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# **Beyond the reef: the influence of seascape structure on the composition and function of tropical fish communities**



Thesis submitted by  
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in June 2021

For the degree of Doctor of Philosophy in Marine Ecology  
in the ARC Centre of Excellence for Coral Reef Studies,  
James Cook University, Australia



This thesis is dedicated to

*Dr Eilir Hedd Morgan, Dr Giulia Cambiè, and Kate Downes*

Marine scientists, educators, mentors, and friends. I am forever indebted to each of you for your friendship, kindness, humour, and wisdom along this journey.

You will never be forgotten.

*Those who dwell, as scientists or laymen, among the beauties or mysteries of the earth are never alone or weary of life... Those who contemplate the beauty of the earth find reserves of strength that will endure as long as life lasts.*

- Rachel Carson, *The Sense of Wonder* (1965)

*The first law of ecology is that everything is connected to everything else*

- Barry Commoner (1971)

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## Statement of the contribution of others

---

This research is the result of collaborations with my Advisors Professor Andrew S Hoey, Dr Mary C Bonin, Professor Graeme Cumming, Dr Stephanie Duce and Professor Serge Andréfouët. Throughout the thesis, I was responsible for the project concept and design, data collection and analysis, interpretation and all written work. My Advisors provided editorial and intellectual input (all), financial support (ASH), equipment (ASH, MCB, and SD), and the benthic habitat map (SA).

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## Ethics and copyright

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This research has been conducted under the James Cook University Animal Research Ethics Committee approval numbers A2455 and A2526.

Every reasonable effort has been made to gain permission and acknowledge the owners of copyright material. I would be pleased to hear from any copyright owner who has been omitted or incorrectly acknowledged.

.....

Katie Sambrook

June 2021

## List of publications arising from this thesis

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**Sambrook K**, Hoey AS, Andréfouët S, Cumming GS, Duce S, Bonin MC (2019) Beyond the reef: The widespread use of non-reef habitats by coral reef fishes. *Fish & Fisheries* 20: 903-920 (**Chapter 2**).

**Sambrook K**, Bonin MC, Bradley M, Cumming GS, Duce S, Andréfouët S, Hoey AS (2020) Broadening our horizons: seascape use by coral reef-associated fishes in Kavieng, Papua New Guinea, is common and diverse. *Coral Reefs* 39: 1187-1197 (**Chapter 3**).

**Sambrook K**, Andréfouët S, Aston EA, Bonin MC, Cumming GS, Duce S, Sievers KT, Hoey AS (in revision) Relative importance of seascape versus within-habitat variables on the distribution of fishes in tropical seagrass beds. *Estuarine, Coastal and Shelf Science* (**Chapter 4**).

**Sambrook K**, Bonin MC, Cumming GS, Duce S, Andréfouët S, Hoey AS (in prep) Influence of seascape and within-reef variables on a key ecosystem process, macroalgal browsing, on coral reefs in Papua New Guinea. *Oecologia* (**Chapter 5**).

### Other outputs during candidature

Torda G\*, **Sambrook K**\*, Cross P, Sato Y, Bourne DG, Lukoschek V, Hill T, Torras Jorda G, Moya A, Willis BL (2018) Decadal erosion of coral assemblages by multiple disturbances in the Palm Islands, central Great Barrier Reef. *Nature Scientific Reports* 8: 11885 \**Shared first authorship*

Jones GP, Barone G, **Sambrook K**, Bonin MC (2020) Isolation promotes abundance and species richness of fishes recruiting to coral reef patches. *Marine Biology* 167: 1-13

## Abstract

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Tropical shallow-water seascapes are heterogeneous environments composed of a mosaic of different habitat patches (e.g., coral reef, mangroves, macroalgal and seagrass beds) that occur in varying shapes, sizes and spatial configurations. These habitat patches are ecologically linked through the flow of materials, nutrients and organisms yet are often studied as discrete entities. While some species of coral reef fishes are known to use multiple habitat types during their lifecycle, we still know very little about the extent of non-reef habitat use among reef fishes, or how the spatial patterning of seascapes might influence fish-habitat relationships or ecological processes associated with coral reef seascapes. Thus, my overarching aim was to examine the influence of seascape structure on the composition and function of tropical fish communities, primarily those associated with coral reef habitats. Specifically, my objectives were to: (1) determine the extent of non-reef habitat use by coral reef fishes at a global scale; (2) examine fish-habitat relationships in tropical seascapes in Papua New Guinea; (3) investigate the relative influence of seascape structure and within-habitat variables on fish distribution and abundance patterns in tropical seagrass beds; and (4) examine the influence of seascape structure and within-reef benthic variables on macroalgae browsing activity on coral reefs.

**Chapter 2** systematically reviews and synthesizes the evidence for non-reef habitat use by coral reef fishes across biogeographic regions, life stages and feeding guilds. Non-reef habitat use was found to be extensive, with at least 670 species of coral reef fish observed away from the reef in a wide range of habitats and study locations. Three-quarters of the fish species for which adult and juvenile data were available occurred in non-reef habitats at both life stages, which suggests that fish interactions with non-reef habitats are complex and multifaceted. In addition, over half of the species recorded from non-reef habitats are potential fisheries targets. In the Caribbean, a similar number of fish species used mangroves and seagrass beds. In contrast, in the Indo-Pacific, seagrass beds emerged as the dominant non-reef habitat used by reef fishes. These differences between biogeographic regions may reflect environmental variation (e.g., tidal regimes) or be a result of unbalanced survey effort across habitats in the Indo-Pacific. Based on the synthesis, I estimate approximately one-fifth of 'coral reef' fishes could be non-reef habitat users, but suggest numbers could be higher given the paucity of data for much of the Indo-Pacific region.

**Chapter 3** builds on some of the research gaps identified in the global synthesis and examines fish-habitat relationships in the remote Tigak Islands, New Ireland Province, Papua New Guinea in the Indo-Pacific. Using remote video cameras, fish communities were surveyed in five habitat types (coral reef slope, coral reef flat, macroalgal beds, mangroves and seagrass meadows). Of the 282 shallow-water reef-associated fish species observed across 360 videos, 99 species (35%) were recorded in non-reef habitats, the majority of which (78 species) were seen on multiple occasions. This suggests that non-reef habitat use by coral reef fishes could be even more extensive than suggested by my synthesis (**Chapter 2**) and reinforces the need to incorporate a range of habitats into study designs in the Indo-Pacific. Of the three non-reef habitats, species richness and relative abundances were highest in macroalgae beds. Fish communities in mangroves and seagrass beds were distinct from each other but contained a similar number of species. Supporting the findings of **Chapter 2**, many of the fish families commonly found in non-reef habitats in this location, particularly from seagrass beds, are targeted by local fishers and are critical for sustaining local livelihoods.

The seagrass meadows around the Tigak Islands cover large areas of shallow water, harbour diverse fish communities, support coastal fisheries and are found in close proximity to a range of other habitat types. In **Chapter 4**, I use a multi-scale approach to examine the relative influence of within-habitat variables and seascape structure on the distribution patterns of 17 fish taxa (i.e., families and species), and life stages (adults and juveniles) for two taxa, in 32 seagrass patches around the Tigak Islands. Of the 21 fish groups examined, two-thirds were influenced more by seascape than within-habitat variables. Site-attached taxa were generally more influenced by within-habitat variables, while taxa with larger home ranges (e.g., rabbitfishes, snappers) and ontogenetic habitat-shifters (e.g., emperors) were generally more influenced by the surrounding seascape. Despite the majority of taxa being reef-associated, proximity to or area of adjacent coral reefs appeared to have limited influence on local abundances. Instead, proximity and area of hardground, macroalgae and mangroves emerged as more influential. My findings show that seagrass patches are not functionally equivalent to each other and that the surrounding seascape can be more important in shaping tropical fish assemblages in seagrass beds than within-habitat variables.

Given the evidence that the surrounding seascape can influence fish distribution patterns, **Chapter 5** explores whether features of the seascape can also impact a key ecological process on coral reefs, specifically macroalgae browsing by reef fishes. I investigated whether seascape

and within-reef benthic variables influenced the rates and agents of macroalgal removal using experimental macroalgae assays and remote video cameras at 24 shallow coral reef sites around the Tigak islands. I predicted that the presence of potential alternative sources of food near to reefs (e.g., neighbouring macroalgal beds), would affect macroalgal removal rates, community composition and/or browsing behaviour of fishes. Contrary to expectations, macroalgae at the seascape or within-reef scale was not a good predictor of macroalgal removal rates. Instead, both rates and agents of browsing were most strongly influenced by distance to oceanic environment, and exhibited differential responses to other seascape (e.g., area of seagrass within 250 m) and benthic variables. My findings provide support for the notion that the surrounding seascape can influence ecological processes on reefs.

Overall, my research highlights that non-reef habitat use among fish typically classified as reef-associated is more common and widespread than has previously been recognised, yet we still know very little about reasons underpinning multi-habitat use. This thesis contributes to the emerging field of seascape ecology by demonstrating that the wider seascape can influence both fish-habitat relationships and ecological processes. Importantly it provides strong support for the need to transition from a habitat- or patch-centric approach to encompass the surrounding seascape. It shows that the spatial context of a habitat patch may affect who uses the patch, from species to life stages, and this information has the potential to be particularly important in terms of identifying and managing critical shallow-water nursery habitats. The adoption of a multi-scale approach to examine fish-habitat relationships and ecological processes in coastal ecosystems can provide a more nuanced perspective, enabling the development of more refined spatial management strategies. Given the widespread loss and degradation of many coastal habitats in the tropics, combined with the importance of coastal fisheries for developing island nations, this thesis highlights the need to scale up research to incorporate the broader seascape if we are to manage and study shallow-water coastal ecosystems effectively in the future.

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# Chapter 1

## General Introduction

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How animals perceive, select and interact with their environment have been longstanding research themes in ecology. Such information can provide valuable insight into how populations and communities are assembled (Hairston et al. 1960; Paine 1966; Estes & Palmisano 1974; Lubchenco & Gaines 1981; Paine & Levin 1981), how species support the functioning of ecosystems (e.g., herbivory, nutrient cycling, Naeem et al. 1994; Tilman et al. 1997; Walker et al. 1999; Burkepile & Hay 2008), maintain ecosystem integrity (e.g., prevention of tipping points, Tilman & Downing 1994; Laurance et al. 2011; Newbold et al. 2016), and sustain ecosystem services (e.g., agriculture, fisheries, coastal protection) that promote human well-being (Scheffer et al. 2001; Elmqvist et al. 2003; Cadotte et al. 2011; Oliver 2016). However, unravelling species-environment relationships is inherently complicated given that such relationships are the product of a complex interplay between numerous abiotic (e.g., light, temperature, precipitation) and biotic (e.g., predation, competition, mutualism) variables operating across a range of spatial and temporal scales (Hutchinson 1957; Connell 1961; Whittaker 1965; Connell 1983; Menge & Olson 1990; Levin 1992; Menge 2000; Harborne et al. 2011).

The spatial distributions of species are shaped by their tolerance to abiotic and biotic conditions (i.e., fundamental niche), as well as a range of intrinsic (e.g., need to find food, find a mate) and extrinsic (e.g., predation, competition) factors that determine the environment where an individual actually occurs (i.e., realised niche; Hutchinson 1957). Several theories have been proposed to explain how individuals are distributed across space (e.g., ideal free distribution, Fretwell & Lucas 1970), and how they make decisions around habitat or patch selection (e.g., optimal foraging theory, Emlen 1966; MacArthur & Pianka 1966; central place foraging, Orians & Pearson 1979; coexistence theory, Chesson 2000; and the landscape of fear, Laundré et al. 2001). For example, optimal foraging theory predicts that individuals seek to maximise fitness by adopting a strategy that maximises energy acquisition, while minimising the costs associated with foraging (e.g., energy acquisition vs. search times, handling of prey, Schoener et al. 1971; Pyke 1984). In contrast, the landscape of fear model predicts that predation risk motivates patch choice and behaviour, for instance by influencing foraging decisions for prey that must select patches across a gradient from risky but profitable, to safe but less profitable

(Brown et al. 1999, Laundré et al. 2010). While these theories have been useful in predicting and understanding spatial distributions within a habitat, they have rarely been applied among habitats, especially in the marine environment.

Primarily, multiple habitat use is thought to be driven by an inability to obtain the full complement of resources required by an individual throughout their life cycle from within a single habitat type (Werner & Gilliam 1984; Dunning et al. 1992; Pope et al. 2000). In this scenario, access to different components of a landscape or seascape may be necessary for survival, such as the need to access specific resources (e.g., shelter or food) (Holbrook et al. 1990; Mittelbach & Osenberg 1993; Dahlgren & Eggleston 2000; Hunt & Seibel 2000). For instance, access to water and suitable grazing habitat underpin the seasonal long-distance migrations of the Serengeti wildebeest (Torney et al. 2018). Atlantic salmon (*Salmo salar*) use freshwater habitats as juveniles, migrate to marine environments to feed as adults, before eventually returning to natal rivers to spawn (Klemetsen et al. 2003). As well as these examples of long-distance migrations which can take place over timescales from months to years, other taxa undertake more frequent (e.g., daily) movements among habitat types, often between feeding and refuge sites (e.g., Ogden & Buckman 1973; Ager et al. 2003; Hitt et al. 2011). Through their cross-habitat movements, multiple habitat users can be an important source of cross-habitat subsidies, for example, through pollination, nutrient transport, and by altering predator-prey interactions as they move around the landscape (Polis et al. 1997; Lundberg & Moberg 2003; Kremen et al. 2007; Heck et al. 2008). Accounting for the range of habitat types found in a landscape and their spatial arrangement can therefore be important for understanding both species distribution patterns and ecological processes.

Recognising that habitats, and therefore species-habitat relationships, rarely function in isolation, has led to a shift away from traditional and reductionist approaches that typically consider a single habitat or patch to a broader landscape perspective that considers the composition and spatial arrangement of habitat patches (Massol et al. 2011). In terrestrial ecosystems, the development of the field of landscape ecology has been gaining momentum since the 1980s (Addicott et al. 1987; Turner 1989; Taylor et al. 1993; Wiens et al. 1993; Wu & Loucks 1995; Turner 2005). Landscape ecology is based around the central tenet that the spatial context of a habitat patch is of ecological importance (Turner 2005), and commonly uses high resolution habitat, vegetation, or land-use maps to relate patterns in landscape structure to biotic patterns and processes. Landscape ecology has provided a useful framework

to evaluate the influence of landscape heterogeneity on biodiversity, species distribution patterns and community structure (e.g., Wiens 1976; Dormann et al. 2007; Tschardt et al. 2012), to understand the effects of disturbance, habitat loss and fragmentation (e.g., Saunders et al. 1991; Andr n 1994; Ricketts 2001; Fahrig 2003; Fischer & Lindenmayer 2007; Turner 2010), and to inform terrestrial conservation planning (e.g., Phua & Minowa 2005; Pressey et al. 2007; Wiens 2007; McAlpine et al. 2015).

In addition, the field of landscape ecology has also been central in demonstrating the value of considering the spatial scale across which species-habitat relationships are examined (Wiens 1989; Levin 1992; Holland et al. 2004; Wheatley & Johnson 2009). For instance, the spatial scale(s) across which a species perceives and interacts with its environment can vary depending on traits such as body size, mobility, and foraging behaviour (Holland et al. 2004; Miguet et al. 2016). Adopting a single spatial scale to study patterns and processes based on human-imposed concepts of habitat or scale, risks missing the true scale of effect (Orians & Wittenberger 1991; Jackson & Fahrig 2015). Instead, landscape ecology promotes studying species-habitat relationships across multiple spatial scales simultaneously (e.g., from mm's to km's) to determine which scale, or combination of factors operating at different scales, best explain ecological patterns and processes (Wedding et al. 2011; Miguet et al. 2016).

Despite the challenges and complexity associated with disentangling species-environment relationships, the need to collect, interpret and synthesise data on such relationships has never been greater. Globally, we are witnessing transformations of both terrestrial and aquatic environments at unprecedented rates due to anthropogenic-induced disturbances (e.g., climate change, development, pollution, overharvesting, Hughes 2000; Jackson et al. 2001; Brooks et al. 2002; Williams et al. 2003; Hoekstra et al. 2005). One of the most visible and quantifiable impacts of human activity can be seen through alterations to the physical structure and composition of habitats (e.g., land-use change, habitat loss, habitat fragmentation, shifts in foundation species, and biotic homogenisation), which can concomitantly affect the population dynamics of fauna that interact with those habitats (McKinney & Lockwood 1999; Cushman 2006; Bonin et al. 2011; Powers & Jetz 2019). For species that are highly dependent on a single habitat type, the effects of habitat change on populations can be rapid (e.g., Warren et al. 2001), and future outcomes for such taxa can be predicted with reasonable confidence. However, for taxa that use multiple habitat types throughout their lives (Pulliam et al. 1992; Schreiber &

Rudolf 2008; Jackson et al. 2019), we know far less about habitat dependencies for these species or how they might be impacted as different habitat types undergo change.

In marine systems, the conventional approach to studying species-habitat relationships has been to focus on a single habitat type in isolation to identify the within-habitat characteristics that best explain biodiversity, species distribution or abundance patterns (e.g., Jones et al. 2004; Graham & Nash 2013). Although this approach may be suitable for those species that complete their entire life cycle within that habitat type (i.e., habitat specialists), it is less suitable for understanding the distribution and abundance patterns of species that use multiple habitats, nor does it consider how external influences operating elsewhere in the seascape might influence biotic and abiotic conditions within a focal habitat patch. Instead, an approach that accounts for the spatial arrangement and composition of habitats and habitat patches within a seascape may provide a more complete picture, and enable us to make more meaningful predictions about future environmental change (Boström et al. 2011; Pittman 2018).

Emerging technologies are now enabling the principles of landscape ecology to be tested in marine systems (Pittman et al. 2004; Boström et al. 2011; Wedding et al. 2011). To date, a seascape ecology approach has been applied to examine how the seascape structure influences the distribution and abundance patterns of a range of taxa (e.g., invertebrates: Turner et al. 1999; Pittman et al. 2004; Carroll & Peterson 2013; fishes: Moore et al. 2011; Berkström et al. 2013a; Ricart et al. 2018), in a variety of habitat types (e.g., coral reefs, Wedding & Friedlander 2008; Kendall et al. 2011; kelp forests, Parnell 2015; Sievers et al. 2016; macroalgae beds, van Lier et al. 2018; Eggertsen et al. 2019; mangrove forests, Pittman et al. 2007; Drew & Eggleston 2008; rocky reefs, Swadling et al. 2019; saltmarshes, Green et al. 2012; and seagrass, Staveley et al. 2017; Olson et al. 2019) and across a range of temperate (e.g., Perry et al. 2018; Rees et al. 2018), subtropical (e.g., Olds et al. 2012a; Gilby et al. 2016) and tropical (e.g., Grober-Dunsmore et al. 2008; Olds et al. 2013; Berkström et al. 2020) ecosystems. Collectively these studies have shown that the seascape surrounding a focal patch can influence species distribution and abundance patterns, although the influence and relative importance of seascape versus within-habitat attributes (e.g., structural complexity) varies among taxa, and between geographic locations. In addition, similar to studies from terrestrial systems (e.g., Thornton et al. 2011), the few seascape studies that have incorporated a range of spatial scales to measure different aspects of the surrounding seascape (e.g., area of habitat within different sized radii)

have found that that the spatial scales considered can influence outcomes (e.g., Kendall 2005; Grober-Dunsmore et al. 2008).

There is increasing recognition that a seascape ecology approach may be particularly beneficial for examining patterns and processes in tropical shallow-water seascapes. Tropical shallow-water seascapes often contain a complex mosaic of different habitat types (e.g., coral and rocky reefs, seagrass meadows, mangroves), in a multitude of spatial configurations, habitat combinations and patch sizes, and these habitats can be interlinked through abiotic and biotic processes (Ogden 1988; Boström et al. 2011; Pittman & Olds 2015). Many of these habitat types are vulnerable to terrestrial influences (e.g., human development, pollution, riverine outflows), and are among the habitats most threatened by climate change (Jackson et al. 2001; Hughes et al. 2003; Orth et al. 2006; Barlow et al. 2018; Halpern et al. 2019). In many locations, widespread changes have already occurred including habitat loss, fragmentation, and degradation (e.g., coral reefs, Hughes et al. 1994; De'ath et al. 2012; mangroves, Richards & Friess 2016; Goldberg et al. 2020; seagrass, Waycott et al. 2009; Rasheed et al. 2014). Alterations to, or the loss of, the foundation species that are responsible for creating the physical habitat structure can directly or indirectly affect species distribution patterns and ecosystem processes (Ellison et al. 2005; Byrnes et al. 2011; Pratchett et al. 2011; Richardson et al. 2017). However, without a more complete understanding about how the spatial arrangement and composition of tropical shallow-water seascapes can influence patterns or processes, our ability to predict future change is limited.

Tropical fishes are an important group to consider within a seascape context. Some species use completely different habitats at different life stages, or may use multiple habitat types on a regular basis for foraging, spawning or refuge (Pittman & McAlpine 2003; Nagelkerken et al. 2015). By using the wider seascape, fish can act as mobile links through nutrient transport, and can affect food web dynamics between different habitat types and habitat patches (Meyer & Shultz 1985; Parrish 1989; Polis et al. 1997; Moberg & Folke 1999; Lundberg & Moberg 2003; Sheaves 2009). Cumulatively, these multi-habitat users could be significant in the transfer of energy, and maintenance of key functions across a range of habitats. However, the prevalence of multiple habitat use among tropical fishes is largely unknown. Furthermore, although several studies have begun to relate tropical fish distribution patterns to spatial patterns of the seascape, this research has typically been restricted to a few geographic locations (e.g., Caribbean, Kendall 2005; Grober-Dunsmore et al. 2007; Pittman et al. 2007; Sekund & Pittman 2017;

Western Indian Ocean, Dorenbosch et al. 2006a; Berkström et al. 2012; Aller et al. 2014). Making generalisations about fish-habitat relationships based on a limited number of localities and without accounting for the spatial context could mean we are making poor decisions about how best to manage individual habitats or seascapes. In addition, few studies have expanded seascape research beyond ecological patterns to examine how the surrounding seascape can influence ecological processes (e.g., herbivory).

Knowledge of both fish-habitat relationships and how the spatial arrangement of habitat types and patches can influence ecological patterns and processes are prerequisites for the effective management of tropical shallow-water seascapes. Without such information, we cannot fully evaluate the effects of habitat loss, degradation, and reconfiguration under different disturbance scenarios, design ecologically meaningful reserve networks, identify and protect critical nursery habitats, or implement appropriate intervention or restoration strategies.

### ***Research aims and thesis structure***

The overarching aim of this thesis was to investigate the influence of seascape structure on the composition and function of tropical fish communities, primarily those considered as coral reef-associated fishes. This aim is achieved through four distinct objectives, each of which relates to a separate chapter.

The first two objectives relate to examining fish distribution patterns in multiple habitats found in tropical shallow-water seascapes, specifically to:

- 1) Synthesise the extent of non-reef habitat use by coral reef-associated fishes at a global scale
- 2) Describe fish-habitat relationships in a tropical shallow-water seascape in Papua New Guinea in the Indo-Pacific.

The second two objectives consider how the spatial configuration and composition of the seascape, and different spatial scales, can influence both fish distribution patterns and ecological processes, specifically to:

- 3) Investigate how within-patch and seascape variables influence fish distribution patterns in tropical seagrass beds
- 4) Examine how seascape and within-reef benthic variables influence rates and agents of macroalgal removal on coral reefs.

**Chapter 2** presents the first global systematic review to quantify the extent of non-reef habitat use among fish species typically recognised as reef-associated and identifies key knowledge gaps. These knowledge gaps inform the direction and focus of the subsequent field-based chapters conducted in the shallow water tropical seascape of the Tigak Islands, New Ireland, Papua New Guinea. **Chapter 3** describes the habitat use patterns of coral reef fish taxa in Papua New Guinea, a relatively unexplored region of the Indo-Pacific. Using remote video cameras, I quantify the occurrence of reef fish taxa across five reef and non-reef habitat types. **Chapter 4** examines fish-habitat relationships across a hierarchy of spatial scales. Specifically, I determine the relative importance of within-patch versus seascape variables as drivers of fish distribution and abundance patterns in tropical seagrass beds, and evaluate the influence of seascape structure at multiple spatial scales. In the final data chapter (**Chapter 5**), I progress from exploring patterns to examining a critical ecological process, macroalgal browsing, within a seascape context. Here, I examine the influence of seascape and within-reef benthic variables on macroalgal removal rates and agents of browsing on coral reefs. Finally the General Discussion (**Chapter 6**) considers how the findings of this thesis can direct future efforts to understand the role of the seascape in driving species distribution patterns and ecological processes.

## Chapter 2

# Beyond the reef: the widespread use of non-reef habitats by coral reef fishes <sup>1</sup>

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### 2.1 Abstract

Marine ecology seeks to understand the factors that shape biological communities. Progress towards this goal has been hampered by habitat-centric approaches that ignore the influence of the wider seascape. Coral reef fishes may use non-reef habitats (e.g., mangrove, seagrass) extensively, yet most studies have focused on within-reef attributes or connectivity between reefs to explain trends in their distribution and abundance. We systematically review the evidence for multi-habitat use by coral reef fishes across life stages, feeding guilds, and conservation status. At least 670 species of ‘coral reef fish’ have been observed in non-reef habitats, with almost half (293 species) being recorded in two or more non-reef habitats. Of the 170 fish species for which both adult and juvenile data were available, almost 76% were recorded in non-reef habitats in both life stages. Importantly, over half of the coral reef fish species recorded in non-reef habitats (397 spp.) were potential fisheries targets. The use of non-reef habitats by ‘coral reef’ fishes appears to be widespread, suggesting in turn that attempts to manage anthropogenic impacts on fisheries and coral reefs may need to consider broader scales and different forms of connectivity than traditional approaches recommend. Faced with the deteriorating condition of many coastal habitats, there is a pressing need to better understand how the wider seascape can influence reef fish populations, community dynamics, food-webs and other key ecological processes on reefs.

### 2.2 Introduction

The problem of pattern and scale has been widely acknowledged in ecology for decades (Levin 1992) but progress in unravelling the causes and consequences of spatial variance and their relevance for many ecological communities has been slow. Marine environments have been particularly challenging to study (Pittman & Olds 2015), because of the depth, inaccessibility,

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and three-dimensional complexity of the ocean; high-resolution, large-extent maps and biological surveys of underwater systems are difficult to obtain (Sbrocco & Barber 2013). Whilst the development of seascape ecology has lagged behind that of landscape ecology because of these issues, the situation has improved over the last two decades. The growing availability of broad-scale datasets (e.g., Sbrocco & Barber 2013) and remote sensing technologies now allow us to apply landscape ecology principles (Pittman & Olds 2015; Grober-Dunsmore et al. 2009) to address questions about the influence of spatial variation in structuring marine communities.

Coastal seascapes are spatially heterogeneous environments (Parrish 1989; Nagelkerken 2009). In the tropics, shallow-water seascapes can include a range of coral reef, mangrove, seagrass, macroalgae and soft sediment habitats. Although these habitats often occur in close proximity to each other (Ogden 1976; Parrish 1989; Grober-Dunsmore et al. 2008), the traditional approach to understanding species distribution and abundance patterns in coastal seascapes has often been confined to observations within a single habitat type (e.g., coral reef, seagrass or mangroves) (Birkeland 1985; Beets et al. 2003; Nagelkerken 2009; Pittman & Olds 2015). Yet it is increasingly acknowledged that habitats rarely function in isolation and that interactions between habitats may be important for productivity and ecosystem functioning (Randall 1965; Ogden 1976; Birkeland 1985; Mumby & Hastings 2008). Marine habitats are linked through the transfer or movement of nutrients, materials and organisms (Nyström & Folke 2001; Lundberg & Moberg 2003; Hyndes et al. 2014; Shantz et al. 2015). As a result, ecological patterns and processes observed within one habitat may, in part, be driven by patterns and processes occurring in another habitat.

Fish are recognised as important mobile links between marine habitats (Nyström & Folke 2001; Lundberg & Moberg 2003; Berkström et al. 2012). For example, the movement of fishes can facilitate trophic subsidies and nutrient transfer between habitats (Meyer & Schultz 1985; Polis et al. 1997; Heck et al. 2008; Shantz et al. 2015). In tropical seascapes, some fishes settle directly to coral reef habitat following their pelagic larval stage and become permanent residents, while others use non-reef habitats (e.g., seagrass, mangrove) throughout their lives. One of the most frequently cited explanations for the use of non-reef habitats by coral reef fishes is that non-reef habitats, such as seagrass beds and mangroves, can be used as nursery habitat for juvenile coral reef fishes (Beck et al. 2001; Nagelkerken et al. 2002; Dahlgren et al. 2006; Nagelkerken 2009; Sheaves et al. 2015). In addition, some coral reef fishes undertake

tidal, diel, monthly, seasonal, and/or yearly movements between coral reef and non-reef habitats to forage, shelter from predators, and/or spawn (Unsworth et al. 2007a; Appeldoorn et al. 2009; Hitt et al. 2011; Taylor & Mills 2013; Honda et al. 2016). Availability of, or access to, non-reef habitats may therefore be important for many coral reef fish species.

Over the last two decades, research into the use of non-reef habitats by fishes in tropical seascapes has expanded considerably, largely because many of the species that use multiple habitat types are ecologically or economically important (see reviews by Adams et al. 2006; Nagelkerken et al. 2008; Appeldoorn et al. 2009; Igulu et al. 2014). However, this growing body of research has not yet been synthesised to determine the number of coral reef fish species that have been observed in non-reef habitats, nor which non-reef habitats are commonly used. Fuelled by growing concerns about global habitat loss and increasing recognition that the wider seascape may influence species distribution and abundance patterns, such a review is both timely and important. Here, a systematic review of the occurrence of coral reef fishes in non-reef habitats is conducted. The findings are separated into two biogeographic regions, the tropical Atlantic and the Indo-Pacific, because they have distinct reef fish assemblages and benthic communities. Specifically, the following themes were examined: 1) how many coral reef fish species have been observed in non-reef habitats; 2) which habitats are used by coral reef fishes in addition to reefs; 3) differences and similarities between the two biogeographic regions; whether habitat use differs depending on 4) life stage or 5) feeding ecology; 6) which multi-habitat fish species are of interest to fisheries; and 7) whether any multi-habitat coral reef fish species are currently endangered, threatened or vulnerable.

### **2.3 Methods**

The literature was systematically searched for original research articles published between 1967 and November 2018 using ISI Web of Science and Scopus. To identify publications that had investigated the use of non-reef habitats by fishes that are generally considered coral reef-associated, the search terms “coral reef\*” and “fish\*” were used in combination with a range of different habitat types namely “seagrass\*”, “grass\*”, “mangrove\*”, “macroalga\*”, “alga\*”, “vegetat\*”, “unvegetat\*”, “mud\*”, “rock\*”, “estuar\*”, “sand\*”, and “soft sediment\*”. Papers were searched for that specifically included these terms in the Title, Keywords and/or Abstract. Outputs from these searches resulted in 2,992 potential papers from the ISI Web of Science database, and 1,419 papers from Scopus. To be included in subsequent analyses, the following selection criteria were applied to each paper: 1) data were collected from a coral reef and also

in at least one other habitat type; 2) non-reef habitats were clearly defined as discrete patches within the study region, rather than as microhabitats within a coral reef; and 3) the paper or supplementary material contained species-specific information. Citations within all papers matching the selection criteria were also examined to identify any additional literature not captured within the literature search. Although many studies will have recorded the presence of coral reef fishes in non-reef habitats (e.g., Unsworth et al. 2009; Chaves et al. 2013; Bradley et al. 2017), only those studies that had surveyed both reef and non-reef habitat were included because this provided unequivocal evidence of the use of both reef and non-reef habitats by a given species within the same study area. In addition, the selection criteria excluded papers that correlated presence or abundance of fishes on the reef with proximity to a non-reef habitat without surveying the non-reef habitat (e.g., Grober-Dunsmore et al. 2007; Olds et al. 2012b).

For each study, fish species were included in the analysis if they were observed both on the reef and in at least one other habitat type, and were classified as ‘reef-associated’ in *FishBase* (Froese & Pauly 2018), hereafter termed ‘coral reef fishes’. Observations of coral reef fish species were recorded as presence/absence data and separated into life history stages (i.e., adults and juveniles) if the data were available. Presence/absence rather than abundance or biomass data was recorded to capture the widest possible range of species that have been observed in non-reef habitats as well as on coral reefs. This decision was driven by the diverse range of reporting variables contained within the literature (e.g., density, presence/absence, proportional abundance, biomass), the different methodologies (e.g., tagging, visual censuses, timed surveys, fishing), as well as the variety of life stages examined by studies (e.g., larvae, early stage juveniles, juveniles, sub-adults and adults). The combination of these factors makes it challenging to present and interpret abundance or biomass data in a meaningful way. Parrotfishes (subfamily Scarinae) were recognised as distinct from the wrasses (Labridae) because of the functional importance of parrotfishes for coral reef ecosystems (e.g., Hoey & Bonaldo 2018).

Each presence/absence observation was assigned to one of seven habitat classes: coral reef, seagrass bed, mangrove, macroalgal bed, sand, estuary or other. Although the benthic composition of seagrass, macroalgae, mangrove and coral reef habitats will differ between regions and studies and could influence coral reef fish use of these habitats, this review does not distinguish between different taxonomic composition in a particular habitat type (e.g., types of seagrass) as this information was not consistently available from the literature. Mangrove

habitat refers only to non-estuarine mangroves. Estuary includes a range of habitats found within the estuary, including mangroves and rocky substrates. The habitat category “other” includes a range of additional non-reef habitats such as channels not dominated by coral communities, rocky boulders and notches, mudflats, gorgonian plains and submerged vegetation. To explore potential biogeographic differences in habitat use by reef fishes, data were separated into two distinct regions: 1) the tropical Atlantic, which includes the Caribbean, Florida Keys, and tropical reefs in Brazil; and 2) the Indo-Pacific, which includes the Indian and Pacific Oceans, as well as the Red Sea.

To examine whether feeding ecology influences the use of multiple habitat types by coral reef fishes, each species was assigned to one of six feeding guilds (i.e., carnivores, corallivores, herbivores, omnivores, piscivores and planktivores) primarily derived from the ‘Diet’ and ‘Food item’ categories in *FishBase* (Froese & Pauly 2018). Here carnivores were defined as species that predominantly consume macro-invertebrates (e.g., crustaceans, molluscs, polychaetes) and/or fish. Fish that consume both animal and plant material were classed as omnivores, and herbivores included species that feed on macroalgae, turf algae, seagrass, microscopic phototrophs and detritus. To explore whether body size might be a factor influencing which taxa were observed in non-reef habitats, and as a proxy for mobility, the maximum total length (cm) recorded for each species was also extracted from *FishBase* (Froese & Pauly 2018). Species were also evaluated based on their importance to fisheries using the five categories reported by *FishBase* (i.e., commercial, minor commercial, subsistence, recreational and, of no interest to fisheries) (Froese & Pauly 2018). Species could be assigned to more than one fisheries category (e.g., commercial and recreational). In addition, multiple habitat-using coral reef fish species were classified as endangered, near threatened or vulnerable or “other” according to the IUCN Red List (IUCN 2018).

## 2.4 Results

There were 107 studies that matched all of the selection criteria. Of these, 56 were based in the tropical Atlantic, 50 were in the Indo-Pacific, and one examined two species, one from each biogeographic region (i.e., Dorenbosch et al. 2006b). The foci of different studies ranged from fish communities (39 out of 107 studies) to specific groups or families (40 studies) down to a single focal species (28 studies). In the tropical Atlantic, 37% of studies (n=21) examined only one non-reef habitat type compared to 59% (n=30) in the Indo-Pacific. Three or more non-reef habitats were examined by 16 studies in the tropical Atlantic, in strong contrast to the Indo-

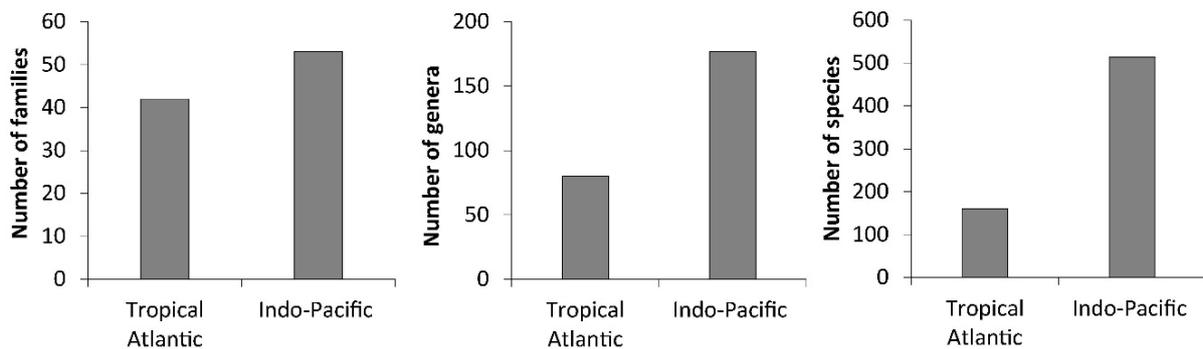
Pacific where there was only a single study (i.e., Kimirei et al. 2011) (Figure A2.1). Seagrass was the most frequently surveyed habitat in both regions, examined in 74% of the 57 studies in the tropical Atlantic, and 63% of the 51 studies in the Indo-Pacific. Mangrove habitats were studied almost twice as much in the tropical Atlantic (70% of the 57 studies) compared to the Indo-Pacific (37% of the 51 studies) (Figure A2.2). The remaining non-reef habitat types (macroalgae, soft sediment, estuary and “other”) were surveyed on far fewer occasions ( $\leq 11$  studies for each habitat) in both regions.

#### **2.4.1 Coral reef fish species and families**

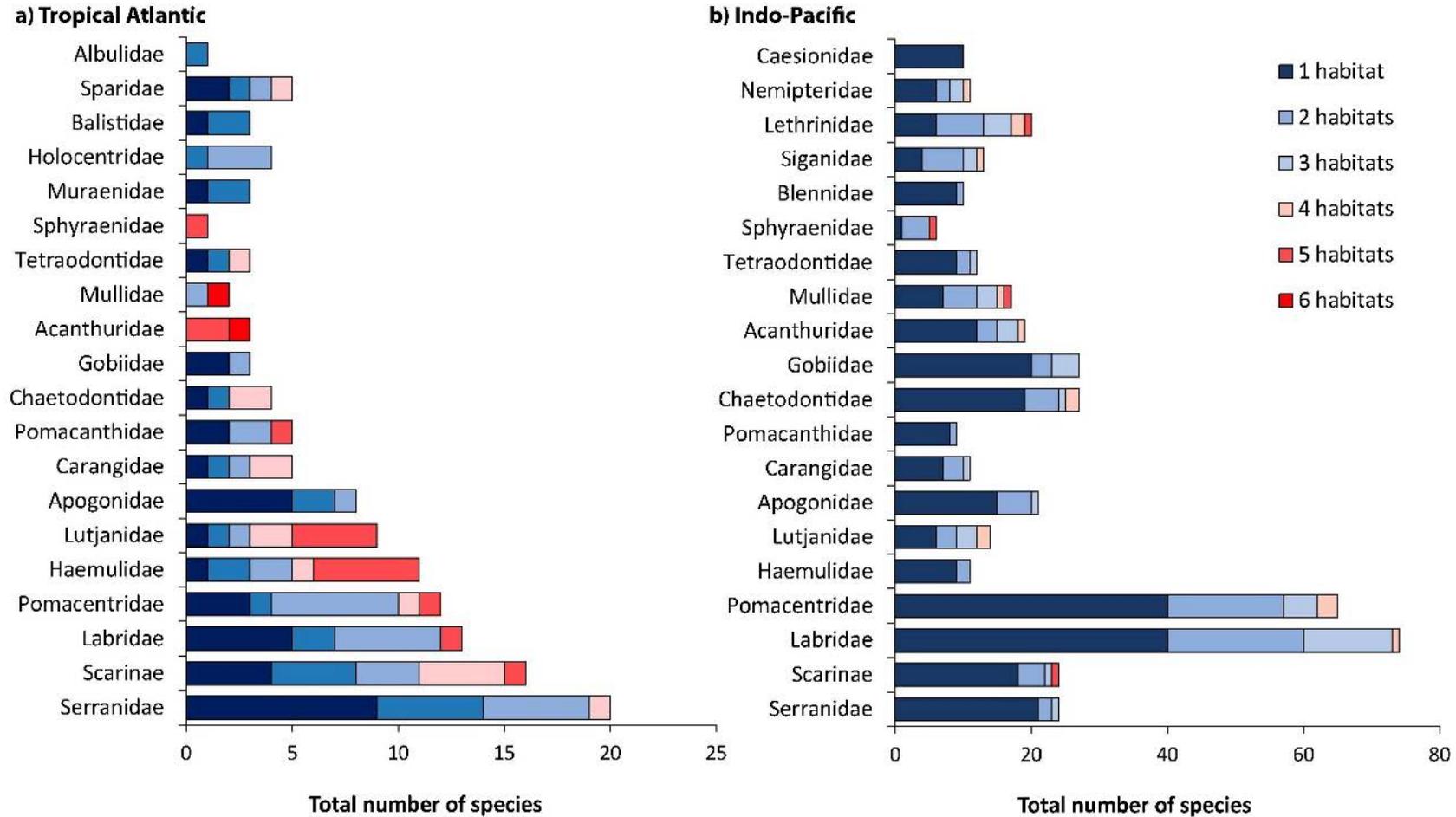
Species observed in multiple habitat types came from a diverse range of reef fish taxa and were recorded in a wide range of non-reef habitats. From these 107 studies, a total of 670 coral reef fish species were observed in non-reef habitats, 160 species from the tropical Atlantic, and 514 species from the Indo-Pacific region (Figure 2.1; Table A2.1). Four species of multi-habitat users (black triggerfish (*Melichthys niger*, Balistidae); spot-fin porcupinefish (*Diodon hystrix*, Diodontidae); red lionfish (*Pterois volitans*, Scorpaenidae) and; great barracuda (*Sphyraena barracuda*, Sphyraenidae)) were common to both regions. In the tropical Atlantic, the highest number of species observed in non-reef habitats were in the Serranidae (groupers) and Scarinae (parrotfishes), with 20 and 16 species respectively (Figure 2.2). In contrast, in the Indo-Pacific, the dominant taxa documented in non-reef habitats were the Labridae and Pomacentridae, with 74 and 65 species respectively (Figure 2.2).

Across both regions, 44% of species (293 of 670) were present in two or more non-reef habitats (Table A2.1). In the tropical Atlantic, 69% of species (111 of 160 species) were noted in two or more non-reef habitats (Figure 2.3), and of these 37% (41 of 111 species) were recorded in every habitat surveyed (e.g., if five non-reef habitats were surveyed, they were found in those five habitats). In particular, 100% of Acanthuridae species (3 out of 3) were observed in five or more non-reef habitats, 73% of Haemulidae species (8 out of 11 species) and 78% of Lutjanidae species (7 out of 9 species) were observed in three or more non-reef habitats in the tropical Atlantic. For example, French grunts (*Haemulon flavolineatum*, Haemulidae) has been recorded in seagrass, mangrove, macroalgae, soft sediment and other non-reef habitats including rocky boulders and mudflats (e.g., Nagelkerken et al. 2000a; Adams & Ebersole 2002; Nagelkerken & van der Velde 2002; Harborne et al. 2008; Burke et al. 2009).

In the Indo-Pacific, 36% of species (186 of 514) were observed in two or more non-reef habitat types (Figure 2.3). Of these, 62% (116 of 186) were recorded in every habitat where their presence or absence was evaluated. In this region, 100% of Mullidae (9 out of 9) and Nemipteridae (5 out of 5) species, 83% of Sphyrnidae (5 out of 6), 70% of Lethrinidae (14 out of 20), 69% of Siganidae (9 out of 13), and 57% of Lutjanidae species (8 out of 14) were recorded in two or more non-reef habitats. For example, the thumbprint emperor (*Lethrinus harak*, Lethrinidae) has been observed in seagrass, mangrove, macroalgae, soft sediment, mudflat and rocky habitats (e.g., Thollot et al. 1990; Ashworth et al. 2006; Barnes et al. 2012; Berkström et al. 2012; Kruse et al. 2016; Wilson et al. 2017).

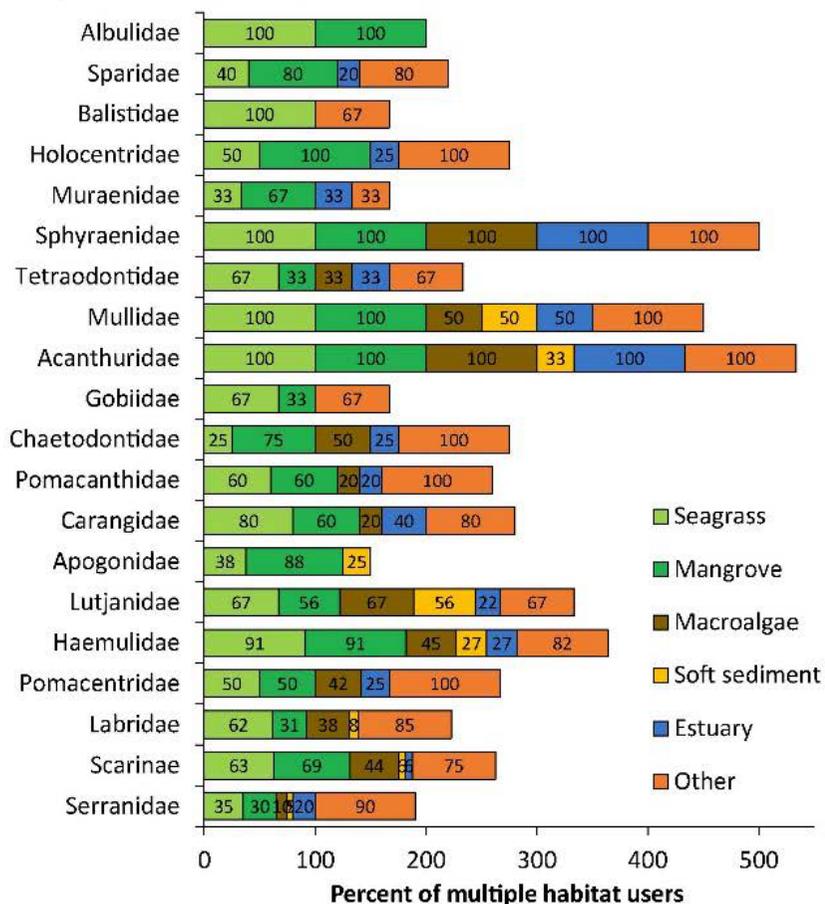


**Figure 2.1.** Number of coral reef fish a) families, b) genera and, c) species observed using both reef and non-reef habitats in the tropical Atlantic and Indo-Pacific regions. Data extracted from 107 publications.

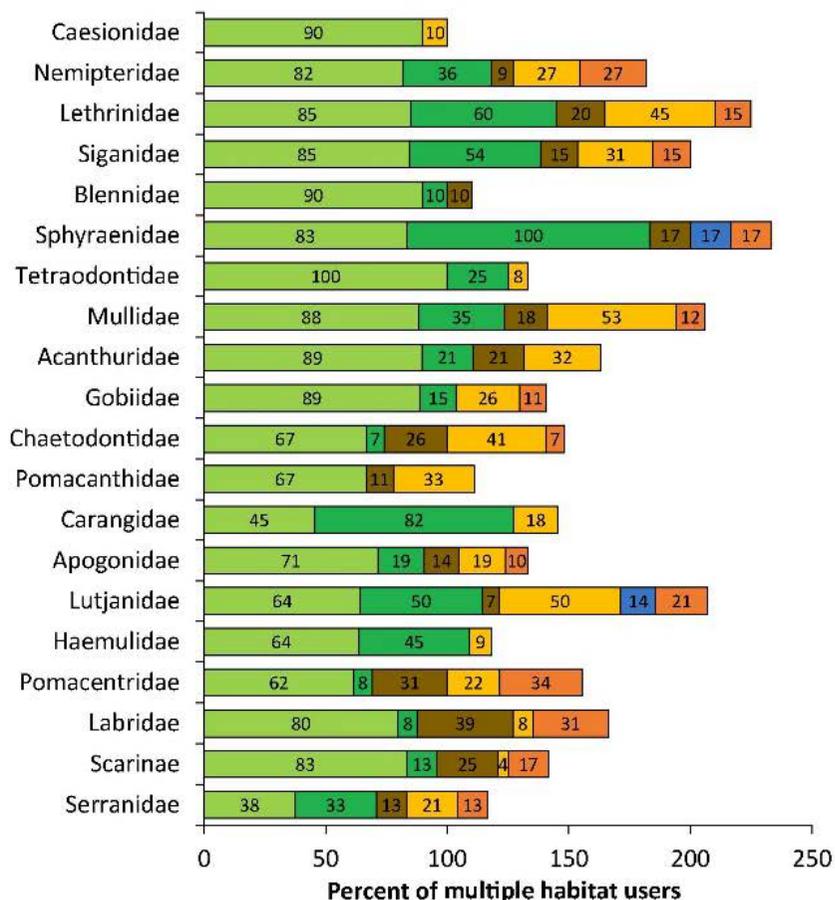


**Figure 2.2.** Prevalence of multiple habitat use among selected coral reef fish taxa in: a) the tropical Atlantic, and; b) the Indo-Pacific. The colours indicate the number of non-reef habitats (1-6) used by each species. In the tropical Atlantic, where a wide range of habitats have been surveyed, many species have been observed in four or more non-reef habitat types. In contrast, in the Indo-Pacific almost 60% of studies only surveyed coral reef fishes in one non-reef habitat.

**a) Tropical Atlantic**



**b) Indo-Pacific**



**Figure 2.3.** Habitat use among selected coral reef fish taxa in: a) the tropical Atlantic, and; b) the Indo-Pacific. Percentages represent the proportion of species within each taxa that are found within each habitat type. For example, for the Acanthuridae in the tropical Atlantic, 100% of the species identified as multiple habitat users were observed in seagrass, mangrove, macroalgae, estuary, and “other” habitat types and 33% of species were observed on soft sediments.

