

### Atoll-dependent variation in depth zonation of benthic communities on remote reefs

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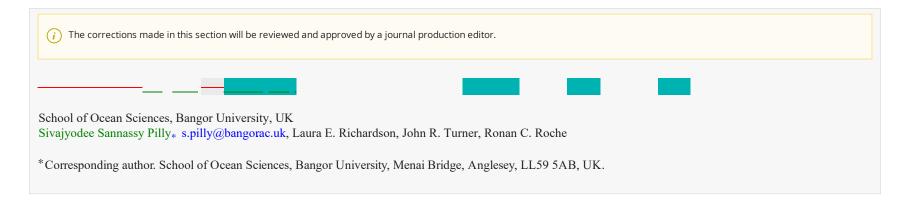
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# Atoll-dependent variation in depth zonation of benthic communities on remote reefs



## **Abstract**

The distribution and organisation of benthic organisms on tropical reefs are typically heterogenous yet display distinct zonation patterns across depth gradients. However, there are few datasets which inform our understanding of how depth zonation in benthic community composition varies spatially among and within different reef systems. Here, we assess the depth zonation in benthic forereef slope communities in the Central Indian Ocean, prior to the back-to-back bleaching events in 2014–2017. We compare benthic communities between shallow (5–10 m) and deep (20–25 m) sites, at two spatial scales: among and within 4 atolls. Our analyses showed the variation in both major functional groups and hard coral assemblages between depth varied among atolls, and within-atoll comparisons revealed distinct differences between shallow and deep forereef slope communities. Indicator taxa analyses characterising the hard coral community between depths revealed a higher number of coral genera characteristic of the deep forereef slopes (10) than of the shallow forereef slopes (6). Only two coral genera consistently associated with both depths across all atolls, and these were *Acropora* and *Porites*. Our results reveal spatial variation in depth zonation of benthic communities, potentially driven by biophysical processes varying across depths and atolls, and provide a baseline to understand and measure the impacts of future global climate change on benthic communities across depths.

### Keywords:

Depth zonation, Coral reefs, Remote systems, Benthic ecology, Community composition, Hard coral assemblages

### Abbreviations

No keyword abbreviations are available

## 1 Introduction

One of the main goals in ecology is to understand how communities occupy space. Biotic (González et al., 2017; Des Roches et al., 2018), abiotic (McGill et al., 2006; Agrawal et al., 2007), and stochastic processes (Hubbell 2005) that are responsible for the organisation of ecological communities, interact and create natural environmental gradients in biophysical resources (Leibold and McPeek 2006; Vellend 2010; Brandl et al., 2019). These naturally occurring gradients combine to limit the distribution, abundance, and diversity of communities (Holt 2003; Peischl et al., 2015). Similar natural variations occur across water depth gradients on tropical reef systems, where biophysical conditions known to influence the physiology of reef organisms co-vary, such as light availability (Brakel 1979; Hoegh-Guldberg and Jones 1999; Cooper et al., 2007), temperature, salinity (Kleypas et al., 1999), and wave and current regimes (Lowe and Falter 2015; Radice et al., 2019). Coral reef species develop different traits to survive within variations of these parameters across depths (Kneitel and Chase 2004; McGill et al., 2006; Darling et al., 2012). As a result, distinct and predictable ecological zonation patterns in coral reef ecosystems can be observed across depths (Sheppard 1982; Done 1983; Roberts et al., 2015; Karisa et al., 2020).

Different zones on shallow coral reef systems are usually defined by the occurrence of one or more dominant organisms that occupy a certain depth in a location (Goreau 1959; Sheppard 1982; Done 1983). For instance, zonation patterns in shallow forereef slope communities are described as being predictably dominated by structurally robust species. Coral species exposed to high light regimes and large temperature fluctuations on shallow areas of the reef slope have adapted morpho-physiologically to this dynamic environment (Titlyanov and Titlyanova 2002; Iglesias-Prieto et al., 2004; Marcelino et al., 2013; Guest et al., 2016). Often prone to high exposure to wave and surge energy (Done 1983), some shallow water species are adapted to thrive in areas where frequent sediment resuspension occurs, forming robust morphologies which can resist high water flow and physical forces (Todd 2008; Duckworth et al., 2017). In contrast, deeper forereefs which are more sheltered from surface wave exposure and receive lower irradiance due to light attenuation with depth (Done 1983; Rex et al., 1995), are characterised by coral communities that frequently adopt encrusting and foliose growth forms to increase light capture efficiency (Titlyanov and Titlyanova 2002; DiPerna et al., 2018a[Instruction: Please delete DiPerna et al 2018a and DiPerna et al.,2018b - the reference has been cited twice. The correct citation should be: Diperna et al., 2018. The system would not allow me to delete one of them to make the appropriate corrections.]). There is also an increase in mixotrophic and heterotrophic traits in marine organisms, such as hard corals, soft corals and sponges to offset limited light availability in deeper reef zones (Fabricius and Klumpp 1995; Fabricius and De'ath 2008; Houlbrèque and Ferrier-Pagès 2009).

Our understanding of the consistency of how coral reef benthic communities vary spatially across depths remains unclear (Edmunds and Leichter 2016; Roberts et al., 2019). This may be due to the complex interactive effects of biophysical processes that drive ecological community structure, which vary across multiple spatial and temporal scales (Hatcher et al., 1987; Magurran 2004; Leibold and McPeek 2006; Vellend 2010; Brandl et al., 2019), resulting in highly heterogenous coral reef communities (Edmunds and Bruno 1996; Huntington and Lirman 2012; Obura 2012; Dalton and Roff 2013; McClanahan et al., 2014; Ford et al., 2020). In addition, much of our current understanding of depth zonation patterns on reefs is based on observations made decades ago (Goreau 1959; Done 1982, 1983; Sheppard 1982), prior to subsequent climate change impacts that have altered coral reef communities (Dubinsky and Stambler 2010; Williams et al., 2019).

