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Lipid consumption in coral larvae differs among sites: a consideration of environmental history in a global ocean change scenario

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The success of early life-history stages is an environmentally sensitive bottleneck for many marine invertebrates. Responses of larvae to environmental stress may vary due to differences in maternal investment of energy stores and acclimatization/adaptation of a population to local environmental conditions. In this study, we compared two populations from sites with different environmental regimes (Moorea and Taiwan). We assessed the responses of *Pocillopora damicornis* larvae to two future co-occurring environmental stressors: elevated temperature and ocean acidification. Larvae from Taiwan were more sensitive to temperature, producing fewer energy-storage lipids under high temperature. In general, planulae in Moorea and Taiwan responded similarly to pCO$_2$. Additionally, corals in the study sites with different environments produced larvae with different initial traits, which may have shaped the different physiological responses observed. Notably, under ambient conditions, planulae in Taiwan increased their stores of wax ester and triacylglycerol in general over the first 24 h of their dispersal, whereas planulae from Moorea consumed energy-storage lipids in all cases. Comparisons of physiological responses of *P. damicornis* larvae to conditions of ocean acidification and warming between sites across the species' biogeographic range illuminates the variety of physiological responses maintained within *P. damicornis*, which may enhance the overall persistence of this species in the light of global climate change.

1. Introduction

Early life-history stages of corals are increasingly threatened by global anthropogenic stressors, such as ocean acidification (OA), the decrease in pH and carbonate ion concentration caused by the absorption of anthropogenic CO_2 by the surface ocean, and ocean warming. For coral larvae, many biological traits and processes vary with pCO$_2$ conditions, with exceptions [1–7]. The biological responses of coral larvae to ocean acidification and concurrent warming will likely affect their dispersal through the depletion of energy stores and a reduction in larval buoyancy [8–10]. Being lecithotrophic, most coral larvae use lipids and protein as endogenous sources of energy throughout dispersal [8,10,11]. Translocated metabolites from *Symbiodinium* are also important sources of energy for dispersing larvae [12,13]. Temperature stress can decouple the physiology of symbiosis in coral larvae (*Porites astreoides*, [14]), and the thermotolerance ceiling could be worsened by the addition of OA. Additionally, in response to OA and warming, upregulation of stress response pathways and increased maintenance of homeostatic processes may necessitate increased consumption of energy stores (i.e. lipid and protein) or shifts in the allocation of

metabolic energy. Trade-offs of these responses to environmental stress include slowed developmental timing, reduced growth and impaired competency to settle (e.g. [14–16]). Furthermore, reduction of lipid stores will decrease larval buoyancy, altering the physical transport of larvae by currents [17]. As a consequence of reduced energy stores and buoyancy, larvae may disperse shorter distances or may terminate in the plankton, with negative consequences for suitable habitat selection, recruitment success, restoration of damaged populations and region-wide connectivity.

At the population level, the effects of OA and warming may be mediated by local environmental conditions. Adaptation to local regimes of pH and temperature occurs over a sufficient spatial scale where environmental conditions differ consistently and where gene flow is low [18–20], and in this context, local adaptation to temperature has been well studied [21,22]. Studies in other systems provide evidence that local adaptation to pH regimes can occur in nature [23–25]. If coral populations are locally adapted to their environmental regime, corals living in a naturally more acidic environment may have developed physiological tools for maintaining normal function in low-pH seawater, as has been shown for elevated temperature [26]. The role of environmental history in shaping the response of coral populations to combined effects of OA and warming remains unknown.

In this study, we assessed the ecological development ('the mechanisms of developmental regulation in real-world environments', [27]) of *P. damicornis* larvae from populations with different environmental histories. Specifically, we examined the effects of OA and temperature on larval performance. To do this, we performed CO_2 and temperature manipulation experiments with *P. damicornis* larvae collected in Moorea, French Polynesia and in Taiwan. These sites were chosen based on our experience working at these locations and *a priori* knowledge of differing oceanographic regimes, the former being a fringing reef in a sheltered lagoon and the latter experiencing tidally driven upwelling events [28]. We used biochemical composition as an index of performance. Lipid composition and biological traits of larvae were measured both upon release and after exposure to experimental conditions in order to assess how OA and warming might alter larval performance and dispersal. Environmental data using autonomous pH and temperature sensors deployed in both locations provided context for experimental conditions. For this study, we asked: (i) do the effects of OA and warming on lipid composition and other physiological traits of *P. damicornis* larvae differ between cohorts released at separate biogeographic locations and (ii) do differences in the effects of OA and warming correspond with the environmental history of each coral population? We hypothesized that responses of coral larvae to conditions of OA and warming would differ between locations and that the responses to pCO_2 and temperature would correspond with environmental history of these parameters.