#### **2.4.2 Comparison of the tropical Atlantic and Indo-Pacific**

Thirty-two fish families were common between the two biogeographic regions. Here four fish taxa of economic or ecological importance were selected to compare and contrast non-reef habitat use trends. Primarily observations were compared between mangrove and seagrass habitats because these habitat types have been most consistently surveyed across the two regions. Taxa with high numbers of species observed in multiple non-reef habitats in both the tropical Atlantic (TA) and Indo-Pacific (IP) include the Haemulidae (11 species in each region), Lutjanidae (9 species in TA, 14 in IP), Scarinae (16 in TA, 24 in IP) and Serranidae (20 in TA, 24 in IP). A similar trend in seagrass and mangrove observations emerged for three taxa (i.e., Haemulidae, Lutjanidae and Scarinae). In the tropical Atlantic, the majority of species from these groups were recorded in both seagrass and mangrove habitats, with fewer species only recorded in one habitat type. However, this pattern differed in the Indo-Pacific, where a greater proportion of species from these three groups were recorded in either mangrove or seagrass habitat types. For example, in the tropical Atlantic, of the 11 Haemulidae species for which data were available, nine species were recorded in both seagrass and mangrove habitat. In contrast in the Indo-Pacific, only one species, the two-striped sweetlips (*Plectorhinchus albovittatus*, Haemulidae), was recorded in both habitat types, five were recorded only in seagrass, and three only in mangroves. Two of the 11 species in the Indo-Pacific were not surveyed in both habitat types. Species from the Scarinae sub-family in the Indo-Pacific were almost sevenfold higher in seagrass habitats compared to mangroves (20 species vs. 3), whereas in the tropical Atlantic, a similar number of species was observed in seagrass and mangrove habitats (10 and 11 species respectively). Species from the Serranidae showed a different pattern to the other three taxa, but shared commonalities between both biogeographic regions. Fewer species of serranids were recorded in both mangrove and seagrass habitat, but comparable numbers of species were split between the two habitat types.

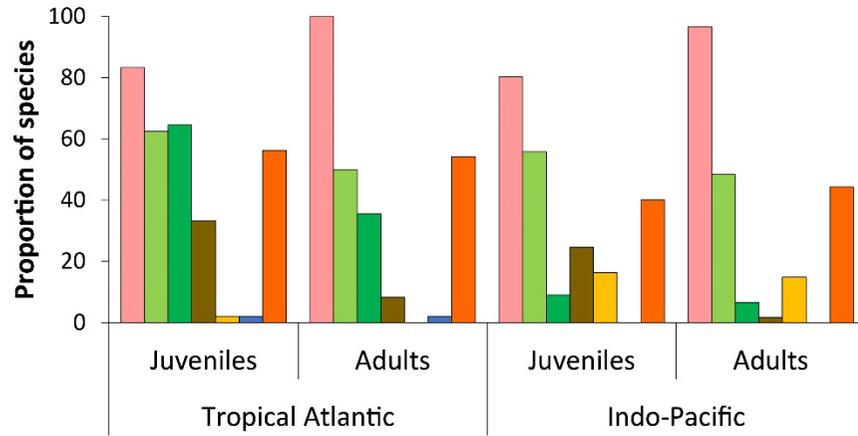
#### **2.4.3 Adult and juvenile use of non-reef habitats**

Of the 107 studies that met the selection criteria, 29 contained information on the habitat use of both adult and juvenile life stages. From these 29 studies, 14 studies covered species in the tropical Atlantic and 16 studies the Indo-Pacific. Data were available for 170 species, 48 species from the tropical Atlantic and 122 from the Indo-Pacific. Both life stages showed a diverse range of habitat use in addition to coral reefs (Figure 2.4), with almost 75% of the 170 species observed as both adults and juveniles in non-reef habitats. In the tropical Atlantic,

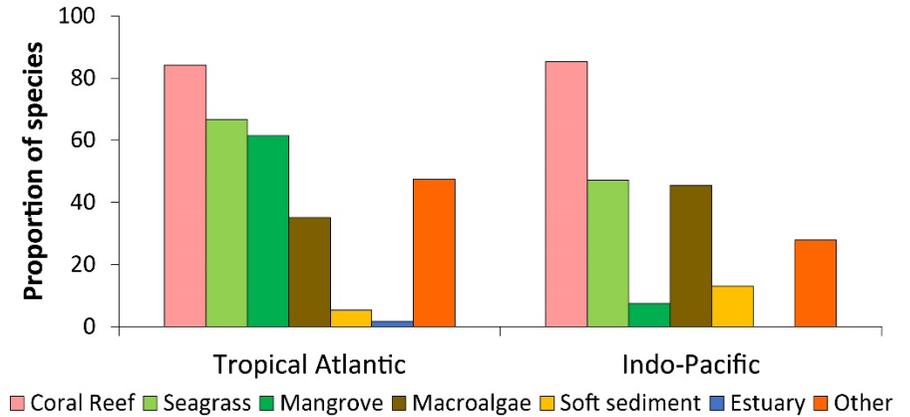
observations of juvenile reef fish species away from the reef were most common in seagrass (62.5%) and mangrove (64.6%) habitats (30 and 31 species, respectively), followed by macroalgae (33.3%, 16 species). In contrast, in the Indo-Pacific, juveniles were most commonly observed in seagrass beds (55.7%, 68 species), followed by “other” non-reef habitat types (40.2%, 49 species), macroalgal beds (24.6%, 30 species), and soft sediment (16.4%, 20 species) (Figure 2.4). For adults, the trends were slightly different. In the tropical Atlantic, adults from 26 species were recorded in “other” non-reef habitat types (54.2%), followed by seagrass (50%, 24 species), and then mangroves (35.4%, 17 species). In the Indo-Pacific, adults followed a similar pattern to that of juveniles with adults from 59 species observed in seagrass beds (48.4%), 54 species in “other” habitats (44.3%), and 18 species in soft sediments (14.8%).

To illustrate the diverse range of habitats used by adults and juveniles, this review focused on observations from six ecologically or economically important taxa, specifically Lutjanidae from both biogeographic regions, the Scarinae and Haemulidae from the tropical Atlantic, and Lethrinidae and Siganidae from the Indo-Pacific (Figure 2.5, Table A2.1). With the exception of the Lethrinidae, juveniles were consistently recorded in a wider range of non-reef habitats than adults from these species. 100% of lutjanids in the tropical Atlantic were observed as juveniles in mangroves, compared to 50% in the Indo Pacific, and 71.4% of juveniles were observed in seagrass habitats compared to 25% in the Indo-Pacific (Figure 2.5). In the tropical Atlantic, all six haemulid species for which adult and juvenile data were available have been recorded as juveniles in mangrove and seagrass habitats, compared to adults from only three species (Figure 2.5). Data were available for eight species of parrotfishes (Scarinae) in the tropical Atlantic (Figure 2.5). Juveniles from these species were more commonly recorded in mangrove and seagrass habitats than adults. In the Indo-Pacific, adults and juveniles from species in the Lethrinidae family were recorded in five non-reef habitats (i.e., seagrass, mangrove, macroalgae, soft sediment and “other”; Figure 2.5). Seagrass was the dominant habitat in which both adults and juveniles from these species were observed (i.e., juveniles from 7 species, adults from 4 species). Adult and juvenile data were available for seven Siganidae species; juveniles of these species have been recorded in five non-reef habitat types, compared to only two for adults (Figure 2.5). Similar to lethrinids, adults and juveniles from the Siganidae were most commonly observed in seagrass beds (i.e., 3 and 5 species respectively).

a) Studies that examined both adult and juvenile life stages



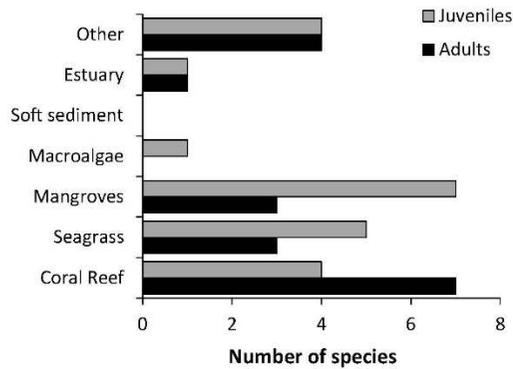
b) All studies that included juvenile life stage



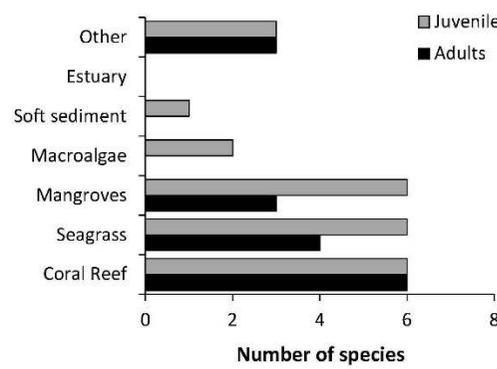
**Figure 2.4.** Proportion of species using coral reef and non-reef habitats by life stage in the tropical Atlantic and Indo-Pacific. (a) Proportion of adult and juvenile coral reef fish species recorded in coral reef and non-reef habitats. Based on data on 170 species from 29 studies that examined both adult and juvenile habitat use. In the tropical Atlantic, data were available for 48 species, and in the Indo-Pacific, data were available for 122 species; b) proportion of juveniles from 233 coral reef fish species observed in reef and non-reef habitats by biogeographic region. Based on the 29 studies that examined adult and juvenile habitat use and an addition 15 studies that examined juvenile habitat use only. In the tropical Atlantic, data were available for juveniles of 57 species, and in the Indo-Pacific, data were available for juveniles from 176 species.

### Tropical Atlantic

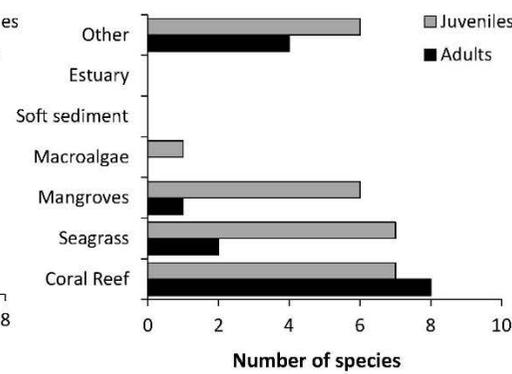
a) Lutjanidae (7 spp.)



b) Haemulidae (6 spp.)

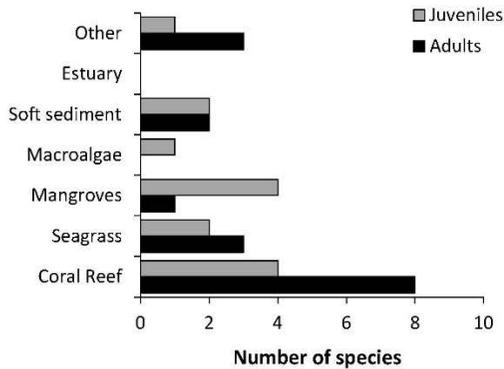


c) Scarinae (8 spp.)

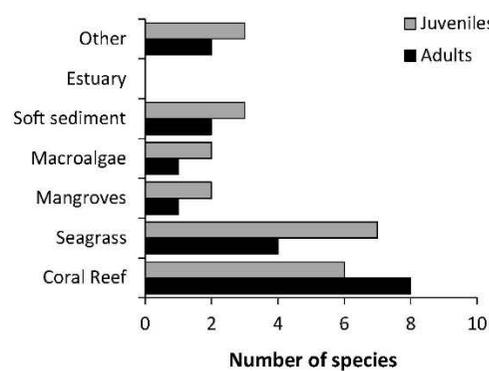


### Indo-Pacific

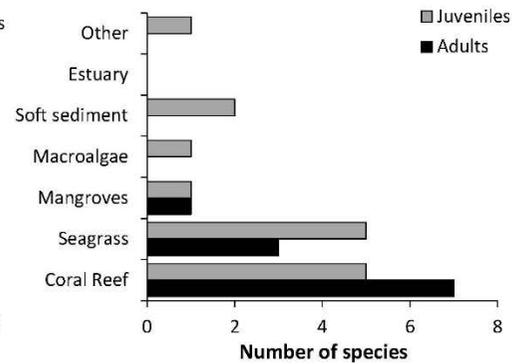
d) Lutjanidae (8 spp.)



e) Lethrinidae (8 spp.)



f) Siganidae (7 spp.)



**Figure 2.5.** Observations of adult and juvenile coral reef fish species on coral reefs and in non-reef habitat for six ecologically and economically important groups in the tropical Atlantic: a) Lutjanidae, b) Haemulidae, c) Scarinae, and; in the Indo-Pacific: d) Lutjanidae, e) Lethrinidae, and f) Siganidae. Based on data from 29 studies that examined both adult and juvenile habitat use. Note that habitats have not been surveyed equitably so absences may be an artefact of sampling.

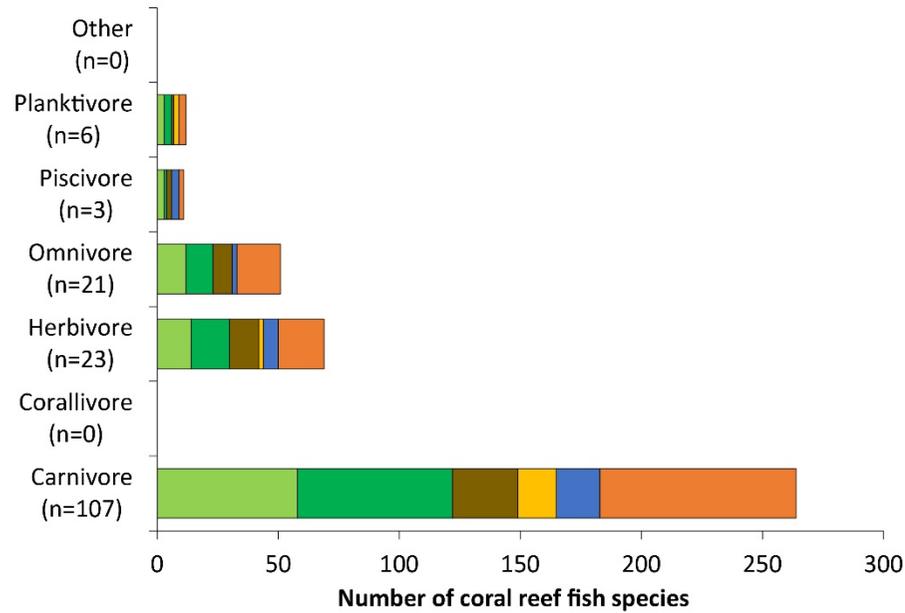
In addition to the 29 studies that collected data on both adult and juvenile reef fishes in non-reef habitats, a further 15 studies documented juveniles only. Two of these focused solely on macroalgal beds as well as coral reef (i.e., Wilson et al. 2010; Evans et al. 2014). When these 44 studies were combined, data were available on juveniles of 233 species, 57 species in the tropical Atlantic and 176 species in the Indo-Pacific (Figure 2.4). Habitat use was diverse; in the tropical Atlantic, juveniles from 66.7% of species were recorded in seagrass, 61.4% in mangroves, 47.4% in “other” non-reef habitat types, and 35.1% in macroalgal beds. In the Indo-Pacific, juveniles from 47.2% of species were recorded in seagrass, 45.5% in macroalgal beds, 27.8% in “other” non-reef habitat types, and 7.4% in mangroves.

#### **2.4.4 Feeding guilds, body size and non-reef habitats**

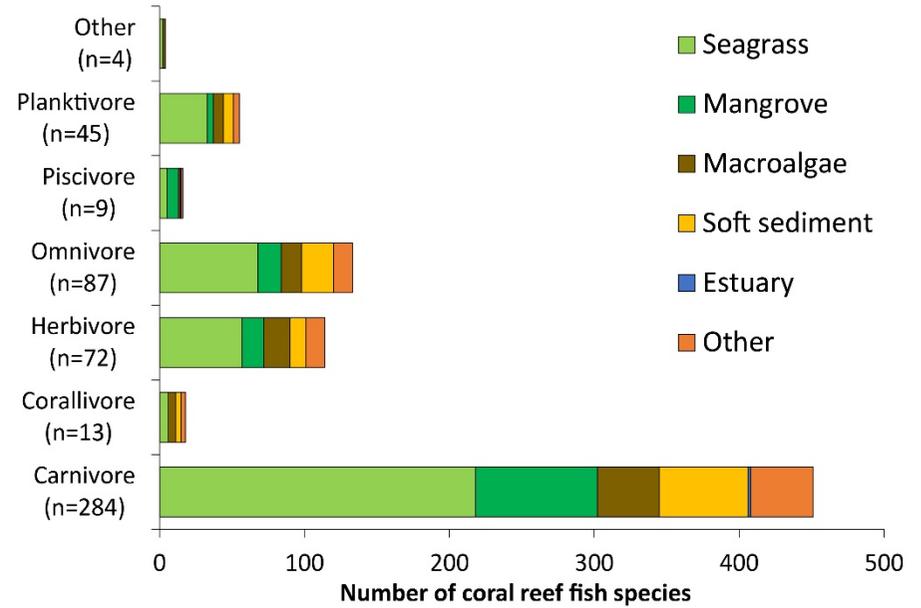
There were similar patterns among regions by feeding guilds, with carnivorous species accounting for over half of all multiple habitat users in both regions (Figure 2.6). In the tropical Atlantic, the highest number of carnivorous species were recorded in “other” habitat types (76% of carnivores), followed by mangroves (60%) and seagrass (54%) (Figure 2.6). In contrast, in the Indo-Pacific, the highest number of carnivorous species was observed in seagrass beds (77%), followed by mangroves (30%), macroalgal beds and “other” habitat types (both 15% of carnivorous species) (Figure 2.6). In the tropical Atlantic, herbivores were relatively evenly distributed between seagrass, mangroves, macroalgae and “other” habitat types. In contrast in the Indo-Pacific, the highest number of herbivores was recorded in seagrass habitats (57 spp.), three- to four-fold higher than macroalgal beds (18 spp.) and mangrove habitats (15 spp.). A similar pattern for each region was observed for omnivores. The number of piscivorous species was low in both regions, but they were recorded in all habitat types except soft sediment. Planktivores were also recorded in all habitat types in the tropical Atlantic and Indo-Pacific except estuarine environments.

Species were recorded across the whole spectrum of body sizes in non-reef habitats, including many smaller-bodied species (Figure A2.4). The modal body size class, based on total length, was 10-19.9 cm. Over a third of the species (36%, 244 spp.) recorded in non-reef habitats have a maximum total length of <20 cm, 38% (257 spp.) can attain a maximum total length of 20-49.9 cm, 16% (110 spp.) can reach 50-99.9 cm, and 8.8% (59 spp.) of species have a maximum total length >100 cm.

**a) Tropical Atlantic**



**b) Indo-Pacific**



**Figure 2.6.** Feeding guilds of coral reef fishes and the non-reef habitats that they have been observed using for the: a) tropical Atlantic, and; b) Indo-Pacific. Feeding guilds assigned from *FishBase* (Froese & Pauly 2018).

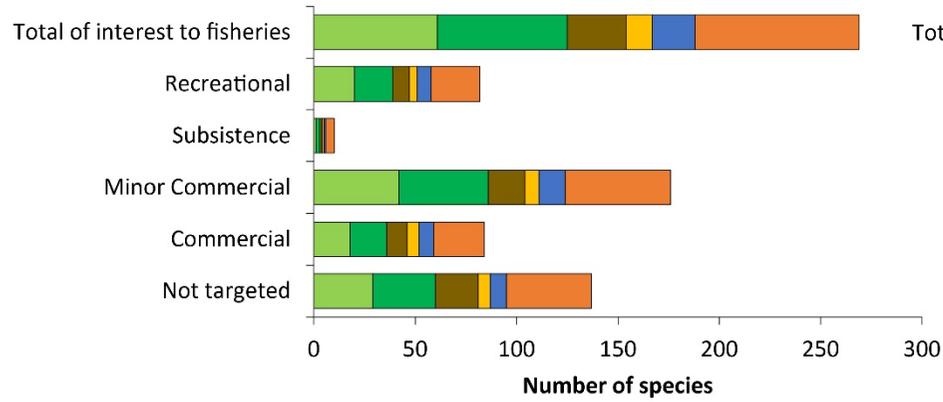
#### 2.4.5 Multi-habitat use by fisheries targets

Across both biogeographic regions, 59% of the coral reef fish species (397 of 670 species) observed in non-reef habitats were potential fisheries targets (e.g., commercial, minor commercial, recreational and/or subsistence) (Figure 2.7). In each region, >90% of the multiple habitat users of interest to fisheries were categorised as being of commercial or minor commercial importance. In the Tropical Atlantic, 31% of fisheries species (30 of 96) were targets for recreational fisheries compared to 20% (66 of 305) in the Indo-Pacific. A limited number of species (>5%) were identified as important for subsistence fisheries in both regions. Patterns in the habitats where fisheries targets were recorded differed between the tropical Atlantic and Indo-Pacific regions, particularly for vegetated habitats. In the tropical Atlantic, fisheries species were evenly observed between mangroves (64 spp.), and seagrass beds (61 spp.), twofold higher than the 29 species observed in macroalgae beds (Figure 2.7). In contrast, in the Indo-Pacific, more than double the number of species were observed in seagrass beds compared to mangroves (238 vs. 109 spp.), and almost fivefold higher than the number observed in macroalgae beds (53 spp.) (Figure 2.7). This equates to 78% of fisheries targets observed in seagrass beds compared to 36% in mangrove habitats, and 17% in macroalgae beds in the Indo-Pacific.

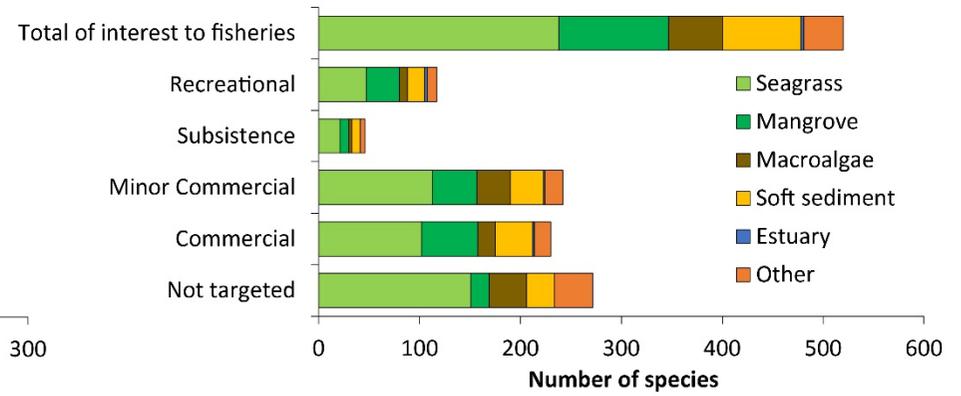
#### 2.4.6 Conservation status of multi-habitat users

Several of the multi-habitat users identified from this synthesis are classified as endangered, near threatened or vulnerable (IUCN 2018). In the tropical Atlantic, 7.5% (12 of 160) of species observed in multiple habitat types are on the IUCN Red List. One species is classified as Endangered: the Nassau grouper (*Epinephelus striatus*, Serranidae), five as Vulnerable: tarpon (*Megalops atlanticus*, Elopidae), masked goby (*Coryphopterus personatus*, Gobiidae), hogfish (*Lachnolaimus maximus*, Labridae), mutton snapper (*Lutjanus analis*, Lutjanidae) and Cubera snapper (*Lutjanus cyanopterus*, Lutjanidae), and six as Near Threatened: bonefish (*Albula vulpes*, Albulidae), queen triggerfish (*Balistes vetula*, Balistidae), rainbow parrotfish (*Scarus guacamaia*, Scarinae), red grouper (*Epinephelus morio*, Serranidae), black grouper (*Mycteroperca bonaci*, Serranidae) and yellowfin grouper (*Mycteroperca venenosa*, Serranidae). In the Indo-Pacific, 3% of species recorded in multiple habitats (16 of 514) are classified on the IUCN Red List. Two species are classed as Endangered: humphead wrasse (*Cheilinus undulatus*, Labridae) and Banggai cardinal fish (*Pterapogon kauderni*, Apogonidae); five as Vulnerable: harlequin filefish (*Oxymonacanthus longirostris*,

**a) Tropical Atlantic**



**b) Indo-Pacific**



**Figure 2.7.** Coral reef fishes observed on coral reefs and in non-reef habitats and their interest to fisheries in the: a) tropical Atlantic, and; b) Indo-Pacific regions by habitat. Data on fisheries extracted using the five fisheries classifications available in *FishBase* (Froese & Pauly 2018).

Monacanthidae), bumphead parrotfish (*Bolbometopon muricatum*, Scarinae), giant grouper (*Epinephelus lanceolatus*, Serranidae), tiger tail seahorse (*Hippocampus comes*, Syngnathidae) and spiny seahorse (*Hippocampus histrix*, Syngnathidae), and nine as Near Threatened: chevron butterflyfish (*Chaetodon trifascialis*, Chaetodontidae), bluespotted ribbontail ray (*Taeniura lymma*, Dasyatidae), blackspot tuskfish (*Choerodon schoenleinii*, Labridae), whitesaddle goatfish (*Parupeneus porphyreus*, Mullidae), orange-spotted grouper (*Epinephelus coioides*, Serranidae), brown-marbled grouper (*Epinephelus fuscoguttatus*, Serranidae), Malabar grouper (*Epinephelus malabaricus*, Serranidae), camouflage grouper (*Epinephelus polyphekadion*, Serranidae), and coral trout (*Plectropomus leopardus*, Serranidae).

## 2.5 Discussion

This review found that a significant number of coral reef fishes, from a wide range of families, life stages and feeding guilds, are present in non-reef habitats; and that over half of those species are of interest to fisheries. To place the number of species into context, the findings suggest that almost 20% of the 2,642 species present in the highly diverse Coral Triangle region (Allen 2015) are likely to occur in reef and non-reef habitats. Furthermore, the estimate of 670 species recorded across the two biogeographic regions is highly likely to be conservative. This is because studies that did not survey reef fish on coral reefs as well as non-reef habitats were excluded (e.g., Nagelkerken et al. 2000c; Bradley et al. 2017; Tano et al. 2017), and summary statistics, rather than complete species lists, were provided for some studies that met the criteria (e.g., Mumby et al. 2008). Importantly, the review also highlights that many species utilise the wider seascape throughout their lives, with 80% of species for which data were available observed as both adults and juveniles in non-reef habitats. Whilst the use of non-reef habitats by reef fishes has been known for some time (e.g., Ogden & Zieman 1977; Birkeland 1985; Ogden 1988; Adams et al. 2006; Appeldoorn et al. 2009), this review is the first to quantify the sheer number of species across a broad spectrum of reef fish taxa that inhabit a diverse array of habitat types in addition to coral reefs.

The difference between biogeographic regions provides strong evidence in support of previous research that has highlighted a potential bias in research effort into multi-habitat use by coral reef fishes (e.g., Nagelkerken 2007; Pittman & Olds 2015; Olds et al. 2016). Over half of the studies were conducted in the tropical Atlantic, yet the Indo-Pacific region covers a much larger area. In the tropical Atlantic, significant research effort has focused on the use of non-reef

habitats by coral reef fishes in the Caribbean (e.g., Nagelkerken et al. 2000b; Nagelkerken et al. 2001; Nagelkerken et al. 2002). As a result, many locations have been sampled on numerous occasions (e.g., Curaçao, Nagelkerken et al. 2000a; Nagelkerken & van der Velde 2002; Cocheret de la Morinière et al. 2002; Nagelkerken & van der Velde 2003; Pollux et al. 2007; Dorenbosch et al. 2009; Grol et al. 2014) and there is a relatively strong knowledge base for some of the species that use multiple habitats (e.g., French grunts, or schoolmaster snapper (*Lutjanus apodus*, Lutjanidae)). In contrast, few locations in the Indo-Pacific have been subjected to detailed examination. This indicates that we still have far more to learn about the use of multiple habitats by coral reef fishes in the Indo-Pacific region.

In addition to the large number of species observed away from reefs, the results highlight that coral reef fishes are using a diverse range of non-reef habitats; 44% of the 670 species were observed in multiple non-reef habitats. However, few studies, particularly in the Indo-Pacific, consider a broad range of habitats when investigating the use of non-reef habitats by coral reef fishes. It is only through synthesis of the data that we can begin to see that coral reef fishes have been observed in a much broader range of habitats across the seascape than previously considered. By extending the search for coral reef fishes to a wider range of habitats, we can build a more complete picture of which habitats are important for each species, and identify any species that may be quite singular in their use of non-reef habitats. For example, the night sergeant (*Abudefduf taurus*, Pomacentridae) has been searched for in habitats dominated by hard structures (i.e., corals, rocky boulders) and vegetation (i.e., seagrass, mangrove and macroalgae), but has only been observed in the hard structure habitats (Nagelkerken et al. 2000a). This indicates that the availability of hard structure, coral or otherwise, is likely to be more important for this species than the presence of vegetated habitat. Identifying the extent of habitat plasticity for a species, and the habitat characteristics that might be important, will provide us with a far greater insight into the impacts of changing seascapes (e.g., habitat loss; Wilson et al. 2010).

The findings are also predominantly based on single daytime observations of species distribution patterns. However, this approach fails to consider how tidal, diel, seasonal or annual variability may affect patterns of habitat use. In instances where studies have included a temporal component, changes to fish communities in the habitats of interest are common (e.g., tidal, Unsworth et al. 2007b; day-night shifts, Kopp et al. 2007; Hitt et al. 2011; seasonal and annual, Kimirei et al. 2011). Including a temporal aspect in studies that examine multiple

habitat use in reef fishes, as well as considering additional non-reef habitats, may reveal even more extensive use of the seascape by certain species of coral reef fish.

Another interesting question is whether differences in habitat use by coral reef fishes exist both between and within the tropical Atlantic and Indo-Pacific regions. Drivers of differences in patterns of habitat use could include the type of non-reef habitats available within study locations (Dorenbosch et al. 2004; Mumby et al. 2004), proximity of habitats (Nakamura et al. 2009a; Nagelkerken et al. 2012; Olds et al. 2013), areal extent of non-reef habitats (Grober-Dunsmore et al. 2004; Gilby et al. 2016), and/or the tidal regime (Unsworth et al. 2007b; Igulu et al. 2014; Harborne et al. 2016a; Davis et al. 2017). For example, seagrass and mangrove habitats are absent from parts of the Indo-Pacific, which could increase the importance of other types of non-reef habitat (e.g., macroalgal beds) to coral reef fishes in this region. Tidal range may also play a role in the use of non-reef habitats by coral reef fishes. In the micro-tidal tropical Atlantic, such habitats are constantly inundated with water and therefore permanently available to fishes (Igulu et al. 2014). In contrast, the Indo-Pacific experiences micro-, meso- and macro-tidal conditions. For locations with larger tidal ranges, some non-reef habitats may be inaccessible at low tide and/or physiologically hostile due to rapid fluctuations in environmental variables (e.g., temperature, salinity; Sheaves 2005; Igulu et al. 2014). Despite this potential temporal constraint on habitat availability, >120 fish species were recorded in mangroves from the 19 studies that examined coral reef fishes in mangroves in the Indo-Pacific. This suggests that mangroves are still extensively used by coral reef fishes in this region. Few studies have, however, investigated how tides might influence the use of non-reef habitats by reef fishes in the Indo-Pacific in combination with surveys on the reef (but see Unsworth et al. 2007b).

More than >230 coral reef fish species were observed as juveniles in non-reef habitats. This figure is likely to be a conservative estimate because many studies did not differentiate between life history stages, and other studies exist that have examined juveniles of coral reef fishes in non-reef habitats but did not simultaneously quantify their use of adjacent coral reef habitat (e.g., Tano et al. 2017) and thus were excluded from this review. Vegetated habitats, particularly mangroves and seagrass beds, have frequently been reputed to provide an important nursery function for many coral reef fish species (Nagelkerken et al. 2000a; Nagelkerken et al. 2001; Sheaves et al. 2015), particularly those species of commercial interest to fisheries (Blaber 2009). However, in addition to observations in mangroves and seagrass

beds, this review also confirms the widespread occurrence of juvenile coral reef fishes in macroalgae beds despite them currently being one of the least studied non-reef habitat types (13% of studies in this review). The potential importance of macroalgal beds as juvenile coral reef fish habitat is further supported by studies from north-eastern Brazil (Chaves et al. 2013) and Zanzibar, Tanzania (Tano et al. 2017). In particular, the use of macroalgal beds seems to be prevalent for labrids, pomacentrids, and siganids (Rossier & Kulbicki 2000; Adams & Ebersole 2002; Wilson et al. 2010; Hoey et al. 2013; Rasher et al. 2013; Evans et al. 2014). These findings are interesting, as it has been suggested that macroalgal-dominated reefs suppress recruitment in many coral reef fishes including pomacentrids, labrids and siganids (Dixson et al. 2014). Yet from this synthesis, it appears that at least for some species, macroalgal beds are a commonly utilised habitat by juveniles. Species of coral reef fish that use macroalgal beds as juveniles could have the potential to become more dominant on reefs experiencing regime-shifts or high macroalgae loads. To understand how this could affect ecosystem functioning on reefs in the future, we need a better understanding of patterns and drivers of macroalgal bed use by coral reef fishes and the contribution such species make to ecological processes on reefs.

There are a wide range of different factors that could influence non-reef habitat use by coral reef fishes, including key traits such as mobility, body size and feeding ecology. An individual's mobility will influence how it interacts with the wider seascape, and across what spatial and temporal scales. For some taxa, body size and mobility can be tightly coupled (Kramer & Chapman 1999; Sale et al. 2005). We might reasonably expect then, that large-bodied highly mobile species are more likely to use multiple habitat types than smaller-bodied, less mobile species. Many of the coral reef fishes using multiple habitats identified in this review were large-bodied, mobile carnivores or piscivores. Such species may have higher energetic demands that require larger home ranges to meet their resource requirements.

Comparing the results to the species list compiled as part of an extensive review on known movement patterns of coral reef fishes by Green et al. (2015), I found that movement data were available for 114 of the 670 species included in the review. Much of the data available explores only a single aspect of movement (e.g., spawning migrations), and hence likely fails to capture the full breadth of potential seascape usage. Moreover, movement data for many species is restricted to a single location which, whilst informative, fails to capture seascape variability. This information is needed, particularly for marine reserve design (Pittman & McAlpine 2003;

Green et al. 2015), because home range or daily movement patterns may vary considerably depending on individual behaviors and the spatial configuration and connectivity of habitat patches in the seascape (Hitt et al. 2011).

Contrary to expectations, a substantial proportion of the species observed in non-reef habitats were small-bodied species. Little, if any attention has been given to species typically thought of as permanent coral reef residents, or species with limited mobility, such as pomacentrids. Yet, large numbers of pomacentrid species were recorded in non-reef habitats, particularly in the Indo-Pacific, and these observations included both adults and juveniles. Moreover, many of these species were recorded using non-reef habitats from multiple studies and/or locations (e.g., beaugregory (*Stegastes leucostictus*, Pomacentridae) in the tropical Atlantic; Adams & Ebersole 2002; Christensen et al. 2003; Aguilar-Perera & Appeldoorn 2008; Harborne et al. 2008; and jewel damselfish (*Plectroglyphidodon lacrymatus*, Pomacentridae) in the Indo-Pacific; Wilson et al. 2010; Berkström et al. 2012; Kruse et al. 2016) indicating that the results are not location specific or due to chance (e.g., recruitment) events. Such species are unlikely to move considerable distances (e.g., pomacentrids, Turgeon et al. 2010) from where they settle. This suggests that there may be another, overlooked, category of coral reef fish species that can occur in non-reef habitats in addition to the standard explanations for non-reef habitat use (e.g., nursery, foraging etc.). Individuals in this alternate group may settle in non-reef habitats deliberately or because of an absence of more suitable habitat at the time of settlement. Once settled in a non-reef habitat, the limited mobility of such individuals would likely constrain them to that habitat for their entire life. Although abiotic and biotic conditions between reef and non-reef habitats are likely to differ substantially, trade-offs in terms of growth, survival and reproduction (Dahlgren & Eggleston 2000) between habitat types have received limited attention (but see Nakamura & Sano 2004a; Nakamura et al. 2007; Grol et al. 2008; Grol et al. 2011a, b; Grol et al. 2014). Given the widespread presence of small-bodied, and presumably less mobile coral reef fish in non-reef habitats documented in this review, the fitness consequences of living in reef versus non-reef habitats warrant further investigation.

Many species of herbivorous fishes were also observed in non-reef habitats. Herbivorous fishes perform a critical role on coral reefs (Nyström & Folke 2001; Scheffer et al. 2001; Rasher et al. 2013), by scraping and excavating the substratum, grazing on algal turfs, or browsing on macroalgae that help maintain a healthy balance between corals and macroalgae. Interestingly, several of the herbivorous species identified as multi-habitat users in this review have been

identified as major contributors to herbivory on reefs, particularly species from the Acanthuridae (Hoey & Bellwood 2009; 2010a; Rasher et al. 2013; Chong-Seng et al. 2014) Siganidae (Fox & Bellwood 2008; Cvitanovic & Bellwood 2009; Cheal et al. 2010), and Scarinae (Bellwood et al. 2003; Mumby et al. 2006; Hoey & Bellwood 2008; Kuempel & Altieri 2017). The frequent presence of herbivores in non-reef habitats raises several interesting questions. For example, whilst larval replenishment from connected reefs following disturbance events has received considerable attention (e.g., Almany et al. 2007; Jones et al. 2009; Hogan et al. 2012), little consideration has been given to the potential role of non-reef habitats as support areas for the replenishment of reefs (but see Mumby & Hastings 2008; Edwards et al. 2010), or the links between larval dispersal and availability of nursery habitats (but see Brown et al. 2016). Such research is important given that the availability of nursery habitats for herbivores near to reefs has been implicated as a key factor on reefs that have repeatedly exhibited rapid recovery rates following coral loss (Adam et al. 2011). Another poorly explored topic is whether non-reef habitats could offer refuge for coral reef fish during disturbance events that differentially impact coral reefs (e.g., coral bleaching events). To date, the search for potential refugia for coral reef fishes has been reef-centric, typically looking at deeper reefs which may experience more stable abiotic conditions than shallow reefs (e.g., Bridge et al. 2013; Jankowski et al. 2015; MacDonald et al. 2016). However, there is evidence that deeper reefs may not be suitable refugia for some species, especially herbivores (Kahng et al. 2010). In light of the deteriorating conditions many reefs are facing, these are important future research subjects.

Much of the research into multi-habitat use has focused on species that use non-reef habitats as juveniles (e.g., Dennis 1992; Nagelkerken et al. 2000a, 2000b; Cocheret de la Morinière et al. 2002; Gillanders et al. 2003; Nagelkerken et al. 2012; Nakamura et al. 2012) or those that make short-term movements between reef and non-reef habitat to feed (e.g., Ogden 1976; Robblee & Zieman 1984; Meyer & Schultz 1985; Burke 1995). Families that contain species commonly associated with ontogenetic shifts from vegetated habitats as juveniles to coral reefs as adults include the Haemulidae (e.g., Tropical Atlantic: Nagelkerken & van der Velde 2004; Grol et al. 2014), Lethrinidae (e.g., Indo-Pacific: Mellin et al. 2007; Kimirei et al. 2013a) and Lutjanidae (e.g., Tropical Atlantic: Nagelkerken & van der Velde 2004; Aguilar-Perera & Appeldoorn 2007; Indo-Pacific: Nakamura et al. 2008; Kimirei et al. 2013b; Paillon et al. 2014), as well as some species from the Scarinae (e.g., Tropical Atlantic: Mumby et al. 2004; Dorenbosch et al. 2006b). Yet until recently, it has been challenging to provide direct, rather

than correlative, evidence for ontogenetic shifts from non-reef habitats to coral reefs (Gillanders et al. 2003). This issue is now being addressed through the use of molecular approaches (e.g., stable isotope analysis) that can begin to quantify the contribution of non-reef habitat-using juveniles to adult populations on the reef (e.g., *Lutjanus ehrenbergii*, Lutjanidae), McMahon et al. 2012; *Lutjanus fulviflamma* (Lutjanidae), Paillon et al. 2014). However, in contrast to the many species of coral reef fishes that display an obligate relationship with reef habitat as juveniles (Jones et al. 2004; Wilson et al. 2008), few, if any, coral reef fishes appear to be functionally dependent on the availability of non-reef habitats as nurseries (Nagelkerken et al. 2000c, but see Mumby et al. 2004).

Another explanation for the presence of reef fishes in non-reef habitats is diel and/or tidal movement between feeding and shelter habitats. Several species of haemulids in the Caribbean have been documented making twilight feeding incursions away from reefs into seagrass beds to feed (see reviews by Appeldoorn et al. 2009; Krumme 2009) but there is little evidence that this behaviour is replicated by haemulids in the Indo-Pacific. Indeed, there is limited empirical data generally for the Indo-Pacific on this topic (but see Honda et al. 2016), although studies demonstrating tidal (e.g., Unsworth et al. 2007b) and diel (e.g., Unsworth et al. 2007a) changes to fish communities on reefs and in non-reef habitats indicate potential for regular feeding movements.

As well as direct uses, non-reef habitats can also play a role in supporting populations of reef fishes and influencing ecological processes on coral reefs through indirect mechanisms. This creates an additional level of complexity when attempting to evaluate the importance of non-reef habitats. For example, the Caribbean rainbow parrotfish (*Scarus guacamaia*) is widely considered to have a strong direct dependency on mangroves as juveniles, with adults absent from reefs more than 10 km away from mangrove habitats (e.g., Mumby et al. 2004; Dorenbosch et al. 2006b), although there are exceptions (Aguilar-Perera & Hernández-Landa 2017). In contrast, the presence of mangroves appears to indirectly support the Indo-Pacific bumphead parrotfish (*Bolbometopon muricatum*) by reducing sedimentation rates on inshore fringing reefs and hence protecting key coral microhabitats for juveniles (Hamilton et al. 2017). Other indirect effects can also be seen through the feeding incursions into vegetated habitats which can contribute nutrients to coral reef systems through defecation (Meyer & Schultz 1985; Hyndes et al. 2014; Shantz et al. 2015). In addition, it appears that whilst proximity to non-reef habitats can result in increased adult abundances of nursery-using species on reefs, it can have

the opposite effect on reef-dependent species and, concomitantly alter key ecological processes (e.g., grazing; Harborne et al. 2016b). The potential for indirect effects (e.g., trophic and/or competitive cascades; Harborne et al. 2016b) to alter fish communities and ecological processes on reefs is therefore likely to be an important but understudied aspect of heterogeneous seascapes.

The findings from this review are highly relevant for fisheries management. Over half of all the species (60%) observed in non-reef habitats are fisheries targets in some capacity, and of these 50% were observed in two or more habitat types in addition to the reef. In particular, many commercially important species from the Carangidae, Haemulidae, Lethrinidae, Lutjanidae and Serranidae were identified as multiple habitat users. Coral reef fisheries provide an important source of income and protein, particularly in developing, or small island nations (Newton et al. 2007). To contribute towards long-term food security and sustainability of stocks, it essential that fisheries species at each stage of their life cycle are supported (Nagelkerken et al. 2015; Baker et al. 2018). Without adequate management of the entire seascape, loss of key non-reef habitats could have negative repercussions for fisheries productivity on coral reefs. However, for many fisheries species identified in this review, we still lack a comprehensive understanding of habitat use patterns and whether dependencies exist on particular habitats for different life history stages. These knowledge gaps limit our ability to effectively manage coral reef fisheries. Focusing management activities solely on reef environments may fail to protect the most vulnerable life stages, or neglect important feeding or spawning grounds which contribute towards population replenishment. Therefore, there is a need to better understand the importance of non-reef habitats for key fisheries species and broaden “coral reef” fisheries management to encompass the wider seascape (e.g., Unsworth et al. 2014; Baker et al. 2018).

From a conservation perspective, the review revealed that 28 of the species observed in non-reef habitats are currently considered at risk of extinction. To protect such vulnerable species, we need data on their direct and indirect habitat and resource requirements for each component of their life cycle, as well as short- and long-term movement patterns (Green et al. 2015). However, detecting rare species using conventional survey techniques can be challenging (Dulvy & Polunin 2004; Pikitch 2018), leading to an incomplete picture of species-habitat relationships. Environmental DNA (eDNA) is a non-destructive molecular technique that can detect the presence of rare species more time and cost efficiently than traditional sampling

techniques (Rees et al. 2014; Pikitch 2018). Recent studies from coastal marine environments indicate that this tool could be applied to detect the presence of rare species across small spatial scales (50-100 m; Port et al. 2016; O'Donnell et al. 2017) which will be required for coral reef seascapes with a range of habitats in close proximity. As a relatively new molecular tool, there are uncertainties around the dynamics of eDNA across spatial and temporal scales (O'Donnell et al. 2017). However, assuming these uncertainties can be resolved satisfactorily, this approach could help in the future to unravel the mysteries of habitat use for species that are of conservation concern. Whilst there is much still to learn about habitat use by vulnerable species, this review provides insight into which 'at risk' species have been recorded in non-reef habitats to date, and provides a starting point for more extensive research to guide conservation efforts in the future.

Information on the four W's (what, when, where and why) is lacking or incomplete for many species. As a consequence, there is still much uncertainty and debate about the importance of non-reef habitats for coral reef fishes and the functional contribution of non-reef habitats to maintain or support ecological processes on coral reefs. Even the ability to group species into categories based on the reasons for use (e.g., nursery, food, shelter), their temporal (e.g., short- or long-term, diel, tidal etc.) or spatial scale of use (e.g., home ranges, spawning or foraging migrations) is restricted to a small number of species. Furthermore, it appears that there are two additional groups of multi-habitat users that have been largely neglected thus far. The first group are transients, specifically, highly mobile individuals or groups of individuals that interact regularly with different components of the seascape. I speculate that the ecological role of these transients should not be underestimated; passing through a habitat patch may also entail some form of habitat use or modification. For instance, transient individuals may specifically use structurally complex habitats, or habitat edges, as corridors for movement between different components of the seascape (Boström et al. 2011; Hitt et al. 2011; Nagelkerken et al. 2015; Davis et al. 2017). The importance of fish as mobile links between habitats has been acknowledged for some time (e.g., Nyström & Folke 2001; Lundberg & Moberg 2003), and highly mobile transients have the potential to be significant contributors to energy transfer between habitats. Opportunistic feeding by transients may influence community structure, individual behaviour and survivorship within non-reef habitats (Rooker et al. 2018), and defecation by transients, particularly for schooling species, could make a substantial contribution to nutrient exchange between habitats.

The second understudied group are individuals that may be able to complete their entire life cycle away from coral reefs. In both of these groups, populations of a species are likely to use multiple habitat types, but in the first instance, individuals will have the capacity to move between different components of the seascape, whereas in the second instance, individuals will settle and remain in a particular patch of habitat. To address the many gaps in our knowledge, a multi-faceted approach incorporating observational, experimental, molecular tools and acoustic telemetry is required. Combining such techniques can rapidly enhance our understanding of seascape usage at the species level (see Baker et al. 2018).

The future of coral reef ecosystems is increasingly uncertain due to a diverse range of stressors, namely elevated sea surface temperatures, severe storms, pollution, disease, and crown-of-thorns starfish outbreaks (Gardner et al. 2003; Osborne et al. 2011; De'ath et al. 2012; Hughes et al. 2017a; Unsworth et al. 2018). However, coral reefs are not the only coastal habitat threatened by anthropogenic and natural stressors. Globally, mangrove losses had reached around 35% two decades ago (Valiela et al. 2001; Duke et al. 2007) with recent estimates showing a continued, but reduced rate of decline (0.16-0.39% per year; Hamilton & Casey 2016), and seagrass has been disappearing at a rate of 110 km<sup>2</sup> per year since 1980 (Waycott et al. 2009). Faced with the deterioration of many coastal seascapes, it is imperative that we expand our understanding of how reef fishes use and depend on the range of habitats in the seascape. A seascape perspective is vital for understanding the strength and significance of connections between habitats, particularly because the loss and fragmentation of various habitats in the seascape is likely to alter functional and structural links between habitat patches in the seascape. Such information will help to more effectively manage coral reef seascapes and their associated fish populations in the future.

## Chapter 3

### Broadening our horizons: seascape use by coral reef-associated fishes in Kavieng, Papua New Guinea, is common and diverse <sup>2</sup>

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#### 3.1 Abstract

There is increasing evidence that non-reef habitats in the seascape surrounding coral reefs are widely used by reef-associated fishes. However, our understanding of seascape use in the Indo-Pacific region is incomplete due to its large geographical range and as a consequence, considerable environmental variation (e.g., tidal regimes). Remote video cameras were used to survey reef-associated fishes within five habitat types (coral reef slope, coral reef flat, macroalgal beds, mangroves and seagrass meadows) around the Tigak Islands, Kavieng, Papua New Guinea. Of the 282 shallow-water reef-associated species observed across 360 videos, 35% (99 species) were recorded in non-reef habitats, the majority (78 species) on multiple occasions. Macroalgal beds dominated by low complexity algal genera (e.g., *Halimeda*, *Caulerpa*) were used extensively by reef-associated fishes, complementing previous research that has documented the use of canopy-forming macroalgae (e.g., *Sargassum*). Mean species richness and relative abundances (MaxN) of reef-associated fishes were two-fold higher in macroalgal beds than mangroves or seagrass. Interestingly, mangroves contained the most distinct fish assemblage of the three non-reef habitats, including several reef-associated species that were not recorded from any other habitat type. This suggests that mangroves possess attributes not shared by other shallow non-reef, or even reef, habitats. Importantly, many of the fish families commonly found in non-reef habitats (i.e., lethrinids, lutjanids) are targeted by local fishers and are thus critical to sustaining local livelihoods. This study demonstrates non-reef habitat use is common for many reef-associated fishes and highlights the need to incorporate a range of habitats into study designs to better understand habitat use patterns in the Indo-Pacific. Given the widespread degradation of coral reefs and other shallow-water habitats, it is becoming more important to recognise that reefs are embedded within a mosaic

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of habitat types that influence patterns and processes, and scale management strategies appropriately.

### **3.2 Introduction**

The influence of the wider landscape or seascape on patterns and processes observed within a habitat patch remains a long-standing and important question in ecology (Miller-Rushing et al. 2019). It has been proposed that the close proximity of particular kinds of habitat may have a range of effects on community composition, species diversity and ecosystem function (Fahrig et al. 2011). The effects of habitat adjacency on species diversity may occur directly, for example if a species requires resources that are found in both habitat types (e.g., different breeding and foraging habitats); or indirectly, via exchanges of nutrients and other subsidies between adjacent habitats (Alsterberg et al. 2017). Habitat adjacency and complementarity have long been recognised as important drivers of species abundance and distribution patterns in terrestrial systems (Fahrig et al. 2011), and a similar effect is increasingly evident from shallow-water coastal ecosystems (e.g., Dorenbosch et al. 2006a; Pittman et al. 2007; Olds et al. 2012a; Berkström et al. 2013a; Aller et al. 2014).

