



# Distinct cross-shelf gradient in mesophotic reef fish assemblages in subtropical eastern Australia

Ryan Pearson\*, Tim Stevens

Griffith School of Environment and Australian Rivers Institute, Gold Coast campus, Griffith University, Queensland 4222, Australia

**ABSTRACT:** The hypothesized importance of mesophotic (30 to 150 m depth) reefs justifies the recent trend in scientific scrutiny of these systems. However, to enable assessment of complex concepts such as connectivity and reef dynamics, baseline assemblage characteristics must first be established. This study used baited remote underwater video (BRUV) technology to investigate the assemblage structure of predatory and scavenging fishes across 4 mesophotic reef bands at ca. 30 to 82 m depth, at 2 locations 25 km apart in subtropical eastern Australia. We aimed to quantify patterns in predatory and scavenging fish assemblage structure at these reefs across the continental shelf and relate this to putative structuring environmental variables. Strong cross-shelf gradients were identified in species richness and overall assemblage composition. While the pattern of latitudinal affiliation did not change across the shelf, predatory and scavenging fish assemblages at non-adjacent reefs were statistically distinct (PERMANOVA interaction term  $p = 0.012$ ), and best (but not well) explained by depth alone (BIOENV  $\rho = 0.396$ ). A high proportion (15 to 45%) of the fish species at each reef band were found only within that band. These cross-shelf trends contrasted with those described from more complex shelf topography such as at the Solitary Islands (250 km south), and did not match published patterns of epibenthic assemblage structure. Our results highlight the need for detailed information on mesophotic reef assemblage structure to support marine conservation and reserve design initiatives, rather than relying on generalised trends from the literature.

**KEY WORDS:** Mesophotic reefs · Subtropical fish · Predator assemblage · Drivers

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## INTRODUCTION

Marine ecosystems worldwide are under well-publicised anthropogenic threats (Hoegh-Guldberg 1999, Hughes et al. 2003, Halpern et al. 2012) that can have strong impacts on marine biodiversity and overall system resilience by impacting the various ecosystem services that biodiversity provides (Duffy 2002). Species within upper trophic levels help to maintain biodiversity and mediate competition (Hixon & Carr 1997) by the top-down process of selective predation, but also tend to be key fishery targets.

The roles of these higher trophic level fish taxa in structuring shallow water (<30 m) ecosystems through competition, predation, and scavenging are relatively

well studied, especially on coral reefs (e.g. McClanahan 1995, Hixon & Carr 1997, Sala 1997, Gomelyuk 2009). However, the influence of these processes on deeper reefs is poorly known, in part because of the cost and logistical difficulties associated with working in areas beyond the practical range of standard SCUBA equipment. In recent years, advances in low-cost and lightweight camera equipment, as well as in diving technology, have helped to fuel growing interest in the ecology of reefs in the mesophotic zone (30 to 150 m depth) (Hughes et al. 2003, Riegl & Piller 2003, Lesser et al. 2009, Puglise et al. 2009, Edgar et al. 2010). Tropical shallow water reefal species may be able to use reefs in the mesophotic zone and/or subtropical reefs as refuges in the face of anthro-

\*Corresponding author: ryan.pearson2@griffithuni.edu.au

pogenic and climate related challenges (Hughes et al. 2003, Riegl & Piller 2003, Bongaerts et al. 2010, Kahng et al. 2014). Mesophotic reefs may also harbour their own unique assemblages, with high levels of depth-endemism reported across multiple taxa, including corals (e.g. Bridge et al. 2012) and fishes (e.g. Pyle et al. 2008, Kane et al. 2014).

Mesophotic reef systems may also play an important role in marine ecosystem dynamics, resilience, and connectivity (Puglise et al. 2009, Tenggardjaja et al. 2014), in ways as yet poorly understood, though some patterns appear consistent. Vertical differences in mesophotic fauna are well established (Kahng et al. 2010, 2014). A transition between characteristic upper and lower mesophotic assemblages in the tropics has been reported at ca. 60 m (e.g. Bridge et al. 2012), and at ca. 50 m in the subtropics (e.g. Malcolm et al. 2010b). Despite these apparent transitions in assemblage structure, Tenggardjaja et al. (2014) found no obvious genetic differences between more than 400 *Chromis verater* specimens collected at shallow and mesophotic reefs (to 113 m) in Hawaii, supporting hypotheses inferring high vertical connectivity and deep refugia. There are also documented links between epibenthic and fish assemblages in the mesophotic zone (Brokovich et al. 2010, Malcolm et al. 2011b), but the strength of this association varies.

A handful of studies have highlighted specific trends in the cross-shelf assemblages of fishes, molluscs, and corals within the eastern Australian subtropics (Edgar et al. 2010, Malcolm et al. 2010a,b, 2011a,b, Harrison & Smith 2012). Most notably, consistent trends exist across all tested taxa in the Solitary Islands Marine Park (SIMP), where the proportion of tropical species increases with distance from shore on shallow reefs (Malcolm et al. 2007, 2010a, Harrison & Smith 2012) but decreases with depth (Malcolm et al. 2011b). The shallow water trends may be driven by the variable influence of conflicting water bodies in the region. The warm East Australian Current (EAC), which flows south along the continental shelf edge, operates mostly offshore in the SIMP and is subject to regular seasonal changes that affect its flow rate, width, and southern extent (Keane & Neira 2008). Inshore however, cooler counter-currents run northward along the coastline, in the opposite direction to the EAC. This creates 3 latitudinally distinct water bodies along the east coast, which correspond well with larval fish assemblages (Keane & Neira 2008, Neira & Keane 2008). The current study area in southeast Queensland (QLD) and northeast New South Wales (NSW) lies

consistently within the most northern, largely EAC-influenced water body (Neira & Keane 2008). Within this region, the distribution of reef biota is poorly known (Richmond & Stevens 2014), and no quantitative studies have examined mesophotic fish assemblages or their relationship to the characteristics of the water body.