2. Material and methods

(a) Collection of coral larvae

During the austral summer in 2012, eight colonies of *P. damicornis* (electronic supplementary material, figure S1) were collected from fringing reef sites in Moorea, French Polynesia and Taiwan, respectively, at approximately 1–3 m depth (17.4803 S,

149.7989 W; 21.9385 N 120.7967 E). At the Richard B. Gump South Pacific Research Station and at the National Museum of Marine Biology and Aquarium (NMMBA) in Taiwan, larvae were collected and pooled following [29] the lunar pattern of reproduction of *P. damicornis* [30]. The 'peak' larvae, determined by the predicted release curve of the colonies (28 February 2012 in Moorea and 25 June 2012 in Taiwan; e.g. [6,31,32]), were used in the experiment. In Moorea, all larvae released were used in the experiments, resulting in unequal genotype ratios. In Taiwan, equal numbers of larvae from each colony were contributed to this pool. The condition of the freshly released larvae was assessed using lipid metrics and other physiological parameters (see below). The remaining larvae in each pool were then randomly assigned to experimental treatments.

(b) Experimental incubations

Larvae were divided among eight tanks containing four treatment combinations of pCO_2 and temperature. Larvae were incubated for 24 h under experimental conditions in two 400 ml containers per aquarium at approximately 0.15–0.25 larva ml^{-1}. These containers had 100 µm mesh sides and a photosynthetically active radiation (PAR)-transparent lid and were anchored in place within the aquarium to ensure that PAR exposure was replicated across tanks. Owing to the time needed to photograph and preserve larvae post-incubation, incubations were staggered by 1 h per aquarium, with the order randomized. At the end of each incubation, larvae within tanks were pooled, and 10 larvae were randomly selected for size measurements ($n = 20$ per treatment). The remaining larvae were frozen at $-80°C$ in aliquots for downstream analyses of lipid classes (3×25 larvae), total protein content (2×5 larvae) and symbiont density (2×5 larvae). Therefore, for comparisons between treatments, $n = 6$ for lipid classes, $n = 4$ for total protein and $n = 4$ for symbiont density.

In laboratories at both sites, two pCO_2 and two temperature treatments were prescribed: ambient temperature–ambient pCO_2 (ATAC), ambient temperature–high pCO_2 (ATHC), high temperature–ambient pCO_2 (HTAC) and high temperature–high pCO_2 (HTHC). The ambient CO_2 treatment (approx. 400–450 µatm pCO_2) approximated current environmental conditions of the water mass bathing the fringing reef where the adult corals were collected (confirmed by environmental data; [32]). The control temperatures (27.5–28°C) approximated the multi-year average temperature for the fringing reefs close to the collection sites for adult *P. damicornis* ([33], T.-Y. Fan January 2007 to March 2009, March 2010 to present, unpublished data). The high CO_2 and high temperature treatments (approx. 900–1000 µatm pCO_2, 30.5–31°C) represent ocean conditions expected by the year 2100 under a business-as-usual scenario [34]. See the electronic supplementary material for detailed information on experimental culturing conditions.

To verify and monitor the physical parameters of the OA × temperature treatments, the chemistry of the seawater in the aquaria was analysed during the experiment. pH, temperature, salinity and total alkalinity of seawater in each aquarium were measured during the incubations following best practices (see the electronic supplementary material).

(c) Assessment of cellular lipids

Lipids were extracted from larval homogenates following [35,36] (see the electronic supplementary material for additional details). Total lipid values were obtained from lipid extracts evaporated at 37°C under nitrogen gas. Lipid classes of wax ester (WE), triacylglycerol (TG) and phospholipid (PL) were quantified using two chromatography-based techniques. For larvae sampled in Moorea, a thin-layer chromatography-flame ionization detection analyser was used (Iatroscan MK-5, Iatron Laboratories, Inc., Tokyo, Japan) following [29]. For larvae sampled in Taiwan, these

Table 1. Summary of physical conditions in experimental aquaria used in Moorea, French Polynesia and in Taiwan. Data are presented as mean $\pm$ s.e. For all parameters, $n = 6$.

