

Research



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Reduced thermal tolerance of massive coral species in a highly variable environment

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Coral bleaching events are increasing in frequency and severity, resulting in widespread losses in coral cover. However, branching corals native to highly variable (HV) thermal environments can have higher bleaching resistance than corals from more moderate habitats. Here, we investigated the response of two massive corals, *Porites lobata* and *Goniastrea retiformis*, from a moderately variable (MV) and a low variability (LV) pool transplanted into a HV pool on Ofu Island in American Samoa. Paired transplant and native ramets were exposed to an acute thermal stress after 6 and 12 months of exposure to the HV pool to evaluate changes in thermal tolerance limits. For both species, photosynthetic efficiency and chlorophyll loss following acute heat stress did not differ between ramets transplanted into the HV pool and respective native pool. Moreover, HV native *P. lobata* exhibited the greatest bleaching susceptibility compared to MV and LV natives and there was no effect of acute heat stress on MV *P. lobata*. There was also a thermal anomaly during the study, where Ofu's backreef thermal regime surpassed historical records—2015 had 8 degree heating weeks (DHW) and 2016 had up to 5 DHW (in comparison to less than or equal to 3 over the last 10 years)—which may have exceeded the upper thermal limits of HV native *P. lobata*. These results strongly contrast with other research on coral tolerance in variable environments, potentially underscoring species-specific mechanisms and regional thermal anomalies that may be equally important in shaping coral responses to extreme temperatures.

1. Introduction

The frequency and magnitude of environmental variation is increasing in the upper ocean [1] as our global climate rapidly warms. Environmental variability strongly influences organismal physiology and behaviour [2,3], community assemblages [4] and ultimately the integrity of ecosystems [5]. Impacts of climate warming are further magnified in marginal/extreme environments, such as low, high, or highly variable (HV) temperature, pH, and/or CO₂ sites [6,7]. However, a number of studies show organisms in variable environments may have enhanced tolerance compared to those in more moderate habitats owing to acclimatization or adaptation [8–12]. Alternatively, warm-adapted species in these extreme environments may be particularly at risk because they live closest to their upper thermal limit and may have limited acclimation capacity [13–15]. Although these populations have probably evolved the greatest thermal tolerance, it is possible an increased cost is involved in maintaining this tolerance [15] compared to other populations with lower tolerances [16]. Such a trade-off is critical for understanding the susceptibility of these populations to climate change.

Tropical reef-building corals live close to their upper thermal limits and are particularly sensitive to periods of elevated sea surface temperatures (SST) [17,18]. Despite coral vulnerability to climate impacts, marginal and extreme reef habitats contain assemblages of corals that have acclimated and/or adapted to survive near or at their thermal thresholds [11,19–21]. Resident coral populations in these environments are continuously exposed to HV abiotic

conditions, yet coral diversity remains high [20] and upper-temperature tolerances are significantly higher than conspecifics from higher latitudes [18] or less variable environments [8–11,21–24]. Mechanisms that contribute to high heat tolerance result from increased prevalence of heat-tolerant photosymbionts (*Durusdinium* spp. family Symbiodiniaceae; [25], but see [26,27]), modifications in gene regulation [28,29], adaptive divergence between coral populations [23,26,30–32] and/or potential epigenetic contributions to thermal tolerance [32,33]. As a result, HV habitats have become popular natural laboratories to understand the capacity of and mechanisms underlying coral stress tolerance [6,8,22].

One such system that has been extensively studied is the network of backreef pools within the National Park of American Samoa on Ofu Island. These backreef pools are nearly identical in species diversity and per cent live coral cover, yet have distinct differences in small-scale environmental variability driven by tidal cycle and pool size [10,20,34]. Coral populations from two pools—a small, HV and a larger, moderately variable (MV) pool—exhibit both fixed and acclimatory responses to HV temperatures that contribute to enhanced thermal tolerance [8]. However, much of the research examining coral resilience in Ofu and elsewhere has been conducted on thermally susceptible branching corals, such as *Acropora* spp. [8,24,35–38]. Thus, there is scant evidence on whether massive, more robust corals exhibit similar responses to increasing environmental variability [22,39].

Additionally, evidence of tolerance trade-offs in organisms from HV habitats has been documented in intertidal porcelain crabs [14,15] and snails [40], diving beetles [16], and seaweeds [41], but the potential negative impacts of extreme environments are largely unknown for tropical reef-building corals. Broadly, trade-offs in stress tolerance can result in reduced fecundity [42,43] and growth [41,44], changes in basal gene expression [42], transgenerational effects on offspring size and metabolism [45], and a limited scope for further acclimation to warmer temperatures [15,16]. For corals, the few documented consequences of elevated heat tolerance trade-offs involve reduced lipids, growth and eggs size (attributed to hosting *Durusdinium*; [46,47]) and reduced larval size [33]. However, we do not know whether similar or extensive trade-offs apply to corals in naturally extreme environments, and what the implications would be for future reef habitats in a warming world.