Many coral reef fish taxa use multiple habitat types throughout their life cycle (Nagelkerken et al. 2000c; Adams et al. 2006; Harborne et al. 2008). An understanding of the use of these habitats and the connections between them is critical to predict and manage the likely impacts of local and global disturbances (Berkström et al. 2012; Pittman & Olds 2015). Although numerous studies have documented the effects of coral loss on coral reef fish communities (e.g., Jones et al. 2004; Pratchett et al. 2011; Richardson et al. 2018), far fewer have considered how disturbances in the wider seascape can influence fish communities, despite many reef systems occurring within a mosaic of highly productive (Birkeland 1985) and extensive (Parrish 1989) non-reef habitat types such as mangroves and seagrass meadows. Much like coral reefs, these shallow-water ecosystems are undergoing rapid systemic change with widespread areal losses and degradation in recent decades (Valiela et al. 2001; Waycott et al. 2009; Hamilton & Casey 2016).

The number of coral reef-associated fish species that have been observed in non-reef habitats is substantial. Globally, at least 670 species of coral reef-associated fishes have been recorded in non-reef habitats in addition to reefs, representing approximately 20% of all coral reef fish species (Sambrook et al. 2019, **Chapter 2**). Reef fishes use the wider seascape for a range of

reasons including foraging (Beets et al. 2003; Hitt et al. 2011), spawning (Pittman & McAlpine 2003), and as juvenile habitat before migrating to reefs as subadults or adults (Dahlgren & Eggleston 2000; Nagelkerken et al. 2000c; Nagelkerken et al. 2001; Adams et al. 2006; Jaxion-Harm et al. 2011). As a result, short- and long-term movements by reef fishes between different components of the seascape can contribute to ecosystem functioning on reefs through nutrient transfer (Meyer & Schultz 1985; Shantz et al. 2015), trophic subsidies and cascades (Heck et al. 2008; Harborne et al. 2016b), and population replenishment (Nakamura et al. 2008; McMahan et al. 2012).

Much of our knowledge about seascape use by reef-associated fishes comes from the Caribbean (Nagelkerken et al. 2000c; Aguilar-Perera & Appeldoorn 2008; Dorenbosch et al. 2009). Less is known about how, why and which reef fishes use non-reef habitats in the Indo-Pacific (Sambrook et al. 2019, **Chapter 2**). Our limited understanding in the Indo-Pacific is, in part, complicated because of its large spatial extent and biophysical variability. For instance, unlike the Caribbean where shallow-water habitats are permanently accessible to fishes, tidal regimes in the Indo-Pacific range from micro- to macro-tidal (Krumme 2009), affecting the accessibility of non-reef habitats to reef fishes (Igulu et al. 2014). As a consequence, understanding fish-habitat relationships in the Indo-Pacific requires exploration across a wider range of locations and tidal regimes.

Understanding habitat use patterns of coral reef fishes is also highly relevant when addressing concerns around long-term food security in the Indo-Pacific (Foale et al. 2013; Blasiak et al. 2017). This is because many of the coral reef fishes that are known to use non-reef habitats are common fisheries targets (Sambrook et al. 2019, **Chapter 2**) and the Indo-Pacific is home to a multitude of small island communities (Brodie et al. 2013) that rely on coral reef fisheries to satisfy daily nutritional requirements (Béné et al. 2007) and as a primary source of income (Bell et al. 2009). Many of these coastal communities are experiencing rapid population growth (Burke et al. 2011) which places increasing pressure on already stretched natural resources (Bell et al. 2009). By expanding our understanding of broader seascape use by coral reef fishes, we can identify essential fish habitats, combinations of habitats and/or target species that require better management or protection, which could contribute towards longer-term sustainable fisheries goals.

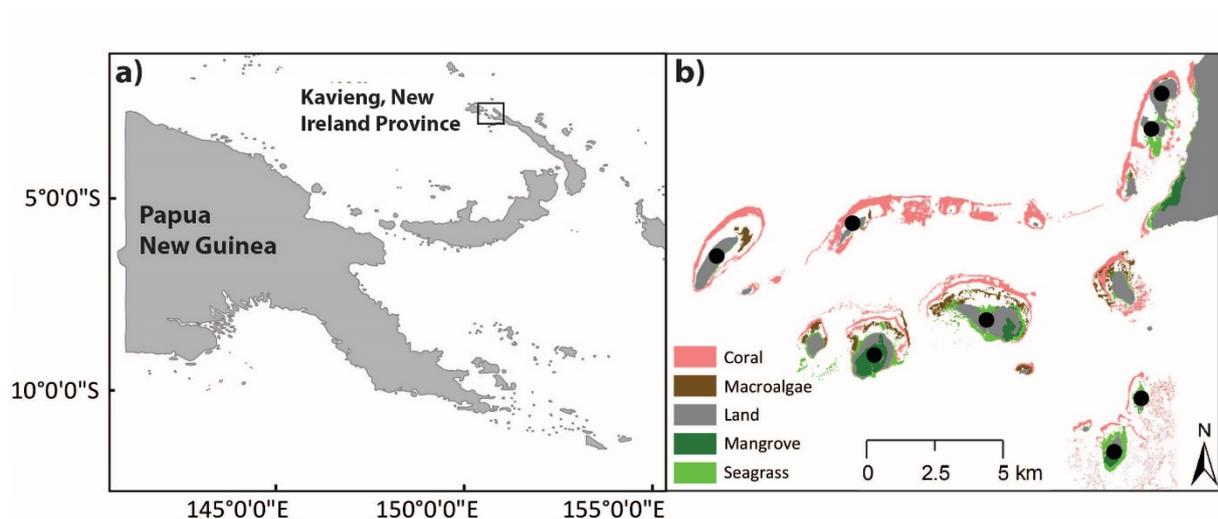
The objective of this study was to describe and compare reef-associated fish communities across five habitat types that are common in coastal tropical marine seascapes (i.e., coral reef

flats, coral reef slopes, mangroves, seagrass meadows and macroalgae beds) in the Indo-Pacific. Specifically, the reef fish assemblages associated with the five habitat types were compared, the overlap in habitat use was quantified and the frequency of use of non-reef habitats identified in Kavieng, Papua New Guinea.

### 3.3 Methods

#### 3.3.1 Study site

The study was conducted around the Tigak Island Group within the Kavieng lagoon, New Ireland, Papua New Guinea ( $2^{\circ} 34' S$ ,  $150^{\circ} 48' E$ ; Figure 3.1). The Kavieng lagoon is  $\sim 380$  km<sup>2</sup> and contains a range of habitats. Extensive reef formations around islands are interspersed with seagrass meadows (predominantly *Enhalus* and *Thalassia* spp.), macroalgae beds (predominantly *Halimeda*, *Caulerpa* spp.) and mangrove forests (*Rhizophora* spp.). The annual water temperature ranges between  $28.7^{\circ}C$  and  $31.6^{\circ}C$  (NOAA 2019). Tides are mixed microtidal (Krumme 2009), with a maximum tidal range of 1.09 m. As a consequence, nearshore habitats (e.g., mangroves) are generally submerged, although inundation depths can be shallow ( $\sim 30$  cm).



**Figure 3.1.** a) Map of Papua New Guinea with location of Kavieng, New Ireland Province, and b) study sites and habitats in the Tigak Island Group, Kavieng. Black circles denote the surveyed islands.

### 3.3.2 Data collection

To quantify habitat use by coral reef-associated fishes, data were collected from five common habitat types, specifically: 1) shallow coral reef slopes (3-6 m depth), 2) reef flats containing hard structure (e.g., coral or rock), 3) macroalgal beds, 4) non-estuarine mangroves and, 5) seagrass meadows. Data on the use of these habitats by reef fishes were collected between 09:00 and 16:00 during April 2018, using unbaited underwater video cameras. This sampling method was chosen as it reduced any bias due to diver presence (Gotanda et al. 2009; Feary et al. 2011) and also due to the presence of saltwater crocodiles (*Crocodylus porosus*) in the area. The lack of baits on the cameras ensured that observations of habitat use were not influenced by attraction of fishes to bait (Bassett & Montgomery 2011).

Single video cameras (GoPros) mounted on steel frames were lowered to the substratum and GPS points were recorded for each camera drop. The use of single, as opposed to stereo, video systems precluded the collection of accurate body size data for fishes and prevented the separation of individual fishes into life stages based on body size. For mangrove and reef slope habitats, care was taken to ensure that cameras faced toward the habitat as opposed to adjacent open waters. Replicate camera drops were separated by a minimum of 50 m, both among and between the five habitat types. Each camera was deployed for a minimum of 20 minutes to enable high replication across broad-spatial scales within a relatively short time frame (e.g., Burge et al. 2012; Bradley et al. 2017; Pereira et al. 2017). The depth of the camera drops ranged from 0.3 to 5.5 m. In total 86 reef slope, 75 reef flat, 41 macroalgae, 58 mangrove and 100 seagrass videos were analysed. This variation was due to differences in the availability of each habitat type, and the exclusion of replicates with low video quality (camera fogging and limited underwater visibility).

### 3.3.3 Video analysis

For each video, a 15-minute segment was analysed. Each segment began at least one minute after the camera had stabilised on the bottom and any sediment disturbed during placement had settled. From each video, species presence and the maximum number of individuals of a species observed in a single frame ( $MaxN$ , *sensu* Cappo et al. 2004) were recorded.  $MaxN$  is a common metric used as a conservative measure of relative abundance (Campbell et al. 2015). Cryptic (e.g., Blenniidae, Gobiidae) and surface-dwelling (e.g., Hemiramphidae) taxa were excluded because they were not able to be consistently counted using video. Individuals were identified to genus or species where possible. *FishBase* (Froese & Pauly 2019) was used to provide an

objective assessment of which species were considered reef-associated, hereafter termed “reef fishes”, or not reef-associated (following Sambrook et al. 2019, **Chapter 2**).

### **3.3.4 Data analysis**

Differences in the composition of fish assemblages between the five habitats were compared with a one-way Permutational Multivariate Analysis of Variance (PERMANOVA) using Type III sum of squares and 9999 permutations (Anderson et al. 2008). A zero-adjusted Bray-Curtis similarity matrix was used to account for the high number of zeros present in the Max $N$  data and applied a fourth-root transformation (Clarke et al. 2006). Pair-wise tests were used to examine differences between habitats and visualised the data using non-metric multidimensional scaling (nMDS). The similarity percentages routine (SIMPER) was used to identify characteristic species for each habitat type.

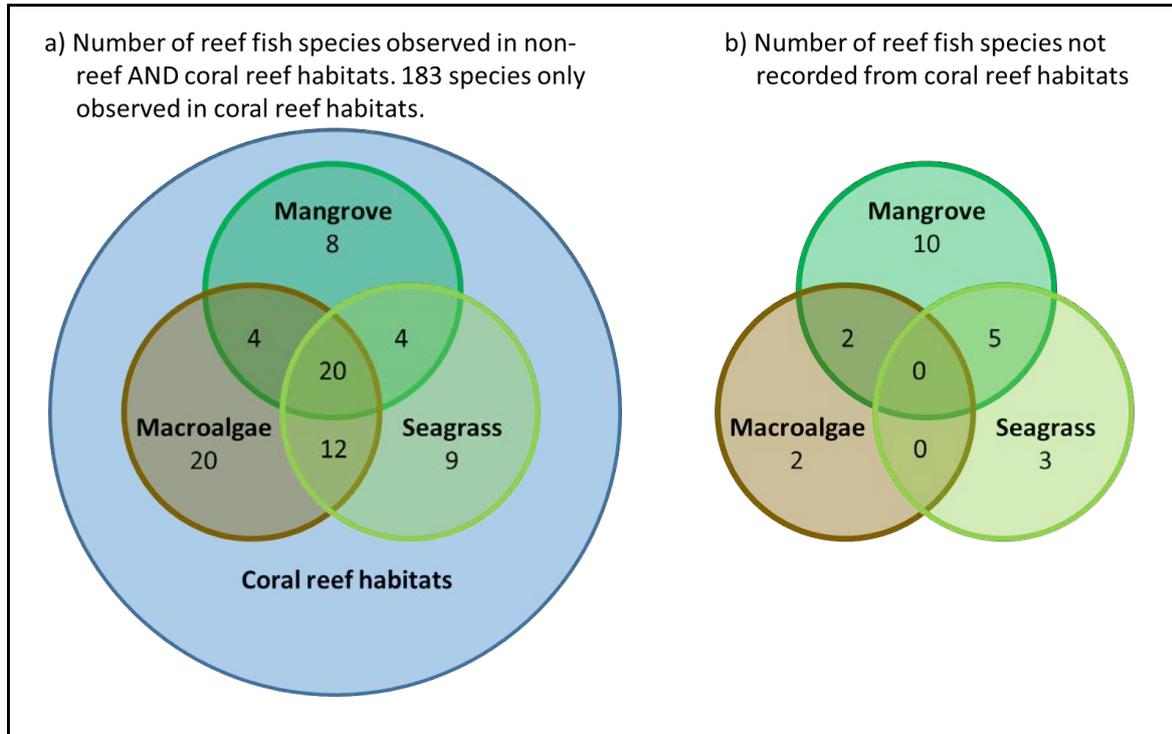
Differences in the mean Max $N$  (i.e., relative abundance) and species richness between habitats were compared with one-way ANOVAs, followed by Tukey’s HSD post hoc tests using R. In addition, one-way ANOVAs were used to compare differences in mean Max $N$  for common multi-habitat users to explore whether group sizes might differ between habitat types.

## **3.4 Results**

Across the five habitat types (reef flat, reef slope, macroalgae, mangrove and seagrass), 15,492 individuals were recorded from a total of 319 taxa, of which 288 were identified to species. Across the five habitats surveyed, 282 out of the 288 species (98%) recorded were classed as reef fishes, with only six species observed on the videos considered non-reef associated. These six non-reef associated fish species were excluded from all further analyses.

In total, 35% of the reef fishes (99 of 282 species) observed in this study were recorded in non-reef habitats, with 20 species recorded in all three non-reef habitats (Figure 3.2a; Table A3.1). Although less speciose compared to the coral reef slope and flat habitats, the total number of reef fish species observed in each non-reef habitat was considerable (Table 3.1). A total of 53 reef fish species were observed in both seagrass meadows and mangroves and 60 species were observed in macroalgae beds (Figure 3.2; Table 3.1). While the total number of species observed was similar across the three non-reef habitat types, the mean species richness and relative abundance (i.e., mean Max $N$ ) of reef fish per video was approximately twofold higher in macroalgae beds compared to mangroves or seagrass meadows (Table 3.1; ANOVA  $F_{2,196}$

= 15.002,  $p < 0.001$ ). The type and/or combination of non-reef habitats used by each of the 99 reef fish species varied widely. Over half of the species were observed in a single non-reef habitat (Figure 3.2), most commonly in macroalgae (22 spp.), followed by mangroves (18 spp.) and seagrass (12 spp.). However there was also considerable overlap in habitat use with almost half (47 of 99 species) recorded from two or more non-reef habitats (Figure 3.2).



**Figure 3.2.** a) Number of reef fish species recorded in reef and non-reef habitats and overlap between non-reef habitats, and b) the number of species not recorded from coral reef habitats (reef slope and/or reef flat).

**Table 3.1.** The number of replicate video samples ( $n$ ), total number of reef fish species observed across all videos, as well as the mean species richness and relative abundance (i.e., total MaxN) of reef fish observed in each habitat type. Total MaxN calculated by summing the MaxN of each species recorded per video

	Sample size $n$	Total no of reef fish species observed	Mean species richness per video $\pm$ SE (range)	Mean relative abundance (total MaxN) per video $\pm$ SE (range)
Reef slope	86	243	35.05 $\pm$ 1.19 (14-59)	95.56 $\pm$ 7.07 (16-420)
Reef flat	75	155	21.76 $\pm$ 0.83 (8-46)	46.20 $\pm$ 2.57 (9-143)
Macroalgae	41	60	13.12 $\pm$ 0.90 (3-36)	31.24 $\pm$ 3.09 (4-70)
Mangrove	58	53	6.84 $\pm$ 0.52 (1-19)	16.69 $\pm$ 1.78 (1-56)
Seagrass	100	53	5.43 $\pm$ 0.39 (0-16)	15.6 $\pm$ 1.54 (0-72)

### 3.4.1 *Fish community differences between habitats*

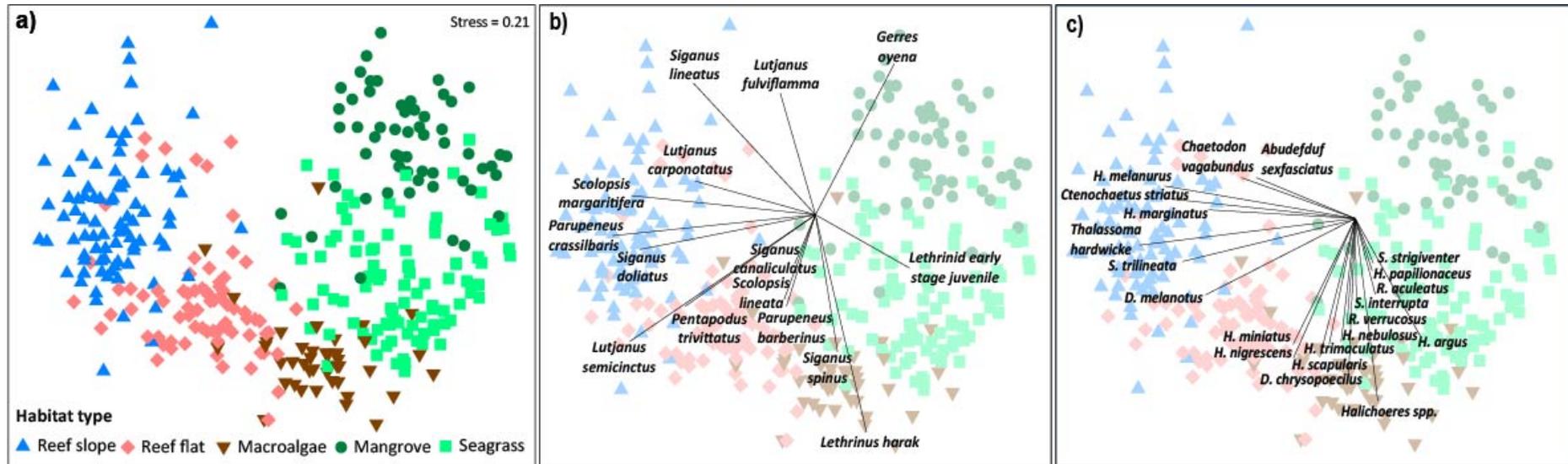
Reef fish assemblages differed between the five habitats (PERMANOVA pseudo- $F_{4,355} = 38.926$ ,  $p=0.0001$ , Figure 3.3), with each habitat type containing a distinct assemblage of reef fishes. Macroalgae beds were broadly characterised by several species of *Halichoeres*, the tuskfish *Choerodon anchorago*, the emperor *Lethrinus harak*, and the damselfish *Dischistodus chrysopoecilus*. *Lethrinus harak*, *C. anchorago* and *Halichoeres* spp., together with the rabbitfish *Siganus canaliculatus* were characteristic of seagrass habitats. Mangroves were characterised by a different suite of species including *Gerres oyena*, *Lutjanus ehrenbergii*, *Lutjanus fulviflamma*, the rabbitfish *Siganus lineatus* and the cardinalfish *Sphaeramia orbicularis*.

### 3.4.2 *Habitat use patterns by reef fish family*

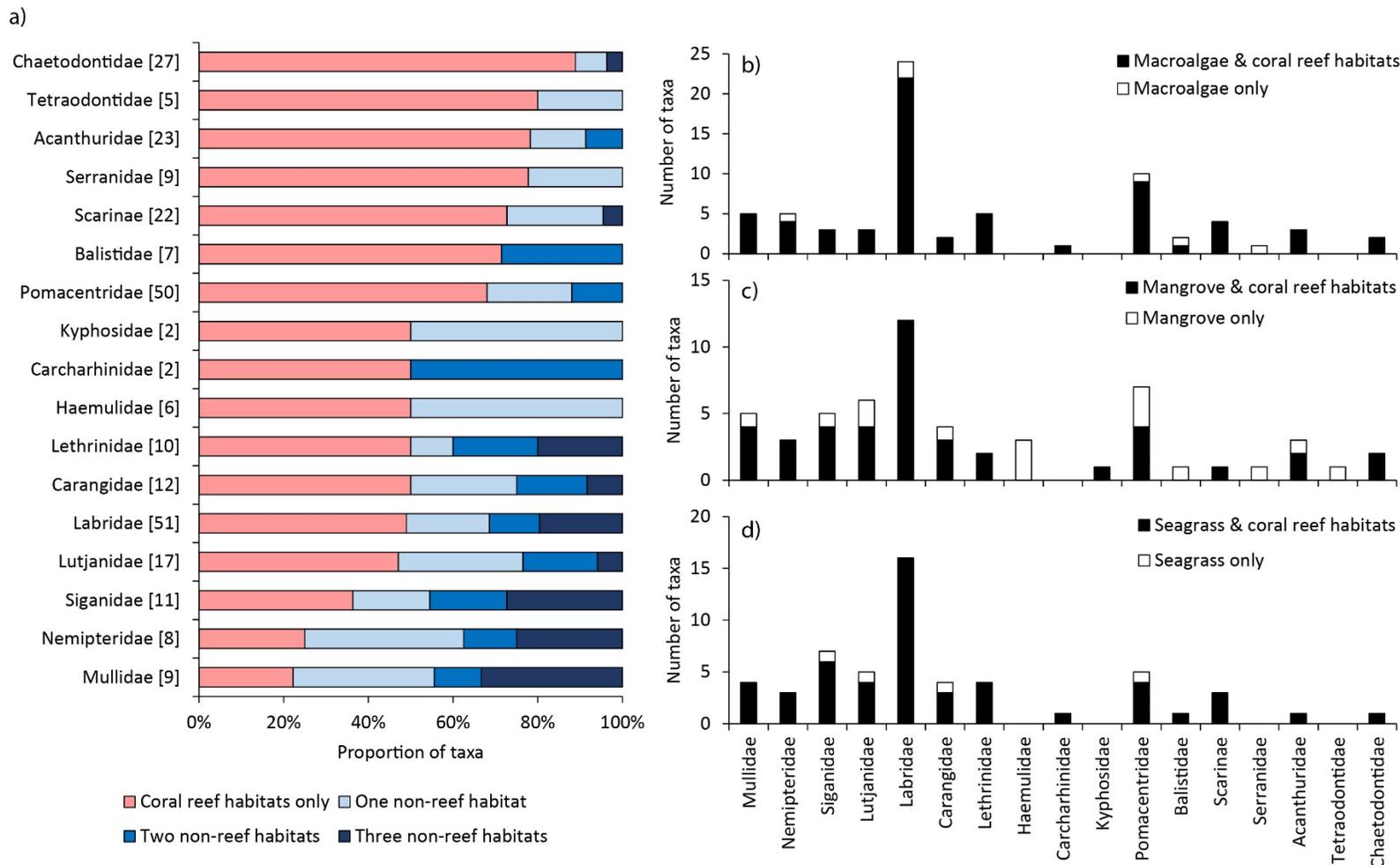
Two-thirds of the 41 families recorded during the surveys contained species that used non-reef habitats (27 families). Ten of these reef fish families contained a high proportion ( $\geq 50\%$ ) of species that were recorded in non-reef habitats including jacks (Carangidae), rabbitfishes (Siganidae), snappers (Lutjanidae), emperors (Lethrinidae) and sweetlips (Haemulidae) (Figure 3.4a). Patterns of habitat use (i.e., type and number of habitats) varied both among and within families. For example, species of snapper (Lutjanidae) ranged from being only recorded in coral reef habitats to being observed in both coral reef habitats and all three non-reef habitat types. In addition, several families contained species that were not recorded from either of the coral reef habitats (Figure 3.4b-d).

### 3.4.3 *Species-level habitat use patterns*

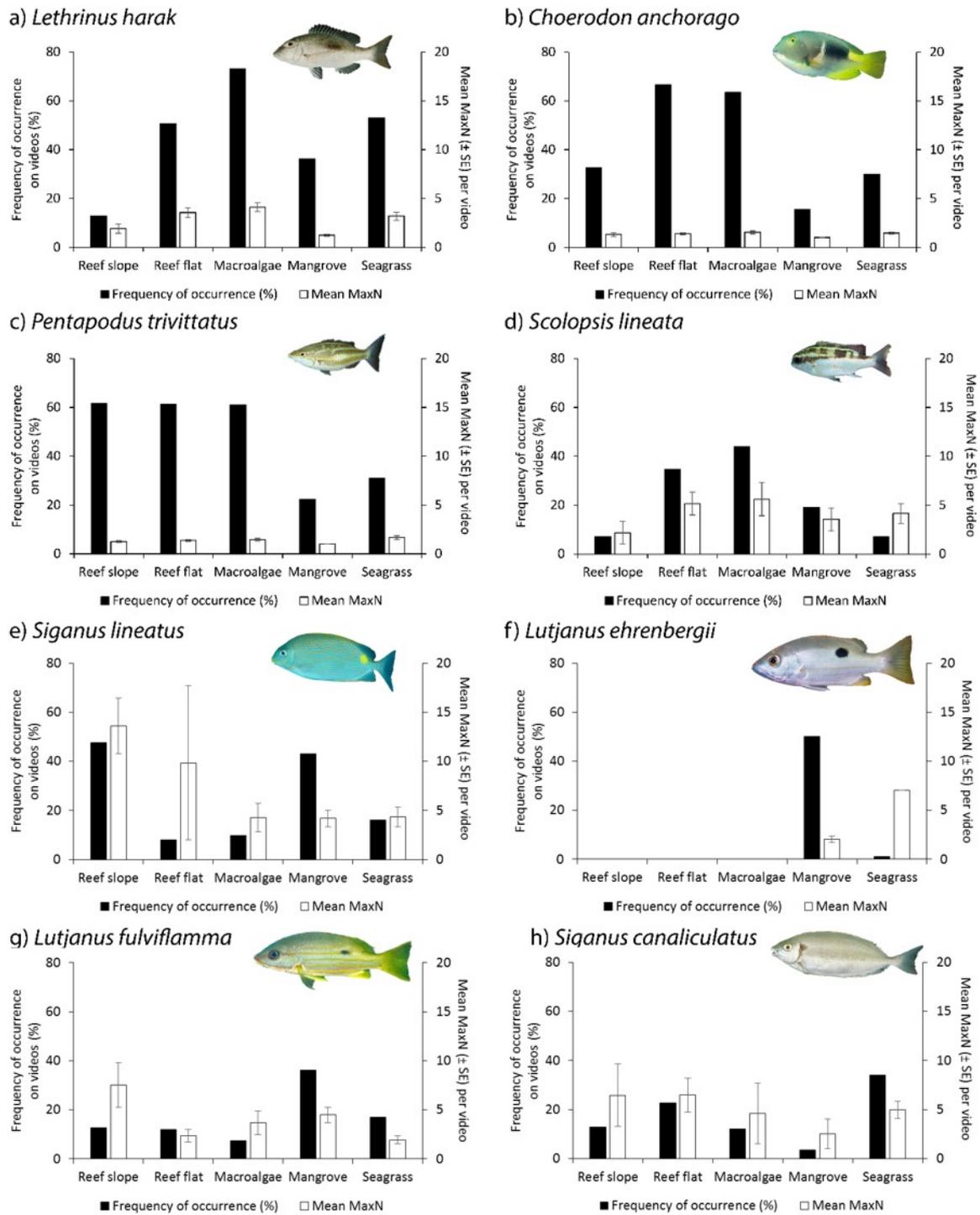
The data revealed a wide variety of habitat usage patterns by the 99 reef fish species that were observed in non-reef habitats. Over three-quarters of the species were recorded on multiple occasions away from coral reef habitats. In addition, over half were observed more frequently in at least one non-reef habitat compared to either of the coral reef habitat types (e.g., Figure 3.5a, d, f-h; Table A3.2), while others occurred in similar frequencies across a range of reef and non-reef habitats (e.g., Figure 3.5b-c, e). Eighteen species were identified as widespread multi-habitat users and occurred in all five habitat types (Table A3.2). These ranged from larger-bodied species such as the emperor *Lethrinus harak*, the snapper *Lutjanus fulviflamma*, and the rabbitfish *Siganus canaliculatus* to smaller-bodied species such as the butterflyfish *Chaetodon vagabundus*, and the wrasses *Halichoeres scapularis* and *Stethojulis strigiventer*.



**Figure 3.3.** a) Non-metric multi-dimensional scaling plot showing the variation in the community structure of reef fishes recorded from remote underwater video in five habitat types (reef flat, reef slope, macroalgae, mangrove and seagrass). Each point represents a single video based on MaxN data. Species where adults: b)  $\geq 25$  cm max length and c)  $< 25$  cm max length recorded on  $> 10\%$  of videos across all habitats or  $> 25\%$  of videos in a non-reef habitat type. Abbreviations of genera are *D.* = *Dischistodus*, *H.* = *Halichoeres*, *R.* = *Rhinecanthus*, *S.* = *Stethojulis*.



**Figure 3.4.** a) Habitat use patterns of common reef fish families showing proportion of taxa within each family only recorded from coral reefs and those recorded from one, two or three non-reef habitats. [x] denotes number of species observed in the study, b) reef fish taxa observed in macroalgal beds, c) mangroves, and, d) seagrass. Black bars represent species recorded in a non-reef habitat type and in coral reef habitats. White bars show the number of reef fish species that were not observed in coral reef habitats.



**Figure 3.5.** Species-specific habitat use patterns for eight common multi-habitat users. Black bars show the frequency of occurrence (%) on videos. White bars show the mean MaxN ( $\pm 1$  SE) calculated from videos where the species was recorded.

Comparing the relative abundance (i.e., MaxN) for some of the most frequently observed and abundant multi-habitat users highlighted potential among habitat differences in group size

(Figure 3.5). For instance, the mean relative abundance for *Lethrinus harak* was significantly lower in mangroves ( $1.24 \pm 0.10$  SE) compared to reef flat, macroalgae and seagrass habitats (ANOVA  $F_{4,158} = 4.68$ ,  $p=0.01$ ). The mean relative abundance for the rabbitfish *Siganus lineatus* was threefold higher on reef slopes compared to mangroves (ANOVA  $F_{1,64} = 6.39$ ,  $p=0.01$ ), despite occurring on a similar number of occasions in each habitat, indicating that *S. lineatus* may occur in larger groups on reef slopes. Similarly, the snapper *Lutjanus fulviflamma* had a higher mean relative abundance on reef slopes ( $7.55 \pm 2.26$  SE) compared to all other habitat types, almost double the mean relative abundance in mangroves ( $4.48 \pm 0.79$  SE).

### 3.5 Discussion

There is increasing evidence of the widespread use of non-reef habitats by reef-associated fishes, with a recent systematic review suggesting that ~20% of reef fish species use non-reef habitats (Sambrook et al. 2019, **Chapter 2**). By comparing fish assemblages across five habitat types in Kavieng, Papua New Guinea, this study found that percentage to be even higher, with over a third (35%) of reef-associated fish species recorded in non-reef habitats, many of which occurred in multiple non-reef habitats. In addition, many of the species identified using multiple habitats are ecologically (e.g., the macroalgae browser *Siganus canaliculatus*) or economically important. These findings thus provide additional support for claims of widespread use of multiple habitat types by reef fishes and for the importance of better understanding habitat complementarity in coral reef ecosystems. The study also demonstrates the value of examining species distributions across a wider range of habitats at each study location in the Indo-Pacific, as has previously been noted for the Caribbean (Nagelkerken et al. 2000a; Harborne et al. 2008). Observed species-specific patterns of habitat use would have been incomplete had a more limited range of habitats been sampled.

Macroalgal beds contained, on average, more species and higher relative abundances of reef fish compared to mangroves or seagrass meadows, albeit considerably lower than the two reef habitats. This supports the growing number of studies that have documented high abundances and diversity of coral reef fishes in macroalgae beds (e.g., Rossier & Kulbicki 2000; Wilson et al. 2010; Chaves et al. 2013; Evans et al. 2014; Eggertsen et al. 2017; Tano et al. 2017). The majority of these studies have focused on beds of canopy-forming macroalgae, such as *Sargassum* (but see Rossier & Kulbicki 2000), finding that the structural complexity and canopy height of algae are important factors influencing its use by coral reef fishes, particularly juvenile life stages (Wilson et al. 2014; Fulton et al. 2019; Fulton et al. 2020; Tang et al. 2020).

In contrast, the macroalgal sites in this study were predominantly a mixture of *Halimeda* and *Caulerpa*, which are smaller and less structurally complex than *Sargassum*. Despite these differences, a similar suite of families (e.g., Labridae, Lethrinidae, Siganidae) to that reported from *Sargassum*-based studies were recorded. This suggests that factors other than structural complexity of the algae, such as the availability of food resources, visibility of predators, and proximity to other habitat types (e.g., van Lier et al. 2018) may also influence the suitability of macroalgae beds for coral reef fishes. Although fish assemblages in macroalgal beds were typically more speciose than seagrass meadows or mangroves, there was considerable overlap in species using macroalgae beds and seagrass meadows, indicating these two habitat types could act as complementary habitats for some fishes (Dunning et al. 1992). Seagrass meadows and macroalgal beds can be structurally similar (Gratwicke & Speight 2005), and contain comparable resources and refugia opportunities (Macreadie et al. 2017) which could drive similarities in habitat use.

In contrast, mangroves contained a distinct assemblage of reef fishes including some species (e.g., *Lutjanus argentimaculatus*) that were exclusive to this habitat. Although the methodology precluded the separation of the data into life stages based on body size, several individuals, particularly snappers (i.e., lutjanids), were observed with juvenile markings within mangrove habitats. These findings are interesting, but require additional research, given the debate about whether mangrove habitats in the Indo-Pacific are as important for coral reef fishes, particularly juveniles, as they are in the Caribbean (Blaber & Milton 1990; Thollot 1992; Dorenbosch et al. 2005a; Nakamura et al. 2008; Unsworth et al. 2009; Barnes et al. 2012; Kimirei et al. 2013a; Dubuc et al. 2019). Mangrove systems in the Indo-Pacific vary considerably depending on tidal regime (Krumme 2009; Igulu et al. 2014), geomorphological and spatial context (Blaber 2007; Unsworth et al. 2008; Olds et al. 2013; Bradley et al. 2019), as well as the size, composition and structural complexity of mangroves forests (Laegdsgaard & Johnson 2001; Nanjo et al. 2014), all of which can influence habitat use patterns (Sheaves 2017). Here, non-estuarine mangroves in a microtidal location were examined and it appears that under these conditions, mangroves might possess certain attributes (e.g., refuge, food availability) that are not provided by the other shallow non-reef, or even reef, habitats surveyed. Therefore the impact of mangrove loss or degradation could be greater for coral reef fishes that appear to selectively use mangroves compared to species that appear to use multiple habitats interchangeably.

Importantly, a substantial number (>50%) of the species using multiple habitat types belong to families (e.g., Carangidae, Lethrinidae, Lutjanidae, and Siganidae) caught by small-scale fisheries in the region (Papua New Guinea National Fisheries Authority 2005, 2007). Reef fishes have historically been, and continue to be, an important source of animal protein for Pacific Island communities (Dalzell et al. 1996; Pinca et al. 2012). However increasing human populations have placed pressure on reefs, and many island nation reef fisheries are considered to be operating at unsustainable levels (Newton et al. 2007). As reefs become more degraded, it has been suggested that the availability of non-reef habitats could play an important role in maintaining the productivity of reef fisheries (Rogers & Mumby 2019). Fishing around the Tigak Islands is largely restricted to inshore waters surrounding the islands, particularly during the monsoon season (Lawless & Frijlink 2016), with the only commercial fishing targeting tuna and other pelagic fishes in offshore waters. My findings suggest that this dependence on inshore waters for several months each year, combined with growing populations, requires careful management of the entire seascape, not just reefs, by local communities to protect food security into the future.

“Coral reef fishes” is a widely used term to describe fish assemblages that occupy waters in the vicinity of coral reefs, yet over one third of the fishes recorded from shallow-water habitats in the Kavieng Lagoon were present in one or more non-reef habitats. Many of these species were frequently encountered away from coral reef habitat and could be considered as “seascape users” or “habitat generalists” as opposed to “coral reef fish”. Terminology aside, being flexible in habitat use could be advantageous given the widespread degradation of many shallow-water coastal habitats. As has been demonstrated from terrestrial landscapes (e.g., in birds, Salido et al. 2011), populations of such habitat generalists might be less vulnerable to the degradation of one habitat type. In contrast, species that obtain complementary resources from different habitats (e.g., food vs. shelter, Ries et al. 2004), may be negatively impacted by habitat disturbance or loss. However, the drivers of multiple habitat use are not well understood for many of the species identified here as multi-habitat users.

It is now widely recognised that coral reefs are moving into uncertain territory. However, efforts to predict how reefs might function in the future rarely consider that many reefs are embedded within, and consequently influenced by, a mosaic of other habitat types. Such connections may become increasingly important in the future both for supporting key

ecological functions on reefs, and providing food security for nations with strong dependencies on coral reef fisheries.

## Chapter 4

### Relative importance of seascape versus within-habitat variables on the distribution of fishes in tropical seagrass beds <sup>3</sup>

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#### 4.1 Abstract

Indo-Pacific seagrass meadows, particularly those in the Coral Triangle, cover large areas of shallow water, harbour diverse fish communities and support coastal fisheries. These meadows are often located in complex seascapes containing a mosaic of habitat types. However, the relative influences of within-habitat variables (e.g., shoot density, canopy height) and the surrounding seascape (e.g., distance to coral reef or mangroves) in shaping fish communities in seagrass meadows are largely unknown. I used a multi-scale approach to examine the relative influence of within-habitat variables and seascape structure on the distribution patterns of 17 fish taxa (i.e., families and species), and life stages for two taxa, in 32 seagrass patches around the Tigak Islands, New Ireland, Papua New Guinea. Of the 21 fish groups examined, two-thirds were influenced more by seascape than within-habitat variables. Site-attached taxa were generally more influenced by within-habitat variables, while taxa with larger home ranges (e.g., rabbitfishes, snappers) and ontogenetic habitat-shifters (e.g., emperors) were generally more influenced by the surrounding seascape. Despite the majority of taxa being reef-associated, proximity to or area of adjacent coral reefs appeared to have limited influence on local abundances. Instead, proximity and area of hardground, macroalgae and mangroves emerged as more influential. The findings show that individual seagrass patches are not functionally equivalent and that the surrounding seascape is important in shaping tropical fish assemblages in seagrass beds. Adopting a multi-scale approach to examine species-habitat relationships will provide a more nuanced perspective, improving our ability to predict which seagrass patches are highly productive or important nurseries which can contribute towards to more refined spatial management strategies.

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<sup>3</sup> Data from this chapter has been submitted as: **Sambrook K**, Andréfouët S, Aston EA, Bonin MC, Cumming GS, Duce S, Sievers KT, Hoey AS (in revision) Relative importance of seascape versus within-habitat variables on the distribution of fishes in tropical seagrass beds. *Estuarine, Coastal and Shelf Science*

## 4.2 Introduction

Understanding how distribution patterns of organisms are influenced and at what spatial scales is a fundamental goal of ecology (Levin 1992; Schneider 2001). An organism's habitat requirements are often dependent on traits such as body size, mobility, behavior, and vulnerability to predation, and these requirements can change during a species' lifecycle (Mittelbach 1981; Rayor & Uetz 1993; Nakazawa 2015). Furthermore, individuals may respond to variables operating across different spatial scales, ranging from within-habitat variables (e.g., structural complexity, food availability) to land- or seascape variables (e.g., composition and spatial configuration of habitats or patches surrounding a focal patch; Menge & Olson 1990; Dunning et al. 1992; Ries et al. 2004; Mayor et al. 2009; Thornton et al. 2011). Consequently, selecting ecologically meaningful scales at which to analyse influences on local distribution patterns can be challenging (Wiens 1989; Wheatley & Johnson 2009; Jackson & Fahrig 2015). One widely applied solution is to adopt a multi-scale approach that incorporates both within-habitat and broader land- or seascape variables (Kotliar & Wiens 1990; Kendall 2005; Mellin et al. 2009; Yeager et al. 2011).

Tropical shallow marine environments are composed of a mosaic of different habitat patches (e.g., coral reefs, mangroves, macroalgae beds, and seagrass meadows) interspersed across the seascape. Many fishes are known to use multiple habitats for a variety of reasons including short-term movements to foraging or spawning sites, and ontogenetic shifts among habitats (Adams et al. 2006; Appeldoorn et al. 2009; Nagelkerken 2009; Sambrook et al. 2019, **Chapter 2**). Although the use of multiple habitats by fishes has long been recognised (Ogden 1976; Parrish 1989), scaling up research from single habitats to incorporate the composition and configuration of multiple habitat types and patches in the surrounding seascape has begun to reveal the complexities of fish-habitat relationships (Grober-Dunsmore et al. 2009; Pittman & Brown 2011; Rees et al. 2018; van Lier et al. 2018). To date, the majority of this research in tropical marine environments has focused on coral reefs. It has found that seascape variables such as proximity to mangrove forests, seagrass meadows, and/or macroalgal beds can influence the distribution of some reef fish taxa on adjacent reefs (e.g., Dorenbosch et al. 2005b; Grober-Dunsmore et al. 2008; Berkström et al. 2013a; Martin et al. 2015). This is particularly evident for taxa that undertake ontogenetic shifts between habitats, and/or species with larger home ranges (e.g., sweetlips, snappers, emperors) that move between foraging and resting sites (Beets et al. 2003; Berkström et al. 2013b). In contrast, the distribution of more

site-attached (e.g., damselfishes) or sedentary (e.g., groupers) taxa appears to be better explained by within-habitat variables (e.g., structural complexity, percent hard coral cover) (Grober-Dunsmore et al. 2007). For those fish taxa whose populations are influenced by the surrounding seascape, proximity to or area of seagrass meadows has often been identified as important (Dorenbosch et al. 2005b; Kendall 2005; Grober-Dunsmore et al. 2007; Olds et al. 2014).

Seagrasses occur in temperate and tropical ecosystems in both estuarine and marine environments. In temperate and tropical Atlantic systems, seagrass meadows are well-established as important fish habitat, particularly because of their role as nurseries for several key fisheries species (e.g., Atlantic cod (*Gadus morhua*), Lilley & Unsworth 2014; Tuya et al. 2014; McDevitt-Irwin et al. 2016). Until relatively recently, the value of seagrass habitat for fishes in tropical Indo-Pacific seascapes had received less attention (Orth et al. 2006; Duarte et al. 2008; Unsworth & Cullen 2010). Emerging social-ecological research has highlighted the significance of Indo-Pacific seagrass habitat in sustaining small-scale fisheries (de la Torre-Castro et al. 2014; Unsworth et al. 2014; Nordlund et al. 2018; Brodie et al. 2020), and hence its likely importance for populations of some tropical fishes.

The Indo-Pacific region is known for its high seagrass diversity, particularly in Pacific small island developing states such as Papua New Guinea and the Solomon Islands (Short et al. 2007; Waycott et al. 2009). In these areas the composition and structure of seagrass beds can be highly variable, even across relatively small areas (10s to 100s of metres) (McKenzie et al. 2020). While some studies have shown the importance of within-habitat features (e.g., canopy height, shoot density) in shaping fish assemblages within seagrass beds (Bell & Westoby 1986; Hyndes et al. 2003; Boström et al. 2006; Horinouchi 2007; Pogoreutz et al. 2012; Yeager & Hovel 2017), others have demonstrated the importance of neighbouring habitat types (e.g., coral reefs: Dorenbosch et al. 2005a; mangroves: Unsworth et al. 2008). Studies examining the relative influence of both within-habitat and seascape variables on fish distribution patterns in seagrass meadows are rare and confined to temperate systems (e.g., Staveley et al. 2017; Scapin et al. 2018; Olson et al. 2019) or the tropical Indian Ocean (Gullström et al. 2008; Aller et al. 2014). Given the potential ecological and economic value of many fishes that use tropical Indo-Pacific seagrass beds, identifying the variables that best predict their distribution merits greater consideration (Sambrook et al. 2020, **Chapter 3**). As well as improving our understanding of species distribution patterns, adopting a multi-scale approach has considerable value for spatial

management strategies (e.g., reserve networks) that typically operate at broader spatial scales (i.e., 10s to 100s km<sup>2</sup>).

I used a multi-scale approach to examine the relative influence of within-habitat variables and seascape structure on distribution patterns of fishes in seagrass beds in the Tigak Island Group, New Ireland Province, Papua New Guinea. Based on the underlying hypothesis that species occurrences will reflect a full set of resource needs, I predicted that within-habitat structure would exert a greater influence on the distribution patterns of less mobile and/or site-attached fish taxa (e.g., damselfishes), whereas the surrounding seascape would be more influential on taxa with larger home ranges and/or taxa that shift between habitats during ontogeny (e.g., emperors, snappers).

### **4.3 Methods**

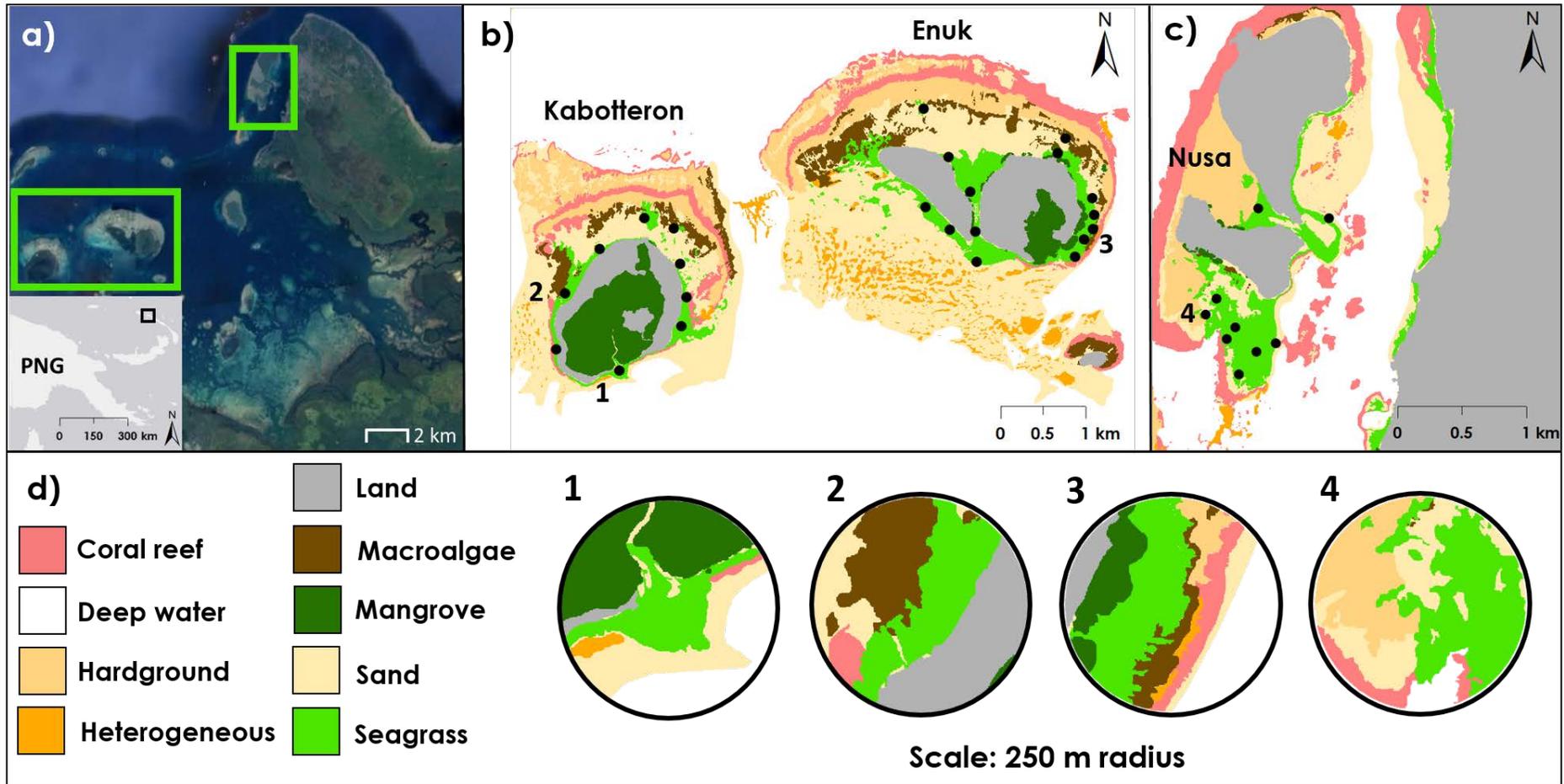
#### ***4.3.1 Study location***

The study was conducted around three island complexes (i.e., Eruk, Kabotteron, and Nusa) of the Tigak Island Group, Kavieng, New Ireland Province, Papua New Guinea (Figure 4.1). Each island has extensive seagrass meadows, with distinct patches, often in close proximity, that vary in their species composition, structural attributes (e.g., canopy height, shoot density, percent cover), and water depth. In addition to large expanses of seagrass, each island has a range of other habitat types (e.g., coral reefs, non-estuarine mangroves, macroalgal beds, sand) that vary in size and spatial configurations. This presents an ideal system to examine the influence of seascape and within-habitat variables on fish taxa in seagrass beds. The tidal regimen is mixed microtidal (Krumme 2009), with a tidal range of ~1.09 m.

#### ***4.3.2 Habitat mapping***

A benthic habitat map for the seascape surrounding each island was produced using high-resolution multispectral Worldview 2 imagery acquired in 2017 (Figure 4.1). The mapping process followed the user-oriented workflow described in Andréfouët (2008). Specifically, polygons of homogeneous colour and texture were photo-interpreted against the satellite images, and labelled based on a preliminary survey that investigated the diversity of shallow habitats throughout the Kavieng area in August 2014 (Andréfouët, unpublished data). Extensive ground-truthing using geo-referenced photographs (n = 1,470 photographs across 126 sites) was subsequently undertaken in September 2017 to verify the accuracy of initial habitat classifications. When necessary, the initial maps were corrected using the 2017 ground-

truthing data to minimise any errors in habitat identification that could bias conclusions. I focused on eight broad benthic habitat categories: coral, deep water, hardground, heterogeneous, macroalgae, mangrove, seagrass and sand (Table 4.1). As benthic habitat maps derived from satellite imagery are restricted to shallow-water only (<20 m), I classified any unmapped regions surrounding islands as 'deep water'. The classification 'hardground' included areas dominated by reef pavement, rock or rubble, and 'heterogeneous' encompassed predominantly small low relief patches of mixed composition including rubble, hard and soft coral, seagrass and algae. In addition, the high spatial resolution of the maps allowed for the identification of unique spectral signatures for different types of seagrass patches, predominantly based on seagrass density.