site	treatment	temperature (°C)	salinity (psu)	pH$_{total}$	A$_T$ (μ.mol kg^{-1})	pCO$_2$ (μ.atm)
Moorea	ATAC	27.99 $\pm$ 0.10	35.82 $\pm$ 0.02	8.004 $\pm$ 0.003	2362 $\pm$ 1	459 $\pm$ 4
	HTAC	30.57 $\pm$ 0.11	35.85 $\pm$ 0.02	7.970 $\pm$ 0.007	2363 $\pm$ 1	496 $\pm$ 9
	ATHC	28.12 $\pm$ 0.18	35.83 $\pm$ 0.02	7.719 $\pm$ 0.008	2364 $\pm$ 1	1002 $\pm$ 22
	HTHC	30.77 $\pm$ 0.20	35.88 $\pm$ 0.02	7.692 $\pm$ 0.012	2368 $\pm$ 2	1068 $\pm$ 33
Taiwan	ATAC	27.75 $\pm$ 0.18	30.61 $\pm$ 0.18	7.971 $\pm$ 0.007	2234 $\pm$ 4	494 $\pm$ 9
	HTAC	30.53 $\pm$ 0.05	30.63 $\pm$ 0.23	7.978 $\pm$ 0.005	2253 $\pm$ 6	487 $\pm$ 9
	ATHC	27.70 $\pm$ 0.20	30.26 $\pm$ 0.29	7.733 $\pm$ 0.006	2204 $\pm$ 15	923 $\pm$ 14
	HTHC	30.43 $\pm$ 0.04	30.93 $\pm$ 0.14	7.748 $\pm$ 0.004	2243 $\pm$ 6	900 $\pm$ 9

lipid classes were quantified using thin-layer chromatography (TLC; see the electronic supplementary material). Phospholipids were quantified using a spectrophotometric assay for phosphorus determination (modified from [37], see the electronic supplementary material).

(d) Characterization of other larval physiological parameters

To assess other aspects of physiology of incubated larvae, total protein, density of *Symbiodinium* and larval size were quantified. The Bradford assay was used to quantify total protein ([38,39], following [6]). *Symbiodinium* density was quantified using a haemocytometer (see the electronic supplementary material). Larval circumference (area) and maximum length were determined from photographs of larvae [40]. For comparisons to other publications, larval volume was calculated following [41] and is included in electronic supplementary material, table S1.

(e) Statistical analyses of biological data

All data were analysed using R version 3.0.1 (R Core Team 2013). Statistical assumptions of normality and homogeneity of variance were met based on Q-Q plots and Levene's test, sometimes following an inverse transformation. Physical experimental conditions were compared using type III sum of squares, with pCO$_2$ and temperature treatments as fixed factors and tank as a random factor. All pre- and post-incubation quantities of lipid classes were standardized to total lipid. For initial conditions of larvae, lipid and other physiological metrics were analysed for effect of Site using a one-way ANOVA. Because initial biological metrics of larvae often differed by site, we compared the change in these metrics over the first 24 h of larval duration between pCO$_2$ exposures, temperature exposures, and sites. To do so, mean pre-incubation levels were subtracted from post-incubation levels, henceforth referred to as Δ, 'delta'. Therefore, a negative ΔWE per larva represents a net consumption of WE during the 24-h incubation. Absolute values of lipids and other parameters are provided in the electronic supplementary material, tables S1 and S2.

Linear mixed-effect models (nlme package in R; [42]) were used to estimate effects on response variables, with pCO$_2$, temperature and site as fixed factors and aquarium (tank) as a random factor. Model selection was performed incrementally, following [43]. Likelihood ratio tests (type III sum of squares) were conducted on selected models fit using maximum likelihood in order to compare the effects of fixed factors [44,45]. Post-hoc analyses were performed using orthogonal contrasts (multcomp package in R; [46]). General linear hypothesis tests with Bonferroni corrections (GLHT) were used for multiple comparisons, and Tukey's HSD was used for single comparisons.

(f) Environmental data collection

pH and temperature time series were generated on fringing reefs in Moorea, French Polynesia and in Taiwan, within 33 m from the collecting locations of adult *P. damicornis* parents (see the electronic supplementary material). A full comparison of environmental regimes between sites can be found in [32].

3. Results

At two sites, we measured the effects of OA and warming on lipid composition and other physiological traits of *P. damicornis* larvae, following a 24-h experimental exposure under laboratory conditions where temperature and seawater chemistry were controlled (table 1). Treatment levels were statistically distinct at each site (see the electronic supplementary material, table S3). Although some incidences of larval death were observed during the incubations, these events were aquarium-specific and were not consistent across treatments or locations.