Here, we test the scope for thermal tolerance in two dominant massive coral species, *Porites lobata* and *Goniastrea retiformis* in the Ofu backreef during an extremely warm year. We compare growth, bleaching sensitivity and endosymbiont species assemblage (Symbiodiniaceae) of coral samples transplanted into the HV pool compared to corals in the neighbouring MV and an additional nearby backreef pool of lesser thermal variability, the low variability (LV) pool. Corals were exposed to controlled, acute heat stress experiments at 6 and 12 months following transplantation to characterize upper thermal limits, acclimation capacities and trade-offs in this extreme environment.

2. Material and methods

(a) Coral collection and transplantation

In July 2015, corals were sampled from three backreef sites (HV, MV and LV) within the National Park of American Samoa of

Ofu Island (14.1780765°S, 169.660109°W). Thirty colonies ($n = 5$ genets site⁻¹ species⁻¹) of two common massive coral species, *P. lobata* and *G. retiformis*, were sampled to remove 24 cores/ramets from each genet in each site ($n = 360$ cores total species⁻¹). Cores were measured for initial buoyant weight, secured to transplant grids via nylon bolts (approx. 36–40 cores grid⁻¹), and returned to the respective native site for a one-week recovery. Ramets were then divided equally and transplanted into either the HV pool common garden or returned to the native reef site ($n = 12$ cores genet⁻¹ site⁻¹ species⁻¹; figure 1). HOBO pendant temperature loggers (Onset Computer Corp.) were deployed on native and transplant grids at all three sites and collected temperature data every 15 min. During January 2016, the LV native sample grid was dislodged by a cyclone but found a few days later and re-secured, precluding the six-month native versus transplant pairwise comparisons.

(b) Acute heat-stress assays

At each time point—6 months (January 2016) and 12 months (July 2016) after transplantation—2 ramets genet⁻¹ species⁻¹ were collected from the grids in each backreef pool (5 genets * 2 ramets = 10 ramets species⁻¹ * 2 species = 20 ramets origin_dest⁻¹ * 5 origin_dest [LV_LV, LV_HV, MV_MV, MV_HV and HV_HV] = 100 ramets total). Cores were scrubbed to remove algal and epiphyte growth prior to buoyant weight measurements. Coral growth was calculated by subtracting initial weight from final weight and then divided by the number of weeks since transplantation to determine weekly growth rate.

Coral ramets were placed in our Coral bleaching automated stress system (CBASS), constructed from sets of head and sump tanks (42 l volume treatment⁻¹), resulting in four experimental tank systems—two heat and two control. A pump provided a flow of 88.9 ml s⁻¹ to each head tank, which was also fitted with six LED bulbs (500 μmol photons m⁻² s⁻¹ ± 20 μE as measured via a Li-COR Li192 spherical quantum sensor) and 12 h 7.00 light/19.00 dark photoperiod. A flow-through drip system provided 9 l h⁻¹ of local seawater throughout the duration of the experiment.

Following previous experiments by Palumbi *et al.* [8], 60 ramets (approx. 30 cm³; four from each genet) were randomly assigned to one of two control and two heat treatment tanks ($n \sim 10$ –15 ramets tank⁻¹) and then subjected to a customized temperature-controlled ramp programme [49]. All ramets from a single species were assayed in 1 day, with the second species assayed the following day. Beginning at 11.00, temperature increased over 3 h from 28 to 36.5°C for *P. lobata* and to 35.5°C for *G. retiformis*, followed by a 3 h incubation at the maximum temperature, then a ramp down to and hold at 28°C for 16 h (electronic supplementary material, figure S4). The control tank was set to remain stable at 28°C for 22 h. The two maximum temperatures were chosen: based on preliminary trials to elicit a visible bleaching response in greater than 50% of fragments, to represent acute thermal exposures above the local bleaching threshold, and to be approximately 1°C above the HV pool's mid-day low tide average maximum temperature.

(c) Symbiodiniaceae physiology under heat stress

The maximum quantum yield (F_v/F_m) of Photosystem II was measured using a pulse amplitude modulation (PAM) fluorometer (Junior-PAM, Walz, Germany; settings in the electronic supplementary material, Information). Following 30 min of dark adaptation, tops of coral ramets were measured in triplicate at the beginning (0 h) and before the end (21 h) of the experiment. Normalized F_v/F_m values (21–0 h)/0 h were used for statistical analyses to correct for between ramet variation in starting values. F_v/F_m values measured at the end of each assay were used for plotting for simplicity to allow for easy comparison to previous studies.