**Figure 4.1.** a) Map of Kavieng, New Ireland Province, with inset showing location within Papua New Guinea; b & c) the three focal island complexes and surrounding benthic habitats. Black circles denote the 32 seagrass survey sites, d) examples of different seascapes surrounding seagrass survey sites based on 250 m radii.

**Table 4.1.** Predictor variables used to examine the influence of within-habitat and seascape variables on the distribution of fishes in seagrass beds.

<b>Metric</b>	<b>Description</b>	<b>Unit</b>	<b>Mean</b>	<b>SE</b>	<b>Range</b>
<i>Within-habitat variables</i>					
Depth	Measured at start & end of each transect	m	0.80	0.04	0.25-2.30
<b>Transect (25 m)</b>					
Coral	Percent cover	%	1.8	0.58	0-39.22
Macroalgae	Percent cover	%	12.66	1.58	0-62.75
Rock	Percent cover	%	1.35	0.47	0-27.45
Rubble	Percent cover	%	0.61	0.29	0-21.57
Sand	Percent cover	%	14.30	1.80	0-80
Seagrass	Percent cover	%	69.26	2.59	5.88-100
<b>Quadrat (5 x 0.25 m<sup>2</sup> per 25 m transect)</b>					
Shoot density	Seagrass shoot density in 5 x 0.01 m <sup>2</sup> sub-quadrats		19.10	0.97	0.33-42.67
Canopy height	Tallest 5 seagrass blades in 5 x 0.01 m <sup>2</sup> sub-quadrats	cm	13.64	0.97	4.03-60.68
Algae	Mean percent cover	%	15.78	1.51	0-60
Seagrass	Mean percent cover	%	41.91	2.84	0-96.67
Epiphytes	Mean percent cover of epiphytes on seagrass blades	%	34.53	2.01	10-90
<i>Seascape variables</i>					
<b>Distance to nearest patch</b>					
Coral reef	Measured using Near function in ArcGIS. Edge-to-edge distance from seagrass patch to X habitat type avoiding crossing land	m	256.41	25.26	4.76-1041.37
Hardground		m	520.07	51.80	11.05-1728.45
Heterogeneous		m	437.25	32.16	17.87-1168.57
Macroalgae		m	457.37	48.42	0-1771.61
Mangrove		m	415.25	34.88	17.34-1368.30
<b>% area of habitat in 100, 250, 500 &amp; 1000 m radii (shown for 100 m scale only)</b>					
Coral reef		%	3.84	0.81	0-27
Deep water		%	0.88	0.28	0-11
Hardground		%	3.09	0.82	0-30
Heterogeneous		%	0.38	0.15	0-6
Macroalgae		%	5.56	1.14	0-44
Mangrove		%	1.88	0.63	0-34
Seagrass		%	56.06	2.57	0-100
Sand		%	22.34	2.28	0-80
<b>Habitat richness</b>					
100 m	Number of habitat types within buffer	N	3.06	0.10	1-6
250 m		N	4.38	0.14	2-7
500 m		N	5.72	0.11	3-7
1000 m		N	6.66	0.07	5-7

### **4.3.3 Surveys**

I surveyed fish and benthic communities in 32 seagrass patches around the three island complexes (Figure 4.1) during April and May 2018. Seagrass patches were selected *a priori* from the initial benthic habitat map to represent a range of seagrass densities and seascape attributes (e.g., distance to mangroves). In each seagrass patch, fish and benthic communities were quantified along three replicate 25 m transects, with adjacent transects separated by a minimum of 10 m. Transects were positioned parallel to shore and placed  $\geq 5$  m away from the boundary of each patch to reduce potential confounding effects of edge habitats on fish communities. All surveys were carried out during daylight hours between 10:00 and 16:00.

All non-cryptic fishes were surveyed in seagrass patches using belt transects. Fish  $> 10$  cm total length (TL) were identified, their TL estimated to the nearest 5 cm, and recorded within a 5 m wide belt as the transect was being placed to minimise any potential diver effects on fish counts. Fish  $< 10$  cm were identified, their TL estimated to the nearest 2.5 cm, and recorded within a 2 m wide belt during a return swim along the transect (Thompson & Mapstone 2002). Individuals were identified to genus or species where possible, and abundances standardised to 250 m<sup>2</sup>.

### **4.3.4 Within-habitat structure**

Within-habitat variables were quantified along the same transects used to survey the fish community. I first quantified benthic composition using point-intercept transects, recording the substratum directly below the transect tape at 50 cm intervals (51 points per transect). The substratum was classified as seagrass, macroalgae, live coral, sand, rubble, or rock/pavement. Secondly, I placed five 0.25 m<sup>2</sup> quadrats at random points along each transect and estimated percent cover of seagrass, macroalgae and the percent epiphyte cover on seagrass blades. Within each quadrat, I also surveyed five smaller 'sub-quadrats' (0.01 m<sup>2</sup>) to quantify the shoot density (i.e., number of shoots) and canopy height (i.e., height of the five longest seagrass blades per sub-quadrat) (Aller et al. 2014; Eggertsen et al. 2017). The depth at the start and end of each transect was recorded.

### **4.3.5 Seascape composition and configuration**

To explore the influence of seascape variables, I extracted spatial pattern metrics from the benthic habitat map using ArcGIS (v10.6.1). I quantified the proportion (area) of eight different habitat types (i.e., coral reef, deep water, hardground, heterogeneous, macroalgae, mangrove,

seagrass, sand) and the number of habitat types (i.e., habitat richness) within 100 m, 250 m, 500 m and 1000 m radii of the survey sites (Figure 4.1, Table 4.1). Given the limited information on home ranges for common fish species at my study sites, radii were selected based on previous studies that have related fish abundance or diversity in other habitat types with the presence of seagrass beds (e.g., Grober-Dunsmore et al. 2007; Olds et al. 2012a; Van Wynsberge et al. 2012; Berkström et al. 2013a). I also measured the proximity (edge-to-edge distance) of the seagrass patch to the nearest polygon of one of five different habitat types (i.e., coral reef, hardground, heterogeneous, macroalgae and mangrove) using the Near function in ArcGIS (Table 4.1). Where the shortest route between a seagrass patch and a different habitat type crossed land, the shortest edge-to-edge distance between patches avoiding land was measured manually.

#### **4.3.6 Spatial autocorrelation**

All response and predictor variables were tested for spatial autocorrelation using a standardised inverse distance weighted Global Moran's I with the threshold distance defined by the average distance from each site ( $n = 32$  sites) to its eight nearest neighbours (average distance = 1,394 m). For variables where significant (pseudo-significance  $P \leq 0.01$  calculated from 999 permutations) spatial autocorrelation was detected with a Moran's I value exceeding 0.4 (Hamylton & Barnes 2018), spatial patterns were further examined using cluster and outlier analysis (Anselin Local Moran's I) in ArcGIS. Two predictor variables at the seascape scale exhibited positive spatial autocorrelation, specifically, area of coral reef (Moran's I = 0.57,  $P < 0.001$ ) and area of mangroves (Moran's I = 0.61,  $P < 0.001$ ) within 1000 m radius. Cluster and outlier analysis revealed that these values were driven primarily by clustering of high values around the island of Nusalik that had a small area of mangroves and a larger area of coral reef habitat compared to sites around the other two islands (i.e., Eruk and Kabotteron). Only three of the 22 response variables were weakly spatially autocorrelated: adult parrotfishes (Scarinae; Moran's I = 0.182,  $P = 0.003$ , *Siganus spinus* (Moran's I = 0.251,  $P = 0.001$ ) and *Stethojulis strigiventer* (Moran's I = 0.165,  $P = 0.000$ ) (Table A4.2). The limited evidence of spatial autocorrelation in the data suggests that survey sites were spaced beyond the zone of spatial dependence (Hamylton 2017; McGarvey et al. 2010) and thus provide a representative characterisation of the spatial heterogeneity of the seascape.

#### 4.3.7 Data analysis

I used boosted regression trees (BRTs) to evaluate the influence of 12 within-habitat and 14 seascape predictor variables (Table 4.1) on the abundance and/or presence of each of 21 fish groups in seagrass beds (Table A4.1). BRTs are a powerful and increasingly popular technique to explore complex ecological datasets (Sekund & Pittman 2017) because they are capable of managing collinearity and can fit non-linear relationships (Elith et al. 2008). BRTs were fitted individually by fish group using functions from the *dismo* package (Hijmans & Elith 2017) in R v4.0.0 (R Core Team 2020).

Although BRTs are capable of managing multicollinearity, I followed (Zuur et al. 2010) and generated a collinearity matrix using Pearson's correlation coefficients in the R package *corrplot* to identify highly collinear ( $>0.7$ ) predictor variables (Figures A4.1 & A4.2). Where high collinearity was identified, I ran a reduced BRT containing only the collinear variables (e.g., area of sand and area of seagrass within 100 m radius) using the *gbm.step* function for each response variable. From the reduced BRT, I retained the predictor variable with the highest relative influence for inclusion in subsequent modelling steps. Models were fitted with a Poisson distribution for count data for fish groups where  $>1,000$  individuals were recorded across all surveys (e.g., parrotfishes, rabbitfishes, snappers, wrasses). Data for less abundant taxa (e.g., threadfin breams, goatfishes) were modelled using a Bernoulli distribution for presence.

An initial model was generated for each of the 21 fish groups allowing all non-collinear predictor variables to be evaluated using the *gbm.step* function in the *dismo* package. Following this initial model run, I used the function *gbm.simplify* (Elith & Leathwick 2017) to identify uninformative predictor variables. This process involves an iterative backwards stepwise removal using 10-fold cross-validation to identify the number and identity of predictor variables that can be removed from the model without affecting the model's predictive performance. Finally, I ran these simplified models through an optimisation loop in which three parameters were varied: bag fraction (0.5, 0.7, 0.8), learning rate (0.001, 0.0001, 0.00001) and tree complexity (1-5). Optimising these three parameters and using slow learning rates helps to counter overfitting issues associated with the use of multiple predictor variables and the generation of large numbers of trees (Elith et al. 2008). This optimisation process enabled the identification of the combination of parameters that produced the 'best' or lowest cross validation deviance (Aston et al. 2019) with the minimum number of trees set to 1,000 and

maximum set to 10,000 trees (Elith et al. 2008). Models that explained <20% deviance were deemed unsatisfactory and excluded from further analysis. In total, models are reported for 17 fish taxa (Table A4.1) including family and species/genera, as well as juveniles and adults from two families (i.e., emperors and parrotfishes). Emperors and parrotfishes were the only taxa where adults and juveniles were easily distinguishable and occurred with sufficient frequency to explore distribution patterns discretely.

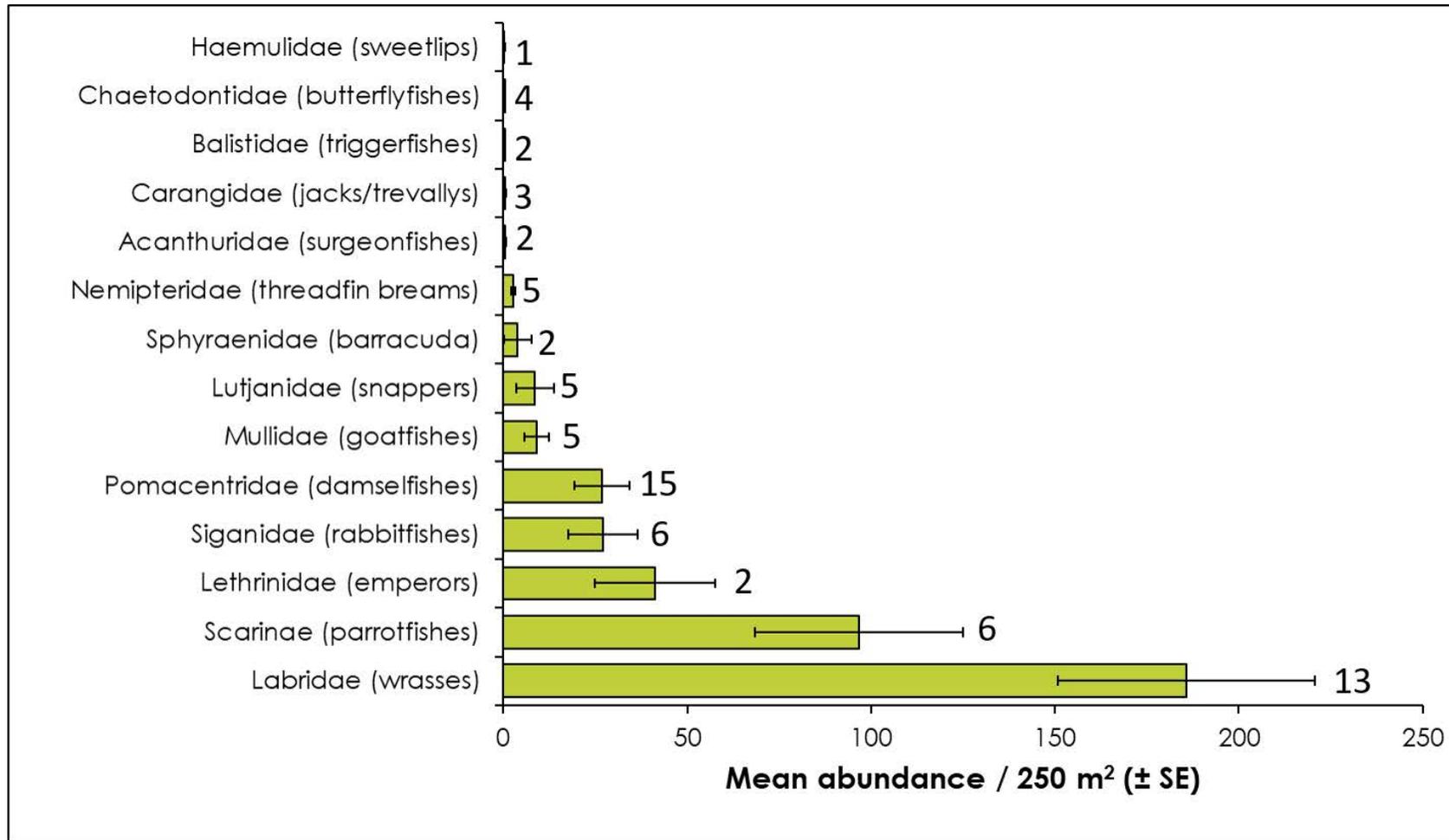
BRTs compute the relative influence of each predictor variable, resulting in a cumulative contribution of predictors summing to 100%. Influential predictor variables are identified by those selected in models more frequently than would be expected by chance. A higher relative influence (%) is therefore indicative that a predictor variable is influencing the response variable under investigation (Elith et al. 2008). On average,  $5.9 \pm 0.25$  SE predictor variables were identified per model as influencing the distribution patterns (abundance or presence) of particular fish groups. For all subsequent analyses, I extracted only those predictor variables identified as more influential than expected by chance. Relationships between a response variable and influential predictor variables were visualised with partial dependency plots using the `gbm.plot` function (Elith & Leathwick 2017).

## **4.4 Results**

A total of 9,746 individuals were recorded from 71 species (or species groups) representing 14 families (including the Scarinae; i.e., parrotfishes) across the 32 seagrass patches. Wrasses (Labridae) were the most abundant taxa, accounting for 50% of all individuals recorded, followed by parrotfishes (22%) and emperors (Lethrinidae: 9.5%; Figure 4.2). The majority of parrotfishes and emperors recorded were juveniles.

### **4.4.1 Seascape vs within-habitat variables**

The influence of seascape and within-habitat variables varied between fish groups. Of the 21 fish groups (e.g., family, species and life stage) examined, 13 (62%) were more strongly influenced by the surrounding seascape compared to within-habitat predictor variables, conversely within-habitat variables were more influential for the remaining eight groups.



**Figure 4.2.** Mean abundance of individuals / 250 m<sup>2</sup> by family of selected taxa observed across 32 seagrass patches in Kavieng, Papua New Guinea. Note: parrotfishes from the subfamily Scarinae are presented separately from the Labridae family. Numbers above bar show number of taxa recorded in each family.

#### 4.4.2 Family responses

Of the five families for which patterns of abundance were analysed, three families (i.e., rabbitfishes, wrasses, snappers) were primarily influenced by seascape variables while parrotfishes and damselfishes were more influenced by within-habitat attributes (Figure 4.3). For rabbitfishes and wrasses the influence of seascape characteristics on abundance was particularly strong, accounting for 82% and 76% of the most influential predictor variables. Specifically, the abundance of wrasses was predominantly influenced by distance to and area of macroalgae (Figure 4.3), being positively related to the proportion of macroalgae within 250 m. In contrast, rabbitfishes and snappers were primarily influenced by the proximity of mangroves. The abundance of both parrotfishes and damselfishes were positively related to the cover of live coral within seagrass patches.

Of the three families for which patterns of presence were analysed, emperors and threadfin breams were more strongly influenced by seascape variables (87% and 67% respectively), whereas goatfishes were more influenced by within-habitat characteristics. Threadfin breams were more likely to be present in seagrass patches proximal to mangroves, whereas emperors were less likely to be present as the proportion of mangrove within 500 m increased. Water depth was the most important predictor for goatfish presence (Figure 4.3), with this taxon more likely to be present as depth increased. All three families were influenced by distance to hardground and were less likely to be present when the seagrass patch was adjacent (<100 m) to hardground.

#### 4.4.3 Species responses

Due to their high relative abundance, responses were compared for three species of wrasse (*Halichoeres trimaculatus*, *H. scapularis* and *Stethojulis strigiventer*), and a group of ‘other’ *Halichoeres* (including *H. argus*, *H. nebulosus* and *H. nigrescens*) that were difficult to differentiate in the field (Figure 4.4). The relative influence of seascape versus within-habitat variables on the abundance of these four wrasse taxa varied considerably (Figure 4.4). ‘Other’ *Halichoeres* responded most strongly to seascape variables (92%), with abundance negatively related to the area of sand within 100 m and positively related to either near (<100 m) or far (>1000 m) patches of macroalgae (Figure 4.4). In contrast, *H. trimaculatus* was more influenced by within-habitat variables (78%), particularly depth, percent cover of seagrass at the quadrat level, and canopy height (Figure 4.4). Seascape and within-habitat variables exerted broadly similar influences on the abundance of *H. scapularis* (43% vs. 57%) and *S. strigiventer*

(53% vs. 47%). *H. scapularis* abundance was negatively related to the canopy height of seagrass and positively related to the area of macroalgae within 250 m whereas *S. strigiventer* was negatively related to the area of mangrove habitat within 1000 m and positively related to the cover of macroalgae at the transect level.

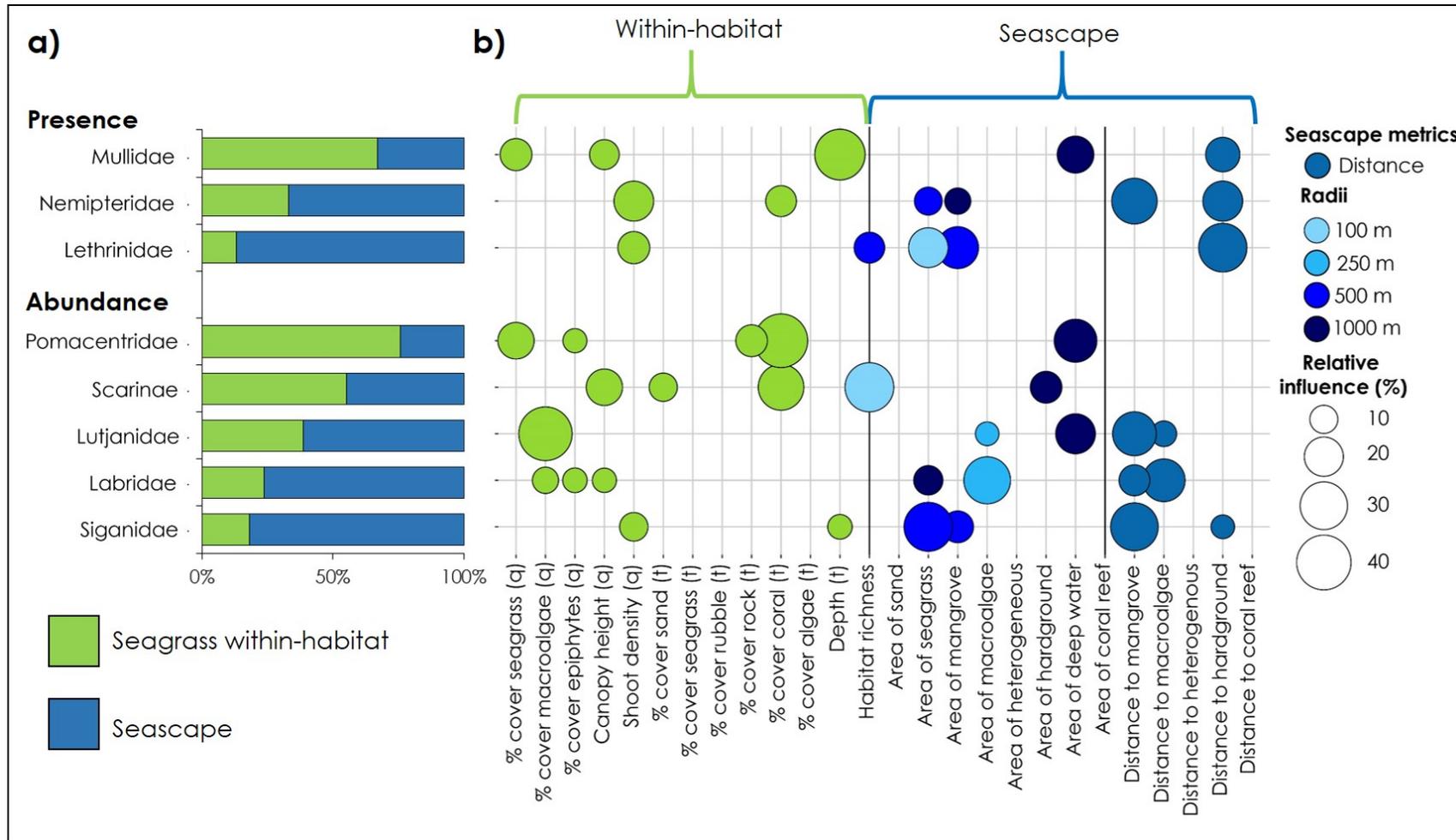
The influence of seascape and within-habitat variables on the presence of three species of rabbitfishes were broadly similar (i.e., influence of seascape: *Siganus spinus* 64%, *S. canaliculatus* 60%, and *S. lineatus* 51%; Figure 4.4). Despite subtle differences in the relative importance of individual variables, the presence of all three species was influenced by distance to hardground (Figure 4.4), with *S. canaliculatus* and *S. spinus* being negatively related and *S. lineatus* positively related to the proximity of hardground. All three species were also influenced by the cover of macroalgae at the quadrat scale, however the nature of the relationship differed with *S. spinus* being positively related and, *S. canaliculatus* and *S. lineatus* negatively related to macroalgal cover. The presence of *Lutjanus fulviflamma* was more influenced by seascape than within-habitat variables (67% vs 33%), and was more likely to be present with increasing distance from coral reef and macroalgal habitat, and as the cover of seagrass increased. The presence of *Lethrinus harak* was more influenced by within-habitat variables (65%), and was positively related to the cover of seagrass and negatively related to the cover of macroalgae and sand within a seagrass patch.

#### **4.4.4 Responses by life stage**

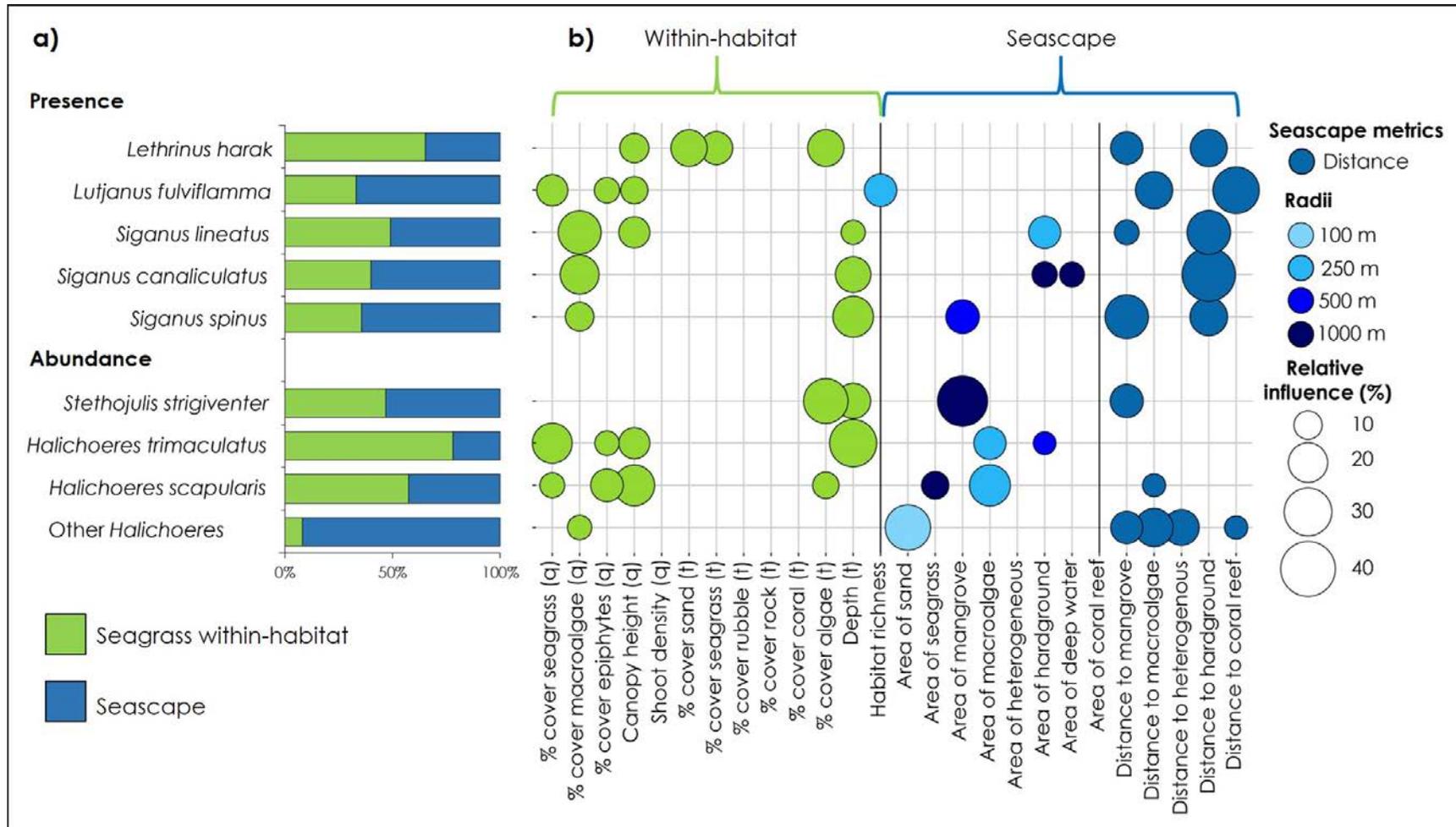
Within-habitat variables had a greater influence on the presence of both adult and juvenile parrotfishes in seagrass beds (55% and 66% respectively) compared to seascape variables, however the relative importance of individual variables differed between life stages (Figure 4.5). Transect-level coral cover, and to a lesser extent water depth, were the important predictors for adult parrotfish presence, whereas quadrat-level macroalgae cover, seagrass density and epiphyte cover on seagrass blades were the most important predictors of juvenile parrotfish presence with juveniles being more likely to be present as these variables increased (Figure 4.5; Figure A4.3).

In contrast to parrotfish, the presence of adult and juvenile emperors were more influenced by seascape compared to within-habitat predictor variables (Figure 4.5). Juvenile emperors were more likely to occur with either low (<10%) or high (>45%) proportions of deep water habitat within 1000 m, and as depth of the seagrass patch increased but less likely to occur as the amount of coral reef within a 250 m radius increased (Figure A4.4). Adult emperors were more

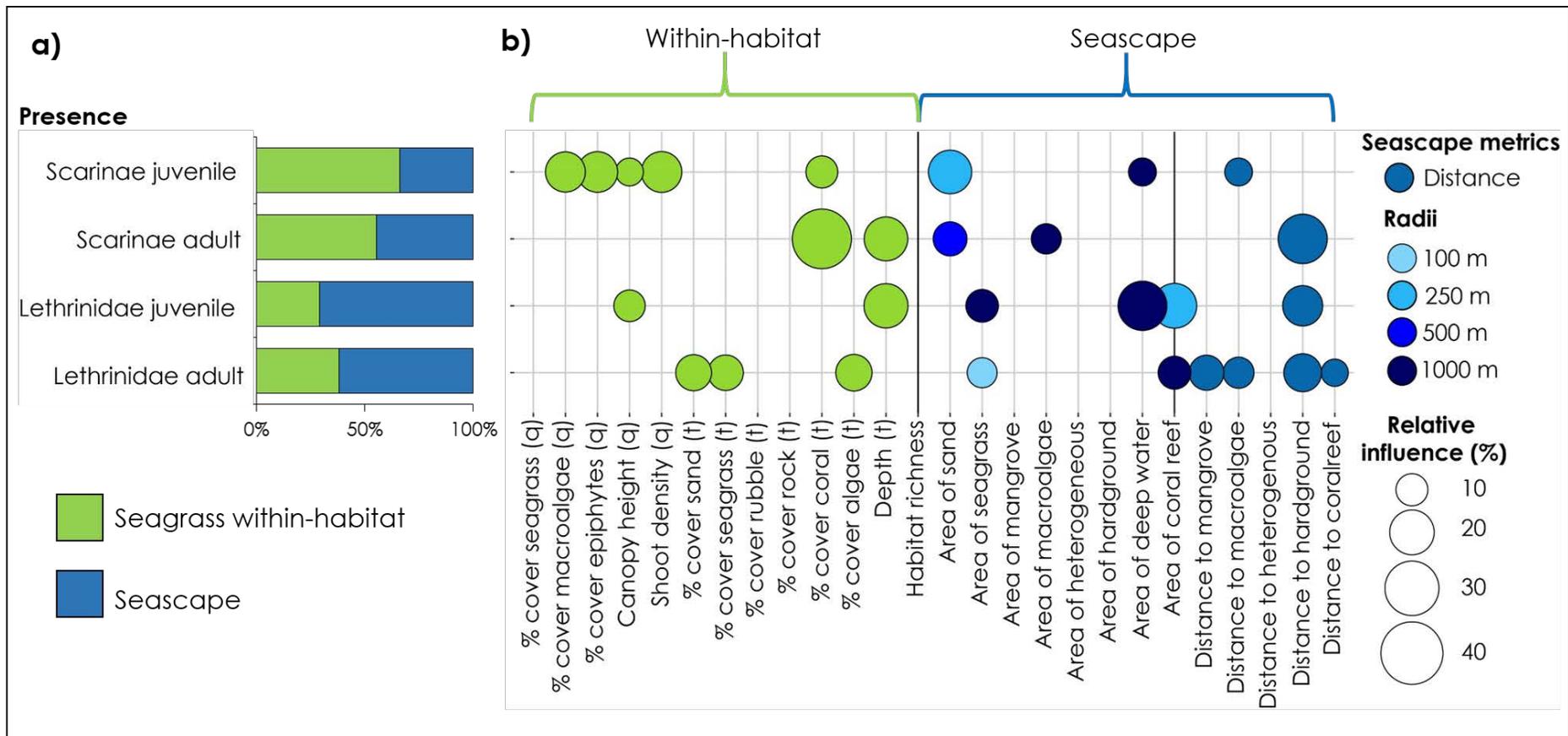
likely to be present in seagrass patches as the proportion of coral reef habitat within 1000 m increased.



**Figure 4.3.** Relative influence of predictor variables contributing to the distribution of fish taxa in seagrass beds more than would be expected by chance: a) proportion of within-habitat and seascape variables and, b) relative influence of individual predictor variables. Circle sizes represent % relative influence. Green shows within-habitat and blue shows seascape predictor variables. Shades of blue represent different spatial scales identified as important. ‘q’ denotes variables quantified at quadrat level, and ‘t’ variables quantified at transect level.



**Figure 4.4.** Relative influence of predictor variables contributing to fish species or species group abundance or presence in seagrass beds more than would be expected by chance: a) proportion of within-habitat and seascape variables and, b) relative influence of individual predictor variables. Circle sizes represent % relative influence. Green shows within-habitat and blue shows seascape predictor variables. Shades of blue represent different spatial scales identified as important. ‘q’ means measured at quadrat level, ‘t’ at transect level.



**Figure 4.5.** Relative influence of predictor variables contributing to the presence of adult and juvenile emperors (family: Lethrinidae) and parrotfishes (subfamily: Scarinae) in seagrass beds more than would be expected by chance: a) proportion of within-habitat and seascape variables and, b) relative influence of individual predictor variables. Circle sizes represent % relative influence. Green shows within-habitat and blue shows seascape predictor variables. Shades of blue represent different spatial scales identified as important. ‘q’ means measured at quadrat level, ‘t’ at transect level.

## 4.5 Discussion

Almost two-thirds of the fish groups recorded in seagrass beds were more strongly influenced by surrounding seascape configuration than within-habitat variables. As predicted, site-attached taxa (e.g., damselfishes) were more influenced by within-habitat variables such as coral cover, while taxa with larger home ranges (e.g., rabbitfishes, snappers) and those known to undertake ontogenetic shifts (e.g., emperors) were generally more influenced by the surrounding seascape, such as distance to mangrove. Interestingly, however, the differing habitat relationships of closely related taxa (e.g., species of *Halichoeres*) with similar movement capacities (Wainwright et al. 2002) suggested that underlying ecology, not simply mobility, also influences how species respond to the wider seascape.

Despite the majority of fish taxa recorded being considered to be reef-associated, proximity to or area of adjacent coral reefs appeared to have limited influence on fish distribution patterns within seagrass patches. Instead, other seascape features such as proximity and area of hardground, macroalgae and mangroves emerged as more influential. These results could be interpreted in two different ways. One perspective could be that these fish are less dependent on coral reefs than previously assumed. An alternative view could be that of hierarchical control, where the presence of these fish are determined by coral reef habitat in the surrounding seascape, but the nuanced dynamics of fish abundance and assemblage structure is subsequently driven by the within-habitat characteristics of seagrass and other aspects of the surrounding seascape.

I predicted that home range area and level of site attachment would influence how taxa responded to seascape and within-habitat variables. Variation in the abundance of damselfishes among seagrass patches was consistent with the hypothesis that site-attached, less mobile species would be shaped more by within-habitat variables. Damselfish typically settle directly to their adult habitat and remain in close association with their settlement site throughout juvenile and adult life stages (e.g., McCormick & Makey 1997). The abundance of damselfish within seagrass patches was primarily influenced by the amount of hard structure (i.e., coral and rock) within the seagrass patches. This is consistent with studies of damselfish on coral reefs that have demonstrated the importance of the structural complexity of habitats in providing refugia from predation and shaping adult populations (e.g., Beukers & Jones 1997; Nemeth 1998). Therefore features at scales greater than a few metres from their settlement site may have limited influence on their local abundances.

In contrast to damselfishes, wrasses (i.e., Labridae) were strongly influenced by the surrounding seascape and were more abundant when macroalgal beds were in close proximity or occupied a large proportion of the seascape within 250 m of the focal seagrass patch. Of the 13 taxa of wrasses recorded from seagrass beds, the most abundant were from the genera *Halichoeres* and *Stethojulis*. While many species of wrasse within these genera are considered highly mobile (Wainwright et al. 2002), home range area estimates for taxa similar to the most abundant species from the present study (i.e., *Halichoeres scapularis*, *H. trimaculatus*, ‘other’ *Halichoeres* and *Stethojulis strigiventer*) are typically small (e.g., <10 m, Jones 2005; Overholtzer-McLeod 2005; Green et al. 2015). Given this information, the strong collective response of wrasses to seascape factors was initially unexpected. However, the findings are similar to research from other tropical and temperate seascapes that also found that the spatial arrangement of the surrounding seascape can influence wrasse distribution and abundance patterns (e.g., Staveley et al. 2017; van Lier et al. 2018; Sievers et al. 2020a). Most attempts to quantify home ranges of wrasses have been conducted in coral reef habitats, where ready access to food and suitable hard structure as refugia may be important given the high predation rates associated with coral reefs (Hixon & Beets 1993; Beukers & Jones 1997). Home ranges may vary between habitat types due to differences in food and refugia availability. For instance, in macrophyte habitats, where the major habitat-forming organisms (i.e., seagrass, macroalgae) are flexible, and densities of predators are lower (Nakamura & Sano 2004a; Aller et al. 2014), individuals may not be as tightly linked to specific refuge sites and therefore may move greater distances. In this context, the characteristics of the immediate local habitat may become less critical than the attributes of the surrounding seascape.

Macroalgal and seagrass beds harbour diverse and abundant invertebrate communities that serve as a potential food source for fishes (Heck & Wetstone 1977; Nakamura & Sano 2005), but the composition of those communities can be quite distinct (e.g., Tano et al. 2016). Proximity to macroalgal beds, combined with less constrained movement, might allow wrasses to access supplementary dietary resources within adjacent macroalgal beds (Dunning et al. 1992). Alternatively, given the similar structural properties of seagrass and macroalgae (Gratwicke & Speight 2005), wrasses may not differentiate between these two habitat types and simply consider adjacent macroalgal patches to be an extension of the seagrass habitat. If this were true, I would have expected to find that area of seagrass also had a positive effect on wrasse abundance, yet the opposite was found. This could be because trophic spatial subsidies (e.g., transfer of invertebrate prey, Polis et al. 1997; Fagan et al. 1999) from macroalgal beds

contribute towards wrasse abundance in adjacent seagrass patches. Density-dependent processes (e.g., inter- or intra-specific competition) operating in macroalgal beds could also contribute to the observed patterns. Macroalgae beds have recently gained recognition as important juvenile habitat for many reef fishes (Chaves et al. 2013; Eggertsen et al. 2017; Tano et al. 2017; Fulton et al. 2020; Tang et al. 2020), including wrasses (Evans et al. 2014; van Lier et al. 2018). As a consequence, high densities could lead to emigration of individuals (Grüss et al. 2011) from macroalgal beds into adjoining seagrass patches.

While wrasses were collectively most influenced by the proximity to and area of macroalgal beds, differential responses to seascape and within-habitat variables became apparent when distribution patterns for the most abundant species were examined. Here *H. scapularis* and *H. trimaculatus* were more influenced by within-habitat variables (e.g., canopy height and seagrass cover), ‘other’ *Halichoeres* was more influenced by seascape variables, and *Stethojulis strigiventer* was almost equally influenced by both. As one of the most diverse fish families, wrasses display a wide range of swimming, feeding and behavioural strategies (Fulton & Bellwood 2002; Wainwright et al. 2002; Bellwood et al. 2006a) which can include ontogenetic habitat (Green 1996) and diet (Nakamura et al. 2003) shifts. Such differential responses of wrasses to seascape and within-habitat variables in the present study may therefore be a reflection of the ecological versatility of particular taxa (Bellwood et al. 2006a).

Limited research has been conducted on the life history and ecology of threadfin breams (Nemipteridae) making it challenging to predict their response to seascape or within-habitat variables. One of the few studies to examine this taxon identified diel differences in foraging patterns between reef-associated adult and juvenile *Scolopsis bilineata* on Australia’s Great Barrier Reef (Boaden & Kingsford 2012). Adults used daytime shelter sites on coral reefs and moved to nearby sand patches at night to feed on benthic invertebrates, whereas juveniles were associated with daytime feeding within sandy/rubble habitats. In the present study, threadfin breams (i.e., *Pentapodus trivittatus*, *Scolopsis lineatus*) were more likely to be present in seagrass beds close to mangroves, far from hardground and with low density seagrass. This information, combined with a previous study at this location that found threadfin breams were frequently observed in both seagrass and mangrove habitats during daylight hours (Sambrook et al. 2020, **Chapter 3**), suggests that these taxa may use seagrass or mangroves as shelter sites, with reef habitat being less important. In addition, by selecting for low density seagrass patches,

there is also potential for daytime feeding in sandy interstices between seagrass shoots which could be important for juveniles.

I also found support for the hypothesis that taxa with larger home ranges are more influenced by seascape than within-habitat variables. Emperors, rabbitfishes, and snappers all responded strongly to seascape variables. Home ranges for some species from these families can exceed 1 km diameter (e.g., *Lethrinus harak*, Léopold et al. 2017), which could encompass up to seven different habitat types in my study system. In addition, individuals from these families have been previously recorded from multiple habitats at this study location (Sambrook et al. 2020, **Chapter 3**). Although many of these taxa could theoretically move between habitats in this study system given the distances involved, this still needs to be formally tested and would be a valuable area of future research. Foraging behaviour (e.g., diet, timing, shelter) may partially contribute towards explaining the strong responses by some of these taxa to seascape structure. Foraging behaviour can be driven by a range of factors including access to spatially separated resources (Dunning et al. 1992; Ries et al. 2004), resource partitioning (Berkström et al. 2013b), tracking of prey movements (e.g., during flood tides; Davis et al. 2017), predator avoidance (Rooker et al. 2018) or landscape structure (Redhead et al. 2016). With the exception of a few well-studied taxa (e.g., grunts, Haemulidae in the Caribbean, Beets et al. 2003; Appeldoorn et al. 2009), we know very little about cross-habitat foraging patterns, particularly in the Indo-Pacific (Sambrook et al. 2019, **Chapter 2**). Irrespective of the mechanisms involved, the strong responses to a wide range of seascape attributes indicate that these taxa could play an important role as mobile links (Lundberg & Moberg 2003) among habitats.

Given that some taxa move from juvenile to adult habitats, sometimes over considerable distances (e.g., >30 km, *Lutjanus ehrenbergii*, McMahon et al. 2012), I predicted that taxa known to undertake ontogenetic habitat shifts (e.g., emperors, Nakamura & Tsuchiya 2008; parrotfishes, Sievers et al. 2020b; rabbitfishes, Henderson et al. 2017; snappers, Nakamura et al. 2008; and wrasses, van Lier et al. 2018) would be more influenced by seascape characteristics than within-habitat variables. However, the findings suggest that the strength of these responses might differ depending on life stage. Within-habitat variables could be more important for juveniles as many are relatively site-attached and hence reliant on the quality of resources within a patch (e.g., food, refuge), whereas the presence of adults may be a function of both adult and juvenile habitats. I found contrasting results for the two taxa for which I was able to examine adult and juvenile distribution patterns (i.e., emperors and parrotfishes).

As expected, juvenile parrotfishes were more influenced by within-habitat rather than seascape variables, and contrary to expectations, adult parrotfishes responded similarly. However, juvenile parrotfishes were more influenced by small-scale seagrass variables (e.g., shoot density, percent cover epiphytes and canopy height), whereas adult parrotfishes were more likely to be present in seagrass beds that contained some hard coral cover and were close to hardground. These responses appear to reflect the feeding ecology of these distinct life stages in parrotfishes. For instance, newly settled juvenile parrotfishes from the *Scarus* genus are initially carnivorous feeding primarily on harpacticoid copepods (Bellwood 1988) which can be abundant in seagrass beds (Nakamura & Sano 2005) particularly those with high seagrass biomass (Vonk et al. 2010). Older juveniles then transition to an herbivorous feeding strategy where they target protein-rich epiphytes (Nakamura et al. 2003; Clements et al. 2017) which also supports my findings. In contrast, the presence of hard coral or hardground would be expected to be more important for adult parrotfish as most feed from carbonate surfaces (Bonaldo et al. 2014).

Adult and juvenile emperors were predominantly influenced by the surrounding seascape, but responded to different seascape metrics. Juvenile emperors were associated with deeper seagrass beds surrounded by limited coral reef habitats, whereas adults responded to a different suite of seascape metrics (e.g., distance to coral reef, hardground, mangrove) and in an opposite direction to juveniles (e.g., positive response to increased area of coral reef habitat). Emperors are known to use seagrass beds for feeding (Espadero et al. 2020), but can occur in a range of habitat types as adults. For juveniles, their presence in seagrass patches located away from coral reef habitats may reflect the higher predation risk associated with reefs compared to seagrass or mangrove habitats (e.g., Shulman 1985; Lugendo et al. 2006; Grol et al. 2014). In well-connected seascapes such as the Tigak Islands, the presence of seagrass beds away from coral reef habitat may therefore be important for juvenile survivorship and contribute to the replenishment of adults on reefs.

In addition to emperors and parrotfishes, juvenile rabbitfishes, snappers and sweetlips (Haemulidae) were also recorded in seagrass beds on several occasions (pers obs). All these taxa are important for local subsistence fisheries in Papua New Guinea (Papua New Guinea National Fisheries Authority 2007), as well as in other Indo-Pacific regions (Nordlund et al. 2018; Brodie et al. 2020). Fishing effort can be higher in seagrass compared to coral reefs, particularly during monsoon season where proximity to shore makes seagrass patches easier to

access (de la Torre-Castro et al. 2014). As has occurred in Tanzania, this presents a sustainability risk whereby juvenile fish are extracted from seagrass beds and adults are taken from reefs (de la Torre-Castro et al. 2014). This process can erode fish populations and hence needs careful management. Improving our ability to predict which seagrass patches function as nurseries will lead to more evidence-based, spatially-refined management strategies (e.g., seasonal closures, restricted fishing gear) in the future.

Multi-scale studies are still relatively rare in aquatic environments, particularly in tropical seagrass ecosystems. However, as I demonstrate in this study, adopting a multi-scale approach to examine species-habitat relationships can provide a more nuanced perspective on factors that influence species distribution patterns. Given that two-thirds of the fish groups examined were influenced more by seascape than within-habitat variables, these findings highlight that seagrass patches are not functionally equivalent, can support quite distinct communities, and that the broader seascape is important in shaping tropical fish assemblages. Moreover, the taxa influenced by seascape variables included several ecologically and economically important taxa (e.g., emperors, parrotfishes and, rabbitfishes), which could have implications for ecological processes, both in seagrass patches and potentially the wider seascape, as well as for fisheries and spatial management strategies.

## Chapter 5

# Influence of seascape and within-reef variables on a key ecosystem process, macroalgal browsing, on coral reefs in Papua New Guinea

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### 5.1 Introduction

Tropical coral reefs are facing substantial changes to their structure and composition driven by climate change (e.g., thermal stress, ocean acidification) and other anthropogenic stressors (e.g., pollution, invasive species, overfishing, Aronson & Precht 2001; Hughes et al. 2003; Bellwood et al. 2004; Gardner et al. 2005; Hughes et al. 2018). Many of these stressors are predicted to increase in both frequency and intensity in the future (Hoegh-Guldberg et al. 2007; Frieler et al. 2013; Hoegh-Guldberg et al. 2017), placing growing pressure on an already vulnerable ecosystem. One of the major concerns is that collectively these pressures will markedly decrease the cover of live corals on reefs, disrupting the delicate balance between corals and macroalgae, and pushing reefs away from coral-dominated conditions towards alternative states, such as macroalgae-dominated reefs (Hughes et al. 2017b). As a result, taxa capable of removing macroalgae are considered a critical component of reef resilience (Hoey & Bellwood 2009; Graham et al. 2013a).

Herbivory has long been recognised as an important process on coral reefs, however the majority of herbivorous reef fish have limited capacity to remove tall canopy-forming macroalgae (e.g., *Sargassum*) that often dominates on degraded reefs. Instead it appears only a few species (i.e., macroalgal browsers) are capable of removing mature stands of macroalgae (e.g., Bellwood et al. 2006b; Hoey & Bellwood 2009). Moreover, the biomass of macroalgae removed by browsers (i.e., removal rates) and the composition of the browsing community appears to be context and site dependent. Differences in both the rates and agents (i.e., the species responsible for the consumption of macroalgal biomass) of browsing on reefs have been reported across a range of spatial scales including biogeographic region (Tebbett et al. 2020), latitude (Bennett & Bellwood 2011), inshore to offshore gradients (Hoey & Bellwood 2010a; Plass-Johnson et al. 2015; Bauman et al. 2017), among reef zones (e.g., Fox & Bellwood 2008; Hoey & Bellwood 2010b; Vergés et al. 2011; Loffler et al. 2015), and among sites (Cvitanovic & Bellwood 2009; Cvitanovic & Hoey 2010). Although these spatial differences in browsing activity have been described extensively, far less is known about what factors

underpin these differences (Vergés et al. 2011; Nash et al. 2016; Roff et al. 2019). Identifying the factors that contribute towards this variability is a fundamental requirement for effective management of the browsing function on coral reefs.

The majority of studies that have investigated the potential drivers of variation in browsing fish assemblages, and/or their functional impact, have focused on within-reef variables. For instance, the composition of the coral community (Cvitanovic & Hoey 2010; Richardson et al. 2020), the cover of live hard coral or macroalgae (Cvitanovic & Hoey 2010; Chong-Seng et al. 2014; Bauman et al. 2017), and the structural complexity of reef habitats (Vergés et al. 2011) have all been linked to differences in browsing rates. Species-specific responses to such variables may be caused by preferences for particular types of reef habitat and/or the associated environmental conditions through their influence on resource availability (Nash et al. 2016), predation risk (Rizzari et al. 2014; Gil et al. 2017; Rasher et al. 2017), refuge potential (Bauman et al. 2019), as well as the provision of suitable recruitment habitat (Hoey et al. 2013).