(a) Lipid composition of *Pocillopora damicornis* larvae

Larvae in Moorea contained on average 32.00 μg total lipid per larva, including 58% WE, 11% TG, and 17% PL, and a mean of 11.65 μg total protein per larva. Larvae in Taiwan contained on average 20.47 μg total lipid per larva, including 39% WE, 18% TG and 6% PL, but a larger total protein fraction, a mean of 22.37 μg per larva.

In *P. damicornis* larvae immediately after release, initial levels of WE and PL, as a proportion of total lipid, did not vary significantly with Site (see the electronic supplementary material, table S4). However, initial TG varied significantly by Site ($p < 0.001$; electronic supplementary material, table S4). Larvae in Moorea had significantly less TG than those in Taiwan (Tukey's HSD, electronic supplementary material, table S4 and figure S2). Total lipid varied significantly by Site ($p = 0.009$; electronic supplementary material, table S4); levels in Moorea were greater than those in Taiwan (Tukey's HSD, electronic supplementary material, table S4 and figure S2).

In response to conditions of pCO$_2$ and temperature, ΔWE ranged from -11.47 μg per larva (MCR HTHC; 62% depletion of initial levels) to $+17.64$ μg per larva (TWN ATAC; more than 100% increase from initial levels). When normalized by total lipid, ΔWE varied significantly only by

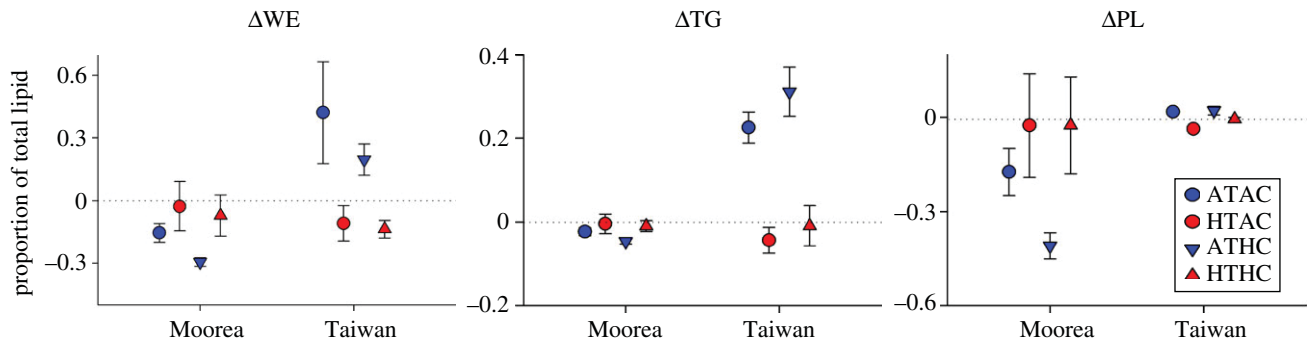**Figure 1.** Changes in lipid composition of *P. damicornis* larvae over 24-h exposures to combinations of pCO_2 and temperature. Mean $\pm$ s.e. ($n = 6$) changes in abundance of WE, TG and PL classes for larvae released in Moorea, French Polynesia and Taiwan. Negative Δ values indicate net decreases in per cent lipid composition over the experimental exposure. Lipid quantities are standardized by total lipid. Experimental treatments: Ambient-T, Ambient-pCO_2 (ATAC), High-T, Ambient-pCO_2 (HTAC), Ambient-T, High-pCO_2 (ATHC), High-T, High-pCO_2 (HTHC).

T × Site and Site ($p < 0.001$ for each; see the electronic supplementary material, table S5). Larval WE content increased significantly at Ambient-T for larvae in Taiwan, but not in Moorea (Tukey's HSD, see the electronic supplementary material, table S6; figure 1).

ΔTG varied between $-2.38\,\mu g$ per larva (TWN HTHC; 42% depletion of initial levels) and $+7.57\,\mu g$ per larva (TWN ATHC; more than 100% increase from initial levels). Only the effect of Site was significant for ΔTG, which was standardized by total lipid ($p = 0.005$; electronic supplementary material, table S5). TG became a larger portion of total lipid for larvae in Taiwan over larvae in Moorea (Tukey's HSD, electronic supplementary material, table S6), particularly noticeable under Ambient-T (figure 1).

