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Functional niche partitioning in herbivorous coral reef fishes

Thesis submitted by:

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Chapter 2:

- Simon Brandl: concept of study, data collection, writing of manuscript
- David Bellwood: concept of study, writing of manuscript

Chapter 3:

- Simon Brandl: concept of study, data collection, data analysis, writing of manuscript
- David Bellwood: concept of study, writing of manuscript

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Abstract

Ecological niche theory predicts that the diversity of life hinges on differences in the ways in which species exploit available resources, i.e. their ecological niche. An organism's niche can be divided into the fundamental and the realized niche. The fundamental niche refers to a species' inherent potential capabilities (based on, for instance, its morphology), while the realized niche represents a species' behaviour when it can interact with the biotic and abiotic components of its environment. Tropical coral reefs stand out as one of the most diverse ecosystems on Earth, but patterns of niche partitioning among coral reef organisms are poorly understood. Herbivorous coral reef fishes are frequently considered to perform one of the most critical ecosystem processes on coral reefs, the removal of algal and detrital material from the reef substratum. However, while several classification schemes have previously been employed to characterize the functional role of different herbivores, detailed holistic investigations of niche differences among herbivore species are rare. Therefore, in this thesis, I use social, morphological, behavioural, and environmental factors to disentangle the functional niches of herbivorous fishes on coral reefs.

One of the most prominent social systems in animals is the association between two individuals, commonly termed a 'pair'. However, given that sexual reproduction in higher animals commonly requires the association between two partners, reproductive connotations frequently override the social-ecological benefits that a partner can provide and thus, the potential effects of pairing on an animal's functional niche. In teleost fishes, pairing is common but little is known about the potential drivers and consequences of this behaviour. My first objective, therefore, was to quantify the extent of pair-formation in coral reef fishes and to examine potential ecological/reproductive correlates of pair-formation. Of a total of 1,981 species of Indo-Pacific reef fishes, 341 (17.2%) are reported to form pairs, with pair-forming species being particularly common (more than 50% of species) in five families. Pair-

forming species had few commonalities with regards to their reproductive strategies. Instead, two ecological traits appear to be prevalent in pairing species: 1) the consumption of small, benthic, and relatively immobile prey items, and 2) the maintenance of permanent burrows. Based on these results, I conclude that pairing may have important ecological benefits in coral reef fishes.

To further explore this hypothesis, I performed an ecomorphological assessment of pair-forming fishes, evaluating whether pairing fishes are also morphologically similar. Using a suite of six morphological traits, predominantly describing the cranial region and overall body shape of reef fishes, I demonstrate that morphology can accurately predict the prevalence of pairing behaviour in 47 species of benthos-feeding, reef fishes and that there is a strong relationship between morphology and pairing behaviour in three common families of reef fishes, the Acanthuridae, Chaetodontidae, and Siganidae. Basically, pair-forming fishes are characterized by concave foreheads, pointed snouts, deep bodies, and large eyes, attributes that relate to the feeding on small prey in topographically complex environments.

I then sought to examine specific benefits of pair-forming behaviour using four species of pairing rabbitfishes as model organisms. In doing so, I revealed evidence for a reciprocal cooperative system in rabbitfishes. In all four species, vigilance was strongly coordinated between pair members, with one individual assuming a head-up position elevated vertically above the substratum, while the partner was feeding, often in small cracks and crevices. Pair members altered their positions more often than one fish feeding continuously. Furthermore, compared to individual fishes, fishes in pairs exhibited longer vigilance bouts, more bites per foray, and deeper penetration of the substratum. By evaluating these findings against a set of six requirements for reciprocal cooperation, I show that the observed behaviour in rabbitfishes may be based on the reciprocal exchange of food and safety

between pair members and that the presence of a partner permits the foraging in concealed microhabitats, unavailable for solitary fishes.

Based on these findings, I examined the fine-scale utilization of foraging microhabitats in herbivorous coral reef fishes. To do so, I introduced a novel individual-based analysis of functional niche overlap in animal communities, which uses convex hull volumes in order to assess niche breadth, overlap, and turnover vs. nestedness in multiple dimensions. Using this analysis on *in situ* behavioural data of the foraging behaviour of 21 species of herbivorous reef fishes, I show that there is limited functional redundancy among reef fish species, with an average niche overlap of only 15.2%. I further reveal a clear distinction between species that utilize predominantly flat, exposed surfaces of sand or bare rock, and species that feed on a wide range of different microhabitats, including cracks and crevices in dead corals and other complex substrata. Thus, the utilization of foraging microhabitats appears to represent an important axis of niche partitioning in herbivorous coral reef fishes.

Using the family Acanthuridae as a study subject, I then quantitatively explored ecological specialization in reef fishes with regards to both fundamental and realized niches. Specifically, I examined ecological specialization in ten surgeonfish species with regards to morphology and two realized niche axes associated with diet and foraging microhabitat utilization, and investigated the relationships between morphological and behavioural specialization. These relationships differed markedly from the traditional ecomorphological paradigm. While morphological specialization showed no relationship with dietary specialization, it exhibited a strong relationship with foraging microhabitat specialization. However, this relationship was inverted: species with specialized morphologies were microhabitat generalists while generalized morphotypes were microhabitat specialists. Thus, I highlight the potential importance of including niche axes beyond dietary specialization into ecomorphological frameworks and suggest that, on coral reefs, morphotypes commonly

perceived as most generalized may, in fact, be specialized in exploiting flat and easily accessible microhabitats.

To investigate the role of concealed microhabitats for benthic organisms, I then performed an observational study on the distribution patterns of juvenile corals and early life-stage macroalgae. Specifically, I examined the microhabitat occupation of juvenile acroporid and pocilloporid corals and early life-stages of the macroalgae *Turbinaria ornata*, revealing that both corals and macroalgae were more prevalent in concealed microhabitats when compared to open or semi-concealed microhabitats. Corals were more common on the reef crest, while macroalgae were more abundant on the reef flat. The distribution patterns of herbivorous fishes also showed a distinct difference between the flat and crest, with species utilizing concealed microhabitats being more common on the crest. In particular, high abundance of pairing rabbitfishes and surgeonfishes of the genus *Zebrasoma* were positively related to high numbers of juvenile corals. Overall, this study suggested that coral reef crevices might be an important microhabitat for corals and algae alike and that the feeding activity of herbivorous fishes capable of feeding in crevices may influence this relationship.

Finally, I experimentally investigated the effects of micro-topographic refuges on grazing dynamics on coral reefs. Micro-topographic refuges decreased overall grazing pressure more than ten-fold and permitted access to only few species of cropping herbivores. In contrast, grazing pressure from detritivorous fishes was virtually absent, therefore precluding an entire ecosystem-process, the removal of particulates. As a consequence, benthic communities differed markedly between microhabitats, with micro-topographic refuges exhibiting longer algal filaments and a more diverse community, including scleractinian corals, which were completely absent from exposed microhabitats. Thus, although occurring on the scale of a few centimetres, micro-topographic refuges appear to generate fundamentally different dynamics between grazing fishes and the benthos.

Overall, the results of this thesis demonstrate that herbivory on coral reefs is not a uniform process, but a multifaceted interplay between morphological, behavioural, social, and environmental factors. This thesis has begun to disentangle the complex differences among herbivorous fish species, their drivers, and the consequences of their feeding behaviour for coral reefs. However, the results of this work suggest that micro-topographical complexity on the scale of a few centimetres represents an important ecological axis, along which fishes appear to partition their functional niches. Future research may focus on examining foraging micro-habitat utilization in reef fishes with an even higher resolution, the effects of reducing the grazing pressure exerted by crevice-feeding fishes, and the long-term consequences of the loss of functional diversity in herbivorous fishes on diversity patterns in the coral reef benthos.

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Chapter 1: General Introduction

In the middle of the 20th century, Hutchinson (1957) asked why there are so many species inhabiting planet Earth. This question has reverberated through the field of ecology throughout the second half of the century (Paine 1966; Whittaker et al. 1973; Brown 1981), and up until now, questions concerning the diversity of organismal communities have a firm hold in the ecological literature (Brown 2014; Laliberté et al. 2014; Stork et al. 2015).

Perhaps the most important legacy of Hutchinson's seminal essay was the notion that one can portray a species' ecological niche as an abstract volume in a synthetic niche space, in which the axes consist of biotic and abiotic factors that affect the species' survival (Grinnell 1917; Hutchinson 1957). Due to incongruence between the space over which a species can theoretically persist, and the realized distribution patterns of a given species, Hutchinson (1957; 1959) suggested a distinction between a species' fundamental niche (the space over which it can possibly persist), and its realized niche (the space over which a species occurs once all biotic and abiotic interactions are accounted for). This concept has been a cornerstone of niche theory throughout its existence (Whittaker et al. 1973; Holt 2009), and with the rise of environmental niche modelling, as a function of the accessibility of global environmental data, research on species' environmental, or Grinnellian, niches across the globe has flourished (Guisan & Thuiller 2005; Warren et al. 2008).

While Hutchinson proposed his niche concept in the context of a Grinnellian niche (i.e. a species' niche defined as its tolerance to a set of environmental factors), ecologists have also adapted his ideas and applied them in an Eltonian niche framework (i.e. a species' niche defined by its functional interactions with biotic and abiotic ecosystem components; Elton 1927; Leibold 1996). Hutchinson's ideas of fundamental and realized niche spaces apply to the Eltonian niche, enabling distinctions between the functional niche space a species could theoretically occupy vs. its functional niche space in real life (Whittaker et al.

1973; Chase & Leibold 2003). Yet, due to the often laborious and field-intensive methods necessary to collect locality-specific data, and the intricate interrelationships between axes in a multidimensional Eltonian niche space, realized Eltonian niches and their relationships on a community scale are often difficult to examine (Araujo & Guisan 2006; Soberon 2007; Devictor et al. 2010).

Meanwhile, however, the increasing human-mediated pressure on biodiversity has positioned species' functional niches on the centre stage of ecological research (Tilman et al. 2014). Towards the end of the 20th century, a clear consensus emerged among ecologists that a diversity of species and their functional niches is crucial for the maintenance of ecosystem functioning (Naeem et al. 1994; Loreau et al. 2001). The establishment of this paradigm was largely due to long-term experiments conducted in relatively low-diversity grassland ecosystems (Tilman & Downing 1996; Tilman et al. 1997; Diaz & Cabido 2001) in which the functional diversity of communities, measured as differences in ecological traits, is easily manipulable and ecosystem responses (by means of, for instance, primary production) are distinctly measurable. Following this, trait-based assessments of functional diversity have spread beyond grassland communities, leading to the establishment of a wide range of literature across ecosystems (Zak et al. 2003; Duffy et al. 2003; Slade et al. 2007; Flynn et al. 2009; reviewed in Hooper et al. 2005; Lefcheck et al. 2015). This has contributed to a deeper understanding of the effects of human activities on entire ecosystems and has resulted in a strong call for the conservation of diversity in the interest of humanity and the biosphere (Hooper et al. 2012; Cardinale et al. 2012; Tilman et al. 2014). However, trait-based studies rarely acknowledge that only species' fundamental niches are considered. This is problematic, given the explicit Hutchinsonian theory that a species' realized niche can be distinctly different from its fundamental niche. While recent years have seen calls for a stronger consideration of realized functional niches (Schleuter et al. 2010), this is difficult to

achieve due to the obstacles imposed by fine-scale variability in the factors governing a species functional niche, and the limits to empirical observations.

In marine ecosystems, these problems are further exacerbated by the nature of aquatic environments, which often preclude sustained observation periods and continuous measurement of abiotic and biotic variables influencing a species functional niche (Hooper et al. 2005). In this context, coral reefs are particularly challenging. In these extremely diverse environments (Connell 1978), there are countless biotic and abiotic interactions (Huston 1985), ranging from associational refuges among primary producers (Hay 1991; Pfister & Hay 1988; Roff et al. 2015), to symbioses between fishes and crustaceans (Karplus 1987; Thompson 2005) or collaborative interspecific foraging benefits for herbivores and carnivores alike (Robertson et al. 1976; Bshary et al. 2006; Vail et al. 2013). Nevertheless, the steep decline of coral reef systems due to anthropogenic disturbances has necessitated the investigation of the relationship between biodiversity and ecosystem functioning, for which researchers relied predominantly on trait-based classifications of species' fundamental functional niches (Bellwood et al. 2003; Mouillot et al. 2013; D'Agata et al. 2013; Mouillot et al. 2014). Although the results of these studies are already sobering, with dramatic declines in ecosystem functioning following moderate decreases in species diversity (Burkepile & Hay 2008; 2010; Bellwood et al. 2012; Rasher et al. 2013), there is good evidence that current trait-based classifications used to identify species' fundamental niches are insufficient (Biro et al. 2010; Fox & Bellwood 2011; Brandl & Bellwood 2014a).

One group of coral reef organisms that has received particular attention throughout the last two decades are nominally herbivorous fishes (*sensu* Choat et al. 2002). Herbivorous fishes exert intense grazing pressure on the coral reef benthos (Poore et al. 2012), and therefore strongly affect patterns of primary production, species diversity, succession, and community composition (Hixon & Brostoff 1996). As a consequence, severe reductions in

the grazing pressure of herbivorous fishes have resulted in catastrophic shifts in the benthic communities on coral reefs worldwide (Hughes 1994; Ledlie et al. 2007; Cheal et al. 2010), with dramatic implications for coral reef ecosystems (Jackson et al. 2001; Bellwood et al. 2004; Graham et al. 2006; Hughes et al. 2010; Graham et al. 2015). Due to the critical importance of herbivorous reef fishes in these ecosystem shifts, a focus has been placed on the functional categorization of herbivorous fishes (Bellwood & Choat 1990; Bellwood et al. 2004; Nyström 2006; Green & Bellwood 2009; Burkepile & Hay 2008; 2010; Rasher et al. 2013), resulting in several complementary schemes to classify the impact of herbivorous fishes on the benthos. Yet, aside from coarse divisions of fishes based on their feeding mechanics and targeted prey, variation in species' functional niches has been given little consideration, and realized niches are rarely quantified.

Two recent observations have provided an additional impetus to re-consider the functional niches of herbivorous fishes and their relationships. The first observation relates to the social systems found in herbivorous fishes, and, in particular, the formation of pairs. Brandl & Bellwood (2013b) investigated the nature of pairs in the herbivorous rabbitfish *Siganus doliatus*, one of the dominant herbivore species on the inner Great Barrier Reef (Fox & Bellwood 2007; Bennett & Bellwood 2011; Loffler et al. 2015). The prevalence of pairs in this species increased with size, and pair members maintained a tight association over extended periods of time, thus corroborating previous observations of monogamy in reef fish species (Whiteman & Côté 2003, 2004). However, 25% of all pairs in this species were found to be same-sex pairs, suggesting that pair-formation in *S. doliatus*, and possibly other pairing herbivorous reef fishes, may have other, non-reproductive, ecological benefits (Brandl & Bellwood 2013b). The second observation relates to the importance of micro-topographic complexity for the foraging of herbivorous fishes. While recognized early as a potentially important ecological axis (Robertson et al. 1979; Robertson & Gaines 1986; Choat &

Bellwood 1985; Bellwood & Choat 1990), a recent study by Fox & Bellwood (2013) emphasized the importance of micro-topographic complexity for herbivorous fishes. Interestingly, Fox & Bellwood (2013) identified three species of pairing rabbitfishes that exploited micro-topographically complex microhabitats (i.e. cracks and crevices on the reef) more often than most other herbivorous species, suggesting that foraging microhabitat utilization patterns (i.e. the ability to exploit micro-topographically complex environments) may be an important axis of niche partitioning in herbivorous reef fish communities on the Great Barrier Reef.

1.1 Thesis aim and outline

The primary aim of this thesis, therefore, was to investigate the potential of microhabitat utilization as an axis of niche partitioning in herbivorous coral reef fish communities and to explore social, morphological, behavioural, and environmental factors that drive foraging microhabitat utilization. To do so, I considered both the fundamental and realized niches of herbivorous fish species and employed a wide range of different techniques.

In **Chapter 2**, I examine the literature on pair-formation in coral reef fishes, in order to assess the likelihood of pairing as an indicator of distinct functional niches. In **Chapter 3**, I expand on these results by performing an ecomorphological study on a large community of benthic-feeding coral reef fishes and relating species' morphologies to their social system.

In **Chapter 4**, I focus on the realized benefits of pair-formation in four species of rabbitfishes and present evidence for a reciprocal cooperative system between pair members, in which partners coordinate their foraging and vigilance behaviour. In **Chapter 5**, I present a community-scale analysis of the realized niches of 21 herbivorous fish species. By introducing a novel, individual-based analysis of functional niche overlap, I present an analytical framework for the investigation of realized niche overlap in diverse communities.

In **Chapter 6**, I use this knowledge to examine ecological specialization in surgeonfishes (f. Acanthuridae).

In **Chapter 7**, I evaluate the significance of micro-topographic complexity for benthic organisms, focusing on juvenile corals and macroalgae as two of the most strongly affected organismal groups. Finally, in **Chapter 8**, I support these conclusions through an experimental assessment of the effects of micro-topographic refuges on the dynamics between grazing fishes and the coral reef benthos.

Chapter 2: Pair-formation in coral reef fishes: an ecological perspective

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2.1 Introduction

Of all the ecosystems on the planet, coral reefs are among the most diverse. Providing structural complexity and heterogeneous habitats for countless organisms, coral reefs harbour approximately 5,000 species of fishes (Bellwood et al. 2012), 1,400 species of corals (Baird et al. 2009) and at least 165,000 species of reef-associated invertebrates (Stella et al. 2011). This exceptional diversity is matched by a comparable variety of behavioural traits and almost every known social interaction can be observed in reef-dwelling organisms.

Although social and reproductive behaviours have elicited intensive theoretical and empirical examination, they are inherently difficult to study. As opposed to morphological characteristics, which are usually conspicuous in a species' external appearance and relatively consistent among individuals, social and reproductive traits are more variable and responsive to localized environmental factors (Yahner 1979). These traits thus exhibit the potential for extremely high plasticity within narrow taxonomic units such as species or families (e.g., Colin & Bell 1991; Morgan & Kramer 2004; Wong et al. 2005). Nevertheless, social and reproductive behaviour are important determinants of an organisms' ecology and a sound understanding of these traits, including their underlying environmental drivers and functional consequences, is crucial for organismic and community ecology (Orians 1961). Meaningful evaluations of social and reproductive behaviours, however, are often stymied by the close relationship between social organizations and reproduction, and the resulting ambiguity of the terms 'social system' and 'mating system' (Emlen & Oring 1977; Robertson

& Hoffmann 1977; Neudecker & Lobel 1982; Hourigan 1989; Reynolds 1996). Social groupings commonly pave the way for reproductive activity throughout the animal kingdom (e.g., Fricke 1980; Getz & Hofmann 1986). Yet, there are well-known examples in which social associations appear to be unrelated to reproductive behaviour, and instead appear to be linked to ecological benefits such as increased safety or foraging success (e.g., Robertson et al. 1976; Parrish & Edelstein-Keshet 1999; Morgan & Kramer 2004). Although invariably interrelated, this review will distinguish between ecological factors (i.e. factors related to survival and energy intake) and reproductive factors (i.e. factors enhancing reproductive output, fertilization rate or ensuring mate availability).

The distinction between social and mating systems is particularly unclear in cases where the social unit is a pair, i.e. two individuals of the same species. Sexual reproduction, by default, requires two individuals, and many animals form transient breeding pairs whose sole purpose is reproduction (e.g., Liske & Davis 1987). It may be for this reason that, in the current literature, pair-formation (i.e. the prolonged association with only one other conspecific) is, with few exceptions (e.g. Gwinner et al. 1994; Pratchett et al. 2006; Young et al. 2008; Brandl & Bellwood 2013a), predominantly linked to a monogamous mating system and circumstances that led to the evolution of monogamy (Emlen & Oring 1977; Wittenberger & Tison 1980; Barlow 1984; Fricke 1986; Reavis & Barlow 1998; Whiteman & Côté 2003, 2004; Reavis & Copus 2011). The social or ecological aspects of pair-formation, the environmental circumstances that have led to the formation of a pair, and the evolutionary factors underpinning pairing behaviour are often not considered.

Pair-formation is described in many taxa, ranging from unicellular organisms to higher vertebrate species. However, understanding of the significance of pairing varies markedly among taxa (**Table 2.1**). Pairing has been extensively studied in birds where the manifest link between pairing and monogamous mating is well established (Orians 1969;

Wittenberger & Tilson 1980; Black 2001; but see Westneat & Stewart 2003). The significance of prolonged pairing beyond the act of reproduction in birds is easily explained by biparental care (e.g. Wittenberger & Tilson 1980; Reynolds 1996; Adkins Regan 2002). In contrast, most marine fishes display virtually no characteristics commonly associated with monogamous mating (Wittenberger & Tilson 1980; Reynolds 1996) and evidence for “genetic monogamy” (i.e. exclusive reproduction with only one mate) is rare. A comprehensive review of monogamy in marine fishes (Whiteman & Côté 2004) found evidence for “genetic monogamy” in only 14.6% of the 164 species of marine fishes that were described as monogamous in the literature, while the majority (64%) display “social monogamy”, described as “a social coalition with no implications for exclusive mating” (Whiteman & Côté 2004). While this definition suggests that these pairs are a ‘social coalition’ rather than a mating pair, environmental factors leading to the formation of social pairs and the possible ecological consequences beyond reproductive advantages or constraints are largely unexplored. This is particularly intriguing, as pair-formation appears to be a common social system in marine fishes, especially on tropical coral reefs (Randall et al. 1997; Allen et al. 2003; Froese & Pauly 2012).

Table 2.1: Pair-formation in animals. Pair-formation is a common behaviour from protists to mammals.

Kingdom	Phylum	Class	Order	Example	Source	
Protista	Ciliophora	Ciliatea	Peniculida	<i>Paramecium primaurelia</i>	Corrado et al. 1997	
		Oligohymenophorea	Hymenostomatida	<i>Tetrahymena pyriformis</i>	Bruns and Brussard 1973	
Animalia	Acanthocephala	Eoacanthocephala	Neoechynorhynchida	<i>Neoechynorhynchus pseudomydis</i>	Cable and Hopp 1954	
	Platyhelminthes	Trematoda	Strigeidida	<i>Schistosoma spp.</i>	Beltran and Boissier 2008	
	Annelida	Polychaeta	Phyllodocida	<i>Harmothoë imbricata</i>	Daly 1973	
	Arthropoda	Merostomata	Crustacea	Xiphosura	<i>Limulus polyphemus</i>	Botton and Loveland 1992
				Malacostraca	<i>Homarus americanus</i>	Atema et al. 1979
				Mecoptera	<i>Panorpa spp.</i>	Thornhill 1979
	Mollusca	Gastropoda	-	-	<i>Ovula ovum</i>	Kei 2010
	Chordata	Chondrichthyes	-	Carcharhiniformes	<i>Carcharhinus melanopterus</i>	Johnson and Nelson 1978
				Perciformes	<i>Amatitlana nigrofasciata</i>	Mackereth and Keenleyside 1992
		Amphibia	-	Anura	<i>Rana sylvatica</i>	Howard and Kluge 1985
Reptilia		-	Squamata	<i>Tiliqua rugosa</i>	Leu et al. 2011	
Aves		-	Passeriforma	<i>Taeniopygia guttata</i>	Adkins-Regan 2002	
Mammalia	-	Rodentia	<i>Galea monasteriensis</i>	Adrian et al. 2008		

The formation of long-term pairs in coral reef fishes has been a widely studied phenomenon. The majority of studies examining reef fish pairs are case studies that seek explanations as to why species that lack bi-parental care and exhibit potential for polygamous mating restrict themselves to a single reproductive partner (e.g., Fricke & Fricke 1977; Gore 1983; Fricke 1986; Barlow 1987; Donaldson 1989; Herold & Clark 1993; Kuwamura et al. 1993; Reavis 1997a,b; Reavis & Barlow 1998; Takegaki 2000; Harding et al. 2003; Whiteman & Côté 2003; Pratchett et al. 2006; Sogabe et al. 2007; Reavis & Copus 2011). The most commonly cited reasons for the evolution of monogamy are: 1) environmental factors that prevent the sequestration of multiple mates; 2) paternal egg tending leading to mutual mate guarding; and 3) sparse populations. In contrast, few studies examine possible ecological correlates of pairing as a social system (Robertson et al. 1979; Hourigan 1989; Pratchett et al. 2006; Brandl & Bellwood 2013a,b). These studies suggest increased feeding efficiency, benefits for territorial defence, and increased vigilance, as possible ecological benefits of pair-formation. However, no holistic approach to examine common ecological traits among pairing species has been made to date. Yet, such an approach may enable inferences about environmental determinants of pair-formation to be made and will provide more detailed insights into the functional ecology of pairing species. This review, therefore, aims to provide an overview of pairing in coral reef fishes following a social-ecological definition of the term pair-formation. To do so, 1) an ecologically orientated definition of the term 'pair-formation' is offered, and 2) a meta-analysis of pair-formation, ecological traits, and reproductive modes in coral reef fishes of the Indo-Pacific is conducted to identify common characteristics of pairing species. These characteristics are then discussed from an ecological perspective.

2.2 The definition of 'pair-formation'

Initially, the definition of the term 'pair-formation' appears to be simple, describing an association between two individuals. In its simplicity, this is only appropriate for molecular pair bonds. However, this definition fails when projected on animals, as every interaction between two individuals would be considered pair-formation. To identify the most common use of the term 'pair-formation' in the literature, a basic meta-analysis was performed, searching for studies with the term 'pair-formation' in the title. The search was specified by the word 'animals' in the topic to exclude a large body of literature about pair-formation in molecules. Using the ISI Web of Knowledge database, a total of 168 studies were found (from 1969 to 2012), with 67 being off-topic or duplicates. The remaining 101 studies were divided into four different categories: 1) articles, describing copulation or spawning and associated courtship or pre-mating behaviours, 2) articles, describing pair-formation beyond copulation or courtship, including post-mating behaviour or the function of long-term pairs, 3) articles about the conjugation of unicellular organisms, including all studies on unicellular eukaryotes and their form of 'mating', and 4) articles, describing pairs without any reproductive background.

Of the 101 studies, half (50.5%) were restricted to the process of copulation or spawning as well as courtship or pre-copulatory mate guarding. Another 33.7% described reproductive pairs, but examined post-copulatory behaviour or long-term pair-formation (where the pair bond is maintained over more than one reproductive season). A further 11.9% of articles described conjugation of unicellular organisms and just 4% of all 101 studies described cooperative 'pairs', without a clear reproductive purpose (**Figure 2.1**). Because more than half of the studies on pair-formation described only copulation or pre-copulatory behaviour, referring to a 'mating system' rather than a 'social system', a more precise definition of pair-formation is needed that incorporates both ecological and reproductive

components. Pair-formation must be distinguished from pure mating and pre-mating behaviour such as courtship or pre-copulatory mate guarding (Kortlandt 1995), as the latter only pave the way for copulation (e.g. Johnson & Nelson 1978; Hartnoll & Smith 1978; Burpee & Sakaluk 1993). As an example, in spawning aggregations of numerous families of reef fishes (e.g. Mullidae, Serranidae, Siganidae), actual spawning occurs between members of a pair, temporarily separating from the aggregation (Johannes 1981; Samoily & Squire 1994; Domeier & Colin 1997). Likewise, many other species (e.g. Acanthuridae) spawn in pairs but remain solitary throughout their lives within their own individual feeding territories (Robertson et al. 1979; Robertson 1984). These 'pairs' are therefore restricted to brief encounters that only serve to fertilize eggs and have no implications for daily survival.

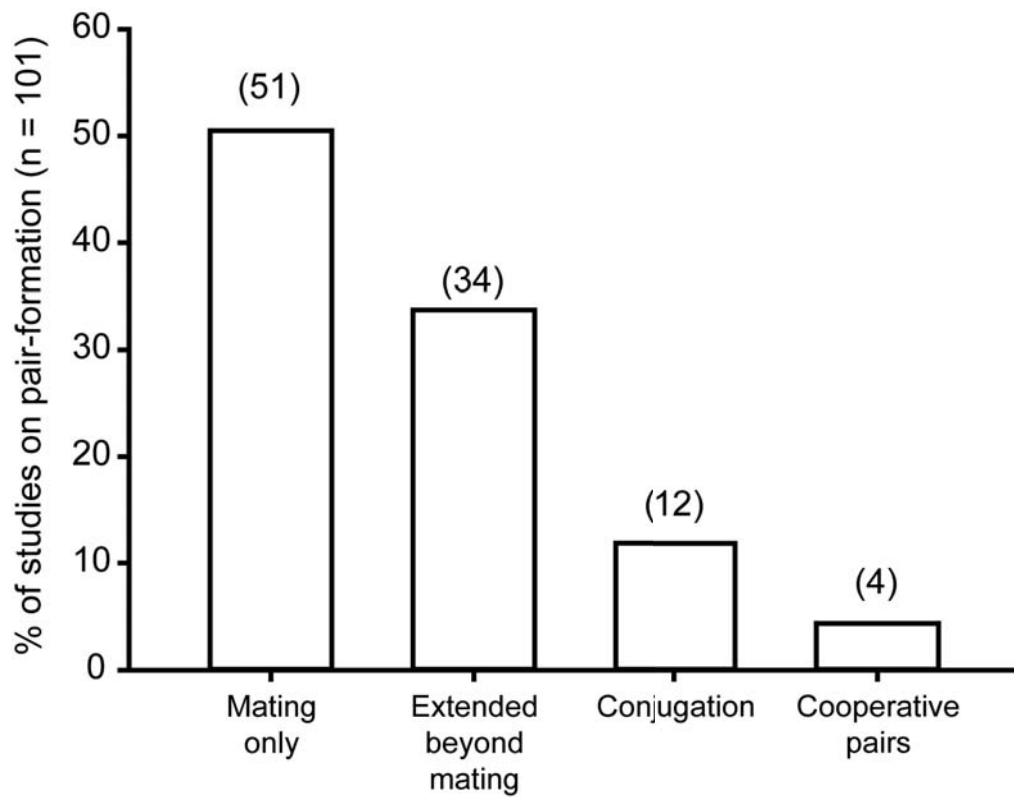


Figure 2.1: Proportions of studies investigating ‘pair-formation’ in animals. Papers are assigned to different categories, reflecting the use of the word pair-formation. Numbers are based on the ISI Web of Knowledge database in November 2011.

With few exceptions (see Ahlgren et al. 2011), there are no clear ecological implications of these transient pairing associations. Hence, if the formation of a pair is restricted to copulation only, the term ‘pair-formation’ may be misleading, as it does not describe a social system but simply the process of reproduction, i.e. a mating system. Therefore, I suggest that the term ‘pair-formation’ be limited to associations that are maintained beyond the process of reproduction, such as 1) cooperation in rearing offspring (e.g. Robertson 1973; Cox et al 1993; Adrian et al. 2008; Young et al. 2008), 2) mutual maintenance of a dwelling or territory (e.g. Linsenmaier and Linsenmaier 1971; Fricke 1986; Mathews 2002), 3) increased predator avoidance through shared vigilance (Gwinner et al. 1994; Pratchett et al. 2006) or 4) improved feeding efficiency (e.g. Robertson et al. 1979; Hourigan 1989; Gwinner et al. 1994). In contrast, pairing for copulation alone is perhaps more appropriately termed a ‘mating-pair’.

An apparent issue arises for pairs in which mate-guarding occurs after reproducing, as it initially appears to solely benefit the guarding individual in terms of its reproductive success. However, pair-formation in these species can be ecologically meaningful. For example, female lobsters (*Homarus americanus*) experience lower predation rates after moulting for reproduction (Atema et al. 1979) and female water-striders (*Gerris remigis*) exhibit higher foraging rates when guarded by males after copulation (Wilcox 1984). Both of these benefits are seen in the Australian sleepy lizard, *Tiliqua rugosa*, where the female can dedicate more time to foraging efforts and predation risk is reduced through increased vigilance of the male (Bull and Pamula 1998; Leu et al. 2011). Thus, in order to address ecological implications of associating with a single individual, I suggest that pair-formation be defined as ‘an association between two conspecific individuals, maintained beyond the time required for reproductive activity leading to fertilization’. With this definition of pair-

formation as a social system, one can begin to explore the role of this behaviour in coral reef fishes.

2.3 Pair-formation in coral reef fishes

To explore pair-formation in reef fishes in greater detail, the Indo-Pacific coral reef fish fauna was examined. The primary goal was to understand the basis of pair-formation in fishes. As a first step, I sought to identify ecological traits that are shared by pair-forming species. Often, ecological traits are correlated and, if evolutionary successful, they occur repeatedly among distantly related taxonomic groups (Westneat et al. 2005). For example, foraging in large aggregations appears to be a beneficial trait for species that feed on pelagic zooplankton. As such, aggregations frequently occur in distantly related taxa such as the Caesionidae, Labridae, Pomacentridae or Serranidae (e.g., Hamner et al. 1988; Hobson 1991). Similarly, cryptobenthic fishes are almost exclusively known to spawn adhesive, demersal eggs that are deposited in a cave, burrow or nesting site (e.g. Fishelson 1976; Thresher 1984; Hernaman & Munday 2007). I therefore reviewed a comprehensive set of traits among Indo-Pacific reef fishes (extracted from Randall et al. 1997; Allen et al. 2003; Randall 2005). The traits encompassed social systems, trophic affiliation, strategies to avoid predators and reproductive characteristics (**Appendix C**). Specifically, each species was classified as either pairing or non-pairing, assigned to a trophic category and classified with regards to other ecological characteristics (maximum size, maintenance of burrows, nocturnal or diurnal activity). In addition, reproductive characteristics such as the spawning mode (pair or group spawning), the nature of gamete release (broadcast, demersal, pouchbrooding, mouthbrooding, egg-scattering, gelatinous egg-mass spawning), and the mating system were recorded.

2.3.1 The prevalence of pairing behaviour in reef fishes

Of 1,981 species of coral reef fishes in 79 families, 341 species are reported to form pairs, while 1,561 species have no records of pair-formation. No information was available for 79 species, which are predominantly small, cryptbenthic species. An average of $18.7\% \pm 0.3$ (mean \pm SE) of species within a family are reported to form pairs, ranging from 100% in the Malacanthidae, Microdesmidae, Monocentridae, Pegasidae, Solenostomidae and Zanclidae to 0% in 42 families (**Figure 2.2**). To account for extreme proportions in families with low numbers of species, families with less than five species were not considered subsequently.