Coral reefs around the world are increasingly vulnerable to more intense and frequent climate-driven disturbances (Anthony 2016; Hughes et al. 2017, 2018; Perry and Alvarez-Filip 2019). However, the extent at which coral reefs are affected by anthropogenic stressors is not uniform across space and varies across depths (Bongaerts et al., 2010; Bridge et al., 2013; Baird et al., 2018). Examining depth zonation of benthic communities in the context of systemic disturbance has provided important insights into complex dynamics such as diverging ecological trajectories and regime shifts, but these studies have been generally limited to shallow (3–10 m) depth ranges (Graham et al., 2015; Gouezo et al., 2019). This focus on shallow depth ranges has limited our understanding of how benthic community composition change across different depths within contemporary shallow tropical reefs (<30 m) (Bridge et al., 2014; Edmunds and Leichter 2016).

Here we examined depth zonation in benthic community composition at two spatial scales (among and within atolls) in the Chagos Archipelago, a relatively isolated reef system in the Central Indian Ocean, prior to pan-tropical bleaching events in 2014–2017 (Eakin et al., 2019) and post 1998 bleaching (Sheppard 1999a). Isolated reefs can

be used an ecological reference point (Sandin et al., 2008; Smith et al., 2008; Williams et al., 2013; Heenan et al., 2017; Head et al., 2019) and examining benthic communities, prior to the back-to-back bleaching event in 2014–2017, establishes a baseline to measure the extent of further climate change impacts and how these vary across depths. Specifically, we compare benthic composition among and within four atolls, between two depth ranges: 5–10 m and 20–25 m on the forereef slopes. With our current knowledge of depth dependent zonation on coral reefs (Done 1982, 1983; Sheppard 1982), we expected to find zonation across depth in benthic community composition that was consistent amongst atolls.

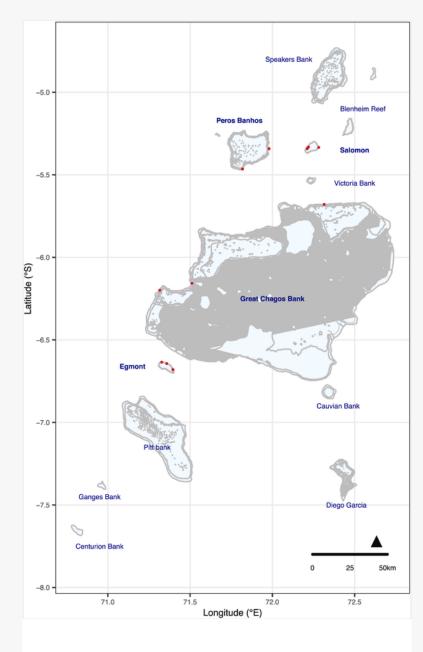
# 2 Methods

### 2.1 Study sites

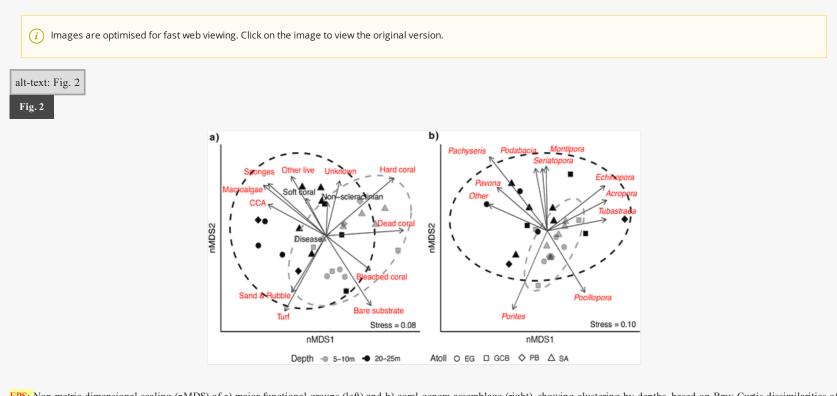
The Chagos Archipelago is located in the centre of the Indian Ocean, at the remote southern end of the Laccadives-Maldives-Chagos ridge, ~500 km from south of Maldives (Sheppard 1999b). The archipelago is comprised of 5 atolls, 52 islands, and constitutes 9400 km² of submerged shallow reefs (<40 m depth) (Dumbraveanu and Sheppard 1999; Sheppard et al., 2013a). The archipelago has been largely uninhabited since the early 1970s ((Sheppard, 1999b)Sheppard, 1999b), with the exception of Diego Garcia (DG), the southern-most atoll that hosts a US naval facility, where strict environmental regulations are enforced, prohibiting all commercial fishing and extractive activities at sea (Purkis et al., 2008). In this study, a total of 26 sites; including 13 at 5–10 m depth (hereafter 'shallow' reefs) and 13 at 20–25 m depth (hereafter 'deep' reefs), were surveyed on forereef slopes across 4 atolls: Peros Banhos (PB), Salomon (SA) (northern atolls), Great Chagos Bank (GCB) and Egmont (EG) (southern atolls) (Fig. 1, Table S1) (see Fig. 3) (see Fig. 2).

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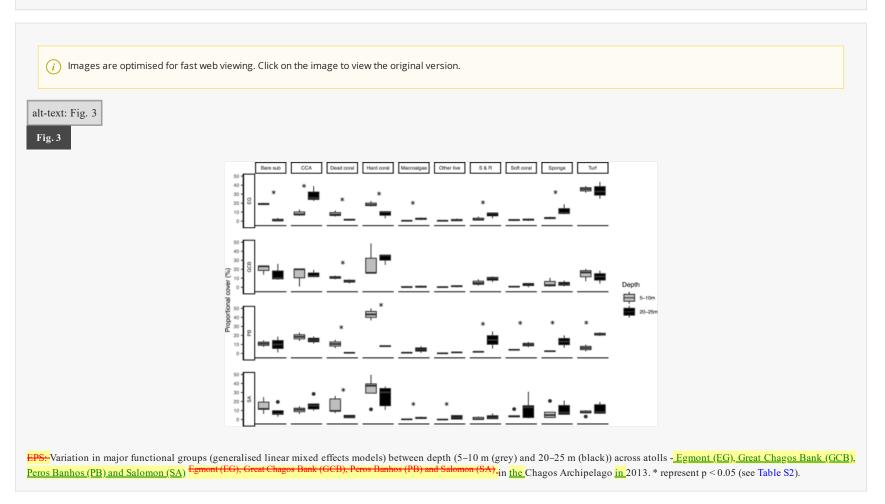
alt-text: Fig. 1
Fig. 1



EPS: Map of Chagos Archipelago showing sampled sites (red points) around surveyed atolls (in bold) – northern atolls: Peros Banhos (PB), Salomon (SA) and southern atolls Great Chagos Bank (GCB) and Egmont (EG) - see Table S1 for list of sites and coordinates. (For interpretation of the references to colour in this figure legend, the reader is referred to the Web version of this article.)



