and have higher levels of CO₂ than surrounding natural areas (Grimm *et al.*, 2008; Lahr
394 *et al.*, 2018). In our study area, Kiel fjord, for example, is characterized by strong seasonal pCO₂,
395 variability (Thompson *et al.*, 2012). Similarly, large estuaries, which are heavily inhabited by
396 humans and contain numerous shipping ports, such as the Saint Lawrence estuary in Canada, or
397 the Chesapeake Bay and Mississippi and Atchafalaya River estuaries in the United States, reveal
398 similar patterns (Dinauer & Mucci, 2017; Lohrenz *et al.*, 2018; Chen *et al.*, 2020). Moreover,
399 those locations are also characterized by huge salinity and temperature variations associated with
400 anthropogenic environmental change (Casties *et al.*, 2016; Pansch *et al.*, 2018; Hinson *et al.*,
401 2022; Kazanavičiūtė *et al.*, 2024). On land, a similar pattern could be observed in impacted and
402 protected sites between stressors such as temperature (i.e., urban ‘heat islands’) and pollutants
403 (Flory & Borden, 2021).

404 Following a multiple-stressor framework with factorial stressor arrangements, our
405 experiments demonstrated greater resistance to these stressors and higher phenotypic plasticity
406 by the populations from the impacted habitats, particularly those more closely located to the city
407 and experiencing the greatest disturbance connected to, for instance, the shipping industry. We
408 demonstrate that populations inhabiting urban environments not only endure human-altered
409 habitat conditions, but they are also more resistant to diverse stressors associated with global
410 environmental change. Therefore, we speculate that those populations would be suitable for
411 genetic rescue to increase population fitness through the introduction of new genetic material to
412 endangered populations/species (Tallmon *et al.*, 2004; Whiteley *et al.*, 2015; Fitzpatrick *et al.*,
413 2023). However, caution is needed in translocation of these populations, as they have a higher
414 probability of establishing in non-native areas, especially other urban environments (Hufbauer *et al.*,
415 2012; Borden & Flory, 2021; Perry & Göttert, 2024). Finally, additional studies testing
416 resistance to various stressors of diverse taxa from multiple geographic locations and
417 environments are needed to confirm the generality of our findings.

418

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424

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629

630 Table 1. Outline of experimental parameters used for a) temperature - pCO₂ *Mytilus* sp., *G.*
 631 *locusta*, and *G. salinus*, b) temperature-salinity - *Mytilus* sp., and c) temperature-salinity - *G.*
 632 *locusta* and *G. salinus* experiments.

a) Temperature – pCO ₂ experiments – <i>Mytilus</i> sp., <i>G. locusta</i> , and <i>G. salinus</i>								
Treatment	Ambient	T1	T2	T3	T4	T5	T6	T7
Temperature (°C)	16	16	16	16	24	24	24	24
pCO ₂ (µatm)	400	1 600	2 700	3 500	400	1 600	2 700	3 500
b) Temperature – salinity experiments – <i>Mytilus</i> sp.								
Treatment	Ambient	T1	T2	T3	T4	T5		
Temperature (°C)	16	20	25	16	20	25		
Salinity (g/kg)	16	16	16	27	27	27		
c) Temperature – salinity experiments – <i>G. locusta</i> and <i>G. salinus</i>								
Treatment	Ambient	T1	T2	T3	T4	T5		
Temperature (°C)	17	20	24	17	20	24		
Salinity (g/kg)	14	14	14	25	25	25		

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634

635 **Figure legends:**

636 Figure 1. Sampling locations used in the study, where individuals of species populations were
637 collected. Schleimünde served as the protected site, whereas Falckenstein beach, Kiel canal,
638 and downtown Kiel served as human-altered sites.

639 Figure 2. Percentage mortality for *Mytilus* sp. from protected and human-altered habitats for the
640 temperature-pCO₂ experiment (left) and the temperature-salinity experiment (right).

641 Figure 3. Percentage mortality for *Gammarus locusta* from protected and human-altered habitats
642 for the temperature-pCO₂ experiment (left) and the temperature-salinity experiment (right).

643 Figure 4. Percentage mortality for *Gammarus salinus* from protected and human-altered habitats
644 for the temperature-pCO₂ experiment (left) and the temperature-salinity experiment (right).

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