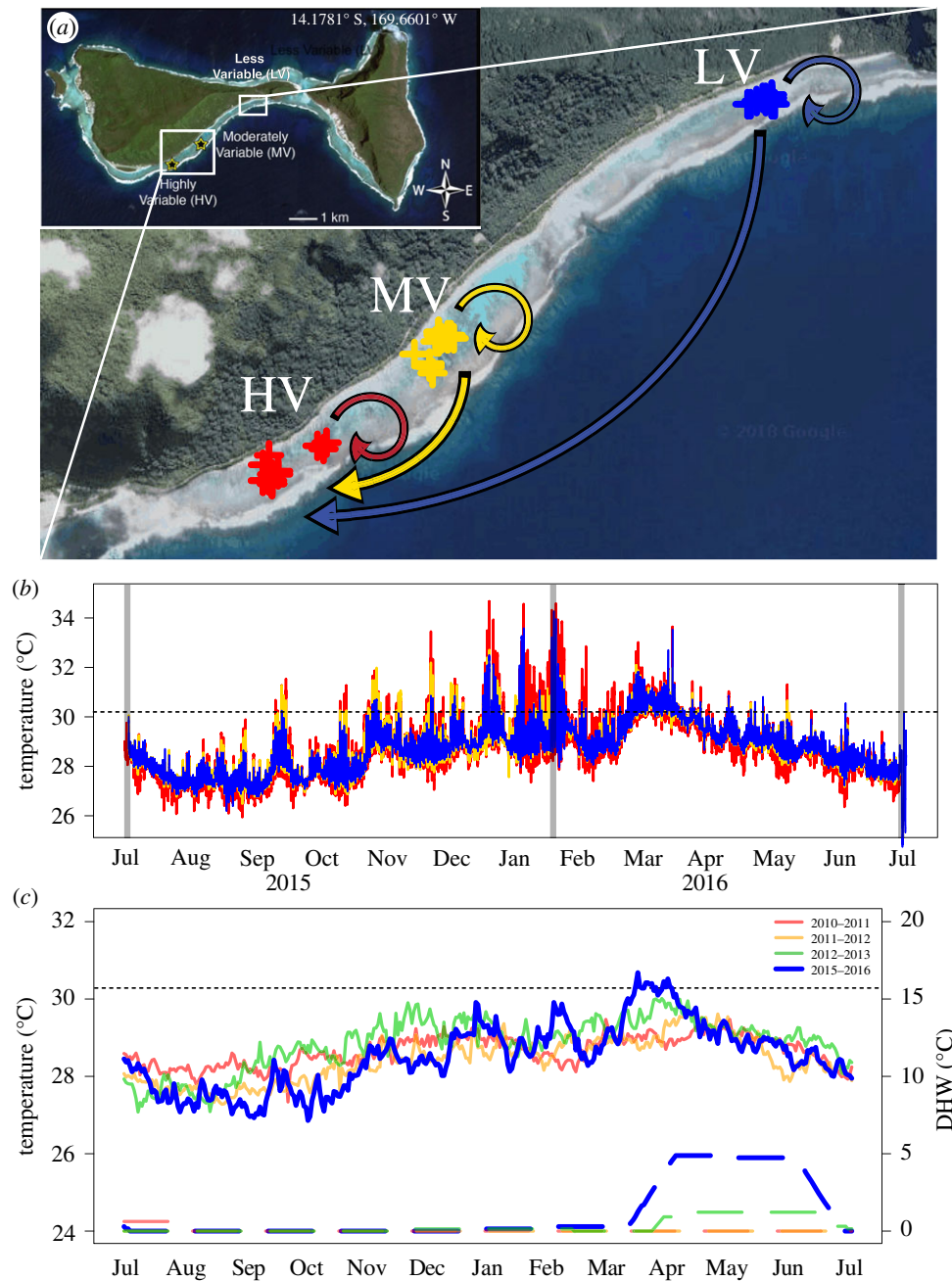


Figure 1. (a) Map of Ofu Island, American Samoa. Arrows show transplant experiment design within three backreef pools—HV (red), MV (gold), LV (blue). (b) *In situ* site temperatures during the study period. Vertical grey lines represent the start of the experiment and data collection time points. (c) Comparison of sea surface temperatures (solid lines) and degree heating weeks (DHW; dashed lines) during years of this and a prior experiment. Temperatures were extracted from a National Oceanic and Atmospheric Administration, Coral Reef Watch 5 km Ofu Island dataset spanning the years 2010–2012 [8] and 2015–2016 (this study). Dotted lines represent the regional bleaching threshold, 30.2°C [48]. (Online version in colour.)

Following experiments, coral tissue was airbrushed from the skeleton using 35 ppt artificial seawater, and the resulting slurry was homogenized, centrifuged and resuspended in 5 ml of seawater. For chlorophyll determination, slurry samples were homogenized using 90% acetone, a glass tissue homogenizer, and a 25 mm GF/F filter, and then stored at 4°C for 24 h. Absorbance spectra were measured using an Ocean Optics Spectrometer, and chlorophyll *a* and *c* values calculated using the Jeffrey & Humphrey [50] equation. Total chlorophyll (*a* + *c*) absorbance was normalized to acetone volume and then scaled to the surface area of each ramet, as measured using the paraffin wax method [51].