Therefore, this study aims to (1) classify and test patterns in predatory and scavenging fish assemblages on mesophotic reefs across the continental shelf of subtropical eastern Australia; (2) evaluate differences in the latitudinal affiliations of derived fish assemblages across the continental shelf for comparison with trends noted elsewhere (especially in the SIMP); and (3) quantify the role of environmental drivers in structuring these assemblages.

## MATERIALS AND METHODS

### Study area

The Southport Seaway and Tweed River mouth are key entry points to ocean waters, situated 65 and 93 km, respectively, south of Brisbane in the vicinity of the Gold Coast on the east coast of Australia. Several discontinuous bands of rocky reef are found from ca. 3 to 35 km seaward of the Gold Coast, in depths of 30 to 82 m. These are well known to local recreational and commercial fishers, but have received little scientific attention (Richmond & Stevens 2014), and have not been accurately mapped.

### Sampling design

Sampling of predator and scavenger assemblage structure was conducted in winter 2013 using baited remote underwater video (BRUV) deployed for 30 min at 4 bands of rocky reef across the continental shelf directly east of the Southport Seaway (153° 25' E, 27° 56' S) and Tweed River mouth (153° 33' E, 28° 10' S) (Fig. 1). The BRUV method has known biases in that it attracts predatory and scavenging species and under-represents herbivores (see Colton & Swearer 2010 for example); nevertheless, it is acknowledged as one of the most precise methods available for fish abundance surveys (Bernard & Götz 2012) since it is not affected by diver avoidance or errors associated with various capture techniques. Moreover, it is low-impact and useable at any depth, on any substrate. Throughout this paper (unless otherwise stated), we refer to the predatory and scavenging reef fish as-

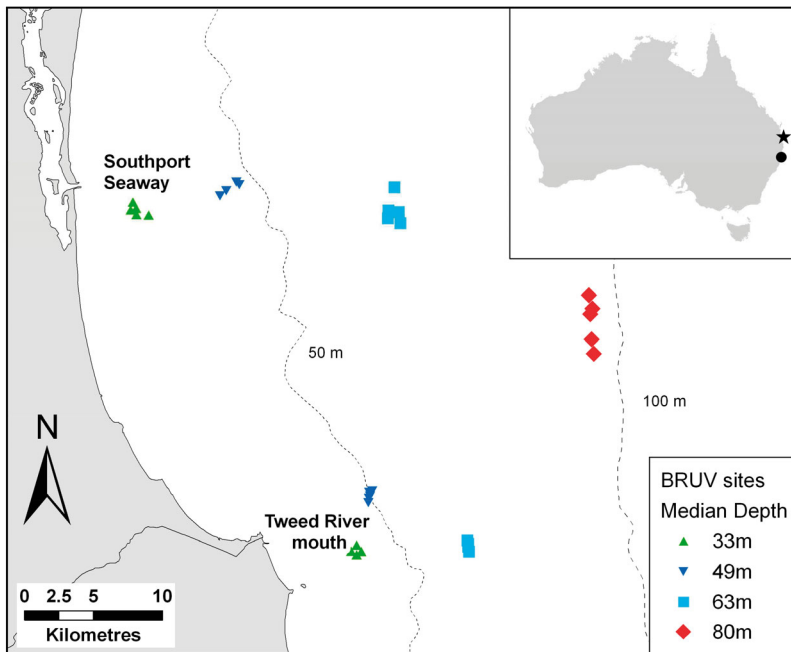


Fig. 1. Study area and sampling sites at Southport Seaway and Tweed River locations. The star in the inset map shows the study position, while the dot shows Solitary Islands Marine Park (SIMP). BRUV: baited remote underwater video

semblage sampled by this technique (Cappo et al. 2004).

The depth range for the reef bands sampled was as follows: band 1, 30 to 34 m depth (median 33 m); band 2, 46 to 51 m (median 49 m); band 3, 62 to 65 m (median 63 m); and band 4, 79 to 82 m (median 80 m). We use the median depth figures to refer to the reef bands throughout the paper (Fig. 1). At the deepest band, we were not able to locate reef offshore from the Tweed River mouth, so the design is unbalanced in this respect. Analyses were interpreted with this in mind, with most weight placed on the pairwise contrasts.

Waypoint data (not shown) obtained from the local charter fishing association was used to locate the reefs, and then BRUVs were deployed on reef patches with a linear extent of more than 200 m (estimated using the onboard echo sounder) to minimise bias associated with variation in reef patch size. At each site, nominally 5 replicate BRUVs were deployed in rotation for 30 min each, spaced at least 500 m apart to minimise overlap between bait plumes (Willis & Babcock 2000, Harvey et al. 2007, Colton & Swearer 2010). Each rig was baited with 2 pilchards *Sardinops sagax* (ca. 100 g) in the bait bag; one cut into 5 equal sections and the other placed into a nylon stocking and crushed to maximise release of oils. Each deployment was recorded on a GoPro2 camera with 2

iTorch 5 LED video lights to provide accurate and consistent colour information for species identification. Two 5 cm squares of Perspex (white with a black border to aid visibility) were attached to the bait pole, 15 cm apart on either side of the bait bag. These were used to estimate the size of a fish at the bait bag. The change in size on screen of this same individual was then used to estimate a distance ca. 2 m behind the bait bag to standardise the sampled volume for variations in water clarity between deployments, and to estimate distance from the bait bag for adjacent habitat descriptors as per Cappo et al. (2004).