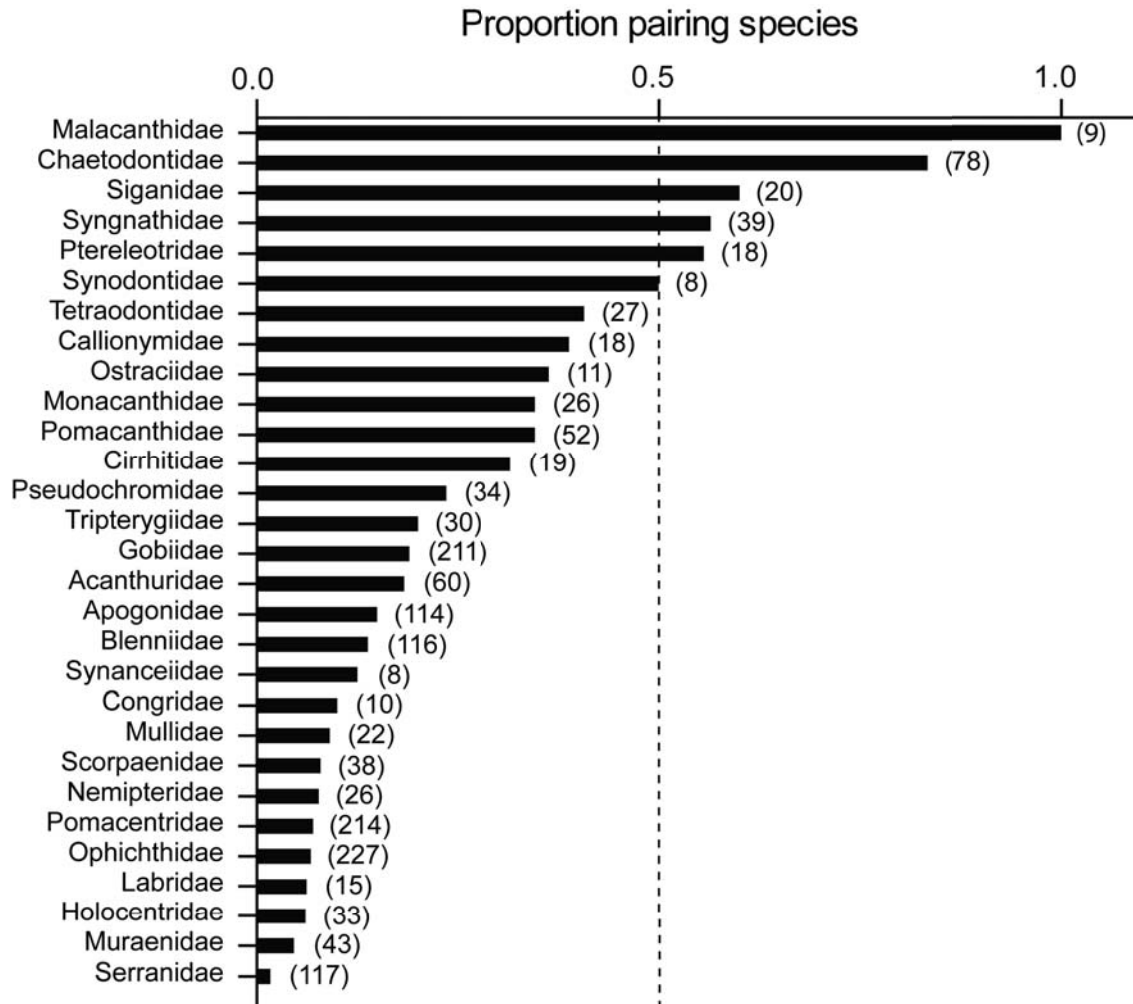


Figure 2.2: The prevalence of pair-formation in coral reef fish families. Only families with more than five species are considered. Dashed line marks the 50% mark. The number of species in each family is given in parentheses.

Five families of reef fishes contain more than 50% of the total pair-forming species. Belonging to two different orders (Perciformes and Syngnathiformes), these include tilefishes (Malacanthidae, 100% of the examined species in pairs), butterflyfishes (Chaetodontidae, 83.3% pairs), rabbitfishes (Siganidae, 60.0%), seahorses and pipefishes (Syngnathidae, 56.4%) and dartfishes (Ptereleotridae, 55.5%) (**Figure 2.3**). The two following families, lizardfishes (Synodontidae, 50.0%) and pufferfishes (Tetraodontidae, 40.7%), belong to different orders, the Aulopiformes and Tetraodontiformes, respectively. Given that the seven families with the highest proportions of pairing species belong to four different orders, the tendency to arrange in pairs appears to have evolved independently across several phylogenetically distinct lineages. Furthermore, there are major differences between families within the same order. For example: triggerfishes (Balistidae) and porcupinefishes (Diodontidae) do not comprise any pairing species (0%), whereas pair-formation is relatively common in pufferfishes (Tetraodontidae 40.7%), box- and cowfishes (Ostraciidae, 36.3%) and filefishes (Monacanthidae, 34.6%). All these families belong to a single order, the Tetraodontiformes. A similar situation occurs in the Syngnathiformes, where seahorses and pipefishes (56.4%) are in marked contrast to trumpet- (Aulostomidae, 0%) and cornetfishes (Fistulariidae, 0%). Overall, based on the most recently assembled phylogeny of fishes (Near et al. 2012), pairing appears to have arisen at least 13 times among reef fishes.



Figure 2.3: Examples of pair forming species in the Chaetodontidae, Siganidae, Pomacanthidae, Gobiidae, Synodontidae, Syngnathidae, Malacanthidae and Monacanthidae. Photographs by J.P. Krajewski, B. Halstead and S.J. Brandl.

Narrowing the taxonomic scale reveals that there is also significant variation within families. In some cases, entire lineages are pair-forming and there are clear distinctions between genera. In the Malacanthidae, for example, pairing is restricted to highly reef-associated, tropical Indo-Pacific species in the genera *Hoplolatilus* and *Malacanthus* (Clark & Pohle 1992; Clark et al. 1998); temperate and Atlantic species of the genera *Branchiostegus*, *Caulolatilus*, *Lopholatilus* and *Malacanthus* are solitary or live in colonies (Able et al. 1982; Ross 1982; Baird & Baird 1992; Mitamura et al. 2005). In the Tetraodontidae, pairing is restricted to the genus *Canthigaster* (Kobayashi 1986; Sikkell 1990), the smallest genus in the family. In other families, the prevalence of pairing behaviour varies within genera. For example, there is a clear dichotomy in *Siganus*, the single genus of the Siganidae, with pair-formation being reported only for reef-associated, colourful tropical species, while drab-coloured, mangrove-associated, estuarine and subtropical species form schools (Woodland 1990; Borsa et al. 2007; Brandl & Bellwood 2013b). Likewise, in the Chaetodontidae, there are pairing and non-pairing species in the dominant genus *Chaetodon*, regardless of their phylogenetic relationships (Hourigan 1989; Roberts & Ormond 1992; Kelley et al. 2013). In the Syngnathidae, both pipefishes (genera *Corythoichthys*, *Dunckerocampus*, *Doryrhamphus*) and sea horses (genus *Hippocampus*) comprise several species that form pairs, while others live solitary or in groups (Allen et al. 2003; Foster & Vincent 2004; Sogabe & Yanagisawa 2008). Given the occurrence of pairing behaviour in a diverse array of orders and families and the high variation within closely related taxa, it appears that pair-formation has arisen repeatedly over many millions of years. With such diverse groups involved, there is ample opportunity for a critical evaluation of the ecological and reproductive role of pair-formation in coral reef fishes.

In ecology, the three main drivers of social behaviour are associated with feeding, predation avoidance and reproduction. To identify ecological and reproductive traits that are

associated with pair-formation in reef fishes, I will explore each of these components separately.

2.3.2 Trophic ecology of pair forming fishes

The acquisition of food is a crucial process in the life history of animals and their prey. Consequently, species are often classified within certain trophic groups (e.g. Williams & Hatcher 1983; Green & Bellwood 2009; Cheal et al. 2012), where species with similar foraging strategies are grouped based on their major prey items. Interestingly, there is a clear pattern with regards to the prevalence of pairing species within trophic groups (**Figure 2.4**). In three trophic groups, spongivores, corallivores and micro-invertevores, more than half of the species form pairs. This is a remarkable proportion considering the relatively small number of pairing species (341) compared to non-pairing species (1,640, i.e. 17.2% pairing species). Two additional groups, omnivores and herbivores, also had relatively high proportions of pairing species (> 25%), while the proportion of pairing species is low in planktivores, macro-invertevores, piscivores, carnivores and detritivores.

The major pattern that emerges from these results is the link between small and / or immobile, sedentary prey items and pairing behaviour (**Figure 2.4**), as all trophic groups that forage on such prey have high proportions of pairing species. Their prey includes coral (polyps, mucus), micro-invertebrates (e.g., harpacticoid copepods, amphipods, small polychaetes, molluscs), filamentous or fleshy algae, sponges, or all of the above (omnivores). All these prey items require relatively high visual acuity and are located in topographically complex microhabitats on the coral reef benthos. Consequently, the majority of pairing species are also reported to be diurnal (96.4%) and are, on average, smaller than non-pairing species (mean maximum total length \pm SE: $162.6 \pm 6.4\text{mm}$ vs. $278.0 \pm 8.6\text{mm}$, respectively). Likewise, in 19 of 29 families that contain both pairing and non-pairing species, pairing

species were on average smaller than their non-pairing counterparts, suggesting that size differences are not solely due to phylogenetic effects.

The relationship between pair-formation and selected trophic groups may be due to the relatively immobile nature and distribution of prey. Although benthic micro-invertebrates are sometimes motile, their movements are restricted to small scales, making these organisms a largely stationary food source on a whole-reef scale (Klumpp et al. 1988; Kramer et al. 2012). Widely distributed prey has been associated with the evolution of monogamous mating systems in animals, as such prey prevents the monopolization of multiple mates (Emlen & Oring 1977; Whiteman & Côté 2004; Reavis & Copus 2011). Such a relationship has been proposed for butterflyfishes, distinguishing between mobile, planktivorous schooling species and benthos-associated, corallivorous species that occur predominantly in pairs (Reese 1975; Fricke 1986), leading to the assumption that environmental factors, food availability in particular, can predict the social system in this family (Hourigan 1989). Several studies argue that evenly distributed, stationary and predictable food sources promote pairing behaviour in butterflyfishes, as males are not able to sequester more than one female (e.g., Hourigan 1989; Reavis & Copus 2011). This is in accordance with theoretical expectations based on the evolution of monogamy (e.g. Emlen & Oring 1977) and is consistent with the high proportion of corallivores that occur in pairs (**Figure 2.4**).

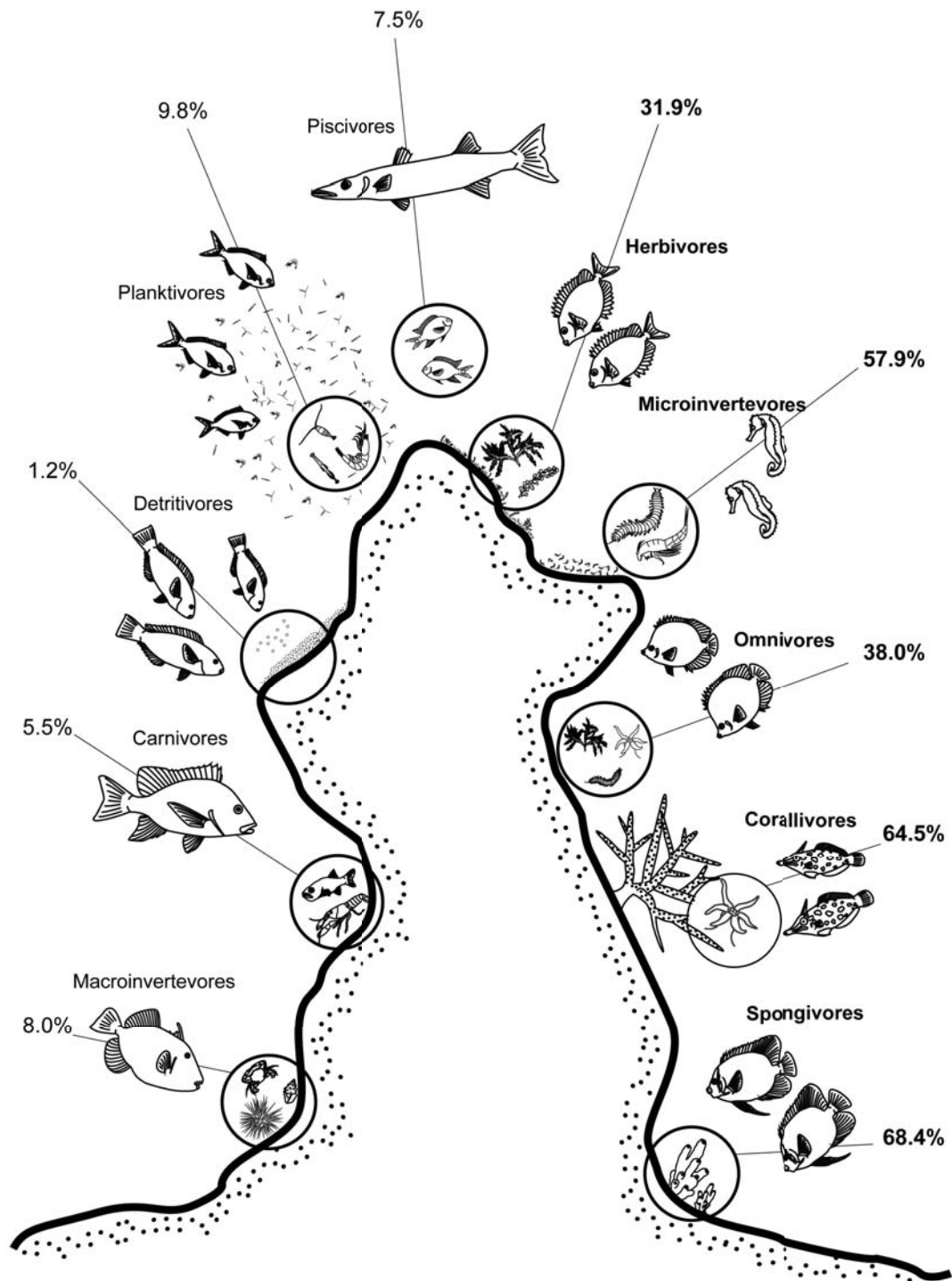


Figure 2.4: The prevalence of pair-formation in major trophic groups of coral reef fishes. Percentages indicate the proportion of pair-forming species within each

trophic group. Groups with more than 25% pairing species are marked are in bold.

Black circles indicate prey items.

However, there are other factors that may also favour the formation of pairs in these trophic groups. One key aspect may be the size and benthic nature of prey items. With the exception of some large species of algae or sponges, the majority of organisms preyed on by pair-rich trophic groups are relatively small (e.g., coral polyps, filamentous algae, harpacticoid copepods) and are located in a micro-topographically highly complex and heterogeneous environment (Mundy 2000). Therefore, species foraging on these items may require highly dexterous movements and fine-scale interactions with the substratum to obtain their prey. This has been documented in several pair-forming groups such as butterflyfishes or rabbitfishes (Motta 1988; Ferry-Graham et al. 2001; Fox & Bellwood 2013) and other studies have highlighted the ability of some species to exploit complex microhabitats (Hobson 1975; Robertson & Gaines 1986; Motta 1988; Montgomery et al. 1989; Ferry-Graham et al. 2001; Fox & Bellwood 2013). Interestingly, many of the latter species are known to form pairs. This may be due to rapid decreases in the feeding efficiency of larger groups (White & Warner 2007a). A loss of foraging efficiency may be exacerbated by increases in the inaccessibility of prey or increased handling time of small, benthic prey (Pratchett et al. 2006; Fox & Bellwood 2013; Brandl & Bellwood 2013a,b). In such circumstances, one would anticipate selection against large groups and more frequent pair-formation (Ford & Swearer 2013b).

Another specialized trophic mode that appears to be related to pair-formation is the maintenance of cleaning stations. Several species of reef fishes clean clients and perform their cleaning tasks as cooperative pairs (Bshary et al. 2008). Besides the most widely known species in the labrid genus *Labroides*, which commonly live in harems or pairs (Allen et al. 2003), there are species within the Gobiidae (e.g. *Elacatinus evelynae*) and Syngnathidae (e.g. *Dunckerocampus pessuliferus*), which are commonly found in pairs and maintain cleaning stations (Harding et al. 2002; Whiteman & Côté 2003; Allen et al. 2003). In most of

these species, it appears that the cleaning service provided by a pair is superior to cleaning by solitary individuals (Whiteman & Côté 2003; Bshary et al. 2008). This benefits both individuals in a simple way, as clients are more likely to visit high quality cleaner stations (Whiteman & Côté 2003). While this has well known ramifications for the mating system and the reproductive success (Bshary et al. 2008), it also provides a good example of cooperative, social pairing that ecologically benefits both pair members.

Overall, although there may be benefits in pairing, it is not clear why all these species occur as pairs rather than in small groups. Perhaps there are other factors operating that limit social group sizes such as the suggestion that larger groups and shoaling behaviour may enhance the overall predation risk from different predator guilds (Ford & Swearer 2013a,b).

2.3.3 Predation avoidance in pair forming fishes

Mortality due to predation is another important force shaping species' behaviour, and is tightly linked to an animal's ecology (Holbrook & Schmitt 2004; Almany & Webster 2006; Holmes & McCormick 2011). Predation risk can influence behaviour in different ways, restricting the movement of prey fishes to small areas around shelter sites (Reavis 1997b; Welsh & Bellwood 2012a; Madin et al. 2012), away from unstable, complex habitats (Hoey & Bellwood 2011), or drive prey fishes to form large aggregations to avoid predation on the individual (Pitcher & Parrish 1993; White & Warner 2007b). Thus, the threat of predation, in synergy with the need to forage and reproduce, can either decrease (e.g., restriction of movements; Welsh & Bellwood 2012a) or increase (e.g., formation of large schools; Welsh & Bellwood 2012b) the size and movement of social groups. This raises the question: are there any circumstances, where a pair may be the preferred group size for avoiding predation?

Pair-formation in reef fishes has rarely been linked to anti-predatory strategies. However, several lines of evidence suggest that it may play a role. For mobile species, the individual risk of predation is likely to increase with decreasing group size (but see Ford & Swearer 2013a,b). In large groups, vigilance is shared between numerous individuals, providing security for each group member (Pitcher & Parrish 1986; White & Warner 2007b). Thus, in theory, in terms of overall predation risk, individuals in pairs should be more vulnerable than those in schools but less vulnerable than solitary individuals (Pratchett et al. 2006). One way that pairing fishes may compensate for the higher risk is with morphological and / or behavioural adaptations. It is striking that virtually all families of mobile (non-burrowing) reef fishes with high proportions of pairing species exhibit conspicuous physical adaptations to avoid predation. This includes large venomous spines (Siganidae, Pomacanthidae), exceptionally deep bodies and bright colouration (Chaetodontidae, Pomacanthidae) or body inflation, extremely tough skin or bony plates encasing the body and/or toxicity (Tetraodontidae, Ostraciidae, Monacanthidae) (Hixon 1991). Although some of these features are also found in solitary or schooling species (e.g., Acanthuridae, Balistidae), families with few pairing species often appear to have fewer morphological adaptations to avoid predation (e.g., Mullidae, Nemipteridae, Labridae, Caesionidae). In these predominantly schooling species, speed appears to be the major determinant of individual survival, as an individual often only needs to be faster than a single adjacent individual to avoid predation. Within families, there appears to be a similar trend. For example, within the Chaetodontidae, Acanthuridae and Siganidae, pairing species possess deeper bodies (Brandl & Bellwood 2013b) and may exhibit brighter coloration and more anti-predatory features, such as eye stripes, than non-pairing species (Kelley et al. 2013).

Behavioural adaptations offer an additional means of decreasing vulnerability to predation in pairs. Most mobile, pairing species are relatively slow swimmers (Fulton 2007),

which makes predator avoidance through high-speed escape rather unlikely, although a deep body (Brandl & Bellwood 2013b) may make rapid direction changes a possible predator avoidance mechanism. The most commonly cited predation avoiding benefit of pair-formation in mobile reef fishes is to spawn with the respective partner. The permanent availability of a reproductive partner releases these species from the necessity to undertake dangerous, predation-prone ventures to spawning sites in search of a mate (Robertson et al. 1979; Herold & Clark 1993). However, many pairing species still migrate to spawning sites (e.g., *Siganus punctatus*, Johannes 1981; *Chaetodon lunulatus*, Yabuta 1997).

It has been suggested that the close association between pairing butterflyfishes and rabbitfishes and the reef matrix (i.e. swimming within the complex interstices of the reef) reduces vulnerability to predation (Hourigan 1989; Borsa et al. 2007). Although the mechanistic basis for this was not explored, recent observations provide an indication of the possible link between pairing, reef complexity and predation avoidance. A recent study of rabbitfishes raised the possibility of shared vigilance between pair-members while feeding, with one individual being observed ‘hanging’ tail-down vertically in the water column, scanning the environment (Fox & Bellwood 2013; Brandl & Bellwood 2013a) (**Figure 2.5a**). Interestingly, this exact same posture and behaviour have been described for pairing tilefishes, with one member of a pair hanging tail-down in the water column while the other one engages in other activities and escaping immediately prior to the arrival of a predator (Clark et al. 1998). The close resemblance of the shared vigilance behaviour observed in rabbitfishes and tilefishes suggests that it may be a common predator avoidance behaviour in pairing species. A similar type of shared vigilance (although lacking the tail-down posture), has been described in the gobies *Valenciennea helsdingenii* and *Valenciennea longipinnis*, in which pair members seem to alternate between vigilance and foraging (Takegaki & Nakazono 1999; Clark et al. 2000). This has also been suggested for the pairing butterflyfish

Chaetodon melannotus (Pratchett et al. 2006). In addition, several species of pair-forming rabbitfishes (*Siganus puellus*, *S. doliatus*, *S. punctatus*) have been observed swimming in a synchronized fashion, aligning their bodies in a manner that creates the illusion of one, significantly larger fish (also *S. stellatus*; Choat, pers. comm.) (**Figure 2.5b**). This is likely to decrease the risk of predation by smaller predators but needs to be explored in further detail. Thus, it appears that pairing reef fishes may escape predation through a range of behavioural responses that are tailored to their pairing social system.

Further support for a non-reproductive role for pair-formation is provided by the presence of homosexual and immature pairs in pairing reef fishes. Homosexual pairs have been found in the butterflyfishes *Chaetodon capistratus* (Gore 1983), *C. chrysurus* (Fricke 1986), *C. lunulatus* and *C. melannotus* (Pratchett et al. 2006), the surgeonfish *A. triostegus* (Robertson et al. 1979), the gobies *Valenciennesa muralis* and *V. strigata* (Pratchett et al. 2006), and the rabbitfish *Siganus doliatus*, where 25% of pairs were homosexual (Brandl & Bellwood 2013a). This is also consistent with reports of non-reproductive, mixed-species pairs. Mixed species pairs have been reported in the blennies *Petroscirtes fallax* and *Meiacanthus lineatus* (Allen et al. 2003) and the tilefishes *Hoplolatilus cuniculus*, *H. chlupatyi*, *H. marcosi* and *H. purpureus* (Clark et al. 1998). All these examples question a purely reproductive function of pair-formation in the respective species and suggest a crucial role of pairing for daily survival. Mortality rates of paired vs. non-paired individuals have yet to be reported for any of these species but predation risk may be the most important driver of this behaviour (Reavis & Barlow 1998).

While anti-predatory behaviour in mobile pairing species is poorly understood, the relationship is much clearer in bottom-dwelling or burrowing species (Hixon 1991; Forrester & Steele 2004; Depczynski & Bellwood 2004; Hernaman & Munday 2005). Of the 102 species that inhabit and maintain permanent burrows, 50 commonly occur in pairs, with many

of them being strongly paired and / or monogamous (Reavis 1997a,b; Reavis & Barlow 1998; Clark et al. 1998; Takegaki & Nakazono 1999; Clark et al. 2000; Pratchett et al. 2006; Hernaman & Munday 2007). While almost half of the burrowing species form pairs, it is noteworthy that 47 out of the remaining 52 species live in close association with shrimps of the family Alpheidae. In these latter, shrimp-associated species, there is a clear division of labour between the shrimp and the goby, where the shrimp is responsible for digging and maintaining the burrow (**Figure 2.5c**). Meanwhile, the goby acts as a sentinel, dedicating extensive time to vigilance and warning the shrimp through rapid tail flicks if a predator is approaching (Karplus 1987; Thompson 2004, 2005). This relationship is mutualistic and obligate. Solitary gobies experience rapid mortality through predation, and solitary shrimps exhibit slower growth rates due to decreasing foraging time (Karplus et al. 1972; Thompson 2004; 2005). While this relationship should be considered a symbiotic relationship rather than pair-formation, it demonstrates a key aspect of the ecology of burrowing species: the maintenance of a permanent burrow is greatly facilitated by the presence of a second individual. Overall, 95.1% of all fish species that maintain permanent burrows do so with a partner, be it fish or shrimp. Division of labour between pair members of burrowing fishes is well documented (Reavis and Barlow 1998; Clark et al. 1998; Takegaki & Nakazono 1999; Clark et al. 2000) and mainly involves the partitioning of circadian tasks (burrow maintenance, vigilance, foraging) (**Figure 2.5d**). Thus, for these species, the pair bond may be the essential feature for daily survival, suggesting a strong ecological basis for the establishment of pairs. This is further reinforced by the presence of non-reproductive pairs in burrowing gobies (Pratchett et al. 2006) and tilefishes (Clark et al. 1998). It is also seen in other organisms that maintain burrows as pairs, including shrimps of the genus *Lysiosquilla* (Christy & Salmon 1991).

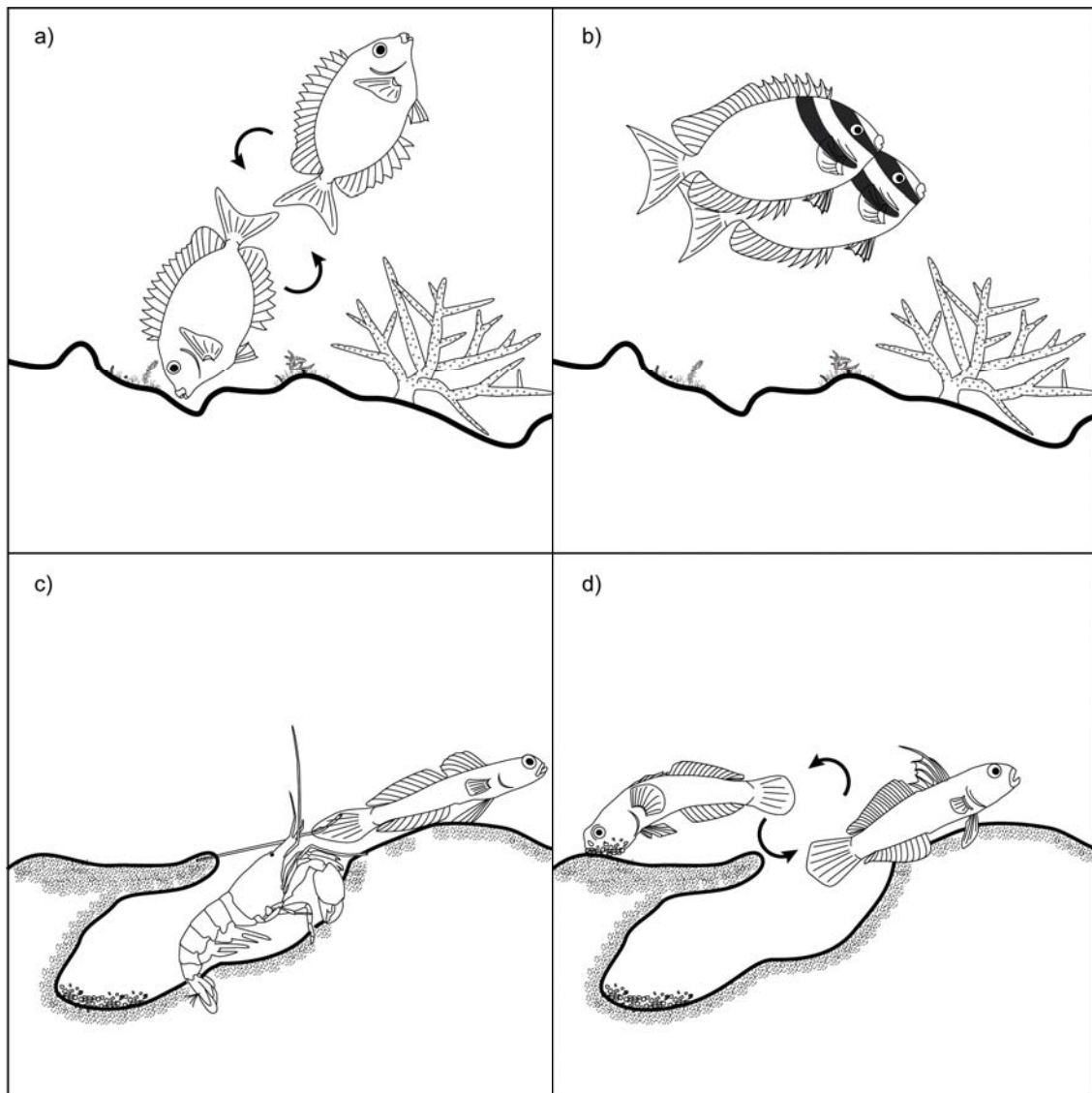


Figure 2.5: Observed behavioural adaptations of pair-forming species in mobile (a & b) and burrowing (c & d) species; a) shared vigilance in rabbitfishes. One individual is assuming a ‘tail-down’ position, scanning the environment, while the other individual is feeding; b) synchronized swimming in rabbitfishes. Individuals arrange in a manner that creates the illusion of a single, larger fish; c) shared labour in shrimp-associated, burrowing gobies. The fish is vigilant while the shrimp is performing maintenance of the burrow, maintaining contact with the goby through its antennae;

d) shared labour in burrowing gobies. One individual is remaining close to the burrow entrance, looking out for predators while the other individual is feeding further away from the burrow.

Thus, in summary, predation appears to be a significant factor that may influence the formation of pairs in numerous species and there appears to be a clear link between pairing and predation in species that maintain a permanent burrow. Although several behavioural traits also suggest anti-predatory benefits of pairing in free-living mobile species, the role of pairing behaviour for predator avoidance in these species is poorly understood.

2.3.4 Reproductive characteristics of pair forming fishes

Coral reef fishes exhibit virtually every form of mating system known in animals, from monogamous mating to mass spawning aggregations, where up to 100,000 individuals of a single species spawn simultaneously (Robertson 1983; Colin & Bell 1991; Sadovy de Mitcheson et al. 2008). Spawning modes include broadcast spawning, demersal clutch spawning, egg-scattering, pouchbrooding, mouthbrooding, the release of gelatinous floating egg rafts and even live bearing (Thresher 1984). If pairing behaviour were solely due to reproduction, one would expect to find several unifying reproductive traits among pairing species, resulting in the following three hypotheses: first, prolonged association with a single partner should result in a mating system restricted to pair members (i.e., pair-spawning), which, second, should lead to high proportion of monogamy in pairing species. Third, given the close link between parental care and pairing (Barlow 1981) and the advantages of aggregations for broadcast spawning (Thresher 1984; Sadovy de Mitcheson & Colin 2012), pairing species should predominantly spawn benthic, demersal eggs with a high incidence of prerequisites for parental investment.

Initially, it appears that these hypotheses are supported. Among pairing reef fishes, 284 out of 341 (83.3%) are reported to have a mating system based on pairs, i.e. courtship and fertilization occurs between two individuals, while the remaining 57 species spawn in groups or aggregations. However, the majority of non-pairing species (69.1%) also mate in

pairs. Pairing species that are reported to spawn in aggregations rather than pairs are predominantly rabbitfishes, some surgeonfishes and few butterflyfishes. In surgeonfishes, this may be the result of relatively high plasticity in social and mating systems, observed throughout the family (Robertson et al. 1979; Robertson 1983). In butterflyfishes, pair spawning has been observed in *Chaetodon nippon* (Suzuki et al. 1980), *C. multivittatus* (Lobel 1989), *C. chrysurus* (Fricke 1986), *C. citrinellus*, *C. unimaculatus*, *C. ornatissimus* (Sancho et al. 2000), *Chaetodon lunulatus* (Yabuta 1997) and *Chaetodon rainfordi* (Thresher 1984), providing relatively good support for the hypothesis that permanently paired species should have a mating system based on pairs (Emlen and Oring 1977; Whiteman and Côté 2004). In contrast, anecdotal reports of spawning aggregations in several strongly paired species such as *C. ephippium*, *C. lunula* and *C. melannotus*, appear to be incongruous (Claydon 2004; Yabuta 2007). However, although these species may aggregate, they may ultimately spawn in pairs with numerous other pairs, utilizing a common location with favourable currents for egg dispersal (Bell & Colin 1986; Hixon 1991). In addition, the total fertilization rate of such pairs might be even higher in an aggregation (Petersen et al. 2001), particularly as interference in pair matings appears to be common in butterflyfishes (Suzuki et al. 1980; Neudecker & Lobel 1982; Lobel 1989).

Reliable records for rabbitfish spawning are not yet available (Woodland 1990). However, the few reports of spawning in rabbitfishes suggest that large spawning aggregations may be the common way of mating in this family (Johannes 1981; Hara et al. 1986; Domeier & Colin 1997; Hoque et al. 1999; Harahap et al. 2001; Sadovy de Mitcheson & Colin 2012). This is particularly surprising, because rabbitfishes are one of the most commonly and strongly paired families (Woodland 1990; Borsa et al. 2007; Brandl & Bellwood 2013a,b). As for butterflyfishes, it may be possible that siganid pairs also spawn in aggregations (Johannes 1981; Woodland 1990). However, there are no benefits in terms of

egg dispersal, because rabbitfishes spawn negatively buoyant demersal eggs (Woodland 1990). Thus, given the reports of homosexual pairs (Brandl & Bellwood 2013a) and the suggested anti-predatory and feeding strategies of pairing rabbitfishes, reproductive factors may only play a partial role for pairing in this family.

Overall, most pairing species also reproduce in pairs. However, any links between reproduction and pairing may need to be interpreted with caution, given the few noteworthy exceptions as well as the general tendency of reef fishes to reproduce in pairs (69.1% of non-pairing species). Reproduction is probably an important factor of pairing in many species but it appears to be only one of a range of potential drivers.

The second hypothesis suggests that pairing leads to monogamy. However, only 25.2% of all species that are known to form pairs are monogamous. True genetic monogamy has rarely been reported in reef fishes (Barlow 1981; Whiteman & Côté 2004), although many studies infer exclusive mating with a single partner from field observations, replacement experiments or aquarium studies (e.g., Barlow 1987; Herold & Clark 1993; Hess 1993; Reavis & Barlow 1998; Whiteman & Côté 2003; Reavis & Copus 2011). Interestingly, some species, such as jawfishes of the genus *Opisthognathus* (Hess 1993) or the anemonefishes *Amphiprion clarkii*, *A. frenatus* and *A. perideraion* (Hirose 1995), appear to be monogamous but live either solitarily (jawfishes) or in groups (anemonefishes). These species are not pair forming *per se* and represent 15.2% of all monogamous species. Therefore, while monogamy appears to be a relatively good indicator of pair-formation, it is not an obligate requirement. In turn, pairing species are not necessarily monogamous, as the majority (74.8%) of pair forming species are not reported to be monogamous.

The third hypothesis predicts that pairing species should predominantly spawn demersal eggs, while spawning in aggregations should be the predominant spawning mode in non-pairing fishes. Guarding or breeding eggs requires high investments by the parents, and

is usually rewarded by a stronger control of parentage and higher survival rates in offspring (Jones & Avise 1997). This high investment is regarded as a characteristic of pair mating, leading to the evolution of monogamy (Wittenberger & Tilson 1980). Thus, if pair-formation were driven by reproduction alone, one would expect all species with parental care to form pairs. Demersal clutch spawners have a high potential for parental care, as eggs are usually attached to the substratum within a restricted territory (e.g. Balistidae; Kawase 2002), cave (e.g. Blenniidae; Fishelson 1976) or burrow (e.g. Gobiidae; Reavis 1997a,b) and require intensive parental care (e.g. Hernaman & Munday 2007). Likewise, the specialized systems of mouthbrooding and pouchbrooding require high parental investment (e.g. Barlow 1981) and could therefore be assumed to have a high potential for pair-formation, while egg-scattering and broadcast spawning species appear to have a low capacity for pair-formation (Johannes 1981; Thresher 1984).