(d) Symbiodiniaceae genotyping

A 1 cm² biopsy was sampled from each *G. retiformis* and *P. lobata* genet ($n = 5 \text{ species}^{-1} \text{ site}^{-1}$) at the end of the experiment (control

ramets only) and in the field at each time point. Samples were incubated for 1–1.5 h at 65°C in a 1% sodium dodecyl sulfate in DNABuffer [52] and then transported back to Old Dominion University. During both time points, similar sized biopsies were sampled from control ramets after acute experiments to characterize Symbiodiniaceae ITS2-level assemblages over time (0, 6 and 12 months). DNA was extracted from the archived coral samples using a guanidinium-based extraction protocol [52].

Symbiodiniaceae were identified from both native coral colonies and corresponding HV pool transplanted replicates at 6 (January) and 12 months (July). We used a 350 bp segment of the internal transcribed spacer region 2 (ITS2) ribosomal DNA (rDNA) for amplification. The ITS2 region was amplified using Symbiodiniaceae specific primers, ITS-Dino-forward [53] and its2rev2-reverse [54]. Each primer also contained a universal linker, for downstream incorporation of Illumina adapters and

barcodes. Primer sequences and polymerase chain reaction (PCR) are specified in the electronic supplementary material. Following barcoding PCR, samples were pooled and sequenced on ODU's Illumina MiSeq (250 bp paired-end Reagent Nano Kit v.2).

Sequenced raw reads were demultiplexed, then trimmed of barcodes, adapters, linkers, ITS2 primers and degenerate bases. Sequences were identified via comparisons against available Symbiodiniaceae databases (<http://webhome.auburn.edu/~santosr/sequencedatasets.htm>; [55]) and confirmed against NCBI's nucleotide BLASTn reference database. Symbiodiniaceae abundance counts for each amplicon sequence variant (ASV) per sample were produced using the R program DADA2.1.8.0 [56], and analysed using the MCMC.OTU model as described in Green *et al.* [57], with fixed effects for origin, destination and time. Pairwise differences between all fixed effect combinations were calculated and adjusted using the false discovery rate (FDR). Count data were further filtered to retain ASV's detected in greater than 10% of all samples. A PERMANOVA was carried out on transformed ASV counts using the ADONIS function of the R package vegan [58].

(e) Statistical analyses

All statistical analyses were performed using R.3.4.3 [59]. Daily maximum, minimum, mean and daily range of temperatures were calculated from the *in situ* data, further divided into seasons: winter (July 2015–October 2015 and April 2016–July 2016), and summer (October 2015–April 2016), and tested using ANOVA, with site and season as fixed effects. *Post hoc* comparisons of significant effects were tested using the *lsmeans* function [60]. We collected time-series data from the National Oceanic and Atmospheric Administration Coral Reef Watch (NOAA CRW) global 5 km product for Ofu Island [48]—SST, sea surface temperature anomalies (SSTA) and degree heating weeks (DHW)—from 2010 to 2012 and from 2015 to 2016. These years were chosen to compare Ofu temperatures between previous 'normal' years—the Palumbi *et al.* [8] study (2010–2012)—and recent mass bleaching years. ANOVA (*lm* function; [61]) and Tukey's *post hoc* comparisons (*lsmeans*) were used to determine whether SST, SSTA and DHW differed between the aforementioned years.

For each coral species, differences in weekly growth, total chlorophyll and normalized F_v/F_m were evaluated with respect to time point (levels: winter and summer), origin (levels: HV, MV, LV), transplantation (levels: HV common garden, native MV and LV) and treatment (levels: heat and control). Sample sizes for each factorial group (origin * transplantation) were five ($n = 5$ genets), with an occasional reduction to 4 or 3 genets owing to sample loss (exact sample sizes for each variable/comparison are in the electronic supplementary material, tables S2 and S3). Effects were tested using a mixed model ANOVA, where time, a combined origin_destination site variable (owing to the unbalanced design (i.e. not all origins in each destination), and treatment were modelled as fixed factors, and colony identity was nested within experimental tank designation as a random factor. Multiple comparisons across factors and interaction terms were assessed *post hoc* using general linear hypothesis testing and multiple comparisons (*glht* function; [62]) for linear mixed-effects models, specifying Tukey's test. To satisfy model assumptions, normality was examined using the *shapiro.test* and homoscedasticity via the *bartlett.test* in R, as well as plotting residuals.

3. Results

(a) Anomalously high Ofu temperatures

In situ backreef temperatures of Ofu Island reveal greater daily maximum and minimum temperatures, and consequently a greater daily range in the HV pool than the MV and LV pool (figure 1*b* and the electronic supplementary material,

figure S1; tables S1 and S4), specifically during the summer. Thermal anomalies were calculated as the total number of days during the experimental duration (July 2015 to July 2016) when temperatures exceeded the NOAA CRW 50 km regional bleaching threshold of 30.2°C [48]. The HV pool had a total of 125 days in which the daily maximum exceeded the bleaching threshold, versus 93 and 81 days over the threshold for the MV and LV pools, respectively. Moreover, the HV pool had 72 and 27 days above 31 and 32°C, versus 38 and 8 for the MV pool, and 33 and 12 days for the LV pool. By contrast to daily fluctuations and high-temperature events, overall mean temperature did not differ among the three pools (electronic supplementary material, figure S1; tables S1 and S4).