ΔPL ranged from $-4.00\,\mu g$ per larva (MCR HTAC; 75% depletion) to $+2.99\,\mu g$ per larva (MCR HTAC; 56% increase). Normalized to total lipid, ΔPL varied significantly by T × Site, T and Site ($p = 0.010$, $p = 0.002$, $p < 0.001$; respectively; electronic supplementary material, table S5). Larvae in Moorea experienced greater depletion of PL at Ambient-T than at High-T, while in Taiwan, ΔPL was similar between temperatures (GLHT, electronic supplementary material, table S6; figure 1).

ΔTL varied between $-18.22\,\mu g$ per larva (MCR HTAC; 57% depletion) and $+21.65\,\mu g$ per larva (MCR ATAC; 68% increase); $pCO_2 \times T \times$ Site and $pCO_2 \times T$ were significant effects ($p = 0.013$, $p = 0.041$, respectively; see the electronic supplementary material, table S7).

(b) Other larval physiological parameters

In *P. damicornis* larvae immediately after release, average *Symbiodinium* density was 7850 cells per larva in Moorea and 9008 cells per larva in Taiwan. Neither total protein (TP) nor symbiont density differed by Site (electronic supplementary material, table S4). Larval area and length varied significantly by Site ($p < 0.001$ for each; electronic supplementary material, table S4), with larger larvae released in Taiwan (Tukey's HSD, electronic supplementary material, table S4 and figure S2).

Changes in response to different pCO_2 and temperature levels were measured in the form of TP, symbiont density and larval size. ΔTP ranged from $-7.43\,\mu g$ per larva (TWN HTHC; 56% depletion) to $+7.53\,\mu g$ per larva (TWN ATAC; 34% increase). Total protein content of larvae did not vary significantly by pCO_2, T or Site (electronic supplementary material, table S7). Changes in symbiont abundance ranged from -6614 cells per larva (TWN ATAC; 73% decrease) to

$+8558$ cells per larva (TWN HTAC; 95% increase). However, symbiont density did not vary significantly by any of the effects tested (electronic supplementary material, table S7). Changes in larval area varied between 69% growth (TWN ATAC) and 64% decrease (TWN HTAC). Only the effect of Site explained a significant amount of variation in this parameter ($p < 0.001$ electronic supplementary material, table S7), with significantly more growth in larval area in Moorea than in Taiwan (Tukey's HSD, electronic supplementary material, table S8; figure 2). Changes in larval length ranged from $-864\,\mu m$ (TWN ATAC) to $+785\,\mu m$ (MCR ATHC); significant effects included pCO_2 and Site ($p = 0.043$, $p = 0.038$, respectively; electronic supplementary material, table S7). Larvae became significantly shorter in length in Taiwan versus Moorea (Tukey's HSD, electronic supplementary material, table S8; figure 2) and at Ambient-pCO_2 versus High-pCO_2 (Tukey's HSD, electronic supplementary material, table S8; figure 2).