Surprisingly, the prevalence of pairing in the different spawning modes did not support the third hypothesis. Mouthbrooding had the lowest proportion of pairing species, followed by broadcast spawning and demersal clutch spawning. Pairing prevalence was highest in egg-scattering and pouchbrooding species (**Figure 2.6**). Mouthbrooding is largely restricted to cardinalfishes (f. Apogonidae) and few other lineages such as jawfishes (f. Opisthognathidae; Hess 1993). Most apogonid species live in aggregations, only forming ‘transient’ breeding pairs (Kuwamura 1985). In aggregating species (e.g. *Apogon notatus*, Allen et al. 2003), the females frequently desert the males after spawning, resulting in a solely courting and copulating pair, and hence a ‘mating pair’ (Kuwamura 1985). While in some pair-forming species, the female may engage in the defence of the eggs after spawning (e.g. *A. doederleini*, Kuwamura 1985), thus providing protection for the male, most pairs in cardinalfishes are probably ‘mating pairs’. Similarly, in jawfishes, individuals are solitary and only pair for courtship and spawning, after which the solitary male orally incubates the

eggs (Hess 1993). This is contrary to the suggestion that high parental investment will favour pair-formation. Similarly, the almost equal proportions of pairing and non-pairing species in broadcast and demersal spawning species, and the high prevalence of pairing in egg-scattering species, suggests that pairing is not significantly linked with parental investment. The two most commonly pairing families, tilefishes and butterflyfishes, are both broadcast spawn pelagic eggs (Thresher 1984; Clark et al. 1988), while egg-scattering is the most widespread spawning mode in rabbitfishes, the family with the third highest prevalence of pairing (Thresher 1984; Woodland 1990).

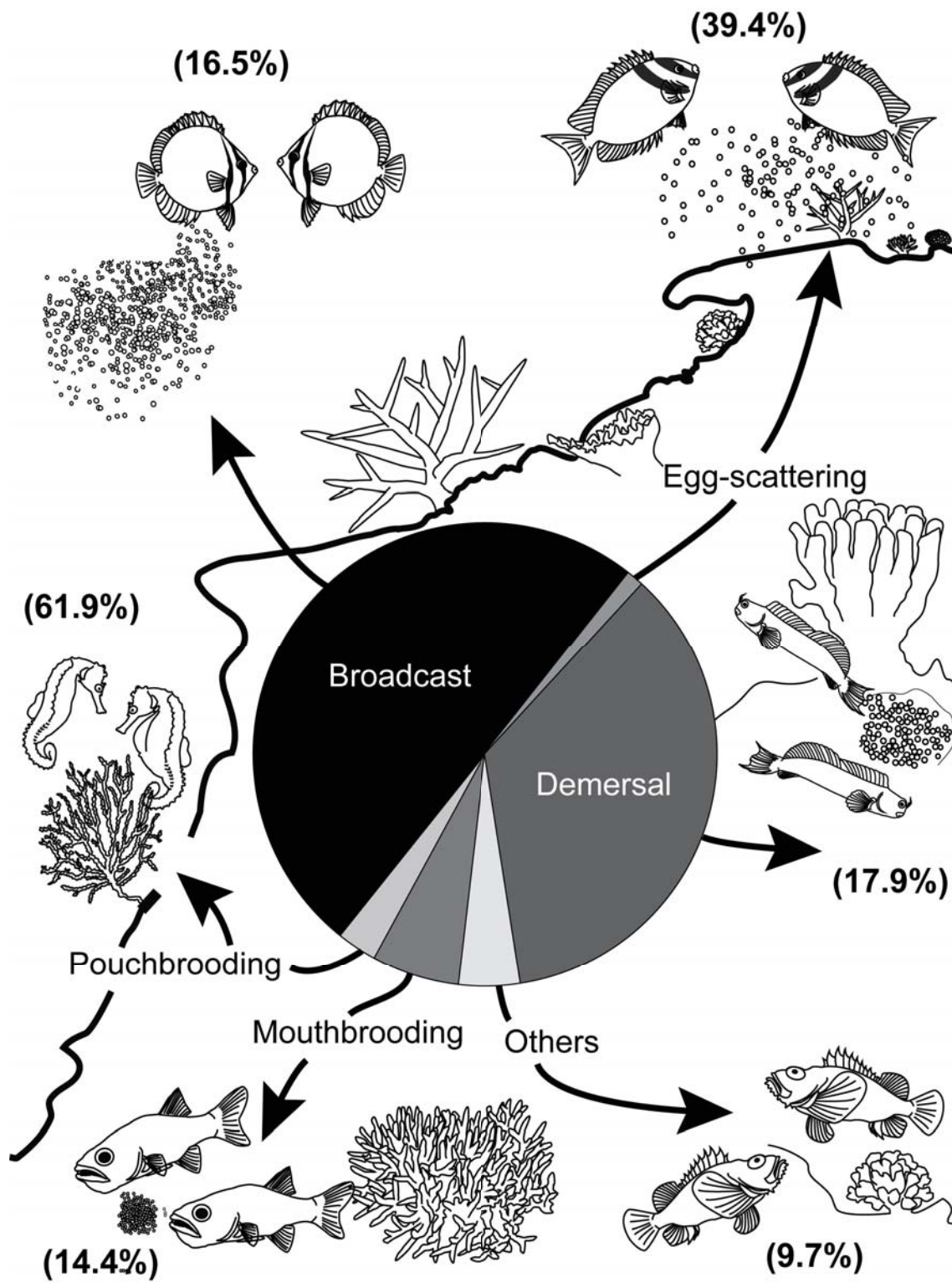


Figure 2.6: The prevalence of pair-formation in different spawning modes.

Proportions indicate the percentage of pair-forming species within each spawning

mode. The pie chart indicates the contribution of each spawning mode to the overall species pool.

The only spawning mode that appears to support the hypothesis that high parental investment should lead to pairing and, subsequently, monogamy, is pouchbrooding, a specialized form of breeding in the Syngnathidae, where the male incubates egg clutches in a pouch located on its ventral surface (Vincent & Sadler 1995). Many syngnathids form pairs (Allen et al. 2003) and there is substantial evidence that numerous species are monogamous, maintaining their pair bond beyond the reproductive season (Vincent & Salder 1995; Jones et al. 1998; Kvarnemo et al. 2000; Sogabe & Yanagisawa 2008). However, overall, with the exception of the Syngnathidae, there is limited evidence in support of a direct correlation between pair-formation and the spawning mode of reef fishes.

2.4 Conclusion and future directions

The formation of pairs is a common social system for animals. Yet, the definition of pair-formation as a social system is ambiguous and often confused with reproduction between only two individuals, which may be more appropriately termed a 'mating pair'. I therefore suggest restricting the term 'pair-formation' to 'an association between two conspecific individuals, maintained beyond the time required for reproductive activity leading to fertilization'. Following this definition, pair-formation is identified as a common trait among coral reef fishes, occurring across a wide range of phylogenetic lineages. However, the processes that have led to the evolution of this social system are poorly understood. This is particularly true for ecological factors that may be correlated with pairing behaviour. Most research to date has focused on the evolution of monogamous mating (i.e. a specialized mating system) in pairing species rather than the implications of pair-formation for daily survival, including food acquisition or reducing predation risk. Furthermore, the majority of studies on pair-formation in reef fishes focus on three families, the Chaetodontidae, Gobiidae

and Syngnathidae, while other families with a high proportion of pairing species such as the Malacanthidae, Siganidae, Ptereleotridae or Synodontidae are rarely investigated.

This review revealed several similarities in the ecology of pairing reef fishes, identifying two major traits. These appear to be common among pair forming species, regardless of phylogenetic relationships. First, pair-formation appears to be beneficial for species that forage on small, benthic, and relatively immobile prey, including coral polyps, sponges, filamentous algae or micro-invertebrates. This is consistent with the theoretical framework for the evolution of monogamous mating, explaining monogamy as a result of widely distributed resources and the inability of males to sequester multiple females. On reefs, the size and benthic nature of these resources may also influence the feeding efficiency of species that forage on such prey. As a consequence, associations with only one individual may be the preferred group size in these species. Second, the maintenance of permanent burrows appears to be almost exclusively restricted to species that live in association with a partner. While not all of these species can be considered to be pairing *per se*, these findings highlight the apparent necessity of a cooperative partner for the maintenance of permanent burrows. In both cases, the allocation of tasks between pair members appears to be important to avoid predation and maximize daily energy intake.

In contrast, there were few aspects of the reproductive biology of reef fishes that helped explain pairing species. Three hypotheses, based on the assumptions that pairing species should exhibit similarities in their reproductive behaviour, were not supported. While most pairing species also reproduce in pairs, there are some notable exceptions to this pattern and pair mating appears to be a common mating system among coral reef fishes in general. Thus, there is no clear link between pair-formation and reproduction in pairs. Monogamous mating, while being a good indicator for pair-formation, is reported in only a quarter of all pair-forming fishes. Finally, although parental investment appears to favour pairing in

pouchbrooding species, there was no clear correlation between these two traits across a broad range of families.

Based on the finding of common ecological traits among pairing species and the unexpectedly weak correlations between pair-formation and reproductive traits, I suggest that pairing behaviour in reef fishes may be strongly linked to ecological factors, benefitting daily survival and food acquisition. However, as opposed to well-defined theories and numerous empirical studies investigating the reproductive biology of pairing species, the environmental circumstances and ecological benefits of pairing species remain poorly understood. Given the potential importance of these ecological aspects, this promises to be an interesting and exciting avenue for future research.

Chapter 3: Morphology, sociality, and ecology: can morphology predict pairing behaviour in coral reef fishes?

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3.1 Introduction

Few groups of animals exhibit the morphological diversity seen in coral reef fishes. To understand the ecological importance of different morphologies, many reef fish studies have used an ecomorphological framework, which seeks to predict an organism's performance of ecological tasks using morphological attributes (Motta 1988; Wainwright 1991; Dumay et al. 2004; Mouillot et al. 2007; Goatley & Bellwood 2009; Ferry-Graham & Konow 2010).

Morphological attributes involved in prey acquisition have attracted much attention among ichthyologists and numerous reliable morphological predictors of foraging ecology have been identified. This includes dentition patterns (Motta 1989), jaw osteology or myology (Wainwright 1988; Alfaro et al 2001; Ferry-Graham et al. 2001a), bite kinematics (Ferry-Graham et al. 2002a; Konow & Bellwood 2005; Ferry-Graham & Konow 2010) and other associated functional complexes such as the visual sensory system (Job & Bellwood 1996; Goatley & Bellwood 2009; Schmitz & Wainwright 2011a) or the locomotory system (Bellwood & Wainwright 2001; Fulton & Bellwood 2002; Collar et al. 2008).

However, while morphological features can serve as proxies for the fundamental ecological niche (usually based on limits to mechanical performance; Hutchinson 1957), the realized ecological niche of a species or individual is determined by a variety of other biotic and abiotic factors (Hutchinson 1959; Ferry-Graham et al. 2002b), which restrict the realized niche that an organism can occupy. These factors can include prey availability (Beukers-Stewart & Jones 2004; Berumen et al. 2005), intra- and interspecific competition (Osenberg

et al. 1992; Berumen & Pratchett 2006), predation pressure (Werner et al. 1983; Madin et al. 2010) and the quest for high reproductive success (Reese 1975; Sogabe et al. 2007; Reavis & Copus 2011). As a response, fishes display various behavioural traits, which further modify their ecological niches and thus their functional impact on the ecosystem (Ferry-Graham et al. 2002b). Behavioural traits commonly show relatively high plasticity (Mittelbach 1984; Robertson 1987; Gardiner & Jones 2010), exacerbating the difficulty of forecasting a realized ecological performance using morphology. Nevertheless, there are some clear examples of morphology predicting behavioural adaptations and thus allowing for an appraisal of realized ecological roles. For example, eye size has been used to make predictions about nocturnal or diurnal activity patterns of reef fishes (Goatley & Bellwood 2009; Schmitz and Wainwright 2011b), while fin aspect-ratios have been found to be strong predictors of habitat utilization behaviour in labrid fishes (Bellwood & Wainwright 2001; Fulton 2007) and good indicators of foraging behaviour in sunfishes (e.g., Robinson et al. 1993). This raises the question: can morphology predict other behavioural traits and therefore aid in understanding ecological performances of reef fishes?

One well-documented behavioural trait in coral reef fishes is the formation of social associations (for instance pairing behaviour or schooling) but to date, no study has examined a potential linkage between morphological attributes and different social associations. However, social systems are tightly linked to the life history of reef fishes, markedly influencing their realized ecological niche. The formation of inter- or intraspecific schools, for instance, has been shown to alter the range of exploitable resources, thereby modifying the realized niche of these schooling individuals (Robertson et al. 1976; Foster 1985; Welsh & Bellwood 2012). In contrast to schooling, pairing has rarely been linked to ecological factors and has largely been examined with regards to reproductive benefits or constraints, especially monogamous reproduction (Barlow 1987; Whiteman & Côté 2004; Reavis &

Copus 2011). Monogamous reproduction is arguably the most common underlying driver of pair-formation in reef fishes and there is strong evidence that the evolution of monogamy has involved changes in various behavioural traits (Emlen & Oring 1977). One trait includes the foraging ecology of species that regularly form pairs: it is argued that, in cases where biparental care is absent, monogamous reproduction arose as a response to feeding on dispersed, small prey items of high nutritional quality, which restricted males to reproduction with just one female due to alterations in female home-ranges (Emlen & Oring 1977; Whiteman & Côté 2004). Accordingly, previous studies have found correlations between foraging ecology, prey types, pairing behaviour and monogamous reproduction (e.g. Barlow 1987; Hourigan 1989).

Thus, assuming that pairing and monogamous reproduction have evolved for similar reasons in most reef fishes (except, for instance, burrowing gobies [f. Gobiidae]; e.g. Reavis & Barlow 1998; Pratchett et al. 2006), it appears reasonable to assume that there are parallels in the ecology of species that form pairs. Given that ecological traits are often manifested in morphological features (e.g. Motta 1988; Schmitz & Wainwright 2011b), one would predict that pairing reef fishes exhibit certain morphological features, shared between phylogenetically distinct species. Specifically, when considering previous findings relating ecological traits to monogamous reproduction in reef fishes, the question arises if species that occur in pairs display morphological adaptations that enable the foraging on small prey items.

Reef fishes that obtain their food by biting or scraping the reef substratum represent a promising group to explore this issue for several reasons. Firstly, a distinct set of lineages display this foraging strategy including the butterflyfishes (Chaetodontidae), angelfishes (Pomacanthidae), surgeonfishes (Acanthuridae), rabbitfishes (Siganidae), parrotfishes (Labridae, tribe Scarini), tetraodontiform fishes and several smaller lineages (e.g., Ehippidae, Kyphosidae) (Wainwright & Bellwood 2002; Konow et al. 2008), offering a

diverse range of taxa. Secondly, their close association with the substratum and the associated benthic community has allowed these lineages to diversify in their cranial morphology and diet. This provides a broad spectrum of different morphologies (Konow et al. 2008; Goatley & Bellwood 2009). Finally, there is high variation in terms of the social associations formed by species within these lineages, making comparisons possible both within and between lineages.

Given that in the past, ecomorphological studies have contributed greatly to our understanding of reef fish ecology (Motta 1988; Wainwright 1991; Bellwood & Wainwright 2001; Ferry-Graham et al. 2001a; Fulton & Bellwood 2002; Collar et al. 2008; Konow et al. 2008), the goal of the present study was to compare the morphology of biting/scraping coral reef fish species in different social systems to identify possible ecomorphological patterns and determine if pair-forming species share any common morphological traits. Specifically, the aims were: 1) to examine the gross morphology of the head, snout, eyes and basic body shape of biting and scraping coral reef fishes to determine, if pair-forming, biting reef fishes share a similar morphospace and, if so, 2) to provide a hypothesis for the functional basis of such ecomorphological segregation.

3.2 Materials and methods

A total of 47 substratum biting or scraping species within the families Chaetodontidae, Pomacanthidae, Acanthuridae, Siganidae, Kyphosidae and scarine Labridae were examined. These groups of biting/scraping taxa were chosen as they are abundant on coral reefs and show a broadly similar foraging ecology, in that they feed by grasping or scraping items from the benthos (Konow et al. 2008).

3.2.1 Social systems

In order to compare the ecomorphology of species in different social systems, the most prevalent social system of each species was determined among 12 different sites on Lizard Island, Great Barrier Reef (GBR), Australia. At each site, the first encountered individual of each focal species was designated as being either solitary (no conspecifics within five metres of the focal individual), paired (two conspecifics in close proximity with coordinated movements and no antagonistic behaviour, no other conspecific individuals in close proximity), a trio (three conspecifics, coordinated movements, no antagonistic behaviour, no other conspecific individuals in close proximity), or in a school (more than three individuals in close proximity with coordinated movements). Each individual was classified on first observation, and then followed for 30 seconds after detection to ensure continuity of behaviour (all individuals retained the original designation). Furthermore, only one individual in each association was assessed, and only species with more than five observations were considered in the data analysis (mean of 58.6 ± 6.8 (SE) observations per species). A total of 2,753 focal observations were recorded from 47 species in the six families. Morphological measurements were then obtained for the observed species. The 47 species include the vast majority of biting and scraping teleosts found at Lizard Island.

3.2.2 Morphology

Morphological measurements were taken from published ‘Randall-style’ images of dead, pinned fishes (Woodland 1990; Froese and Pauly 2012). These have a standard presentation and permit accurate measurements of fishes in lateral presentation, with a mean of 2.1 images per species. All specimens were freshly caught and only mature individuals were considered. Five different morphological measurements were taken from each replicate: 1) the “snout angle” (S_A , **Figure 3.1**). After defining the midline (passing horizontally through the mid caudal peduncle), an eye-line was drawn perpendicular to the midline touching the front of

the eye. A second line, perpendicular to the midline, was drawn half-way between the vertical eye-line and the anterior tip of the premaxilla. From this line, lines were drawn to the tip of the premaxilla from the point where the second line crossed the upper and lower margins of the snout. The angle formed by those two lines marks the snout angle; 2) the “head angle” (H_A , Fig. 1) is formed by the upper snout angle line and a line connecting the upper margin of the snout to the crossing of lines II and IV (parallel to midline touching the dorsal rim of the orbit) in front of the orbit; 3) body depth (B_D) is measured along a line, perpendicular to the midline, drawn from the first dorsal spine to the lower margin of the body; 4) eye diameter (E_D): Measured parallel to the midline, through the centre of the pupil and, 5) snout length (S_L) measured from a line drawn vertically through the centre of the eye and along a perpendicular, horizontal line to the tip of the premaxilla. All variables were expressed as residuals from measurements regressed against standard length (SL) of the respective measured specimens. A mean of each variable was calculated for each species. In addition, the maximum size of each species was obtained from the literature (Randall et al. 1997). Measurements were taken using vernier callipers and a protractor.

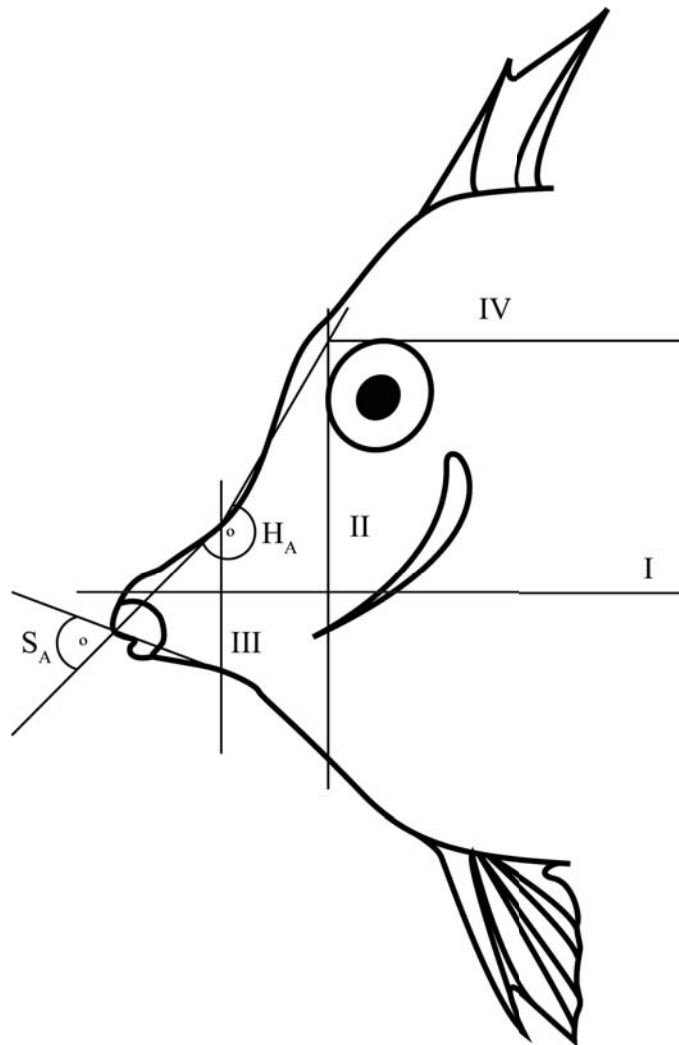


Figure 3.1: Morphological measurements of the head. From the midline (I), a perpendicular line (II) is drawn touching the anterior rim of the orbit. A second perpendicular line (III) is drawn half way between (II) and the anterior tip of the premaxilla. Where this line meets the upper and lower rim of the snout, lines are drawn to the tip of the premaxilla. The angle formed by these two lines defines the snout angle (SA). A third line is drawn parallel to the midline touching the dorsal margin of the orbit (IV). The head angle (HA) is the angle between the upper line of the snout angle and a line that connects the snout intercept with the crossing of lines (II) and (IV).

3.2.3 Statistical analysis

To explore morphological variation among species, a principal components analysis (PCA) was performed, using the mean of the residuals of each of the five morphological measurements for each species, and maximum size as variables. Species on either side (negative and positive) side of PC1 were compared with regards to the prevalence of solitary individuals, pairs, trios and schools using a non-parametric Mann-Whitney U test (to allow for unequal sample sizes). In order to test for the effect of size, each morphological variable was tested using analyses of covariance (ANCOVA) with standard length (SL) as continuous, pair-formation as categorical, and the respective morphological measurements as dependent variables. As the species used for this analysis represent non-independent samples due to their phylogenetic relationships, I used phylogenetic analysis to evaluate the consistency of patterns when accounting for phylogenetic relationships (Felsenstein 1985). This was only possible within families, as no published phylogeny exists for all of the species used in this study. For the Chaetodontidae and Siganidae, pruned phylogenies were obtained from Fessler and Westneat (2007) and Kuriwa et al. (2007), respectively. These phylogenies contained all of the species examined in this study (except for *Chaetodon lunulatus*). No significant linear or non-linear trends were found in the data, leading to the assumption that all branch lengths were consistent with a Brownian motion model. To evaluate the relationship between morphology and pair-formation the scores on PC1 (as a representative summary of the morphological features) and the percentage of individuals in pairs were compared in linear regression analyses using the phylogenetically independent contrasts (PICs) for both variables (R Development Core Team 2012, packages ‘ape’ and ‘caper’). Due to the lack of well resolved phylogenies that include the majority of species examined in this study, simple linear regression analyses were performed for the Acanthuridae and Labridae without accounting for phylogenetic biases. In the Chaetodontidae, *Chaetodon baronessa* was

excluded from this analysis due to extensive collection by other researchers in the main observation sites immediately before the observation period. Thus, local populations of *C. baronessa* were not considered to be natural.

3.3 Results

The prevalence of pairing varied considerably within and between families: in the Siganidae (n = 731 observations), 53.1% of all individuals were observed in pairs, with the highest proportion in *Siganus vulpinus* (80.4%) and the lowest in *Siganus canaliculatus* (0%). This is followed by the Chaetodontidae (33.5%, n = 314), ranging from *Chaetodon lunulatus* with 61.2% of individuals in pairs to *Chaetodon plebeius* at 7.1%. The Pomacanthidae (n = 145) showed a comparable prevalence of pairing (30.6%), but with less variation; from the highest in *Centropyge vroliki* (35.7%) to lowest in *Pomacanthus sexstriatus* (25.8%). Within the Acanthuridae (n = 736), 13.6% of all fishes were encountered in pairs, with the highest prevalence in *Zebrasoma scopas* (43.3%) and the lowest proportions in *Acanthurus lineatus* and *Acanthurus blochii* (0%). In the Kyphosidae, only *Kyphosus vaigiensis* had sufficient observations (n = 26) and only 3.8% of these individuals were found in pairs. Similarly, only 3.5% of the scarine Labridae (n = 799) were found to be paired, with *Scarus rubroviolaceus* exhibiting the highest prevalence of pairing (13%), while *Scarus flavipectoralis*, *Scarus ghobban*, *Scarus globiceps* and *Scarus psittacus* were not observed to form pairs (0%).

3.3.1 Morphology

The PCA of the morphology of the 47 species showed a clear segregation of species along PC1, with the first axis explaining 40.8% of the variation (**Figure 3.2a,b**). Species with a high prevalence of pairing were located on the positive end of PC1. Except for the Labridae (exclusively negative), all families contained species that were located on both sides of

PC1. The second axis (PC2) explained 22.3% of the variation, with the Acanthuridae being exclusively on the negative side and the Pomacanthidae and Chaetodontidae on the positive side. All other families were scattered on both sides. Scores on PC1 were mostly driven by the eye diameter and the head angle (convexity or concavity of the forehead) and to a lesser extent by the snout angle, body depth, and maximum size. PC2 was predominantly driven by the snout length and to a lesser extent by maximum size (**Figure 3.2c, Table 3.1**). When comparing the prevalence of pairing on the negative ($n=22$ species) and positive ($n=25$) side of PC1, pair formation was significantly more prevalent in species on the positive ($35.8 \pm 4.7\%$ (mean \pm SE) of individuals in pairs) than on the negative ($9.0 \pm 3.2\%$ (mean \pm SE) of individuals in pairs) side (Mann-Whitney U test: $p < 0.001$). Schooling showed an inverse pattern ($13.9 \pm 4.7\%$ (mean \pm SE) individuals in schools on the positive side of PC1, $25.3 \pm 4.8\%$ (mean \pm SE) on the negative side; Mann-Whitney U test: $p = 0.03$). Likewise, solitary individuals were significantly less prevalent on the positive side ($47.4 \pm 4.5\%$ SE) than on the negative side ($63.3 \pm 4.3\%$ SE) (Mann-Whitney U test: $p = 0.04$). The prevalence of trios was low on both sides of PC1 ($2.9 \pm 0.7\%$ and $2.4 \pm 1.3\%$ SE, respectively; Mann-Whitney U test: $p > 0.05$).

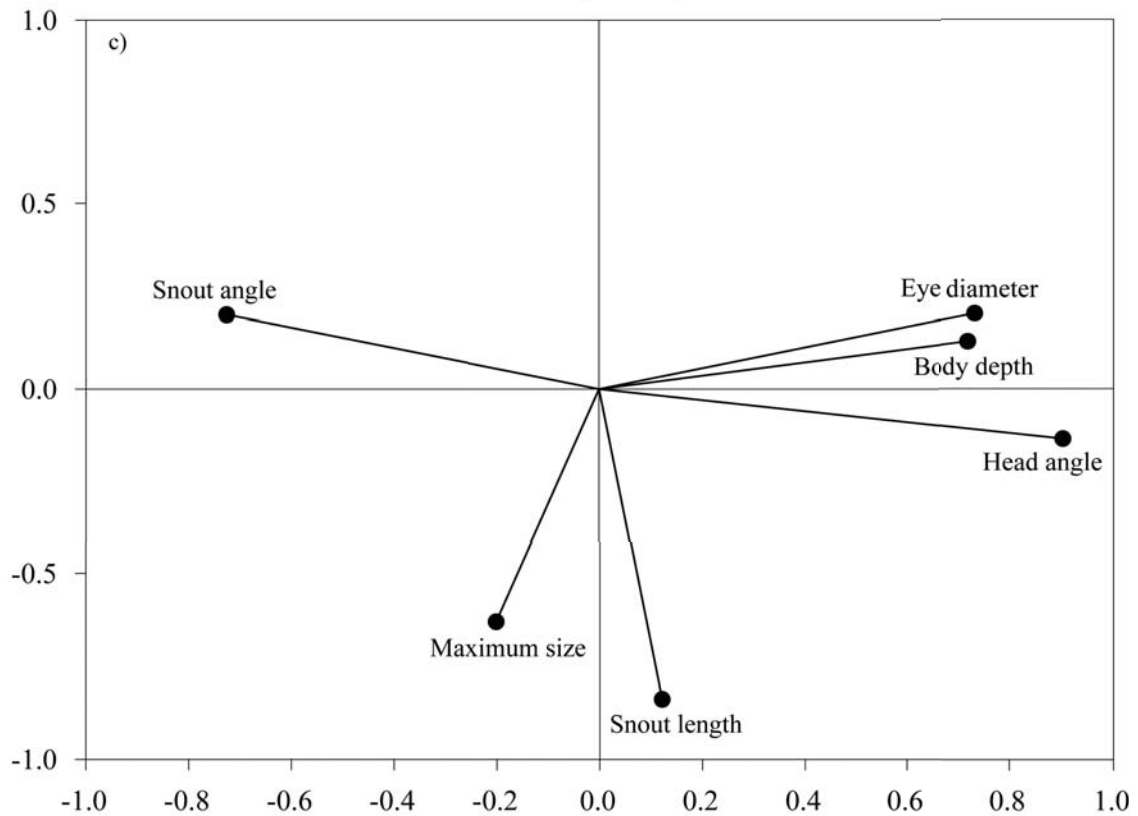
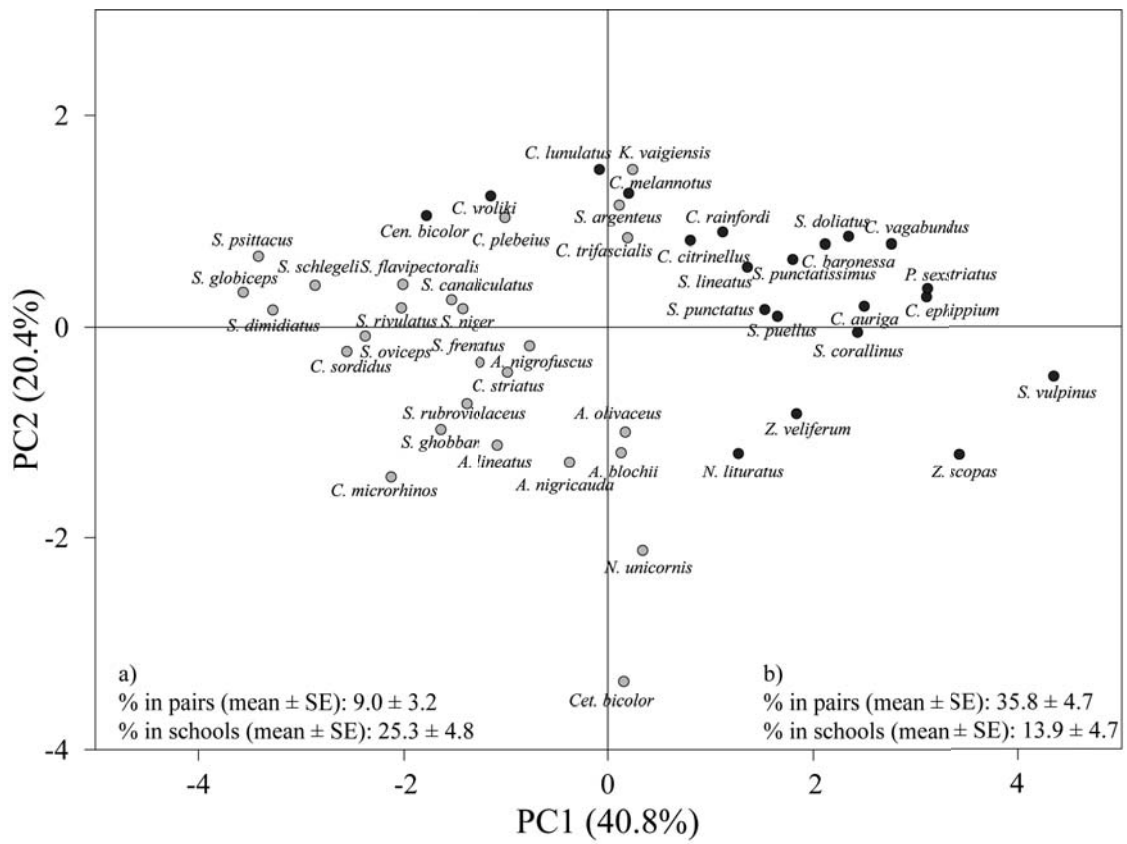


Figure 3.2: Principal component analysis of morphological features and the distribution of species in morphospace. a) Species with low prevalence of pairing behaviour are located predominantly on the negative side of PC1 whereas b) species with high proportions of pairing are found on the positive side of PC1. Schooling proportions show an inverse pattern. Pairing species are represented by black dots, whereas non-pairing species are in black. c) shows the morphological attributes driving the pattern, with eye diameter, the head angle, snout angle and body depth being the main factors driving differentiation along PC1.

Table 3.1: Eigenvalues and factor loadings for the principal components analysis performed on residuals of morphological traits compared to standard length.

	Eigenvalue	% total variance	Cumul Eigenvalue	Cumul %			
Factor 1	2.447455	40.79092	2.447455	40.7909			
Factor 2	1.223603	20.39339	3.671059	61.1843			
Factor 3	0.930829	15.51382	4.601888	76.6981			
Factor 4	0.666249	11.10415	5.268137	87.8023			
Factor 5	0.483022	8.05036	5.751158	95.8526			
Factor 6	0.248842	4.14736	6.000000	100.0000			
	Factor 1	Factor 2	Factor 3	Factor 4	Factor 5	Factor 6	
S_A	-0.728549	0.197066	0.035396	0.554128	-0.321502	0.136773	
H_A	0.900723	-0.137511	0.001422	-0.008800	-0.122124	0.393440	
B_D	0.717580	0.126305	-0.177412	0.567338	0.324868	-0.101190	
E_D	0.731271	0.202678	0.472616	0.033127	-0.389908	-0.218340	
S_L	0.119335	-0.841167	-0.431204	0.105151	-0.255069	-0.127060	
M_S	-0.203592	-0.633456	0.699140	0.158389	0.205222	0.035829	

The ANCOVAs revealed that, size differences between species had no influence on the morphological segregation of pairing and non-pairing species. On average, pair-forming species had significantly larger eyes ($F_{1,90} = 33.3$; $P < 0.001$), wider head angles (occasionally exceeding 180° , i.e. concave, $F_{1,90} = 86.1$; $P < 0.001$), narrower snout angles (ANCOVA: $F_{1,90} = 19.5$, $P < 0.001$), and deeper bodies ($F_{1,90} = 25.4$; $P < 0.001$) than non-pairing species. Snout length did not vary significantly ($F_{1,90} = 0.58$; $P = 0.448$). Only the head angle showed a weak negative relationship to standard length in pairing species (slope = -0.43 , $r^2 = 0.18$), while no such relationship was found in non-pairing species (slope = 0.01 , $r^2 < 0.01$). All other morphological features showed no significant size relationship.