All deployments were analysed by the same observer, and maxN recorded for each species; with MaxN defined as the highest number of individuals of a given species visible within a single frame of video throughout each 30 min deployment (Malcolm et al. 2011b). Individuals were identified to the finest

level of taxonomic classification possible using standard and locally relevant guides, principally Kuitert (1996), Allen et al. (2003), Allen (2009), and Davie et al. (2011) as well as online sources [www.fishbase.org](http://www.fishbase.org) and <http://australianmuseum.net.au>.

### Drivers for assemblage structure

Data on a range of putative drivers for fish assemblage structure were collected or assembled from existing data sources (Table 1). Temperature and salinity data were measured *in situ* using a conductivity, temperature and depth (CTD) probe (model RBR XR-420, CTD Marine) deployed simultaneously with the BRUVs. Incident light levels were assessed on most of the deployments (25 of 32; see 'Results') using a HOBO Pendant temperature and light logger (Model UA-002-64) attached to the CTD. Light intensity (in lux) was averaged over 30 s at the top (ca. 2 m depth) and bottom (ca. 2 m above the substrate, to avoid any disturbed sediment) of the water column, and used to calculate percentage light intensity remaining at the bottom.

Geographic attributes to assess the influence of land- or river-sourced inputs (distance from shore and distance to the nearest estuary) were derived using GIS software (ArcGIS v.9.2). Attributes of the

Table 1. Putative drivers for predator and scavenger fish assemblage structure

Driver	Source	Units
Depth	On-board echo sounder	m
Benthic relief	From video footage, within 2 m of bait bag	Categorical 1 = <10 cm 2 = 10 cm–1 m 3 = >1 m
Benthic complexity	From video footage, within 2 m of bait bag	Categorical 1 = <5 visible holes or crevices (> 5 cm diameter) 2 = 5–10 holes or crevices 3 = >10 holes or crevices
Benthic biota density	From video footage, within 2 m of bait bag	Categorical 1 = <20 individuals 2 = 20–40 individuals 3 = >40 individuals
Distance to estuary	ArcGIS v.9.2: distance to nearest river mouth or estuary opening	km
Distance to shore	ArcGIS v.9.2: distance to nearest part of the shoreline	km
Temperature at bottom	Measured <i>in situ</i> , CTD	°C
Salinity at bottom	Measured <i>in situ</i> , CTD	‰
Light remaining at bottom	Measured <i>in situ</i> , light logger	%, calculated from mean light in bottom 2 m / mean light at surface (lux)

local benthic community were derived from the field-of-view of the BRUV. Published quantitative methods for measuring habitat complexity (Luckhurst & Luckhurst 1978, Friedlander & Parrish 1998, Wilson et al. 2007) proved unworkable due to both the distortion inherent in the GoPro wide-angle lens and the depth of the reefs (which made manual measurement impossible). Therefore, the estimated distance behind the bait bag (2 m) was used as the outer limit for assigning 3 complexity descriptors: benthic relief, benthic complexity, and benthic biota density, which were each separated into 3 categories (Table 1). Benthic relief categories were based on vertical heights: <10 cm, 10 cm to 1 m, and >1 m. Benthic complexity categories were based on the number of visible holes ~5 cm diameter or greater: <5, 5 to 10, and >10. Lastly, benthic biota density categories were based on the number of individual benthic organisms visible within 2 m of the bait bag: <20, 20 to 40, and >40 (Table 1).

### Analytical design

The design tests for differences between the 4 reef bands (fixed: 33, 49, 63, and 80 m), and between the 2 locations (fixed: Seaway and Tweed), using both univariate (species richness, total abundance) and

multivariate (predatory and scavenging fish assemblage structure) metrics.

Univariate metrics were tested using 2-way (band × location) general linear models (GLMs) in SPSS v.21. A range of data transformations and model parameters were tested to meet the assumptions of linearity, normality, and homoscedasticity; those selected for the final analysis minimised goodness-of-fit statistics (–2 log likelihood; Akaike's information criterion, AIC and Bayesian's information criterion, BIC). Post hoc tests were conducted on significant terms to determine where the differences lay.

We tested for differences in species richness between latitudinal affiliations and reef bands. Latitudinal affiliations of each species were categorised as tropical, subtropical or temperate based on information in [www.fishbase.org](http://www.fishbase.org) (Froese & Pauly 2013). We acknowledge that latitudinal affiliations of some species in FishBase may be ambiguous or poorly known, therefore conclusions around these categories are drawn with caution.

For each transect, species richness of predatory and scavenging fish within each category was determined, and differences between categories and reef bands were tested using a linear mixed model (LMM) approach. The model tested the fixed factors band and affiliation and the random factor site

nested within band. As above, the data transformation was selected to minimise goodness-of-fit statistics while satisfying the assumptions of the test.

