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1 A coralline alga gains tolerance to ocean acidification over multiple generations of exposure

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13

14 Crustose coralline algae (CCA) play a crucial role in the building of reefs in the photic zones
15 of nearshore ecosystems globally and are highly susceptible to ocean acidification¹⁻³. Yet the
16 extent to which CCA can gain tolerance to ocean acidification over multiple generations of
17 exposure is unknown. We show that while calcification of juvenile CCA is initially highly
18 sensitive to ocean acidification, after 6 generations of exposure the effects of ocean
19 acidification disappears. A reciprocal transplant experiment conducted on the 7th generation
20 where half of all replicates were interchanged across treatments confirmed that they had
21 acquired tolerance to low pH and not simply to laboratory conditions. Neither exposure to
22 greater pH variability, nor chemical conditions within the micro-scale calcifying fluid
23 internally, appeared to play a role in fostering this capacity. Our results demonstrate that reef-
24 accreting taxa can gain tolerance to ocean acidification over multiple generations of
25 exposure, suggesting that this cosmopolitan species could maintain its critical ecological role
26 in reef-formation.

27

28 **Introduction:**

29 A growing body of evidence suggests ocean acidification will greatly reduce the capacity of
30 major reef-forming species to continue calcifying and hence to accrete the calcium carbonate
31 building blocks of reefs². CCA are ecologically important calcareous red macroalgae that
32 help create and cement reefs from the tropics to the poles and provide a settlement substrate
33 for marine invertebrate larvae such as corals, abalones and sea urchins. Coralline algae
34 precipitate high magnesium calcite, one of the most soluble forms of calcium carbonate, and
35 are therefore expected to be highly susceptible to increased dissolution rates under ocean
36 acidification⁴. Juvenile coralline algae are at particular risk, with extremely reduced
37 abundance under ocean acidification^{5,6}. Ocean acidification decreases both linear extension
38 and calcification (hereafter termed growth) of CCA by altering the calcifying fluid chemistry
39 of adult CCA to become less favorable for precipitation of calcium carbonate⁷⁻⁹. Adult
40 coralline algae that are most resistant to ocean acidification tend to be those that can alter
41 their mineralogy¹⁰ or resist changes to their calcifying fluid⁷⁻⁹.

42 It is currently unknown whether benthic calcifying organisms gain tolerance to ocean
43 acidification after exposure for multiple generations. Our understanding of these processes in
44 marine species is limited to either evolutionary responses of short-lived, non-reef-building
45 species such as phytoplankton and non-calcareous polychaete worms over multiple (five to
46 hundreds) generations¹¹⁻¹³, or acclimatization responses of longer-lived species over 1 or 2
47 generations^{14,15}. Most reef-forming species have slower growth rates and generation times
48 than microalgae, which has limited our ability to test whether they can gain tolerance to
49 ocean acidification. However, some tropical reef-building and cementing CCA grow
50 extremely fast to colonize available space after disturbance¹⁶, occupying a broad range of
51 habitats that requires high phenotypic plasticity¹⁷. Some CCA also reproduce sexually or
52 asexually within weeks of settlement, even while still microscopic¹⁸. These traits make CCA
53 valuable model organisms to test whether reef-accreting calcifying species can potentially
54 gain tolerance to ocean acidification over multiple generations of exposure.

55 In many shallow-water reef environments, night-time declines in pH due to local
56 metabolic activity often exceed the changes in open-ocean pH predicted to occur by 2100
57 under business-as-usual CO₂ emissions scenarios (e.g. Representative Concentration Pathway
58 8.5, 0.4 pH units)^{7,20,21}. It has been proposed that this natural diurnal variability may impart
59 tolerance to low pH for resident calcifying species via selection for individuals that are better
60 adapted to survive in these environments^{20,22}. Additionally, transgenerational acclimation can
61 occur via a variety of epigenetic or maternal effects after exposure of subsequent generations