Annual temperatures also differed over the course of our study, where 2015 had greater max, min and average *in situ* temperatures in comparison to 2016 (figure 1*c*). In comparison to temperatures of the previous study by Palumbi *et al.* [8] (e.g. a non-anomalous year), this study had a greater number of DHW than 2010, 2011 and 2012 (figure 1*c*, electronic supplementary material, table S4). 2015 had up to 8 DHW over five months (six months prior to the first sampling point), 2016 had less than or equal to 5 DHW that spanned four months, while 2010 had less than or equal to 3 DHW over 2.5 months (figure 1*c*). In addition, SST and SSTA from 2016 were higher than in 2011–2012, as well as 2015.

(b) Coral host growth over time

For both coral species, weekly growth rate was influenced by the two-way interaction between origin_destination transplant site and time. Averaged across both time points, *P. lobata* from the HV pool grew approximately 2.5 times more than MV and LV corals transplanted into the HV pool (figure 2*a*; electronic supplementary material, tables S2 and S5). By July 2016, growth was greatest in HV corals, and MV and LV native corals grew twice that of transplanted paired ramets. Additionally, the growth of native *P. lobata* ramets was higher in July than January (figure 2*a*). For *G. retiformis*, weekly growth in July 2016 was 2–3 times higher in corals native to the MV pool compared to MV transplants and both LV groups (figure 2*d*; electronic supplementary material, tables S2 and S6), but not different than corals native to the HV pool. Similar to *P. lobata*, there were no growth differences in January. Growth of *G. retiformis* native to the MV pool was two times greater in July than January (figure 2*d*; electronic supplementary material, tables S2 and S6).

(c) Symbiodiniaceae photophysiology under acute heat stress

Photophysiological responses of *in hospite* Symbiodiniaceae following heat stress varied by coral host species. For *P. lobata*, acute heat stress reduced F_v/F_m (calculated as loss normalized to starting value; see Material and methods) for HV and LV natives and MV and LV corals transplanted into the HV pool ($p < 0.0001$; figure 2*b*, denoted with '*'). However, MV native *P. lobata* were not affected by acute heat stress (electronic supplementary material, tables S3 and S5), and F_v/F_m values were approximately 1.2–1.8 times higher in MV heated corals than heated HV and LV corals for both time points (figure 2*b*; electronic supplementary material, table S5). For *G. retiformis* there were no differences in F_v/F_m values among native and transplanted groups, nor was there an effect of heat treatment in January. Photochemical efficiency of heat-treated samples varied by a