The apparent relationship between morphology and pairing among all taxa is also evident when the species within families are tested in a phylogenetic analysis to account for non-independence of samples. For both, the Chaetodontidae and the Siganidae, linear regressions exploring the relationship between morphology and pair-formation showed a relationship when performed on the traits (Chaetodontidae: $r^2 = 0.59$, $P = 0.026$; Siganidae: $r^2 = 0.72$, $P = 0.004$; **Figure 3.3**). This was consistent when the regression was performed on phylogenetically independent contrasts of the traits (Chaetodontidae: $r^2 = 0.76$, $P = 0.014$; Siganidae: $r^2 = 0.49$, $P = 0.022$) (**Figure 3.4**). As the results of the uncorrected data were congruent with the results of the phylogenetically corrected data in two families, it is unlikely that phylogenetic effects are confounding the overall results. A significant relationship between morphological attributes and pair-formation was also found in the Acanthuridae ($r^2 = 0.82$, $p < 0.001$), while in the Labridae, where pair-formation is uncommon, no such relationship was found ($r^2 = 0.15$, $p = 0.166$). As *Chaetodon lunulatus* was a strong outlier showing high pairing prevalence and a comparably low score on PC1, and was not present in the phylogeny used for the phylogenetically independent contrasts, this species was excluded from the regression analysis ($r^2 = 0.20$; $P = 0.23$ including *C. lunulatus*).

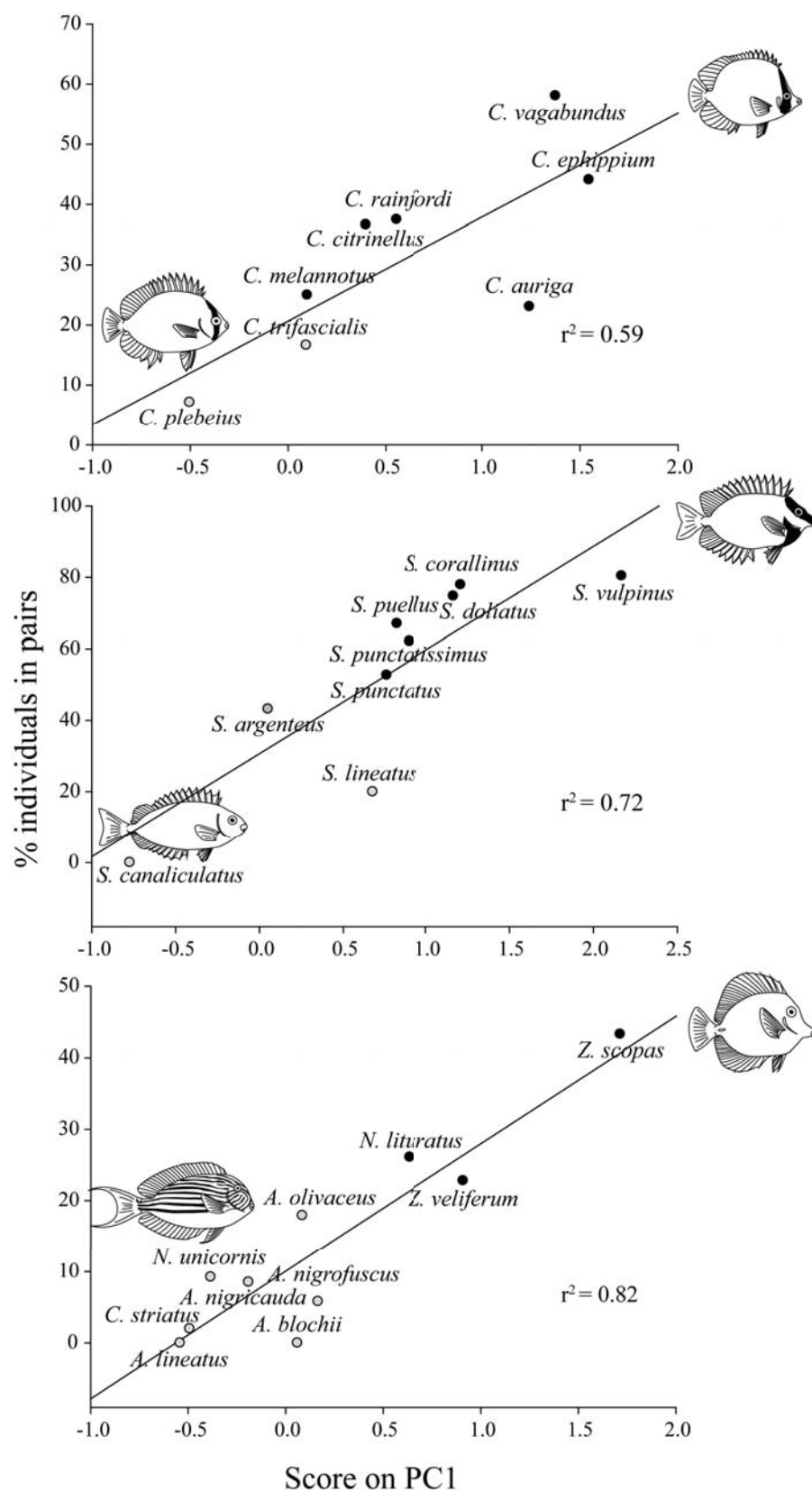


Figure 3.3: The relationship between pair-formation and morphology (scores on PC1) in the Siganidae, Chaetodontidae and Acanthuridae. All families show a significant relationship between their overall morphology (score on PC1) and their tendency to form pairs (% individuals in pairs).

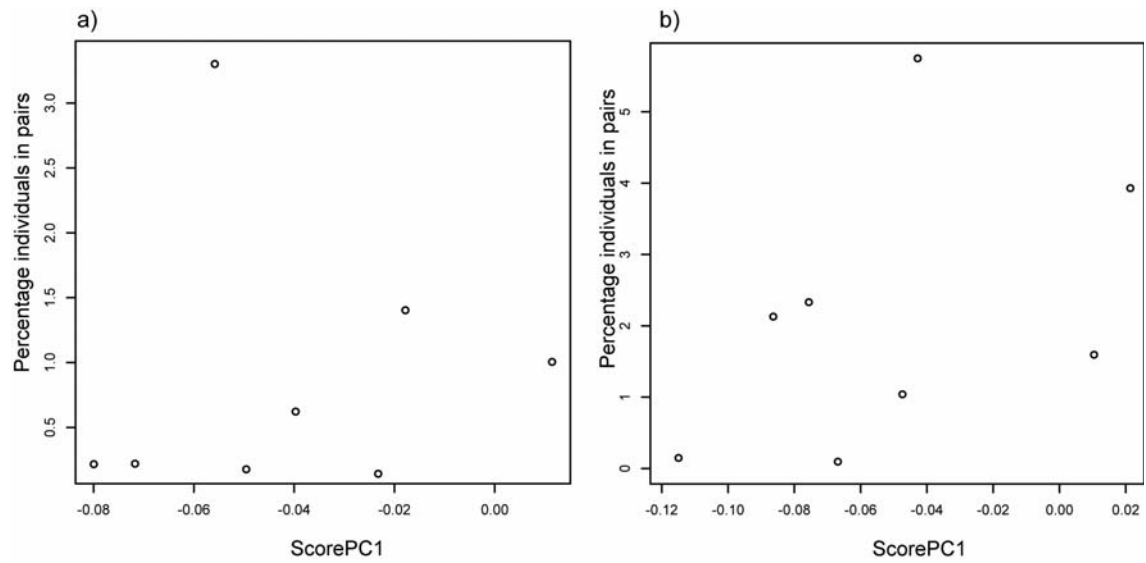


Figure 3.4: Plots of the linear regressions performed on phylogenetically independent contrasts of the score on PC1 and the prevalence of pairing for (a) the Chaetodontidae and (b) the Siganidae. Points represent respective nodes in the phylogeny.

3.4 Discussion

Among biting and scraping teleosts, the prevalence of pairing varied considerably. Pair-formation was most common in the Siganidae, followed by the Chaetodontidae, Pomacanthidae and Acanthuridae. It is negligible in the Kyphosidae and scarine Labridae. When mapped onto the morphological ordination, pair-forming taxa of all families occupied a similar morphospace, characterized by large eyes, an obtuse head angle (concave forehead), an acute snout angle (pointed snout), a deep body, and a relatively small maximum length. Interestingly, snout length was not correlated with pair-formation. A significant relationship between the species' position in morphospace and the proportion of individuals in pairs was also revealed within the Acanthuridae, Chaetodontidae and Siganidae. Phylogenetic relationships do not appear to be driving this pattern, as no phylogenetic effects were found in the Chaetodontidae and the Siganidae when examined using phylogenetically independent contrasts of the data. There was no relationship between pairing and morphology in the scarine Labridae.

The results raise two major questions: 1) what is the putative functional significance of the observed morphological features in pairing species and 2) how does this relate to the role of pair-formation in these species?

3.4.1 Functional significance of pair-forming morphology

The five morphological traits associated with pairs (large eyes, concave foreheads, pointed snouts, deep bodies, and small maximum sizes) are examined separately below. Larger eyes have been shown to improve the general visual capabilities of fishes, enhancing both acuity and sensitivity (Li et al. 1985; Fernald 1991; Miller et al. 1993; Schmitz & Wainwright 2011a). Thus, based on superficial morphology, and without considering physiological differences, pair-forming species appear to display relatively well-developed visual

capacities. An enhanced visual performance is known to favour the detection and discrimination of small, delicate prey items (Wetterer 1989; Schmitz & Wainwright 2011a) especially in dim, weakly illuminated reef microhabitats like caves, crevices or sub-rubble systems (Marshall et al. 2003; Schmitz & Wainwright 2011b). Larger eyes would therefore allow for high foraging selectivity and may have important implications for the feeding behaviour, as prey detection is the first crucial step of a successful foray (Ferry-Graham et al. 2002b).

The concave forehead might also play a critical role in foraging. This relates to depth perception. Consumption of attached and motionless food sources requires considerable coordination, starting with targeting and approaching the prey at the right angle and distance (Rice & Westneat 2005). Depth perception is crucial at this stage for estimating distances and in the recognition of three-dimensional substratum structures. In fishes, depth perception is usually achieved with binocular visual fields, i.e. stereopsis (Wetterer 1989; Job & Bellwood 1996; but see Pettigrew et al. 2000). Stereopsis is established by an overlap of the monocular visual fields of each eye. The extent of this overlap, hence the binocular visual field, depends on the convergence distance from the centre of the eye to the convergence point (McComb & Kajiura 2008). Thus, shorter convergence distances yield larger overlapping areas, a broader binocular visual field, and ultimately, a larger visual field in which depth and three-dimensional structure can be recognized. This field is usually largest along the line of the visual axis (Tamura 1957). A concave forehead may substantially decrease the binocular convergence distance (**Figure 3.5**), thus yielding an improved depth perception as a function of a larger binocular field for these species, especially as the visual axis in substratum feeding species is expected to be antero-ventral (Tamura 1957). While other factors, such as head width or lateral protrusion of the eyes, are likely to influence the convergence distance, it appears plausible that a concave, rather than a convex forehead, may decrease the

convergence distance. Similarly, the alignment of the snout with the visual axis, in the binocular field, will be facilitated by a concave forehead and will help maintain accurate visual control of snout positioning during foraging (Martin & Katzir 1999; Guillemain et al. 2002). Thus, the concave forehead, as seen in pairing species, might enable improved visual perception of substratum topography and allow for precise close-range evaluation of the benthos in topographically complex microhabitats.

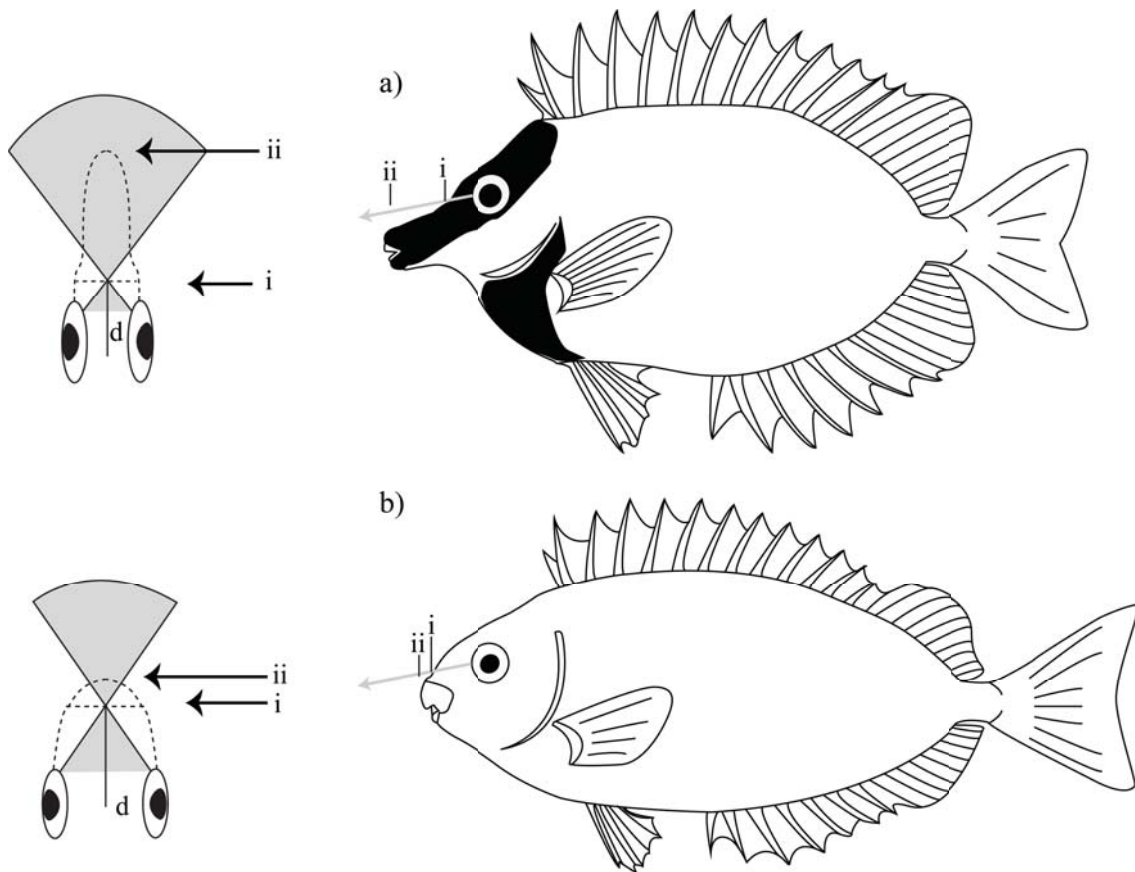


Figure 3.5: An estimate of possible differences in binocular visual fields of fishes with different morphologies. Pair-forming species' (example a, *Siganus vulpinus*) morphology may allow for a shorter convergence distance (d) and a convergence point closer to the eyes (i) despite maintaining an elongated snout (ii), located in the centre of a larger binocular field. Schooling species (example b, *Siganus canaliculatus*) lacking the morphological attributes of pairing species appear to show inverse patterns. Both fishes are drawn from photographs at the same scale. Distances and angles of the visual fields are estimates based on photographs, osteological and preserved material, without accounting for other factors that can influence the visual field such as lateral eye protrusion or aphakic apertures.

The pointed snout allows for delicate, dexterous movements and can function like a pair of forceps thus facilitating a selective foraging behaviour (Ferry-Graham et al. 2001b; 2008; Hernandez et al. 2009). Furthermore, this snout configuration grants access to small crevices and interstices, where concealed prey items, which cannot be exploited by other species, can be utilized. This is a well-known trait of some long-jawed butterflyfishes (Motta 1988; Ferry-Graham et al. 2001a, 2001b) and has recently been identified for pairing species within the Siganidae (Fox & Bellwood 2012). However, despite the pointed snout, some species would still be able to produce efficient gape expansion even while having their snout in interstices. Pair-forming Siganidae (*Siganus doliatus*) and Acanthuridae (*Zebrasoma veliferum*) both exhibit exceptional gape expansion due to intramandibular flexion (Konow et al. 2008), permitting them to achieve relatively large gapes even with a small mouth. High body depths might also contribute to this foraging pattern in enhancing manoeuvrability (Webb 1997; Gerstner 1999). Similarly, a smaller maximum size in pairing species would be beneficial in terms of accessing topographically complex habitats.

Overall, the large eyes, the concave forehead, pointed snout and general body proportions that unite pair forming fishes in a similar morphospace all appear to favour foraging on small, discrete and potentially concealed prey items. The prey can be detected and distinguished due to enhanced visual performance (large eyes), approached and targeted using improved depth perception due to the concave forehead, and ultimately reached and gathered utilizing a pointed snout.

This suite of morphological traits is found in all three families where pairing is common and shows a significant relationship to the prevalence of pairing in the examined species. In the Siganidae, *Siganus corallinus*, *Siganus puellus* and *Siganus vulpinus* exhibit all the mentioned morphological traits in a pronounced form, are strongly paired, and have recently been identified to forage selectively within small reef crevices and interstices (Fox &

Bellwood 2012). In contrast, siganid species lacking this morphology (*Siganus canaliculatus*, *Siganus argenteus* and *Siganus lineatus*) occur predominantly in groups and are reported to feed unselectively on a wide range of large, erect macroalgae or detrital matter (Woodland 1990; Paul et al. 1990; Fox & Bellwood 2008, 2011; Fox et al. 2009). In the Chaetodontidae, all examined species are known to feed on small, benthic prey items (Pratchett 2005).

Generalists and micro-invertebrate feeders (e.g. *Chaetodon ephippium*, Pratchett 2005) appear to be more frequently paired and possess a more pronounced morphology.

Accordingly, the extreme long-snouted genera *Forcipiger* and *Chelmon* are usually found in pairs (Allen et al. 1998; Ferry-Graham et al. 2001a, 2001b) and are reported to exploit reef crevices and interstices (Motta 1988). There are no dietary differences between the remaining species, despite marked variation in their pairing prevalence. This suggests that there may be other factors in the foraging ecology of the Chaetodontidae (such as the choice of feeding microhabitats), underpinning the differences in pairing prevalence. *Chaetodon lunulatus* was excluded from my analyses as it was the only strong outlier. Belonging to the crown-subgenus *Corallochaetodon*, this species has a different morphology, dentition patterns and greatly exceeds other *Chaetodon* species in intramandibular flexion (Motta 1988, 1989; Konow et al. 2008), suggesting different bite kinematics and a different foraging strategy.

Finally, in the Acanthuridae, species in the genus *Zebrasoma* are most frequently paired and possess the most modified morphological attributes. Correspondingly, these species are reported to differ from other acanthurids in their foraging ecology in exploiting a wide range of microhabitats (Robertson et al. 1979; Robertson & Gaines 1986; Montgomery et al. 1989; Fouda & El-Sayed 1994). In contrast, most members of the genus *Acanthurus* rarely form pairs and appear to have a morphology unsuitable for selectively foraging on small, benthic prey items. Matching this, most members of the genus *Acanthurus* are described as relatively unselective grazers of the open EAM (epilithic algal matrix) (Russ 1984; Fox & Bellwood

2012). Likewise, parrotfishes are known to relatively unselectively scrape algae, detritus and associated matter off planar EAM-covered substrata (Bellwood & Choat 1990; Bonaldo & Bellwood 2011; Fox & Bellwood 2012), suggesting that their foraging strategy does not entail the need for pairing. There was no significant relationship between morphological features and pair-formation in parrotfishes.

These results are highly consistent with one of the prevailing hypothesis for the evolution of monogamy in reef fishes, based on the restricted ability of males to guard more than one female, as a function of altered home-ranges which are shaped by foraging on small, widely dispersed, high quality food items (Emlen & Oring 1977). Thus, given that this way of foraging appears to be a consistent feature of the majority of pair-forming species examined in this study, it appears likely that the species that form pairs may also do so for reproductive reasons and that they mate monogamously. This is supported by observations on, for instance, *Zebrasoma scopas*, *Zebrasoma veliferum* and *Chaetodon lunulatus*, which are reported to reproduce in pairs and are probably monogamous (Robertson 1983; Yabuta 1997; Whiteman & Côté 2004). However, for other species, including all species within the Siganidae and the butterflyfish *Chaetodon ephippium*, there is no record of pair spawning. Conversely, the latter species and the rabbitfish *Siganus punctatus*, both strongly pairing and with the associated morphological features, are reported to spawn in aggregations (Johannes 1981; Yabuta 2007), offering little potential for monogamous reproduction (Emlen & Oring 1977). While the lack of data makes it difficult to assess the importance of monogamous reproduction as a driving force behind pairing, it does appear to be a significant factor in some species. Overall, my observations support the suggestion that pairing arose as a response to foraging on small, high quality, but dispersed, food items, which may also be associated with, or gave rise to, monogamous reproduction.

3.4.2 An ecological role for pair-formation?

In species that forage on small, inaccessible prey items, vulnerability to predation may be a crucial factor. Vulnerability to predation appears to be significantly decreased when feeding in groups (Ryer & Olla 1998; Overholtzer & Motta 2000; White & Warner 2007a; DeMartini et al. 2011). In contrast, vulnerability increases with handling-time of prey (Popp 1988), restricted visual fields (Guillemain et al. 2002), and an increased body angle during foraging (Krause & Godin 1996). A foraging strategy targeting small, concealed, benthic prey appears to be particularly dangerous by virtue of time- and attention-consuming dexterous movements and restricted visual fields when in close proximity to the substratum. Thus, although deep bodies, as exhibited by pairing species, may decrease vulnerability while foraging (e.g., Persson et al. 1996), feeding on small and concealed prey is likely to increase the overall vulnerability of the fish to predation (cf. Krause & Godin 1996; Ferry-Graham et al. 2001b, 2008; Rice & Westneat 2005). Given high selectivity for small prey items coupled with inaccessible, structurally complex microhabitats and a patchy distribution of prey, large group sizes may not be compatible with efficient feeding because of decreasing feeding efficiency (White & Warner 2007b). Accordingly, schooling is largely absent in species occupying the ‘pairing’ morphospace (highest prevalence of schooling in *Naso lituratus*, 13%) and is significantly more common among species with ‘non-pairing’ morphological traits ($25.3\% \pm 4.8\%$ (mean \pm SE) of individuals in schools).

Associating with only one individual would result in increased vigilance (vs. solitary feeding) while limiting reductions in feeding efficiency and prey accessibility (vs. schooling) (Ryer & Olla 1998; White & Warner 2007b). In theory, in pairs, one fish can function as an observer while the other fish forages. Performed in an alternating pattern, this benefits both pair members (Wickler 1985). While previous studies have not found a significant increase in feeding rate of paired vs. solitary individuals (Bonaldo et al. 2005; Gregson et al. 2008), it is

possible that feeding on small, concealed prey is only performed effectively when fishes are arranged in a pair, whilst solitary individuals display a more conservative foraging strategy. Hence, species, which are morphologically equipped to forage in this way, might execute this feeding mode only when they are able to rely on the safety provided by a partner.

Overall, the pattern of pairing species occupying a similar ecological morphospace appears to be consistent both among and within families. Using only superficial morphological traits, it appears that species with higher pairing proportions occupy a similar morphospace characterized by large eyes, concave foreheads, pointed snouts, deep bodies, and small maximum sizes. This relationship held for the Acanthuridae, Chaetodontidae, Pomacanthidae and Siganidae, whereas the scarine Labridae and Kyphosidae show no such pattern. This appears to be the first study to reveal a distinct ecomorphospace for pairing reef fish species. It is hypothesized that the morphological attributes allow for selective foraging on small, delicate prey items on micro-topographically complex substrata including reef caves, crevices and interstices. Given this, it is likely that pairing arose for ecological reasons and subsequently gave rise to shared reproductive traits, with several species exhibiting a monogamous mating system. The presence of pairs in species with apparently non-monogamous reproductive strategies suggests that there may be additional factors underpinning the formation and maintenance of pairs. While at this stage, I am unable to resolve the evolutionary history of foraging ecology, pair-formation and monogamy, there does appear to be a strong link between morphology, sociality and ecology in biting coral reef fishes that warrants further investigation.

Chapter 4: Coordinated vigilance provides evidence for direct reciprocity in coral reef fishes

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4.1 Introduction

Cooperation is widespread among animals (Dugatkin 1997), and it is now widely accepted that cooperation is also present among unrelated individuals (Trivers 1971; Mesterton-Gibbons & Dugatkin 1992; Dugatkin & Mesterton-Gibbons 1996; Nowak 2006). In this context, reciprocity or ‘reciprocal altruism’, which involves a costly action beneficial for another individual, based on an expected future payoff through reciprocation, has garnered particular interest (Trivers 1971; Axelrod & Hamilton 1981; Schino & Aureli 2010). This interest has arisen from the notion that cooperative individuals should be prone to exploitation by their respective partners if the latter defects after having received help (Ghoul et al. 2014), therefore leading to fitness declines in cooperating individuals. However, it has been suggested that reciprocity can be evolutionarily stable, even if modelled under an iterated prisoner’s dilemma (IPD), in which both players employ a strategy called ‘tit-for-tat’ (Axelrod & Hamilton 1981; Nowak & Sigmund 1992). The IPD and several other game-theoretical models have subsequently provided frameworks for the evaluation of reciprocity in animals and throughout the last few decades, the presence of reciprocity has been suggested in fishes (Zöttl et al. 2013; Taborsky 1984; Fischer 1988; Milinski et al. 1990; Milinski 1987), birds (Godard 1993; Krama et al. 2012; Krams et al. 2008), and mammals (Carter & Wilkinson 2013; Wilkinson 1984; Rutte & Taborsky 2008; Cheney et al. 2010; Hauser et al. 2003).

However, almost all reported occurrences of direct reciprocity in animals have now been challenged (Clutton-Brock 2009), as many aspects of reciprocity are thought to require a suite of complex cognitive abilities. This includes the recognition of individual partners, the capacity to recall their previous actions, or the ability to make intentional investments under the expectation that it will entail a future reward (Stevens & Hauser 2004; Pfeiffer et al. 2005). For this reason, it has been questioned whether direct reciprocity exists in animals which are assumed to lack complex social and cognitive skills (Clutton-Brock 2009; Clements & Stephens 1995; Milinski & Wedekind 1998; Russell & Wright 2009; Connor 2010; Melis & Semmann 2010), and most evidence of direct reciprocity to date is confined to a few cases in birds and mammals (Carter & Wilkinson 2013; Rutte & Taborsky 2008; Hauser et al. 2003; Taborsky 2013). Yet, in response to these criticisms, it has also been posited that most of the cognitively demanding actions of reciprocity stem predominantly from the theoretical framework in which reciprocity is investigated (Schino & Aureli 2010; Taborsky 2013; Raihani & Bshary 2011; Brosnan et al. 2010). Specifically, many aspects of game-theoretical models such as tit-for-tat in the IPD have been questioned over the last two decades (Connor 1995; Noë 2006). Most recently, the discrete time structure of the IPD and other models, as well as their incapacity to allow for the exchange of information among partners during cooperative interactions, have been identified as a major weakness of these models (Sirot 2012; van Doorn et al. 2014). These weaknesses undermine our capacity to adequately judge whether or not reciprocity is present (Carter 2014), particularly given the often emphasized need for continuous information exchange among cooperating partners in a natural setting (Boesch & Boesch 1989; Milinski et al. 1990; Krams et al. 2008).

Teleost fishes have contributed substantially to the debate about cooperation and reciprocity in animals, and a variety of systems have been discussed in the context of the reciprocal exchange of commodities. This includes 1) egg-trading in hermaphroditic hamlets

(f. Serranidae), which describes the exchange of costly eggs for fertilization by the partner (Fischer 1984, 1988), 2) helping behaviour in cooperatively breeding cichlids (f. Cichlidae), which involves the trading of resource access against the provision of brood care and territorial defence (helpers ‘pay-to-stay’; Bergmüller & Taborsky 2005; Bergmüller et al. 2005), 3) interspecific cleaning behaviour by pairs of cleaner wrasses (f. Labridae), which involves the removal of parasites from client fishes (Trivers 1971; Bshary & Grutter 2006; Bshary & Schäffer 2002), and 4) predator inspection in sticklebacks (f. Gasterosteidae) and other small fishes (Dugatkin 1997; Dugatkin & Alfieri 1991), which involves a pair of fishes approaching a predator in order to assess the threat arising from its presence, for which the presence of a partner provides a safer situation than if the fish were to approach the predator alone (Milinski et al. 1990; Dugatkin & Alfieri 1991). While most of these systems were initially identified as cases of reciprocity, subsequent assessments and syntheses have argued that they are based on pseudo-reciprocity and by-product mutualism rather than direct reciprocity (Clutton-Brock 2009; Connor 1992, 1996, 2010; Stephens et al. 1997), invoking the limited potential of teleost fishes to cope with the cognitive and social demands of reciprocity. However, there is now substantial evidence for many complex social processes in fishes (Bshary et al. 2002, 2014; Brown 2015; Brown et al. 2011; Ward et al. 2008, 2011), including image scoring (Bshary & Grutter 2006), punishment (Bshary & Grutter 2005), pre-emptive appeasement (Bergmüller & Taborsky 2005; Bergmüller et al. 2005), or partner preference (Croft et al. 2006). Thus, teleost fishes appear to provide a promising group for the investigation of reciprocity (Taborsky 2013; Bshary et al. 2002).

Rabbitfishes (f. Siganidae) are an Indo-Pacific family of teleost fishes, which includes 28 species and is separated in two major groups (Woodland 1990). The first group consists of predominantly schooling species, which are drab-coloured and commonly associated with mangrove and estuarine habitats, while the second group consists of species which occur

mostly in stable pairs, are brightly coloured, and are commonly associated with coral reefs (Brandl & Bellwood 2013a,b). In the latter group, the presence of same-sex pairs has spurred research exploring the potential ecological role of pairing behaviour in this family, and it has been proposed that their foraging habits, which entail the penetration of cracks and crevices in the reef matrix, may necessitate the presence of a vigilant partner (Brandl & Bellwood 2014a,b). Consequently, it has recently been suggested (based on photographic evidence) that pair members may cooperate in order to achieve high levels of vigilance during foraging (Fox & Donelson 2014). However, the potential presence of cooperation among pair members in rabbitfishes has not been evaluated quantitatively and individual costs and benefits are unknown, undermining our ability to judge whether vigilance behaviour in pairing rabbitfishes may be based on reciprocity.

The goal of the present study, therefore, was to quantify the major aspects of pairing behaviour in rabbitfishes, and to assess whether the behaviour exhibited by rabbitfish pairs may satisfy the basic requirements of reciprocal cooperation, such as reciprocal alternation between feeding bouts and a costly investment by one individual that directly benefits the partner.

4.2 Materials & Methods

4.2.1 Data collection

All fieldwork was conducted on coral reefs around Lizard Island, a granitic mid-shelf island in the northern Great Barrier Reef, Australia (14°40'08"S 145°27'34"E). Four different species of pairing rabbitfishes were considered (*Siganus corallinus*, *S. doliatus*, *S. puellus*, and *S. vulpinus*), as they represented the most abundant pairing species in the study area (Big Vicki's Reef). The four examined species differ in their dietary preferences, with *S. corallinus* and *S. doliatus* feeding predominantly on red corticated and red filamentous algae,

while *S. puellus* feeds mainly on sponges, and *S. vulpinus* predominantly on cyanobacteria (Hoey et al. 2013). However, all species are similar in their foraging behaviours, as all four species have been reported to exploit cryptic reef micro-habitats such as cracks and crevices in the substratum (Brandl & Bellwood 2014a). In addition, all four species are strongly pair-forming, with approximately 80% of all adult individuals occurring in pairs in *S. corallinus*, *S. doliatus*, and *S. vulpinus*, and approximately 70% in *S. puellus* (Brandl & Bellwood 2013a).

Haphazardly encountered pairs of the four species were followed while SCUBA diving and video footage of their behaviour was collected for a period of 12 minutes for each pair (using Sony DCR-SR300E camcorders). Care was taken to sample different sites on the reef in order to avoid re-sampling of the same pairs. For each pair, the size was estimated, and video recording commenced one minute after the fish were observed to feed, in order to prevent behavioural biases due to the presence of the observer (foraging was interpreted as a sign that fish had acclimatized to the observer). Observers aimed to keep a constant distance from the recorded fish; video sequences in which the distance to the fish resulted in unreliable examination of fish behaviour or obvious behavioural responses to the observer were discarded from the analyses. Videos were collected during three different times of day (0600 to 1000h; 1000 to 1400h; 1400 to 1800h) and subsequently analysed in 5-second intercepts. Specifically, videos were paused every five seconds to determine the current behaviour of the partners as a point measure. For every 5-second point intercept at which both members of the pair were visible in the video, the angles of both individuals (θ_A and θ_B) relative to a vertical line perpendicular to the substratum were measured using a protractor that was superimposed on the computer screen (**Figure 4.1**). Angles were assessed quantitatively to describe a conspicuous behaviour, henceforth termed ‘vigilance position’, in which one individual was found hovering high above the substratum with its head oriented

upwards (**Figure 4.1c**). In addition, the fish's height above the substratum (cm), the distance between pair members (cm), the complexity of the surrounding microhabitat (1 = lowest complexity to 5 = highest complexity, Wilson et al. 2007), and the behaviour of individuals (swimming, foraging, hovering, or displaying) were recorded. These parameters were recorded because of their potential influence on the behaviour exhibited by the partners (for instance, higher complexity of the surrounding environment may reduce the threat arising from predation through the provision of shelter (Beukers & Jones 1998). Swimming was defined as active, directional movements, while foraging was defined as at least one of the two individuals engaged in active feeding (i.e. biting from the substratum). Hovering entailed both individuals being stationary and motionless (similar to the vigilance position), and displaying was noted when individuals engaged in displays towards other pairs of the same species. The incidence of fin-flicks (rapid flashing of dorsal, pelvic, or anal fin spines) was also noted, along with specific information on the identity of the fin-flicking fish and the subsequent behaviours of both individuals. This was recorded based on evidence from other families of fishes, that suggests that fin-flicks serve as a communicative signal (Murphy & Pitcher 1987; Brown et al. 1999; Tricas et al. 2006; Parmentier et al. 2011), and the potential importance of communication in reciprocal cooperation (Sirot 2012; van Doorn et al. 2014). All occurrences of presumed flight behaviour (i.e. an individual rapidly abandoning its position followed by extensive swimming behaviour) were likewise recorded, specifying which individual initiated the behaviour, whether the fish's visual fields were obstructed or not, and if the partner followed the flight. I observed 15 pairs each for three species (*S. corallinus*, *S. doliatus*, *S. vulpinus*), and 14 pairs for *S. puellus*, resulting in a grand total of 59 pairs, all of which were filmed for a period of 12 minutes. As only data points were included in which both pair members were in the video frame, and not visibly disturbed by the observer, the average number of non-independent data points (5-second intercepts) extracted

per pair was 83.8 (± 5.1 SE) for *S. corallinus*, 71.7 (± 8.1 SE) for *S. doliatus*, 53.4 (± 5.9 SE) for *S. puellus*, and 71.0 (± 7.2 SE) for *S. vulpinus*.

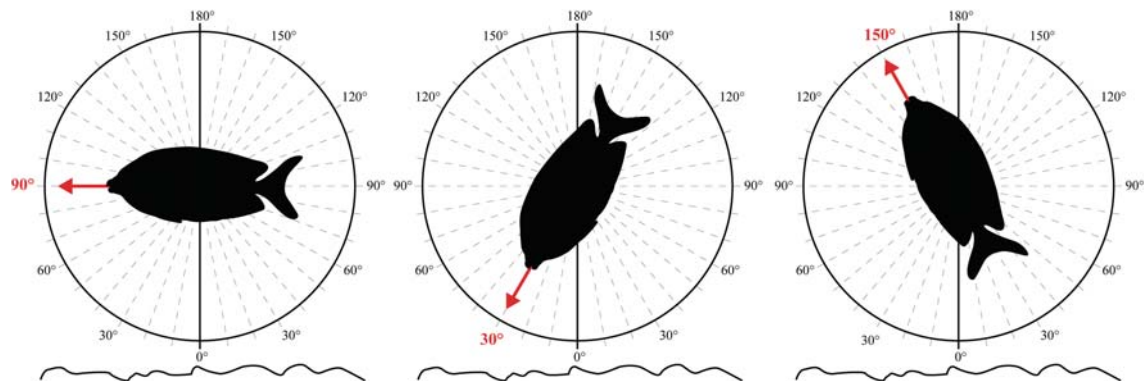


Figure 4.1: Schematic description of the assessment of angles exhibited by rabbitfishes. Angles were assessed based on a vertical line perpendicular to the substratum using a protractor superimposed on the screen. 90° denotes a horizontal position parallel to the substratum (a), while 30° mark a head-down position (b), and 150° mark a head-up position (c). The head-up position (ranging from ~90° to 180°) was identified as a vigilance position.