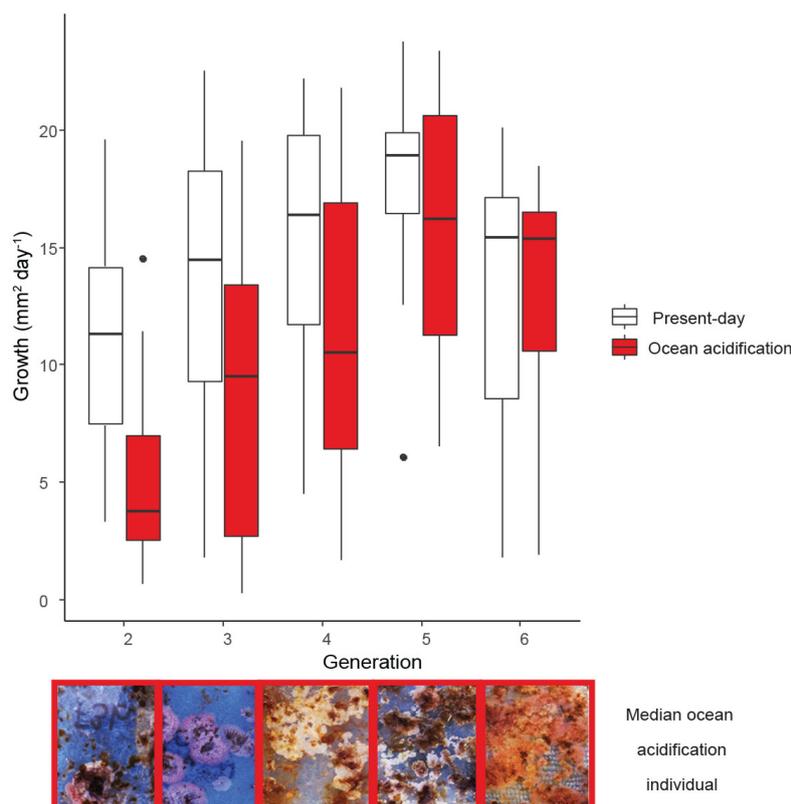
62 to stressors^{14,23}, and could be promoted by prior exposure of individuals to greater variability
63 in that stressor²⁴.

64 We investigate whether the cosmopolitan reef-accreting and cementing species
65 *Hydrolithon reinboldii* (synonymous with *Hydrolithon boergesenii*) can increase its tolerance
66 to ocean acidification after multiple generations of exposure via mineralogical changes in
67 skeletal formation or conversely the ability to resist seawater induced chemical changes to
68 their calcifying fluid. We also test whether this tolerance can be gained faster in individuals
69 with prior or regular exposure to low pH. We grew CCA for 8 generations under present-day
70 (pH 8.00) and future ocean acidification treatments (pH 7.70), with both high (± 0.37) and
71 low (± 0.07) pH variability, using populations from sites with either naturally low or high pH
72 variability (see materials and methods and Table S1) and tracked complex changes in their
73 calcification mechanisms. For the first time, we demonstrate that a species of coralline algae
74 can gain tolerance to ocean acidification after 6 generations of exposure, but neither exposure
75 to greater pH variability, nor changes in their calcification physiology related to mineralogy
76 or chemistry in the calcifying fluid influence this capacity.

77

78 **Changes in sensitivity to ocean acidification**

79 The resistance of juvenile CCA to ocean acidification increased incrementally over
80 each generation during this 423-day experiment. Second generation CCA grew 56.1% slower
81 across all ocean acidification treatments relative to the present-day treatments (Fig. 1). This
82 sensitivity to lower mean pH declined over subsequent generations to 33.1, 29.7, 12.8, and
83 2.0% between generation 2 to 6, supported by the significant interaction between mean pH
84 and generation in the general linear model (Table S2, $F_{1,224} = 8.60$, $p = 0.004$). The difference
85 further decreased to 0.6% when compared to the generation 7 populations kept in their
86 original treatments (Fig. 2).