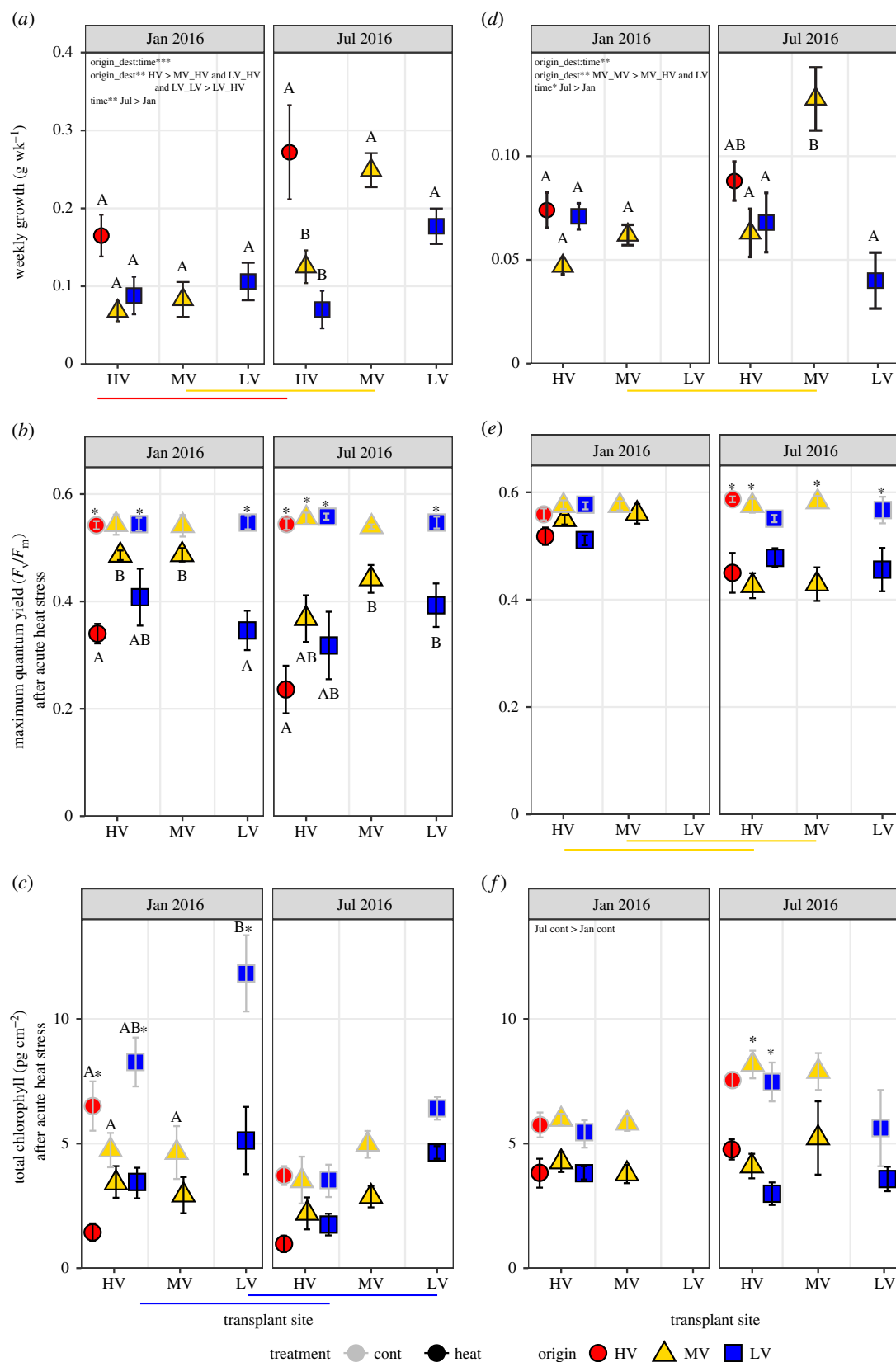


Figure 2. Weekly growth rate (g week⁻¹; top panel), maximum quantum yield (F_v/F_m ; middle panel) and total chlorophyll (pg cm⁻²; bottom panel) of Symbiodiniaceae following acute heat stress (mean \pm s.e.) in *P. lobata* (a–c) and *G. retiformis* (d–f) with respect to transplant destination and time. Significant *post hoc* comparisons within each time panel are represented by letters (differences among transplant groups) and asterisks (effect of acute heat stress). Coloured horizontal lines represent significant differences within paired transplant groups over time. (Online version in colour.)

time and treatment interaction, with higher F_v/F_m values in January than July, but only for MV heated corals ($p < 0.0001$; figure 2e, electronic supplementary material, tables S3 and S6). For both species, there were no significant tank effects.

Total chlorophyll (a + c) differed by either native pool or time. For *P. lobata*, native LV corals had approximately two

times higher control than HV and MV corals during January (time * origin_dest * trt $p = 0.047$; figure 2c; electronic supplementary material, tables S3 and S5). In January, acute heat stress reduced total chlorophyll values in LV and HV corals (electronic supplementary material, tables S3 and S5). Similar to F_v/F_m , there was no effect of treatment on total chlorophyll

content in *P. lobata* from the MV pool (figure 2c; electronic supplementary material, tables S3 and S5). For *G. retiformis*, there was an interactive effect of treatment and time, where total chlorophyll control values were greater in July than January ($p < 0.0001$; figure 2f; electronic supplementary material, tables S3 and S6). Similar to F_v/F_m , there was no effect of treatment in January, but heat stress reduced total chlorophyll values in MV and LV corals transplanted into the HV pool in July ($p \leq 0.0001$; electronic supplementary material, table S3, denoted with *).

(d) Stable symbiodiniaceae composition

Symbiodiniaceae ITS2 rDNA was analysed for distinct ASVs, and resulted in two ASVs for *P. lobata* and six ASVs for *G. retiformis*. Dominant Symbiodiniaceae were all *Cladocopium* spp. (formerly Clade C; [27]) and species varied between *P. lobata* and *G. retiformis*. In *P. lobata*, *Cladocopium* ITS2 type C15 (NCBI accession no. AY239369.1) was dominant at greater than 99%, but a few coral individuals contained background proportions (less than 1%) of *Cladocopium* ITS2 type C40 (AY258485.1; electronic supplementary material, figure S4A; table S7). For *P. lobata*, *Cladocopium* community composition did not change over time.

Unlike *P. lobata*, *G. retiformis* corals contained mostly *Cladocopium* ITS2 type C40 at 50–73%, types C15 and C3 (AF499789.1) at 6–27% and 3–6%, respectively, and types C1 (AF333515.1), C15b (AY258491.1) and C21 (AY239372.1) were detected at background proportions (less than 1%; electronic supplementary material, figure S4B, table S7). *Goniastrea retiformis* community composition varied by native backreef pool and time. ITS2 type C3 varied by origin, where it was present (5–10%) in *G. retiformis* from the HV and MV pool but absent from LV corals (PERMANOVA FDR < 0.05 $p = 0.0062$). ITS2 type C15 was present (2–30%) in HV and LV *G. retiformis* but absent in MV corals (PERMANOVA FDR < 0.05 $p = 0.0014$) until January 2016, when type C15 increased to 40–50% in MV corals and became absent from LV corals.