Differences in predatory and scavenging fish assemblage structure for the same 2-way (band  $\times$  location) design were tested using permutational multivariate analysis of variance (PERMANOVA) (Anderson et al. 2008) within the PRIMER 6 software package (Clarke & Gorley 2006). Multivariate analyses used the conventional Bray-Curtis similarity, because it ignores conjoint absences in a multi-species matrix (Anderson et al. 2008). Patterns in assemblage structure were visualised using non-metric multidimensional scaling (nMDS), and were very consistent across the range of data transformations. Results of PERMANOVA analyses were similarly consistent across data transformations; those presented are based on untransformed data, which had the lowest stress value in the nMDS. Significant relationships were further explored by pairwise testing, using Monte Carlo randomisation if the number of available permutations was low ( $<1000$ ). The distribution of fish species across reef bands was also examined to determine what proportion of observed species were confined to only 1 band, or ranged across several. SIMPER analysis was used to determine which individual taxa had the greatest influence on the derived patterns.

The influence of putative environmental drivers, both singly and in combination, on derived assemblage pattern was examined using the BIOENV analysis in PRIMER (Clarke & Gorley 2006), which compares the matrix of biological similarity with matrices derived from different combinations of abiotic variables. The abiotic variables were cross-correlated and, where variable pairs were highly autocorrelated (Spearman's  $\rho > 0.9$ ), the variable with the lower individual BIOENV correlation was removed to avoid overweighting in the multi-factor BIOENV models.

## RESULTS

### Overview

Between 11 June and 25 August 2013, 32 successful 30 min BRUV deployments were conducted on the 4 bands of rocky reef at 2 locations across the inner continental shelf, with the exception of the deepest reef band where only 1 location was available. Five replicate deployments were sampled at each reef band  $\times$  location, except in 3 instances where only 4 landed on reef substrate. The dataset

comprised 1917 individuals (based on MaxN) from 101 taxa, including 97 bony fish, 2 sharks and 2 rays (see Table S1 in the Supplement at [www.int-res.com/articles/suppl/m532p185\\_supp.pdf](http://www.int-res.com/articles/suppl/m532p185_supp.pdf)).

The most abundant species (444 ind.) was the Indian scad *Decapterus russelli*, although this species was not widespread (5 deployments). The Australian mado *Atypichthys strigatus* was arguably the most prevalent, being both widespread (25 deployments) and found in high abundance (391 ind.). The snapper *Pagrus auratus* was the most widely occurring (30 deployments), though present in comparatively low abundances (58 ind.) (Table S1).

In total, 38 fish families were represented; Kyphosidae were the most abundant (604 ind.), closely followed by Carangidae (588). Fewer than 87 individuals were observed from all other families. Sparidae was the most ubiquitous family, appearing in all but 1 deployment, though in relatively low abundance (86 ind.) (Table S1).

### Species richness and abundance

Species richness declined from inshore to offshore reef bands (Table 2, Fig. 2a), and this was consistent for the 2 locations. Species richness at adjacent reef bands (33 and 49 m, 49 and 63 m, 63 and 80 m) was not significantly different ( $p > 0.05$ ), but was different for bands more widely separated, supporting the roughly linear decline from inshore to offshore (Fig. 2a).

Table 2. Two-way (band  $\times$  location) general linear model (GLM) analyses for species richness and total abundance of predatory and scavenging fishes on continental shelf reefs. Data were  $\log_{10}(x + 1)$  transformed to meet the assumptions of the test. Significant differences ( $p < 0.05$ ) are given in **bold**

Fixed effects	Numerator df	Denominator df	F	p
<b>Species richness</b>				
Band	3	25	3.789	<b>0.023</b>
33 vs. 49 m				0.679
33 vs 63 m				0.065
33 vs 80 m				<b>0.006</b>
49 vs 63 m				0.144
49 vs 80 m				<b>0.014</b>
63 vs 80 m				0.181
Location	1	25	0.297	0.591
Band $\times$ Location	2	25	0.359	0.702
<b>Total abundance</b>				
Band	3	25	0.721	0.549
Location	1	25	2.216	0.149
Band $\times$ Location	2	25	2.054	0.149

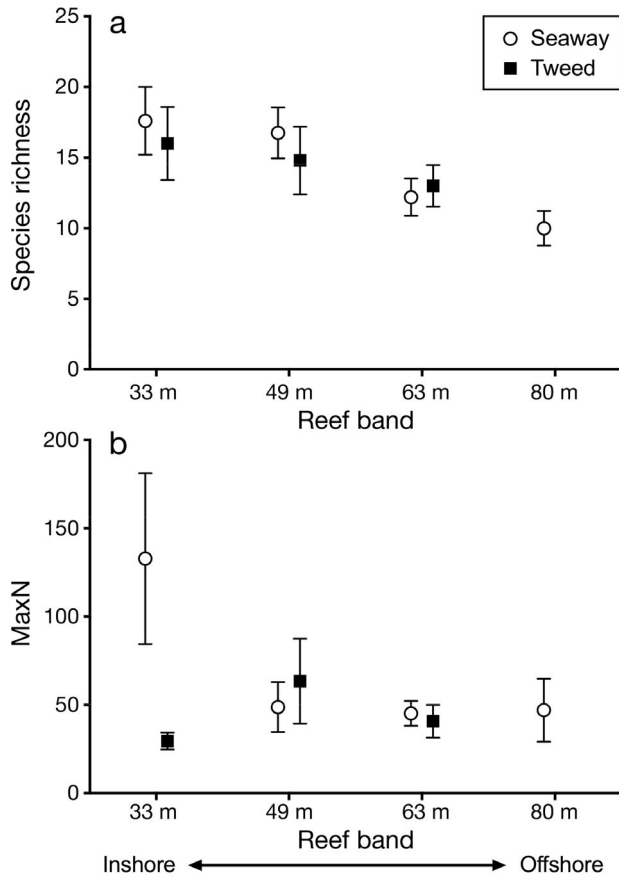


Fig. 2. Mean ( $\pm$ SE) (a) species richness and (b) total abundance (MaxN) from baited remote underwater video (BRUV) sampling of predatory and scavenging fishes on continental shelf reefs