To determine whether pair members sequentially alternated their roles between feeding bouts, I also counted the occurrence of alternating (A to B, B to A) and repeated (A to A, B to B) foraging bouts, with the latter also including instances where both pair members were feeding simultaneously (AB to A, AB to B, A to AB, B to AB, AB to AB). One foraging bout was defined as active foraging behaviour by either or both of the pair members (A, B, or AB) at a given 5-second intercept until it was interrupted by a different behaviour (or a change of the identity of the forager) at the next 5-second intercept. The next feeding bout commenced at the next 5-second intercept at which one of the individuals (or both) engaged in foraging behaviour. If the individual feeding during the previous bout was feeding again, repeated foraging behaviour was recorded. If the individual not engaged in foraging at the last 5-second intercept engaged in foraging, alternated foraging was recorded. Data were collected for all pairs in all species ($n = 59$).

In addition to the videos of pairing fish, 24 videos of solitary individuals in all four species (average of 6.0 ± 0.44 SE individuals per species; *S. corallinus*: $n = 8$; *S. doliatus*: $n = 3$; *S. puellus*: $n = 7$; *S. vulpinus*: $n = 6$; average observation period of 459.8 ± 41.6 SE seconds per individual) were collected and analysed to quantify the length of vigilance bouts, the number of bites per foray (foray defined as a continuous sequence of bites from the substratum), and the maximum substratum penetration depth during forays. Vigilance bouts refer to the duration (in s) over which a fish was observed in the assumed vigilance posture (a stationary 'head-up' position in the water column, exhibiting an angle $> 90^\circ$), without interruption by feeding or active swimming. The number of bites per foray was quantified as the number of consecutive bites taken by a fish without engaging in other behaviour such as vigilance (defined above). The penetration depth was estimated as the extent (in cm) to which a foraging fish penetrated into cracks in the substratum (Brandl & Bellwood 2014a). The same protocol was performed with 32 randomly selected videos of pairs (eight per species),

where one haphazardly selected individual of the pair was selected for the duration of the video.

4.2.2 Statistical procedures

I used linear and additive mixed effects models to separately analyse the angles of pair members during swimming and foraging (which, when combined, accounted for 92.8% of the behaviours displayed) for each species. I tested whether the angle of one individual in a pair (θA) was independent from predictor variables, including environmental factors (time of day, microhabitat complexity), or variables associated with the partner (the angle of the partner [θB], its height above the substratum, and the distance between pair members). For all analyses, pair ID was included as a random factor to account for non-independence of points taken from the same pair. For the data gathered during foraging activity, residual plots indicated non-linearity for θA as a dependent variable in all species. Thus, data were analysed using generalized additive mixed effects models (GAMMs) with a Gaussian error distribution and a cubic regression spline smoother, calculated by automatic cross-validation, for θB and *heightB* during foraging (hastie & Tibshirani 1990). Due to temporal non-independence of behaviours (i.e. an individual might be more likely to assume an angle close to the angle from the previous data point), a temporal autocorrelation function was also added (Zuur et al. 2009). The analysis was repeated for θB as the dependent variable for all species. Variables included in the final model were selected using likelihood ratio tests, and model fits were assessed using residual plots. Angles during swimming were analysed in the same fashion but using generalized linear mixed effects models (GLMMs) with pair ID included as a random factor and incorporating a temporal autocorrelation coefficient. This was performed in order to demonstrate that the observed behavioural patterns during foraging are not simply a random behaviour, which is also present during other aspects of the fishes' daily activity.

To test whether pair members alternated their roles more often than they performed the same role consecutively over the 12-minute observation period, occurrences of sequential changes in the identity of the foraging individual (either alternating or repeated foraging) were analysed. All transitions to or from bouts where both individuals were feeding simultaneously were assigned to be repeated in order to yield conservative estimates. The occurrence of alternating or repeated foraging bouts was analysed using four species-wise zero-inflated GLMMs with a negative binomial error distribution, using counts of alternated and repeated foraging events within pairs as dependent variables and pair ID as a random factor to account for the non-independence of data collected from the same pair. For all GLMMs, model fits were assessed using residual plots, all of which were satisfactory. For each pair, I also calculated the deviation from a balanced (0.5) proportion of 5-second intercept points at which individual A or B were foraging, and tested the overall distribution of feeding by pair members in each species against a balanced distribution using Pearson's Chi-squared tests. The relative occurrence of fin-flicks during different behaviours (standardized as the number of fin-flicks per 5-second intercept during which a given behaviour was displayed) was analysed using a frequency test (Pearson's Chi-squared test).

To examine potential behavioural differences between solitary and paired fish, solitary and paired individuals were compared for each species, separately, using GLMMs with the respective individual fitted as a random factor to account for non-independence of repeated measures for each fish. All data were modelled using a negative binomial error distribution as non-normality and overdispersion were detected during the modelling process. I tested the effects of the social status (solitary or paired) on the time spent in the vigilance posture (seconds), the number of consecutive bites per foray (bite counts), and the maximum extent to which individuals penetrated the substratum during foraging (cm). The value 1 was subtracted from the count dataset in order to prevent inaccurate estimates due to zero-

truncation (i.e. as at least one bite was necessary to constitute a foray, there were no zeros in the count dataset, possibly resulting in inappropriate model estimates in a Poisson or negative binomial model; Zuur et al. 2009). For the time spent in vigilance posture, seconds were transformed to centiseconds to yield integer values. When the anterior structures of individuals were concealed due to penetration of the substratum, bites per foray were determined by the occurrence of caudal and pectoral fin-movements, which precede food acquisition in rabbitfishes (Pink & Fulton 2014). All analyses were performed using the software R and the packages *mgcv*, *nlme*, and *glmmADMB*.

4.3 Results

In all four rabbitfish species, when foraging, one pair member commonly assumed a stationary, upright position in the water column above the substratum (entailing an angle of 90° or larger), while the partner was feeding. The feeding individual often penetrated deep into cracks and crevices in the substratum with substantial obstructions to its visual field (**Figure 4.2**). Possible flight responses (entailing rapid abandonment of vigilance position or foraging activity) were almost exclusively initiated by the individual positioned head-up in the water column (95.1% of instances), which always had an unobstructed field view of the surrounding environment. In contrast, at the onset of flights, the forager's eyes were often not visible (15.7%), but it followed the fleeing individual in 94.1% of cases, suggesting that individuals in the water column were more vigilant than the forager and that information was rapidly communicated to the foraging fish. It is possible that some of the presumed flight responses were not due to the threat imposed by potential predators but rather in order to engage in territorial defence or simply to move on to another foraging location. However, the high density of predatory fish in the study area (and the presence of the observer as a

potential threat) suggest that at least a proportion of the observed responses were associated with predator-oriented vigilance.



Figure 4.2: Foraging and vigilance postures in four species of pairing rabbitfishes. The foraging individual (in the head-down position) feeds in cracks and crevices in the substratum, while the vigilant individual is positioned in the water column with its head up. Note the obstructions to the visual field of the forager, suggesting high vulnerability to predation and the unobstructed field of perception of the vigilant fish. (a) *Siganus corallinus*, (b) *S. vulpinus*, (c) *S. doliatus*, (d) *S. puellus*.

Pair members strictly coordinated their vigilance efforts, which is reflected by the angles assumed during foraging (**Figure 4.3**). A low angle (= head down, foraging) in individual A (θ_A) was complemented by a large angle (= head up, vigilant) in individual B (θ_B) and *vice versa* in all four species. The GAMMs confirmed that pair members' angles were non-independent and arranged in a contrasting manner (GAMM_{AB}: *S. corallinus*: $edf = 5.781$; $F = 11.23$; $P < 0.0001$; *S. doliatus*: $edf = 3.512$; $F = 11.94$; $P < 0.0001$; *S. puellus*: $edf = 3.802$; $F = 5.105$; $P = 0.0009$; *S. vulpinus*: $edf = 5.116$; $F = 4.438$; $P = 0.0005$), suggesting that individuals take turns in being vigilant. The height of individual B likewise showed a significant inverse relationship with θ_A (GAMM_{AB}: *S. corallinus*: $edf = 3.865$; $F = 13.96$; $P < 0.0001$; *S. doliatus*: $edf = 3.716$; $F = 10.39$; $P < 0.0001$; *S. puellus*: $edf = 2.576$; $F = 2.576$; $P < 0.0001$; *S. vulpinus*: $edf = 3.668$; $F = 20.057$; $P < 0.0001$), suggesting that an unobstructed field of perception (i.e. a large angle and a position high above the substratum) in one fish represents the best predictor for foraging (i.e. a low angle) in the partner (*S. corallinus*: $adj. R^2 = 0.428$; *S. doliatus* = $adj. R^2 = 0.433$; *S. puellus* = $adj. R^2 = 0.570$; *S. vulpinus* = $adj. R^2 = 0.397$). Except for *S. corallinus*, in which the time of day exhibited a small effect on the angle of individual A ($P = 0.004$; $adj. R^2 = 0.443$), the angle and height of the partner were the only variables retained, as likelihood ratio tests indicated that the inclusion of other variables did not significantly improve the model fit. The analyses yielded similar results when repeated using θ_A and height_A as predictors for θ_B (GAMM_{BA}: *S. corallinus*: $adj. R^2 = 0.413$; *S. doliatus* = $adj. R^2 = 0.462$; *S. puellus* = $adj. R^2 = 0.529$; *S. vulpinus* = $adj. R^2 = 0.327$). In contrast to the angles during foraging, the angles of pair members while swimming showed a clear positive, linear relationship (GLMM_{AB}: *S. corallinus*: $\beta = 0.718$; $t = 22.56$; $P < 0.0001$; *S. doliatus*: $\beta = 0.761$; $t = 28.87$; $P < 0.0001$; *S. puellus*: $\beta = 0.644$; $t = 18.01$; $P < 0.0001$; *S. vulpinus*: $\beta = 0.673$; $t = 23.22$; $P < 0.0001$), indicating that individual angles are non-independent and linearly synchronized during movement, with no other variables

eliciting a significant effect in any of the four species (**Figure 4.4**). Thus, pairs of rabbitfishes travelled together synchronously, but performed contrasting, complementary roles during foraging, which were strongly coordinated with minimal overlap in vigilance behaviour.

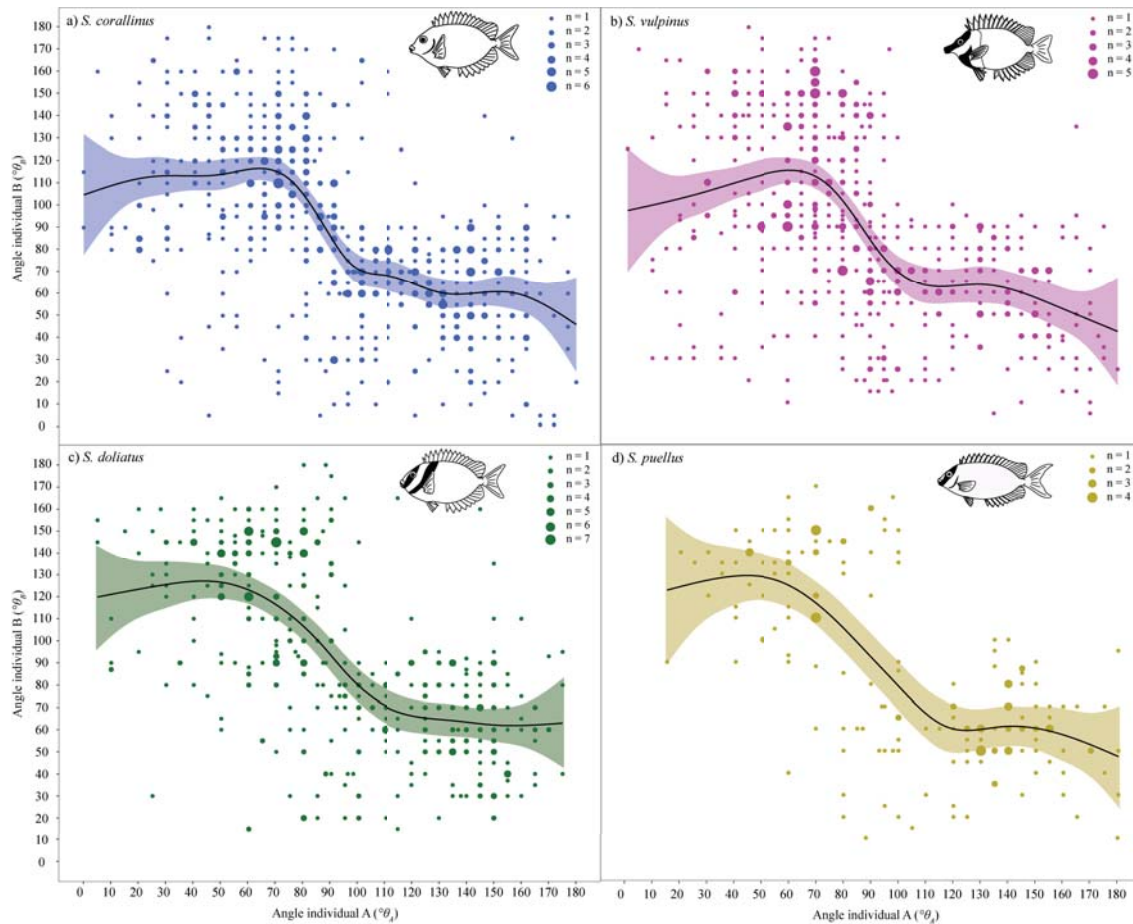


Figure 4.3: Graphical representation of coordination between foraging and vigilance in four species of rabbitfish pairs. Axes mark the angles of individuals in pairs (θ_A , θ_B), while each point represents the angle assumed at a given 5-second point intercept with the size varying according to the number of overlying points. The four predicted smoothed lines mark the predicted fits from generalized additive mixed effects models (GAMMs) and their upper and lower 95% confidence interval. In all species, data are predominantly spread between the upper left and lower right quartile of the plot. The smoothing function, fit by automatic cross-validation, suggests that individuals assume contrasting angles ($<90^\circ$ and $>90^\circ$); however, the extent of the angle is negligible once a certain threshold is reached ($\sim 120^\circ$ and 60° ,

respectively). The observed pattern was consistent among species. a) *S. corallinus* ($n = 15$), b) *S. vulpinus* ($n = 15$), c) *S. doliatus* ($n = 15$), d) *S. puellus* ($n = 14$). $n =$ number of independent pairs represented in the plot.

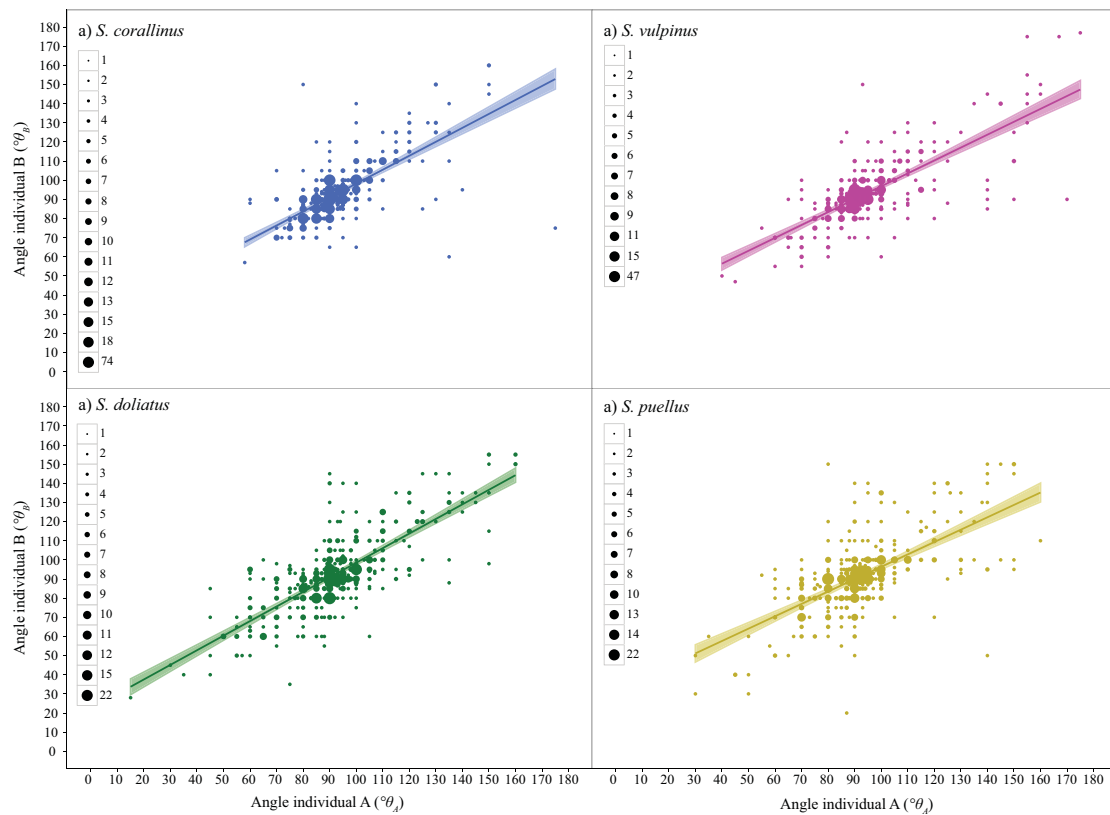


Figure 4.4: Angles exhibited by pair members during swimming behaviour. In all four species, angles are synchronized linearly, with the angle of individual B being the best linear predictor of the angle of individual A. The size of dots marks the number of overlying data points. Data represent repeated observations on 15 independent pairs in a) *S. corallinus*, b) *S. vulpinus*, c) *S. doliatus* and 14 independent pairs in d) *S. puellus*. Trendlines represent the predicted fit ($\pm 95\%$ confidence intervals) from GLMMs performed for each species separately.

Across all pairing rabbitfish species examined, a vigilant individual (i.e., a stationary individual not engaged in foraging activity and exhibiting an angle $>90^\circ$) was present during 82.6% of foraging activity (i.e. when at least one individual was feeding). When feeding, pair members alternated their roles significantly more often than they continued in the same role (**Figure 4.5a**), with a higher proportion of alternated foraging bouts compared to repeated foraging bouts in all species (**Figure 4.5b**; parameter estimates for repeated counts compared to alternated counts: *S. corallinus*: $\beta = -1.538$; $z = -12.6$; $P < 0.0001$; *S. doliatus*: $\beta = -0.826$; $z = -5.61$; $P < 0.0001$; *S. puellus*: $\beta = -1.858$; $z = -7.99$; $P < 0.0001$; *S. vulpinus*: $\beta = -1.094$; $z = -7.70$; $P < 0.0001$). The average deviation from a balanced proportion of feeding events per individual (0.5) was relatively small across pairs in all species (0.14), and within species, the distribution of feeding events was not statistically different from an expected balanced distribution (Pearson's Chi-squared test; *S. corallinus*: $\chi^2 = 2.419$, $df = 1$, $P = 0.120$; *S. doliatus*: $\chi^2 = 2.294$, $df = 1$, $P = 0.130$; *S. puellus*: $\chi^2 = 2.630$, $df = 1$, $P = 0.105$; *S. vulpinus*: $\chi^2 = 3.480$, $df = 1$, $P = 0.062$), although these estimates have to be interpreted with care due to the relatively small sample size.

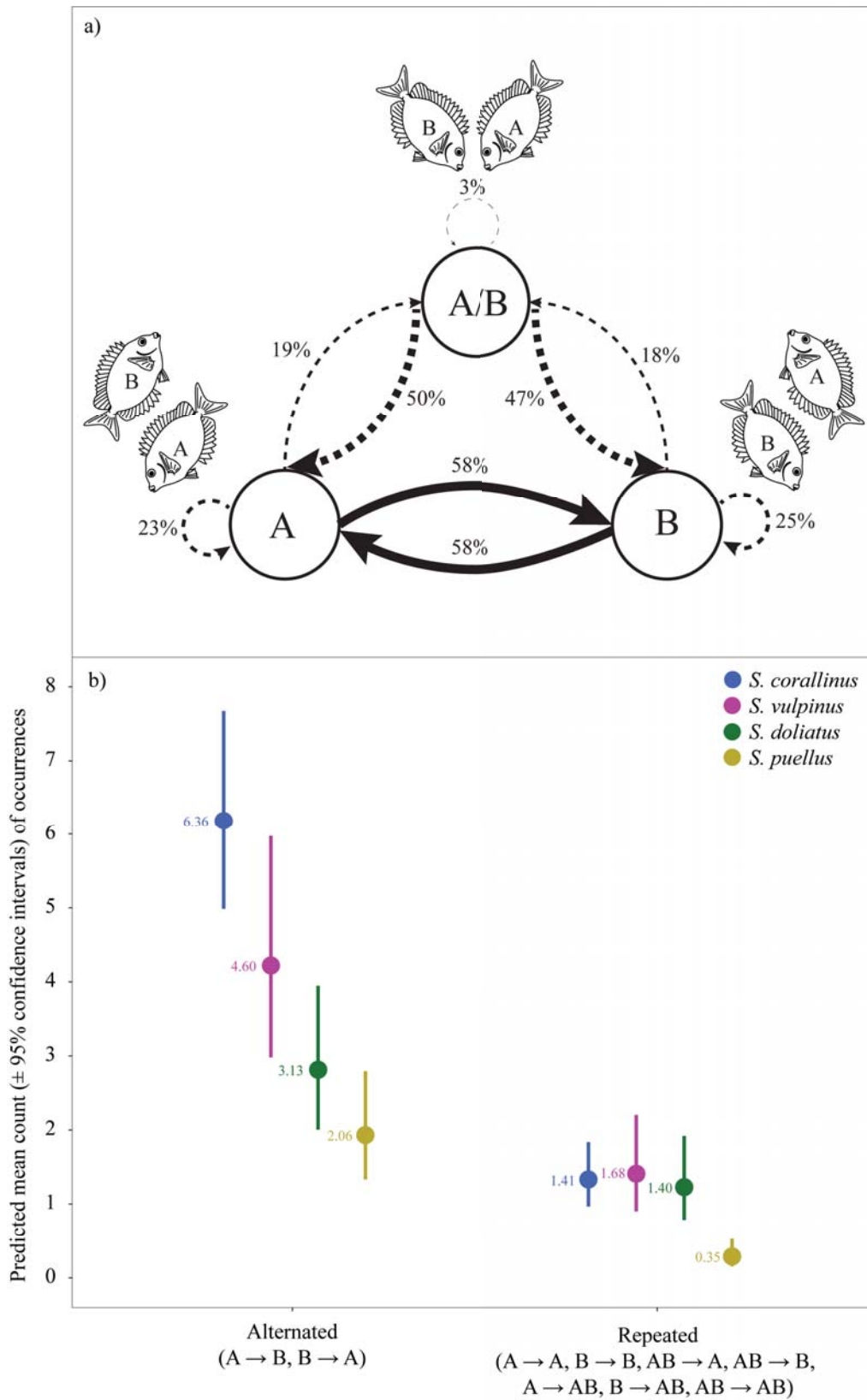


Figure 4.5: Patterns of alternated or repeated foraging bouts in pair members. a) Schematic figure showing the percentage of different pathways pooled for all pairs in all species ($n = 59$). In cases where only one of the pair members is foraging (A or B), a subsequent foraging bout by the partner (B or A) is most common (58% and 58% of all sequential foraging bouts; solid arrows). In contrast, repeated foraging bouts were significantly less common (23% and 25%, respectively; dashed arrows) and so were changes to foraging bouts performed simultaneously by both individuals (19% and 18%, dashed lines). Foraging bouts performed by both pair members simultaneously were almost always succeeded by foraging bouts performed by a singular fish (50%A and 47%B). b) The occurrence of alternated or repeated foraging bouts by pair members in all four species. Circles represent average values for a 12-minute period, predicted from zero-inflated generalized linear mixed effects models (GLMMs) with lines marking the upper and lower 95% confidence intervals. While the average number of both alternating and repeated foraging bouts varied among species as a function of the overall foraging activity, alternated foraging bouts (i.e. individuals taking turns) were significantly more common than repeated foraging bouts in all species (see text for statistical results). Observed values are provided to indicate model fits. Blue = *S. corallinus* ($n = 15$), magenta = *S. vulpinus* ($n = 15$), green = *S. doliatus* ($n = 15$), gold = *S. puellus* ($n = 14$). n = number of independent pairs represented in the plot.

The behaviour of solitary and paired rabbitfish individuals differed in all species (**Figure 4.6**). Compared to solitary rabbitfishes, paired rabbitfishes exhibited significantly longer vigilance bouts except in *S. doliatus*, in which estimates followed the same trend but fell outside the α -level of 0.05 (parameter estimates for solitary individuals compared to paired individuals: *S. corallinus*: $\beta = -0.733$; $z = -4.12$; $P < 0.0001$; *S. doliatus*: $\beta = -0.621$; $z = -1.64$; $P = 0.09$; *S. puellus*: $\beta = -0.934$; $z = -6.37$; $P < 0.0001$; *S. vulpinus*: $\beta = -0.695$; $z = -7.34$; $P < 0.0001$). In all species, paired individuals took significantly more bites per foray (parameter estimates for solitary individuals compared to paired individuals: *S. corallinus*: $\beta = -0.360$; $z = -2.48$; $P = 0.013$; *S. doliatus*: $\beta = -0.792$; $z = -2.29$; $P = 0.022$; *S. puellus*: $\beta = -0.935$; $z = -7.14$; $P < 0.0001$; *S. vulpinus*: $\beta = -0.461$; $z = -1.96$; $P = 0.05$). In *S. corallinus* and *S. doliatus*, paired individuals penetrated deeper into the substratum than solitary individuals and while the same trend was visible in *S. puellus* and *S. vulpinus*, estimates in the latter two species were not statistically significant (parameter estimates for solitary individuals compared to paired individuals: *S. corallinus*: $\beta = -0.578$; $z = -3.99$; $P < 0.0001$; *S. doliatus*: $\beta = -1.665$; $z = -2.58$; $P = 0.0099$; *S. puellus*: $\beta = -0.323$; $z = -1.75$; $P = 0.08$; *S. vulpinus*: $\beta = -0.308$; $z = -1.73$; $P = 0.084$). Overall, differences between solitary and paired individuals were highly consistent among species. All species showed the same trends, varying only slightly in extent. The lack of significance in some variables may have arisen from small sample sizes in solitary individuals, which were both rare and exceptionally easily disturbed.

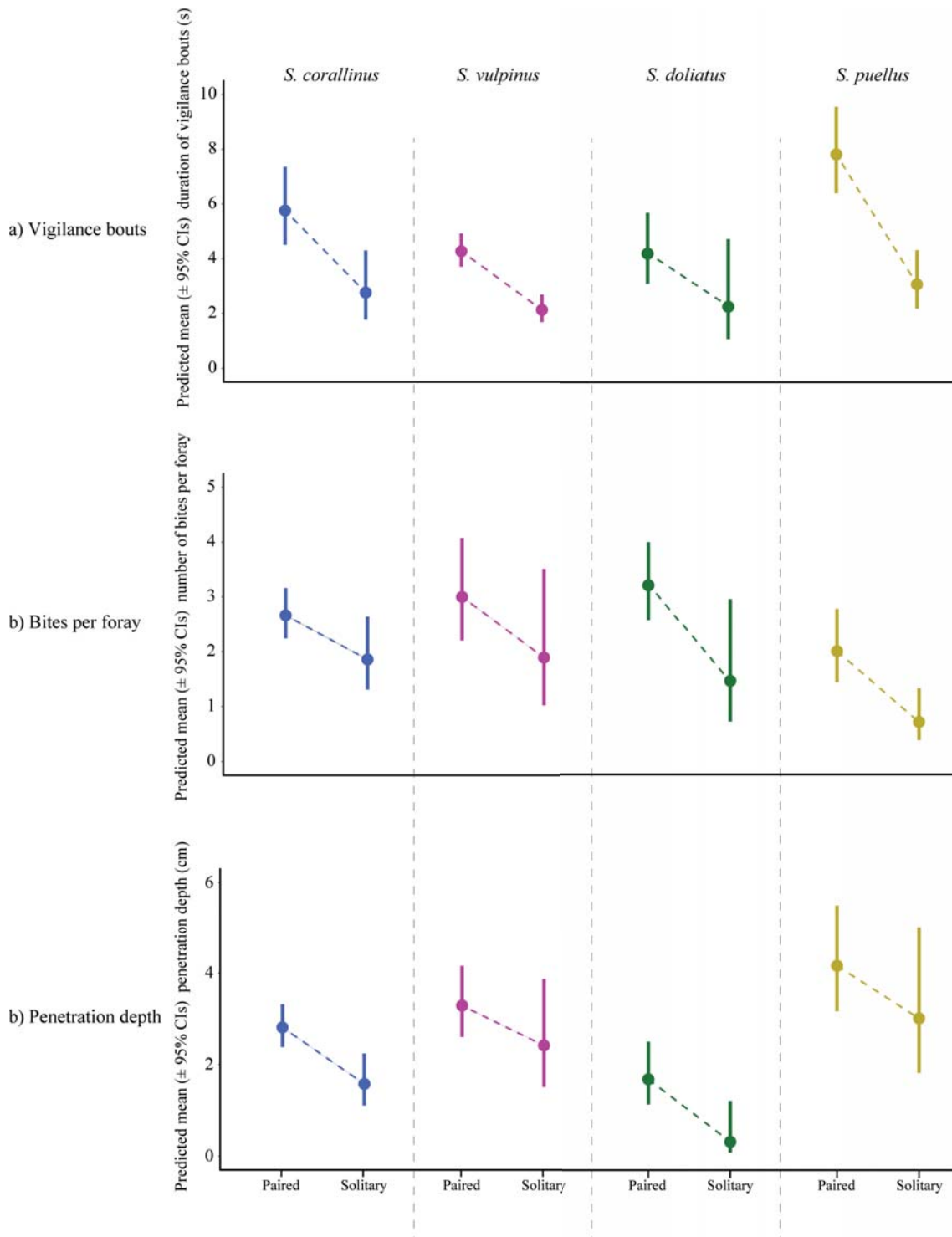


Figure 4.6: Behavioural differences between paired and solitary individuals in all four species. a) Average predicted duration of vigilance bouts (\pm 95% confidence

intervals) in paired and solitary individuals. b) Average predicted number of bites per foray (\pm 95% confidence intervals). c) Average predicted maximum penetration into cracks and crevices in the reef substratum during forays (\pm 95% confidence intervals). All values were obtained from GLMMs performed separately on all four species. Blue = *S. corallinus* (pair $n = 8$; solitary $n = 8$), magenta = *S. vulpinus* (pair $n = 8$; solitary $n = 6$), green = *S. doliatus* (pair $n = 8$; solitary $n = 3$), gold = *S. puellus* (pair $n = 8$; solitary $n = 7$). n = number of independent individuals on which observations were performed.

The relative frequency of fin-flicking differed significantly among behaviours (Pearson's Chi-squared test: $\chi^2 = 231.250$, $df = 3$, $P < 0.001$), occurring significantly more often during foraging activity (i.e. when at least one of the fishes was engaged in foraging) and while displaying to other pairs (aggressive interaction) and hovering, while significantly less fin-flicks were performed during swimming. When one fish was in the vigilance position, while its partner was foraging, almost all observed fin-flicks were produced by the vigilant fish (92.4% of recorded fin-flicks during foraging activity) and subsequent actions (abandonment of current positions, switching positions, chasing) were taken by individuals on average 2.3 ± 0.2 seconds after the fin-flicking, suggesting that feeding individuals may respond to fin-flicks by the vigilant fish. This is further supported by the observation that 61.8% of all observed flight responses were preceded by fin-flicks.

4.4 Discussion

In this study, I provide field-based observational evidence for a coordinated, cooperative vigilance system in four species of pairing rabbitfishes. Specifically, I demonstrate that during foraging, pair members strongly coordinate their positions: while one individual forages with its head down, its partner assumes an elevated, upright position in the water column, allowing for an unobstructed visual field to scan the surrounding environment. Paired fishes alternate frequently between foraging and the vigilance position. Compared to solitary individuals, individuals in pairs exhibit longer vigilance bouts than their solitary counterparts, but appear to benefit from the presence of the partner by exhibiting more consecutive bites per foray and deeper penetration into crevices in the substratum.

The posture assumed by one of the rabbitfishes while its partner is foraging closely matches reports in birds and mammals, where vigilant individuals are commonly identified by raised heads and/or elevated positions (Radford et al. 2009; McGowan & Woolfenden

1989; Favreau et al. 2010; Pays et al. 2009), a behaviour that has only recently been reported for teleost fishes (Fox & Donelson 2014). For rabbitfishes, the upright position is likely to favour vigilance as it potentially enables a greater ability to detect predators (and possibly also competitors) compared to foraging fishes, probably due to a less obstructed visual field (Fernández-Juricic et al. 2004; Arenz & Leger 1997). The assumed angle may allow an unobstructed field of view while remaining close to the reef and the partner. While I am unable to demonstrate that the upright position serves primarily for the detection of predators, my observations and previous evidence suggest that scanning the surroundings for competitors, potential new partners, or food are of limited importance when compared to predator detection. While rabbitfishes do occasionally engage in aggressive behaviour with other pairs, these interactions are infrequent (<1% of the total behaviours observed in this study) and home-ranges of pairs are non-exclusive (Brandl & Bellwood 2013b), suggesting that scanning the surroundings for competitors may only play a minor role. In addition, pair bonds between rabbitfishes are relatively stable (Brandl & Bellwood 2013a,b), questioning the need to continuously look out for a new partner. Finally, the foraging strategy of rabbitfishes along with their dietary preferences for small and cryptic algae, sponges and cyanobacteria (Brandl & Bellwood 2014a; Hoey et al. 2013), which require careful and close examination of concealed micro-habitats, make it unlikely that an elevated position will be beneficial for the detection of food. However, as I am unable to quantify the relative contributions of these various roles, I use the term ‘vigilance’ in its widest sense as being aware of the surrounding environment.

Whenever the pair was observed to rapidly abandon its positions (i.e. engaged in rapid directional swimming behaviour), the vigilant individual had an unobstructed view of the surrounding environment, while the visual field of the forager was often blocked by the reef substratum. Upon the vigilant individual abandoning its position (which was far more

frequent than the forager abandoning its position first), the forager consistently trailed the vigilant individual, suggesting that foraging individuals reliably (94.1% of all cases) respond to actions or potential warning cues generated by the vigilant individual. While peripheral vision and social monitoring by the forager may facilitate the reaction to the vigilant partner's behaviour (Fernández-Juricic et al. 2005), the frequent occurrence of fin-flicks prior to abandoning the vigilance position may indicate intentional communication (Hollén et al. 2008; Wickler 1985; Manser 1999). Although the significance of fin-flicks has not yet been investigated in rabbitfishes, fin-flicks are known to generate an acoustic signal in another family of reef fishes in which pairing is prevalent (f. Chaetodontidae; Tricas et al. 2006; Parmentier et al. 2011) and fin-flicks have been described as an important warning signal in other fish species (Brown et al. 1999; Murphy & Pitcher 1987). Thus, although the role of fin-flicking remains to be determined in rabbitfishes, it seems likely that fin-flicks serve as a communication signal and that the forager is able to perceive these signals despite visual restrictions. My findings that the vast majority of fin-flicks in rabbitfishes occurred in situations in which communication is beneficial (i.e. when one fish was foraging while the partner was vigilant or while displaying to other pairs) support a role of fin-flicks in the maintenance of coordination between the forager and the vigilant fish.