4. Discussion

We tested whether exposure to HV temperatures increased or decreased stress resistance in two massive coral species from distinct backreef environments. Corals transplanted for one year into the site with the highest variability (the HV pool common garden) did not increase growth or improve photo-physiological responses following acute heat stress, as observed in previous studies [8]. Instead, growth and stress tolerance responded differently to spatial and temporal variation in temperature regimes, and differently in *P. lobata* and *G. retiformis*. Unexpectedly, *P. lobata* native to the HV pool, the site with the highest thermal variability, were most sensitive to experimental bleaching. Previous work in Ofu found increased stress tolerance following acclimation to the greater thermal variability of the HV regime [35], yet we observed a negligible effect of this variability on thermal performance for corals transplanted into and a deleterious effect for corals from the HV pool. Our results suggest that not all coral species may respond positively (or similarly) to HV thermal habitats.

High magnitudes of temperature variation have recently been recognized as a significant promoter of reef-building coral thermal tolerance over small spatial scales (less than 10 km) and could increase resilience to anticipated ocean

warming (e.g. [8,9,22,63]). Coral populations from inshore/protected habitats with high diurnal fluctuations consistently exhibit greater growth and/or natural bleaching tolerance than conspecifics from offshore/exposed habitats, a paradigm congruent across the Caribbean [9,23,64], Red Sea [65], Ofu Island in the South Pacific [22,34], northwest Australia [11] and Great Barrier Reef [24]. By contrast, coral growth in the present study was not different among the three backreef populations in their native environments (except lower growth in LV native *G. retiformis* in July 2016) despite differences in thermal regimes, although MV and LV *P. lobata* transplants in the HV pool had lower growth than paired native ramets and HV native genets in July 2016. Moreover, HV native corals and corals transplanted into the HV pool were susceptible to acute bleaching stress during one or both time points. We also observed no effect of acute heat stress on native and transplanted MV *P. lobata* corals (except for F_v/F_m values in MV transplants during July 2016). This starkly contradicts previous studies examining coral species from or transplanted into the HV pool, which found higher: thermal tolerance limits [8,10,22], the prevalence of heat-tolerant *Durussdinium trenchii* [66] and transcription of heat responsive genes [29] than MV pool corals. Despite persistent high magnitudes of thermal variability, the HV pool did not increase heat tolerance of massive coral species during our study, which complicates the notion that HV thermal habitats are universally beneficial for increasing the adaptive and acclimatory potential of all coral species.

The most obvious distinction between previous experiments and ours is that prior research has predominantly focused on corals in the genus *Acropora* [24,35,38,67]. Biological traits such as colony morphology, growth rate and reproductive mode separate branching corals such as *Acropora* spp. from massive coral species into ‘competitive’ and ‘stress-tolerant’ life-histories, respectively [68]. Large, slow-growing massive corals are thought to be more thermally tolerant to chronically variable and disturbed habitats than branching species in both the Caribbean [69] and Indo-Pacific [70], given life-history traits such as increased tissue thickness and energy surplus [36,37,71]. The HV population of *P. lobata* has previously exhibited higher growth (versus MV corals) and stress resistance (versus forereef corals; [22]), but here, *P. lobata* in the HV pool demonstrate reduced stress tolerance compared to MV and LV populations. These massive coral species are naturally abundant within the HV pool [20], thus, their common occurrence, as well as the increased growth and stress resistance shown previously in HV *P. lobata* makes it unlikely that the taxonomic difference between the present and previous studies is the main explanation for contrasting results of minimal growth differences and reduced thermal tolerance of HV corals seen herein.

Although both *P. lobata* and *G. retiformis* are clustered into the stress-tolerant life-history strategy [68], species-specific responses are apparent under acute bleaching stress. For both photochemical efficiency (F_v/F_m) and total chlorophyll, we found opposing effects of time, where heat stress affected *P. lobata* in January but *G. retiformis* corals were more affected in July 2016. In addition, stronger effects of pool of origin were evident for *P. lobata* bleaching responses and July 2016 growth versus *G. retiformis*. Ofu backreef *Acropora* populations harbour pool-specific Symbiodiniaceae communities, where *Acropora* spp. in the HV pool predominantly host *D. trenchii*, while MV corals host both *D. trenchii* and *Cladocopium* type

C2 [25]. By contrast, we observed similar Symbiodiniaceae communities within *P. lobata* (type C15) across the backreef, site-specific assemblages within *G. retiformis* (type C40, C15 and C3), and distinct species-specific assemblages. While it is unclear whether different Symbiodiniaceae *Cladocopium* assemblages could be driving the observed species-specific seasonal variation in photophysiological responses to bleaching stress [72], both intra- and interspecific host and symbiont variation is known to shape growth and thermal tolerance limits in corals (e.g. [36,73,74]).