In contrast, total abundance of predatory and scavenging fishes was not different between reef bands or locations (Table 2, Fig. 2b). In spite of the apparent difference between locations inshore (33 m reef band), when transformed to meet the assumptions of the test there was no statistical difference. This was confirmed by pairwise testing ( $p > 0.05$  for all band  $\times$  location combinations), and was consistent across different valid transformations.

**Latitudinal affiliations**

Analysis of the latitudinal affiliations associated with species richness within each band of reef revealed no significant interaction ( $p = 0.661$ ) in the reef band  $\times$  affiliation term (Table 3, Fig. 3), indicating that there was no change in latitudinal affiliation across the shelf. As suggested by Fig. 3, there were differences within the affiliation term ( $p < 0.001$ ), with pairwise analysis revealing that subtropical spe-

Table 3. Linear mixed model (LMM) analyses for species richness by latitudinal affiliation (tropical, subtropical, temperate) and reef band (33, 49, 63, and 80 m) of predatory and scavenging fish species on continental shelf reefs. Species richness was  $\log_{10}(x + 1)$  transformed to meet the assumptions of the test. Significant differences ( $p < 0.05$ ) are given in **bold**

	Numerator df	Denominator df	F	p
<b>Fixed effects</b>				
Reef band	3	28	3.688	<b>0.024</b>
Affiliation	2	28	16.914	<b>&lt;0.001</b>
Reef band $\times$ Affiliation	6	56	0.686	0.661
	Estimate	SE	Z	p
<b>Random effects</b>				
Site (reef band)	0.004804	0.004113	1.168	0.243

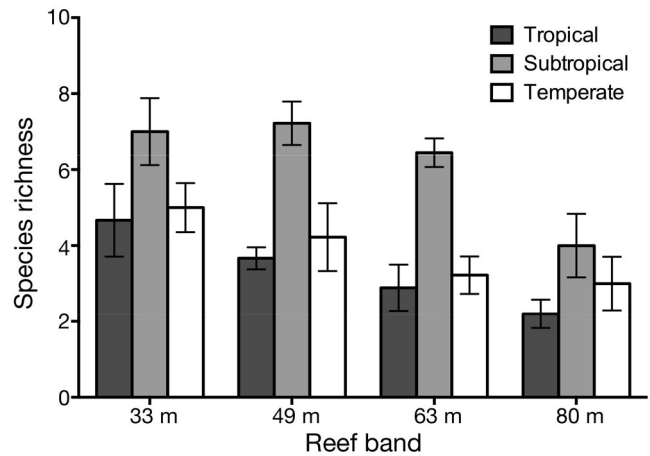


Fig. 3. Latitudinal affiliation of predator and scavenger fishes at each reef band as mean ( $\pm$ SE) species richness in each category

cies richness was significantly different ( $p < 0.001$ ) from both temperate and tropical species richness, which were not different from one another ( $p = 0.294$ ). There were significant differences between reef bands, consistent with the previous analysis.

**Assemblage structure**

There were clear patterns in predatory and scavenging fish assemblage structure (Fig. 4), confirmed by the significant PERMANOVA interaction term (Table 4). Pairwise test results showed that the assemblage at the furthest offshore sites (80 m band) was clearly distinct. Of the shallower 3 bands, assemblage structure at adjacent sites was not significantly

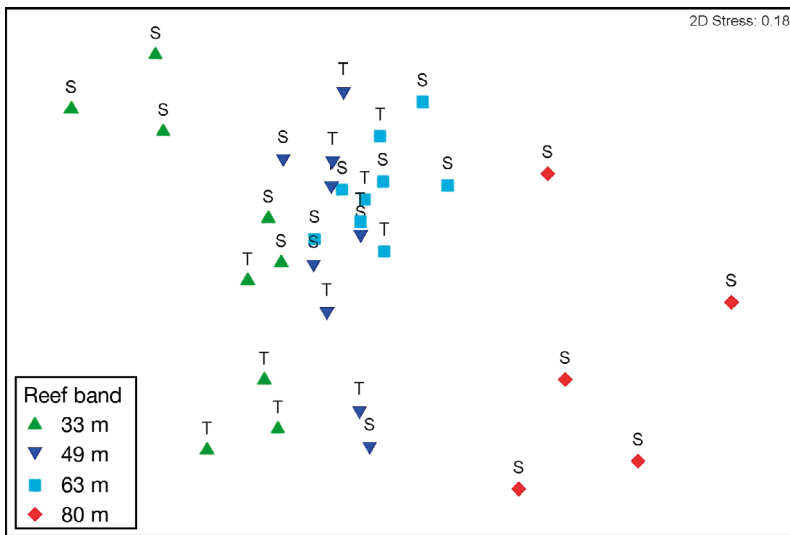


Fig. 4. Non-metric multidimensional scaling (nMDS) of predator and scavenger fish assemblage structure from baited remote underwater video (BRUV) sampling on continental shelf reefs. Symbols correspond to the sites illustrated in Fig. 1; S: Southport Seaway; T: Tweed River. Results are based on Bray-Curtis similarity from untransformed data; other transformations showed very similar patterns, but with higher stress levels