Interestingly, the described scenario, in which foraging severely compromises vigilance while information is readily available from a vigilant conspecific, precisely matches the conditions under which coordinated vigilance should be favoured (Sirot 2012; Fernández-Juricic et al. 2004). This is strongly supported by the angles rabbitfishes assume during foraging episodes where one fish's angle and height above the substratum are the best predictors for complementary behaviour in the partner (**Figure 4.3**). Clearly, pairs of rabbitfishes coordinate their positions during foraging and possibly do so through communication via fin-flicks.

Given this, the question then arises whether the coordinated behaviour in rabbitfish pairs represents a cooperative system based on by-product mutualism/pseudo-reciprocity (Connor 1995, 2010), or if rabbitfishes may satisfy the requirements of direct reciprocity. Several recent papers have emphasized the lack of evidence supporting the fundamental characteristics of direct reciprocity in animals (Clutton-Brock 2009; Connor 2010; Melis & Semmann 2010). These include: i) continuous cooperation between the same individuals, ii) behavioural adaptations to assist the partner, iii) adjustment of assistance provided according to received assistance, iv) cooperation not restricted to kin or potential mates, v) assistance entailing momentary net fitness costs to the assisting individual, and vi) cooperative behaviour being found in wild populations (after Clutton-Brock 2009).

For rabbitfishes, there is evidence supporting all of these requirements. (i) As rabbitfish pairs are stable and remain together for extended periods of time (Woodland 1990; Brandl & Bellwood 2013b), cooperation is likely to occur continuously between the same individuals. (ii) My results show that individuals prolong the length of vigilance bouts in the presence of a partner. This may represent a behavioural adaptation to assist the partner, as paired fishes exhibited an increased number of bites per foray (*S. doliatus*, *S. puellus*, *S. corallinus*) or deeper substratum penetration (*S. corallinus*, *S. doliatus*), therefore increasing the likelihood of a higher overall food intake (as reported for pied babblers *Turdoides bicolor*; Hollén et al. 2008). (iii) Pair members frequently alternate between foraging and vigilance and the ratio between assistance provided and assistance received appears to be well balanced. While feeding was unevenly distributed in a few of the observed pairs, this may be a consequence of the length of observations, and a more balanced distributions may be observed if fish were monitored over an entire day. (iv) The observation that feeding within pairs is not generally skewed toward one individual suggests that cooperation is not solely based on male mate-guarding (as found in sleepy lizards, *Tiliqua rugosa*; Bull &

Pamula 1998), and the common occurrence of same-sex pairs in rabbitfishes (Brandl & Bellwood 2013b) suggests that cooperation is not limited to reproductive pairs (Annett et al. 1999). Due to the reproductive strategies of reef fishes and their pelagic larval stage, cooperation limited to kin is also highly unlikely.

(v) Fitness costs may include predation risk and lost foraging opportunities during vigilance behaviour but such costs are inherently difficult to quantify (Carter 2014; Bednekoff 1997). As in other animals where mutual vigilance has been described as a potential cooperative system (Radford et al. 2009; Hollén et al. 2008; Wright et al. 2001; Ridley et al. 2013), the vigilant individual in rabbitfish pairs is positioned above the underlying substratum with its head elevated, making vigilance beneficial for the overall awareness of the surroundings and therefore potentially self-serving rather than costly (provided predators selectively target foraging individuals). However, given the nutritive constraints of marine herbivory (or spongivory), which necessitate constant and intensive grazing, prolonged vigilance bouts are probably nutritionally costly rather than self-serving (Choat & Clements 1998). This is further supported by the low levels of simultaneous vigilance (both pair members hovering motionless above the substratum, 6.6% of behaviours across all pairs), which would indicate competition for vigilance in a scenario where predators preferentially target foragers (Sirot 2012; Bednekoff 1997; Sirot et al. 2009; Clutton-Brock et al. 1999). Thus, while gaps in foraging activity associated with prolonged vigilance are likely to represent a significant cost, the deeper penetrations into crevices, as well as the higher number of bites per foray appear to be an intuitive reward for the partner. The lack of a clear difference between solitary and paired individuals in *S. puellus* and *S. vulpinus* in terms of penetration depth may point towards differences in the dietary preferences of these two species or their morphological adaptations. While *S. corallinus* and *S. doliatus* feed predominantly on filamentous and corticated red algae, food that is readily

used by other herbivorous fish species (Hoey et al. 2013; Choat et al. 2002), only a few reef fishes feed on cyanobacteria, which are the main food source for *S. vulpinus*. Thus, the latter might be more readily available in more accessible microhabitats. In addition, *S. vulpinus* exhibits the morphological characteristics most suited for the exploitation of crevices among the four investigated species, suggesting that even solitary individuals might be able to penetrate into the substratum with no substantial obstructions to the visual field (Brandl & Bellwood 2013a). *S. puellus*, in turn, exhibits the most fusiform morphology among the examined species (Brandl & Bellwood 2013a), suggesting that quick escape from predators might play a bigger role in this species, possibly permitting solitary individuals to penetrate into the substratum despite the lack of a vigilant partner. However, given the relatively small sample size, the lack of significance in these comparisons should be interpreted with caution. These minor differences notwithstanding, there appear to be clear costs (vigilance bouts) and benefits (foraging efficiency) associated with cooperative vigilance in all four rabbitfish species examined. Finally, vi) all observations were conducted on the reef, indicating that the described vigilance system occurs in wild populations. Thus, the coordination of foraging and vigilance in rabbitfish pairs appears to satisfy all the basic requirements for reciprocal cooperation.

While I cannot hope to fully resolve the question of reciprocity with observational evidence alone, my findings are consistent with direct reciprocity. Thus, my study corroborates the tenor of several recent studies, which posit that reciprocity may be a lot more common under natural settings than when forced into the stringent rules of game-theoretical models (Taborsky 2013; Raihani & Bshary 2011; Brosnan et al. 2010; Carter 2014; van Doorn 2014). Indeed, my results help us to understand why we may find reciprocity in animals, which lack the presumed cognitive requirements for reciprocity (Clutton-Brock 2009). First, cooperative interactions among rabbitfishes are restricted to just

one partner at a time. This alleviates frequently-cited cognitive issues arising from recognizing a range of individuals and remembering their previous actions in a large group of animals (Milinski et al. 1990; Stevens & Hauser 2004; Milinski & Wedekind 1998) in order to repay for the behaviour of a previous partner, as has been posited for predator inspection and egg-trading in fishes (Clutton-Brock 2009; Connor 1992, 1996). Second, continuous foraging activity, immediate alternation, and the similar and simultaneous needs (food and safety) for both pair members in rabbitfishes prevent long time-lags between rounds in an IPD, as often found in, for instance, primates (Brosnan et al. 2010; De Waal & Brosnan 2006). This again relaxes the need for complex cognitive abilities, such as memory, to underpin reciprocal cooperation (Rutte & Taborsky 2008). Third, rabbitfishes cooperate continuously with the same partner over an extended period of time. There is now considerable evidence that such interdependence between social partners can foster high levels of cooperation in an IPD, as individuals do not systematically surrender to the short-term temptation of cheating on the partner (Roberts 2005; St-Pierre et al. 2009), therefore making tit-for-tat (or, more specifically 'generous tit-for-tat') a strategy with high levels of cooperation. Finally, my study provides preliminary evidence for the continuous exchange of social information between partners by means of fin-flicks and suggests that individuals quickly react to the behaviour displayed by the partner, as indicated by high levels of coordination (Sirot 2012). Such elimination of discrete rounds in the IPD and the continuous exchange of information have been proposed as a major factor in favouring cooperation (either by coercion or reciprocity), as it lowers the cost for cooperating individuals (van Doorn et al. 2014). Given the low levels of simultaneous vigilance observed in the present study, reciprocity appears more likely to operate in rabbitfishes than coercion, which would incline individuals to simply copy the partner's behaviour (van Doorn et al. 2014).

In summary, my study identifies pairing rabbitfishes as an intriguing group of animals in the context of reciprocity, cooperation, and cognition. Although limited to observational data, I provide strong evidence for a clear coordination of foraging and vigilance behaviour in pairs and demonstrate that pair members frequently alternate their foraging. I further show that rabbitfish pairs have the potential to satisfy all the basic requirements of reciprocity and discuss a range of conditions, which may favour reciprocal exchange in animals. My evidence suggests that the complex cognitive and social skills, frequently assumed to be necessary for the evolution of direct reciprocity in animals, may be advantageous but, as in fishes, may not be essential.

Chapter 5: Individual-based analyses reveal limited functional overlap in a coral reef fish community.

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5.1 Introduction

All over the world, ecosystems are changing. Human domination is altering both the biotic diversity and habitat structure of ecosystems, leading to fundamental changes in ecosystem processes (Chapin et al. 1997; Hooper et al. 2005). This can ultimately result in the loss of goods or services provided by an ecosystem (Vitousek et al. 1997). Biodiversity in all its facets (e.g. taxonomic diversity, phylogenetic diversity and functional diversity) is accepted as a key factor for the persistence of ecosystems (Duffy et al. 2003; Hooper et al. 2005; Gamfeldt et al. 2008). Functional diversity, in particular, has been the focus of many recent studies evaluating stability or changes in ecosystem processes (Petchey & Gaston 2006; Mouillot et al. 2012). In this context, the functional role of a species is frequently treated as an equivalent to its ecological niche (Halpern & Floeter 2008; Mouillot et al. 2011).

The ecological niche of a species is commonly described as the volume occupied in a multidimensional hyperspace, symbolizing the ecosystem (Hutchinson 1957; Whittaker et al. 1973; Devictor et al. 2010). If this volume is based on a species' impact on the ecosystem, the niche is usually described as a species' *functional niche* (Hutchinson 1957; Rosenfeld 2002). Beyond this, the classification of a species' niche can be extended to its fundamental or realized niche. The *fundamental niche* refers to the potential volume along all axes that a species could theoretically occupy, if species-specific, intrinsic characteristics, such as morphological or physiological attributes, were the only determinants. In contrast, the *realized niche* is the volume a species actually occupies when constrained by biotic and

abiotic factors (Hutchinson 1957; Whittaker et al. 1973; Devictor et al. 2010). Most studies that seek to evaluate species' ecosystem roles use a functional-diversity framework with multiple phenotypic traits from which a classification of the respective species niches are derived (Petchey & Gaston 2006; Villéger et al. 2008; Villéger et al. 2011), thus measuring species' fundamental niches (Hutchinson 1957; McGill et al. 2006; Wright et al. 2006). Realized niches, however, can only be assessed using data on the species *in situ*, where restricting abiotic or biotic agents are operating. Unfortunately, there is a notable lack of empirical studies that incorporate field observations to determine the realized functional niche of species across communities, particularly in high diversity marine ecosystems (Hooper et al. 2005).

Among marine systems, coral reefs harbour exceptional diversity but are also subject to intense human pressure (Vitousek et al. 1997; Hooper et al. 2005). Despite providing vital goods and services to millions of people, human impacts are progressively altering the biodiversity, structure and processes on coral reefs (Hughes et al. 2003). One critical functional process on coral reefs that is under threat is the removal of algae by herbivorous fishes (Bellwood et al. 2004; Nyström 2006). Herbivorous species are traditionally classified into distinct functional groups according to their taxonomic affiliation (e.g. families Acanthuridae, Labridae, Siganidae), dietary preferences (e.g. macroalgae, turf algae, detritus) or their way of acquiring food (e.g. scraper, excavator, grazer) (Nyström 2006; Green & Bellwood 2009; Cheal et al. 2012). Using these classifications, many studies have investigated links between herbivorous species assemblages and the health or resilience of coral reefs (Burkpile & Hay 2008; Green & Bellwood 2009; Cheal et al. 2010). However, these classification schemes are inherently problematic.

While facilitating a broad overview of the functional composition of a community, functional-group approaches have several shortcomings leading to a significant loss of

information. Many studies measure only the fundamental niche and, to date, most studies only consider *which species* feeds on *what* and *how* they do it (Nyström 2006; Johansson et al. 2013), while other components such as *where* species feed are rarely accounted for (but see Fox & Bellwood 2013). Furthermore, existing classification schemes are often equivocal and consensus about functional group membership has yet to be achieved (Green & Bellwood 2009; Cheal et al. 2012). Most importantly, arranging species in a broad classification scheme discounts variability among individuals, a widely neglected but potentially important component of biodiversity that can contain valuable information on species' functional niches (Messier et al. 2010; Violle et al. 2012).

To address some of these issues, I compare the realized functional niches of herbivorous coral reef fishes, with regards to fine-scale spatial characteristics of their functional impact (i.e. *where* the fish feed). I present a novel way of assessing the realized functional niche of species that uses behavioural data, and takes individual variation into consideration, to examine patterns of redundancy and complementarity in functionally important species. Based on field data on the feeding behaviour of 21 herbivorous coral reef fish species, I use a recently developed multidimensional framework (Villéger et al.; Villéger et al. 2011; Mouillot et al. 2012) to provide estimates of the functional impact of each species (the volume occupied in functional niche hyperspace) and the functional overlap between species (the volume shared between species in this hyperspace). I offer this method as an alternative way of assessing the functional niche of species that allows for a more nuanced assessment of the functional role of species beyond traditional functional-group frameworks. Specifically, the objectives of my study were to: 1) develop a multidimensional framework to examine functional niche overlap between species using a high-diversity herbivorous reef fish community as a model, 2) compare the results of my analyses to traditional classification schemes to assess the sensitivity and applicability of the analyses, 3) provide a novel,

continuous rather than categorical representation of the functional niches of herbivorous reef fishes based on their microhabitat utilization patterns and 4) evaluate the ecological implications of these findings.

5.2 Materials and methods

5.2.1 Data collection


The study was conducted during April and May 2012 on Lizard Island (14°40'08"S 145°27'34"E), a granitic mid-shelf island in the northern Great Barrier Reef (GBR). Lizard Island has a rich herbivorous fish fauna, encompassing the vast majority of herbivorous fishes found on the GBR (Cheal et al. 2012). Behavioural observations were made on the crest and upper slope (1-8 m) of two fringing reefs on the sheltered side of the island, in Mermaid Cove and off Turtle Beach (**Figure 5.1**).



Figure 5.1: Lizard Island and the two sample sites, Mermaid Cove and Turtle Beach. Sites are located on the sheltered side of the island.

Observations were carried out on adult individuals of 21 species from three families (Acanthuridae, Labridae, Siganidae). The 21 species represent the majority of roving herbivores present on these reefs. No other roving herbivorous species were sufficiently abundant on the sampled reefs to be included in the analyses. A haphazardly selected individual was followed until it took a bite from the reef. I then recorded the surface type (open horizontal [A], open vertical [B], underside [C], concealed vertical [D] or concealed horizontal [E]) and the substratum type (branching coral, encrusting coral, massive coral, dead coral skeleton, macro-algae, coral rubble, soft coral, sand, sponge, turf covered reef matrix or other substratum types such as giant clams). Surfaces were considered A or B if they were exposed, planar microhabitats, C was on the underside of overhanging surfaces, while surfaces D and E were recorded if herbivores fed in concavities that required the penetration into holes between 1 and 30 cm across in the substratum. These latter microhabitats ranged from small holes, crevices or fissures in the reef matrix to the microhabitat between the branches of branching, corymbose corals. A further criterion for surfaces D and E was that the concavity depth had to equal or exceed its smallest width. I also recorded the extent of penetration into the substratum exhibited by individuals, distinguishing between feeding in concavities with its snout (from the premaxilla tip to the anterior rim of orbit) or with its body (from anterior rim of orbit to the caudal fin margin) (Fox & Bellwood 2013). Vernier callipers were used to measure the penetration depth. To minimize the likelihood of recording the same individual within a 24 h period, reefs were sampled on alternating days and different sampling areas of the reef were used throughout the day. Observations were discarded if fish exhibited a detectable response to the observer. To ensure consistent recordings, all data were collected by two observers throughout the study. Overall, a total of 1,734 observations were recorded (with a mean of 82.5 observations per species) (**Table 5.1**).

Table 5.1: Microhabitat utilization patterns of herbivorous reef fishes. FG = Traditional functional group affiliation; GD = grazer/detritivore; AB = algal browser.

Family	Species	FG	Surface						Substrate						Penetration			
			A	B	C	D	E	DC	MA	SD	RB	TR	$\sum_{i \neq j} (Others)$	P _B %	P _S %	P _S x (mm)		
	<i>A. nigricauda</i>	GD	89	6.5	0	3.8	0.8	1.2	0	47.3	12	38.5	1	0	0	0	0	
	<i>A. nigrofuscus</i>	AB	69.6	18.3	0	4	8.1	19.5	2.5	0	17.3	59	1.7	0	0	17.5	18.6	
	<i>A. olivaceus</i>	GD	85.1	14	0	0	0.9	0	0	21.1	10.7	68.2	0	0	0	0	0	
	<i>C. striatus</i>	GD	66.3	32.3	0	1.3	0	12.6	3.4	4.8	5.5	73.6	0	0	0	0	0	
	<i>N. unicornis</i>	AB	45	41.7	0	1.7	11.7	0	0	17.2	0	71.1	0	0	0	0	7.5	23.3
	<i>Z. scopas</i>	AB	26.1	23.5	0.6	24.4	25.4	21.5	0	0	0.4	77	1.1	1	40	37.1	17.8	
	<i>Z. veliferum</i>	AB	34.9	30.6	2.3	20	12.2	14.7	0	0	8.4	75.5	1.4	0	0	29.2	14.4	
Labridae	<i>C. microrhinos</i>	EX	64	36	0	0	0	14	0	0	3.7	82.4	0	0	0	0	0	
	<i>C. sordidus</i>	EX	72.6	18.7	2.3	5.6	0.8	34.1	0	0	11.2	52.2	2.5	1.2	300	2.4	25	
	<i>S. dimidiatus</i>	SC	74.9	20.7	0	0	4.4	19.6	0	0	9.9	70.5	0	1.8	100	1.8	30	
	<i>S. frenatus</i>	SC	59.3	37	0	1.6	2	17.5	0	0	6.9	75.6	0	0	0	3.1	20	
	<i>S. niger</i>	SC	52.4	32.9	5.7	6	3.1	25.1	0	0	6	65	4	2.4	85	3.7	16.7	
	<i>S. oviceps</i>	SC	65.3	32.7	0	0	2	29.3	0	0	2	65.3	3.4	0	0	1.5	30	
	<i>S. rivulatus</i>	SC	69.2	25.4	0	3.5	2	7.5	0.8	2.1	5.4	84.2	0	0	0	2.6	35	
	<i>S. schlegeli</i>	SC	67.7	25.1	0.9	3.7	2.5	3.8	1.1	15.8	10.4	68.5	0.5	1.7	60	3.4	25	
Siganidae	<i>S. argenteus</i>	AB	63.6	22	0	3	11.4	7.5	20.8	1.2	21.4	49.1	0	0	0	15.5	16.7	
	<i>S. corallinus</i>	AB	7.9	8.7	2.2	30.7	50.5	46.6	1.5	0	22.2	28	1.8	11.3	106.4	70.7	24.9	
	<i>S. dotiatus</i>	AB	25.2	16.2	1.3	29.2	28.1	31.1	4.2	1.1	14	48.9	0.7	8.3	160.7	45.3	20.5	
	<i>S. puellus</i>	AB	24.5	22.8	5	31	16.6	14.6	8.5	4.4	11.5	52.4	8.5	23.5	130.4	29.6	23.4	
	<i>S. punctatus</i>	AB	27.9	20.8	1.5	25.3	24.5	33.5	7.1	0.9	11.7	43.1	3.7	9.2	70	33.7	19.4	
	<i>S. vulpinus</i>	AB	13.2	6	2	21.2	57.6	67.4	0	0	13.2	15.4	3.9	5.3	87.1	74	31.1	

5.2.2 *Individual-based analyses of species' realized functional niches*

To assess the realized functional niche of species, a multidimensional functional niche space can be created using the behavioural observations made on individuals. In this study, I used the number of bites taken by an individual during a feeding foray, the surface position, substratum type and the extent of body- or snout-penetration. As behavioural observations often include data of different types (such as categorical and numeric data), I used Gower's metric (Gower 1966) to calculate a distance matrix that serves as the basis for an ordination. Based on a Gower's distance matrix, a principal co-ordinate analysis (PCoA) can be performed using appropriate corrections for negative eigenvalues (Legendre & Legendre 1998). PCoA is a useful way of representing patterns from pairwise distance measures and is designed to function with multiple distance metrics. The values yielded by the PCoA serve as synthetic trait values (i.e. new trait values based on the relative importance of behavioural traits in the initial dataset). These new synthetic trait values are then used to establish the distribution of individuals in multidimensional space (Laliberté & Legendre 2010), thus creating a synthetic functional niche hyperspace (i.e. a multidimensional space that encompasses all possible trait combinations), from which individual coordinates can be extracted.

Based on these individual coordinates, a species' functional niche volume can be calculated that encompasses all individuals of a species within a given ecosystem (Cornwell et al. 2006; Villéger et al. 2011). The niche volume (a convex hull volume) is delimited by the individuals with the most extreme positions in multidimensional space (called vertices). In practice, this means that species with large niche volumes have vertices that are widely dispersed in multidimensional space, suggesting that individuals in this species exhibit a high degree of variation in their realized functional niche. In contrast, the niche volume is small

when individuals of the species deliver only a restricted range of functions. In my study, this would equate to the use of a limited range of microhabitats.

The overlap among species' functional niche volumes can also be calculated. To describe distances between species in assemblages, the term "functional dissimilarity" has been proposed (Villéger et al. 2011). For coral reef fishes, recent studies stress the concept of complementary functions for herbivorous fishes (Burkepile & Hay 2008, 2011; Rasher et al. 2013). I therefore use the term functional complementarity as a measure of the overlap between species' volumes in functional niche space. Functional complementarity is the inverse of functional redundancy (species with high overlap have high redundancy but low complementarity and *vice versa*) and can be expressed as the percentage of overlap between the niche volumes of two species. This is calculated as the ratio between the volume of the overlap (intersection) and the overall volume of the respective niches. Thus, if individuals in two species implement the same functions, the volume shared by those two species would be expected to be large (> 50% overlap). This can be considered functional redundancy. In contrast, little or no overlap between species provides a measure of complementarity (< 50% overlap).

In addition, the effect of niche volume on the overlap of species should be considered. Specifically, a small niche volume can only have limited overlap with another volume. Thus, it is desirable to know whether little overlap between two species is the result of two large volumes overlapping only marginally (species A and B) or a consequence of a small volume (species C) being entirely or partially nested within a larger volume (species D). In both cases, the volume of overlap is small, yet the ecological consequences are fundamentally different. In the first case, individuals of the two species perform highly dissimilar functions and only few individuals are located in the same area of multidimensional space. In contrast, in the second case, most individuals of species C are located within the volume occupied by

species D but perform only a small fraction of the functions executed by individuals of species D, which, in turn, performs all functions administered by species C. This distinction has originally been proposed for decomposing β -diversity, where the respective contributions of either scenario have been termed “turnover” (the scenario of species A and B) and the “nestedness” (C and D) components (Baselga 2010, 2012; Villéger et al. 2013). Using the distribution of individuals in functional niche space and the resulting niche volumes, the contribution of the turnover and nestedness components to the functional niche overlap between species can be calculated (Villéger et al. 2013).

5.2.3 Realized functional niches and complementarity in herbivorous coral reef fishes

Using Gower’s distance metric and a PCoA, I calculated the realized functional niche volumes of 21 herbivorous reef fish species based on their microhabitat utilization patterns during foraging. To account for different sample sizes, subsamples of 30 individuals were used to calculate functional impacts using 999 randomized permutations within each species. The niche volume of each species was then standardized against the total volume of niche space occupied by all species. While this provides insights into the functional impact of a species, it is also desirable to know whether or not the volume of a species is significantly lower than expected based on the total niche space volume. To test this, I used randomized permutation testing (Manly 1997), running 999 randomized iterations where 30 individuals were chosen from the whole community regardless their taxonomic identity (to represent hypothetical species). The volumes occupied by individuals of the focal species were then compared to the distribution of the hypothetical species volumes, to determine if the volumes of observed species differ significantly from hypothetical species (based on a random draw from individual values across the entire community).

I also calculated the overlap between species' niche volumes. As the overlap is highly dependent on the extent of niche volumes, values were standardized against the total volume occupied by the two species (Villéger et al. 2011). These values were then compared to values yielded by 999 randomized iterations to evaluate if the calculated overlap differs significantly from a null expected value based on random volumes in total niche space. To construct a null-model, two sets of 30 individuals (hypothetical species), drawn randomly from the total pool, were used to create an overlap between two niche volumes (i.e. overlap between two hypothetical species, again standardized against the total volume of the two hypothetical species volumes). To avoid bias toward increased overlap between the two hypothetical species, the permutations were coded to automatically exclude individuals that have been assigned to one of the two hypothetical species.

Using pairwise overlaps of the 21 focal species, a non-metric multidimensional scaling (nMDS) analysis and Ward's hierarchical clustering analysis were performed to visualize the obtained pairwise distance matrix. Finally, I calculated the contribution of the turnover and nestedness component to the overlap between species following the equations given by Baselga (2012) and Villéger et al. (2013) to account for the different sizes of niche volumes among species.

All analyses were carried out on four dimensions of the functional niche hyperspace as a result of a trade-off between computation time and the percentage of variation within the original dataset expressed on these four dimensions (69.8% explained by four dimensions). Statistical computations were performed using the packages *ape*, *cluster*, *geometry*, *rcdd* and *vegan* as well as the functions *CHVintersect* and *betapart* (Villéger et al. 2011, 2013; Baselga 2012) in the software R (R Development Core Team 2012), using the HPC unit at James Cook University, Townsville, Australia.

5.3 Results

5.3.1 Realized functional niche volumes

The functional niche volumes of the 21 species varied markedly. Volumes varied from just 0.8% of the total niche space in the excavating parrotfish *Chlorurus microrhinos* to 58.8% in the rabbitfish *Siganus punctatus* (**Figure 5.2**). The functional volume of *Chlorurus microrhinos* was thus only 1.4% of the volume occupied by *Siganus punctatus*. Aside from *S. punctatus*, nine other species had niche volumes that were not found to be significantly different from expected volumes (i.e. compared to hypothetical species based on a random draw; randomized permutation testing: $P > 0.05$). These species included all rabbitfishes (*Siganus*) and surgeonfishes in the genera *Zebrasoma* and *Naso*, as well as *Acanthurus nigrofuscus*. The functional niche volumes of all other species were significantly lower than expected under the null model. This was highly significant ($P < 0.001$) in the grazing, detritivorous surgeonfishes in the genus *Acanthurus* and in *Ctenochaetus striatus*, as well as the parrotfishes *Chlorurus microrhinos* and *Scarus oviceps*. Moderate significance levels ($P = 0.01 - 0.001$) were found for all scraping parrotfishes (*Scarus*, **Figure 5.2**).

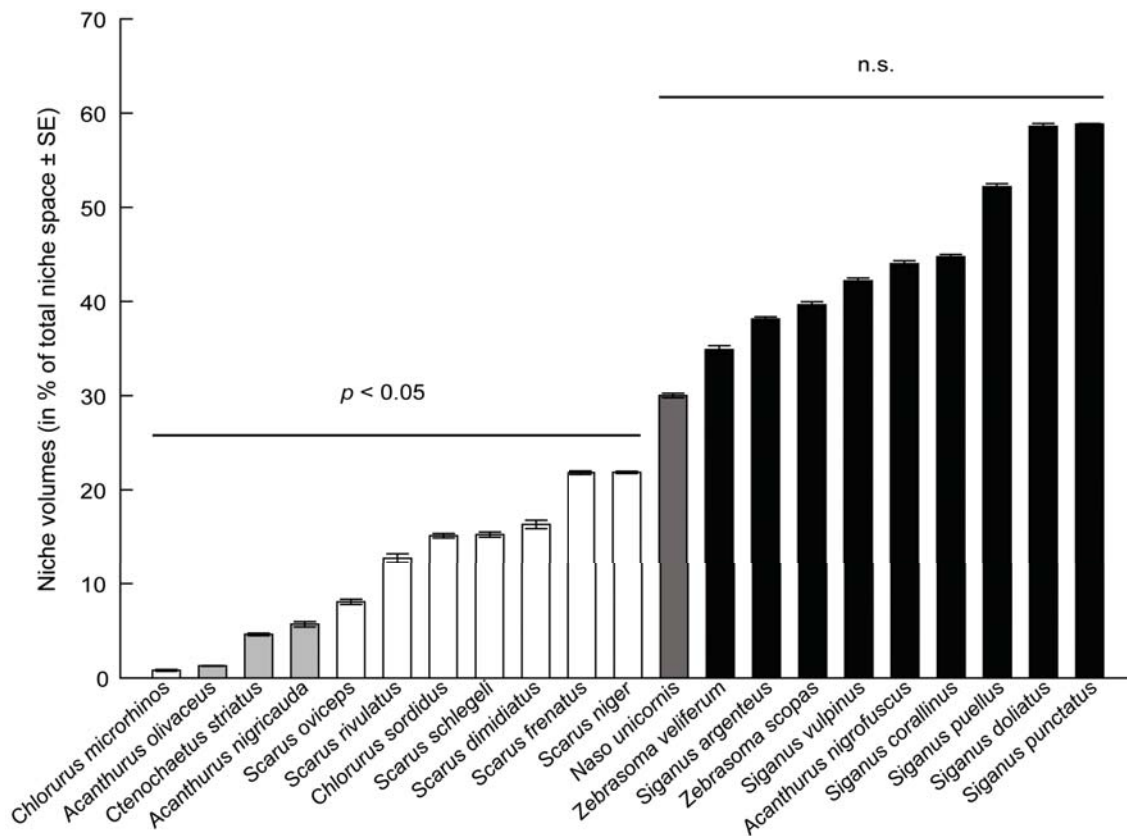


Figure 5.2: The relative volumes of herbivorous fishes in functional niche space based on feeding microhabitat preferences. Bars mark the percentage of niche space occupied relative to the total niche space available. Niche volumes with $p < 0.05$ are significantly lower than expected, based on hypothetical volumes (drawn randomly from the total niche space; $n = 999$). Black bars = croppers; dark grey = browsers; light grey = detritus feeders; white = scrapers/excavators.

5.3.2 Functional redundancy vs. complementarity

Based on the occupied niche volumes of species, the overlap between all species pairs was calculated, supplying a measure of pairwise, functional complementarity (i.e. a pairwise, functional dissimilarity matrix; Table **5.2, 5.3**). There was extensive complementarity among herbivorous fish species. The mean overlap of functional niche volumes among species was just $15.2\% \pm 0.1$ SE, and the highest overlap between two species amounted to only 42.6% between the rabbitfishes *Siganus doliatus* and *Siganus punctatus*. Based on my 50% cut-off, no species pair displayed redundancy in microhabitat utilization.

Table 5.2: Percentages of non-overlapping volumes between species of herbivorous species resulting in a pairwise dissimilarity matrix. Note that values indicating zero would have to be interpreted with caution as zero overlap on a single dimension may mask existing ecologically important overlap on other dimensions.

	Agri	Agro	Aoli	Cstr	Scor	Sdol	Stus	Sdim	Sriv	Ssch	Snig	Sovi	Sfre	Zsco	Zvel	Svul	Spue	Sarg	Nuni	Cmic	Csor
Agri																					
Agro	0.917																				
Aoli	0.825	0.977																			
Cstr	0.897	0.908	0.941																		
Scor	0.995	0.677	1.000	0.970																	
Sdol	0.932	0.677	0.995	0.978	0.622																
Stus	0.981	0.625	0.996	0.945	0.682	0.574															
Sdim	0.884	0.874	0.988	0.731	0.964	0.931	0.861														
Sriv	0.866	0.841	0.985	0.872	0.953	0.902	0.848	0.766													
Ssch	0.845	0.811	0.966	0.867	0.911	0.833	0.913	0.826	0.861												
Snig	0.947	0.815	0.992	0.856	0.859	0.836	0.741	0.833	0.877	0.805											
Sovi	0.968	0.888	0.974	0.839	0.941	0.918	0.950	0.905	0.921	0.854	0.925										
Sfre	0.850	0.740	0.996	0.909	0.896	0.756	0.733	0.816	0.751	0.699	0.754	0.884									
Zsco	0.970	0.698	0.995	0.945	0.741	0.701	0.676	0.897	0.916	0.880	0.833	0.955	0.746								
Zvel	0.958	0.672	0.989	0.981	0.810	0.706	0.614	0.815	0.890	0.755	0.744	0.893	0.707	0.639							
Svul	0.957	0.804	0.999	0.987	0.671	0.635	0.678	0.891	0.903	0.846	0.935	0.980	0.834	0.780	0.751						
Spue	0.978	0.653	0.996	0.967	0.633	0.642	0.580	0.831	0.859	0.852	0.798	0.864	0.753	0.667	0.638	0.691					
Sarg	0.957	0.657	0.985	0.953	0.747	0.740	0.617	0.748	0.868	0.841	0.789	0.905	0.737	0.751	0.710	0.655	0.664				
Nuni	0.943	0.686	0.991	0.981	0.706	0.700	0.683	0.846	0.919	0.842	0.842	0.847	0.735	0.719	0.710	0.699	0.776	0.672			
Cmic	0.965	0.987	0.998	0.933	0.998	0.988	0.991	0.937	0.962	0.962	0.970	0.932	0.964	0.995	0.985	0.995	0.992	0.984	0.981		
Csor	0.948	0.827	0.996	0.867	0.930	0.849	0.901	0.787	0.793	0.793	0.744	0.904	0.783	0.802	0.843	0.780	0.927	0.858	0.853	0.946	

Table 5.3: Associated *P*-values obtained under the null model ($n = 999$). *P*-values > 0.95 indicate significantly less overlap than expected given the sample size and the total niche volume.