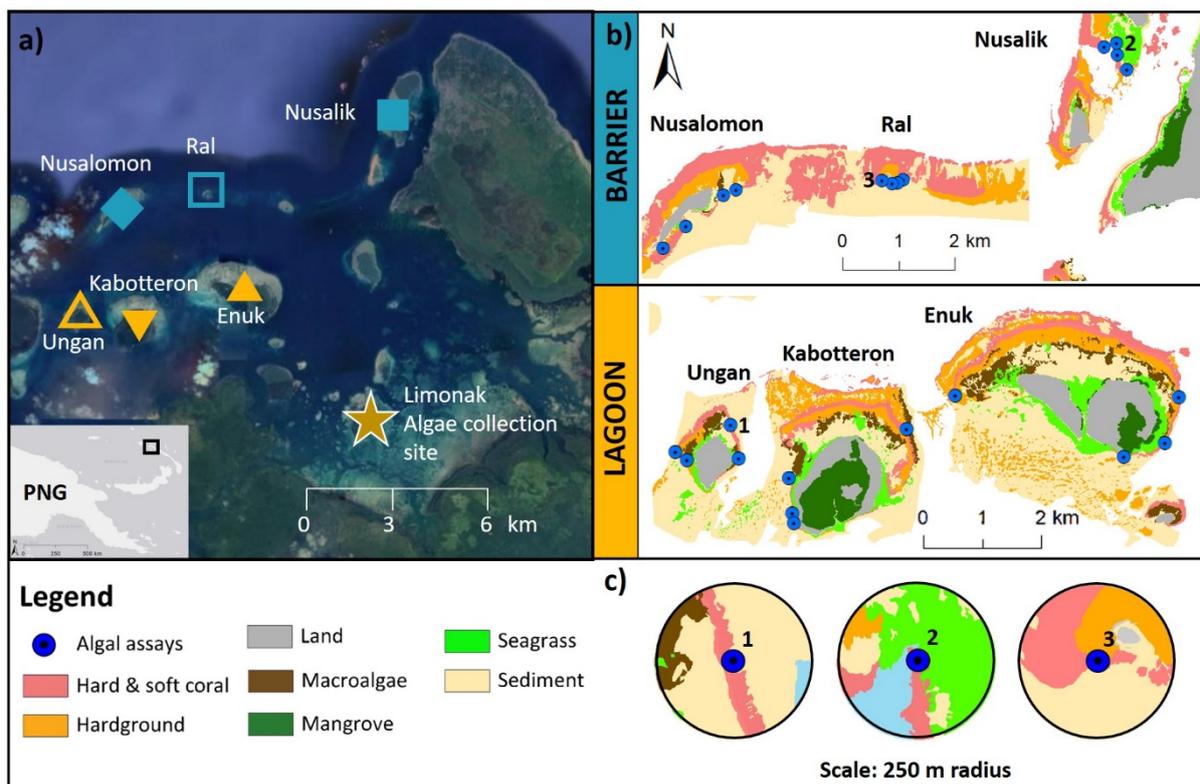
Potential drivers of variation in browsing activity beyond within-reef variables have received limited attention. There is, however, a growing body of research showing that the seascape surrounding coral reefs (e.g., the presence of seagrass beds, mangroves) can exert a considerable influence on the composition and abundance of reef fish communities (e.g., Grober-Dunsmore et al. 2007; Olds et al. 2012a; Pittman & Olds 2015). More recently, research has shown that the seascape composition and configuration can also influence the delivery of certain ecological functions by fishes (e.g., grazing and bioeroding by parrotfishes, Eggertsen et al. 2019; 2020a; 2020b), and a similar response could occur in browsing fishes. For instance, the availability of alternative food sources for browsing fishes in close proximity to reefs (e.g., neighbouring macroalgal and seagrass beds) may affect the foraging behaviour and hence the realised rates and agents of browsing on reefs. Indeed, differences in browsing activity have been linked to seascape structure on subtropical reefs off Australia's east coast, with browsing rates higher on isolated reefs compared to reefs in close proximity to each other (Martin et al. 2018), and further away from mangrove habitat (Yabsley et al. 2016). Given that many browsing species in tropical coral reef ecosystems are highly mobile (Meyer & Holland 2005; Welsh & Bellwood 2014), and are often observed in non-reef habitats as adults (e.g., siganids, Lugendo et al. 2005; Evans et al. 2014; **Chapters 3 & 4**), or juveniles (e.g., siganids, Hoey et al. 2013; Tang et al. 2020), it is reasonable to expect that the composition and spatial configuration of the seascape may directly influence browsing activity on coral reefs.

Here I investigate whether the seascape structure and within-reef benthic variables affect browsing activity on tropical coral reefs in the Tigak Islands, Papua New Guinea. Specifically, I examine which seascape and within-reef variables influence: 1) macroalgal removal rates, and 2) the agents of macroalgae removal among reef sites.

## 5.2 Methods

### 5.2.1 Study area

The study was conducted on reefs surrounding six islands in the microtidal location of the Tigak Islands, Kavieng, New Ireland Province, Papua New Guinea ( $2^{\circ} 38'S$ ,  $150^{\circ} 43'E$ ; Figure 5.1) during August 2019. At each island, four reef crest sites (1-3 m depth) were haphazardly selected from sheltered positions (i.e., away from the wave and wind exposed crest) (Figure 5.1b).



**Figure 5.1.** a) Map of the Tigak Islands, Kavieng, New Ireland Province with inset showing location in Papua New Guinea (PNG). The six study islands are shown with blue symbols for barrier reef islands and yellow symbols for lagoon islands (see results for distinction between island types). The star shows the macroalgae collection site, b) shallow water habitat types and macroalgal assay sites (blue circles) around each island and, c) examples of different seascapes (250 m radius) surrounding assay sites.

### 5.2.2 Macroalgal assays

Assays of two locally abundant canopy-forming brown macroalgae (*Sargassum* sp. and *Cystoseira trinodis*) were used to quantify rates of macroalgae removal at each of the 24 reef sites. These taxa were selected because canopy-forming brown algae are often implicated in phase shifts from coral- to macroalgal-dominated reefs (Hughes 1994; Bellwood et al. 2006b; Hughes et al. 2007a; Hoey & Bellwood 2009). Algae were collected from the reef flat of an inner lagoon island, Limonak (Figure 5.1a), returned to the Nago Island Mariculture and Research Facility within 1 hour of collection and maintained in large (~1000L) circular aquaria with flow-through seawater and supplemental aeration for up to 3 days. Prior to deployment, thalli were spun for 30 seconds in a salad spinner to remove excess water, and the wet weight (g) and maximum height (cm) of each thallus was recorded. The mean initial weight and height were  $57.2 \text{ g} \pm 1.7 \text{ SE}$  and  $50.2 \text{ cm} \pm 10.1 \text{ SE}$  respectively for *Cystoseira* and  $75.9 \text{ g} \pm 2.7 \text{ SE}$  and  $57.6 \text{ cm} \pm 13.4 \text{ SE}$  for *Sargassum*.

Four randomly selected thalli of each macroalga (Chong-Seng et al. 2014) were deployed in pairs (i.e., one thallus of each macroalga) between 09:00 and 10:00 at each of the 24 sites. Three assay pairs were exposed to local herbivore communities, and one assay pair was placed inside an exclusion cage (~55 x 30 x 20 cm, 1 cm square mesh) to quantify any losses due to handling, translocation and hydrodynamic conditions at the site (following Hoey & Bellwood 2010b). Each thallus was attached to the substratum (i.e., dead coral or rock) using galvanised wire and an elastic band, with a small numbered plastic tag used to identify individual thalli (following Hoey & Bellwood 2010b). After four hours (i.e., 13:00-14:00) assays were collected, spun in a salad spinner for 30 seconds, and thallus weight and height recorded. To account for reductions in algal weight due to handling rather than herbivory, the weight of macroalgae consumed from the exposed assays was standardised using Equation 1:

$$[T_o \times (C_f/C_o) - T_f] \qquad \text{Equation 1.}$$

where  $T_o$  and  $T_f$  are the initial and final weights for the exposed thalli, and  $C_o$  and  $C_f$  are the initial and final weights for the cage control (Cronin & Hay 1996).

### 5.2.3 Agents of macroalgal removal

To identify the agents of macroalgal consumption and quantify their contribution to macroalgal removal, a small video camera (GoPro 4) mounted on a dive weight was positioned approximately 1 m from one of the assay pairs at each of the 24 sites. At the start of each

recording, a small scale bar was held adjacent to each assay and used to calibrate the size of fishes in the video footage. For each video, the first two hours of video footage were viewed, and the identity, total length (to nearest cm) and number of bites on each macroalga by individual fish were recorded. Given that the bite impact of an individual is influenced by body size, bites were subsequently converted to mass standardised bites (total number of bites per species x biomass) hr<sup>-1</sup> (following Bellwood et al. 2006b) using published length-weight relationships (Kulbicki et al. 1993; Kulbicki et al. 2005; Froese & Pauly 2021).

To quantify the herbivorous fish assemblages at each site the first two hours of video footage from each assay was viewed and the presence and maximum number of individuals observed on a single frame (MaxN) for species belonging to the families Acanthuridae, Kyphosidae, Siganidae and subfamily Scarinae (family Labridae) were recorded. MaxN was used as a conservative measure of relative abundance of species observed during the video recording period (Cappo et al. 2004).

#### ***5.2.4 Characterising the within-reef benthic community composition and seascape structure***

To examine the influence of within-reef benthic variables on rates and agents of macroalgal removal, the benthic community was surveyed along four 25-m point-intercept transects at each site (depth 1-3 m). Each transect was placed along the reef crest separated by a minimum of 5 m between adjacent transects. The substratum directly beneath the transects at 25 random points was recorded. The substratum was recorded as hard (scleractinian) coral, macroalgae, soft coral, epilithic algal matrix (EAM), crustose coralline algae (CCA), sand, rubble, and 'other' which included bare rock, recently dead coral, seagrass and other sessile invertebrates. Hard corals were identified to genus and morphotype where possible, and macroalgae were identified to genus.

Features of the seascape surrounding each of the 24 sites were quantified from a benthic habitat map of each island (see **Chapter 4** for detail of map production). The proportion of five broad habitat types identified as influencing different aspects of the reef fish community in **Chapter 4**, specifically hard and soft coral reef structure, hardground (e.g., pavement), macroalgae beds, sand, and seagrass beds in the surrounding seascape was quantified. Mangroves were excluded from analysis due to their limited distribution among study sites (i.e., present around only six sites compared to a minimum of 13 sites for other habitat types), and because browsing species were not recorded from mangrove habitat in this study region (**Chapter 3**), and are rarely

observed in mangrove habitats elsewhere (**Chapter 2**). ArcGIS (v10.6.1) was used to quantify the proportion by area of each habitat type within a 250 m radius of each site. The 250 m radius size was chosen based on available home range data for browsing species (e.g., browsing siganids, Ebrahim et al. 2020; *Naso unicornis*, Meyer & Holland 2005), and given the close proximity of some sites around islands that would have resulted in spatial overlap of seascape characteristics leading to high spatial autocorrelation. In addition, distance from each site to open ocean was measured using the Near function in ArcGIS. If the route crossed land, the shortest distance was measured manually avoiding land.

### 5.2.5 Data analysis

Distance-based linear models (DISTLM, McArdle & Anderson 2001; Anderson et al. 2008) and distance-based redundancy analyses (dbRDA) were used to identify and visualise which of the 16 benthic and 6 seascape variables (Table A5.1) best predicted spatial patterns of macroalgal removal and browsing activity. No predictor variables were identified as collinear (Pearson's  $r > 0.7$ ) based on pairwise Pearson's correlations obtained from draftsman plots. Predictor variables were  $\log(x+1)$  transformed and normalised to account for different scales of measurement. All possible combinations of predictor variables were fitted using the *Best* procedure (Anderson et al. 2008). Models were selected using the Akaike's Information Criterion adjusted for small sample sizes (AICc). In addition, given the potential risk of overfitting associated with AICc values, model outcomes were checked for consensus against the more conservative Bayesian Information Criterion (BIC) (Anderson et al. 2008).

Following analysis of model outputs, the six islands were subsequently separated into two groups, based on their position and proximity to open ocean. Three islands, hereafter termed "barrier reef islands" were directly adjacent to and near to open ocean (<2.2 km) and three islands, termed "lagoon islands", were further from open ocean (>5 km) and situated behind the barrier reef islands (Figure 5.1b).

The amount of macroalgae removed from assays was compared using a univariate three-factor nested permutational analysis of variance (PERMANOVA) with reef position (i.e., barrier or lagoon) as a fixed factor, island as a nested random factor within reef position, and site as a nested random factor within island. Analyses were based on zero-adjusted Bray-Curtis similarity resemblance matrices that were constructed from untransformed data. The PERMANOVA (Type III sums of squares) analysis was performed using 9999 permutations of residuals under a reduced model (Anderson et al. 2008). The PERMDISP routine was used

to verify that any differences were primarily due to location rather than dispersion effects. Similar patterns of removal were observed for both *Cystoseira* and *Sargassum* (Figure A5.1) and data were subsequently pooled for analysis and visualisation.

Species that accounted for >1% of mass standardised bites on assays were identified, and their macroalgal consumption patterns (i.e., mass standardised bites) were subsequently analysed to explore the influence of within-reef and seascape variables on browsing community activity. Differences in the browsing community (MaxN) and function (i.e., mass standardised bites) were compared using a two-factor nested permutational analysis of variance (PERMANOVA) with reef position as a fixed factor and island as a nested random factor within reef position, following the same procedure described above. Analyses were based on zero-adjusted Bray-Curtis similarity resemblance matrices that were constructed using square-root transformed data. Where the number of possible permutations was <100, results were interpreted cautiously using the more conservative Monte Carlo p-value. Results were visualised using principal coordinates analysis (PCO) plots. A SIMPER (similarity percentage) analysis was used to identify the browsing taxa driving any differences in the browsing community between lagoon and barrier reef islands. The relationship between the relative abundance of each browsing species and mass standardised bites was examined using Pearson's correlation coefficient ( $r$ ).

Univariate analyses were conducted in R v4.0.0 (R Core Team 2020) and all multivariate data analyses were performed using the software PRIMER v7 with PERMANOVA+ (Anderson et al. 2008).

### **5.2.6 Spatial autocorrelation**

Each response and seascape predictor variable was checked for spatial autocorrelation using a standardised inverse distance weighted Global Moran's I conceptualisation of spatial relationships based on the average distance to each site's four nearest neighbours (average distance = 3,459 m; Table A5.2). Where spatial autocorrelation values were significant (pseudo-significance  $P \leq 0.01$  calculated from 999 permutations) and Moran's I values >0.4 (Hamylton & Barnes 2018), relationships were visually examined using cluster and outlier analysis (Anselin Local Moran's I) in ArcGIS. Only one seascape predictor variable was significantly spatially autocorrelated suggesting that our sampling sites were far enough apart to each represent independent samples. The area of sand within 250 m radius was significantly, but not strongly, spatial autocorrelated (Moran's I = 0.48,  $P = 0.001$ ). Cluster and outlier analysis showed one high-low outlier at Eruk (lagoon) indicating the seascape surrounding one

site had a high proportion of sand within 250 m compared to other neighbouring sites, one low-high outlier (i.e., low value for one site compared to neighbouring sites) and two high-high clusters at Nusalomon (barrier) and two low-low clusters at Nusalik (barrier). Three response variables exhibited positive spatial autocorrelation, specifically, total mass standardised bites (Moran's I = 0.626, P <0.001), *Naso lituratus* mass standardised bites (Moran's I = 0.598, P <0.001) and mean % macroalgae consumed (Moran's I = 0.686, P <0.001). These were driven by differential consumption rates by species among islands (see Results).

### 5.3 Results

The proportion of macroalgae removed was lower and more variable on lagoon reefs, than on barrier reefs (Figure 5.2). For instance, algal consumption at sites around Eruk ranged from  $28.1\% \pm 4.9$  (mean  $\pm$  SE) to  $70.0\% \pm 5.7$ , and from  $16.4\% \pm 5.7$  to  $66.6\% \pm 9.9$  at Ungan. Macroalgae removal was two-fold higher on barrier ( $71.2\% \pm 2.0$ ) compared to lagoon reefs ( $35.0\% \pm 3.3$ ) (Figure 5.2). However, one barrier reef island, Nusalik, had considerably less macroalgae consumed on average ( $57.1\% \pm 3.9$ ) compared to the other two other barrier reef islands (i.e., Nusalomon  $81.4\% \pm 1.6$ , Ral  $75.8\% \pm 2.1$ ). The mean handling loss was  $12.2\% \pm 1.0$  ( $6.3\text{ g} \pm 0.7$ ) for *Cystoseira* and  $7.1\% \pm 0.8$  ( $4.9\text{ g} \pm 0.7$ ) for *Sargassum*.

The best fit distance-based linear model for macroalgal removal rates included eight predictor variables. Distance to open ocean was the only seascape variable included in the top model, with the remaining seven related to within-reef benthic community composition (Figure 5.3). Together these eight predictor variables explained 90% of the variation in macroalgal removal rates among sites (Table 5.1). Lower macroalgal removal rates were strongly correlated with increasing distance from open ocean and this variable alone explained 51% of the variance. The cover of rubble, sand and massive *Porites* at each site was negatively related to removal rates, while the cover of crustose coralline algae (CCA) and *Acropora* (digitate, staghorn, and tabulate growth forms) was positively related.

From the video footage, 23 fish species were observed taking a total of 61,014 bites from the macroalgal assays (4,086.25 total mass standardised bites). Six species were responsible for >99% of the mass standardised (ms) bites: the orange-spine unicornfish *Naso lituratus* (59.2%), the rabbitfishes *Siganus canaliculatus* (13.5%) and *Siganus doliatus* (10.3%), the blue-spine unicornfish *Naso unicornis* (7.0%), and two species of chub, *Kyphosus cinerascens* (5.3%) and *Kyphosus vaigiensis* (3.9%). A total of 61.6% of the variance was explained by the best fit

DISTLM for mass standardised bites of the browsing community (Table 5.1). The best fit model included five predictor variables: distance from open ocean, area (%) of seagrass within a 250 m radius, and the cover of encrusting corals, sand, and other *Acropora* at each site (Figure 5.4). Distance to open ocean explained the majority (25.9%) of the variance. Lower browsing activity was typically recorded further away from open ocean and at sites with higher cover of sand, with *S. canaliculatus*, *K. cinerascens* and *K. vaigiensis* absent from sites with this combination of attributes (Figure 5.4). Area of seagrass within 250 m of each site also influenced the browsing community bite rates. At sites with higher amounts of seagrass within 250 m, *S. canaliculatus* and the two kyphosid species were responsible for more bites than *N. lituratus* or *N. unicornis* which were either absent or took few bites on assays (Figure 5.4).

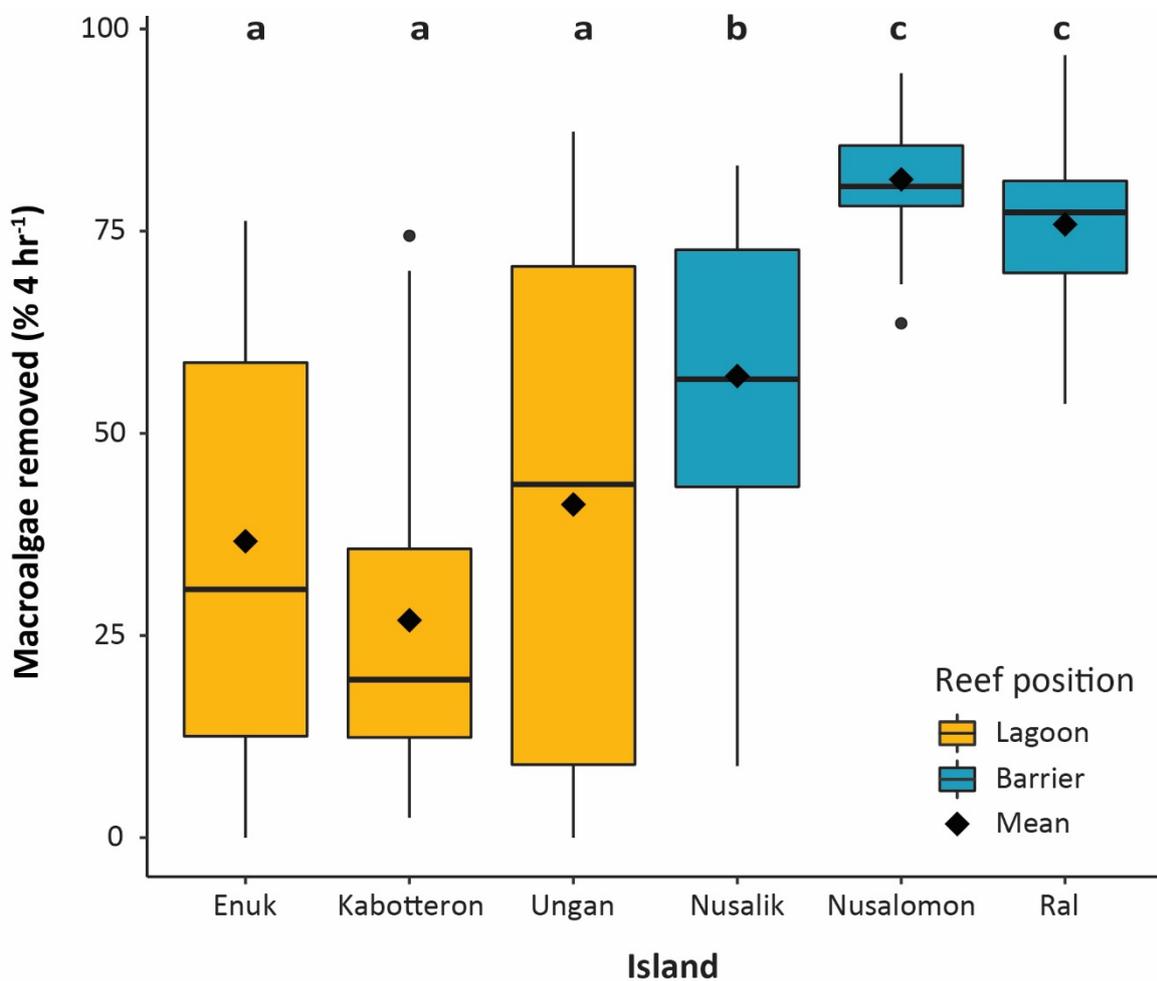
The browsing community structure and feeding of browsing fish assemblages differed between lagoon and barrier reef islands (Figure 5.5, Table 5.2, Figure A5.2). The structure of the browsing communities around lagoon islands was variable with low similarity among sites (SIMPER average similarity = 26.9), compared to barrier reef sites (average similarity = 54.9). Only a single species contributed towards the similarity between lagoon islands, the rabbitfish *S. doliatus*, whereas browsing fish communities around barrier reef islands were characterised by three species, primarily *N. lituratus*, followed by *S. doliatus* and *S. canaliculatus*.

The browsing species most frequently recorded from video footage were: *S. doliatus* (83% of videos), *N. lituratus* (67%), and *S. canaliculatus* (58%). Five of the six key browsing species were more frequently observed at barrier compared to lagoonal reef sites. The dominant browsing species, *N. lituratus*, was six times more abundant (MaxN = 12.3 individuals  $\pm$  3.0 SE) and took 73 times more ms bites (198.8 ms bites  $\pm$  60.3 SE) at barrier reef sites compared to lagoon sites (MaxN = 2.0  $\pm$  1.1 SE, ms bites = 2.6  $\pm$  1.2 SE). Similarly, the relative abundance of the rabbitfish *Siganus canaliculatus* was four-fold higher, and the bite rate was 30 times more, on macroalgal assays at barrier reef sites compared to lagoon islands. *K. vaigiensis* was the only species that had similar abundances and browsing rates among barrier reef and lagoon islands.

Importantly, the presence of a browsing species at a site did not always equate to browsing activity (Figure 5.6). On several occasions, individuals or groups of browsers were observed on the video footage but not recorded feeding on assays, the only exception being *N. unicornis* which was always recorded browsing on assays whenever it was observed (Figure 5.6). While browsing activity was related to relative abundance for three of the six browsing species (*K.*

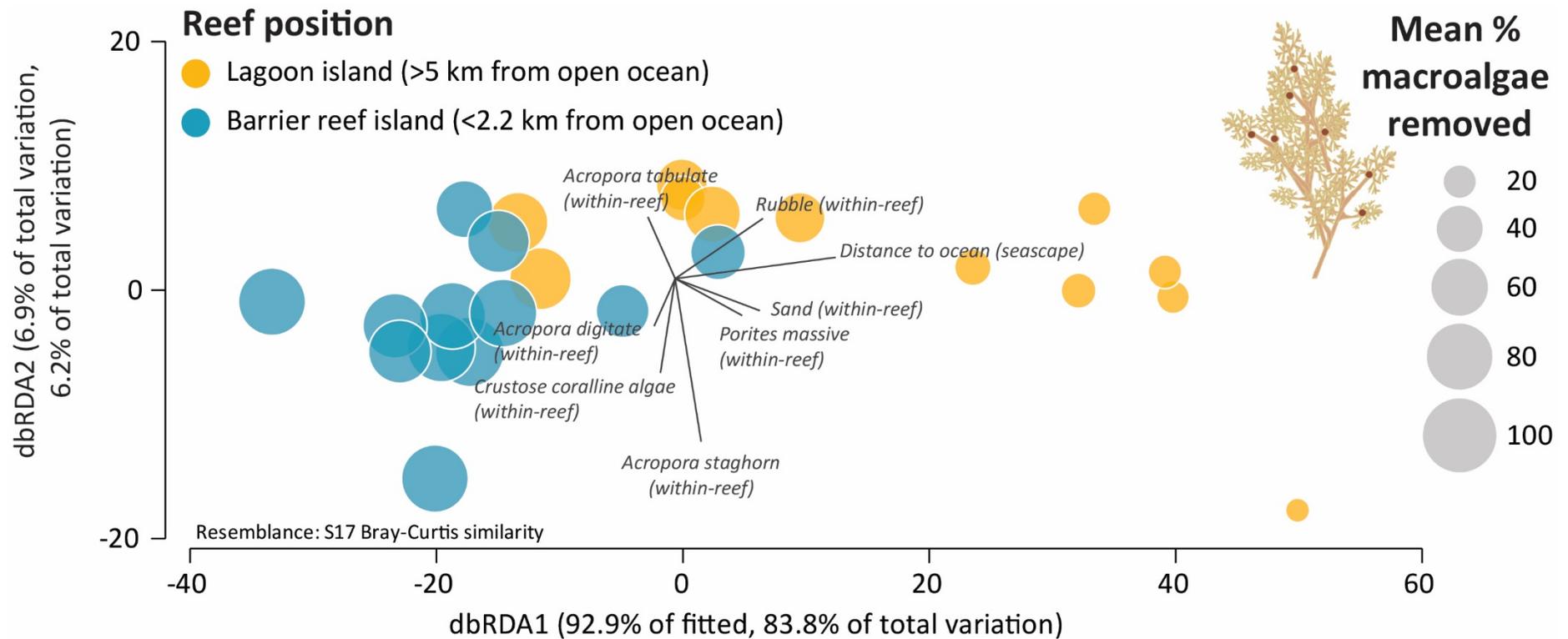
*cinerascens*, *N. unicornis* and *S. doliatus*), the remaining three browsing species (*K. vaigiensis*, *N. lituratus* and *S. canaliculatus*) showed no significant relationship between relative abundance and browsing activity (Figure 5.6) indicating that presence was not necessarily a strong predictor of function for all browsing taxa. In addition, the positive correlation between relative abundance and bites for *K. cinerascens* was driven by a single outlier.

Despite the overall dominance of *N. lituratus* in terms of ms bites, its dominance was largely attributable to extensive feeding at two barrier reef islands, Nusalomon and Ral. Elsewhere, *K. cinerascens* took, on average, the highest number of ms bites at Nusalik (barrier reef), *N. unicornis* was responsible for the most ms bites at Ungan (lagoon reef), and *S. doliatus* took the most bites at EnuK and Kabotteron (lagoon reefs).

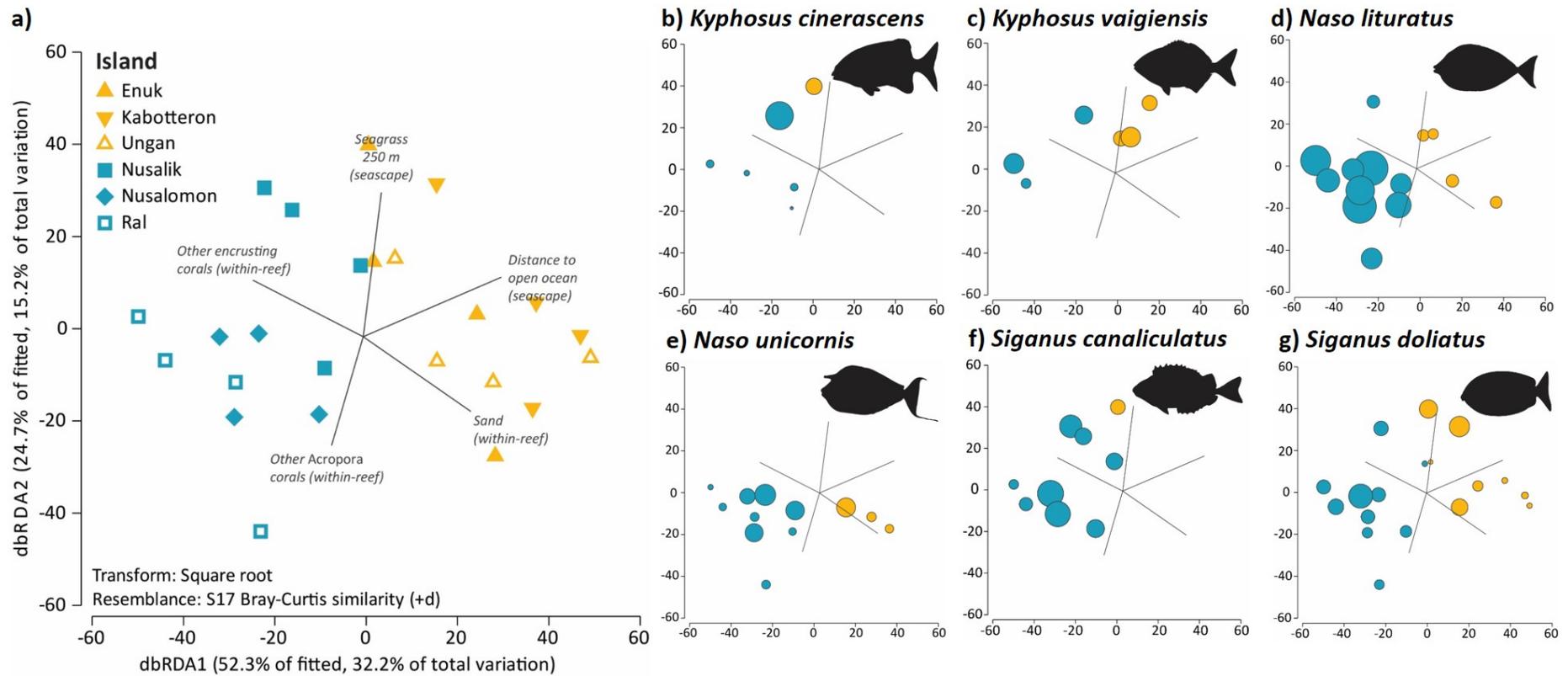


**Figure 5.2.** Differences in macroalgal removal (%) between lagoon and barrier reefs. Median and 25% quantiles are shown on the boxplot with black dots as outliers. Diamond symbol represents the mean macroalgal removed (%) from assays for each island. Letters above

boxplots denote similar groups in terms of mean macroalgal removal identified by PERMANOVA pairwise tests ( $p < 0.01$ , Table A5.3).



**Figure 5.3.** Distance-based redundancy analysis (dbRDA) ordination showing the predictor variables identified in the top model (DISTLM) as influencing macroalgal removal rates. Yellow = lagoon sites, blue = barrier reef sites. Image credit: Tracey Saxby, Integration and Application Network.



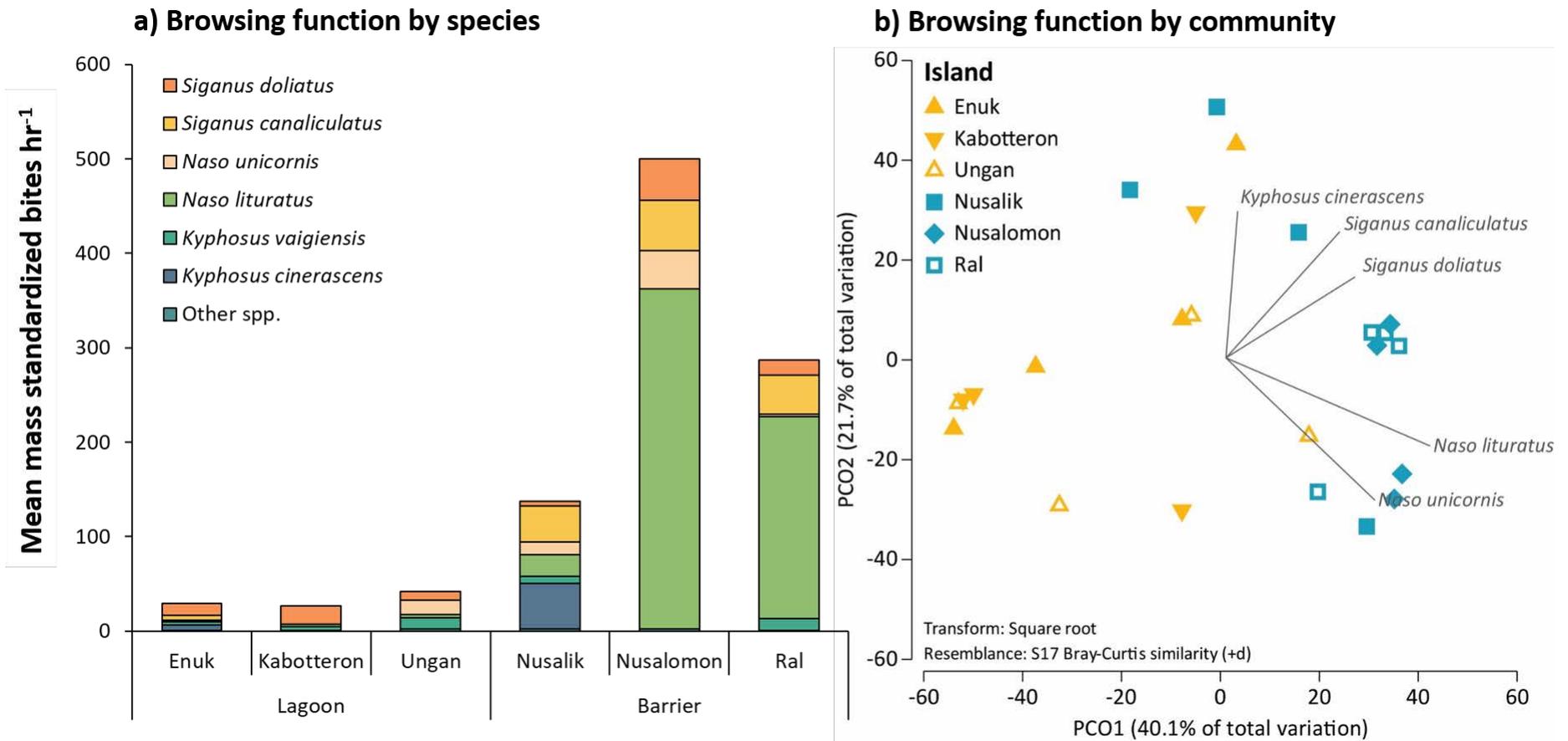
**Figure 5.4.** Distance-based redundancy analysis (dbRDA) ordination showing the predictor variables identified in the best fit model (DISTLM) as influencing browsing community mass standardised bites at each site. Smaller panels (b-g) show presence of six main browsing taxa in relation to vectors described in panel (a). Bubble size represents the square-root transformed ms bites for each individual taxa at sites where they were observed feeding. Yellow = lagoon sites, blue = barrier reef sites.

**Table 5.1.** Top model of benthic and seascape variables influencing macroalgal removal rates and mass standardised bites of browsing community. Results given for AICc and BIC models from distance-based linear modelling (DISTLM).

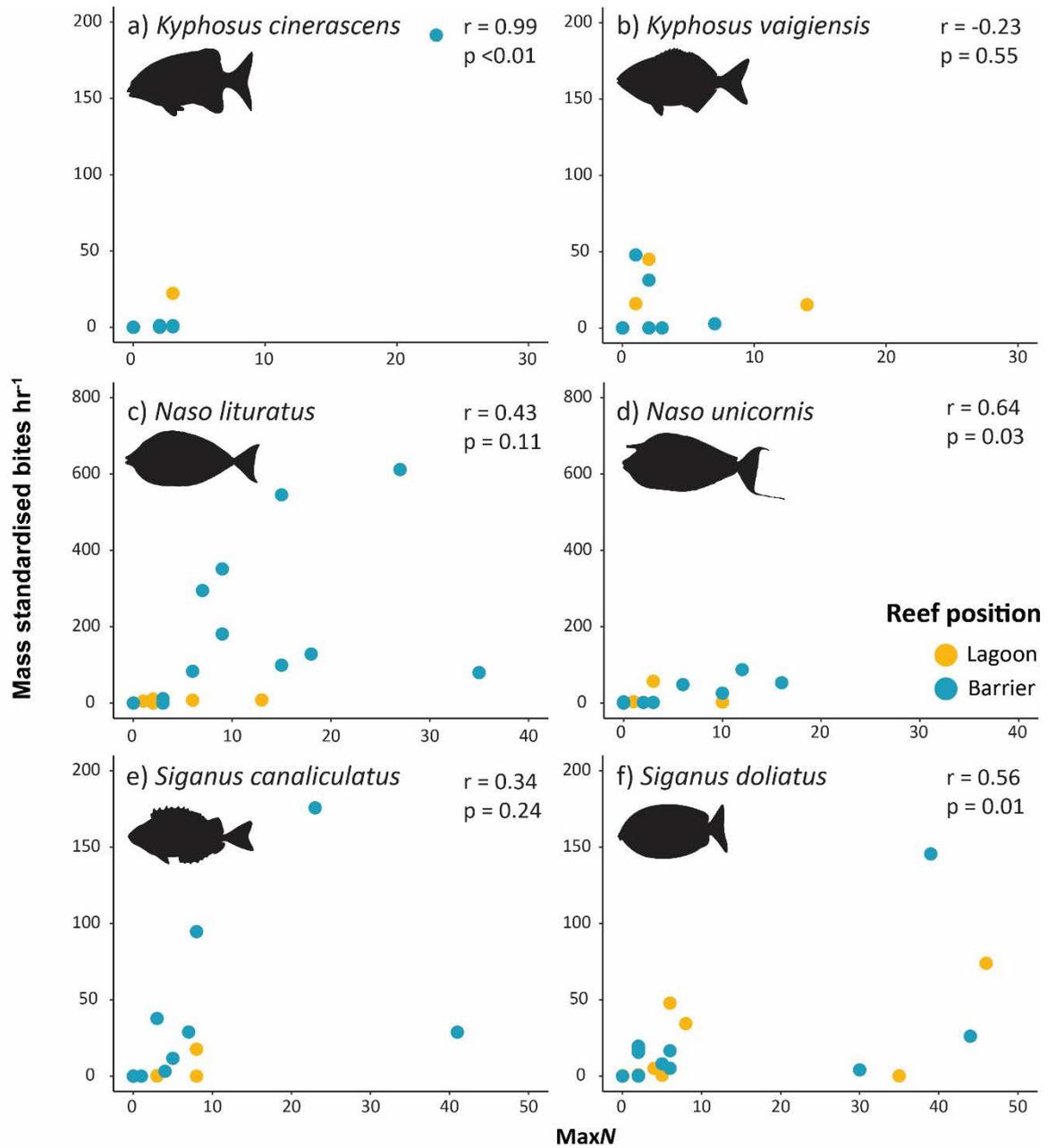
Model	R <sup>2</sup>	AICc	BIC
<b>Macroalgal removal rates</b>			
<i>Seascape variables:</i> Distance to open ocean	0.90	130.7	128.44
<i>Within-reef benthic variables:</i> sand, rubble, crustose coralline algae, <i>Porites</i> massive, <i>Acropora</i> staghorn, <i>Acropora</i> tabulate, <i>Acropora</i> digitate			
<b>Mass standardised bites of browsing community</b>			
<i>Seascape variables:</i> Distance to open ocean, area of seagrass 250 m	0.62	181.52	186.47
<i>Within-reef benthic variables:</i> other encrusting corals, other <i>Acropora</i> , sand			

**Table 5.2.** PERMANOVA results for comparing differences between reef position (fixed), islands (random nested within reef position) and sites (random nested within islands) for macroalgal removal rates, and differences between reef position (fixed) and islands (random nested within reef position) for mass standardised bites of the browsing community. \*ECV denotes percent estimated components of variation.

	df	SS	MS	Pseudo- <i>F</i>	P(perm)	ECV*
<b>Macroalgal removal rates</b>						
Reef position	1	38942	38942	16.179	<b>0.0423</b>	36.5%
Island(Reef position)	4	9636.3	2409.1	1.419	0.2202	2.1%
Site(Island(Reef position))	18	30664	1703.6	2.351	<b>0.0003</b>	11.6%
Residual	115	83338	724.68			
Total	138	163950				
<b>Mass standardised bites of browsing community</b>						
Reef position	1	15376	15376	9.4826	<b>0.0011</b>	36.8%
Island(Reef position)	4	6486.2	1621.5	0.7344	0.7344	3.7%
Residual	18	37599	2088.8			
Total	23	59461				
<b>Relative abundance (MaxN) of browsers</b>						
Reef position	1	5452.9	5452.9	3.7883	<b>0.0336</b>	19.1%
Island(Reef position)	4	5757.7	1439.4	1.0197	0.4531	0.4%
Residual	18	25409	1411.6			
Total	23	36619				



**Figure 5.5.** Browsing rates by: a) the six species each accounting for >1% of mass standardised bites and all other species (n = 17 spp.), and b) PCO plot to visualise the browsing function at the browsing community level. Vectors show species contributing to community patterns. Yellow = lagoon sites, blue = barrier reef sites.



**Figure 5.6.** Relationship between relative abundance (MaxN) of the six most common browsing species and their realised impact (i.e., mass standardised bites hr<sup>-1</sup>). a) *Kyphosus cinerascens*, b) *K. vaigiensis*, c) *Naso lituratus*, d) *N. unicornis*, e) *Siganus canaliculatus*, and f) *S. doliatus*. Yellow = lagoon sites, blue = barrier reef sites. R value = Pearson's correlation coefficient.

## 5.4 Discussion

Both rates and agents of macroalgal removal on shallow water reefs were most strongly influenced by distance to oceanic environment, but exhibited differential responses to other seascape and benthic variables. Macroalgal removal rates were higher on reefs closer to open ocean compared to those further inside the lagoon, and greater at sites containing higher cover of *Acropora* spp. and crustose coralline algae than sites with higher cover of sand, rubble and/or massive *Porites*. Similarly, the agents of macroalgal removal (i.e., fish species recorded feeding on the macroalgal assays) were influenced by distance to open ocean, the area of seagrass within 250 m, and encrusting corals, *Acropora* spp., and sand. Assay consumption on two of the reefs closest to open ocean was dominated by a single browser, the orange-spine unicornfish *Naso lituratus*, however, there was considerable variation in the species responsible for browsing across the study system, particularly around the three lagoon reefs.

### 5.4.1 Environmental drivers of rates and agents of browsing

Distance to open ocean was the most important variable explaining differences in both macroalgal removal rates and browsing activity among sites, with the highest removal rates at reefs proximal to the open ocean. Environmental and physical gradients are frequently evident between inshore and offshore reef systems including changes in pollutant loads, turbidity, sedimentation, wave energy, depth, and productivity (Bellwood & Wainwright 2001; Larcombe et al. 2001; De'ath & Fabricius 2010; Fisher et al. 2015; Bainbridge et al. 2018). These gradients can have concomitant effects on both benthic (Wismer et al. 2009; De'ath & Fabricius 2010; Baum et al. 2015; Draisma et al. 2018) and fish communities (Williams 1982; Williams & Hatcher 1983; Mallela et al. 2007) on coral reefs, and therefore it seems reasonable to expect that these differences would also translate to differential delivery of a key ecological function, such as browsing. Yet few studies have explicitly examined how browsing patterns on reefs can change across an inshore-offshore gradient (for exceptions see Hoey & Bellwood 2010a - Great Barrier Reef (GBR); Plass-Johnson et al. 2015 - Indonesia). Comparing findings among studies does however begin to reveal some consistent patterns across regions. Similar to this study, others have shown that outer reefs appear to support a higher biomass or relative abundance of browsers and browsing communities are often dominated by species from the acanthurid genus *Naso* (Hoey & Bellwood 2010a; Plass-Johnson et al. 2015). In contrast, the browsing community composition on inshore reefs appears to be more variable, as observed in this study and others (e.g., Cvitanovic & Bellwood 2009). Potential mechanisms for these

inshore-offshore differences in browsing fish assemblages are difficult to ascertain, but may be related to differences in dietary resource availability, predation, and the availability of preferred habitats.

Rates of macroalgal removal may be driven by the availability of macroalgal resources on reefs (e.g., Cvitanovic & Hoey 2010; Hoey & Bellwood 2010a; Nash et al. 2016), and/or the availability of alternative resources elsewhere in the seascape (e.g., Hoey & Bellwood 2010a; Bennett & Bellwood 2011; Bellwood et al. 2018). In the present study, there was no relationship between macroalgal removal rates and within-reef macroalgae cover, potentially because macroalgae cover was generally low (<10%) on reefs across all sites. Similarly, no relationship emerged between removal rates and the presence of macroalgal beds in the surrounding (250 m buffer) seascape. This was despite lagoon reefs, where removal rates were consistently lower than barrier reefs, often being situated in the vicinity of macroalgal beds. One explanation for this lack of relationship may be that several, but not all, of the macroalgal beds around the lagoon islands are dominated by green algae (e.g., *Halimeda*, *Caulerpa*) which may be a less attractive or palatable resource than brown macroalgae (e.g., *Sargassum*) for many browsing taxa (e.g., *Kyphosus vaigiensis*, Loffler et al. 2015; *Naso unicornis*, Rasher et al. 2013). Future steps to gain greater insight into the influence of resource availability on browsing patterns should include examining whether different types of macroalgal beds in the surrounding seascape can affect outcomes, and conducting experimental assays across a breadth of vegetated habitats (e.g., seagrass, green and brown macroalgal beds) as well as on reefs.

The rapid rate of assay consumption at the barrier reef sites compared to reefs around the lagoon islands could indicate resource limitation. The two barrier reefs with the highest macroalgal removal rates (i.e., Nusalomon and Ral), had little vegetated habitat, and instead were surrounded by a low complexity matrix dominated by sand and hardground which browsers may be reluctant to move across (Turgeon et al. 2010; Gil et al. 2017) to find alternative resources. As a consequence, browsers located at these two reefs may be less dispersed and have fewer foraging options leading to higher intensity of browsing where macroalgae assays are offered.

While the availability of sufficient resources is essential for survival, foraging decisions also need to be carefully balanced against the risk of predation. Habitat structure quantified at seascape and within-reef scales could influence predation risk by altering predator-prey

dynamics (Catano et al. 2016; Gaynor et al. 2019). How browsers respond to the threat of predation will be contingent on species identity, body size, foraging behaviour and propensity to form groups (Rizzari et al. 2014; Catano et al. 2016). In the current study, browsers, particularly *Naso lituratus* and *N. unicornis*, tended to occur in larger groups around the barrier reefs compared to lagoonal reefs, which is a common behavioural strategy in predator-rich environments (Magurran 1990; Michael et al. 2013; Gil et al. 2017). High predation risk can in turn affect foraging behaviour (Rizzari et al. 2014; Nash et al. 2016) by leading to faster feeding rates (Catano et al. 2016) when suitable resources are located which may be the case around two of the barrier reef islands (i.e., Nusalomon and Ral). However, elevated predation risk can also suppress feeding in less structurally complex habitats by limiting refuge potential (Fox & Bellwood 2007; Hoey & Bellwood 2011; Graham & Nash 2013; Gil et al. 2017), and/or requiring increased vigilance (Brown 1999), and this may be contributing towards the lower browsing activity around the lagoonal reef sites with higher amounts of sand.

#### ***5.4.2 Differential responses of browsing agents across the seascape***

Species-specific habitat preferences may disproportionately influence the maintenance or delivery of browsing on coral reefs (Ruttenberg et al. 2019). Although 23 species were recorded taking bites on macroalgae, only six species accounted for the majority of browsing activity (99% of bites). Importantly, the spatial distribution of those six browsing species differed considerably across the seascape and their differential distribution patterns translated into variation in the structure of the browsing community and browsing rates.

The orange-spine unicornfish *Naso lituratus* was responsible for the highest browsing rates on reefs but its distribution, and therefore realised impact, was largely confined to two reefs that were close to open ocean (i.e., Nusalomon and Ral). Such apparent habitat specificity by *N. lituratus* for outer, as opposed to inshore, reefs has been reported elsewhere (e.g., GBR, Cheal et al. 2012; McClure et al. 2019; Indonesia, Plass-Johnson et al. 2015) although in the present study this habitat preference was evident across a relatively compressed inshore-offshore system (e.g., <10 km between reefs near and far from open ocean as compared to >50 km in the GBR and Indonesia studies). In addition, *N. lituratus* occurred predominantly at sites that supported benthic communities typically found in higher energy reef environments (e.g., dominated by CCA, *Acropora* spp., encrusting corals, Dollar 1982; Fabricius & De'ath 2001; Wismer et al. 2009; Gove et al. 2015). These findings lend strong support for the notion that

*N. lituratus* has a tightly coupled relationship with the physical or environmental properties associated with outer reef habitats (Cheal et al. 2012).

The blue-spined unicornfish *N. unicornis* also followed a similar distribution pattern to *N. lituratus* but was less abundant and took fewer bites. However, these findings were unexpected given that *N. unicornis* is found in a wide range of reef environments on other Indo-Pacific reefs and is often the dominant browser (e.g., Hoey & Bellwood 2009; Vergés et al. 2012; Rasher et al. 2013; Plass-Johnson et al. 2015; Tebbett et al. 2020). Lower abundances may be an artefact of overexploitation as, although both *Naso* species are heavily targeted by fisheries across the Indo-Pacific, *N. unicornis* appears less resilient to fishing pressure (Bejarano et al. 2013; Ford et al. 2016; but see Taylor et al. 2019).

The white-spotted rabbitfish *Siganus canaliculatus* is generally considered an inshore species (Fox & Bellwood 2008; Cvitanovic & Bellwood 2009; Hoey et al. 2013) and is rarely recorded from outer reefs on the GBR (Cheal et al. 2012; Hoey et al. 2013; McClure et al. 2019). However, that pattern was reversed around the Tigak Islands where *S. canaliculatus* was more abundant and took more bites at sites that were closer to open ocean (<2 km) and/or surrounded by extensive seagrass beds (>17% of 250 m radii). The difference in distribution patterns may be a result of the compressed inshore-offshore gradient enabling *S. canaliculatus* individuals to occupy habitats otherwise unavailable to them. For instance, the configuration of habitat patches in Kavieng may facilitate movement from juvenile seagrass nursery habitats (Grandcourt et al. 2007; Lin et al. 2019) to preferred open ocean spawning grounds (Hasse et al. 1977) which may not be feasible across the cross-shelf distances of the GBR. The findings that this species appeared to use two reef habitats with differing seascape structures in this location (e.g., close to open ocean and/or surrounded by extensive seagrass beds) could be indicative of such an ontogenetic habitat shift but detailed size-based studies would be needed to test this hypothesis. The fewer bites and observations of *S. canaliculatus* around lagoon islands could also reflect the availability of other dietary resources in adjacent habitats (Hoey & Bellwood 2010a; Bennett & Bellwood 2011) given that *S. canaliculatus* can feed on seagrass as well as macroalgae (Lam 1974), and vegetated habitat patches are more common inside the lagoon compared to the barrier reef sites. Future research that compares juvenile abundances and feeding activity across a range of habitats (e.g., coral reefs, macroalgal and seagrass beds) may help us to obtain a more complete picture of distribution and browsing patterns for this species.