EPS:-Non-metric dimensional scaling (nMDS) of a) major functional groups (left) and b) coral genera assemblage (right), showing clustering by depths, based on Bray-Curtis dissimilarities of square-root transformed data in a total of 26 sites in the Chagos Archipelago. Coloured ellipses represent dispersion of depth centroids at 95% confidence limit – greyblue: shallow reefsites (5–10 m) and blackred: deep reefsite (20–25 m). Symbols represent surveyed atolls – Egmont (EG), Great Chagos Bank (GCB), Peros Banhos (PB) and Salomon (SA). Bottom: Vectors represent a) major functional groups (left) and b) coral genera (right) distribution to the patterns on the ordination plot. Red labels in in (a) indicate a significant contribution and black labels indicate a non-significant contribution. (For interpretation of the references to colour in this figure legend, the reader is referred to the Web version of this article.)



## 2.2 Benthic composition

At each site, benthic composition was quantified from 30 digital photo-quadrats taken in February and March 2013 (total across all sites, n = 780). Digital photo-quadrats were randomly extracted as still images from 10 min continuous video-swims at both 5–10 m and 20–25 m depths. The housing was equipped with two spotlights and two red laser pointers set at 10 cm apart to provide a consistent scale-measurement of the benthos and to adjust for lower-light levels at greater depth to facilitate benthic image analysis. The camera was maintained approximately 0.5 m above the substrate and at a  $45^{\circ}$  angle to capture benthic organisms under overhangs and canopies (Goatley and Bellwood 2011).

Each video was converted into an image sequence (25 frames per sec; in Pinnacle Studio, v22.2.0). To ensure that images selected for analysis did not contain the same section of the forereef, frames were randomly selected, but separated by a minimum of 80–100 frames (Matlab, R2018a.Ink). Benthic image analysis was carried out by SSP, RR, and LR using Coral Point Count with excel extensions (CPCe) (Kohler and Gill 2006). To account for any variation in observer bias in identification of benthic composition, the frames were equally divided among observers and analysed (10 images each per person per site). Proportional cover of benthic categories was quantified by identifying substrate and benthic organisms under fifteen randomly allocated points on each image, with one point assigned within a 3 x 5 grid cell stratification (Suchley 2014).

Substrate type and benthic organisms were categorised as: hard coral (identified to genus), soft coral (identified to family), non-scleractinian coral (*Millepora*, *Heliopora* and *Distichopora*), sponge, crustose coralline algae (CCA), macroalgae, turf algae, bare substrate with algal film (hereafter bare substrate), sand, rubble (<10 cm maximum length), dead coral, bleached coral, diseased coral, 'other live' and unknown. 'Other live' included all sessile invertebrates such as bryozoans, tunicates, bivalves, giant clams, corallimorphs, anemones and zoanthids. Soft corals were identified and grouped within the four most common families occurring in the archipelago – Alcyoniidae, Xeniidae, Nepthtiidae and Nidaliidae (Schleyer and Benayahu 2010), or other soft coral. Where image quality limited identification of hard corals to genus level or morphology (approximately 0.1–0.5%), corals were classified as 'other'. The classification of benthic categories for this study was based on NOAA Coral Reef Information System Instruction: Please amend this reference so it is cited as "Coral reef Ecosystem Division, 2014".] (Coral Reef Ecosystem Division,)(Coral Reef Ecosystem Division)

[42014], Denis et al. (2017) and the CATAMI classification system (Althaus et al., 2015).

## 2.3 Data analysis

To assess whether benthic composition varied across depths, and if depth zonation varied among atolls, benthic community composition was first visualised at two levels: 1) proportional cover of major functional groups (hard coral, soft coral, sponge, CCA, macroalgae, turf algae, bare substrate with algal film, sand and rubble, non-scleractinian coral, bleached coral, dead coral, diseased coral, 'other live'), and 2) proportional cover of hard coral assemblages, identified to genera. Benthic composition was visualised across: a) depth, and b) atoll, using non-metric Multidimensional Scaling (nMDS: vegan package; Oksanen et al., 2012), based on a Bray-Curtis dissimilarity matrix of square root transformed data. A scree plot was used to evaluate ordination stress and a Shepard stress plot to confirm correlation between the original dissimilarity matrix

and the distances on the final nMDS plot. The nMDS was computed on 3 dimensions (k = 3) with a stress value  $\leq 0.1$ . The envfit function (vegan package) was used to fit vectors of major functional groups and coral genera, to their respective nMDS ordinations.

To assess differences in benthic composition across depth and atolls, we performed two-way nested permutational multivariate analyses of variance (PERMANOVA; Anderson, 2017) on: 1) major functional groups, and 2) coral genera, as a function of the interaction between atolls (4 atolls; fixed factors) and depths (2 depths; fixed factors), with sites (random factors) nested in atoll (9999 permutations; adonis2 function: vegan package). Average within-group dispersion was examined using a multivariate homogeneity test (betadisper: vegan package). Where a significant interaction between depth and atoll was found, we tested for variation in 1) major functional groups, and 2) coral genera across depths at each individual atoll using one-way nested PERMANOVAs. Data were square-root transformed and analysed using Type III sum of squares to accommodate an unbalanced design, in both one-way and two-way PERMANOVA.