	Agri	Agro	Aoli	Cstr	Scor	Sdol	Stus	Sdim	Sriv	Ssch	Snig	Sovi	Sfre	Zsco	Zvel	Svul	Spue	Sarg	Nuni	Cmic	Csor
Agri																					
Agro	0.997																				
Aoli	0.989	0.999																			
Cstr	0.997	0.999	0.999																		
Scor	0.999	0.689	0.999	0.999																	
Sdol	0.999	0.781	0.999	0.999	0.359																
Stus	0.999	0.415	0.999	0.999	0.766	0.128															
Sdim	0.997	0.989	0.999	0.822	0.999	0.999	0.995														
Sriv	0.975	0.993	0.999	0.985	0.999	0.998	0.998	0.946													
Ssch	0.990	0.953	0.999	0.980	0.997	0.985	0.995	0.989	0.983												
Snig	0.999	0.979	0.999	0.982	0.995	0.975	0.935	0.965	0.970	0.979											
Sovi	0.999	0.996	0.999	0.989	0.999	0.999	0.997	0.973	0.997	0.995	0.996										
Sfre	0.989	0.924	0.999	0.997	0.996	0.970	0.925	0.992	0.926	0.736	0.971	0.996									
Zsco	0.999	0.822	0.999	0.998	0.864	0.778	0.724	0.993	0.994	0.991	0.982	0.998	0.915								
Zvel	0.999	0.731	0.999	0.999	0.944	0.829	0.356	0.987	0.995	0.881	0.953	0.995	0.898	0.514							
Svul	0.999	0.927	0.999	0.999	0.740	0.471	0.671	0.998	0.997	0.995	0.995	0.999	0.996	0.944	0.936						
Spue	0.999	0.636	0.999	0.999	0.495	0.566	0.170	0.997	0.996	0.994	0.983	0.994	0.935	0.710	0.515	0.823					
Sarg	0.999	0.644	0.999	0.999	0.925	0.917	0.351	0.839	0.994	0.990	0.978	0.991	0.962	0.889	0.870	0.612	0.648				
Nuni	0.999	0.812	0.999	0.999	0.818	0.736	0.756	0.988	0.991	0.991	0.987	0.984	0.898	0.887	0.911	0.786	0.818	0.604			
Cmic	0.999	0.999	0.999	0.999	0.999	0.999	0.999	0.999	0.999	0.999	0.999	0.998	0.999	0.999	0.999	0.999	0.999	0.999	0.999		
Csor	0.999	0.978	0.999	0.989	0.999	0.951	0.987	0.982	0.968	0.984	0.933	0.998	0.974	0.976	0.984	0.937	0.990	0.991	0.993	0.998	

An ordination based on non-metric multidimensional scaling of the dissimilarity matrix revealed two distinct groups (**Figure 5.3**), characterized by differences in the extent of their functional niche volumes and functional complementarity. The groupings were consistent with the hierarchical clustering analysis (**Figure 5.4**) and reflect the differences in the functional impact among species with a major division between species with limited overlap (mean overlap: $11.5\% \pm 1.1$ SE) and small volumes (detritus feeders, scrapers and excavators) and species with higher overlap ($31.5\% \pm 0.8$ SE) and large volumes (croppers and browsers).

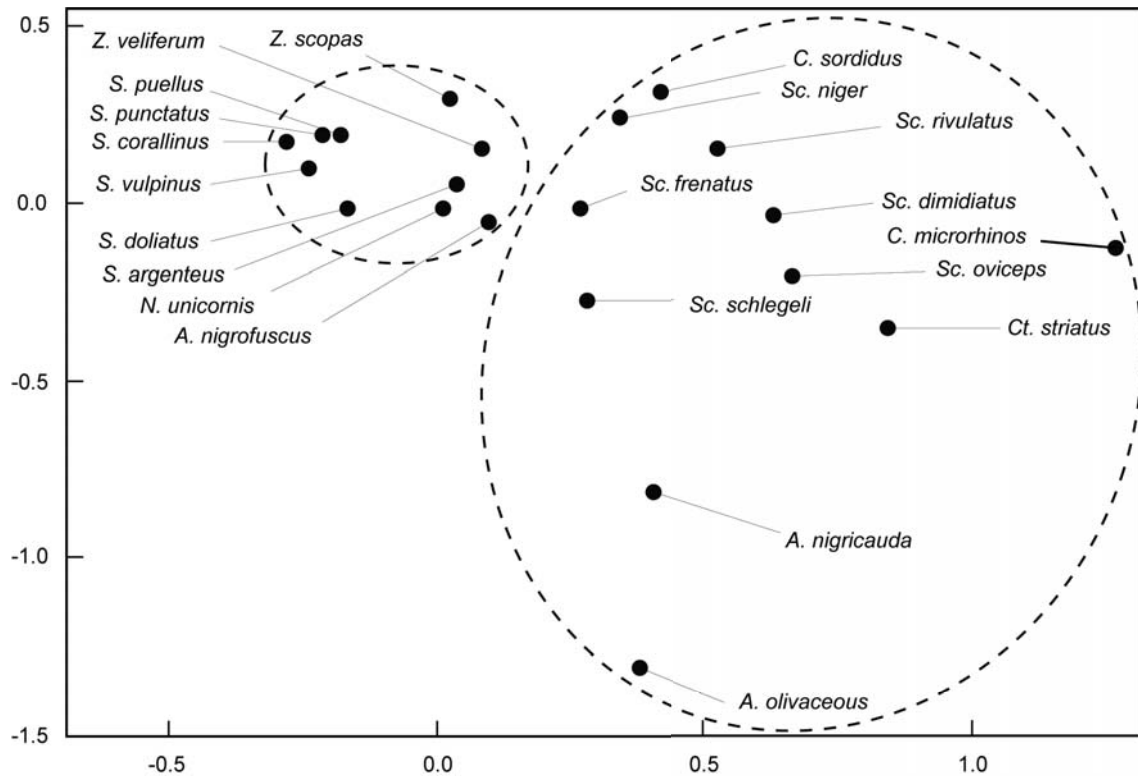


Figure 5.3: Functional distances between herbivorous reef fish species. The ordination plot represents the results from a non-metric multidimensional scaling analyses, performed on the pairwise dissimilarity matrix yielded by the analysis of overlap between niche volumes. Groupings are based on Ward's hierarchical clustering and indicate the functional overlap/distance between herbivore species, showing two distinct groups that differ markedly in their functional impact. The larger group on the right includes species with small niche volumes with minimal overlap, the smaller group on the left is characterized by species with large niche volumes and more overlap.

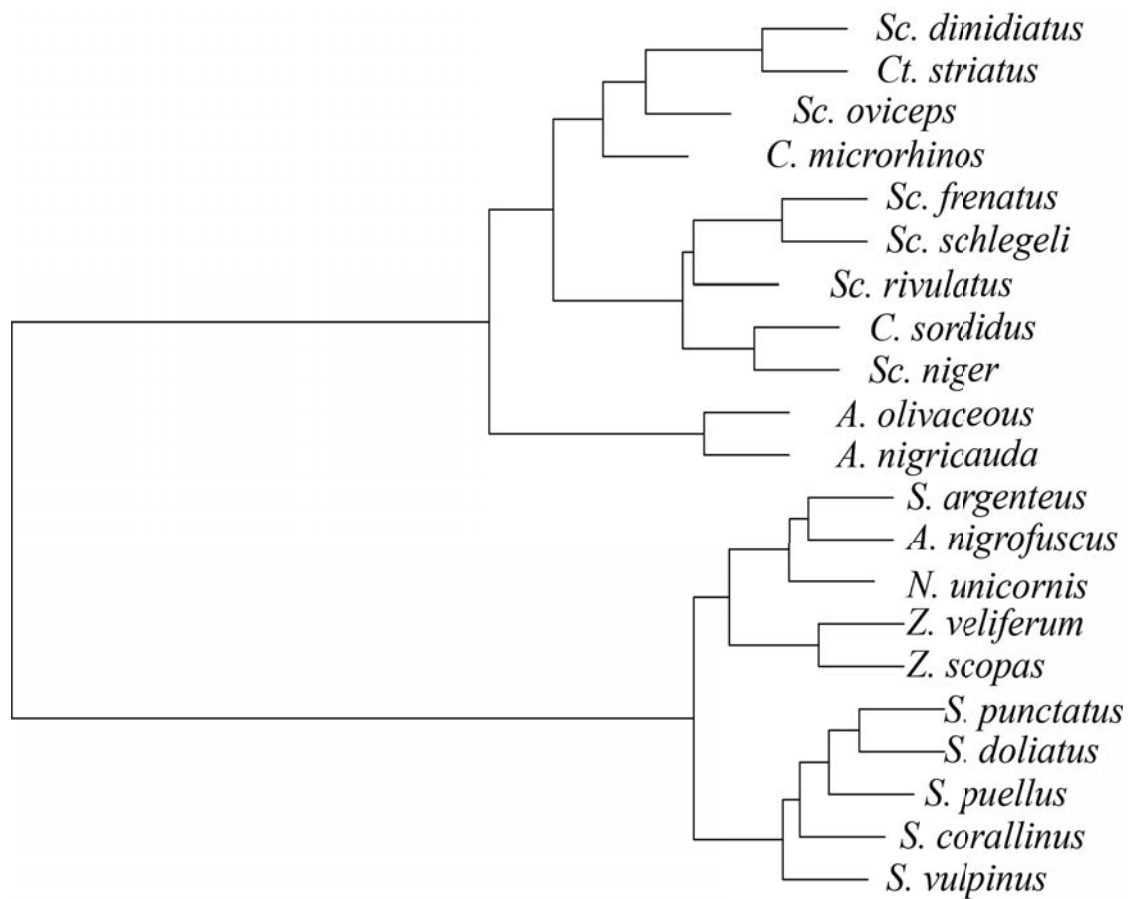


Figure 5.4: Ward's hierarchical clustering analysis, revealing a split into two distinct groups of herbivorous reef fishes based on their preferred feeding microhabitats.

These patterns are best exemplified by focusing on three species: the surgeonfish *Acanthurus nigricauda*, the parrotfish *Scarus schlegeli* and the rabbitfish *Siganus doliatus* and their utilization of different surface types (A, B, C, D and/or E) during foraging. These species show marked differences in their functional niche volumes (*Acanthurus nigricauda*, $4.6\% \pm 0.1$ SE; *Scarus schlegeli*, $15.1\% \pm 0.2$ SE; *Siganus doliatus*, $58.6\% \pm 0.3$ SE; values as percentages of the total niche space occupied). The volume occupied by *Acanthurus nigricauda* amounts to only 7.9% of the space occupied by *Siganus doliatus*. As one would expect, these species also show varying degrees of functional complementarity (**Figure 5.5**). *Acanthurus nigricauda* and *Scarus schlegeli*, both characterized by small niche volumes and united in the same group, overlap by 15.5% of their (combined) functional volumes. In contrast, *Acanthurus nigricauda* and *Siganus doliatus* only show an overlap of 6.8%. These differences strongly reflect the foraging patterns of the respective species (**Figure 5.6**). When considering targeted reef surfaces, for example, individuals of *Acanthurus nigricauda* almost exclusively target horizontal, open surfaces [A] ($88.9\% \pm 0.8$ SE of all bites) and do not feed on any other surfaces to a significant extent. Individuals of *Scarus schlegeli* show slightly more variation by also feeding on vertical, open surfaces [B] ($25.1\% \pm 1.3$ SE) while still targeting predominantly horizontal, open surfaces ($67.7\% \pm 1.4$ SE). In contrast, individuals of *Siganus doliatus* exhibit the broadest niche on this axis by feeding on almost all available surfaces on the reef (except undersides [C]) (**Table 5.1**). This segregation is reinforced by differences in the targeted substratum types and the extent of substratum penetration. Both *A. nigricauda* and *S. schlegeli* forage predominantly on the reef matrix or sandy substrata while *S. doliatus* utilizes a wide range of substrata and frequently penetrates the substratum when feeding.

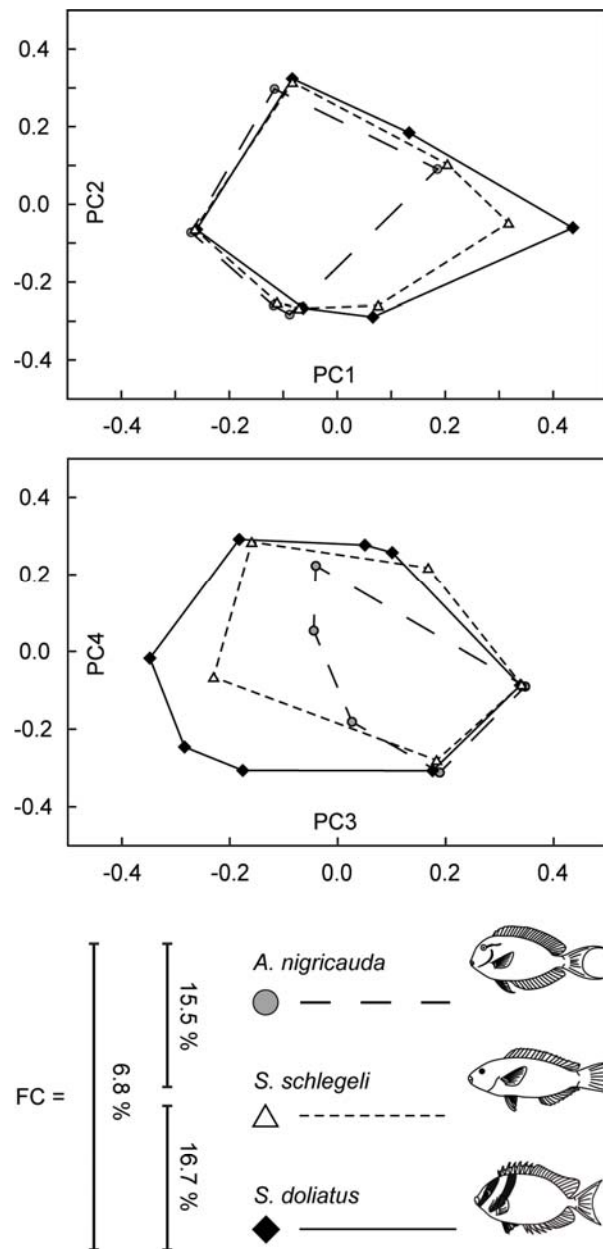


Figure 5.5: Multidimensional niche volumes of *Acanthurus nigricauda*, *Scarus schlegeli*, and *Siganus doliatus* calculated over four axes. Symbols mark the most extreme individuals (vertices) from which niche volumes (convex hull volumes) are calculated. Note the changes in niche volumes when considering more than two axes. Functional complementarity (FC) between species is expressed in % overlap between niche volumes of the respective species.

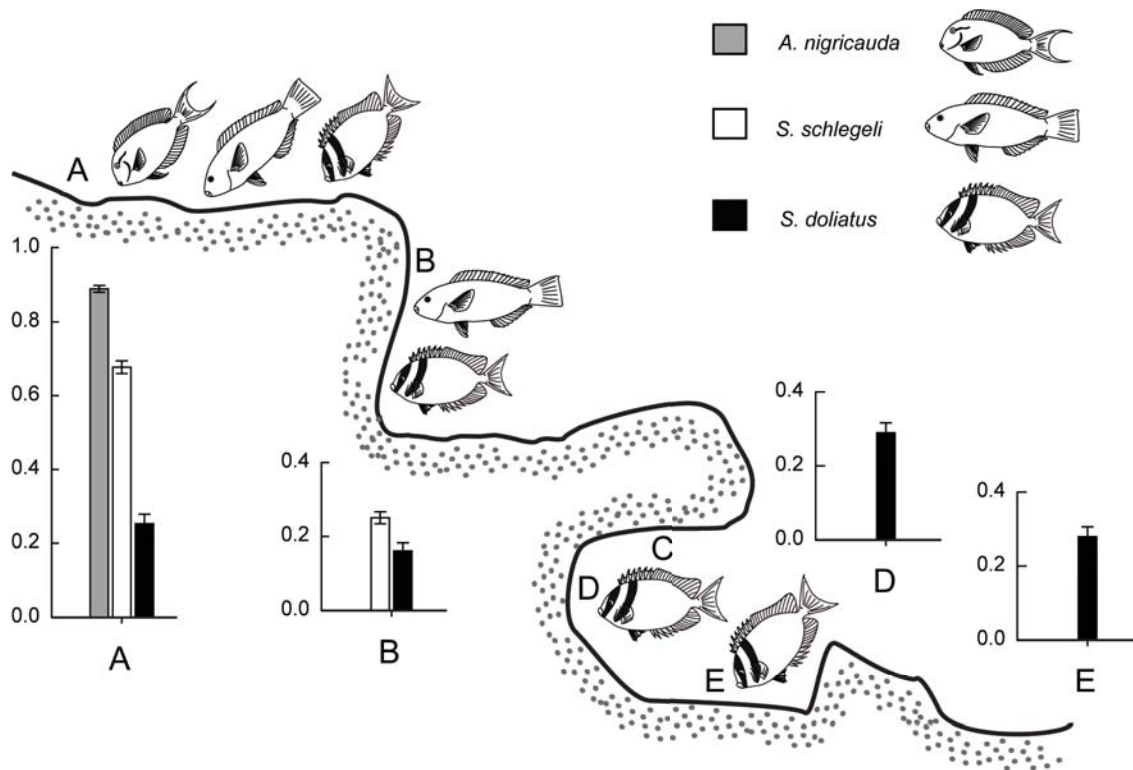


Figure 5.6: Microhabitat utilization patterns in three species exemplified by surface type use of *A. nigricauda*, *S. schlegeli* and *S. doliatus*. Barplots indicate proportions of bites taken from different surfaces (only surfaces with > 0.15 of the total bites illustrated). Observed patterns show low variation in the choice of feeding microhabitats in *A. nigricauda* and slightly more variation in *S. schlegeli*, while *S. doliatus* targets almost all surfaces to a roughly equal extent. Surfaces are coded according to the main text (A = open, horizontal; B = open, vertical; C = underside; D = concealed, vertical; E = concealed, horizontal).

Decomposing the dissimilarity into its turnover and nestedness components further reveals the nature of the niche partitioning among the three species. The overlap between *A. nigricauda* and *S. schlegeli* is predominantly due to the nestedness of the small niche volume of *A. nigricauda* within the volume of *S. schlegeli* (72.3% overlap due to nestedness; **Figure 5.7**) and both species' niches are nested within the large volume of *S. doliatus* (52.5% and 73.7% nestedness, respectively).

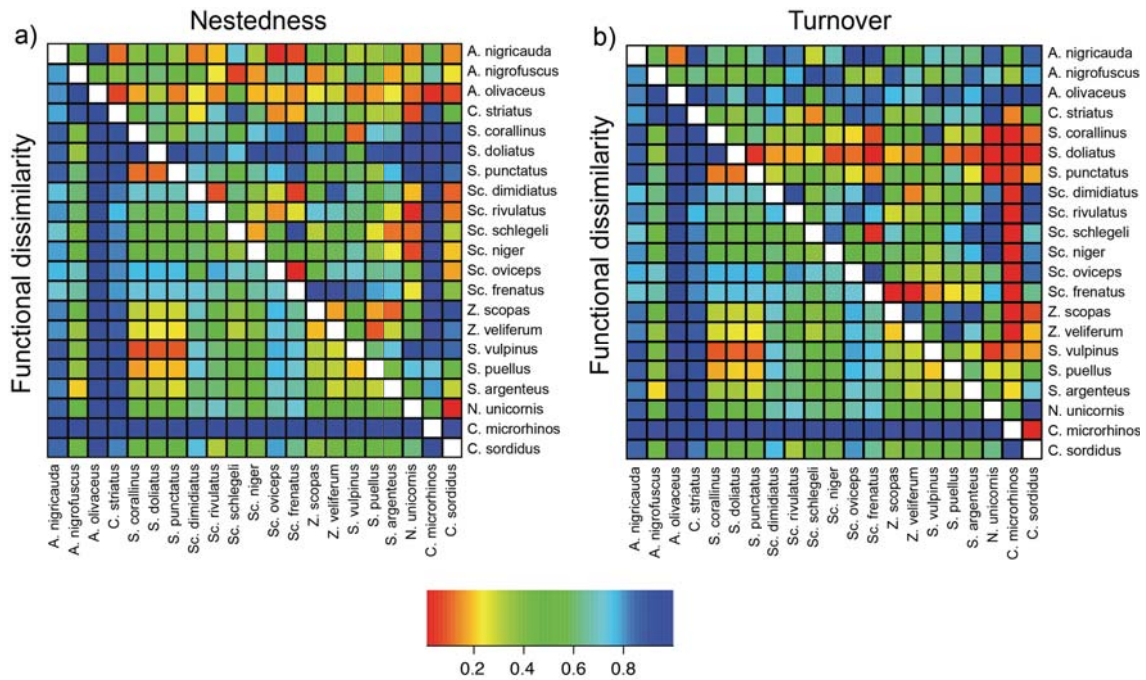


Figure 5.7: Contribution of turnover and nestedness components to functional overlap among herbivorous fishes. In both heatmaps, the lower left matrix marks the functional dissimilarity, while the upper right triangular matrix denotes the contribution of, a) the nestedness component and b) the turnover component of the overlap.

5.4 Discussion

Using a new multidimensional analysis of species' functional niches based on individual observations, I provide evidence for extensive complementarity among herbivorous reef fishes when considering their choice of feeding microhabitats. Some species deliver their function over a wide range of microhabitats while other species appear to be restricted in their microhabitat utilization patterns, occupying less than 5% of the largest niche volume. In calculating the functional complementarity (i.e. the overlap between niche volumes of species), I show that herbivorous species can be broadly divided into two distinct groups, which are not revealed in traditional functional classifications. These groups, characterized by the choice of feeding microhabitats, show striking differences. One group exhibits little individual variation, leading to small functional niches and extensive complementarity, while species in the other group show high variance among individuals, large functional niche volumes and, as a result, less complementarity. These results may have important implications for our understanding of herbivorous processes on coral reefs and highlight the benefits of the new approach in complementing and extending traditional functional groupings.

5.4.1 Individual-based analyses vs. traditional groupings

The assessment of the foraging ecology of herbivorous reef fishes using individual-based analyses provides a finely-graded perspective of functional niche occupation and the extent of overlap among species within a herbivore community. Traditional classification schemes provide a broad, categorical framework. In contrast, my analysis presents a largely concordant but finer-scaled picture that places all species and individuals within a continuous range of functional niches, thus permitting a deeper understanding of functional variation within and among herbivorous species and within traditional functional groups. As a

consequence, we can quantitatively explore the extent of functional redundancy or complementarity within functional groups of herbivorous reef fishes.

The differences between traditional groupings and the results yielded by my analyses are apparent when comparing, for example, the functional role of rabbitfishes (f. Siganidae). All 28 members of this family are united in one genus, *Siganus*, and are considered to exhibit limited variation in their taxonomy and general morphology (Borsa et al. 2007). As a result, their functional niche is assumed to be relatively uniform (Green & Bellwood 2009; Cheal et al. 2012). While this view is partially supported when examining the foraging microhabitats of rabbitfishes (all species are in the same cluster), the results of my analyses suggest that there are marked differences between species, as the maximum functional overlap between the two most similar species (*S. doliatus* and *S. punctatus*) was only 42.6%. The extent of separation is even more apparent when examining the functional overlap between *S. corallinus* and *S. argenteus*, which only have an overlap of 25.3%. Such functional differences have been suggested based on morphological traits (Brandl & Bellwood 2013a) and dietary analyses (Hoey et al. 2013). However, it is only by using *in situ* observations that we are able to reveal the extent of segregation in the functional niches of rabbitfishes.

Another advantage of the new analyses compared to traditional groupings is the species-specific assessment based on individuals. This can be illustrated using the rabbitfish *S. doliatus*. Occupying more than half of the available functional niche space with regards to feeding microhabitat selection, *S. doliatus* appears to exhibit remarkable flexibility and the ability to exploit virtually all available locations. Although some studies suggest an exceptional functional role for *S. doliatus* on coral reefs (Fox & Bellwood 2007; Cheal et al. 2010; Brandl & Bellwood 2013b), the capabilities of *S. doliatus* to utilize a range of different microhabitats and, consequentially, its large functional niche volume have not been considered in traditional groupings. The large occupied niche volume in *S. doliatus* can be

attributed to a high degree of individual variation within this species. Despite the fact that previous individual-based observations have shown that coral reef fishes display high flexibility in their foraging behaviour (Dill 1983; Biro et al. 2010; Fox & Bellwood 2011), traditional groupings often neglect this aspect. My results emphasize the potential importance of assessing variance among individuals within a species when determining its functional niche (Messier et al. 2010; Violle *et al.* 2012).

An additional benefit of the analyses is the potential to incorporate multiple axes (Rosenfeld 2002). Traditional groupings or assessments of niche overlap are commonly based on a single factor, e.g. diet or feeding mode (Nyström 2006; Green & Bellwood 2009; Cheal et al. 2012; Fox & Bellwood 2013). However, redundancy is likely to decrease with the incorporation of multiple axes, which can reveal important information when assessing the functional structure of a species assemblage (Rosenfeld 2002). Again, rabbitfishes provide a good example. The only previous study on the feeding microhabitats of rabbitfishes (Fox & Bellwood 2013) reported *S. corallinus*, *S. puellus* and *S. vulpinus* to overlap almost entirely in their occupied niche (calculated solely by the extent to which fishes penetrated the substratum). I find that while there is significant overlap between these three species, the consideration of surface type, substratum type and penetration depth reveals a marked reduction in their functional overlap (e.g. 30.9 % between *S. puellus* and *S. vulpinus*, with a turnover component of 38.8%). In addition, factors such as dietary specialization (Hoey et al. 2013) are likely to further differentiate functional niches, underlining the importance of multiple axes in disentangling functional niches within a species community.

Our approach also permits comparisons between traditional functional groups. For instance, the surgeonfish *A. nigricauda* and the rabbitfish *S. doliatus* have previously been classified in either the same ('grazers/detritivores'; Green & Bellwood 2009) or different functional groups ('grazers/detritivores' and 'browsers'; Cheal et al. 2012). My analysis

identified their functional niche volumes to be markedly different (4.6 ± 0.1 % SE and 58.6 ± 0.3 % SE of the total niche space, respectively with just 6.8 % of overlap in volumes). Thus, irrespective of their functional group membership, these species differ in their microhabitat utilization and my analyses allow for a quantitative estimate of the disparity between species.

Even more detail can be revealed when considering the components contributing to the overlap between species. For example, *Z. scopas* and *S. argenteus*, both classified as browsers (Cheal et al. 2012), exhibit a 24.9% overlap in their niche volumes and are grouped in the same cluster. My analyses reveal that 89.7% of the dissimilarity between these two species is due to high turnover, i.e. functionally dissimilar individuals (**Figure 5.7**). Thus, even if both are browsers, both species comprise of individuals that feed in microhabitats not utilized by the other species. A similar picture emerges within parrotfishes, which have been assumed to exhibit relatively similar functional niches (Bellwood & Choat 1990; Cardoso et al. 2006; Green & Bellwood 2009; Price et al. 2010; Cheal et al. 2012). However, when examining the two scraping parrotfishes *Scarus frenatus* and *S. oviceps*, these species appear to differ considerably in their functional niches (11.2 % of overlap), with *S. frenatus* targeting more coral rubble and concealed surfaces than *S. oviceps*. This is again predominantly due to high turnover. In marked contrast, while the two excavating parrotfishes *Chlorurus microrhinos* and *C. sordidus* also only overlap in 5.4% of their niche volumes, 97.2% of this dissimilarity is due to nestedness, with the relatively small niche volume of *C. microrhinos* almost entirely nested within the volume of *C. sordidus*. This suggests that, based on microhabitat utilization patterns, *C. sordidus* may be able to compensate for the loss of *C. microrhinos* but not *vice-versa*. Although parrotfishes are clustered in the same broad group, the detail my analyses can provide emphasizes the variation within the cluster (Fig. S2) and provides an indication of the extent and nature of functional complementarity.

The differences within all three families highlight the broad pattern seen in microhabitat use. While siganids are generally croppers and parrotfishes generally scrapers, the former exhibit broad microhabitat utilization whereas the latter only use a limited range of microhabitats. Thus, parrotfishes may not show the extent of redundancy one may assume. In all these cases, the quantitative estimates yielded by individual-based analyses of functional niche overlap show that traditional groupings offer only a first step in disentangling the complex functional structure of high diversity assemblages. My approach may be of benefit to ecologists in other fields who seek to resolve functional niche partitioning in highly diverse species communities. Thus, it may be useful to tease apart the functional niches of, for instance, birds (e.g. Şekercioğlu 2006), insects (Andersen 1997), or across various groups of animals with similar ecosystem functions, such as pollinators (Clark et al. 2001).

In summary, my approach offers a high-resolution quantitative method to explore functional niche occupation and the extent and nature of functional overlap among species. Several recent studies have emphasized that there are areas of potential weaknesses that can arise from assuming redundancy in key functional groups (Petchey et al. 2007; Cheal et al. 2013; Johansson et al. 2013). My analyses provide means to go beyond functional interpretations based on the abundance of species in an ecosystem to quantitatively evaluate species' actions and activities.

5.4.2 Ecological implications

From an ecological perspective, the results of the present study point to two distinct herbivore groups with extensive niche partitioning within both these and traditional functional groups. This calls for caution when assessing reef resilience based on broad functional groupings. Several species, mostly surgeon- and parrotfishes, appear to graze heavily on open, exposed

reef surfaces and sandy substrata. There are, however, subtle differences in the foraging behaviour of these species. This is particularly evident within scraping parrotfishes, which appear to partition the available, exposed substratum very finely by having relatively distinct foraging behaviour characterized by little intraspecific variation and high functional turnover, leading to extensive complementarity. This is consistent with observations on parrotfish foraging behaviour in the Caribbean (Cardoso et al. 2006). In the parrotfishes, explaining the coexistence of so many morphologically and trophically similar species in a single ecosystem has been a challenge (Choat & Bellwood 1990; Streelman et al. 2002; Price et al. 2010; Choat et al. 2012). My analyses suggest that fine-scale niche partitioning of feeding microhabitats may be an important factor in explaining these patterns. This implies that scraping parrotfishes, which are generally assumed to have very similar functional niches (thus showing high functional redundancy), may provide yet another example of limited functional redundancy in herbivorous reef fishes (Bellwood et al. 2003; Hoey & Bellwood 2009), as their ability to remove algae appears to depend strongly on species-specific microhabitat utilization patterns.

In contrast to the first group (small niches, predominantly parrotfishes), only a limited suite of species (mostly rabbitfishes and several surgeonfishes) graze on concealed reef microhabitats. However, individuals in these species are highly flexible in their microhabitat utilization patterns; all feed on a wide range of different microhabitats, including those targeted by species with limited niche volumes. Flexibility in the use of different microhabitats has been suggested previously for some rabbitfishes and surgeonfishes (Robertson et al. 1979; Robertson & Gaines 1986; Fox & Bellwood 2013) and may suggest that either, a) concealed microhabitats do not provide enough resources to maintain these species if they feed solely in concealed areas, or b) that species feeding on open microhabitats competitively exclude the more flexible species from feeding on easily

accessible surfaces (Robertson et al. 1979; Robertson & Gaines 1986; Alwany et al. 2005). The latter scenario may be particularly important for reef management as species with high flexibility in their foraging microhabitat selection may be critical for the resilience of reef ecosystems because of their ability to remove algae from inaccessible microhabitats (Brandl & Bellwood 2013a; Fox & Bellwood 2013).

Overall, using a new multidimensional framework to analyse feeding microhabitat utilization among herbivorous reef fishes, my study demonstrates the sensitivity of individual-based *in situ* behavioural observations when evaluating a species' realized functional niche, especially in high diversity ecosystems. I find that herbivorous species vary greatly in their functional impact with regards to their microhabitat utilization patterns. Indeed, I found no functional redundancy (based on a 50% cut-off) among herbivorous fish species in their microhabitat use. Herbivores broadly divide into species that feed predominantly on open surfaces and species that also exploit concealed microhabitats. Species within the first group appear to partition their feeding microhabitats very finely, resulting in small functional niche volumes with high complementarity. In contrast, individuals of species feeding on concealed microhabitats show high variation in their foraging behaviour, leading to higher overlap of niche volumes (mostly due to a strong turnover component, thus maintaining functional complementary rather than redundancy). These results underscore the potential sensitivity of the analyses and their ability to reveal details of species' functional niches and their role in ecosystem processes.

Applied to coral reef fish communities, my results emphasize the need for caution when assessing reef resilience based solely on assessments of biodiversity and broad functional groups and stress the importance of complementarity among herbivorous reef fish species. Studies using traditional functional groups may find high redundancy; however, this can arise as a result of either true overlap in functions or excessively broad functional

categories. My results suggest that the latter may be more common than previously thought. Given the global threat of changing environmental conditions and the progressive loss of biodiversity, the key question now is: to what extent can a species' functional niche shift in response to changing environments or changing patterns of biodiversity?

Chapter 6: Exploring the nature of ecological specialization in a coral reef fish community: morphology, diet, and foraging microhabitat use

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6.1 Introduction

The fundamental characteristics of species' niches shape ecological communities across the globe. Within communities, species differ not only in their positions in niche space, but also in the respective sizes of the volumes they occupy (Whittaker et al. 1973). Specialist species, often considered as taxa that occupy the smallest niche volumes, have evoked particular interest in a wide range of fields (Futuyma & Moreno 1988). Ecological specialization is accepted as a key factor for countless facets of organismal biology, affecting speciation (Schluter 2000), species distribution patterns (Emery et al. 2012), and ecosystem functioning and stability (Clavel et al. 2010). Yet, as ecological specialization can occur on various levels (e.g. fundamental vs. realized or individuals vs. populations; Bolnick et al. 2003; Poisot et al. 2011; Ferry-Graham et al. 2002a), many aspects of its ecosystem impacts remain unresolved.

Two of the most commonly studied features of ecological specialization are morphological/physiological attributes and behavioral traits relating to foraging (Futuyma & Moreno 1988), and their respective relationships. By identifying morphological traits that directly relate to prey capture or procurement capacities, numerous studies have established strong correlations between morphological and dietary specialization in birds (Grant & Grant 1996), reptiles (Henderson et al. 2013), fishes (Huckins 1997), and mammals (Aguirre et al. 2002), suggesting a tight link between fundamental and realized niches and permitting the

inference of simple directional evolutionary pathways. Consequently, relating morphological features to dietary preferences has become one of the most widespread techniques used to extrapolate ecological function (Wainwright & Reilly 1994). Ichthyologists, in particular, have made extensive use of ecomorphological techniques to interpret the ecology of fishes, especially when sustained observational studies are unfeasible (Wainwright 1988, 1996; Douglas & Matthews 1992; Wainwright & Bellwood 2002; Pouilly et al. 2003; Snorrason et al. 1994; Sturmbauer et al. 1992; Hulsey & García de León 2005).

However, in the last two decades, there have been a number of intriguing studies that have revealed high behavioral plasticity in morphologically specialized fish species (Liem 1980; Motta et al. 1995; Barnett et al. 2006; Bellwood et al. 2006; Sanderson 1990). It appears that morphological specialization in teleost fishes does not always entail dietary specialization and that, in many cases, a species' realized (dietary) niche is largely unrelated to its morphology (Alfaro et al. 2005; Wainwright et al. 2005). One of the most striking examples of this scenario was described in African cichlids, where the apparent mismatch between morphological features and dietary niches in ecologically specialized species has been termed "Liem's Paradox" (i.e. why are morphological specialists often dietary generalists; Liem 1980; Robinson & Wilson 1998). The most common explanation for the paradox is that specialized morphological traits enable exploitation of "fallback foods" when preferred resources are low. It is posited that this has arisen as a result of competition-driven morphological specialization during low-resource situations, while retaining the capacity to exploit a range of preferred resources under more typical circumstances (Robinson & Wilson 1998). This hypothesis has been supported by both empirical and theoretical work on a wide range of taxa (Fontaine et al. 2008; Ungar et al. 2008), and appears to be the most prevalent explanation in teleost fishes (Bouton et al. 1998; Janovetz 2005).