Of the six key browsing species, only the rabbitfish *Siganus doliatus* was widely distributed across all reefs around the Tigak Islands. Indeed *S. doliatus* had no apparent preference for a particular benthic community or seascape type, which supports the notion that *S. doliatus* is more of a habitat generalist on reefs than other browsing species (Bennett & Bellwood 2011; Richardson et al. 2020). Such flexibility is likely attributable to its varied diet as it is known to feed on a diverse range of algae (Fox et al. 2009; Hoey et al. 2013; Loffler et al. 2015), as well as its small body size compared to the other key browsers which may enable it to access more cryptic alga in structurally complex habitats (Bennett & Bellwood 2011; Richardson et al. 2020). *S. doliatus* was also the only species consistently recorded browsing around lagoon islands, and accounted for the majority of browsing activity at two of the lagoon islands (i.e., Eruk and Kabotteron). This suggests that this species might be important for maintaining the coral-algal balance around lagoon sites. However, *S. doliatus* is recognised as primarily a cropper of red and green algae, and its efficacy in the removal of brown, fleshy macroalgae has been questioned on several occasions (Fox et al. 2009; Streit et al. 2015). Rather, it is hypothesised that its bites are targeting epiphytes on the macroalgae rather than the algae (Streit et al. 2015). Therefore, its ability to make a significant contribution to macroalgal removal on lagoonal reefs is likely to be limited.

The third key family of browsers recorded from this location were kyphosids. Of the two species recorded (i.e., *Kyphosus cinerascens* and *K. vaigiensis*), both were patchily distributed and no clear patterns emerged to indicate whether these species were responding to certain aspects of the benthic community or seascape structure. Data on the ecology of tropical kyphosids is relatively limited but *K. vaigiensis* is often a significant contributor to macroalgal removal on reefs (e.g., Hoey & Bellwood 2011; Bennett & Bellwood 2011; Vergés et al. 2012; Michael et al. 2013) and is recognised as a true browser of brown macroalgae (Clements & Choat 1997; Choat et al. 2004). In contrast, little is known about the ecology or movement patterns of *K. cinerascens*, but its diet on the GBR has been reported to primarily consist of filamentous and foliose red and green algae (Choat et al. 2004). For *K. vaigiensis*, its patchy distribution around the Tigak Islands may be a reflection of its high mobility (>2 km daily movements, Welsh & Bellwood 2014) and efficiency at finding cryptic resources throughout the seascape (Hoey & Bellwood 2010a). For highly mobile species with such opportunistic feeding strategies, more temporal and spatially extensive sampling approaches will be necessary to identify whether benthic or seascape variables influence spatial distribution and browsing patterns.

### 5.4.3 *Limitations*

The study was conducted over a relatively short timeframe (2 weeks), and as such provides a snapshot into the process of macroalgal removal in this system. Trophic subsidies, primary productivity and physical conditions can change between habitats and seasons (Polis et al. 1997; Marczak et al. 2007; Bejarano et al. 2017) which may in turn alter browser behaviour or distribution patterns. Extending the study across a longer time period would provide greater insight into the consistency of the patterns observed, and the stability of browsing fish assemblages. However, despite the limited temporal scale, it was still possible to detect trends in the data such as the dominant species, and higher macroalgal removal rates around barrier reef islands, and these results were comparable with studies elsewhere. In addition, only one spatial scale at the seascape level (i.e., 250 m) was considered here, however, individuals can respond to their environment across different spatial scales depending on life history stage, ecology and movement capabilities (Jackson & Fahrig 2015; **Chapter 4**). Evaluating browsing activity across a greater range of spatial scales in the future may provide additional insights into macroalgal removal on coral reefs.

### 5.4.4 *Conclusions*

Browsers perform a critical function on reefs by removing the large, fleshy macroalgae that have been linked to phase shifts on coral reefs (e.g., Bellwood et al. 2006b). However, browsing activity is often patchily distributed across space (Cvitanovic & Bellwood 2009) with limited understanding about the drivers behind this variability. Here, I show that processes operating across different spatial scales, including the surrounding seascape, contribute to the function of browsing on tropical coral reefs. Distance to open ocean appeared particularly important for browsing activity with macroalgal removal higher at sites closer to open ocean. However, this was strongly influenced by a single species (*Naso lituratus*) with a high bite rate that exhibited a clear preference for higher energy outer reef sites. As coral reefs reconfigure and reassemble under an increasing number and intensity of anthropogenic stressors (Graham et al. 2013b; Williams & Graham 2019), it is imperative to better understand habitat preferences of key browsing species to identify what factors might affect the critical ecosystem function of browsing. Adopting a seascape perspective will provide additional insight into what factors drive spatial patterns of both rates and agents of browsing on tropical coral reefs.

## Chapter 6

### General Discussion

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#### 6.1 Thesis contribution and key findings

Coral reef ecosystems are recognised as hyperdiverse environments (Barlow et al. 2018), with high cultural, ecological and economic values (Moberg & Folke 1999), but they face an uncertain future due to the effects of anthropogenic-induced disturbances (e.g., climate change, Hughes et al. 2017b). In many locations, significant changes to coral reef communities have already occurred, including coral loss (Bruno & Selig 2007; Hughes et al. 2018), flattening of reef structures (Alvarez-Filip et al. 2009) and the reconfiguration of coral communities into novel assemblages (Graham et al. 2013b; Williams & Graham 2019). These changes to coral reef habitats have caused substantial changes in the composition and functional structure of fish communities (Jones et al. 2004; Graham et al. 2006; Pratchett et al. 2008; Pratchett et al. 2011; Richardson et al. 2018; Stuart-Smith et al. 2018). But coral reefs rarely exist in isolation and are typically surrounded by a range of other habitat types (e.g., mangrove forests, seagrass beds, Ogden 1988; Parrish 1989) which are also subject to ongoing losses, fragmentation and degradation (Halpern et al. 2019). Yet we still know comparatively little about how reef fish taxa interact with, and are influenced by, other habitats in the seascape. This thesis provides novel insights into the ubiquity of multiple-habitat use among tropical reef fishes, the relative importance of seascape versus within habitat patch attributes in shaping tropical fish assemblages and how the spatial context of a reef can influence the delivery of a key ecological function. Collectively the findings of this thesis build upon an emerging body of research on seascape ecology (e.g., Boström et al. 2011; Pittman et al. 2011; Olds et al. 2016; Pittman 2018) and highlights the need to adopt a broader seascape perspective when evaluating ecological patterns and processes on coral reefs.

Over the last few years, seascape ecology has burgeoned into an exciting, fast-paced discipline that has the potential to transform the way we think about and manage marine ecosystems. Narrowing our perspective to a single habitat type using a patch-centric approach (Luckhurst & Luckhurst 1978; Heck & Orth 1980; Roberts & Ormond 1987; Sale 1991) is becoming outdated. Indeed, studies using a seascape ecology framework have revealed that the spatial arrangement and composition of habitat types and patches can have a significant influence on species distribution and abundance patterns (e.g., Pittman & Olds 2015; Pittman 2018; van Lier

et al. 2018; Berkström et al. 2020). Despite its utility, there has been a relatively slow uptake of seascape ecology perspectives among coral reef ecologists, where the majority of research continues to focus almost exclusively on reef habitat, either within-reef (e.g., coral cover, Russ et al. 2021; structural complexity, Darling et al. 2017; Agudo-Adriani et al. 2019) or between-reefs (e.g., larval dispersal, Abesamis et al. 2017; Almany et al. 2017; Harrison et al. 2020) when measuring the effects of habitat composition and structure on fish community structure and ecosystem functioning.

A potential hurdle to the widespread adoption of a broader seascape perspective in the discipline of coral reef fish ecology is a limited understanding of the prevalence of multiple habitat use among reef fishes. Prior to this thesis, data on multiple habitat use had never been synthesised to examine the diversity of reef fish taxa that are observed outside of coral reef habitats or to describe species-specific habitat use patterns. **Chapter 2** addressed this fundamental knowledge gap by providing the first global synthesis of non-reef habitat use among fishes commonly associated with coral reef habitats. The synthesis revealed that non-reef habitat use is extensive and widespread among reef fish taxa, and includes ecologically important herbivores (e.g., parrotfishes and rabbitfishes), economically important fishes (e.g., emperors, grunts, and snappers) as well as species threatened with extinction (e.g., Nassau grouper and Māori wrasse). I estimated that approximately 20% of fish species classified as “reef-associated” are also using non-reef habitats, and suggest that this figure is likely conservative and may be considerably higher. In **Chapter 3**, the conservative nature of this estimate of non-reef habitat use was further supported by an in-depth examination of shallow-water habitat use in a diverse seascape in Kavieng, Papua New Guinea. In this study location, 35% of the reef-associated fish species were observed in non-reef habitats. Findings from these two thesis chapters (**Chapters 2 & 3**) suggest that we are almost certainly underestimating the importance of non-reef habitats and the broader seascape on coral reef systems, and consequently, predictions about the future outlook of coral reefs and their associated fish communities could be biased by an incomplete picture.

One of the common hypotheses to explain the observed use of multiple habitats by reef fishes is that non-reef habitats are primarily the domain of juveniles, with these habitats serving as nurseries (Nagelkerken et al. 2002; Dorenbosch et al. 2006b). My thesis provided evidence to support the use of non-reef habitats by juvenile fishes through both the global synthesis (**Chapter 2**), and video and diver-based surveys (**Chapters 3 & 4**) that showed juveniles from

a wide variety of taxa occur in non-reef habitats, often frequently. Significantly, by applying seascape ecology principles that consider the spatial configuration of different habitats and adopting a multi-scale approach, **Chapter 4** is one of the first studies to show that the distribution patterns of both juvenile emperors and parrotfishes within seagrass patches is influenced, at least to some degree, by the spatial arrangement and type of habitats within the surrounding seascape. Moreover, juveniles from these two families responded to different seascape variables suggesting that nursery habitat selection is more complex than simply locating and settling into a particular habitat type. Adopting a more mechanistic approach to examine why juveniles are using different habitats and habitat patches (e.g., enhanced growth, refugia from predation, Grol et al. 2011b, Grol et al. 2014), will help us to better understand the drivers of settlement across a broader range of fish taxa and can help design strategies for the potential replenishment of reef fish populations.

In temperate reef environments, macroalgae has long been recognised as important habitat for fishes, particularly as settlement habitat for some species (e.g., Anderson 1994; Carr 1989). Yet in tropical coral reef seascapes, macroalgae is widely viewed as a sign of reef degradation. Areas of macroalgae cover on reefs are often associated with deteriorating reef conditions or phase-shifts (Hughes 1994; McCook 1999; Cheal et al. 2010; Graham et al. 2015) and macroalgae cover impedes coral recruitment and hence recovery of coral populations (e.g., Kuffner et al. 2006; Hughes et al. 2007b). However, its natural occurrence as a habitat in its own right is sometimes overlooked (but see Tano et al. 2017; Fulton et al. 2020), particularly by coral reef scientists. Throughout this thesis macroalgal beds were identified as important habitat for some fishes or were influential in explaining species distribution patterns across the seascape (**Chapters 2, 3 & 4**). Interestingly, while the global synthesis (**Chapter 2**) identified differences in habitat use patterns for seagrass and mangroves between the Tropical Atlantic and Indo-Pacific, there was consensus in terms of macroalgal bed use. Macroalgae beds were predominantly used by juvenile rather than adult reef fishes, indicating that macroalgal beds could play an important, though currently underappreciated, nursery role for tropical fishes. In **Chapter 3**, using the Tigak Island seascapes in Papua New Guinea, I showed that macroalgal beds contained more species and higher abundances than mangrove and seagrass habitat types, and the presence of macroalgal beds had a positive effect on the abundance of wrasses in neighbouring seagrass patches (**Chapter 4**). Together, these findings lend strong support to a growing body of research (Eggertsen et al. 2017; Fulton et al. 2019; Fulton et al. 2020) showing that naturally occurring macroalgal beds are a key habitat for many fish species and that their

presence can have positive effects on fish abundance in multiple habitat types found in tropical seascapes.

## **6.2 Conservation and management implications**

The sizeable number of fish species identified as multi-habitat users, including many that are ecologically important, fisheries targets, and/or threatened with extinction, highlights the importance of holistic approaches that consider multiple aspects of the seascape to inform the effective conservation and management of shallow-water tropical seascapes. Scaling up research to understand fish-seascape relationships (**Chapters 2, 3 & 4**), how these might influence ecological processes (**Chapter 5**) and contribute to ecosystem integrity will be critical in the development and implementation of spatial management strategies that are fit for purpose.

This thesis provides new insights into why it is so critically important to include a range of habitat types in spatial management plans. Designing marine protected area networks based on the CAR principles (Comprehensiveness, Adequacy and Representative) has become common practice (e.g., Great Barrier Reef Marine Park, Australia, Fernandes et al. 2005). The underlying theory in this approach is that by incorporating multiple habitat types and patches of different sizes, spatial configurations and locations, there is a greater chance of capturing the full complement of biodiversity within a seascape, protecting critical support areas for growth, reproduction or survival (e.g., juvenile nurseries), maintaining ecological integrity, and supporting population viability (Margules & Pressey 2000; Day et al. 2002). Until recently, we knew little about the extent to which these assumptions were true. The widespread use of multiple habitat types by such a significant portion of the fish community documented in this thesis provides strong support for including a range of habitat types in spatial management plans. However, it also revealed new insights that could be used to refine spatial planning strategies that incorporate multiple habitats. Discrete patches of the same habitat type can support quite distinct fish communities (**Chapter 4**), both in terms of taxonomic and functional composition, and differences in the spatial context of a patch can flow through to the delivery of key ecological functions (e.g., browsing, **Chapter 5**). The findings of this thesis indicate that the positioning and placement of marine protected areas, or the choice of which habitat patch to protect will need to be tailored to the objectives of the reserve or reserve network (e.g.,

to conserve biodiversity, maintain ecosystem integrity, protect essential fish nursery habitat or to manage fisheries).

The findings from this thesis are also relevant for fisheries management. For example at my study site in the Tigak Islands, seagrass meadows, macroalgal beds and mangrove habitats were extensively used by both adult and juvenile reef fishes (**Chapters 3 & 4**), and the majority of multiple habitat-using taxa are fisheries targets (e.g., emperors, snappers). Moreover, my findings also suggest that some habitat patches may support higher abundances and diversity of fisheries species than other patches of the same habitat type (**Chapter 4**). Human communities in this location, like many across the Indo-Pacific, are heavily reliant on healthy marine ecosystems and fisheries stocks (Bell et al. 2009; Gillet 2014). Although a lot of attention is given to coral reef fisheries, it is vital to recognise that other habitats are also important in their own right, and in the context of coral reef fisheries management. Both seagrass beds and mangroves support highly productive fisheries (e.g., mangroves, Sheaves 2017; Carrasquilla-Henao et al. 2019; seagrass, Nordlund et al. 2018), and are accessible year round unlike coral reefs which can be challenging to access during adverse weather conditions, making sheltered vegetated habitat types a more predictable source of fish protein (e.g., during monsoon season). In addition, if we assume connectivity between habitat patches, then mangroves, macroalgal and seagrass beds are not only a valuable resource for fisheries in their own right, but could also enhance fish populations on reefs. Therefore, adopting a seascape perspective for fisheries management may be critical for longer term food security, particularly for the Pacific Island Countries and Territories that have extensive nearshore vegetated habitats.

The long-term viability of many tropical shallow-water habitats is uncertain given the diverse range of stressors these habitats are exposed to (e.g., coastal developments, elevated sea surface temperatures, severe storms, pollution, Jackson et al. 2001; Bellwood et al. 2004; Halpern et al. 2019). These threats have led conservation and management efforts to focus on maintaining key functions to enhance resilience and support recovery processes (e.g., Nyström & Folke 2001; Harun-or-Rashid et al. 2009; Graham et al. 2011; Rasheed et al. 2014). While some approaches to build resilience have considered connectivity between different habitat types (e.g., MPA networks, Green et al. 2009; Bernhardt & Leslie 2013; Torres-Pulliza et al. 2013), others have focused on implementing measures within a single habitat type. For instance, one measure that has been repeatedly proposed is to restrict fishing of taxa that perform key

ecological roles (e.g., herbivores, McCook 1999; Hughes et al. 2007b; Graham et al. 2013a; MacNeil et al. 2015; Williams et al. 2019; Mumby et al. 2021), but solely on reefs rather than the range of habitats used by herbivores. Yet the findings from this thesis suggest quite strongly that there will be considerable value in thinking about herbivore management, and other resilience building strategies, within a seascape context. For example, herbivores such as parrotfishes and rabbitfishes were revealed to use multiple habitat types, not just coral reefs, throughout their lives (**Chapters 2 & 3**), and their spatial distribution (**Chapter 4**) and the functional impact of browsers was influenced by seascape structure (**Chapter 5**). Management strategies that seek to conserve herbivores, for example threatened species that have distinct juvenile habitats (e.g., the bumphead parrotfish *Bolbometopon muricatum*, Hamilton et al. 2017; and the rainbow parrotfish *Scarus guacamaia*, Mumby et al. 2006), in order to promote reef resilience will be more effective if they incorporate connections between different habitat types in the seascape.

As well as resilience building, there is a growing need for active interventions to restore habitats or critical ecosystem functions (Bosire et al. 2008; Hein et al. 2021; Sinclair et al. 2021). However until recently, restoration was focused on a single habitat type rather than considering the value of interactions among multiple habitat types found in the seascape mosaic (Gilby et al. 2018; Pittman et al. 2021). One prominent effect of habitat loss is that habitat patches may become smaller, more fragmented and increasingly isolated, which can cause the breakdown of critical relationships between habitats (Saunders et al. 1991; Steffan-Dewenter & Tschardtke 1999; Fischer & Lindenmayer 2007; Fahrig 2013). As we enter the UN Decade of Ecosystem Restoration (Waltham et al. 2020), spatially explicit approaches that consider aspects such as the composition of the surrounding matrix may be more effective in restoring habitats and associated fauna than single habitat approaches by re-establishing functional connections (Hale et al. 2020), such as seascape nurseries (Nagelkerken et al. 2015). However, the success of seascape restoration approaches will ultimately depend on whether we address the dominant stressor of climate change given it will particularly impact shallow-water ecosystems, especially those in close proximity to the coast (Duarte et al. 2020; Kleypas et al. 2021).

### **6.3 Future research**

The findings of this thesis open up an interesting interface between traditional ecological theory and landscape perspectives. Our understanding about how multiple habitat or patch use, and

spatial context can affect individual fitness is limited. As a consequence, incorporating spatially explicit approaches into empirical tests of key ecological theories (e.g., optimal foraging, landscape of fear) may reveal new and valuable insights into population and community dynamics. As a relatively new field, there are also many fruitful avenues of seascape research to be pursued which will progress the field from testing landscape ecology principles, to developing a framework more suitable for marine environments, and ultimately facilitating the transition into a solutions-oriented science that can support the long-term persistence and health of marine environments. Indeed, a recent horizon scan has identified some of the most pressing seascape-related questions among academics and practitioners that need to be addressed to advance the field (Pittman et al. 2021). Themes identified as important include changing seascapes, connectivity, ecosystem-based management (EBM), restoration, and seascape goods and services (Pittman et al. 2021), with knowledge gained from this thesis directly contributing to several of these. The findings from this thesis have provided some novel insights and made a valuable contribution to expanding our appreciation about the influence of the seascape on the structure and function of tropical fish communities. However, the thesis has also identified some key knowledge gaps that should form the basis of future research to more fully understand the implications of changing seascapes.

### ***6.3.1 Multiple habitat users, early life history, and the seascape mosaic***

One sizeable and longstanding knowledge gap is our limited understanding about the reasons for multiple habitat use for the vast majority of fish species. I found evidence of extensive use of multiple habitat types in many species but we do not yet understand the extent to which the use of multiple habitat types makes these species more or less vulnerable to habitat loss and changing seascapes. If species that use multiple habitats have obligate dependencies on specific resources in each habitat type then we might expect to see detrimental effects on populations if critical habitats or habitat patches are lost, whereas if the use of multiple habitats in the seascape is an indication of high levels of resource flexibility then changing seascapes may have little impact. This is likely to differ by species, such that species with greater habitat flexibility will be more likely to cope with changing conditions (e.g., Colossi Brustolin et al. 2019; Kingsbury et al. 2019; Stuart-Smith et al. 2021), and importantly may hold the key to future diversification across longer evolutionary timeframes (Gajdzik et al. 2019). Multiple habitat users may increase response diversity by responding to environmental perturbations at different spatial scales than species constrained to a single habitat type (Elmqvist et al. 2003), with the potential to relocate away from patches if conditions become suboptimal (Lenihan et

al. 2001), only returning when conditions become more favourable. Through their habitat flexibility, some multiple habitat users could contribute towards recovery processes through larval supply, population replenishment into depauperate habitat patches, or by performing an essential role that can support recovery (e.g., grazing).

Processes operating during early life stages may be important drivers of multiple habitat use in many taxa. Juveniles from a wide range of taxa were recorded using multiple habitat types (**Chapter 2**), particularly from families that are economically (e.g., emperors) or ecologically important (e.g., herbivorous fishes). In addition, juveniles from separate families (e.g., emperors and parrotfishes) appeared to use different types of seagrass beds depending on the seascape context and within-patch attributes (**Chapter 4**). However, it is currently impossible to determine whether these patterns are a result of active habitat patch selection at settlement and/or differential post-settlement survival in different habitat types and patches. Few studies have directly quantified settlement of fishes into a range of habitat types (but see Rooker et al. 1996; Nakamura et al. 2009b, Hedberg et al 2018), and even fewer within a seascape context (but see Nakamura et al. 2009a). In part, progress in this area has been impeded due to the challenge of accurate identification to species level for some early life stages (e.g., juvenile parrotfishes). However, advances in DNA analyses are helping to overcome this issue and have begun to reveal interesting insights into habitat use patterns of early stage juveniles (e.g., Sievers et al. 2020a).

Collecting data from a wider range of habitats, habitat patches, and seascapes situated in different environmental and spatial contexts will be important in furthering our understanding of multiple habitat use patterns by tropical fishes. One way this is likely to be achieved is through the increased use of remote video cameras in combination with the rapidly advancing field of computer vision. Deploying cameras across a more extensive range of habitats and habitat patches will be less time intensive than traditional diver-led surveys, can capture habitat use patterns of diver-averse taxa, and is easily replicable across different seascapes (Mallet & Pelletier 2014; Weinstein 2018). Moreover, a number of projects are currently underway to improve battery performance and night-vision capabilities so that data collection can occur across longer time periods (e.g., Coro & Bjerregaard Walsh 2021), particularly useful for species that undertake day-night shifts between habitat types. Progress in computer vision technology (e.g., onboard processing, motion detection for more efficient data capture, automated detection of species, body sizes, and relative abundances) will also considerably

reduce the time traditionally involved in manual data extraction from videos (Wäldchen & Mäder 2018; Lopez-Marcano et al. 2020; Coro & Bjerregaard Walsh 2021). In addition to video data, molecular techniques will also help unravel habitat use patterns of rare species (e.g., eDNA, Budd et al. 2021) and cryptic life stages (e.g., Sievers et al. 2020a). Scaling up research using more spatially extensive approaches such as remote cameras will lead to more accurate predictive models about species distribution patterns that can inform future spatially explicit conservation and management strategies.

More intensive research that explores the underlying drivers of habitat use patterns is essential in order to interpret findings emerging from seascape research in an ecologically meaningful way and to fully comprehend the implications of changing seascapes. For instance, delving into why discrete taxa and life stages were using different types of seagrass patches (**Chapter 4**) is an important next step emerging from this thesis. Identifying mechanisms underpinning multiple habitat use will necessitate a combination of field, laboratory, and modelling techniques such as quantifying larval supply into a range of habitats and habitat patches, testing habitat preferences; evaluating survival and growth rates of post-settlement fishes (e.g., food availability, predation, competition) and; examining how the intervening matrix between juvenile and adult habitats influences thresholds of connectivity. Progressing beyond observational studies towards more mechanistic approaches will help us to evaluate whether coral reef-associated taxa will be able to switch to, or persist in, non-reef habitat types, and better predict the consequences of habitat loss or fragmentation on individual taxa.

### **6.3.2 Behavioural seascape ecology**

Another key area for future research is improving our knowledge about how the seascape structure can affect patterns of movement. This thesis found that different taxa and life stages responded to different spatial scales (e.g., from metres to 1 km radii), similar to other seascape studies (e.g., Yeager et al. 2011; Olds et al. 2012a; Sievers et al. 2020b). Although the spatial scales examined in seascape studies are often similar, there can be considerable variability in the spatial scale identified as influential for species distribution and abundance patterns, between study locations and even taxa. These differences among studies may be caused by differences in the spatial arrangement and composition of habitat patches within the focal seascape, however this topic has not yet been explored. Greater information on species-specific movement patterns, as well as intra-specific variability may therefore be valuable to help us identify features that might facilitate or impede movement (Taylor et al. 1993). Acoustic

tracking capabilities have become increasingly refined in recent years and will be useful to shed light on this topic given that manually (i.e., divers) following individual fishes over long periods is not feasible. However, despite recent advances in satellite, underwater and aerial drone imagery capture and processing, it remains challenging to produce high resolution, accurate mapping of seascape habitats across broad areas (Benoist et al. 2019; Joyce et al. 2019; Lyons et al. 2020). Therefore few movement ecology studies have explicitly examined movement between different habitats (but see Hitt et al. 2011; Rooker et al. 2018; Ebrahim et al. 2020; Heidmann et al. 2021), instead quantifying movement patterns in terms of migration distances or home ranges. More study designs that enable individuals to be tracked between different habitats are therefore needed. One of the few studies to explore movement in two common reef fishes within a seascape context found that predator avoidance influenced movement patterns and patch occupation (Rooker et al. 2018). Moreover, few studies have formally tested for matrix effects on movement patterns, despite the fact that the type of matrix has been implicated in declines in species numbers away from nursery habitats (e.g., deep water, sand, Berkström et al. 2020 but see Pagès et al. 2014), as well as the potential for certain types of matrix to act as a movement corridor between two spatially separated patches (Davis et al. 2017). Clearly, there is much to learn about the factors that influence movement patterns for fishes in coastal marine seascapes.

### ***6.3.3 Seascape ecology and ecosystem functioning***

Understanding how the spatial arrangement and composition of habitat patches influences ecosystem dynamics and functioning across seascapes, and differing spatial scales, should be a priority area for future research. To date, few studies have considered how the wider seascape can influence ecosystem functioning, and even in terrestrial systems, matching ecological processes with spatial patterns remains a challenging research frontier (Turner & Gardener 2015). Rather, there is a tendency to make assumptions based on fish-seascape patterns which may not be a robust substitute for function. This thesis found that the spatial context of a focal reef influences macroalgal browsing rates on coral reefs, with higher browsing rates on outer reefs with little vegetated habitat nearby (**Chapter 5**). Based on these findings, future areas of research should include examining macroalgal removal on reefs from similar spatial contexts (e.g., similar distances from open ocean) but with differing availabilities of vegetated habitat types (e.g., seagrass, macroalgal beds), as well as examining feeding behaviour across a range of habitat types including reefs to determine foraging flexibility among browsing taxa. Expanding this approach to examine other types of herbivorous fishes should be a priority for

future research. For instance, grazing parrotfishes perform a critical role by preventing the establishment of macroalgae (Bellwood et al. 2004; Mumby et al. 2006; Burkepile & Hay 2008; Adam et al. 2011), commonly use non-reef habitats (Campbell et al. 2011; Sambrook et al. 2020; Sievers et al. 2020a; **Chapters 3 & 4**), and have been implicated in reef recovery following disturbance events (Mumby & Hastings 2008; Adam et al. 2011; Harborne et al. 2016b). Therefore understanding how the surrounding seascape can influence grazing patterns on reefs warrants greater attention.

The application of big data approaches (e.g., global datasets) will allow us to address some of these knowledge gaps and rapidly advance our understanding of patterns, processes, and scales within a seascape context. One of the major challenges for advancing seascape ecology is the difficulty associated with collecting and analysing sufficient amounts of data from underwater environments. This can limit our ability to detect relationships, identify commonalities and differences between seascapes, and test predictions. Much of the data that could enable us to test out seascape-related hypotheses across greater spatial extents may already exist but will require collaboration, multi-disciplinary teams and access to historical satellite imagery. For instance, herbivory on both coral reefs and in seagrass beds has been extensively explored across multiple locations (e.g., coral reef: Great Barrier Reef & Ningaloo, Australia, Vergés et al. 2012, Fiji, Rasher et al. 2013; Indonesia, Plass-Johnson et al. 2015; Singapore, Bauman et al. 2017; Tanzania, Eggertsen et al. 2019; Palau, Roff et al. 2019; and seagrass herbivory: Indonesia, Unsworth et al. 2007c; Puerto Rico, Swindells et al. 2017; Tanzania, Eggertsen et al. 2019; Australia, Vanderklift et al. 2021). Bringing these data together and examining different types of herbivory within a seascape context may reveal valuable insights and hidden connections between different elements of the seascape.

#### **6.4 Concluding remarks**

Overall, this thesis has demonstrated that multiple habitat use by tropical fishes is widespread, perhaps more so than previously recognised, and that by looking in a wider range of habitat types, we can gain a deeper understanding of fish-habitat relationships. In addition, this thesis has shown that patches of the same habitat type may not be functionally equivalent, with the taxa and life stage of fish occupying a patch, and the delivery of their function also moderated by seascape context. Finally, this thesis demonstrated that the spatial context of a reef is an important driver of a key ecological process (i.e., macroalgae browsing) on reefs. As a body of

work, this thesis demonstrates that the seascape matters and supports growing calls to transition from a habitat- or patch-centric approach, to approaches that encompass the surrounding seascape. Adopting a seascape ecology approach will give us a more complete picture of connections between habitats, how such connections might influence patterns and processes, and what that might mean as seascapes continue to change into the future.

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## Appendices

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## Appendix A: Supplementary materials for Chapter 2

### *Beyond the reef: the widespread use of non-reef habitats by coral reef fishes*

**Table A2.1.** Coral reef fish species observed in non-reef habitats by life stage. Habitat categories: SG = seagrass, MG = mangrove, MA = macroalgae beds, SS = soft sediment, E = estuarine environments, OT = Other (includes non-coral dominated channels, gorgonian plains, mudflats, boulders, notches and transition habitat between reef and seagrass). Observations were recorded as follows: X = observed in habitat, 0 = habitat surveyed but species not observed, - = habitat not surveyed. Criteria for inclusion in table: 1) in each study, individuals were observed on coral reefs AND in non-reef habitat, and; 2) species were classed as coral reef-associated (*FishBase*; Froese & Pauly 2018).

Species	All						Juveniles						Adults						References
	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	
<b><u>Tropical Atlantic</u></b>																			
Acanthuridae																			
<i>Acanthurus bahianus</i>	X	X	X	0	X	X	X	X	X	-	-	X	X	0	0	-	-	0	1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19
<i>Acanthurus chirurgus</i>	X	X	X	X	X	X	X	X	X	-	-	X	X	X	0	-	-	X	1, 2, 3, 4, 5, 6, 7, 8, 10, 11, 12, 13, 14, 15, 16, 18, 19, 20
<i>Acanthurus coeruleus</i>	X	X	X	0	X	X	X	X	X	-	-	X	0	0	0	-	-	0	2, 4, 6, 7, 8, 10, 12, 13, 14, 15, 16, 19, 21, 22
Albulidae																			
<i>Albula vulpes</i>	X	X	-	-	-	-	X	X	-	-	-	-	-	-	-	-	-	-	23
Apogonidae																			
<i>Apogon binotatus</i>	0	X	-	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	24
<i>Apogon lachneri</i>	X	X	-	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	24
<i>Apogon maculatus</i>	X	X	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	24
<i>Apogon planifrons</i>	0	X	-	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	24
<i>Apogon townsendi</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	18

	All						Juveniles						Adults						
Species	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	References
<i>Phaeoptyx conklini</i>	0	X	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	24
<i>Phaeoptyx pigmentaria</i>	0	X	-	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	24
<i>Phaeoptyx xenus</i>	0	X	-	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	24
Atherinidae																			
<i>Atherinomorus stipes</i>	X	X	-	X	-	-	X	X	-	X	-	-	-	-	-	-	-	-	25
Aulostomidae																			
<i>Aulostomus maculatus</i>	X	X	0	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	8,14,26
Balistidae																			
<i>Balistes vetula</i>	X	0	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	4,8
<i>Melichthys niger</i>	0	0	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	8
Blennidae																			
<i>Ophioblennius trinitatis</i>	0	-	X	-	-	-	0	-	0	-	-	-	0	-	X	-	-	-	6
Carangidae																			
<i>Carangoides bartholomaei</i>	X	-	X	-	X	X	X	-	X	-	-	-	0	-	0	-	-	-	4,6,8,16
<i>Caranx crysos</i>	X	X	-	-	-	0	-	-	-	-	-	-	-	-	-	-	-	-	7,8
<i>Caranx latus</i>	X	X	0	-	X	X	-	-	-	-	-	-	-	-	-	-	-	-	14,16
<i>Caranx ruber</i>	X	X	0	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	4,7,8,14,18
<i>Seriola rivoliana</i>	0	0	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	8
Chaetodontidae																			
<i>Chaetodon capistratus</i>	X	X	X	-	-	X	X	X	-	-	-	X	X	X	-	-	-	X	2,4,7,8,9,12,13,14,15,18,19,22,27
<i>Chaetodon ocellatus</i>	0	X	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	7,8,27
<i>Chaetodon striatus</i>	0	X	X	0	X	X	0	-	X	-	-	-	0	-	0	-	-	-	1,2,6,7,8,16,27
<i>Prognathodes aculeatus</i>	0	0	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	8
Cirrhitidae																			
<i>Amblycirrhitis pinos</i>	0	0	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	8
Dasyatidae																			
<i>Hypanus americanus</i>	X	0	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	7,8
Diodontidae																			
<i>Diodon holocanthus</i>	X	X	X	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	2,14
<i>Diodon hystrix</i>	X	X	0	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	2,14,21

	All						Juveniles						Adults							
Species	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	References	
Engraulidae																				
<i>Anchoa lyolepis</i>	X	X	-	X	-	-	X	X	-	X	-	-	-	-	-	-	-	-	-	25
Ephippidae																				
<i>Chaetodipterus faber</i>	-	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	16
Gerreidae																				
<i>Gerres cinereus</i>	X	X	X	X	X	X	X	X	-	-	-	X	X	X	-	-	-	-	X	1,2,4,8,9,12,13,14,16,20,21,22,28
Gobiidae																				
<i>Coryphopterus glaucofraenum</i>	X	X	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	4,7,18
<i>Coryphopterus personatus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	18
<i>Gnatholepis thompsoni</i>	-	0	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	4
Grammatidae																				
<i>Gramma loreto</i>	0	0	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	8
Haemulidae																				
<i>Anisotremus surinamensis</i>	0	X	0	-	-	0	0	-	0	-	-	-	0	-	0	-	-	-	-	6,8,14
<i>Anisotremus virginicus</i>	X	X	0	-	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	2,4,7,14,16,18
<i>Haemulon album</i>	X	0	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	8
<i>Haemulon aurolineatum</i>	X	X	X	-	X	X	X	X	0	-	-	-	0	0	0	-	-	-	-	2,3,4,6,7,9,10,16
<i>Haemulon carbonarium</i>	X	X	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	8,10,27
<i>Haemulon chrysargyreum</i>	X	X	0	-	-	X	X	X	-	-	-	X	X	X	-	-	-	-	X	12,13,14
<i>Haemulon flavolineatum</i>	X	X	X	X	-	X	X	X	-	X	-	X	X	X	-	-	-	-	X	1,2,4,5,7,8,9,10,11,12,13,14,15,18,19,20,21,22,27,28,29,30,31,32,33,34,35
<i>Haemulon macrostomum</i>	X	X	-	-	-	0	-	-	-	-	-	-	-	-	-	-	-	-	-	4,7
<i>Haemulon parra</i>	X	X	X	-	X	X	X	X	X	-	-	0	X	X	0	-	-	-	0	2,6,7,8,12,16,20,22,28
<i>Haemulon plumierii</i>	X	X	X	X	-	X	X	X	X	-	-	-	0	0	0	-	-	-	-	1,2,4,6,7,8,9,10,11,13,14,21,22,27,29,30,36,37
<i>Haemulon sciurus</i>	X	X	X	X	-	X	X	X	-	-	-	X	X	X	-	-	-	-	X	2,3,4,5,7,8,9,12,13,14,15,20,21,22,27,28,29,30,31,33,38,39
Holocentridae																				

	All						Juveniles						Adults						
Species	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	References
<i>Holocentrus adscensionis</i>	0	X	0	-	X	X	0	0	-	-	-	X	0	0	-	-	-	X	2,8,14,16,20
<i>Holocentrus rufus</i>	X	X	0	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	2,4,8,14
<i>Myripristis jacobus</i>	0	X	0	-	-	0	-	-	-	-	-	-	-	-	-	-	-	-	4,8,14
<i>Sargocentron vexillarium</i>	X	X	0	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	2,14
Kyphosidae																			
<i>Kyphosus sectatrix</i>	0	X	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	2,7,8
Labridae																			
<i>Bodianus rufus</i>	0	0	0	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	8,14
<i>Clepticus parrae</i>	0	0	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	8
<i>Halichoeres bivittatus</i>	X	X	X	X	-	X	X	X	-	-		X	X	X	-	-		X	1,2,4,7,8,9,12,14,18,40,41
<i>Halichoeres brasiliensis</i>	0	-	X	-	-	-	0	-	0	-	-	-	0	-	0	-	-	-	6
<i>Halichoeres garnoti</i>	X	X	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	2,4,8
<i>Halichoeres maculipinna</i>	X	0	X	0	-	X	X	0	-	-	-	X	X	0	-	-	-	X	1,8,12,14
<i>Halichoeres pictus</i>	0	0	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	8
<i>Halichoeres poeyi</i>	X	0	X	0	-	X	X	0	X	-	-	-	X	0	X	-	-	-	1,2,4,6,8,9,14
<i>Halichoeres radiatus</i>	X	X	0	0	-	X	-	-	-	-	-	-	-	-	-	-	-	-	1,7,8,14
<i>Lachnolaimus maximus</i>	X	0	0	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	2,7,8,14
<i>Thalassoma bifasciatum</i>	X	X	0	-	-	X	X	X	-	-	-	X	X	X	-	-	-	X	2,4,7,8,9,12,14,18,41
<i>Xyrichtys martinicensis</i>	-	0	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	4
<i>Xyrichtys splendens</i>	X	-	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	1
Labrisomidae																			
<i>Labrisomus nuchipinnis</i>	0	-	X	-	-	-	0	-	X	-	-	-	0	-	0	-	-	-	6
<i>Malacoctenus triangulatus</i>	0	X	X	-	-	-	0	-	0	-	-	-	0	-	X	-	-	-	4,6,18
Lutjanidae																			
<i>Lutjanus alexandrei</i>	-	X	X	X	-	-	-	X	-	-	-	-	-	0	-	-	-	-	42,43
<i>Lutjanus analis</i>	X	X	X	X	-	X	X	X	-	-	-	0	0	0	-	-	-	X	1,2,7,8,13,14,20,29
<i>Lutjanus apodus</i>	X	X	X	X	-	X	X	X	-	-	-	X	X	X	-	-	-	X	2,3,4,5,7,8,9,12,13,14,15,17,18,20,21,22,26,27,28,31,44,45
<i>Lutjanus cyanopterus</i>	0	X	-	-	-	0	-	-	-	-	-	-	-	-	-	-	-	-	8

	All						Juveniles						Adults						
Species	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	References
<i>Lutjanus griseus</i>	X	X	X	-	-	X	X	X	-	-	-	X	X	X	-	-	-	X	2,3,4,5,7,8,9,12,13,14,20,21,28,46
<i>Lutjanus jocu</i>	0	X	-	-	X	-	-	X	-	-	X	-	-	0	-	-	X	-	2,16,29,42,47
<i>Lutjanus mahogoni</i>	X	X	X	X	-	X	X	X	-	-	-	X	X	X	-	-	-	X	1,2,4,8,10,11,12,13,14,15,19,20,22,31
<i>Lutjanus synagris</i>	X	X	-	-	X	X	-	-	-	-	-	-	-	-	-	-	-	-	4,7,8,16,27,36
<i>Ocyurus chrysurus</i>	X	X	X	X	-	X	X	X	X	-	-	X	0	0	0	-	-	0	1,2,4,5,6,7,8,9,10,11,12,13,14,15,17,20,22,27,29,33,36,48
Malacanthidae																			
<i>Malacanthus plumieri</i>	X	0	X	X	-	X	-	-	-	-	-	-	-	-	-	-	-	-	1,4,8
Megalopidae																			
<i>Megalops atlanticus</i>	0	X	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	7
Monacanthidae																			
<i>Aluterus schoepfii</i>	0	0	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	8
<i>Cantherhines macrocerus</i>	0	0	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	8
<i>Monacanthus tuckeri</i>	X	0	-	-	-	0	-	-	-	-	-	-	-	-	-	-	-	-	7,49
Mullidae																			
<i>Mulloidichthys martinicus</i>	X	X	0	-	-	X	X	X	-	-	-	0	X	X	-	-	-	X	2,4,8,12,14
<i>Pseudupeneus maculatus</i>	X	X	X	X	X	X	0	-	X	-	-	-	X	-	X	-	-	-	1,2,4,6,8,14,16
Muraenidae																			
<i>Gymnothorax funebris</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	2
<i>Gymnothorax moringa</i>	0	X	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	2,8
<i>Gymnothorax vicinus</i>	-	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	16
Ophichthidae																			
<i>Myrichthys ocellatus</i>	-	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	16
Opistognathidae																			
<i>Opistognathus aurifrons</i>	-	0	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	4
Ostraciidae																			
<i>Acanthostracion polygonius</i>	0	X	0	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	8,14
<i>Lactophrys bicaudalis</i>	X	X	X	0	-	X	-	-	-	-	-	-	-	-	-	-	-	-	1,8,14
<i>Lactophrys triqueter</i>	0	X	X	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	14
Pomacanthidae																			

	All						Juveniles						Adults						
Species	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	References
<i>Holacanthus bermudensis</i>	0	0	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	7
<i>Holacanthus ciliaris</i>	X	X	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	2,7
<i>Holacanthus tricolor</i>	0	0	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	8
<i>Pomacanthus arcuatus</i>	X	X	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	2,4,7,8
<i>Pomacanthus paru</i>	X	X	X	-	X	X	0	-	X	-	-	-	X	-	0	-	-	-	1,6,7,8,14,16
Pomacentridae																			
<i>Abudefduf saxatilis</i>	X	X	X	-	X	X	0	X	X	-	-	-	0	0	0	-	-	-	2,4,6,7,8,9,13,14,16,18,19,27,28
<i>Abudefduf taurus</i>	0	0	0	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	14
<i>Chromis cyanea</i>	0	0	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	8
<i>Chromis multilineata</i>	0	0	0	-	-	X	0	0	-	-	-	0	0	0	-	-	-	X	8,12,14
<i>Microspathodon chrysurus</i>	0	0	X	-	-	X	0	-	X	-	-	-	0	-	0	-	-	-	6,8
<i>Stegastes adustus</i>	X	X	0	-	-	X	0	X	-	-	-	0	0	X	-	-	-	X	2,4,8,12,14,18
<i>Stegastes diencaeus</i>	X	X	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	4,8
<i>Stegastes fuscus</i>	0	0	X	-	X	X	0	-	X	-	-	-	0	-	X	-	-	-	6,7,16
<i>Stegastes leucostictus</i>	X	X	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	1,2,4,8
<i>Stegastes partitus</i>	X	X	0	-	-	X	0	0	-	-	-	X	0	0	-	-	-	X	1,2,4,7,8,12,14,41
<i>Stegastes planifrons</i>	X	X	0	0	-	X	0	0	-	-	-	X	X	X	-	-	-	X	2,4,8,9,12,14,41,50
<i>Stegastes variabilis</i>	0	0	X	-	X	X	0	-	X	-	-	-	0	-	X	-	-	-	6,7,8,16
Scarine																			
<i>Cryptotomus roseus</i>	-	0	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	4
<i>Nicholsina usta usta</i>	0	X	-	-	-	0	-	-	-	-	-	-	-	-	-	-	-	-	8
<i>Scarus coeruleus</i>	X	0	-	-	-	X	X	0	-	-	-	X	0	0	-	-	-	0	12,13,20
<i>Scarus guacamaia</i>	X	X	-	-	-	X	X	X	-	-	-	X	0	0	-	-	-	0	7,8,12,13,28,39,51
<i>Scarus iseri</i>	X	X	X	0	-	X	X	X	-	-	-	X	X	X	-	-	-	X	1,3,4,5,7,8,9,10,11,12,13,18,20,31,33
<i>Scarus taeniopterus</i>	X	X	X	0	-	X	0	X	-	-	-	-	0	0	-	-	-	-	1,2,5,7,8,10,22
<i>Scarus vetula</i>	0	X	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	7,8,21
<i>Sparisoma amplum</i>	0	-	X	-	-	-	0	-	X	-	-	-	0	-	0	-	-	-	6
<i>Sparisoma atomarium</i>	X	X	0	0	-	X	-	-	-	-	-	-	-	-	-	-	-	-	1,4,8
<i>Sparisoma aurofrenatum</i>	X	X	X	0	-	X	X	X	-	-	-	X	X	0	-	-	-	X	2,3,7,8,9,10,12,18,31
<i>Sparisoma axillare</i>	X	-	X	-	X	-	X	-	X	-	-	-	0	-	0	-	-	-	6,16,52
<i>Sparisoma chrysopterus</i>	X	X	-	-	-	X	X	X	-	-	-	X	X	0	-	-	-	X	4,7,8,12,13,20,27

	All						Juveniles						Adults						
Species	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	References
<i>Sparisoma frondosum</i>	0	-	X	-	-	-	0	-	X	-	-	-	0	-	0	-	-	-	6
<i>Sparisoma radians</i>	X	X	X	X	-	X	-	-	-	-	-	-	-	-	-	-	-	-	1,4,7,8
<i>Sparisoma rubripinne</i>	0	X	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	2,7,8,27
<i>Sparisoma viride</i>	X	X	X	0	-	X	X	X	-	-	-	X	0	0	-	-	-	X	1,2,3,4,7,8,12,13,15,18,19,21,50
Sciaenidae																			
<i>Equetus punctatus</i>	0	0	0	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	14
<i>Pareques acuminatus</i>	X	0	X	-	-	X	0	-	X	-	-	-	0	-	0	-	-	-	6,7
Scorpaenidae																			
<i>Pterois volitans</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	53,54,55,56
<i>Scorpaena plumieri</i>	0	X	0	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	14
Serranidae																			
<i>Cephalopholis cruentata</i>	0	0	0	-	-	X	0	0	-	-	-	0	0	0	-	-	-	X	8,12,14
<i>Cephalopholis fulva</i>	X	0	0	-	X	X	X	-	0	-	-	-	-	-	-	-	-	-	8,16,57
<i>Epinephelus adscensionis</i>	X	-	0	-	X	-	X	-	0	-	-	-	-	-	-	-	-	-	16,57
<i>Epinephelus guttatus</i>	X	0	X	0	-	X	-	-	-	-	-	-	-	-	-	-	-	-	1,8
<i>Epinephelus morio</i>	0	0	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	7
<i>Epinephelus striatus</i>	X	X	X	0	-	X	X	-	0	-	-	-	-	-	-	-	-	-	8,36,50,57
<i>Hypoplectrus aberrans</i>	0	0	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	8
<i>Hypoplectrus chlorurus</i>	0	X	0	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	2,4,8,14
<i>Hypoplectrus guttavarius</i>	0	0	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	8
<i>Hypoplectrus indigo</i>	0	0	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	8
<i>Hypoplectrus puella</i>	X	X	0	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	2,4,7,8,14,18
<i>Hypoplectrus unicolor</i>	X	X	0	-	-	X	X	X	-	-	-	X	X	X	-	-	-	X	2,12,14
<i>Mycteroperca bonaci</i>	0	X	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	7,8
<i>Mycteroperca tigris</i>	0	0	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	8
<i>Mycteroperca venenosa</i>	0	0	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	8
<i>Rypticus maculatus</i>	0	X	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	7
<i>Rypticus saponaceus</i>	0	0	0	-	X	X	-	-	-	-	-	-	-	-	-	-	-	-	14,16
<i>Serranus flaviventris</i>	-	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	16
<i>Serranus tabacarius</i>	0	0	0	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	14
<i>Serranus tigrinus</i>	X	0	0	X	-	X	-	-	-	-	-	-	-	-	-	-	-	-	1,8,18
Sparidae																			

	All						Juveniles						Adults						
Species	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	References
<i>Archosargus rhomboidalis</i>	X	X	-	-	X	X	-	-	-	-	-	-	-	-	-	-	-	-	2,7,16,21,23
<i>Calamus bajonado</i>	0	X	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	7,8,21
<i>Calamus calamus</i>	X	X	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	8,21
<i>Calamus pennatula</i>	-	0	0	0	0	X	-	-	-	-	-	-	-	-	-	-	-	-	4
<i>Lagodon rhomboides</i>	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	28
Sphyraenidae																			
<i>Sphyraena barracuda</i>	X	X	X	-	X	X	X	X	-	-	-	X	X	X	-	-	-	0	2,3,8,12,13,14,15,16,20,28
Synodontidae																			
<i>Synodus intermedius</i>	X	0	0	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	2,4,14
Tetraodontidae																			
<i>Canthigaster rostrata</i>	X	X	X	0	-	X	0	0	-	-	-	X	0	0	-	-	-	X	1,2,4,8,12,14,18
<i>Sphoeroides greeleyi</i>	-	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	16
<i>Sphoeroides spengleri</i>	X	0	0	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	2,4,14
<b><u>Indo-Pacific</u></b>																			
Species	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	
Acanthuridae																			
<i>Acanthurus auranticavus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Acanthurus blochii</i>	X	X	X	-	0	-	X	X	-	0	-	-	0	0	-	0	-	-	58,59,60,61,62
<i>Acanthurus dussumieri</i>	X	X	X	X	-	-	-	-	X	-	-	-	-	-	-	-	-	-	58,60,61,63,64,65
<i>Acanthurus grammoptilus</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	65
<i>Acanthurus leucosternon</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58,66,67
<i>Acanthurus mata</i>	X	X	-	X	-	-	X	-	-	0	-	-	0	-	-	0	-	-	58,61,64,68,69
<i>Acanthurus nigricauda</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Acanthurus nigrofuscus</i>	X	0	X	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	58,60,67,70,71
<i>Acanthurus thompsoni</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Acanthurus xanthopterus</i>	X	X	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	61,64,72
<i>Ctenochaetus binotatus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	67
<i>Ctenochaetus striatus</i>	X	0	-	X	-	-	X	-	-	-	-	-	-	-	-	-	-	-	58,67,71,73,74
<i>Ctenochaetus strigosus</i>	X	0	-	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	58