4. Discussion

(a) Variation in maternal investment of *Pocillopora damicornis* larvae

Measurements of biochemical and biological traits of newly released *P. damicornis* larvae provide critical contextual information about the physiological condition of freshly released larvae and inform the link between maternal investment and larval performance. *P. damicornis* larvae in Moorea received more total lipid, though energy-rich lipid classes were equal to or less dense than for larvae released in Taiwan. Based on these initial conditions, particularly for WE, larvae released at both locations contained similar long-term energy stores, which may influence buoyancy and dispersal potential. Additionally, the proportion of total lipid concentrated in long-term storage (WE) versus short-term storage (TG) indicates that in order to mount a physiological response to environmental changes, such as OA, larvae from Moorea will likely need to mobilize part of their WE pool, sacrificing their buoyancy in the process. Larvae from corals of the Taiwan population contained more TG upon release. Maternal investment may prime these larvae for more immediate response to rapidly changing environmental conditions [29]. If utilization of long-term lipid stores is required as well, their dispersal distance may be diminished. In Moorea and elsewhere, brooded *P. damicornis* larvae commonly disperse several kilometres
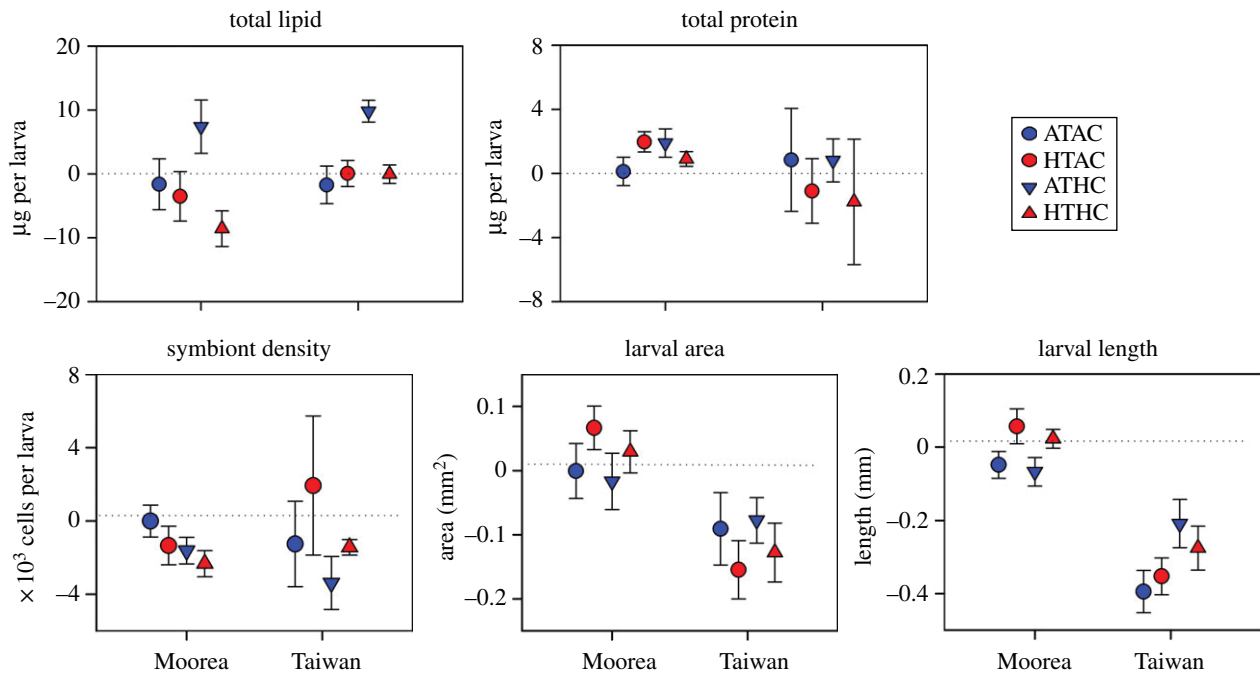
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Figure 2. Changes in larval physiological condition for *P. damicornis* larvae following 24-h exposures to $p\text{CO}_2$ and temperature. Mean $\pm$ s.e. ($n = 6$) changes in larval composition and size for larvae released in Moorea, French Polynesia and Taiwan. Negative Δ values indicate net consumption over the experimental exposure. Physiological condition traits are standardized per larva (μg per larva), unless otherwise noted. Experimental treatments: Ambient-T, Ambient-$p\text{CO}_2$ (ATAC), High-T, Ambient-$p\text{CO}_2$ (HTAC), Ambient-T, High-$p\text{CO}_2$ (ATHC), High-T, High-$p\text{CO}_2$ (HTHC).

away from the natal reef, and these populations have genetic similarity [47,48]. However, local retention is also frequently observed [48], and larvae of this species likely do not successfully recruit to other islands in French Polynesia [47].

With equal symbiont densities upon release, larvae of the two populations had similar potential to produce new energy, in the form of fixed carbon, assuming they contained the same clade of *Symbiodinium* with similar photophysiology. However, the clades of *Symbiodinium* may have differed between sites, likely appropriate for the local environmental conditions of each location [49–51].

Larval size correlated with total protein and was greater in larvae from Taiwan. Large *P. damicornis* larvae can survive for a longer period of time in the plankton than small larvae and can therefore potentially disperse farther [41]; however, in this case, it is unknown whether planula size was correlated with lipid or protein content. A longer dispersal period may allow planulae to discover preferred or novel habitats for settlement, but at the cost of extended predation risk and removal from favourable parental habitat [52–54]. Suggested by this study, larvae from Taiwan may not have a longer dispersal distance because their larger size did not correspond to larger stores of long-term energy biomolecules (WE).

(b) Responses of larval lipid consumption and other physiological traits to changes in $p\text{CO}_2$ and temperature

Changes in energy stores of coral larvae under future ocean change may affect coral populations through shifts in dispersal distance and recruitment success. TG can be quickly hydrolysed for immediate energy needs and is the first lipid class to be consumed during larval development or during periods of starvation (e.g. [55]). WE has a slower turnover rate and serves as long-term energy deposits for larvae [8,56,57]. In addition, WE plays an important role in dispersal of coral larvae, governing buoyancy (e.g. [57]) and entraining larvae in surface currents [58]. If TG stores are consumed, larvae will use WE, but their vertical position in the water column will change as a result, with potentially negative consequences for dispersal.