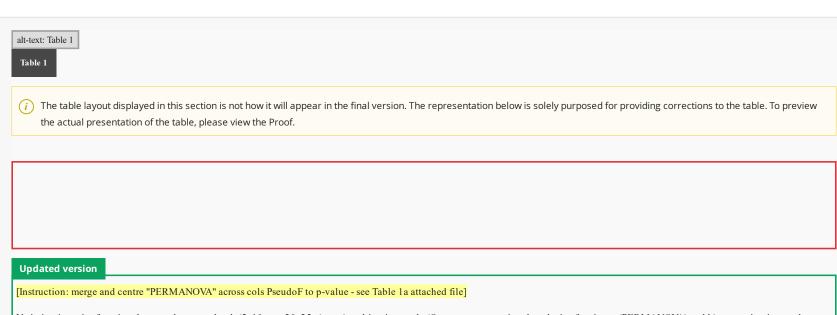
Indicator taxa analyses were used to determine the association between coral genera and the depth ranges at which they occur. Coral genera that significantly associated with shallow and deep reefs across each atoll, were used to infer on the biotic and abiotic state of the environment that prevail at the different depth ranges (Cáceres and Legendre 2009). Prior to analysis, the proportional cover matrix of the coral genera assemblage was converted into presence/absence data. Coral genera characteristic of shallow and deep reefs were identified using a p-value threshold of <0.05 (9999 permutations, indicators: indicspecies package; Cáceres 2020).

Generalised linear mixed-effects models (GLMMs) with a binomial distribution and logit-link function were used to model major functional groups within each atoll, treating depth as a fixed effect and sites as a random effect (glmer: lme4 package, Bates et al., 2015). GLMMs were also performed on coral genera that consistently characterised the hard coral assemblage across both depths and all atolls. When major functional groups or coral genera cover had a high number of zeros and did not fit the standard binomial distribution (>55% of the data consisted of zero values), a zero-inflated generalised linear mixed-effect models with a beta distribution was fitted (depth as a fixed factor and sites as a random factor; glmmtmb: glmmTMB package; Brooks et al., 2017). All analyses were performed using R 3.5.1 (R Development Core Team 3.5.1, 2018).

### 3 Results

#### 3.1 Variation in major functional groups

Within-atoll analyses revealed significant variation in major functional groups between shallow and deep reefs across all four atolls (PERMANOVA, all: p = 0.001, Table 1). There was also an interactive effect of depth and atoll (PERMANOVA, Pseudo  $F_{3,779} = 13.54$ , p = 0.001) and dispersion (Table S4) on benthic composition, indicating the variation in major functional groups between depth varied among atolls; notably at EG and PB (Figure 2, Table 1, Table S5a).



Variation in major functional groups between depth (5–10m vs 20–25m) on a) multivariate scale (One-way permutational analysis of variance (PERMANOVA) and b) on a univariate scale (generalised linear mixed effect models (GLMM)) within atolls: Egmont (EG), Great Chagos Bank (GCB), Peros Banhos (PB) and Salomon (SA) in the Chagos Archipelago in 2013. BS – Bare substrate with algal film, CCA – crustose coralline algae, HC – hard coral, MA – macroalgae, DC- dead coral, OL – other live, SR – sand and rubble, SP – sponge, T – turf, BL – bleached coral, UK – unknown, NS – non-scleractinian coral. In bold are significant p values. df-degree of freedom. (a) b):

Atoll	PERMANOVA								
Aton	PseudoF-ratio	df	p-value						
EG	46.844	1,179	<0.001						
GCB	5.156	1,179	<0.001						
PB	48.943	1,119	<0.001						
SA	28.208	1,229	<0.001						

Atoll	[Instruction: merge and centre GLMM across table i.e from BS to NS - see Table 1b attached file columns with "-" (where analysis could not be done), should be merged and centered as per Table 1b in attached file]GLMM													
		BS	CCA	НС	MA	DC	OL	SR	SC	SP	T	BL	UK	NS
	Coeff	0.628	-1.464	1.052	-2.189	1.742	-1.269	-1.158	-0.393	-1.294	0.078	_	0.032	-
EG	SE	0.210	0.287	0.322	0.607	0.397	1.009	0.521	0.865	0.340	0.215		0.461	
	p-value	0.003*	<0.001	0.001	<0.001	<0.001	0.209	0.026	0.650	<0.001	0.717		0.946	
	Coeff	0.466	-0.573	-0.356	-0.456	0.5418	-1.122	-0.733	-1.021	-0.105	0.321	0.321	0.740	-
GCB	SE	0.384	0.867	0.461	1.204	0.189	0.978	0.418	0.991	0.819	0.499	0.499	1.153	
	p-value	0.225	0.509	0.440	0.705	0.004	0.251	0.079	0.303	0.898	0.520	0.193	0.521	
	Coeff	0.732	0.260	2.165	-1.399	2.864	-0.359	-2.096	-0.944	-1.698	-1.622		0.088	
PB	SE	1.092	0.320	0.244	1.035	0.657	0.624	0.710	0.293	0.543	0.519	_	0.167	_
	p-value	0.503	0.416	<0.001	0.176	<0.001	0.566*	0.003	0.001	0.002	0.002		0.589*	
	Coeff	0.642	-0.490	0.398	-2.119	1.628	-2.966	-1.292	-0.698	-0.624	-0.395		0.150	-0.791
SA	SE	0.392	0.252	0.418	0.516	0.357	0.948	0.840	0.650	0.509	0.303	_	0.335	1.983
	p-value	0.101	0.052	0.341	<0.001	<0.001	0.002	0.124	0.283	0.221	0.192		0.654	0.690

- could not compare as groups and not occur at both deputs, Zero inflated model using GENINTINES.

A decreasing hard coral, bare substrate and dead coral cover was observed with increasing depth compared to CCA, macroalgae, sand and rubble and sponge cover which increased on the deep forereef slopes at EG (Figure 3, Table 1b). At PB, hard coral and dead coral cover also decreased with depth compared to sand and rubble, sponge, soft coral and turf algae cover which increased on the deep forereef slopes (Figure 3, Table 1b). At GCB, only dead coral cover varied significantly with depth (Figure 3, Table 1b).

Table 1b). At SA, macroalgae and 'other live' cover increased significantly with depth while dead coral showed a significant decrease with increasing depth (<u>Figure 3</u>, Table 1b).