Species	All						Juveniles						Adults						References
	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	
<i>Naso brevirostris</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Naso unicornis</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	66
<i>Zebrasoma desjardini</i>	-	0	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	64
<i>Zebrasoma scopas</i>	X	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	67,74
<i>Zebrasoma veliferum</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58,71
<i>Zebrasoma xanthurum</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
Antennariidae																			
<i>Antennarius pictus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Histrio histrio</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
Apogonidae																			
<i>Apogon ishigakiensis</i>	X	-	-	-	-	-	X	-	-	-	-	-	X	-	-	-	-	-	59
<i>Apogon sangiensis</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Apogonichthyoides taeniatus</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Archamia fucata</i>	0	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Cheilodipterus lachneri</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Cheilodipterus macrodon</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Cheilodipterus novemstriatus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Cheilodipterus quinquelineatus</i>	X	0	X	X	-	-	X	-	X	0	-	-	X	-	-	0	-	-	58,59,63,64,65,68,69,72
<i>Fibramia ambionensis</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Fibramia thermalis</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Fowleria aurita</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Jaydia catalai</i>	-	0	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	64
<i>Ostorhinchus aureus</i>	X	0	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58,64
<i>Ostorhinchus compressus</i>	0	-	-	-	-	X	0	-	-	-	-	X	0	-	-	-	-	X	75
<i>Ostorhinchus cyanosoma</i>	X	0	-	-	-	X	X	-	-	-	-	0	0	-	-	-	-	X	58,71,75
<i>Ostorhinchus rueppellii</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	65
<i>Ostorhinchus wassinki</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	65
<i>Pristiapogon fraenatus</i>	-	0	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	64
<i>Pterapogon kauderni</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	76
<i>Sphaeramia orbicularis</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	63
<i>Zoramia leptacantha</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58

	All						Juveniles						Adults						
Species	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	References
<b>Atherinidae</b>																			
<i>Atherinomorus lacunosus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58,71
<b>Aulostomidae</b>																			
<i>Aulostomus chinensis</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58,67,68,72
<b>Balistidae</b>																			
<i>Balistapus undulatus</i>	X	0	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58,74
<i>Balistoides viridescens</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Melichthys niger</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Pseudobalistes fuscus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Rhinecanthus aculeatus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Sufflamen albicaudatum</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Sufflamen chrysopterum</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Sufflamen fraenatum</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<b>Belonidae</b>																			
<i>Strongylura leiura</i>	0	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Tylosurus crocodilus crocodilus</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<b>Blennidae</b>																			
<i>Atrosalarias fuscus</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	65
<i>Ecsenius frontalis</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Ecsenius gravieri</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Meiacanthus grammistes</i>	X	-	-	0	-	-	0	-	-	0	-	-	X	-	-	0	-	-	69
<i>Meiacanthus mossambicus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58,67
<i>Meiacanthus migrolineatus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Petroscirtes breviceps</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Petroscirtes mitratus</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Salarias guttatus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	77
<b>Caesionidae</b>																			
<i>Caesio caerulea</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Caesio lunaris</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58,71
<i>Caesio suevica</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Caesio teres</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58

	All						Juveniles						Adults						
Species	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	References
<i>Caesio varilineata</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Caesio xanthonota</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Pterocaesio chrysozona</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Pterocaesio marri</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Pterocaesio pisang</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Pterocaesio tile</i>	-	0	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	64
Carangidae																			
<i>Alectis ciliaris</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Carangoides ferdau</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Carangoides fulvoguttatus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Caranx ignobilis</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58,61,78
<i>Caranx melampygyus</i>	X	X	-	X	-	-	-	X	-	0	-	-	-	0	-	0	-	-	61,62,72,79
<i>Caranx papuensis</i>	0	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58,61
<i>Caranx sexfasciatus</i>	0	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Gnathanodon speciosus</i>	0	X	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58,80
<i>Scomberoides commersonnianus</i>	0	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Scomberoides tol</i>	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	61
<i>Trachinotus blochii</i>	0	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
Centriscidae																			
<i>Aeoliscus strigatus</i>	-	0	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	64
Chaetodontidae																			
<i>Chaetodon aureofasciatus</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	81
<i>Chaetodon auriga</i>	X	X	X	X	-	-	X	-	X	X	-	-	-	X	-	X	-	-	58,61,62,64,65,66,70,71,73,82
<i>Chaetodon austriacus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Chaetodon baronessa</i>	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	74
<i>Chaetodon bennetti</i>	-	0	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	64
<i>Chaetodon ephippium</i>	X	0	-	X	-	-	X	-	-	-	-	-	-	-	-	-	-	-	64,73
<i>Chaetodon falcula</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Chaetodon fasciatus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Chaetodon flaviviridis</i>	-	0	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	64

	All						Juveniles						Adults						
Species	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	References
<i>Chaetodon guttatissimus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	67
<i>Chaetodon kleinii</i>	X	0	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58,67,74
<i>Chaetodon lineolatus</i>	-	0	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	64
<i>Chaetodon lunulatus</i>	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	74
<i>Chaetodon melannotus</i>	X	0	-	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	58,71,73,82
<i>Chaetodon octofasciatus</i>	0	-	-	-	-	X	0	-	-	-	-	X	0	-	-	-	-	X	75
<i>Chaetodon paucifasciatus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Chaetodon plebeius</i>	X	0	X	X	-	-	X	-	X	-	-	-	-	-	-	-	-	-	64,65,73
<i>Chaetodon speculum</i>	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	73
<i>Chaetodon trifascialis</i>	0	-	X	-	-	X	0	-	X	-	-	0	0	-	-	-	-	X	65,75
<i>Chaetodon trifasciatus</i>	X	0	X	-	-	-	X	-	X	-	-	-	-	-	-	-	-	-	58,65,67,73
<i>Chaetodon vagabundus</i>	X	-	-	X	-	-	X	-	-	0	-	-	X	-	-	0	-	-	68,69,74
<i>Chaetodon xanthocephalus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Chelmon marginalis</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	81
<i>Hemitaurichthys zoster</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Heniochus acuminatus</i>	X	0	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	58,60,61,62,81
<i>Heniochus diphreutes</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Heniochus intermedius</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
Clupeidae																			
<i>Spratelloides delicatulus</i>	0	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
Dasyatidae																			
<i>Neotrygon kuhlii</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Taeniura lymma</i>	X	X	-	-	-	X	0	-	-	-	-	X	0	-	-	-	-	0	58,75
Diodontidae																			
<i>Cyclichthys spilostylus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Diodon hystrix</i>	X	X	0	X	-	X	-	0	-	0	-	-	-	0	-	X	-	-	58,61,62
<i>Diodon liturosus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
Echeneidae																			
<i>Echeneis naucrates</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
Ephippidae																			
<i>Platax orbicularis</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58,61

	All						Juveniles						Adults						
Species	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	References
<i>Platax teira</i>	0	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
Fistulariidae																			
<i>Fistularia commersonii</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58,63,67,71
<i>Fistularia petimba</i>	X	X	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58,64,83
Gerreidae																			
<i>Gerres filamentosus</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Gerres longirostris</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Gerres oblongus</i>	0	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Gerres oyena</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58,73,82
Gobiidae																			
<i>Amblyeleotris steinitzi</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Amblyeleotris sungami</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Amblygobius albimaculatus</i>	X	X	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58,72
<i>Amblygobius phalaena</i>	X	-	-	X	-	X	X	-	-	X	-	-	X	-	-	0	-	-	68,69,83,84
<i>Amblygobius semicinctus</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Asterropteryx semipunctata</i>	X	-	-	X	-	X	X	-	-	0	-	-	X	-	-	0	-	X	68,69,72,83,84
<i>Bathygobius fuscus</i>	0	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Callogobius hasseltii</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	77
<i>Ctenogobiops feroculus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	77
<i>Ctenogobiops formosa</i>	0	-	-	X	-	-	0	-	-	X	-	-	0	-	-	X	-	-	69
<i>Ctenogobiops maculosus</i>	X	-	-	0	-	-	X	-	-	0	-	-	X	-	-	0	-	-	69
<i>Ctenogobiops pomastictus</i>	X	-	-	X	-	-	X	-	-	X	-	-	X	-	-	X	-	-	68,69
<i>Eviota infulata</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	77
<i>Eviota queenslandica</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	77
<i>Eviota sigillata</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	77
<i>Fusigobius aureus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	77
<i>Fusigobius neophytus</i>	X	X	-	X	-	-	X	-	-	0	-	-	X	-	-	X	-	-	58,68,69,77
<i>Gnatholepis anjerensis</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Gnatholepis cauerensis</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	77
<i>Gobiodon oculolineatus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	77
<i>Gobiodon unicolor</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	77
<i>Macrodontogobius wilburi</i>	X	-	-	0	-	-	X	-	-	0	-	-	X	-	-	0	-	-	69

	All						Juveniles						Adults						
Species	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	References
<i>Phyllogobius platycephalops</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	77
<i>Pleurosicya muscarum</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	77
<i>Valenciennea muralis</i>	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	84
<i>Valenciennea puellaris</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<b>Haemulidae</b>																			
<i>Diagramma pictum</i>	X	0	-	X	-	-	-	0	-	X	-	-	-	0	-	0	-	-	58,62,64
<i>Plectorhinchus albovittatus</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58,85
<i>Plectorhinchus chaetodonoides</i>	X	-	-	0	-	-	X	-	-	0	-	-	0	-	-	0	-	-	69
<i>Plectorhinchus flavomaculatus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58,66,67,82
<i>Plectorhinchus gaterinus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58,82
<i>Plectorhinchus gibbosus</i>	0	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58,85
<i>Plectorhinchus lineatus</i>	0	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	61,63
<i>Plectorhinchus obscurus</i>	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	61
<i>Plectorhinchus plagiodesmus</i>	0	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Plectorhinchus schotaf</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58,66
<i>Plectorhinchus vittatus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58,68
<b>Hemiramphidae</b>																			
<i>Hemiramphus far</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	61,79
<i>Hyporhamphus dussumieri</i>	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	61
<b>Holocentridae</b>																			
<i>Myripristis murdjan</i>	X	-	-	-	-	X	0	-	-	-	-	X	0	-	-	-	-	0	71,75
<i>Neoniphon sammara</i>	X	X	-	0	-	-	-	0	-	0	-	-	-	X	-	0	-	-	58,61,62
<i>Sargocentron diadema</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58,71
<i>Sargocentron rubrum</i>	-	X	-	0	-	-	-	0	-	0	-	-	-	X	-	0	-	-	61,62
<b>Kyphosidae</b>																			
<i>Kyphosus cinerascens</i>	0	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Kyphosus vaigiensis</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58

	All						Juveniles						Adults							
Species	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	References	
Labridae																				
<i>Anampses caeruleopunctatus</i>	X	0	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	-	58, 65, 71
<i>Anampses geographicus</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	-	65, 81
<i>Anampses meleagrides</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Anampses twistii</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Bodianus anthioides</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Bodianus axillaris</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Cheilinus chlorourus</i>	X	0	X	X	-	-	X	-	X	0	-	-	X	-	-	0	-	-	-	58, 63, 64, 65, 67, 68, 69, 72, 79
<i>Cheilinus fasciatus</i>	X	-	-	0	-	X	0	-	-	-	-	X	0	-	-	-	-	X	-	72, 75, 82
<i>Cheilinus lunulatus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Cheilinus oxycephalus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Cheilinus trilobatus</i>	X	0	X	-	-	X	0	-	X	-	-	X	0	-	-	-	-	0	-	58, 65, 66, 67, 74, 75, 82
<i>Cheilinus undulatus</i>	X	0	X	-	-	-	X	-	X	-	-	-	-	-	-	-	-	-	-	51, 58, 66, 79, 82, 85, 86
<i>Cheilio inermis</i>	X	0	X	0	-	-	X	-	X	0	-	-	X	-	-	0	-	-	-	58, 59, 63, 65, 66, 68, 69, 71, 72, 79, 82, 83
<i>Choerodon anchorago</i>	X	X	-	0	-	X	X	-	-	0	-	X	X	-	-	0	-	X	-	63, 68, 69, 75, 79, 85, 87
<i>Choerodon cauteroma</i>	X	-	X	-	-	X	X	-	X	-	-	X	X	-	-	-	-	X	-	81, 88
<i>Choerodon cyanodus</i>	X	-	X	-	-	X	X	-	X	-	-	X	X	-	-	-	-	X	-	81, 88
<i>Choerodon monostigma</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	-	81
<i>Choerodon schoenleinii</i>	X	-	X	-	-	X	X	-	X	-	-	X	X	-	-	-	-	X	-	81, 88
<i>Cirrhilabrus rubriventralis</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Coris aygula</i>	X	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	-	65, 71
<i>Coris batuensis</i>	X	-	-	-	-	X	0	-	-	-	-	X	0	-	-	-	-	0	-	68, 75
<i>Coris caudimacula</i>	X	0	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	-	58, 65, 67, 71
<i>Coris formosa</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	67
<i>Coris variegata</i>	X	-	-	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71, 83
<i>Diproctacanthus xanthurus</i>	0	-	-	-	-	X	0	-	-	-	-	X	0	-	-	-	-	X	-	75
<i>Epibulus insidiator</i>	X	-	-	-	-	X	0	-	-	-	-	X	X	-	-	-	-	X	-	75
<i>Gomphosus caeruleus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	67, 71

Species	All						Juveniles						Adults						References
	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	
<i>Gomphosus varius</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	65
<i>Halichoeres argus</i>	X	X	-	-	-	-	X	-	-	-	-	-	X	-	-	-	-	-	59, 63, 68
<i>Halichoeres chloropterus</i>	X	-	-	-	-	X	X	-	-	-	-	X	X	-	-	-	-	X	75
<i>Halichoeres hortulanus</i>	X	0	-	X	-	X	0	-	-	-	-	X	0	-	-	-	-	0	58, 63, 67, 74, 75
<i>Halichoeres leucurus</i>	0	-	-	-	-	X	0	-	-	-	-	X	0	-	-	-	-	X	75
<i>Halichoeres margaritaceus</i>	X	0	-	0	-	X	X	-	-	-	-	X	X	-	-	-	-	X	63, 72, 75
<i>Halichoeres marginatus</i>	X	0	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	58, 65
<i>Halichoeres melanochir</i>	0	-	-	-	-	X	0	-	-	-	-	X	0	-	-	-	-	X	75
<i>Halichoeres melanurus</i>	X	-	-	X	-	X	X	-	-	0	-	0	X	-	-	0	-	X	69, 74, 75
<i>Halichoeres miniatus</i>	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	87
<i>Halichoeres nebulosus</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	65, 81
<i>Halichoeres richmondi</i>	0	-	-	-	-	X	0	-	-	-	-	0	0	-	-	-	-	X	75
<i>Halichoeres scapularis</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 63
<i>Halichoeres trimaculatus</i>	X	-	X	0	-	-	X	-	X	0	-	-	X	-	-	0	-	-	59, 65, 68, 69, 72
<i>Halichoeres vrolikii</i>	0	-	-	-	-	X	0	-	-	-	-	X	0	-	-	-	-	X	75
<i>Halichoeres zeylonicus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Hemigymnus fasciatus</i>	X	0	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	58, 65, 71
<i>Hemigymnus melapterus</i>	X	-	X	-	-	X	X	-	X	-	-	X	X	-	-	-	-	X	65, 71, 81
<i>Hologymnosus annulatus</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	65, 71
<i>Labrichthys unilineatus</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	65
<i>Labroides dimidiatus</i>	X	0	X	-	-	X	0	-	X	-	-	X	0	-	-	-	-	X	58, 65, 67, 71, 75
<i>Larabicus quadrilineatus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Macropharyngodon bipartitus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Macropharyngodon negrosensis</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	81
<i>Macropharyngodon ornatus</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	65
<i>Novaculichthys taeniourus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Oxycheilinus arenatus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Oxycheilinus bimaculatus</i>	X	0	-	X	-	-	X	-	-	0	-	-	X	-	-	0	-	-	58, 59, 64, 69
<i>Oxycheilinus celebicus</i>	0	-	-	-	-	X	0	-	-	-	-	0	0	-	-	-	-	X	75
<i>Oxycheilinus digramma</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 71
<i>Oxycheilinus mentalis</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Paracheilinus octotaenia</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71

	All						Juveniles						Adults						
Species	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	References
<i>Pseudocheilinus evanidus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Pseudocheilinus hexataenia</i>	X	-	-	X	-	X	0	-	-	-	-	0	0	-	-	-	-	X	67, 71, 74, 75
<i>Pteragogus cryptus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Pteragogus flagellifer</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Stethojulis albovittata</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 71
<i>Stethojulis bandanensis</i>	X	0	X	0	-	-	-	-	X	-	-	-	-	-	-	-	-	-	58, 65, 67, 72, 79
<i>Stethojulis interrupta</i>	X	0	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	58, 65, 71, 81
<i>Stethojulis strigiventer</i>	X	X	X	0	-	-	X	-	X	0	-	-	X	-	-	0	-	-	58, 59, 65, 68, 69, 72, 81, 83
<i>Stethojulis trilineata</i>	X	X	-	-	-	X	X	-	-	-	-	X	X	-	-	-	-	X	63, 75
<i>Thalassoma amblycephalum</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	67
<i>Thalassoma hardwicke</i>	X	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	65, 67
<i>Thalassoma hebraicum</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	67
<i>Thalassoma lunare</i>	X	0	X	X	-	X	0	-	X	-	-	0	0	-	-	-	-	X	64, 65, 67, 71, 74, 75
<i>Thalassoma lutescens</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	65
<i>Thalassoma rueppellii</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
Lethrinidae																			
<i>Gnathodentex aureolineatus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Gymnocranius grandoculis</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58,66
<i>Lethrinus atkinsoni</i>	X	-	X	0	-	-	X	-	X	0	-	-	0	-	-	0	-	-	59, 65, 68, 69, 83, 89, 90
<i>Lethrinus erythropterus</i>	0	X	-	-	-	X	0	-	-	-	-	X	0	-	-	-	-	X	75,85
<i>Lethrinus genivittatus</i>	X	X	-	X	-	-	X	-	-	-	-	-	0	-	-	-	-	-	58,59,64,80
<i>Lethrinus harak</i>	X	X	X	X	-	X	X	X	-	X	-	0	X	X	-	X	-	X	58, 59, 61, 62, 63, 66, 67, 68, 69, 79, 82, 85, 87, 89, 91, 92, 93, 94, 95, 96
<i>Lethrinus lentjan</i>	X	X	-	X	-	X	X	X	-	-	-	0	X	0	-	-	-	X	58, 61, 64, 66, 82, 91, 92, 93
<i>Lethrinus mahsena</i>	X	X	-	X	-	-	X	-	-	X	-	-	-	-	-	-	-	-	58, 66, 67, 70, 82
<i>Lethrinus microdon</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 66
<i>Lethrinus miniatus</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58

	All						Juveniles						Adults						
Species	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	References
<i>Lethrinus nebulosus</i>	X	X	X	X	-	-	X	-	X	X	-	-	X	-	-	-	-	-	58, 59, 61, 64, 66, 70, 83, 90, 97
<i>Lethrinus obsoletus</i>	X	X	-	X	-	-	X	-	-	X	-	-	X	-	-	X	-	-	58, 59, 61, 66, 67, 68, 83, 89, 94
<i>Lethrinus olivaceus</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Lethrinus ornatus</i>	X	-	-	-	-	-	X	-	-	-	-	-	0	-	-	-	-	-	59, 68
<i>Lethrinus punctulatus</i>	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	80
<i>Lethrinus rubrioperculatus</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 61
<i>Lethrinus semicinctus</i>	-	0	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	64
<i>Lethrinus variegatus</i>	X	X	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	58, 82, 90
<i>Lethrinus xanthochilus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Monotaxis grandoculis</i>	X	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71, 74
Lutjanidae																			
<i>Lutjanus argentimaculatus</i>	X	X	-	X	-	-	-	X	-	0	-	-	-	X	-	0	-	-	58, 61, 62, 64, 66, 82, 85, 87, 89
<i>Lutjanus bohar</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 66, 67
<i>Lutjanus carponotatus</i>	X	-	X	X	-	X	X	-	X	-	-	0	0	-	-	-	-	X	75, 80, 81
<i>Lutjanus decussatus</i>	X	X	-	-	-	X	0	-	-	-	-	X	X	-	-	-	-	X	63, 75
<i>Lutjanus ehrenbergii</i>	X	X	-	-	-	-	0	X	-	-	-	-	-	-	-	-	-	-	58, 82, 98, 99, 100
<i>Lutjanus fulviflamma</i>	X	X	-	X	-	0	X	X	-	-	-	0	X	0	-	-	-	0	58, 61, 63, 64, 66, 67, 75, 82, 85, 87, 91, 92, 93, 98, 101
<i>Lutjanus fulvus</i>	0	X	-	0	-	-	-	X	-	0	-	-	-	0	-	0	-	-	58, 61, 62, 85, 102
<i>Lutjanus gibbus</i>	X	0	-	X	-	-	X	-	-	X	-	-	X	-	-	X	-	-	58, 59, 66, 68, 69, 82, 83, 94
<i>Lutjanus kasmira</i>	X	0	-	X	X	X	-	-	-	-	-	-	-	-	-	-	-	-	58, 64, 103
<i>Lutjanus lutjanus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 66
<i>Lutjanus monostigma</i>	0	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 63
<i>Lutjanus quinquelineatus</i>	-	0	-	X	-	-	-	0	-	0	-	-	-	0	-	X	-	-	62, 64
<i>Lutjanus russellii</i>	-	X	-	0	-	-	-	X	-	0	-	-	-	0	-	0	-	-	61, 62
<i>Lutjanus vitta</i>	-	0	-	X	-	-	-	0	-	X	-	-	-	0	-	X	-	-	62
Monacanthidae																			
<i>Aluterus scriptus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58

	All						Juveniles						Adults						
Species	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	References
<i>Amanses scopas</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 67, 71
<i>Cantherhines pardalis</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	67, 71
<i>Oxymonacanthus longirostris</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	65
<i>Pervagor randalli</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
Mullidae																			
<i>Mulloidichthys flavolineatus</i>	X	X	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 64, 68, 71, 72, 79, 82
<i>Mulloidichthys vanicolensis</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 66
<i>Parupeneus barberinoides</i>	X	-	X	0	-	-	X	-	X	-	-	-	X	-	-	-	-	-	59, 65, 68, 81, 83
<i>Parupeneus barberinus</i>	X	X	-	X	-	X	X	-	-	0	-	X	X	-	-	0	-	X	58, 59, 63, 66, 68, 69, 72, 74, 75, 79, 82, 83
<i>Parupeneus ciliatus</i>	X	0	-	0	-	-	X	-	-	0	-	-	X	-	-	0	-	-	58, 59, 68, 69
<i>Parupeneus cyclostomus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 71
<i>Parupeneus forsskali</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Parupeneus heptacanthus</i>	-	0	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	64
<i>Parupeneus indicus</i>	X	X	-	X	-	-	X	-	-	X	-	-	X	-	-	X	-	-	58, 59, 61, 62, 64, 66, 68, 69, 82, 94
<i>Parupeneus macronemus</i>	X	X	-	X	-	-	X	-	-	X	-	-	-	-	-	-	-	-	58, 66, 67, 70, 71, 82
<i>Parupeneus multifasciatus</i>	X	-	-	X	-	-	X	-	-	0	-	-	X	-	-	0	-	-	59, 68, 69, 72, 74, 83
<i>Parupeneus pleurostigma</i>	X	0	-	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 66, 72, 82
<i>Parupeneus porphyreus</i>	X	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	72, 104
<i>Parupeneus rubescens</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 61, 66, 82
<i>Parupeneus spilurus</i>	-	0	X	X	-	-	-	0	X	X	-	-	-	0	-	X	-	-	62, 64, 65
<i>Upeneus taeniopterus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	79
<i>Upeneus tragula</i>	X	X	X	X	-	X	X	-	-	X	-	X	X	X	-	X	-	X	58, 59, 60, 61, 62, 64, 75, 87
Muraenidae																			
<i>Echidna nebulosa</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Gymnothorax griseus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Gymnothorax nudivomer</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71

	All						Juveniles						Adults						
Species	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	References
<i>Gymnothorax pictus</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
Nemipteridae																			
<i>Pentapodus trivittatus</i>	X	-	-	-	-	X	X	-	-	-	-	0	X	-	-	-	-	X	75
<i>Scolopsis affinis</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	63
<i>Scolopsis bilineata</i>	X	X	X	X	-	-	-	-	X	-	-	-	-	-	-	-	-	-	60, 63, 65, 74, 79
<i>Scolopsis bimaculata</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Scolopsis ciliata</i>	X	X	-	X	-	0	0	-	-	-	-	0	X	-	-	-	-	0	63, 74, 75
<i>Scolopsis ghanam</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 66, 71, 82
<i>Scolopsis lineata</i>	X	X	-	0	-	X	X	-	-	0	-	0	X	-	-	0	-	X	63, 69, 75, 79
<i>Scolopsis margaritifera</i>	0	-	-	-	-	X	0	-	-	-	-	0	0	-	-	-	-	X	75
<i>Scolopsis monogramma</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	79
<i>Scolopsis temporalis</i>	-	0	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	64
<i>Scolopsis trilineata</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	79
Ophichthidae																			
<i>Myrichthys colubrinus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
Ostraciidae																			
<i>Lactoria cornuta</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Lactoria fornasini</i>	0	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Ostracion cubicus</i>	X	0	-	X	-	-	-	0	-	0	-	-	-	0	-	X	-	-	58, 62, 64, 71
<i>Ostracion cyanurus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
Pinguipedidae																			
<i>Parapercis cylindrica</i>	X	X	X	X	-	-	X	0	-	X	-	-	X	0	-	X	-	-	59, 60, 62, 63, 64, 68, 69, 83
<i>Parapercis hexophtalma</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
Platycephalidae																			
<i>Papilloculiceps longiceps</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Thysanophrys chiltonae</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
Plotosidae																			
<i>Plotosus lineatus</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
Pomacanthidae																			
<i>Apolemichthys xanthotis</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71

Species	All						Juveniles						Adults						References
	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	
<i>Centropyge multispinis</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 71
<i>Centropyge tibicen</i>	-	0	-	X	-	-	-	0	-	X	-	-	-	0	-	X	-	-	62
<i>Centropyge vrolikii</i>	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	74
<i>Genicanthus caudovittatus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Pomacanthus chrysurus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Pomacanthus imperator</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Pomacanthus sexstriatus</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	81
<i>Pygoplites diacanthus</i>	X	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71, 71
Pomacentridae																			
<i>Abudefduf bengalensis</i>	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	87
<i>Abudefduf lorenzi</i>	0	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	63
<i>Abudefduf septemfasciatus</i>	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	61
<i>Abudefduf sexfasciatus</i>	X	X	X	-	-	X	0	-	-	-	-	X	0	-	-	-	-	0	58, 60, 63, 67, 75, 87
<i>Abudefduf sparoides</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Abudefduf vaigiensis</i>	X	X	-	-	-	X	X	-	-	-	-	X	X	-	-	-	-	0	58, 63, 71, 73, 75
<i>Abudefduf whiteleyi</i>	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	60
<i>Amblyglyphidodon batunai</i>	X	-	-	-	-	X	X	-	-	-	-	X	0	-	-	-	-	X	75
<i>Amblyglyphidodon curacao</i>	0	-	-	X	-	X	0	-	-	-	-	X	0	-	-	-	-	X	74, 75
<i>Amblyglyphidodon leucogaster</i>	X	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	67, 71, 74
<i>Amphiprion akallopisos</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	67
<i>Amphiprion allardi</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Amphiprion bicinctus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Amphiprion ocellaris</i>	0	-	-	-	-	X	0	-	-	-	-	X	0	-	-	-	-	X	75
<i>Amphiprion rubrocinctus</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	81
<i>Cheiloprion labiatus</i>	X	-	-	-	-	X	X	-	-	-	-	X	X	-	-	-	-	X	75
<i>Chromis atripectoralis</i>	X	-	-	-	-	X	0	-	-	-	-	X	0	-	-	-	-	X	67, 75
<i>Chromis dimidiata</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Chromis lepidolepis</i>	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	74
<i>Chromis margaritifer</i>	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	74
<i>Chromis opercularis</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Chromis retrofasciata</i>	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	74
<i>Chromis ternatensis</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71

	All						Juveniles						Adults						
Species	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	References
<i>Chromis viridis</i>	X	-	X	-	-	-	X	-	X	-	-	-	-	-	-	-	-	-	60, 65, 67, 71, 73
<i>Chrysiptera annulata</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Chrysiptera cyanea</i>	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	73
<i>Chrysiptera rex</i>	0	-	-	-	-	X	0	-	-	-	-	0	0	-	-	-	-	X	75
<i>Chrysiptera rollandi</i>	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	74
<i>Chrysiptera talboti</i>	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	74
<i>Chrysiptera unimaculata</i>	X	-	-	0	-	-	X	-	-	0	-	-	X	-	-	0	-	-	68, 69
<i>Dascyllus aruanus</i>	X	0	X	X	-	X	X	0	X	X	-	X	X	0	-	X	-	X	58, 60, 62, 64, 65, 70, 71, 73, 75
<i>Dascyllus carneus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Dascyllus marginatus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Dascyllus reticulatus</i>	-	-	X	X	-	-	-	-	X	-	-	-	-	-	-	-	-	-	65, 74
<i>Dascyllus trimaculatus</i>	X	0	-	X	-	-	X	-	-	-	-	-	-	-	-	-	-	-	58, 71, 73, 74
<i>Dischistodus chrysopoecilus</i>	X	-	-	-	-	X	X	-	-	-	-	X	X	-	-	-	-	X	75
<i>Dischistodus melanotus</i>	X	-	-	-	-	X	0	-	-	-	-	0	X	-	-	-	-	X	75
<i>Dischistodus perspicillatus</i>	X	-	X	-	-	0	X	-	X	-	-	0	X	-	-	-	-	0	65, 75
<i>Dischistodus prosopotaenia</i>	X	-	X	0	-	X	X	-	X	0	-	X	X	-	-	0	-	X	65, 68, 69, 75
<i>Neoglyphidodon melas</i>	X	0	X	-	-	X	0	-	X	-	-	X	0	-	-	-	-	X	58, 65, 71, 75
<i>Neoglyphidodon nigroris</i>	0	-	-	-	-	X	0	-	-	-	-	0	0	-	-	-	-	X	75
<i>Neopomacentrus azysron</i>	0	-	-	-	-	X	0	-	-	-	-	0	0	-	-	-	-	X	75
<i>Neopomacentrus cyanomos</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	81
<i>Neopomacentrus filamentosus</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	81
<i>Neopomacentrus fuliginosus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Neopomacentrus miryae</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Plectroglyphidodon lacrymatus</i>	X	0	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	68, 65, 67
<i>Plectroglyphidodon leucozonus</i>	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	73
<i>Pomacentrus adelus</i>	X	-	-	-	-	X	0	-	-	-	-	X	X	-	-	-	-	X	75
<i>Pomacentrus amboinensis</i>	X	0	-	X	-	X	X	-	-	-	-	X	0	-	-	-	-	X	64, 73, 74, 75
<i>Pomacentrus baenschi</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	67
<i>Pomacentrus bankanensis</i>	X	-	-	-	-	X	X	-	-	-	-	0	0	-	-	-	-	X	73, 75
<i>Pomacentrus burroughi</i>	X	-	-	-	-	X	0	-	-	-	-	X	X	-	-	-	-	X	75

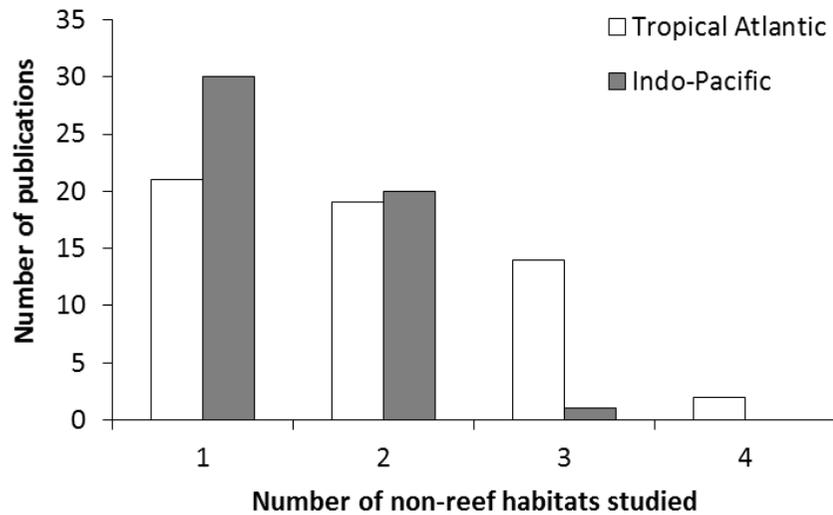
	All						Juveniles						Adults						
Species	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	References
<i>Pomacentrus chrysurus</i>	X	-	-	0	-	X	X	-	-	0	-	0	X	-	-	0	-	X	68, 69, 73, 75
<i>Pomacentrus coelestis</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	65
<i>Pomacentrus limosus</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	81
<i>Pomacentrus milleri</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	81
<i>Pomacentrus moluccensis</i>	X	-	X	X	-	X	X	-	X	-	-	X	0	-	-	-	-	0	60, 65, 73, 74, 75
<i>Pomacentrus nigromanus</i>	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	74
<i>Pomacentrus philippinus</i>	0	0	X	X	-	X	0	-	-	-	-	X	0	-	-	-	-	X	60, 64, 75
<i>Pomacentrus trichourus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 67, 71
<i>Pomacentrus trilineatus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 67
<i>Pomacentrus vaiuli</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	65
<i>Stegastes nigricans</i>	X	0	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	58, 65
<i>Stegastes obreptus</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	65
Priacanthidae																			
<i>Heteropriacanthus cruentatus</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Priacanthus hamrur</i>	X	0	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 64
Pseudochromidae																			
<i>Pseudochromis flavivertex</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Pseudochromis fridmani</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Pseudochromis fuscus</i>	X	-	X	-	-	0	0	-	X	-	-	0	X	-	-	-	-	0	75, 77, 81
<i>Pseudochromis olivaceus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Pseudochromis springeri</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
Scarinae																			
<i>Bolbometopon muricatum</i>	X	X	-	-	-	-	X	X	-	-	-	-	-	-	-	-	-	-	85
<i>Calotomus carolinus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 66, 67, 82
<i>Calotomus spinidens</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 73, 82
<i>Calotomus viridescens</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Chlorurus atrilunula</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	66
<i>Chlorurus gibbus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Chlorurus microrhinos</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	65
<i>Chlorurus sordidus</i>	X	0	X	0	-	X	X	-	X	-	-	X	0	-	-	-	-	X	58, 65, 66, 67, 71, 72, 75, 81, 82

		All						Juveniles						Adults						
Species		SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	References
	<i>Chlorurus strongylocephalus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	82
	<i>Hipposcarus harid</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 66, 82
	<i>Leptoscarus vaigiensis</i>	X	0	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	58, 65, 66, 79, 82
	<i>Scarus dubius</i>	X	-	-	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	72
	<i>Scarus ferrugineus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
	<i>Scarus frenatus</i>	X	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	65, 67, 82
	<i>Scarus ghobban</i>	X	X	X	X	-	X	X	X	X	0	-	X	0	X	-	0	-	0	58, 60, 61, 62, 64, 66, 67, 68, 69, 71, 75, 81, 82
	<i>Scarus niger</i>	X	-	-	-	-	X	0	-	-	-	-	0	0	-	-	-	-	X	71, 75, 82
	<i>Scarus prasiognathos</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	65
	<i>Scarus psittacus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 66, 71, 82
	<i>Scarus quoyi</i>	0	-	-	-	-	X	0	-	-	-	-	X	0	-	-	-	-	X	75
	<i>Scarus rivulatus</i>	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	87
	<i>Scarus russelii</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	66, 82
	<i>Scarus scaber</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	82
	<i>Scarus tricolor</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	67
	<i>Scarus viridifucatus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
Scombridae																				
	<i>Rastrelliger kanagurta</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
Scorpaenidae																				
	<i>Caracanthus unipinna</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	77
	<i>Dendrochirus brachypterus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
	<i>Parascorpaena mossambica</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
	<i>Pterois antennata</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
	<i>Pterois miles</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 71
	<i>Pterois radiata</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
	<i>Pterois volitans</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	63
	<i>Scorpaenopsis venosa</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
Serranidae																				
	<i>Anyperodon leucogrammicus</i>	0	X	-	-	-	X	0	-	-	-	-	X	0	-	-	-	-	0	58, 75

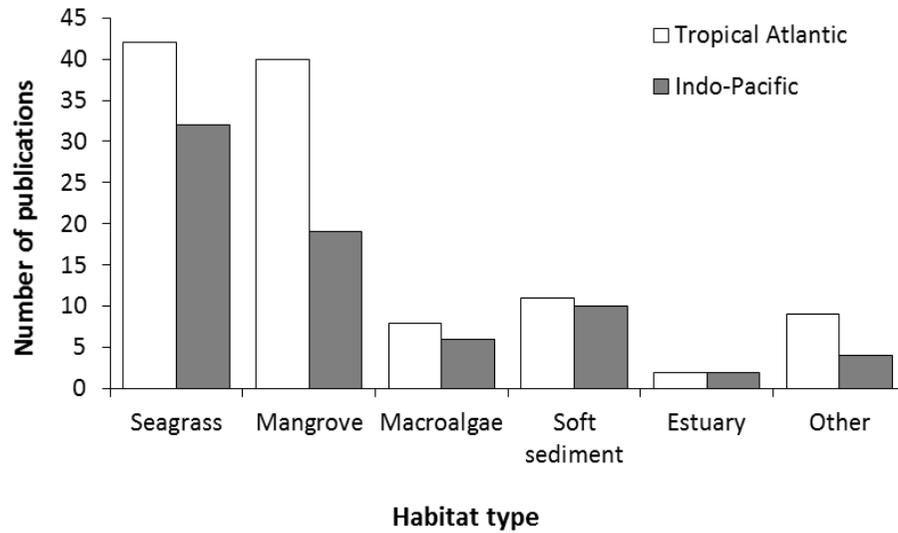
	All						Juveniles						Adults						
Species	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	References
<i>Cephalopholis argus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Cephalopholis boenak</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	81
<i>Cephalopholis cyanostigma</i>	0	-	-	-	-	X	0	-	-	-	-	X	0	-	-	-	-	0	75
<i>Cephalopholis miniata</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Epinephelus areolatus</i>	-	0	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	64
<i>Epinephelus coeruleopunctatus</i>	X	X	-	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 61, 64
<i>Epinephelus coioides</i>	0	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Epinephelus cyanopodus</i>	-	0	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 61, 64
<i>Epinephelus fasciatus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Epinephelus fuscoguttatus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Epinephelus lanceolatus</i>	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	61
<i>Epinephelus maculatus</i>	-	0	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	64
<i>Epinephelus malabaricus</i>	0	X	-	0	-	-	-	X	-	0	-	-	-	X	-	0	-	-	58, 61, 62
<i>Epinephelus merra</i>	X	0	-	-	-	X	0	-	-	-	-	X	0	-	-	-	-	X	63, 64, 75
<i>Epinephelus polyphkadion</i>	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	61
<i>Epinephelus rivulatus</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	81
<i>Epinephelus tauvina</i>	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	61
<i>Epinephelus tukula</i>	0	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Grammistes sexlineatus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 67, 71
<i>Plectropomus leopardus</i>	-	0	-	X	-	-	-	0	-	0	-	-	-	0	-	X	-	-	62, 64
<i>Plectropomus maculatus</i>	-	-	X	-	-	-	-	-	X	-	-	-	-	-	-	-	-	-	81
<i>Pseudanthias squamipinnis</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Variola louti</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
Siganidae																			
<i>Siganus argenteus</i>	-	0	-	X	-	-	-	0	-	X	-	-	-	0	-	0	-	-	62
<i>Siganus canaliculatus</i>	X	X	-	X	-	-	-	X	-	X	-	-	-	X	-	0	-	-	58, 61, 62, 64
<i>Siganus doliatus</i>	X	0	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	60, 85
<i>Siganus fuscescens</i>	X	X	-	X	-	-	X	-	-	-	-	-	X	-	-	-	-	-	59, 63, 80, 105
<i>Siganus guttatus</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	63, 89
<i>Siganus lineatus</i>	0	X	-	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	61, 85, 87, 106
<i>Siganus luridus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 71
<i>Siganus puellus</i>	X	0	-	X	-	-	X	-	-	-	-	-	X	-	-	-	-	-	59, 64

	All						Juveniles						Adults						
Species	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	References
<i>Siganus rivulatus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Siganus spinus</i>	X	X	-	0	-	-	X	-	-	0	-	-	X	-	-	0	-	-	59, 63, 68, 69, 72
<i>Siganus stellatus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 66, 82
<i>Siganus sutor</i>	X	X	-	-	-	-	X	0	-	-	-	0	X	0	-	-	-	0	58, 66, 67, 82, 91
<i>Siganus virgatus</i>	X	X	X	-	-	X	X	-	X	-	-	X	0	-	-	-	-	X	63, 75, 81
Sillaginidae																			
<i>Sillago sihama</i>	0	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
Sphyraenidae																			
<i>Sphyraena barracuda</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Sphyraena flavicauda</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 67
<i>Sphyraena forsteri</i>	0	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Sphyraena jello</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Sphyraena obtusata</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Sphyraena putnamae</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
Synanceiidae																			
<i>Synanceia verrucosa</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
Syngnathidae																			
<i>Corythoichthys flavofasciatus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 67, 71
<i>Corythoichthys haematopterus</i>	X	0	-	0	-	-	0	-	-	0	-	-	X	-	-	0	-	-	63, 68, 69
<i>Corythoichthys intestinalis</i>	X	-	-	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	72
<i>Corythoichthys schultzi</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
<i>Hippocampus comes</i>	X	-	X	-	-	-	0	-	X	-	-	-	X	-	X	-	-	-	107
<i>Hippocampus histrix</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
<i>Syngnathoides biaculeatus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
Synodontidae																			
<i>Saurida gracilis</i>	X	0	-	X	-	-	X	-	-	0	-	-	X	-	-	X	-	0	58, 68, 69, 71, 72
<i>Saurida nebulosa</i>	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	61
<i>Saurida undosquamis</i>	-	0	-	X	-	-	-	0	-	X	-	-	-	0	-	X	-	-	62
<i>Synodus dermatogenys</i>	0	-	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	83
<i>Synodus variegatus</i>	X	X	-	X	-	-	0	-	-	X	-	-	-	-	-	-	-	-	58, 70, 71, 72, 83

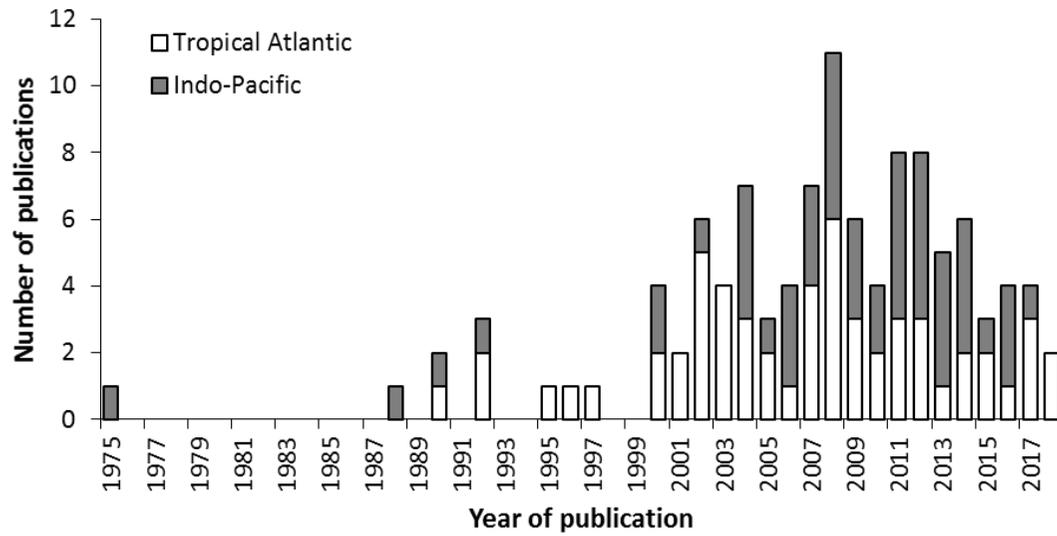
		All						Juveniles						Adults						
Species		SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	SG	MG	MA	SS	E	OT	References
	<i>Trachyrhampus bicoarctatus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
Tetraodontidae																				
	<i>Arothron diadematus</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
	<i>Arothron hispidus</i>	X	X	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 64, 71
	<i>Arothron immaculatus</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
	<i>Arothron mappa</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
	<i>Arothron meleagris</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
	<i>Arothron nigropunctatus</i>	X	0	-	-	-	0	0	-	-	-	-	0	X	-	-	-	-	0	58, 75
	<i>Arothron stellatus</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 71
	<i>Canthigaster bennetti</i>	X	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 67
	<i>Canthigaster coronata</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
	<i>Canthigaster margaritata</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	71
	<i>Canthigaster solandri</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58
	<i>Canthigaster valentini</i>	X	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 67
Tripterygiidae																				
	<i>Enneapterygius tutuilae</i>	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	77
Zanclidae																				
	<i>Zanclus cornutus</i>	X	0	-	X	-	-	-	-	-	-	-	-	-	-	-	-	-	-	58, 66, 74



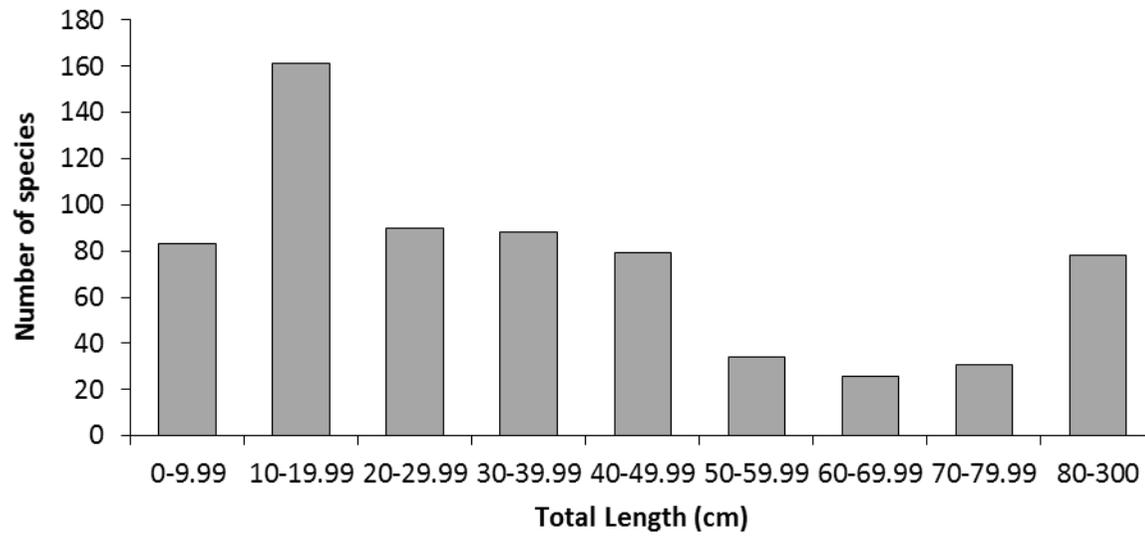
**Figure A2.1.** Number of non-reef habitats surveyed within each of the 107 publications by biogeographic region. In both regions, most studies have examined a single non-reef habitat type. In the Indo-Pacific, far fewer studies have examined three or more non-reef habitats compared to the tropical Atlantic.



**Figure A2.2.** Habitat types examined within 107 publications that recorded coral reef fishes on coral reefs and in at least one non-reef habitat, by biogeographic region. In the tropical Atlantic, the number of publications collecting data from seagrass and mangrove habitats is almost equal. In contrast, in the Indo-Pacific, the number of studies examining seagrass beds is almost double that examining mangrove habitats. The remaining habitat types have all been studied to a far lesser extent to date.



**Figure A2.3.** Number of studies by year (1975-2018) and biogeographic region that surveyed or captured coral reef fishes on coral reefs and in at least one non-reef habitat.



**Figure A2.4.** Maximum total length (cm) of species recorded on coral reefs and in at least one non-reef habitat. Data extracted from *FishBase* (Froese & Pauly 2019).

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## Appendix B: Supplementary materials for Chapter 3

### *Broadening our horizons: seascape use by coral reef-associated fishes in Kavieng, Papua New Guinea, is common and diverse*

**Table A3.1:** Frequency of occurrence (%) of coral reef-associated fishes occurring in non-reef habitats. RS = Reef slopes, RF = Reef flat, MA = Macroalgae, MG = Mangrove, SG = Seagrass.