87

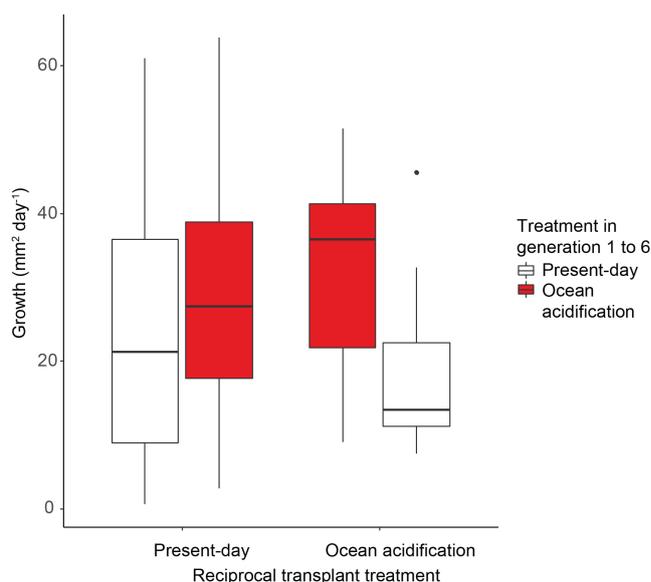
88 Figure 1: Growth of CCA populations after 2 to 6 generations of exposure to present-day (pH
 89 8.00) and ocean acidification (pH 7.70) treatments. Also shown are the median individual
 90 coralline algal plates for each generation under the ocean acidification treatments (out of 24).
 91 Scale 50 by 45 mm of images. Note here responses to constant and variable pH regimes
 92 combined. See supplementary figure Fig. S1 for individual treatment responses. Median and
 93 75% quartiles, mean $n = 24$ per treatment and generation combination.

94

95 Reciprocal transplant experiment

96 A reciprocal transplant experiment was also undertaken to test whether the CCA
 97 gained tolerance to ocean acidification rather than just the aquaria settings. This was
 98 undertaken at generation 7 where half of all replicates were interchanged with replicates of
 99 treatments with the opposite mean pH treatment, while keeping the variability the same as the
 100 previous generation (see methods). Growth was 40.3% lower in generation 7 populations
 101 transferred from the present-day into the ocean acidification treatments compared to those
 102 grown there for 6 generations, and this was 22.2% lower than the extension of control
 103 populations kept within the control treatments (Fig. 2, significant interaction between mean
 104 pH treatment and the treatment of origin: Table S3, $F_{1,79} = 4.6$, $p = 0.035$). The negative
 105 effects of low pH on growth were larger in populations originally from the more variable pH

106 site compared to the stable pH site in early generations over the course of the experiment
 107 (Fig. S1, $F_{1,224} = 4.05$, $p = 0.045$). This is contrary to the hypothesis that individuals from
 108 more variable environments are more tolerant to ocean acidification, or that they gain this
 109 tolerance more rapidly over multiple generations. Trends in total recruit area typically
 110 mirrored those of growth rates, but were inherently more variable (Fig. S2, S3, Table S3–4).
 111



112
 113 Figure 2: Growth of CCA after 7 generations grown in present-day (pH 8.00) and ocean
 114 acidification (pH 7.70) treatments (controls), versus those transferred into these novel
 115 treatments after being in the treatment with the other mean pH for 6 generations. Responses
 116 to constant and variable pH regimes are combined. Median and 75% quartiles, $n = 12$ per
 117 treatment and generation combination.

118

119 **Implications for future nearshore reefs**

120 This is the first demonstration that over multiple generations of exposure benthic
 121 calcifying species can increase their tolerance to ocean acidification, indicating that at least
 122 this species of CCA has the potential to gain tolerance to ocean acidification well within the
 123 timeframe of when its effects will manifest. Although this finding is by-itself critical to more
 124 accurately forecasting reef futures, outstanding questions remain, including the generality of
 125 this finding amongst other coralline algal species and in combination with other stressors
 126 such as ocean warming or in the presence of competitors that could be unaffected by ocean
 127 acidification. However, if coralline alga can develop resistance to ocean acidification over
 128 multiple generations of exposure, it is possible that other calcifying benthic species with

129 similar calcification mechanisms⁹ could have the same capability. This information is crucial
130 for coral reefs, for example, where many coral species are long-lived and experimentally
131 testing their capacity to gain tolerance over multiple generations is logistically challenging,
132 requiring years or decades of exposure to ocean acidification in controlled laboratory settings.