Table 4. Two-way (band × location) PERMANOVA for predatory and scavenging fish assemblage structure on continental shelf reefs. Based on Bray-Curtis similarity matrix from untransformed data. p-values derived by Monte-Carlo randomisation for pairwise tests because the number of unique permutations was low. Significant differences ( $p < 0.05$ ) are given in **bold**

	Numerator df	Denominator df	Pseudo-F	p	Unique permutations
<b>Fixed effects</b>					
Band	3	25	3.265	<b>&lt;0.001</b>	9859
Location	1	25	1.583	0.092	9882
Band × Location	2	25	1.850	<b>0.012</b>	9886
<b>Pairwise</b>					
			t	p	Unique permutations
<b>Seaway</b>					
33 vs. 49 m			1.565	0.068	126
33 vs. 63 m			1.754	<b>0.030</b>	126
33 vs. 80 m			1.831	<b>0.012</b>	126
49 vs. 63 m			1.291	0.156	126
49 vs. 80 m			1.550	<b>0.040</b>	126
63 vs. 80 m			1.702	<b>0.018</b>	126
<b>Tweed</b>					
33 vs. 49 m			1.455	0.085	126
33 vs. 63 m			2.222	<b>0.008</b>	35
49 vs. 63 m			1.253	0.195	126
<b>Seaway vs. Tweed</b>					
33 m			1.894	<b>0.019</b>	126
49 m			1.056	0.354	126
63 m			1.004	0.403	126

different ( $p > 0.05$ ). However, the 2 more spatially separated sites (33 and 63 m) did display significant differences ( $p < 0.05$ ), reinforcing the picture of progressive change in the predatory and scavenging fish fauna from inshore to offshore. Assemblage structure at the most inshore band (33 m) was different between the Tweed and Seaway locations ( $p = 0.019$ ), but this was not the case at the 49 or 63 m bands.

At the most inshore reef band, 45% of fish species were observed in no other reef band (Table 5a). This proportion fell to 15% at the 63 m band, but rose again to 41% at the 80 m band (only sampled offshore from the Seaway), highlighting the distinct fish assemblage in this deepest, furthest offshore reef band, as shown by the nMDS and pairwise results. Overall, more than half the observed fish fauna (55 of 101 species; Table 5b) occurred at only 1 depth band, with only 6 species observed at all 4 bands. A total of 9 species were observed within both the shallowest and deepest reef bands (Table S1).

SIMPER analyses indicated that 3 species (*Decapterus russelli*, *Atypichthys strigatus*, and *Seriola dumerili*) were the highest contributors to dissimilarity between reef bands, and to similarity within the bands. *D. russelli* was the single most abundant species in the study (Table S1), but was only observed within the 33 m band. It therefore represented a large proportion of the difference between the 33 m and all other reef bands (SIMPER: 33 vs. 49 m = 24%; 63 m = 28%; 80 m = 40%). This species also accounted for 30% of the similarity within 33 m reef bands. At the other end of the depth spectrum, *S. dumerili* was the 3rd most abundant fish species, but occurred only in the 80 m reef band, thus accounting for the greatest contribution to assemblage dissimilarity (17%) between the 49 and 80 m reef bands, and 24% of similarity within the 80 m band. *A. strigatus* (the 2nd most abun-

Table 5. Fish species distributions between reef bands, as (a) proportion of all fish species occurring at a reef band that occur only at that reef band, and (b) number of fish species by the number of reef bands in which they occur. A total of 9 baited remote underwater video (BRUV) samples were taken at each reef band except for the 80 m reef band where  $n = 5$ , since this depth was not sampled at the Tweed location

(a) Reef band	Total no. of fish species occurring	No. of fish species occurring only in this band	Proportion occurring only in this band (%)
33 m	55	25	45
49 m	55	14	25
63 m	39	6	15
80 m	27	11	41

(b) No. of bands	No. of fish species
1	56
2	21
3	18
4	6

vdant fish species; Table S1) occurred across all 4 reef bands, but at different densities. It was responsible for 25 % of assemblage dissimilarity between the 49 and 63 m bands, and 20 % of dissimilarity between the 63 and 80 m bands. *A. strigatus* accounted for 34 % of similarity within the 49 m reef band and 58 % of similarity within the 63 m reef band.