	RS	RF	MA	MG	SG
<b>Acanthuridae</b>					
<i>Acanthurus spp.</i>	19.8	12.0	2.4	1.7	0
<i>Acanthurus auranticavus</i>	2.3	0	0	5.2	0
<i>Acanthurus triostegus</i>	1.7	41.3	12.2	0	3
<i>Ctenochaetus spp.</i>	0	0	0	1.7	0
<i>Ctenochaetus striatus</i>	84.9	70.7	7.3	0	0
<b>Apogonidae</b>					
<i>Apogonidae spp.</i>	0	0	0	17.2	0
<i>Cheilodipterus spp.</i>	0	2.7	0	1.7	0
<i>Sphaeramia orbicularis</i>	0	0	0	37.9	0
<b>Balistidae</b>					
<i>Rhinecanthus aculeatus</i>	0	0	22	1.7	0
<i>Rhinecanthus verrucosus</i>	2.3	30.7	29.3	0	4
<b>Carangidae</b>					
<i>Carangidae spp.</i>	8.1	0	0	8.6	0
<i>Caranx ignobilis</i>	0	2.7	7.3	1.7	0
<i>Caranx papuensis</i>	1.2	0	0	0	1
<i>Caranx sexfasciatus</i>	0	0	0	12.1	1
<i>Caranx spp.</i>	4.7	5.3	4.9	1.7	3
<i>Gnathanodon speciosus</i>	1.2	0	0	0	1
<b>Carcharhinidae</b>					
<i>Carcharhinus melanopterus</i>	18.6	6.7	12.2	0	4
<b>Chaetodontidae</b>					
<i>Chaetodon lunula</i>	11.6	2.7	0	6.9	0
<i>Chaetodon rafflesii</i>	18.6	21.3	4.9	0	0
<i>Chaetodon vagabundus</i>	37.2	36	24.4	22.4	3
<b>Gerreidae</b>					
<i>Gerres longirostris</i>	0	0	0	0	3
<i>Gerres oyena</i>	1.2	0	0	51.7	12
<b>Haemulidae</b>					
<i>Plectorhinchus albovittatus</i>	0	0	0	1.7	0
<i>Plectorhinchus gibbosus</i>	0	0	0	5.2	0
<i>Plectorhinchus spp.</i>	0	0	0	1.7	0
<b>Kyphosidae</b>					
<i>Kyphosus cinerascens</i>	14	0	0	1.7	0
<b>Labridae</b>					
<i>Cheilinus chlorourus</i>	2.3	0	9.8	0	0
<i>Cheilinus spp.</i>	0	2.7	0	1.7	0

<i>Cheilinus trilobatus</i>	27.9	28	2.4	3.4	3
<i>Cheilio inermis</i>	15.1	21.3	24.4	0	19
<i>Choerodon anchorago</i>	32.6	66.7	63.4	15.5	30
<i>Coris spp.</i>	0	0	2.4	0	0
<i>Halichoeres argus</i>	0	18.7	36.6	3.4	33
<i>Halichoeres chloropterus</i>	5.8	22.7	2.4	0	1
<i>Halichoeres margaritaceus</i>	0	4	7.3	0	0
<i>Halichoeres marginatus</i>	27.9	22.7	2.4	0	0
<i>Halichoeres melanurus</i>	54.7	25.3	2.4	0	1
<i>Halichoeres miniatus</i>	0	5.3	19.5	0	0
<i>Halichoeres nebulosus</i>	3.5	14.7	46.3	0	0
<i>Halichoeres nigrescens</i>	2.3	18.7	41.5	5.2	1
<i>Halichoeres papilionaceus</i>	0	4	26.8	1.7	2
<i>Halichoeres scapularis</i>	15.1	58.7	73.2	6.9	16
<i>Halichoeres spp.</i>	11.6	45.3	80.5	1.7	36
<i>Halichoeres trimaculatus</i>	12.8	38.7	53.7	5.2	4
<i>Stethojulis bandanensis</i>	16.3	13.3	14.6	0	0
<i>Stethojulis interrupta</i>	4.7	9.3	24.4	1.7	5
<i>Stethojulis spp.</i>	1.2	2.7	4.9	0	2
<i>Stethojulis strigiventer</i>	8.1	30.7	43.9	5.2	28
<i>Stethojulis trilineata</i>	64	66.7	24.4	5.2	0
<i>Thalassoma amblycephalum</i>	4.7	1.3	0	0	1
<i>Thalassoma hardwicke</i>	65.1	66.7	12.2	0	1
<b>Latidae</b>					
<i>Psammoperca waigiensis</i>	0	0	0	6.9	0
<b>Lethrinidae</b>					
<i>Lethrinid early stage juvenile</i>	1.2	0	7.3	0	21
<i>Lethrinus erythropterus</i>	14	1.3	2.4	0	0
<i>Lethrinus harak</i>	12.8	50.7	73.2	36.2	53
<i>Lethrinus obsoletus</i>	0	4	9.8	0	1
<i>Lethrinus spp.</i>	24.4	4	9.8	10.3	10
<b>Lutjanidae</b>					
<i>Lutjanus argentimaculatus</i>	0	0	0	13.8	0
<i>Lutjanus carponotatus</i>	24.4	20	0	0	5
<i>Lutjanus ehrenbergii</i>	0	0	0	50	1
<i>Lutjanus fulviflamma</i>	12.8	12	7.3	36.2	17
<i>Lutjanus fulvus</i>	10.5	4	0	17.2	0
<i>Lutjanus gibbus</i>	18.6	6.7	4.9	0	1
<i>Lutjanus monostigma</i>	4.7	1.3	0	3.4	0
<i>Lutjanus russellii</i>	8.1	0	0	13.8	0
<i>Lutjanus semicinctus</i>	39.5	44	31.7	0	1
<b>Monacanthidae</b>					
<i>Acreichthys tomentosus</i>	0	0	0	0	19
<b>Mugilidae</b>					
<i>Ellochelon vaigiensis</i>	0	0	0	1.7	3
<i>Mugilidae spp.</i>	0	0	0	3.4	2
<b>Mullidae</b>					
<i>Mullidae spp.</i>	0	0	0	1.7	0
<i>Mulloidichthys flavolineatus</i>	14	6.7	2.4	3.4	5

	<i>Parupeneus barberinus</i>	37.2	54.7	22	8.6	29
	<i>Parupeneus crassilabris</i>	33.7	38.7	9.8	0	0
	<i>Parupeneus cyclostomus</i>	11.6	4	2.4	0	0
	<i>Parupeneus indicus</i>	1.2	5.3	19.5	3.4	3
	<i>Upeneus tragula</i>	2.3	0	0	1.7	4
<b>Muraenidae</b>						
	<i>Muraenidae spp.</i>	0	0	0	0	1
<b>Nemipteridae</b>						
	<i>Pentapodus trivittatus</i>	61.6	61.3	61	22.4	31
	<i>Scolopsis bilineata</i>	19.8	6.7	2.4	0	0
	<i>Scolopsis ciliata</i>	10.5	4	0	1.7	2
	<i>Scolopsis lineata</i>	7	34.7	43.9	19	7
	<i>Scolopsis margaritifera</i>	48.8	33.3	14.6	0	0
	<i>Scolopsis trilineata</i>	0	0	9.8	0	0
<b>Pomacentridae</b>						
	<i>Abudefduf lorenzi</i>	0	0	0	22.4	1
	<i>Abudefduf sexfasciatus</i>	26.7	20	7.3	1.7	0
	<i>Abudefduf sordidus</i>	0	0	0	1.7	0
	<i>Abudefduf vaigiensis</i>	5.8	5.3	0	5.2	0
	<i>Chrysiptera biocellata</i>	0	0	2.4	1.7	0
	<i>Chrysiptera unimaculata</i>	0	1.3	2.4	0	0
	<i>Dischistodus chrysopoecilus</i>	0	30.7	65.9	0	3
	<i>Dischistodus melanotus</i>	23.3	62.7	4.9	0	0
	<i>Dischistodus perspicillatus</i>	4.7	16	7.3	0	1
	<i>Dischistodus prosopotaenia</i>	0	2.7	0	0	1
	<i>Pomacentridae spp.</i>	33.7	33.3	9.8	0	0
	<i>Pomacentrus adelus</i>	18.6	14.7	2.4	1.7	0
	<i>Pomacentrus bankanensis</i>	17.4	17.3	4.9	0	0
	<i>Pomacentrus simsiang</i>	0	8	0	0	1
	<i>Pomacentrus tripunctatus</i>	0	1.3	0	3.4	0
	<i>Stegastes albifasciatus</i>	1.2	12	17.1	0	0
<b>Scaridae</b>						
	<i>Chlorurus sordidus</i>	32.6	13.3	0	0	1
	<i>Hipposcarus longiceps</i>	40.7	8	0	0	3
	<i>Scarus globiceps</i>	1.2	1.3	2.4	0	0
	<i>Scarus psittacus</i>	9.3	0	2.4	0	0
	<i>Scarus spinus</i>	3.5	5.3	2.4	0	0
	<i>Scarus spp.</i>	89.5	86.7	17.1	1.7	8
<b>Scatophagidae</b>						
	<i>Scatophagus argus</i>	0	0	0	8.6	0
<b>Serranidae</b>						
	<i>Cephalopholis spiloparaea</i>	0	0	2.4	0	0
	<i>Epinephelus corallicola</i>	0	0	0	1.7	0
<b>Siganidae</b>						
	<i>Siganus argenteus</i>	12.8	6.7	0	0	1
	<i>Siganus canaliculatus</i>	12.8	22.7	12.2	3.4	34
	<i>Siganus doliatus</i>	25.6	32	0	1.7	2
	<i>Siganus lineatus</i>	47.7	8	9.8	43.1	16
	<i>Siganus spinus</i>	1.2	16	56.1	5.2	11

	<i>Siganus spp.</i>	1.2	1.3	0	0	5
	<i>Siganus vermiculatus</i>	0	0	0	5.2	2
<b>Sphyraenidae</b>						
	<i>Sphyraena barracuda</i>	0	0	0	0	3
	<i>Sphyraenidae spp.</i>	0	0	0	0	2
<b>Tetraodontidae</b>						
	<i>Arothron hispidus</i>	0	0	0	1.7	0
<b>Toxotidae</b>						
	<i>Toxotes jaculatrix</i>	0	0	0	17.2	0

**Table A3.2.** Frequency of occurrence (%) and mean MaxN ( $\pm$  SE) for the eighteen coral reef-associated fish species observed in all five habitat types. Bold indicates habitat most frequently observed in.

Family	Species	Reef slope		Reef flat		Macroalgae		Mangrove		Seagrass	
		<i>(n=86)</i>		<i>(n=75)</i>		<i>(n=41)</i>		<i>(n=58)</i>		<i>(n=100)</i>	
		Freq (%)	Mean MaxN $\pm$ SE	Freq (%)	Mean MaxN $\pm$ SE						
Chaetodontidae	<i>Chaetodon vagabundus</i>	<b>37.2</b>	0.5 $\pm$ 0.1	36.0	0.6 $\pm$ 0.1	24.4	0.2 $\pm$ 0.1	22.4	0.3 $\pm$ 0.1	3.0	0.0 $\pm$ 0.0
Labridae	<i>Cheilinus trilobatus</i>	27.9	0.3 $\pm$ 0.1	<b>28.0</b>	0.3 $\pm$ 0.1	2.4	0.0 $\pm$ 0.0	3.4	0.1 $\pm$ 0.0	3.0	0.0 $\pm$ 0.0
Labridae	<i>Choerodon anchorago</i>	32.6	0.4 $\pm$ 0.1	<b>66.7</b>	0.9 $\pm$ 0.1	63.4	1.0 $\pm$ 0.2	15.5	0.2 $\pm$ 0.0	30.0	0.4 $\pm$ 0.1
Labridae	<i>Halichoeres nigrescens</i>	2.3	0.0 $\pm$ 0.0	18.7	0.3 $\pm$ 0.1	<b>41.5</b>	0.8 $\pm$ 0.2	5.2	0.1 $\pm$ 0.0	1.0	0.0 $\pm$ 0.0
Labridae	<i>Halichoeres scapularis</i>	15.1	0.2 $\pm$ 0.1	58.7	1.2 $\pm$ 0.2	<b>73.2</b>	1.9 $\pm$ 0.3	6.9	0.1 $\pm$ 0.1	16.0	0.2 $\pm$ 0.1
Labridae	<i>Halichoeres trimaculatus</i>	12.8	0.2 $\pm$ 0.0	38.7	0.5 $\pm$ 0.1	<b>53.7</b>	0.7 $\pm$ 0.1	5.2	0.1 $\pm$ 0.1	4.0	0.0 $\pm$ 0.0
Labridae	<i>Stethojulis interrupta</i>	4.7	0.1 $\pm$ 0.1	9.3	0.1 $\pm$ 0.1	<b>24.4</b>	0.6 $\pm$ 0.2	1.7	0.0 $\pm$ 0.0	5.0	0.1 $\pm$ 0.1
Labridae	<i>Stethojulis strigiventer</i>	8.1	0.1 $\pm$ 0.0	30.7	0.5 $\pm$ 0.1	<b>43.9</b>	0.8 $\pm$ 0.2	5.2	0.1 $\pm$ 0.1	28.0	1.2 $\pm$ 0.4
Lethrinidae	<i>Lethrinus harak</i>	12.8	0.2 $\pm$ 0.1	50.7	1.8 $\pm$ 0.3	<b>73.2</b>	3.0 $\pm$ 0.4	36.2	0.5 $\pm$ 0.1	53.0	1.7 $\pm$ 0.3
Lutjanidae	<i>Lutjanus fulviflamma</i>	12.8	1.0 $\pm$ 0.4	12.0	0.3 $\pm$ 0.1	7.3	0.3 $\pm$ 0.2	<b>36.2</b>	1.6 $\pm$ 0.4	17.0	0.3 $\pm$ 0.1
Mullidae	<i>Mulloidichthys flavolineatus</i>	<b>14.0</b>	0.8 $\pm$ 0.4	6.7	0.2 $\pm$ 0.1	2.4	0.0 $\pm$ 0.0	3.4	0.0 $\pm$ 0.0	5.0	0.1 $\pm$ 0.0

Mullidae	<i>Parupeneus barberinus</i>	37.2	0.4 ± 0.1	<b>54.7</b>	0.9 ± 0.1	22.0	0.5 ± 0.2	8.6	0.1 ± 0.1	29.0	0.8 ± 0.2
Mullidae	<i>Parupeneus indicus</i>	1.2	0.0 ± 0.0	5.3	0.1 ± 0.0	<b>19.5</b>	0.3 ± 0.1	3.4	0.0 ± 0.0	3.0	0.0 ± 0.0
Nemipteridae	<i>Pentapodus trivittatus</i>	<b>61.6</b>	0.8 ± 0.1	61.3	0.8 ± 0.1	61.0	0.9 ± 0.1	22.4	0.2 ± 0.1	31.0	0.5 ± 0.1
Nemipteridae	<i>Scolopsis lineata</i>	7.0	0.2 ± 0.1	34.7	1.8 ± 0.5	<b>43.9</b>	2.5 ± 0.9	19.0	0.7 ± 0.3	7.0	0.3 ± 0.1
Siganidae	<i>Siganus canaliculatus</i>	12.8	0.8 ± 0.5	22.7	1.5 ± 0.5	12.2	0.6 ± 0.4	3.4	0.1 ± 0.1	<b>34.0</b>	1.7 ± 0.4
Siganidae	<i>Siganus lineatus</i>	<b>47.7</b>	6.5 ± 1.5	8.0	0.8 ± 0.7	9.8	0.4 ± 0.2	43.1	1.8 ± 0.5	16.0	0.7 ± 0.2
Siganidae	<i>Siganus spinus</i>	1.2	0.0 ± 0.0	16.0	0.7 ± 0.3	<b>56.1</b>	4.3 ± 1.0	5.2	0.5 ± 0.3	11.0	0.8 ± 0.3

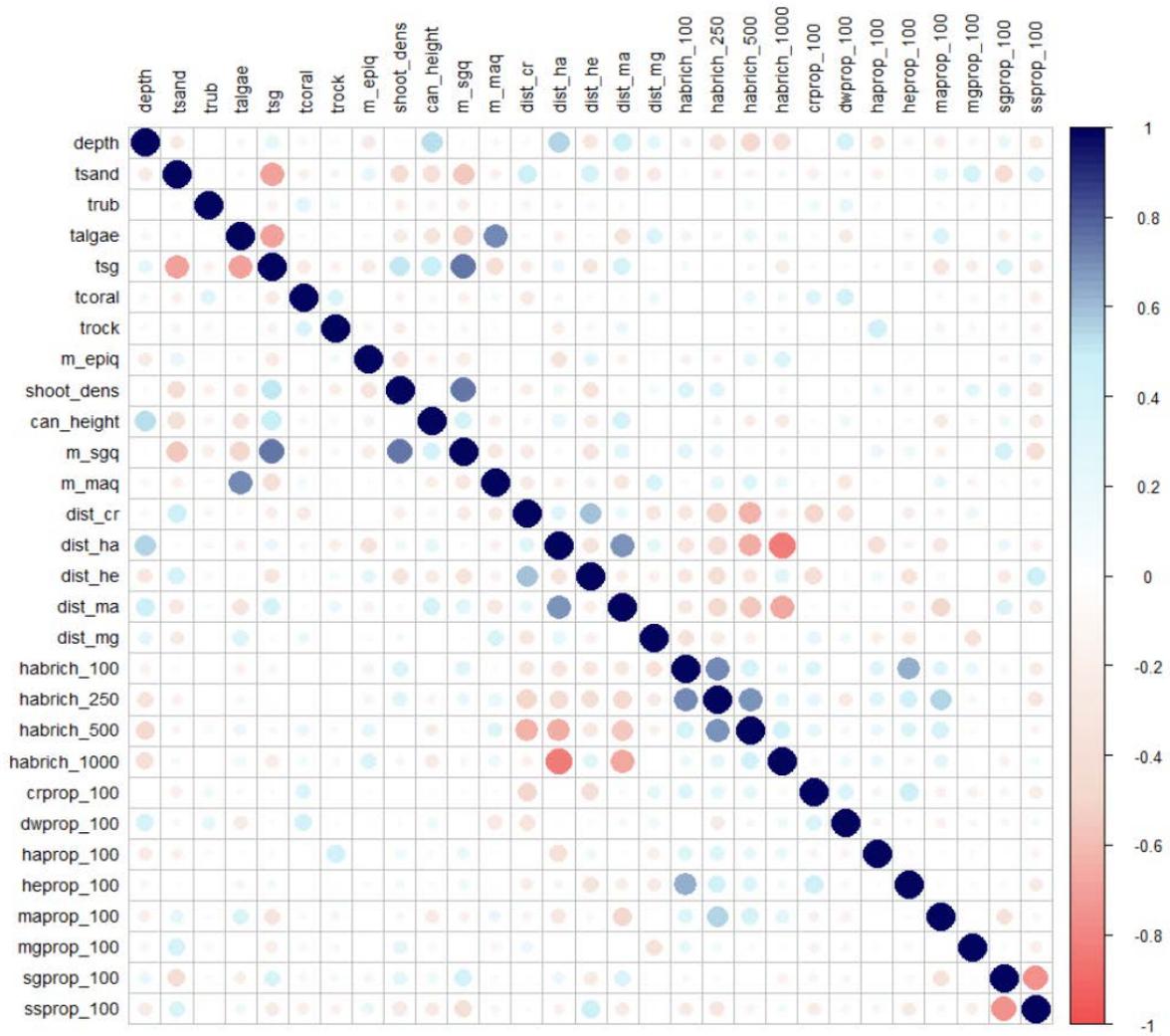
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## Appendix C: Supplementary materials for Chapter 4

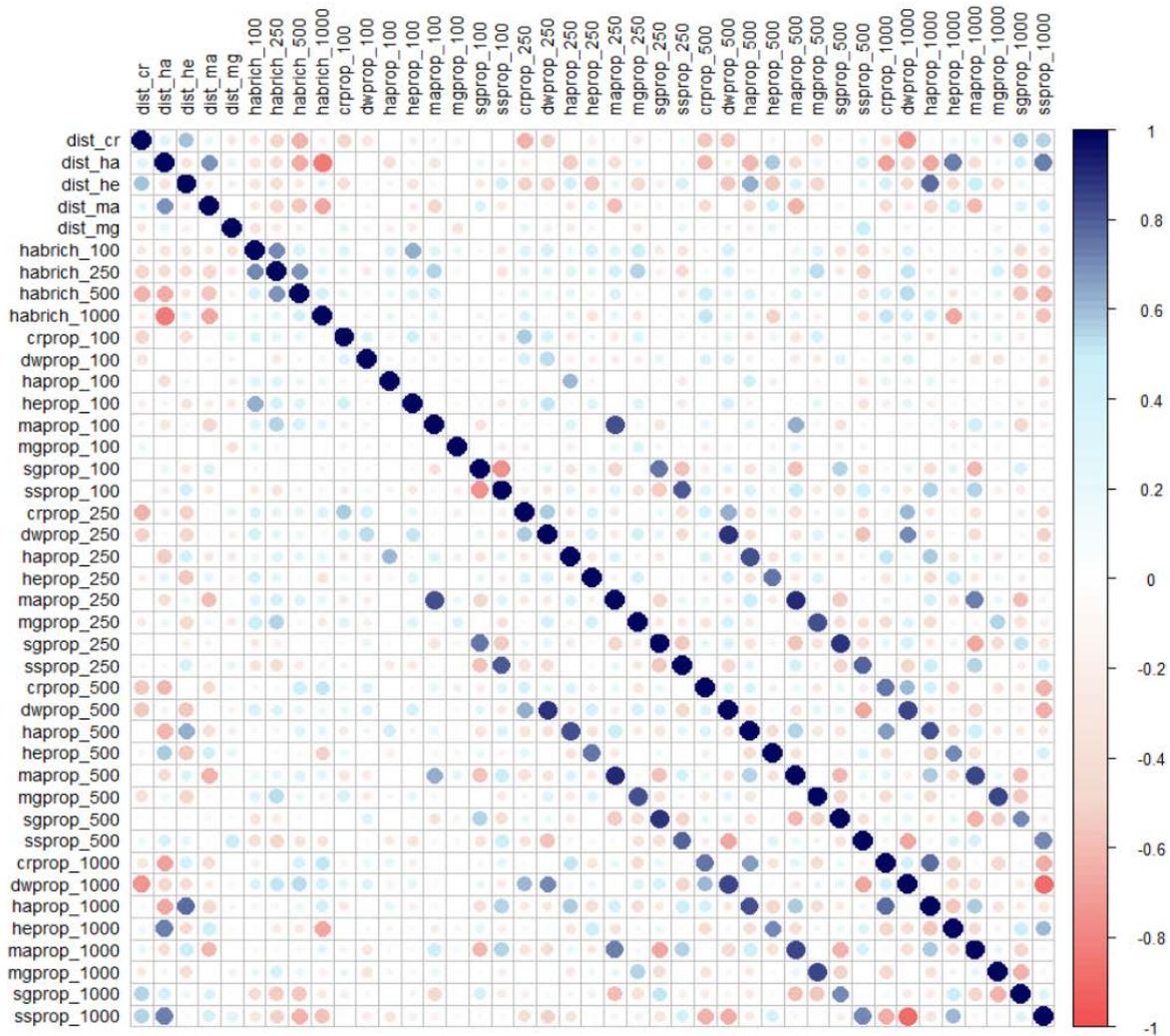
### *Relative importance of seascape versus within-habitat variables on the distribution of fishes in tropical seagrass beds*

**Table A4.1.** Final BRT models for each response variable following model optimisation, optimal number of trees and % deviance explained.

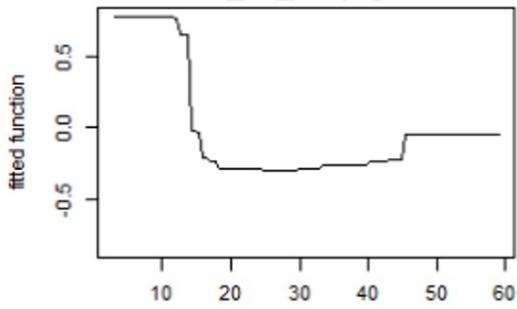
Response variable	Metric	Distribution	Tree complexity	Learning rate	Bag fraction	Number of trees	Deviance explained
<b>Families</b>							
Labridae (exc. Scarinae)	Abundance	Poisson	3	0.001	0.8	5800	0.83
Lethrinidae	Presence	Bernoulli	1	0.001	0.7	7050	0.53
Lutjanidae	Abundance	Poisson	1	0.001	0.5	5850	0.78
Mullidae	Presence	Bernoulli	5	0.0001	0.5	8150	0.20
Nemipteridae	Presence	Bernoulli	1	0.001	0.7	7400	0.48
Pomacentridae	Abundance	Poisson	1	0.001	0.8	2850	0.44
Scarinae	Abundance	Poisson	2	0.001	0.7	4900	0.77
Siganidae	Abundance	Poisson	3	0.001	0.7	4550	0.88
<b>Adults vs juveniles</b>							
Lethrinid Adult	Presence	Bernoulli	5	0.001	0.7	2300	0.54
Lethrinid Juvenile	Presence	Bernoulli	5	0.001	0.7	2200	0.56
Scarinae TP/IP	Presence	Bernoulli	4	0.001	0.8	1650	0.56
Scarinae Juvenile	Presence	Bernoulli	1	0.001	0.5	6900	0.42
<b>Species / genera</b>							
<i>Lethrinus harak</i> (Lethrinidae)	Presence	Bernoulli	3	0.001	0.5	4250	0.59
<i>Lutjanus fulviflamma</i> (Lutjanidae)	Presence	Bernoulli	3	0.001	0.7	5900	0.89
<i>Halichoeres scapularis</i> (Labridae)	Abundance	Poisson	5	0.001	0.5	1500	0.49
<i>Halichoeres trimaculatus</i> (Labridae)	Abundance	Poisson	2	0.001	0.7	2650	0.75
<i>Other Halichoeres spp.</i> (Labridae)	Abundance	Poisson	2	0.001	0.7	7350	0.81
<i>Stethojulis strigiventer</i> (Labridae)	Abundance	Poisson	1	0.001	0.8	2550	0.57
<i>Siganus canaliculatus</i> (Siganidae)	Presence	Bernoulli	2	0.001	0.7	1850	0.40
<i>Siganus lineatus</i> (Siganidae)	Presence	Bernoulli	2	0.001	0.7	2400	0.56
<i>Siganus spinus</i> (Siganidae)	Presence	Bernoulli	5	0.001	0.8	6700	0.94



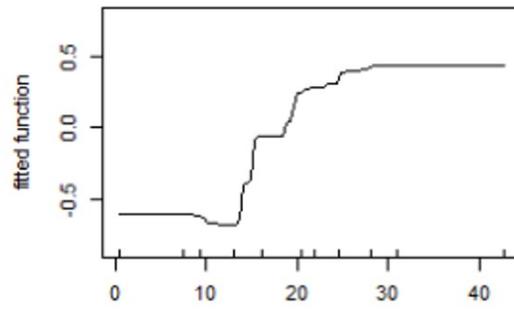
**Figure A4.1.** Multi-collinearity matrices between within-habitat and seascape predictor variables. Only includes area metrics at 100 m radius.



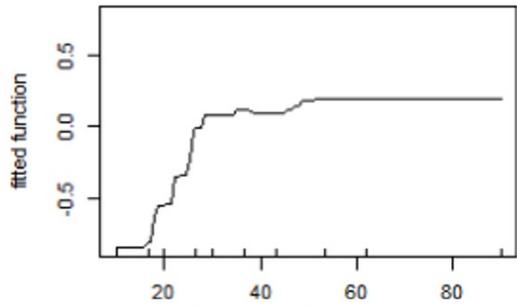
**Figure A4.2.** Multi-collinearity matrices between all seascape predictor variables examined across the four spatial scales (100 m, 250 m, 500 m and 1000 m radii).



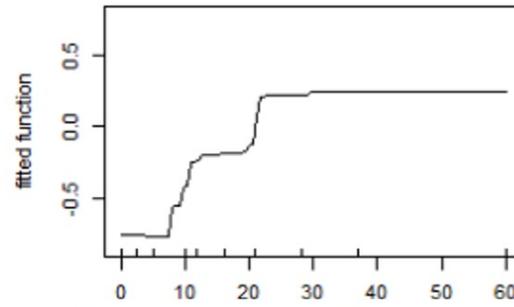
**% area of sand 250 m radius (14.1%)**



**Shoot density (12.3%)**

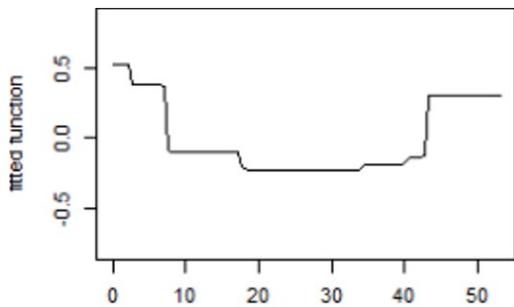


**% cover of epiphytes - quadrat (12.1%)**

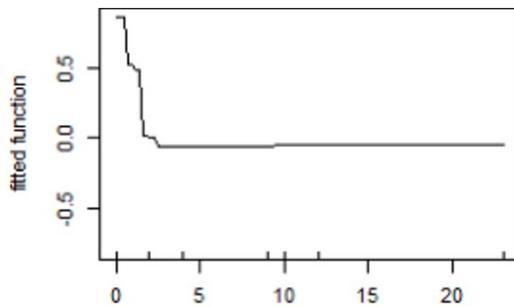


**% cover macroalgae - quadrat (11.1%)**

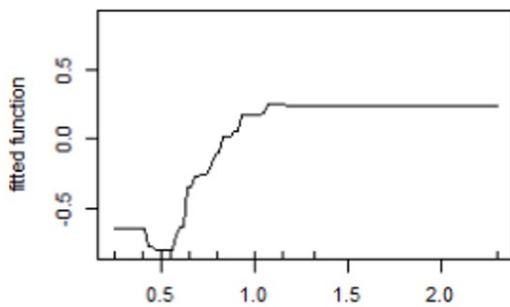
**Figure A4.3.** Partial dependency plots for the four most influential predictor variables explaining the presence of juvenile parrotfishes (subfamily: Scarinae) in seagrass beds.



**% area of deep water 1000 m radius (18.6%)**



**% area of coral reef 250 m radius (15.3%)**



**Depth (15%)**

**Figure A4.4.** Partial dependency plots for four most influential predictor variables explaining the presence of juvenile emperors (family: Lethrinidae) in seagrass beds.

**Table A4.2.** Moran’s I spatial autocorrelation for all response and predictor variables based on the average distance from each site (n = 32 sites) to its eight nearest neighbours. Following Hamylton & Barnes (2018), variables with a Moran’s I value > 0.4 are considered spatially autocorrelated (denoted with \*).

Variable	Type	Description	IDW-Threshold-1394m-Standardised		
			Moran's	z	p-value
Labridae	Response	Taxa abundance/ presence	0.105	1.663	0.096
Lethrinidae			0.116	1.569	0.117
Lutjanidae			0.010	1.701	0.089
Mullidae			0.021	0.558	0.577
Nemipteridae			0.158	1.987	0.047
Pomacentridae			-0.109	-0.894	0.372
Scarinae			-0.064	-0.526	0.599
Siganidae			0.097	1.477	0.140
Scarinae adult			0.182	2.987	0.003
Scarinae juvenile			-0.055	-0.387	0.698
Lethrinidae adult			0.026	0.808	0.419
Lethrinidae juvenile			0.128	1.699	0.089
<i>Lutjanus fulviflamma</i>			-0.016	1.346	0.178
<i>Lethrinus harak</i>			0.026	0.808	0.419
<i>Parupeneus barberinus</i>			0.035	0.709	0.478
<i>Siganus canaliculatus</i>			-0.073	-0.462	0.644
<i>Siganus lineatus</i>			-0.013	1.623	0.104
<i>Siganus spinus</i>			0.251	3.474	0.001
<i>Stethojulis strigiventer</i>			0.165	3.567	0.000
<i>Halichoeres spp.</i>			0.066	1.036	0.300
<i>Halichoeres scapularis</i>	-0.050	-0.202	0.840		
<i>Halichoeres trimaculatus</i>	-0.020	0.120	0.905		
Depth	Predictor	Measured at 25 m transect	0.097	1.278	0.201
% cover sand			0.290	3.376	0.001
% cover rubble			0.002	0.677	0.498
% cover macroalgae			-0.019	0.113	0.910
% cover seagrass			0.032	0.626	0.531
% cover hard coral			-0.025	0.163	0.870
% cover rock			0.042	1.057	0.290
Shoot density	Predictor	Measured by 0.25m <sup>2</sup> quadrats	0.113	1.445	0.149
Canopy height			0.296	3.311	0.001
% cover epiphytes			0.145	1.799	0.072

% cover seagrass			-0.038	-0.079	0.937
% cover macroalgae			-0.120	-0.917	0.359
Distance to coral reef	Predictor	Seascape metrics	0.201	2.413	0.016
Distance to hardground			0.105	1.355	0.175
Distance to heterogeneous			0.217	2.572	0.010
Distance to macroalgae			-0.026	0.044	0.965
Distance to mangrove			0.065	0.980	0.327
Habitat richness (100 m radius)			-0.051	-0.212	0.832
Habitat richness (250 m radius)			0.031	0.617	0.537
Habitat richness (500 m radius)			-0.018	0.121	0.904
Habitat richness (1000 m radius)	0.114		1.449	0.147	
Area of coral reef (100 m radius)	-0.042		-0.131	0.895	
Area of deep water (100 m radius)	0.326		3.943	0.001	
Area of hardground (100 m radius)	-0.079		-0.556	0.579	
Area of heterogeneous (100 m radius)	Only 1 site has heterogeneous habitat within 100 m				
Area of macroalgae (100 m radius)	-0.010		0.234	0.815	
Area of mangrove (100 m radius)	-0.044		-0.163	0.870	
Area of seagrass (100 m radius)	0.115		1.481	0.139	
Area of sand (100 m radius)	0.064		0.941	0.347	
Area of coral reef (1000 m radius)	0.572*		6.045	0.000	
Area of deep water (1000 m radius)	0.364		3.922	0.000	
Area of hardground (1000 m radius)	0.190		2.235	0.025	
Area of heterogeneous (1000 m radius)	0.132		1.644	0.100	
Area of macroalgae (1000 m radius)	0.316		3.490	0.000	
Area of mangrove (1000 m radius)	0.606*		6.481	0.000	
Area of seagrass (1000 m radius)	0.185		2.157	0.031	
Area of sand (1000 m radius)	0.302	3.322	0.001		

## Appendix D: Supplementary materials for Chapter 5

### *Influence of seascape and within-reef variables on a key ecosystem process, macroalgal browsing, on coral reefs in Papua New Guinea*

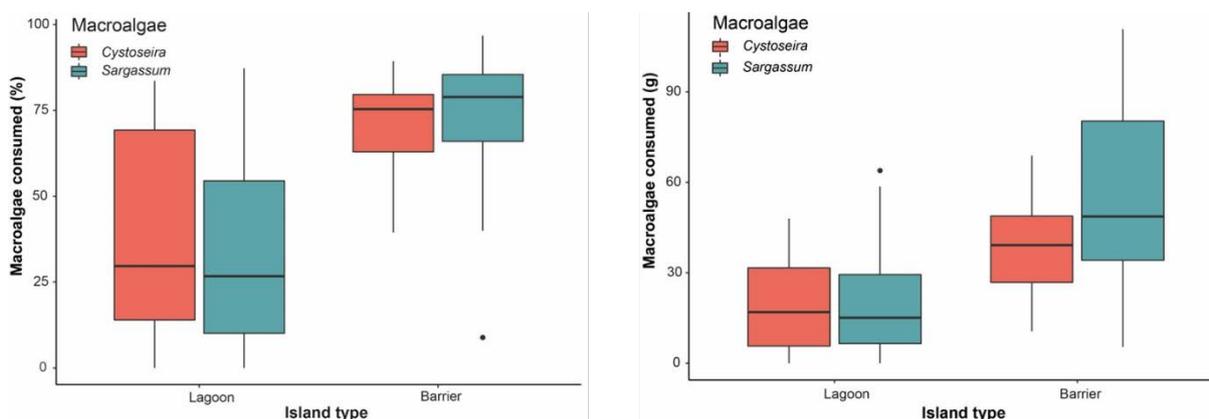
**Table A5.1.** Predictor variables used to examine the influence of seascape structure and within-reef benthic variables on browsing activity on coral reefs.

Seascape variable	Within-reef benthic variables	
Distance to oceanic environment	<i>Acropora</i> tabular	Other massive corals
<i>Area of habitat in 250 m radii</i>	<i>Acropora</i> staghorn	Other encrusting corals
Hard & soft coral	<i>Acropora</i> bottlebrush	Soft coral
Hardground	<i>Acropora</i> digitate	Epilithic algal matrix (EAM)
Macroalgae	Other <i>Acropora</i>	Crustose coralline algae (CCA)
Sand	<i>Pocillopora</i>	Sand
Seagrass	Massive <i>Porites</i>	Rubble
	Branching <i>Porites</i>	Macroalgae

**Table A5.2.** Moran's I spatial autocorrelation for response and predictor variables based on the average distance from each site (n = 24 sites) to its four nearest neighbours. Following Hamylton & Barnes (2018), variables with a Moran's I value >0.4 are considered spatially autocorrelated (denoted with \*).

Variable	Type	IDW-Threshold-3459m-Standardised		
		Moran's	z	p-value
MaxN browsers (>99% of bites spp. only)	Response	0.109	0.992	0.321
<i>Kyphosus cinerascens</i> MaxN		-0.025	0.317	0.751
<i>Kyphosus vaigiensis</i> MaxN		-0.093	-0.449	0.653
<i>Naso lituratus</i> MaxN		0.208	1.710	0.087
<i>Naso unicornis</i> MaxN		0.087	0.880	0.379
<i>Siganus canaliculatus</i> MaxN		0.021	0.550	0.582
<i>Siganus doliatus</i> MaxN		-0.056	-0.080	0.937
Total mass standardised bites (>99% of bites spp. only)		<b>0.626*</b>	4.387	0.000
<i>Kyphosus cinerascens</i> ms bites		-0.036	0.252	0.801
<i>Kyphosus vaigiensis</i> ms bites		-0.193	-1.044	0.297
<i>Naso lituratus</i> ms bites		<b>0.598*</b>	4.428	0.000
<i>Naso unicornis</i> ms bites		0.208	1.747	0.080

<i>Siganus canaliculatus</i> ms bites		0.030	0.533	0.594
<i>Siganus doliatus</i> ms bites		-0.138	-0.800	0.424
Mean macroalgae consumed (%) per site		<b>0.686*</b>	4.563	0.000
Area of hard & soft coral habitat (250 m radius)	Predictor	0.221	1.875	0.061
Area of hardground/heterogeneous habitat (250 m radius)		0.178	1.228	0.119
Area of macroalgal beds (250 m radius)		0.362	2.748	0.006
Area of seagrass beds (250 m radius)		0.367	2.990	0.003
Area of sand (250 m radius)		<b>0.479*</b>	3.252	0.001

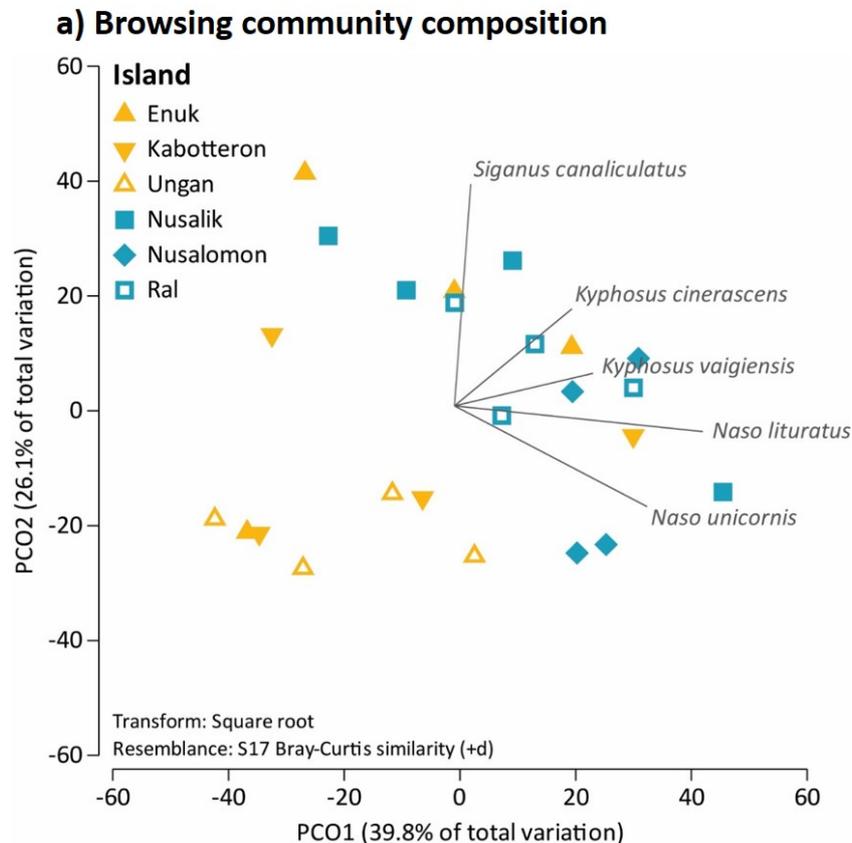


**Figure A5.1.** Comparison of macroalgae assays consumed between reef positions for each genera of macroalgae. Left figure shows percent consumed, right figure shows removal in grams (g).

**Table A5.3.** Pairwise PERMANOVA tests comparing macroalgal removal rates around each island.

Islands	t	P(perm)	Unique permutations
Eruk, Kabotteron	1.4022	0.130	9947
Eruk, Ungan	0.4816	0.832	9942
Eruk, Nusalik	2.5547	0.002	9942
Eruk, Nusalomon	4.1847	0.000	9934
Eruk, Ral	4.0519	0.000	9937
Kabotteron, Ungan	1.6298	0.074	9951
Kabotteron, Nusalik	4.4622	0.000	9954
Kabotteron, Nusalomon	6.6043	0.000	9936
Kabotteron, Ral	6.5076	0.000	9956
Ungan, Nusalik	2.4369	0.006	9941
Ungan, Nusalomon	3.6854	0.000	9956
Ungan, Ral	3.5993	0.000	9948

Nusalik, Nusalomon	4.3651	0.000	9916
Nusalik, Ral	3.5295	0.000	9937
Nusalomon, Ral	2.0799	0.043	9918



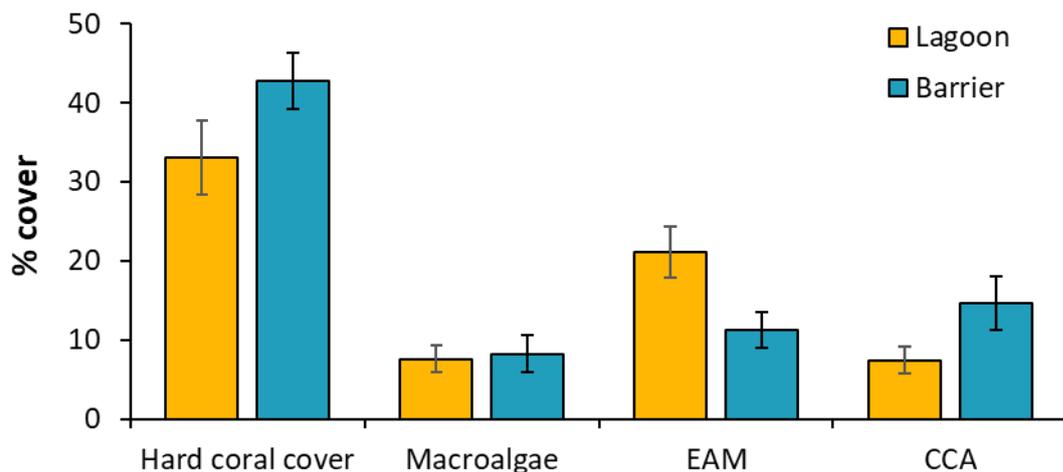
**Figure A5.2.** PCO of relative abundance (MaxN) of browsing community for the six browsing species accounting for 99% of mass standardised bites. Vectors show species contributing to community patterns. Yellow = lagoon island reefs, blue = barrier island reefs.

#### *Characterising the within-reef benthic community composition and seascape structure*

We characterised the assay sites at two different spatial scales: within-reef benthic community composition, and the seascape composition within 250 m radius. Differences in the benthic community along transects and the seascape composition surrounding a site were explored using a two-factor nested permutational analysis of variance (PERMANOVA) with reef position (i.e. barrier vs lagoon) as a fixed factor, and island as a random factor nested within reef position. For the benthic community analysis, transect data were averaged by site and square-root transformed. Seascape data were log transformed. Resemblance matrices were constructed based on Bray-Curtis for the benthic community data and Euclidean distances for the seascape data. Patterns were visualised using principal component analysis (PCA).

### *Within-reef benthic community composition*

Mean live hard coral cover was variable among sites ranging from  $8\% \pm 3.7$  SE to  $62\% \pm 2.6$  SE (overall mean  $37.9\% \pm 3.1$  SE) (Figure A5.3). Macroalgae cover was generally low across all sites ( $7.9\% \pm 1.4$  SE; Figure A5.3), with *Halimeda* spp. dominating the algal assemblage. There was no significant difference in benthic community composition between the barrier and lagoon sites (pseudo- $F = 1.41$ ,  $df = 1$ ,  $p(\text{mc}) = 0.24$ ) (Figure A5.4a, Table A5.4). However, there was a significant difference between islands nested in reef position (pseudo- $F = 1.92$ ,  $df = 4$ ,  $p = 0.006$ ) (Table A5.4). Pairwise comparisons revealed that the difference was attributable to one barrier reef island, Nusalik, which had a different benthic community to the other two barrier reef islands ( $p(\text{mc}) < 0.05$ ). SIMPER analysis identified that this was driven by higher amounts of epilithic algae matrix (EAM) and *Porites* spp. at Nusalik, compared to higher amounts of crustose coralline algae (CCA) at the other two barrier reef islands (i.e., Nusalomon and Ral).



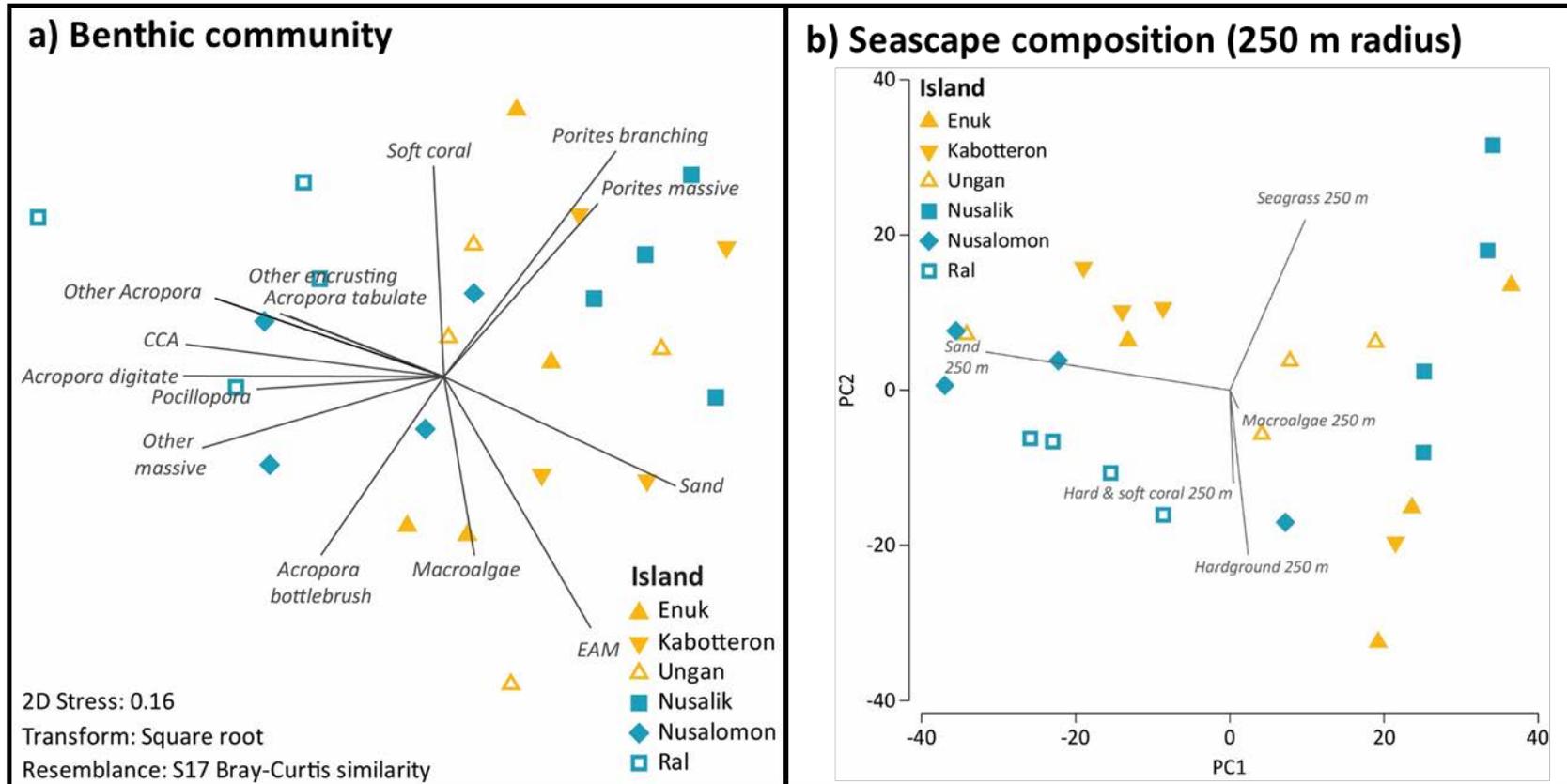
**Figure A5.3.** Comparison of four ecologically relevant benthic habitat categories between lagoon and barrier reef sites.

### *Seascape structure*

The composition of the seascape within a 250 m radius of each site was also significantly different between islands nested in reef position ( $df = 4$ , pseudo- $F = 2.81$ ,  $p = 0.005$ ) but not between the lagoon and barrier reef sites (pseudo- $F = 1.99$ ,  $df = 1$ ,  $p(\text{mc}) = 0.168$ ) (Figure A5.4b, Table A5.4). Pairwise comparisons showed that the barrier reef island, Nusalik, had a different seascape composition to the other two barrier reef islands, Nusalomon and Ral ( $p(\text{mc}) < 0.01$ ). SIMPER analysis identified that this was driven by the higher areal coverage of seagrass around the Nusalik sites compared to the other two barrier reef islands.

**Table A5.4.** PERMANOVA results for comparing differences between reef position (fixed) and islands (random nested within reef position) for the benthic community and seascape composition. \*ECV denotes percent estimated components of variation.

	df	SS	MS	Pseudo- <i>F</i>	P(perm)	ECV*
<b>Benthic community</b>						
Reef position	1	14.874	14.874	1.4091	0.2322	5.1%
Island(Reef position)	4	42.224	10.556	1.9226	<b>0.0041</b>	17.8%
Residual	18	98.825	5.4903			
Total	23	155.92				
<b>Seascape composition</b>						
Reef position	1	20.413	20.413	1.9938	0.168	13.8%
Island(Reef position)	4	40.953	10.238	2.8146	<b>0.005</b>	26.9%
Residual	18	65.475	3.6375			
Total	23	126.84				



**Figure A5.4.** a) nMDS plot of benthic community, and b) PCA plot of the seascape composition to visualise differences between sites, islands and reef positions. Vectors show variables contributing to benthic and seascape patterns. Yellow = lagoon islands, blue = barrier reef islands.