133

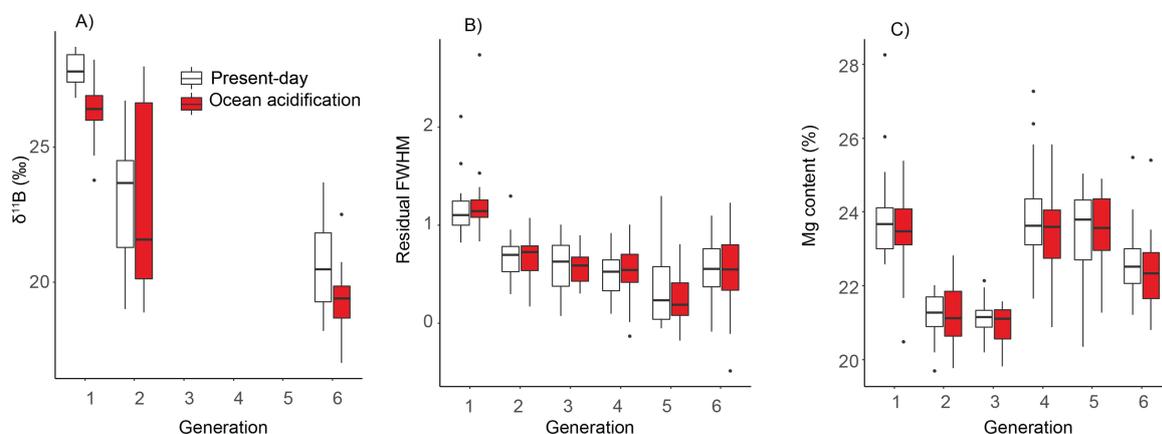
134 **Evolutionary and physiological mechanisms promoting tolerance**

135 The diminishing effect of ocean acidification on growth over each subsequent
136 generation supports the premise that successive cohorts of CCA are subject to strong
137 selection for resistance to ocean acidification, or that transgenerational acclimation occurred
138 on individuals within each tank. We did not aim to determine the underlying evolutionary
139 processes that could alter tolerance to ocean acidification. Previous research indicates that
140 species can gain complete tolerance after a full generation of exposure to climate change
141 stressors¹⁴. Our evidence indicates slower, more incremental increases in tolerance to climate
142 change stressors could also occur in select taxa in the future. However, the trends observed
143 here were also rapid in the context of evolutionary adaptation. The rapid (less than two years),
144 yet incremental, nature of these changes, and the predominately asexual reproduction,
145 indicate that transgenerational acclimatory processes could have played a large role in
146 inducing the changes observed. However, the effects of ocean acidification were particularly
147 strong on the recruitment of the first generations of CCA, and because selective pressure is a
148 combination of the effect size of the stressor selected against and the heritability of the trait,
149 we cannot rule out that selection for individuals better able to calcify under these treatments
150 occurred. While not our aim here, the next step is to further understand the contributions of
151 genetic versus epigenetic processes that contribute to changing tolerance under climate
152 change in reef-building and accreting species.

153 Regardless of the eco-evolutionary processes involved, exposure to greater variability
154 in pH did not foster the capacity to gain tolerance to ocean acidification faster²⁴. This is
155 surprising given that organisms from these habitats tend to respond more positively to ocean
156 acidification^{22,25}, and that phenotypic or transgenerational plasticity is predicted to be higher
157 on these populations^{17,24}. Therefore, transplanting organisms from highly variable pH
158 environments into habitats with constant pH regimes, as has been suggested regarding ocean
159 warming and temperature variability²⁶, might not be a successful strategy to combat against
160 ocean acidification. The CCA also did not alter their calcifying fluid toward conditions

161 traditionally considered favorable for precipitation⁸. Hence, physiological mechanisms that
 162 allow for tolerance in adult CCA could have no bearing on their persistence in future reefs.