### 3.2 Variation in hard coral assemblage

Thirty-seven coral genera were recorded across all sites, with a higher number of coral genera were recorded at deep reefs (34) than shallow reefs (23). SA contained the highest coral genera richness (29), followed by GCB (22 genera), PB (20 genera) and EG (16 genera). Within-atoll variation revealed significant differences in hard coral assemblages between depths (PERMANOVA, all: p = 0.001, Table 2a). However, significant depth-by-atoll interactions (PERMANOVA, Pseudo  $F_{3,650} = 7.40$ , p = 0.001; Figure S1) and dispersion (Table S4), indicate the variation in hard coral assemblages between depth varied among atolls across the archipelago, particularly at SA and PB (Figure 2., Table S5b).

# alt-text: Table 2 Table 2

(i) The table layout displayed in this section is not how it will appear in the final version. The representation below is solely purposed for providing corrections to the table. To preview the actual presentation of the table, please view the Proof.

### **Updated version**

[Instruction: merge and centre PERMANOVA as per Table 2b in attached file]

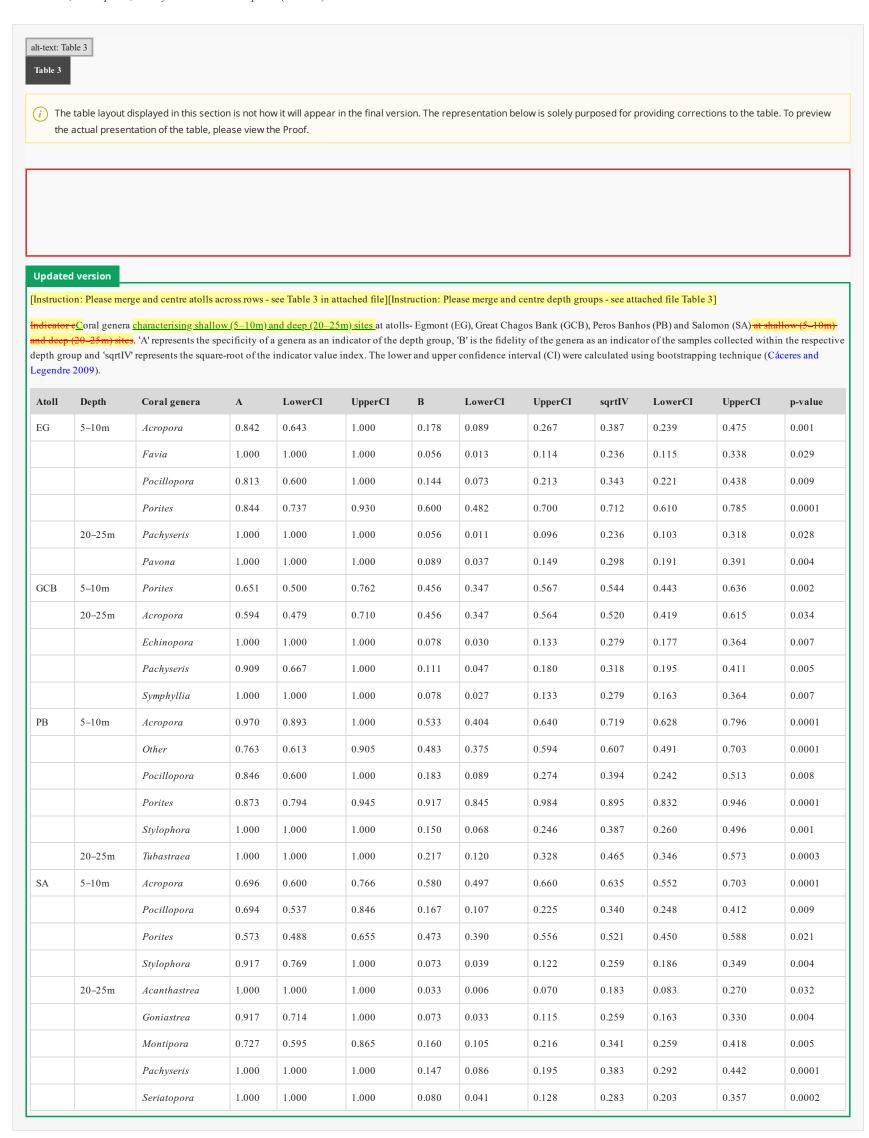
Variation in the hard coral assemblage between depth on a) multivariate scale (One-way permutational analysis of variance (PERMANOVA) and b) variation in common characteristic indicator coral genera on a univariate level (generalised linear mixed effect models (GLMM)) within atolls: Egmont (EG), Great Chagos Bank (GCB), Peros Banhos (PB) and Salomon (SA) in the Chagos Archipelago in 2013. In bold are significant p values. \* indicate where zero-inflated generalised mixed effect models were performed. df – degree of freedom, Coeff- coefficient estimates, SE-Standard error. a) b).

Atoll	PERMANOVA								
Aton	PseudoF-ratio	df	p-value						
EG	17.016	1,116	<0.001						
GCB	6.494	1,160	<0.001						
РВ	25.461	1,95	<0.001						
SA	15.150	1,276	<0.001						

<u>Coral genera Variable</u>	Atoll	Effect	Coeff	SE	z-statistics	p-value
Acropora		Intercept	-5.636	0.758	-7.437	<0.001
		Depth	2.145	0.972	2.208	0.027
Porites		Intercept	-4.725	0.397	-11.900	<0.001
		Depth	2.113	0.493	4.290	<0.001
Pocillopora*	EG	Intercept	-1.647	0.239	-6.886	<0.001
		Depth	-0.285	0.269	-1.058	0.290
Other		Intercept	-3.518	0.267	-13.183	<0.001
		Depth	0.197	0.369	0.533	0.594
Acropora		Intercept	-2.137	1.101	-1.941	0.052
		Depth	-1.607	1.608	-0.999	0.318
Porites		Intercept	-4.339	1.090	-3.981	<0.001
	GCB	Depth	1.062	1.490	0.713	0.476
Pocillopora	ССВ	Intercept	-4.076	0.340	-12.006	<0.001
		Depth	-0.192	0.475	-0.405	0.686
Other		Intercept	-3.205	0.261	-12.297	<0.001
		Depth	0.090	0.365	0.248	0.804
Acropora*		Intercept	-2.245	0.776	-2.891	0.004
		Depth	0.491	0.783	0.627	0.531
Porites	PB	Intercept	-4.719	1.041	-4.532	<0.001
		Depth	3.632	1.341	2.708	0.007
Pocillopora*	гь	Intercept	-1.344	0.259	-5.183	<0.001
	-	Depth	-0.707	0.294	-2.407	0.016
Other		Intercept	-4.563	0.430	-10.614	<0.001
		Depth	1.508	0.531	2.841	0.004
Acropora	SA	Intercept	-3.838	0.625	-6.146	<0.001
		Depth	1.771	0.865	2.047	0.041
Porites		Intercept	-2.980	0.313	-9.529	<0.001
		Depth	0.430	0.435	0.989	0.323
Pocillopora		Intercept	-5.288	0.338	-15.627	<0.001
		Depth	0.987	0.409	2.412	0.016