Additionally, *Symbiodinium* dinoflagellates in brooded larvae, vertically transmitted to larvae prior to release, satisfy a significant portion of larval energy budgets [12]. Changes in lipid content of planulae, like *P. damicornis*, represent the balance between demands of holobiont metabolism and the carbon translocation by *Symbiodinium*. In particular, WE consumption increases when endosymbiont metabolites are not available [12].

When lipid utilization was quantified in *P. damicornis* planulae over 24 h under experimental conditions, a range of responses across treatments and biogeographic sites was observed. For example, in Moorea at High-T, High-$p\text{CO}_2$, planulae depleted up to half of their energy-storage lipids while in Taiwan at Ambient-T, Ambient-$p\text{CO}_2$, other planulae more than doubled their deposits of these lipids. Assuming constant rates of respiration over a 24-h period (Peak larvae, Moorea, approx. $0.08–0.13$ nmol O_2 per larva per min [6]), larvae would need to consume $1.33–2.16$ μg WE over the incubation [59]. Larvae in this study consumed less than or equal to 1 μg WE per symbiont density, but photosynthetic activities and consumption of TG may have accounted for the remaining energy burned during respiration. The net production of WE and TG during experimental exposures was likely due to the photosynthetic activities of *Symbiodinium*. Given the ecological significance that energy-storage lipids play in the successful dispersal of these lecithotrophic larvae [8–10], the point when larvae stop accumulating lipid and start depleting lipid stores may be a useful indicator of stressful environmental conditions in future work.

6

Higher temperatures, but not pCO$_2$, may affect larval dispersal for some populations in the future, due to changes in WE that alter buoyancy of planulae (e.g. [57]). For Peak planulae in this study, WE content was sensitive to temperature, depending on the site, but not to pCO$_2$ (figure 1). In Taiwan, under Ambient-T, WE content increased over the incubation period, while larvae at High-T experienced net depletion. Metabolic rates of planulae increase with temperature [4,6], so utilization of lipids, from stores or from photosynthesis, would also be expected to rise. However, in Moorea, WE content of larvae did not respond to changes in temperature. Increased primary productivity of *Symbiodinium* at High-T or increased rates of translocation may explain this pattern, as *P. damicornis* in Moorea tend to host a more thermotolerant *Symbiodinium* clade [50,51] than *P. damicornis* in Taiwan [49]. Putnam *et al.* [60] observed no change in the photophysiology of *P. damicornis* planulae in response to elevated pCO$_2$ and temperature; however, rates of translocation of fixed carbon can increase in response to low pH and elevated temperature [61,62]. Translocated carbon could be used to satisfy immediate energy demands rather than oxidation of stored WE [12].

The utilization of WE may occur when demand for energy surpasses the pool of TG (e.g. [63]). In general, TG content remained constant with respect to temperature and pCO$_2$ levels. Differences in TG content between sites followed the pattern of maternal investment of this energy-storage lipid. In Moorea, the energetic demands of early dispersal surpassed the combined resources of TG synthesis and translocated fixed carbon, perhaps due to smaller initial TG stores for larvae from Moorea (figure 1). The smaller pool of TG was likely partially maintained through oxidation of WE [64,65]. In Taiwan, planulae had higher TG content, particularly under Ambient-T. Larvae from Taiwan may contain a surplus of TG, due to higher maternal investment. Conversion of TG to WE under Ambient-T may have caused the increase in WE observed under these conditions. Overall, the species-level patterns of lipid utilization and production indicate that *P. damicornis* planulae are launching a physiological response to conditions of warming, but not OA, and this response is at least partially fuelled by the utilization of WE and TG. As a result, planulae store less lipid under warming conditions, particularly planulae in Taiwan, and therefore may have fewer resources for performing the developmental changes associated with metamorphosis and settlement (e.g. [66]).

Our measurements of utilization of energy-storage lipids represent the net activity of the planula holobiont (animal + symbiont). The symbiont cells themselves contain lipid that contributes to the measured amount for the holobiont (e.g. [67]). Because symbiont density did not vary between sites, pCO$_2$ levels or temperature levels, differences in function of the *Symbiodinium*, in addition to changes in host physiology, may have contributed to patterns of lipid utilization and production (figure 1). The photosynthetic activities of *Symbiodinium* as well as their translocation rates of glycerol and lipid bodies containing WE and TG potentially change the amount of fixed carbon received by the host [68,69] and therefore the amount of lipid available.