163 The physiological mechanisms responsible for reduced sensitivity under the ocean
 164 acidification treatments are complex. Raman spectroscopy analyses indicated that changes in
 165 the mineralogy from high magnesium calcite into less readily dissolved polymorphs such as
 166 dolomite did not occur here (Table S5, Figure 3). Similarly, neither $\delta^{11}\text{B}$ (pH_{cf} proxy) or
 167 Raman full width at half maximum (FWHM: proxy of the calcite saturation state in the
 168 calcifying fluid [Ω_{cf}] when precipitation occurs) were elevated in individuals kept for 8
 169 generations under mean low pH compared to generation 2 individuals (Table S6, S7, Figure
 170 3). In fact, both $\delta^{11}\text{B}$ and FWHM decreased between generations 1, 2 and then the
 171 experimental endpoint, and they tended to be lower in ocean acidification treatments
 172 generally. The physiological ability to maintain elevated pH_{cf} or Ω_{cf} under low pH in adult
 173 coralline algae and corals are traits related to resistance to ocean acidification, and we
 174 hypothesized *a priori* that any CCA here that gained resistance to ocean acidification would
 175 have done so via improving these traits. However, changes in these calcification-related
 176 parameters ($\delta^{11}\text{B}$, FWHM, Mg) do not match changes in growth rates over successive
 177 generations under ocean acidification. The inverse relationship between FWHM and growth
 178 throughout the course of the experiment (Figure S5), together with lack of changes in other
 179 geochemical parameters, indicates that CCA tolerant to ocean acidification here were those
 180 which could calcify when carbonate chemistry is suboptimal for precipitation within the
 181 calcifying fluid, when compared to that of adults or controls. This is likely caused by both the
 182 rapid use of calcium and carbonate during calcification in the semi-enclosed space²⁸, and
 183 declining pH_{cf} via ocean acidification⁸. Additionally, traits such as increasing organic matrix
 184 proteins involved in calcification²⁹ could also contribute to trends observed here.



185

186 Figure 3: a) $\delta^{11}\text{B}$ (proxy for pH in the calcifying fluid), b) FWHM (proxy for calcite
187 saturation state in the calcifying fluid), c) and Mg content of CCA over 6 generations under
188 present-day or ocean acidification treatments. Median and 75% quartiles, mean $n = 24$ per
189 treatment and generation combination.

190

191 **Conclusions**

192 These results highlight that we cannot predict which species will be winners or losers
193 under climate change based solely on their responses to short-term exposure to acute
194 stressors. We demonstrate for the first time that some coralline algae are capable of rapidly
195 gaining tolerance to ocean acidification over six generations. If this capacity is present in
196 other select species of coralline algae, then as a group they could be the largest contributors
197 to global reef accretion in future oceans, given their already high abundance in temperate
198 nearshore reefs, tolerance to ocean warming³⁰, and the recent large-scale declines in the
199 abundance of the longer-lived tropical scleractinian corals³¹.

200

201 **Methods**

202 **Collection and treatment design**

203 We collected adult coralline algae, *Hydrolithon reinboldii* (synonymous with *boergesenii*³²),
204 rhodoliths from Western Australia's northern Kimberley locations of Tallon (16°40S,
205 123°14E) and Shell Islands (16°48S, 123°04E) in October 2016. These two sites are
206 characterised by similar thermal regimes, but different pH regimes. Tallon Island is a
207 seaweed and seagrass dominated reef with strong diel pH variability (up to 1.06 units daily).
208 Shell Island is dominated by *Acropora* spp. corals and experiences a low range in pH
209 variability (typically < 0.10 units daily). This species was chosen because it is the most
210 common species at both sites, and because it was the only species that lives at both sites that
211 can be identified using morpho-anatomical methods, hence retaining intact and undamaged
212 individuals that would not occur if molecular sequencing was required prior to the
213 experiment starting. This species is common in the Indo-Pacific region³³, and was identified
214 with assistance by staff at the Western Australian Museum. Samples are lodged both at
215 Victoria University of Wellington and the University of Western Australia. The collection,
216 environmental variables at the collection sites, and experimental design and monitoring are
217 described in detail within the initial segment of the research on the adults in Cornwall et al.⁷.
218 Briefly, we created 4 treatments, within which each of the two source population were
219 placed: 1) control means (8.00) and low pH variability (daily range of 0.14), or 2) high pH