### Drivers of assemblage structure

BIOENV analyses of individual drivers of predatory and scavenging fish assemblage showed that depth, distance to shore, and distance to estuary were the most important factors, all with similar correlation values (Table 6). Not surprisingly, these were strongly autocorrelated, and thus only depth was used in the subsequent combined models. Salinity, benthic relief, and benthic complexity all had a significant influence ( $p \leq 0.004$ ), although correlation values were low (approximately 0.2). Perhaps surprisingly, light, temperature, and benthic density had no detectable influence ( $p > 0.05$ ; Table 6). Separate analyses of light intensity versus depth from the CTD to which it was attached showed the expected strong negative exponential relationship ( $r^2 = 0.747$ ). However, light intensity varied markedly at small spatial and temporal scales, indicating that 'snapshot' readings may be of little value, and that long-term mean values would be required to assess the direct rela-

Table 6. BIOENV analysis: Spearman's correlations of individual drivers and best combined models with observed predator and scavenger fish assemblage structure. Based on ranked Euclidean distance similarity for abiotic data, and Bray-Curtis similarity from untransformed data on fish assemblage structure. Values in **bold** denote significant ( $p < 0.05$ ) correlations.  $n = 32$ , except for 'light remaining' values, which were not available for all deployments; this value is based on a reduced set of 25

	$\rho$	$p$
<b>Individual drivers</b>		
Depth	0.396	<b>&lt;0.001</b>
Distance to estuary	0.366	<b>&lt;0.001</b>
Distance to shore	0.358	<b>&lt;0.001</b>
Salinity	0.218	<b>&lt;0.001</b>
Benthic relief	0.210	<b>0.002</b>
Benthic complexity	0.195	<b>0.004</b>
Light remaining	0.079	0.143
Bottom temp	0.075	0.136
Benthic biota density	0.036	0.224
<b>Best models</b>		
Depth	0.396	<b>&lt;0.001</b>
		in every case
Depth, salinity	0.384	
Depth, relief, salinity	0.347	
Depth, complexity, salinity	0.342	
Depth, relief, bottom temperature, salinity	0.330	

tionship between light and assemblage structure. It also suggests that depth is a suitable proxy for light intensity in short term studies such as this.

The BIOENV model that best explained the observed patterns in predatory and scavenging fishes contained depth alone ( $\rho = 0.396$ ,  $p < 0.001$ ). Adding drivers to the model, while still displaying relatively high correlations, only resulted in a weaker relationship with assemblage structure (Table 6). This was also the case if either of the terms autocorrelated with depth (distance to estuary and distance to shore) was substituted for depth in the analysis (data not shown).

## DISCUSSION

### Summary

This study revealed the existence of a very clear and consistent cross-shelf gradient in species richness of scavenging and predatory fishes on mesophotic rocky reefs that declines approximately linearly over 4 reef bands within a depth range of 30 to 82 m. Perhaps surprisingly, this was not matched by a gradient in overall abundance, which was not signif-



icantly different between depths or locations, in contrast to patterns reported elsewhere (see Kahng et al. 2014 for review).

Multivariate analyses of fish assemblage structure revealed a clear cross-shelf pattern. Assemblages at adjacent reef bands were generally not significantly different, but were different from those further away. The scavenging and predatory fish assemblage at the 80 m band was different from all others, but the latitudinal analysis showed, perhaps surprisingly, that this did not represent a tropical dominated assemblage recruiting from reefs further north, as would be expected if the offshore reefs were influenced by the southward flowing EAC. The fish assemblage was dominated by subtropical fish species at all reef bands. While the 33, 49, and 63 m reef bands were more closely related to one another, each contained a subset of species (ranging from 15 to 45%) that were not observed at other bands.

Depth was the most important individual driver for patterns in assemblage structure, and on its own was better than any combination of drivers. However, distance to shore and to the nearest estuary were nearly as important, and salinity, benthic relief, and complexity were also significantly related to observed assemblage structure.

### Relationship with other fish studies

Studies on reef fish fauna at the SIMP, on the eastern Australian coast about 260 km south of the present study, reveal contrasting trends when comparing distance from shore and depth gradients. Malcolm et al. (2010a) found that species richness on shallow (<25 m) reefs increased with distance offshore, and that this was largely a result of the addition of tropical species. However, species richness decreased with depth down to 65 m (Malcolm et al. 2011b), as was the case in this study. Further afield, fish species richness at subtropical (ca. 28.5°N) mesophotic reefs also declined with depth in the Gulf of Aqaba (Brokovich et al. 2008), as did the density of specific trophic levels (e.g. grazing fishes) (Brokovich et al. 2010).

In tropical coral reef locations, there is a distinct transition between euphotic and mesophotic fish communities, with a general pattern of increasing fish species richness to a maximum at about 25 to 30 m, then decreasing monotonically to the lower limit of the survey, typically 50 to 65 m (e.g. Garcia-Sais 2010, Kahng et al. 2010). At Isla Desecheo in the Caribbean, Garcia-Sais (2010) defined distinct differences in fish assemblages between euphotic (15 to 25 m) and

mesophotic (30 to 50 m) depths, as well as distinct differences within the mesophotic zone, with the deepest (50 m) being different from all other depths. Detailed mesophotic fish assemblage surveys are rare in the literature, but a similar picture is apparent from locations in the southwest Atlantic (Pereira-Filho et al. 2011), the Gulf of Mexico (Dennis & Bright 1988) and western Australia (Fitzpatrick et al. 2012). Our results are consistent with these trends in both tropical and subtropical regions in that fish assemblage is clearly different between upper- and mid-mesophotic depths. Comparable studies in temperate waters typically sample a far greater depth range, in that they sample the mesophotic zone as the upper extent of surveys into aphotic depths. Nonetheless, fish species richness and assemblage structure are also strongly related to depth, both in the mesophotic (Williams & Bax 2001) and euphotic depths (Hyndes et al. 1999).