Other	Intercept	-3.056	0.151	-20.283	<0.001
	Depth	0.245	0.208	1.175	0.240

Indicator taxa analyses identified a total of 14 coral genera as well as the 'other' category that characterised the hard coral community across all atolls and depth combinations (Table 3). A higher number of coral genera (10) characterised the deeper reefs than the shallow reefs (6). Coral genera significantly associated with shallow reefs in EG were *Acropora*, *Porites* and *Pocillopora*. Coral genera significantly associated with deep reefs in EG were *Favia*, *Pachyseris* and *Pavona*. *Porites* was the only genus that significantly characterised the shallow reefs in GCB. The hard coral assemblage on deep reefs in GCB was characterised by *Acropora*, *Echinopora*, *Pachyseris* and *Symphyllia*. The shallow reefs of both northern atolls, PB and SA were characterised by *Acropora*, *Pocillopora*, *Porites* and *Stylophora*. The only coral genus that significantly associated with the deep reefs in PB was *Tubastrea*. In SA, the hard coral community on deep reefs was characterised by *Acanthastrea*, *Goniastrea*, *Montipora*, *Pachyseris* and *Seriatopora* (Table 3).



Acropora and Porites were the only genera that were consistently associated with both depths across all atolls. Univariate analyses showed that the cover of Acropora decreased with increasing depth at EG and SA (Table 2b). Increasing depth also had a negative effect on the cover of Porites at EG and PB (Table 2b).

## 4 Discussion

Our results showed the variation in the benthic community structure of the Chagos Archipelago was driven by significant depth-by-atoll interactions. Although the zonation patterns of the benthic community varied across atolls, a distinct contrast was observed between shallow (5–10 m) and deep (20–25 m) communities, both in terms of the major functional groups and hard coral assemblages, within each atoll. Several earlier coral reef studies report predictable vertical zonation in benthic communities and hard coral assemblages (e.g. Goreau 1959; Done 1982, 1983; Sheppard 1982). However, the among-atoll variation in depth effect observed in this study is consistent with recent research documenting spatial heterogeneity in tropical reefs (Edmunds and Bruno 1996; Williams et al., 2013; Edmunds and Leichter 2016; Karisa et al., 2020; Ford et al., 2020). Spatial variation in depth zonation may be indicative of local biophysical gradients at each atoll that are interacting with depth to structure benthic assemblages (Whittaker 1972; Reice 1994; Vellend 2010). Physical variables that co-vary with depth, such as light (Edmunds et al., 2018), wave exposure (Williams et al., 2013; Gove et

al., 2015), reef slope (Sheppard 1982), seasonal thermocline (Kahng and Kelley 2007), resource availability (Fox et al., 2018; Williams et al., 2018), internal wave activities (Leichter and Salvatore 2006; Radice et al., 2019), and disturbance history, e.g. storms (Hughes and Connell 1999) and climate-induced bleaching events (Bridge et al. 2013, 2014; Adjeroud et al., 2018), have been shown to influence reef communities.

The distinct variation in benthic composition across depth within each atoll, is comparable to changes with depth in several other Indo-Pacific locations: Central Pacific Kingman and Palmyra atoll (Williams et al., 2013), French Polynesia (Edmunds and Leichter 2016), New Caledonia (Adjeroud et al., 2019), the Maldives (Ciarapica and Passeri 1993), and Kenya (Karisa et al., 2020). Coral genera that consistently associated with shallow reefs were *Acropora*, *Pocillopora* and *Stylophora*. These coral genera frequently exhibit arborescent and bushy morphologies in shallow areas (Sheppard 1982) and have better light-scattering properties (Titlyanov and Titlyanova 2002; Marcelino et al., 2013). They are therefore better adapted to the high light regimes that can be experienced in shallow reef environments (Hoegh-Guldberg and Jones 1999; Winters et al., 2003). These species also have effective mechanisms, such as polyp inflation, tentacular action and mucus production to shed sediment particles that are resuspended due to wave action (Erftemeijer et al., 2012; Duckworth et al., 2017) at highly exposed shallow reefs (Fulton et al., 2001).

Pachyseris, Pavona, Echinopora, Acanthastrea, Goniastrea, Montipora and Symphyllia were characteristic of the hard coral assemblage on deep reefs. These coral genera frequently adopt foliose, encrusting and massive growth forms on deeper forereef slopes (Sheppard 1982). As irradiance levels decrease with increasing depths, morphological adaptations including the development of larger surface areas, such as foliose and encrusting growth forms, help increase efficiency in light capture to optimise photosynthetic activities (Done 1983; DiPerna et al., 2018b)[Instruction: Please delete DiPerna et al 2018b and DiPerna et al., 2018a - the reference has been cited twice. The correct citation should be: Diperna et al., 2018. The system would not allow me to delete one of them to make the appropraite corrections.]. Foliose and massive species also contain higher densities of photosynthetic dinoflagellates (Symbiodinium) for maximising food production in low light regimes (Li et al., 2008). Many of the deeper dwelling massive and sub-massive species exhibit heterotrophic traits; with larger polyp sizes and longer tentacles to allow energy requirements to be met by zooplankton feeding (Hoogenboom et al., 2015; Tremblay et al., 2015). The occurrence of the azooxanthellate Tubastrea as a characteristic coral genera across the deep overhangs of the northern atoll, PB (Andradi-Brown et al., 2019) provides evidence of the presence of highly productive waters in the area. We hypothesise that the deeper forereef areas of the archipelago are more likely recipient of cooler, nutrient-rich upwelling waters than shallow reef areas (Sheppard 2009; Sheppard et al., 2017).