220 variability (0.58 daily range); 3) ocean acidification means (7.70) and low pH variability
221 (0.20 range daily), or 4) high pH variability (0.89 daily range). See Table S1 for full details,
222 or Figure 1 from Cornwall et al.⁷ for schematic of details. There were 6 independent
223 experimental tanks per treatment combination per population, for a total of 48 experimental
224 tanks per generation. This allows us to determine the interactive effects of changes in both
225 mean and variability in pH. As recommended by all best practices guides³⁴⁻³⁷, we used CO₂
226 gas to lower pH/elevate CO₂ concentrations, and CO₂-free air to increase pH/lower CO₂.
227 Seawater was pumped from an area of soft-sediment bottom 100m offshore at Watermans
228 Bay, Perth, then filtered to 25 µm using a series of sand filters, and then subsequently ran into
229 a header tank where pH conditions were modified before moving into each experimental
230 tank. There were 3 header tanks per treatment. Temperature was maintained in a series of
231 waterbaths at 26 to 27 °C, 4°C warmer than the summer maximum outside of our
232 experimental facility. No recruitment of coralline algae occurred within this header tank,
233 allowing us to be certain that coralline algal recruits throughout our experiment were not
234 sourced from local populations. This temperature (26 to 27 °C) roughly matches that of the
235 collections site in the Kimberley region in Winter³⁸ or is slightly higher, and was chosen to
236 limit the stress that the CCA here would be exposed to. We chose to hold temperature
237 constant during the duration of the experiment to avoid any confounding effects of
238 temperature variations between generations.

239

240 **Multigenerational experiment**

241 On collection, only healthy adult coralline algae were selected that possessed extensive
242 numbers of conceptacles (i.e. reproductive structures where spores are formed). On day 0 (i.e.
243 the collection date), adult coralline algae were extensively cleaned of epiphytes and scrubbed
244 to limit carry-over of other coralline algal spores from the field into the experimental set up.
245 On day 78, coralline algal recruits were visible and on day 128 the adults were removed and
246 the process of creating the generation 3 and subsequent generations began. This involved
247 cutting out two 4.5 by 5 cm (L × W) sections of the tank containing the highest cover of
248 coralline algae recruits and placing these into completely new tanks, attached vertically (Fig.
249 S7). Four of the generation 2 ocean acidification (both high and low variability) replicate
250 tanks contained no visible recruits. For these tanks, populations from different tanks were
251 used to create the third and subsequent generations. Therefore, we used 44 distinct
252 populations of CCA to source the subsequent generations. We repeated this process once
253 CCA recruits in the control treatment obtained the same estimated size as the 2nd generation,

254 roughly every 41 to 51 days. Once visible, CCA grew rapidly, approaching 1 cm diameter per
255 week. The CCA on plates would then develop conceptacles and release spores which disperse
256 into the water column around 2 weeks after being placed into brand new experimental tanks
257 not previously used. A standardised size was used rather than a standardised time because of
258 slight seasonal differences in the duration required to reproduce during each generation. This
259 was process was continued until generation 7. This resulted in using 336 separate
260 experimental tanks (7 ×48) throughout the course of this study.

261 We acknowledge that it is possible that reproduction could be occurring in recruits even when
262 microscopic (e.g.¹⁸), so in theory each generation could possibly be comprised of small
263 amounts of differing generations. This would result in the number of generations here being
264 more than the 8 recorded. However, we consider that extremely unlikely. This is because the
265 experimental tanks were monitored daily, and we did not observe more rapid recruitment than
266 would be expected if the recruits were reproducing before they became visible (i.e. weeks), or
267 if time to reproductive maturity was variable, as recruits became visible in all tanks within the
268 same weekly time frame when recruitment occurred. This consistent time to visible
269 recruitment also means it was highly unlikely that recruitment occurred in generation 2 from
270 spores of other species carried-over from the field. All coralline algal recruits possessed
271 similar morphoanatomical features, including similarly sized and shaped conceptacles that
272 aligned with the adults' identification³⁹. Thousands of conceptacles were produced in each
273 generation, and all that we analyzed contained multiple spores indicating that they were
274 asexually reproducing⁴⁰. Damaged conceptacles were not analyzed. In other words, the algae
275 were asexually reproducing which is most common in coralline algae⁴¹, especially for this
276 species for which only these life history stages have been noted³⁹. The probability of skipping
277 sexual reproduction in many species has previously been suggested⁴¹. A proportion of
278 conceptacles were not fully formed at the end of each generation or their contents were
279 already emptied or they were damaged, for these we cannot fully confirm the life stage or
280 nature of reproduction.