High levels of apparent depth-endemism were observed in this study as over half of the species occurred in only 1 reef band, similar to observations elsewhere in the Pacific (Pyle et al. 2008, Kane et al. 2014), although it is not possible to separate the depth effect from distance offshore in this study. Of the 101 species encountered, only 9 appeared in both 33 and 80 m bands, and only 6 occurred across all depth bands. In contrast, 56 species were apparent depth specialists. Therefore, the capacity of the deep reefs to act as refuges and sources of recruitment for shallower reefs may be limited to a relatively small pool of species, although investigations over longer time spans would be required to determine this. That said, the deep reef refugia hypothesis relates to the re-supply of euphotic reefs (Bongaerts et al. 2010), which were not examined in this study. Nevertheless, our findings add weight to the importance of mesophotic reefs in maintaining biodiversity through harbouring apparently depth-specialist species.

The sites on the reef bands we surveyed constitute spatially isolated reefs emerging a few metres above the flat sandy sediments of the continental shelf, which fall only 50 m over a distance of more than 30 km. Given that the 80 m reef band is about 16 km from the nearest 63 m reef, it is not surprising that the fish assemblage here is distinct from all others, but nonetheless interesting that latitudinal affiliations are not different. Furthermore, trends in species richness across the depth range of the continental shelf conform to those from the bulk of studies, most of which examined different depths on a continuous rocky or coral reef slope. This has important implications for scales of connectivity in mesophotic reefs (Slattery et al. 2011) and for reserve design.

### Relationship with benthic assemblages

Published trends in macrobenthic species richness, abundance, and assemblage structure of mesophotic reefs vary widely, but do not necessarily correspond to observed patterns in fish assemblages. Richmond & Stevens (2014) documented macrobenthic assemblage structure at the same reefs as the current study (excluding the 80 m sites) as part of a wider appraisal of benthic biodiversity in the region. Unlike the current study, macrobenthic assemblages were distinctly different between the Seaway and Tweed locations for the inner 2 reef bands, but not separated at the 63 m band, which was different from the shallower assemblages. There were also distinct gradients in both species richness and total abundance of macrobenthic taxa, which both increased with depth and distance offshore (Richmond & Stevens 2014). That is, the observed gradients were the opposite of that found for predatory and scavenging fish species richness in this study, indicating that the fish and benthos are not closely coupled in this setting. In addition, assemblages in the 63 m reef band were characterised by the presence of tropical taxa not found at the shallower, more inshore reefs. This is consistent with other eastern Australian benthic studies (e.g. Harrison & Smith 2012), and is attributed to the influence of the south-flowing EAC bringing invertebrate larval forms from tropical waters, which then settle and colonise the offshore reefs, whereas inshore reefs are subjected to a northward flowing counter current. However, it is apparent that these influences do not appear to operate in the same way for scavenging and predatory fish in this situation, although the influence of the EAC is cited as a driver for patterns in fish assemblage structure in the Solitary Islands (Malcolm et al. 2010b) where there is moderately good correlation between fish and dominant benthic assemblages (Malcolm et al. 2011b). Richmond & Stevens (2014) speculated that some near-shore reefs maybe overfished, leading to a trophic cascade, but further work is required to test this assertion.

### Possible drivers of observed fish assemblage patterns

The role of depth as a key driver for fish assemblage patterns fits with findings elsewhere (Malcolm et al. 2010b, 2011b), though it is rarely the sole driver; distance from shore, habitat structure, and species diversity also show positive relationships with fish assemblage structure (Luckhurst & Luckhurst 1978, García-Charton & Pérez-Ruzafa 1998, Gomelyuk 2009, Mc-

Clain & Barry 2010). We found that depth alone is the best indicator of assemblage structure, but in part, this may be because the arrangement of reefs relative to the coast in the current study was linear. Shallow reefs only occur near shore, and deep reefs only occur offshore in the study area. Thus, it is not possible to separate the influence of depth and distance offshore due to simultaneous changes in each. In addition, our study reports on the fish fauna of reefs separated by 10s of kms of sand flats whereas many others report on different depths within the same reefs (e.g. Brokovich et al. 2008, Bejarano et al. 2014).

Given the long established relationship between light penetration and depth (Gordon & McCluney 1975), it should be noted that the depth-based relationship identified here cannot be disentangled from the effect of decreasing light availability. That said, high spatio-temporal variation in water clarity means that depth is generally a more practical proxy for long-term light intensity averages (see Armstrong et al. 2006, Smith et al. 2010, and Amado-Filho et al. 2012 for examples). However, light alone is unlikely to be limiting at most of the depths studied, given that the kelp *Ecklonia radiata* is found at high densities at depths of ca. 70 m offshore from Moreton Island, less than 100 km north of the study area (Marin 2013, Richmond & Stevens 2014). In other mesophotic ecosystems throughout the world, reported shifts from herbivory to carnivory (especially planktivory; Kahng et al. 2010) with depth, even where edible macroalgae is abundant (Garcia-Sais 2010), suggest that other factors are at play. In the local context, these could include localised shelf-break upwellings (Baird et al. 2006, Dela-Cruz et al. 2008, Marin 2013) or fishing pressure (Richmond & Stevens 2014).

### Future research

Much work remains to be done to establish the wider role of mesophotic reefs in continental shelf ecology. In the local context, studies to examine the along-shelf distribution patterns of nektonic taxa are underway. Future studies in the region would benefit greatly from more detailed bathymetric data (e.g. from swath or multibeam acoustic surveys). Such information is necessary to design studies to examine the more pressing questions raised in this study—specifically, to understand scales of connectivity in fish and invertebrate biota between isolated reef outcrops and their relationship with the fauna of inter-reef soft substrates—information that is critical for future conservation initiatives.

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