Acropora and Porites characterised both shallow and deep forereef slopes across all atolls. Previous studies in the Chagos Archipelago have also highlighted the dominance of these two coral genera at both shallow and deep forereef slopes (Sheppard et al., 2008). These typically depth generalist genera contain large numbers of species (Vellend, 2010 Veron et al., 2019) that have different physiological and phenotypical traits which give them the ability to persist in diverse environments across depth gradients (Toda et al., 2007; Darling et al., 2012). Both genera have high larval dispersion rates facilitating settlement across depths (Holstein et al., 2016; Serrano et al., 2016). Acropora and Porites have been found to host diverse endosymbionts to optimise photosynthesis in decreased light (Muir et al., 2015; Ziegler et al., 2015; Tan et al., 2020). Acropora are often characterised as a fast growing weedy species which have overtopping abilities and can encroach neighbouring species (Riegl and Purkis 2009). Porites with massive morphologies can grow taller and larger, and physically outcompete other surrounding species (Potts et al., 1985).

Increasing depth had a positive effect on turf algae at PB and CCA cover at EG. Several previous studies have found that turf algae and CCA were more abundant in shallow reef environments (Williams et al., 2013; Marlow et al., 2019; Karisa et al., 2020). The high cover of turf algae and CCA we observed at deep forereef slopes of the Chagos Archipelago may be related to the grazing pressure across depth (Heenan and Williams 2013). Intensive herbivory promotes the removal of epiphytic turf algae (Rasher et al., 2012; Osuka et al., 2018; Roff et al., 2019) and provides space for the growth of the rapid colonising CCA (Airoldi 2000; Russell 2007). Within EG, high grazing intensity coupled with elevated herbivore and excavator biomass between 8 and 17 m, were associated with CCA dominated habitats on forereef slopes (Samoilys et al., 2018; Sheppard et al., 2013b). A lower biomass of herbivorous fishes has also been reported on the deeper forereefs of the Chagos Archipelago (Andradi-Brown et al., 2019), which could potentially explain the observed higher turf cover on the deep reefs of PB.

Among all measured major functional groups, dead coral was the only group which had consistently higher cover in shallow forereef slopes across all atolls. During the 1998 massive bleaching events, all shallow ocean-facing reefs in the Chagos Archipelago were highly impacted, leaving large quantities of dead corals (Sheppard, 1999). Several subsequent bleaching events in shallow waters in the 2000s also resulted in localised mortality (Harris and Sheppard 2008; Sheppard et al., 2008), which may have inhibited recruitment and growth, explaining the higher dead coral cover at 5–10m across the archipelago.

Soft coral and sponge cover increased with depth, notably at EG and PB. Previous findings in Indo-Pacific reef systems (Reichelt et al., 1986; Barnes and Bell 2002), including the Chagos Archipelago (Sheppard 1981; Schleyer and Benayahu 2010) show similar increase in soft coral and sponge cover with depth. Sponges and soft corals species have phototrophic, heterotrophic and mixotrophic abilities (Wilkinson 1983; Fabricius and Klumpp 1995). Their high cover in deeper forereef can result from photoadaptation of autotrophic species to increased depths (Fabricius and De'ath 2008; Shoham and Benayahu 2017). Mixotrophic and heterotrophic sponges and soft corals can feed from large flows of inorganic nutrients that are delivered by cold water upwelling and/or internal waves that occur on deeper reef sites (Lesser 2006; Pupier et al., 2019). Deep steep walls are also less prone to sedimentation creating favourable habitat for octocorals (Bridge et al., 2011). Accentuated by the steep walls of deep sites of PB (Sheppard 1980; Winterbottom et al., 1989), deep rich waters may upregulate heterotrophic feeding in soft corals and sponges, explaining the observed higher cover in its deep forereef slopes (Schleyer and Benayahu 2010; Sheppard et al., 2013b).

It is important to understand the reef community composition of the Chagos Archipelago in the context of disturbances that have affected its reefs. Classic ecological theories suggest that following a disturbance, communities undergo secondary succession (Horn 1974). A major disturbance event prior to 2013 was the pan tropical bleaching event in 1998 (Wilkinson et al., 1999). Previous studies provided evidence that the benthic community of the Chagos Archipelago has transitioned through different successional stages since the 1998 bleaching event. A recovery occurred from minimal hard coral cover (~12% in 1999; Sheppard, 1999) to a coral-dominated community where no significant divergence was apparent in 2012, relative to the pre-1998 community (Sheppard et al. 2008, 2013b). Despite the overall reassembly towards a coral-dominated reef, there was significant variability in benthic community recovery, across depths and atolls (Sheppard et al., 2008). In 2001, less degradation was observed in deep sites of the northern atolls (PB and SA) (high mortality observed at <10-15 m) while the southern atolls (GCB and EG) showed significant mortality at depths >35 m (Sheppard et al., 2002). In 2006, the hard coral assemblages in shallow sites (4-10 m) recovered faster than deep forereef slopes (20-25 m) across PB, SA and GCB, with the exception of EG where no recovery was observed at either depth (Sheppard et al., 2008). In 2012, a general decrease in hard coral cover and increase in sponge and soft coral cover could be observed with increasing depth (5-25 m) across the archipelago (Sheppard et al., 2013b). This rapid recovery in shallow sites (8-10 m) may be driven by the high growth rates in *Acropora*-dominated shallow communities within the archipelago (Perry et al., 2015).

Our results reveal the spatial variation in depth zonation of benthic communities, potentially linked to previously described biophysical gradients occurring across these depths and atolls (Sheppard 2009; Fasolo 2013; Sheppard et al., 2017). These benthic community data from 2013 will provide insights into the impact of the 2014–2017 back-to-back bleaching events as well as subsequent recovery. It is becoming more important to understand the spatial variability in reef ecosystems in order to infer and predict how communities might respond to the effects of increasing disturbances (Bridge et al., 2014; Hughes et al., 2019). There is therefore a need to incorporate spatial variation in depth effect within ecological studies, as contemporary shallow reefs are changing rapidly across depths.