281 We measured the linear extension of CCA on the created replicate plates using images
282 taken every two weeks and analysed in ImageJ. This step involved analyzing 2496 images.
283 We also assessed the total cover of recruits at the end of every generation. The trends in total
284 area occupied by coralline algae in the tanks, and the growth on monitored plates were
285 similar. However, it became apparent that the total area of recruitment was affected by both
286 the treatment effect (mean pH) on growth and random variability in the number of recruits.
287 Therefore, we used the change in total area of the individuals of created plates as the focal

288 measurement of interest, as it contained one response variable and not two (i.e. growth and
289 recruitment success). See supplementary Figure 2 for these data, which follow similar trends
290 but are inherently more variable. This step involved analyzing 1440 images. A random
291 selection of more than 10% of all images were analyzed twice to check for consistency
292 between multiple assessors.

293

294

295

296 **Reciprocal transplant experiment**

297 The process of exposing subsequent generations to their original treatment was continued in
298 the 48 populations until generation 7, whereupon we conducted a reciprocal transplant to
299 determine whether improved growth in ocean acidification treatments over time were
300 representative of altered response to the treatments themselves, or an interaction between lab
301 effects and altered response to treatments²³. We exchanged half of the ocean acidification
302 replicates with half of the ambient replicates within the same variability treatments, i.e. pH
303 7.70 low pH variability replicates interchanged with pH 8.00 low pH variability replicates
304 and vice versa. We did this because of the observed changing effects of mean pH over
305 subsequent generations and lack of effects of pH variability. Therefore, we were not
306 interested in the effects of switching individuals' treatments with differences in pH
307 variability, only in how switching them between different mean pH treatments would
308 influence fitness. When we create the additional novel or original treatments (i.e.
309 "transferred" into novel treatments or placed in the same treatment as "controls") we have 16
310 different treatment combinations. We assessed growth over the first 2 weeks after transferring
311 the generation 7 individuals into novel and control treatments, before they became limited for
312 further space. We also measured total recruitment of their offspring after 61 days. We
313 removed all standardised coralline algal plates from their tanks for the same length of time
314 irrespective of whether they were transferred into a new treatment or placed into a control
315 treatment.

316

317 **Measurement metrics**

318 We also measured proxies to aid in understanding changes in calcification physiology of the
319 coralline algae at the completion of the experiment. $\delta^{11}\text{B}$, a proxy for pH at the site of
320 calcification, was measured on generations 1, 2, 6, 7 and 8 to determine trends over
321 generations and due to treatments using previously published methods⁷⁻⁹, with the only

322 difference being bleaching durations of 3 times as long. We also measured Raman
323 spectroscopy on every generation as a proxy of the saturation state of calcium carbonate
324 (based on ν_1 Full Width at Half Maximum; FWHM) and magnesium content (based on ν_1
325 wavenumber), following previously described protocols^{7,9}.

326

327

328

329 **Statistical analysis**

330 We used general linear models in R to determine differences in growth, recruit area, and
331 geochemical proxies across all available generations assessing the full interaction between
332 mean pH (control or ocean acidification), pH variability (low or high), generations (2, 3, 4, 5,
333 6), and site of origin (Shell or Tallon). Waterbath was used a random factor in models to
334 account for this potential source of variability. For generations 7 and 8, we used the same
335 model structure, except the generation factor was replaced with the transfer (novel treatment
336 or original control treatment) treatment and each generation was assessed separately, and
337 waterbath was removed from the model. We used a log-fit mixed model for recruit area of
338 generations 2 to 7, due to non-normally distributed data.

339

340 **Data availability**

341 Data is available at dryad: <https://doi.org/10.5061/dryad.pzgmsbcfq>.

342

343 **Code availability**

344 All R code will be available upon request.

345

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355

356 **Author contributions**

357 CEC and SC conceived and ran the experiment and wrote the manuscript. CEC ran the
 358 statistical analysis and conducted the boron isotope measurements. MTM conceived the
 359 experiment, provided vital laboratory equipment, facilities and resources. TMD conducted
 360 the Raman spectroscopy. EL, BM, KG, FP, and QD all ran the experiment and measured
 361 seawater carbonate chemistry. All authors edited the manuscript, provided intellectual input,
 362 and agreed to its submission.

363

364 **Conflicting interests**

365 We have no conflicts of interest.

366

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