Traits of size and growth were not consistently significantly affected by elevated pCO$_2$ and temperature. Planula size in *P. damicornis* may not be uniformly sensitive to OA and warming due to the contribution of *Symbiodinium* to larval energy metabolism; this input of exogenous energy may even allow larvae to grow in size during dispersal, a possibility not

available to aposymbiotic lecithotrophic (non-feeding) larvae. Firstly, PL levels were higher at warmer temperatures, but only for larvae in Moorea. Secondly, larval area and length were greater for larvae released in Moorea versus Taiwan, though this pattern differed from that of PL. If this trend of reduced planula size over the first 24 h of dispersal in Taiwan continues throughout the larval stage, it may carry over to reduce the stringency of habitat selection for settlement, increase post-settlement survival and growth, and increase the age of first reproduction [52,70,71]. However, even though larval size decreased over the 24-h exposures in Taiwan, absolute larval size still remained larger than larvae in Moorea. While smaller planula size at the beginning of settlement would be a fitness disadvantage [72,73], this may not be the case for planulae from Taiwan, which maintained their symbiont density, TL and TP, despite the decrease in larval size.

(c) The role of environmental history and geographical site

The study sites in Moorea and Taiwan had different regimes of temperature and pH [32] that have presumably persisted over the recent history of the study populations. Corals in Moorea experienced, on average, warmer temperatures with a narrower range of values. On the other hand, corals in Taiwan experienced lower mean pH and a greater range of pH values. In both sites, autonomous sensors deployed on natal fringing reefs [32] confirmed that the Ambient treatment conditions were observed within the water mass bathing the fringing reef during the experiment, while the High treatment conditions were not. Overlaid on the between-site differences in environmental conditions are the genetic differences between *P. damicornis* populations [74,75], which may indicate the possibility of local adaptation to environmental conditions. Although not predictive, our results shed light on the possibility that the physiological plasticity of *P. damicornis* planulae to resist future ocean conditions may be influenced by the environmental conditions to which its population is adapted or acclimatized [19]. In general, our results can be used to evaluate some expectations based on a local adaptation scenario: planulae from Moorea should perform better than those from Taiwan under High-T, and planulae from Taiwan should perform better than those from Moorea under High-pCO$_2$. In support, for example, we found that larvae from Taiwan were more sensitive to temperature, producing fewer energy-storage lipids under High-T. In Moorea, planulae at High-T consumed similar amounts of lipid as at Ambient-T, suggesting that their demands for stored energy were not elevated, perhaps offset by increased carbon translocation from their symbionts. Contrary to expectations from a local adaptation scenario, in general, planulae in Moorea and Taiwan responded similarly to pCO$_2$. Additionally, corals in the study sites with different environments produced larvae with different characteristics, which may play a role in the different physiological responses observed.

5. Conclusion

Comparisons of physiological responses of *P. damicornis* larvae to OA and warming between sites across the species' biogeographic range improve our understanding of the possible future success of this species. An overarching outcome of this study is that site-specific aspects of planula physiology provide a framework for describing and understanding the

consequences of environmental stress. For example, the effects of temperature on WE, long-term energy-storage lipids with an important role in larval buoyancy, differed based on biogeographic site. Because larvae from Moorea are endowed with more total lipid and thus have higher absolute amounts of WE upon release, they may be able to accommodate depletion of these stores without sacrificing their dispersal potential or the energy reserves required to complete metamorphosis and settlement. Our results support expectations for performance based on a local adaptation scenario. While there is still much to learn about the interplay of environmental history and population genetics, the variety of physiological responses maintained within *P. damicornis* may enhance the overall persistence of this species in the light of global climate change.

Data accessibility. The biological and environmental datasets supporting this article are publicly available in the LTER Network Data Catalog (portal.lternet.edu) as well as the MCR LTER local data catalogue, with the identifier knb-lter-mcr.5022.

Authors' contributions. E.B.R. conceived of the study, designed the study, carried out the experiments and associated laboratory sample analyses, carried out the statistical analyses and drafted the manuscript. C.-S.C. and T.-Y.F. helped coordinate the study. H.-H.L. participated in the design of the lipid analysis and helped coordinate the study. G.E.H. helped design and coordinate the study and helped draft the manuscript. All authors gave final approval for publication.

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