## **Author contributions**

Sivajyodee S Pilly: Conceptualization, Formal analysis, Validation, Data curation, Writing – original draft, Writing – review & editing, Visualisation. Laura Richardson, Ronan Roche: Validation, Writing – review & editing. John Turner, Ronan Roche: Methodology, Investigation, Supervision, Project administration, Funding acquisition.

## **Declaration of competing interest**

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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## Appendix A Supplementary data

Supplementary data to this article can be found online at <a href="https://doi.org/10.1016/j.marenvres.2021.105520">https://doi.org/10.1016/j.marenvres.2021.105520</a>.

### **Author statement**

All authors have agreed to the revised version of this manuscript.

### **Uncited References**

[Instruction: delete this reference - the system would not let me delete this duplication] Coral Reef Ecosystem Division,; JEN et al., 2019

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## **Highlights**

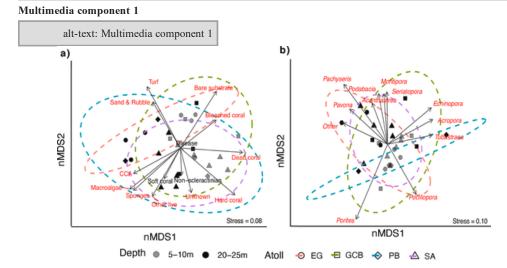
- Significant depth-by-atoll interactions drove benthic community structure.
- CCA, turf algae, sponge and soft coral cover were highest on deep reefs.
- Hard coral, dead coral and bare substrate cover were highest on shallow reefs.
- Acropora, Stylophora and Pocillopora characterised shallow reefs.
- Acropora and Porites consistently associated with both shallow and deep reef zones.

# Appendix A Supplementary data

The following are the Supplementary data to this article:

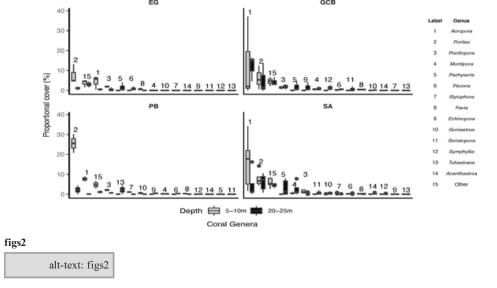


<u>■ Multimedia Component 1</u>



figs1

alt-text: figs1



## **Queries and Answers**

Q1

Query: Please confirm that the provided email "s.pilly@bangor.ac.uk" is the correct address for official communication, else provide an alternate e-mail address to replace the existing one, because private e-mail addresses should not be used in articles as the address for communication.

Answer: Reviewed

Q2

Query: Citation(s) 'Veron et al., 2019; System (2014); Sheppard, 1999; Chagos Archipelago, 2013' have not been found in the **reference list**. Please add the corresponding reference to the reference list.

**Answer:** We have now added/amended the above references:

- 1. Veron et al, 2019 Veron, J.E.N., Stafford-Smith, M.G., Turak, E., DeVantier, L.M., 2019. Corals of the World. Version 001Beta.
- 2. Sheppard 1999 has now been amended to sheppard 1999b: Sheppard, C.R.C., 1999b. Corals of chagos, and the biogeographical role of chagos in the Indian ocean. In: Sheppard, C.R.C., Seaward, M.R.D. (Eds.), Ecology of the Chagos Archipelago. Linnean society of london, pp. 53–66.
- 3. System 2014 has been amended as per newly added reference and updated in section 2.2 Benthic composition, last paragraph Coral Reef Ecosystem Division, 2014

  Coral Reef Ecosystem Division (CRED), Pacific islands fisheries science center (PIFSC), national marine fisheries service (NMFS) and National Oceanic and Atmospheric Administration (NOAA), 2014. Benthic percent cover derived from image analysis for selected locations in the pacific ocean.

https://www.coris.noaa.gov/metadata/records/faq/CRED\_ImageAnalysis.html. The system would not allow me to delete the old reference, kindly delete it please.

Q3

Query: Figs. 2, 3 were not cited in the text. Please check that the citation(s) suggested are in the appropriate place, and correct if necessary.

**Answer:** We do not need to cite Fig.2 and Fig.3 in this sentence. The sentence should be as follows:

"In this study, a total of 26 sites; including 13 at 5–10m depth (hereafter 'shallow' reefs) and 13 at 20–25m depth (hereafter 'deep' reefs), were surveyed on forereef slopes across 4 atolls: Peros Banhos (PB), Salomon (SA) (northern atolls), Great Chagos Bank (GCB) and Egmont (EG) (southern atolls) (Fig. 1, Table S1). Fig2 and 3 has now been cited accordingly in section 3 (Results).

Q4

Query: The citation 'Dennis et al., (2017)' has been changed to match the author name in the reference list. Please check here and in subsequent occurrences, and correct if necessary.

Answer: Reviewed

Q5

**Query:** Have we correctly interpreted the following funding source(s) and country names you cited in your article: Darwin, United Kingdom; Bertarelli Foundation, Switzerland; Bangor University, United Kingdom?

Answer: Yes - Darwin Initiative (Grant no - 19-027) and Bertarelli Foundation (Round 1, Project 6).

Q6

Query: The Uncited References section comprises references that occur in the reference list but are not available in the body of the article text. Please cite each reference in the text or, alternatively, delete it. Any reference not dealt with will be retained in this section.

Answer: Done

Q7

Query: Please confirm that given names and surnames have been identified correctly and are presented in the desired order and please carefully verify the spelling of all authors' names.

Answer: Reviewed

Q8

**Query:** Your article is registered as a regular item and is being processed for inclusion in a regular issue of the journal. If this is NOT correct and your article belongs to a Special Issue/Collection please contact s.hussain.1@elsevier.com immediately prior to returning your corrections.

Answer: No, this paper is not part of a special issue.