Additionally, it could be that corals in these backreef pools are locally adapted to their native thermal conditions. In the Florida Keys, mass gain, protein and lipid levels, and gene expression plasticity of *Porites astreoides* were greater for corals in their native environment in comparison to foreign transplants [9,28]. Similarly in Ofu, backreef (HV and/or MV) *P. lobata* had consistently higher growth, environmental tolerance and cellular responses than corals from or reciprocally transplanted to a nearby forereef [22,30,34]. In Barshis *et al.* [22], HV *P. lobata* grew more than forereef corals, and both HV and MV *P. lobata* exhibited increased tolerance under acute thermal stress compared to forereef corals regardless of acclimation to stable or fluctuating temperatures (though HV and MV did not differ; [22]). Notably, this experiment used a 36 days aquarium-based acclimation versus the 12 months field acclimatization performed herein and observed no differences between HV and MV populations. We also found the highest growth in HV natives versus MV and LV corals transplanted into the HV pool, but only for *P. lobata* during July 2016 and no differences among their native environments. However, differences in stress tolerance between paired native versus transplanted ramets exist for both species: a non-significant then significant reduction in both F_v/F_m for MV native versus transplanted *P. lobata* and total chlorophyll for MV and LV native versus transplanted *G. retiformis* from January to July 2016, suggesting a potentially higher stress level in transplanted ramets. For local adaptation to occur in these backreef populations, individuals would need to perform better at home versus away [75], which is illustrated here for coral growth but not stress tolerance (excepting the instances mentioned above). In addition, HV corals have previously demonstrated increased tolerance owing to the conditions of the HV pool [8,10], yet in this study, F_v/F_m values suggest HV native *P. lobata* were most susceptible to stress. Local adaptation could contribute to the complexity of our results, though it cannot be fully supported, as we did not observe classic patterns of best performance at home versus away, nor did we conduct a full reciprocal transplant moving HV corals into the MV or LV pools.

For HV corals, increased growth but reduced stress tolerance could be evidence of tolerance trade-offs owing to specialization to HV habitats. Skeletal growth records of massive *Porites* colonies along the Great Barrier Reef illustrate progressive accretion rates associated with warming SST followed by precipitous declines following repeated mass bleaching events ([76], but see [77]). We explored the relationship between HV *P. lobata* coral growth and response to acute thermal stress and found a negative, albeit non-significant, correlation between growth and total chlorophyll (Pearson's $R = -0.41$; electronic supplementary material, figure S5) and no correlation between growth and photochemical efficiency. Taken together, our results corroborate recent findings that

coral growth is probably not a good predictor of bleaching responses under extreme temperatures [78].

Compromised bleaching tolerance of HV native corals and a lack of enhanced performance for corals transplanted into the HV pool could also be attributed to the magnitude and duration of maximal summertime temperatures recorded during this study. From 2015 to 2016, a strong El Niño increased SST and triggered the third pan-tropical mass bleaching event [79,80]. This bleaching event was reported to be the most extensive and severe in recent human history; and reefs in American Samoa were predicted to experience intense bleaching conditions [79]. Our experiments were a few months prior to or post maximal bleaching stress on Ofu Island (2015: February–June, 2016: March–June; figure 1c), however in January 2016, we observed sparse paling in some HV pool branching corals but not in our donor or transplanted corals (C. N. Klepac 2016, personal observation). Thus, the patterns observed herein could represent the initial stages of response to or accumulated after-effects of the thermal anomaly. The HV pool regularly experiences brief but frequent temperatures that reach over 35°C, which greatly exceed the regional bleaching threshold of 30.2°C [10,20], and our acute thermal stress assays serve as an experimental analogue to the strong thermal variation in this pool. Much of the thermal tolerance research previously conducted in Ofu used similar thermal stress assay profiles [8], yet these experiments occurred during milder years, where 2 DHW was rarely exceeded in comparison to 5–8 DHW during our study.

It is thus tempting to speculate whether the extreme temperatures in the HV pool during this study could have overwhelmed the physiological performance underlying temperature tolerance of this population of corals. However, this study would need to be repeated during non-bleaching years and during peak summer temperatures to effectively disentangle the effect of recent thermal history versus taxonomic, evolutionary and population-specific drivers of massive coral species upper thermal limits. Indeed, the differences in thermal tolerance limits observed herein are complex, challenging our understanding of how naturally tolerant populations will fare under rapid climate change. Regardless of the complexity, it is clear that higher magnitudes of temperature variation is not a universal promoter of thermal tolerance limits and that species-specific mechanisms and regional thermal anomalies may be equally important in shaping coral responses to extreme temperatures.

Data accessibility. Data will be made accessible through the author's GitHub repository (https://github.com/courtneyklepac/project_reducedtolerance_massivecorals_variablehabitats) [49]. Unique Symbiodiniaceae sequences are available on NCBI's BioProject no. PRJNA647413.

Authors' contributions. C.N.K. and D.J.B. designed the study, C.N.K. performed field and laboratory research and conducted analyses, C.N.K. and D.J.B. wrote the manuscript.

Competing interests. We declare we have no competing interests

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