As the marine analogues to cichlids in African Rift Lakes, coral reef fishes with their extreme diversity offer an outstanding opportunity to investigate ecological specialization (Wainwright & Bellwood 2002). Despite extreme morphological specialization in numerous reef fishes lineages (Ferry-Graham et al. 2002a), attempts to relate these features to dietary niche partitioning have had mixed success (Bellwood et al. 2006; Price et al. 2010). Herbivorous reef fishes *sensu lato* (Clements et al. 2009) provide an excellent group for exploring patterns of ecological specialization. Herbivore communities are of critical importance for the benthic structure of shallow reef environments and have been the subject of intensive research with regards to their diet (Choat et al. 2002; Hoey et al. 2013), foraging behavior (Rasher et al. 2013), and evolutionary history (Choat et al. 2012). However, patterns of ecological specialization in herbivore families have rarely been examined. This is particularly interesting given the variation in morphological diversification among and within herbivorous families (Choat et al. 2012; Brandl & Bellwood 2013a), and the presence of dietary niche partitioning across distinct phylogenetic lineages (Choat et al. 2002; Robertson & Gaines 1986; Robertson et al. 1979). Furthermore, there is evidence for subtle partitioning of feeding microhabitats in many families (Robertson & Gaines 1986; Brandl & Bellwood 2014a), which has been linked to morphological traits (Brandl & Bellwood 2013a; Fox & Bellwood 2013) but has not been considered with regards to ecological specialization, despite its potential importance in aquatic ecosystems (Holbrook & Schmitt 1992). There is, therefore, strong indication of the potential for ecological specialization in herbivorous reef fishes along three distinct but interrelated niche axes: morphology, diet, and foraging microhabitat utilization. In addition, in light of the functional importance of herbivores in shaping benthic dynamics, and the tendency of ecological specialists to be among the first species to succumb to human-induced disturbances (Clavel et al. 2010), detecting ecological specialists among herbivorous fish species may be important for coral reef management,

particularly given current anthropogenic pressures (Bellwood et al. 2012; Edwards et al. 2014).

To assess the relationship between fundamental (morphological) and realized (behavioural) specialization on coral reefs, I examined an assemblage of ten species of surgeonfishes (f. Acanthuridae). I first identified morphological specialization among species using external features, and then considered two distinct behavioural niche axes, diet and foraging microhabitat utilization. For each behavioural axis, I examined behavioral specialization using an individual-based multidimensional framework Brandl & Bellwood 2014a), which allows for a representation of all species along a quantitative generalist-specialist continuum. I examined the relationship between morphological specialization and the two behavioural niche axes based on three underlying hypotheses: 1) no relationship (as a null-hypothesis), 2) a positive relationship following the ‘traditional’ ecomorphological paradigm where morphological specialists are behavioral specialists, and 3) an inverse relationship (potentially consistent with Liem’s Paradox) (**Figure 6.1**).

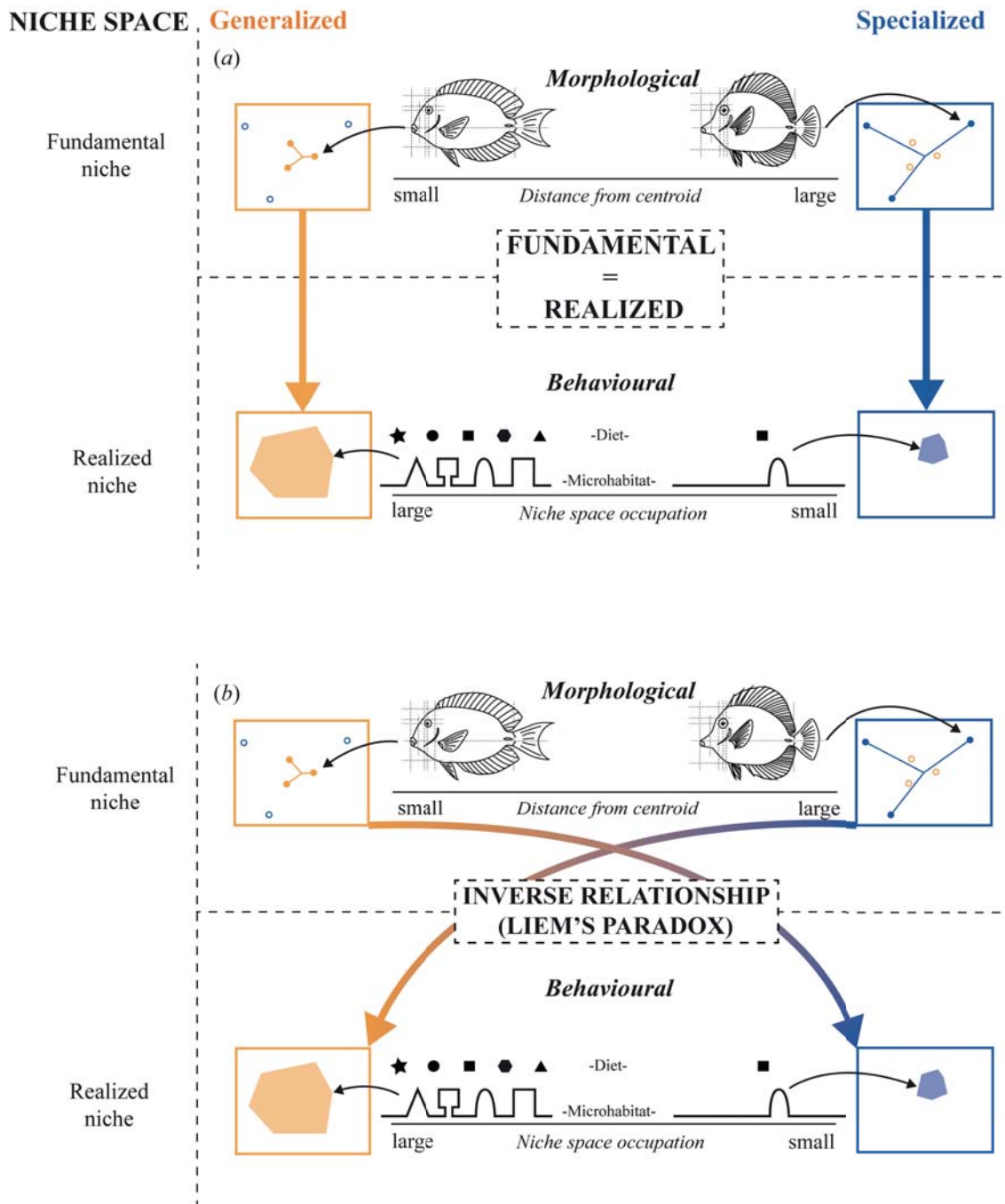


Figure 6.1: A conceptual synthesis of the potential relationships between fundamental (morphological) and realized (behavioural, i.e. diet [filled symbols] or foraging microhabitat utilisation [open symbols]) specialization in surgeonfishes. The three hypotheses were, i) no relationship between morphological and behavioural

specialization (not figured), ii) morphological specialists are behavioural specialists (a), or iii) morphological specialists are behavioural generalists and *vice versa*, potentially consistent with a phenomenon termed Liem's paradox (b).

6.2 Materials and methods

All fieldwork and sampling was conducted on reefs around Lizard Island, a mid-shelf island in the northern Great Barrier Reef (GBR), Australia (14°40'08"S 145°27'34"E). Data were collected from adult individuals of ten surgeonfish species in four genera, *Acanthurus blochii*, *A. lineatus*, *A. nigricauda*, *A. nigrofuscus*, *A. olivaceus*, *Ctenochaetus striatus*, *Naso lituratus*, *N. unicornis*, *Zebrasoma scopas*, and *Z. velifer*. These species represent 40% of the surgeonfish species recorded from the GBR and encompass all numerically abundant species at the study site (Cheal et al. 2012).

6.2.1 Morphological characterization

Species were characterized using a set of morphological traits consisting of 19 distance-based measurements and three angle measurements. All morphological measurements were selected based on previous linkages between the morphology and functional aspects of the species' foraging ecology, such as feeding, locomotion, or sensory abilities (Douglas & Matthews 1992; Bellwood et al. 2014). All values were transformed to ratios by standardizing them against head or standard length, as required (e.g., snout length \div head length). As none of the ratios showed a significant relationship with standard length, analyses were performed using the standardized values rather than residuals. The average number of individuals measured for each species was 3.8 ± 0.3 (SE). Details are provided in Bellwood et al. (2014). A list of morphological features measured and their ecological significance is provided in **Table 6.1**.

Table 6.1: List of morphological traits measured and their ecological significance. All measurements are explained in greater detail in Bellwood et al. 2014 and references therein.

Morphological trait	Measured as	Affects
Body depth	Body depth/standard length	Predator avoidance, maneuverability
Head length	Head length/standard length	Sensory and trophic capabilities
Narrowest point on caudal peduncle	Width caudal peduncle/body depth	Swimming speed
Body aspect ratio	Body depth ² /body area	Predator avoidance, maneuverability
Medial caudal fin ray length	Medial fin ray/longest fin ray	Swimming ability and speed
Snout length	Snout length/head length	Precision in prey acquisition
Eye diameter	Eye diameter/head length	Visual acuity
Head depth	Head depth/body depth	Hydrodynamics, maneuverability
Lower jaw length	Lower jaw length/head length	Range of prey items ingested
Anterior of orbit to forehead	Anterior of orbit/head length	Binocular vision
Horizontal eye position	Distance posterior the orbit/ head length	Field of perception
Vertical eye position	Distance ventral of orbit/head depth	Field of perception
Mouth-eye distance	Orbit centroid to premaxilla/head length	Eye-snout coordination
Mouth position	Ventral of snout/head depth	Prey acquisition
Longest dorsal spine	Longest dorsal spine/body depth	Predator avoidance, maneuverability
Snout angle	Angle characterizing tip of snout	Prey acquisition
Head angle	Angle characterizing anterior cranial region	Prey acquisition
Snout-eye angle	Angle between snout and eye	Eye-snout coordination

6.2.2 Diet

Specimens were collected from the reef using spearguns and barrier nets. Specimens were kept on ice and the whole alimentary tract was removed and preserved in 10% buffered formalin within 1h of capture. Contents were identified using microscopic analysis (4x10 magnification) along transects of a 100×100mm grid on a Perspex plate. For detrital feeders with considerable amounts of fine organic matter in alimentary tracts, examination was performed using 10x40 magnification to allow for the identification of microalgae and other microscopic dietary items. An average of 6.6 ± 0.3 (SE) specimens per species were analyzed. The methods are described in detail in (Choat et al. 2002). A list of dietary categories is provided in **Table 6.2**.

Table 6.2: List of categories of dietary items found in alimentary tracts of the examined species.

Categories of dietary items

Arthropoda
Brown filamentous algae
Brown foliose algae
Brown thallose algae
Calcareous sediment
Chitinous matter
Chitons
Crustacea
Echinoids
Eggs
Fish scales
Foraminifera
Gastropoda
Gelatinous zooplankton
Green filamentous algae
Green foliose algae
Green thallose algae
Heteropod mollusca
Hydroids
Organic matter
Red filamentous algae
Red foliose algae
Red thallose algae
Siliceous material
Spicule
Sponge spicule
Stomatopoda
Sponges
Gelatinous matter

6.2.3 Foraging microhabitat preferences

Observations on foraging microhabitat selection were carried out on SCUBA or while snorkeling (during April/May 2013). Haphazardly chosen individuals were followed until the first foray. Then, the surface orientation and accessibility [open horizontal (A), open vertical (B), underside (C), concealed vertical (D) or concealed horizontal (E)], the substratum type (live coral [LC], dead coral skeleton [DC], macro-algae [MA], coral rubble [RB], sand [SD], turf covered reef matrix [TR]) and the number of bites taken from the respective microhabitat were recorded. In addition, the extent to which an individual penetrated the substratum was measured using vernier calipers. The average number of observations per species was 49.8 ± 1.3 (SE) (details in Brandl & Bellwood 2014b).

6.2.4 Microhabitat availability

The abundance of different microhabitats was also estimated during April/May 2013, using 20 ten-meter chain intercept transects laid parallel to the reef crest (following Goatley & Bellwood 2011). A transect tape was laid conforming to the reef benthos to include concealed microhabitats. The underlying microhabitat was recorded every 20cm, using vernier calipers to measure the depth of crevices and depressions. Measurements were taken perpendicular to the substratum. To match the foraging microhabitat observations, microhabitat characteristics included the surface orientation and exposure (A, B, C, D, E), substratum type (live coral, dead coral, rubble etc.), and depth of interstices and crevices.

6.2.5 Statistical procedures

In terms of morphological specialization, analyses were performed at the species level. Morphological data (traits averaged for each species) were analyzed using a principal coordinate analysis (PCoA) based on Gower's dissimilarity matrix. Distances were calculated

among species based on the average trait measurements for each species. Using the coordinates for each species in the first two dimensions, I calculated the distance from the centroid of the ordination for each species as a proxy for morphological specialization (following Bellwood et al. 2006). While this quantification is dependent on the relative proportions of species morphotypes used, the species were selected to broadly reflect the relative proportions of surgeonfish types in GBR assemblages.

I used an individual-based multidimensional framework for the calculation of behavioural niche spaces yielding quantitative, continuous estimates of niche space occupation for both dietary and microhabitat selection data based on inter-individual variation for each species. First, I calculated the pairwise distances between all individuals of all species using Gower's distance metric. Following this, a PCoA was performed using Cailliez' correction for negative eigenvalues (Laliberté & Legendre 2010), yielding a distribution of all individuals in a synthetic dietary/microhabitat niche space. Subsequently, I computed species niche space occupation for diet and microhabitat utilization in the same fashion, based on individuals with the most extreme coordinates using convex hulls (Ackerly & Cornwell 2007; Cornwell et al. 2006). To account for unequal sample sizes in both datasets (i.e. different numbers of observations among species, which may lead to biased estimates of niche volumes as a matter of chance) and to decrease the effect of few extreme individuals on the overall volume for each species, I performed randomized permutations ($n = 999$) with a constant subsample of individuals for each species and calculated the mean niche space occupation for each species for further analyses (i.e. for each species, 999 volumes were calculated from subsamples of individuals). Thus, my measure of species' degree of specialization was a function of inter-individual variability within populations, a common metric of species' ecological niches (Whittaker et al. 1973; Colwell & Futuyma 1971). Due to restrictions in the dimensionality of the data and because I did not directly compare dietary

and microhabitat specialization, dietary niche space occupation was computed on two dimensions, whereas microhabitat niche space occupation was conducted on both two- and three-dimensional space to maximize extraction of information.

For foraging microhabitat specialization, I also computed Manly's χ^2 log-likelihood statistic (χ_{L2}^2) (Manly et al. 2002), which quantifies a population's selectivity for a given resource based on availability within the ecosystem. I used model design I with sampling protocol A (Manly et al. 2002), which measures resource selection on the population level, to calculate selection ratios for each species (used/available). High ratio values indicate high levels of specialization, while low ratio values suggest a generalist strategy.

I investigated the relationships between morphological specialization (as measured by the distance from the centroid in morphospace), dietary specialization, microhabitat specialization (both of the latter quantified as the dietary/microhabitat niche space occupied by individuals within a species, i.e. inter-individual variability across time and space), and microhabitat selectivity (Manly's χ^2 log-likelihood statistic (χ_{L2}^2)) using Bayesian linear models with morphology as a fixed effect and the respective behavioural variable as response. Uninformative default priors were used on the fixed effect estimates (0 on intercept and regression coefficient, variance of 100,000,000 for both parameters; Hadfield 2010). For the residual variance, I specified uninformative inverse Wishart priors with $V = 1$ and $\nu = 0.002$. Models were run for 2,000,000 iterations with a burnin of 50,000 and a thinning interval of 1,000. Chain convergence was validated using trace chain plots.

6.2.6 Phylogenetic correction and evaluation of assumed relationships

To account for biases due to shared ancestry, I used a phylogenetically corrected principal component analysis (PCA) on the morphological data using phylogenetic distances from a pruned tree (**Figure 6.2**) based on Sorensen et al. (2013) (Martin & Wainwright 2011; Revell

2012). In this ordination, evolutionary independence is achieved by incorporating estimated ancestral states for each character into the analysis, therefore reducing phylogenetic correlation among scores on each axis to zero. Subsequently, I examined the relationship between morphological and dietary/microhabitat specialization using phylogenetic Bayesian Mixed Models (BMMs). Again, phylogenetic relatedness was determined from a pruned tree (Sorensen et al. 2013) and fitted as a random factor under a Brownian motion model of evolution in the BMM (Hadfield 2010). In order to evaluate whether phylogenetic correction altered the results from my previous models, and due to the small sample size of only ten species in three distinct phylogenetic groups, I specified informative priors using the parameter estimates obtained from the previous uncorrected models for each relationship. This was done as my prior belief in the respective relationships was based on the uncorrected model estimates. For both random and residual variance, I again specified uninformative inverse Wishart priors and the number of iterations was specified as above. In addition, I performed a sensitivity analysis in order to assess the effect of the specified priors on the posterior estimates. Specifically, I specified three informative, but arbitrary, priors for each relationship, suggesting the relationship between morphology and the three behavioural variables to be either positive ($\mu = 1, V = 0.2$), non-existent ($\mu = 0, V = 0.2$), or negative ($\mu = -1, V = 0.2$). All analyses conducted in this paper were performed in the software *R* (R Core Team 2014) using the packages *ape* (Paradis et al. 2004), *MCMCglmm* (Hadfield 2010), *phytools* (Revell 2012), *geometry* (Habel et al. 2014), *rcdd* (Geyer & Meeden 2014), *cluster* (Maechler et al. 2015), *adehabitat* (Calenge 2006), and *geiger* (Harmon et al. 2008).

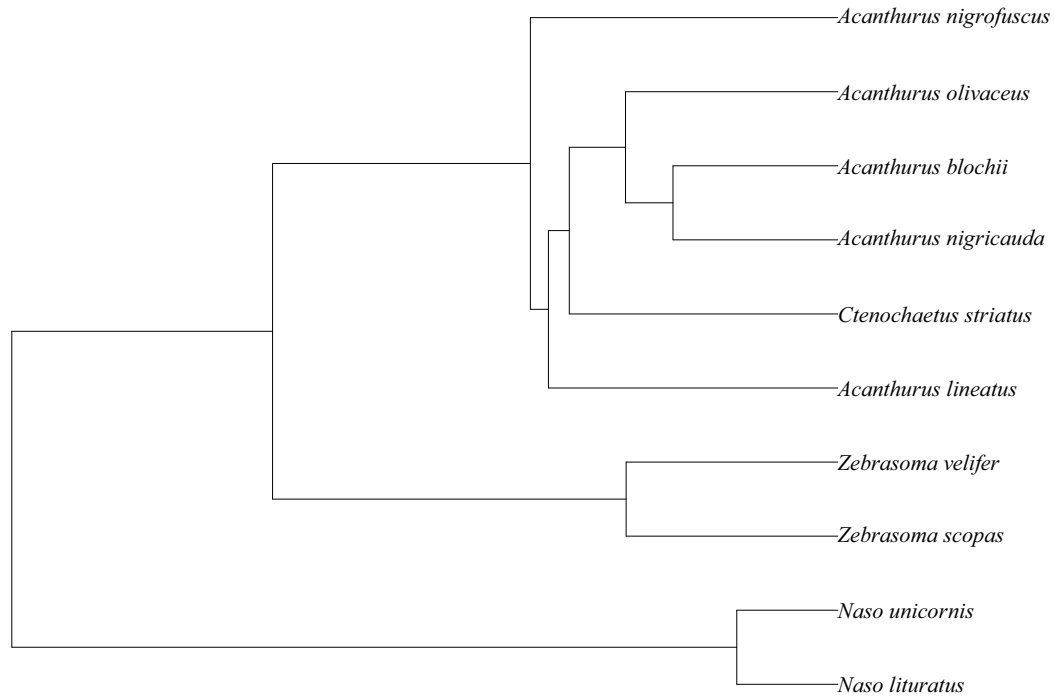


Figure 6.2: Pruned phylogenetic tree after Sorensen et al. 2013, which was used for all phylogenetic corrections.

6.3 Results

I found marked differences in morphological and behavioral specialization patterns among surgeonfish species. Morphologically, *Z. velifer*, *N. unicornis*, and *Z. scopas* were most specialized, exhibiting the greatest distance from the centroid of the biplot (**Figure 6.3**). The least specialized morphologies were found for *C. striatus*, *A. nigricauda*, *A. olivaceus*, and *A. blochii*. In the dietary niche space, *A. blochii* occupied the greatest area, followed by *N. unicornis*. The smallest areas were occupied by *A. olivaceus* and *Z. scopas* (**Figure 6.4a**). In terms of microhabitat utilization (in three dimensions), *Z. scopas* exhibited the largest niche volume, followed by *A. nigrofuscus* and *Z. velifer*. The other members of the genus *Acanthurus* occupied the smaller niche volumes (**Figure 6.4b**). Patterns were similar when using only two dimensions. When accounting for the availability of different microhabitats by using Manly's χ^2 log-likelihood ratio, *Z. velifer* and *A. nigrofuscus* had the lowest specialization scores, while *A. blochii* and *A. nigricauda* had the highest scores, therefore suggesting the latter species to be most specialized (**Figure 6.5**). The most abundant microhabitat in terms of benthic cover was open horizontal rock substratum (29.2%), followed by live coral (16.5%). The most common concealed microhabitat were crevices in horizontal rocky substrata (7.1%). Overall, concealed microhabitats accounted for 24.6% of available foraging microhabitats, while open microhabitats accounted for 75.4%.

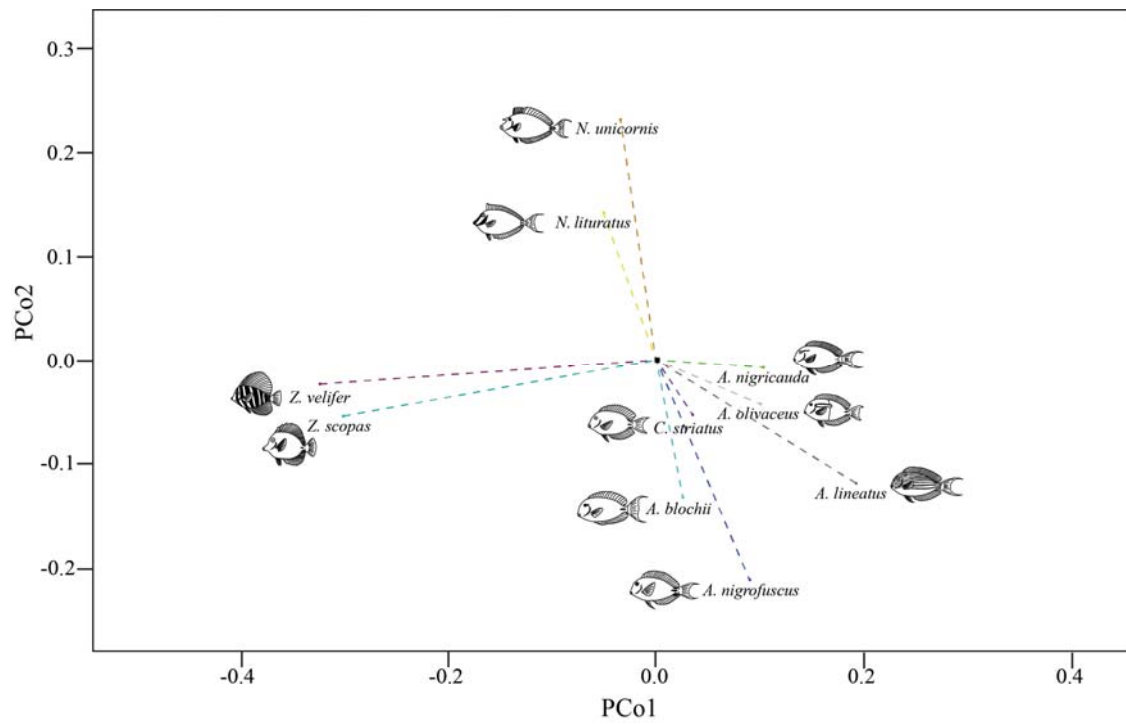


Figure 6.3: Morphological ordination from which species-level morphological specialization was inferred by calculating the distance from the centroid of the ordination.

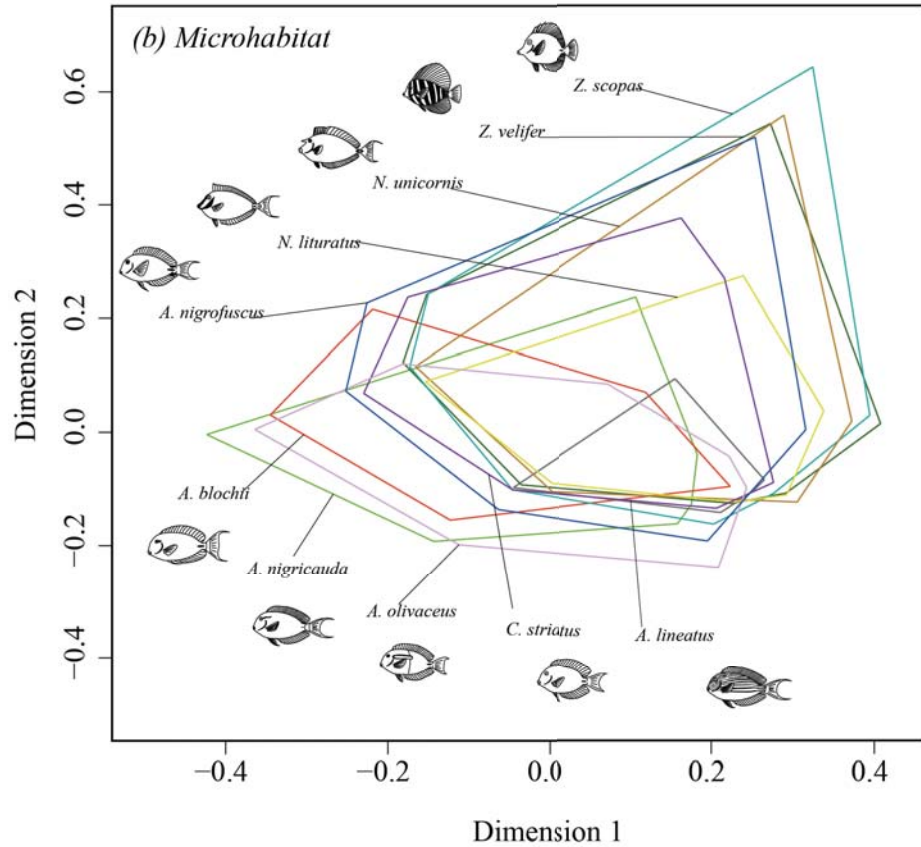
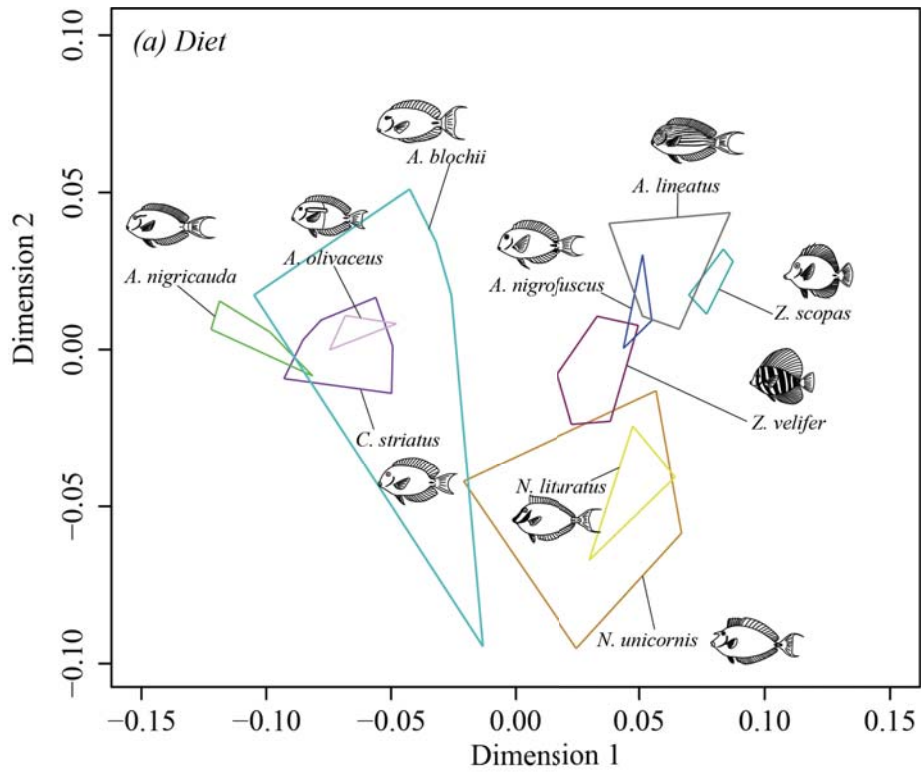


Figure 6.4: Niche volumes for ten species of surgeonfishes in two-dimensional niche space for (b) diet and (c) foraging microhabitat utilization.

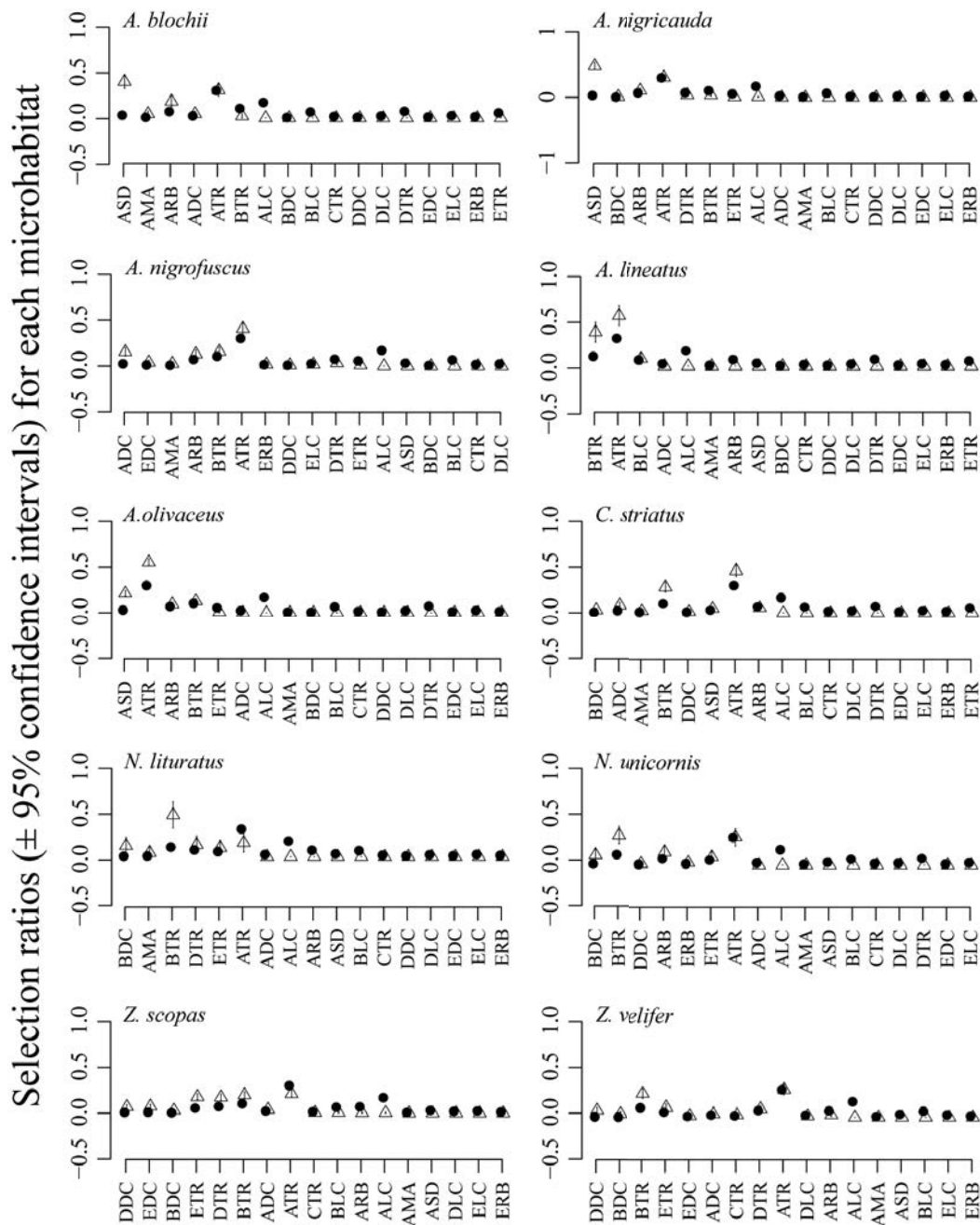


Figure 6.5: Resource selection ratios for respective microhabitats in ten species of surgeonfishes. Open triangles are the usage, while filled circles mark the availability of the respective microhabitat. Acronyms consist of the surface orientation (A,B,C,D,E) and the given substratum type (DC, TR, RB, LC, MA, SD).

Morphological specialization had a markedly different relationship with dietary and microhabitat specialization (**Figure 6.6**), respectively. There was virtually no relationship between morphological and dietary specialization (Bayesian linear model: posterior mean density estimate $\beta = -0.025$, 95% credible interval (CI) = -0.096 to 0.043). In contrast, morphological specialization showed a negative relationship with foraging microhabitat utilization specialization ($\beta = -0.392$, $CI = -0.679$ to -0.106), with morphologically specialized species exhibiting the largest niche volumes, i.e. being the most behaviorally generalized species. There was a similar trend in the relationship between morphology and microhabitat selectivity (Manly's χ^2 log-likelihood statistic), although 95% CIs of the mean posterior estimate also extended beyond zero ($\beta = -1.020$, $CI = -2.876$ to 1.172).

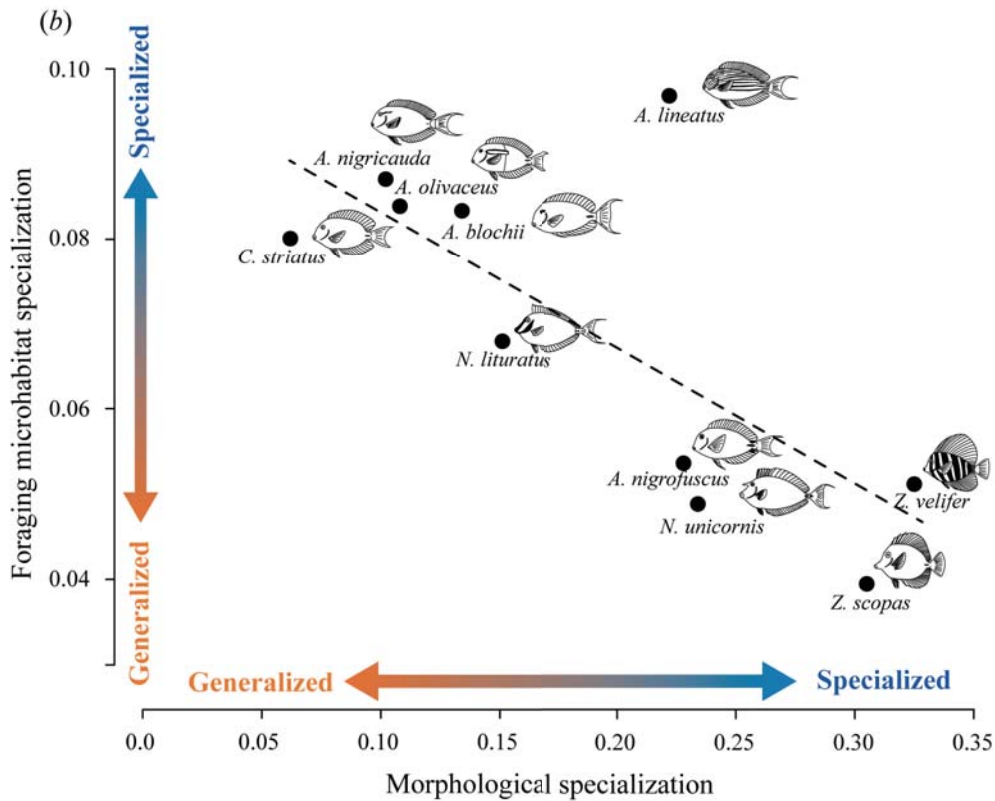
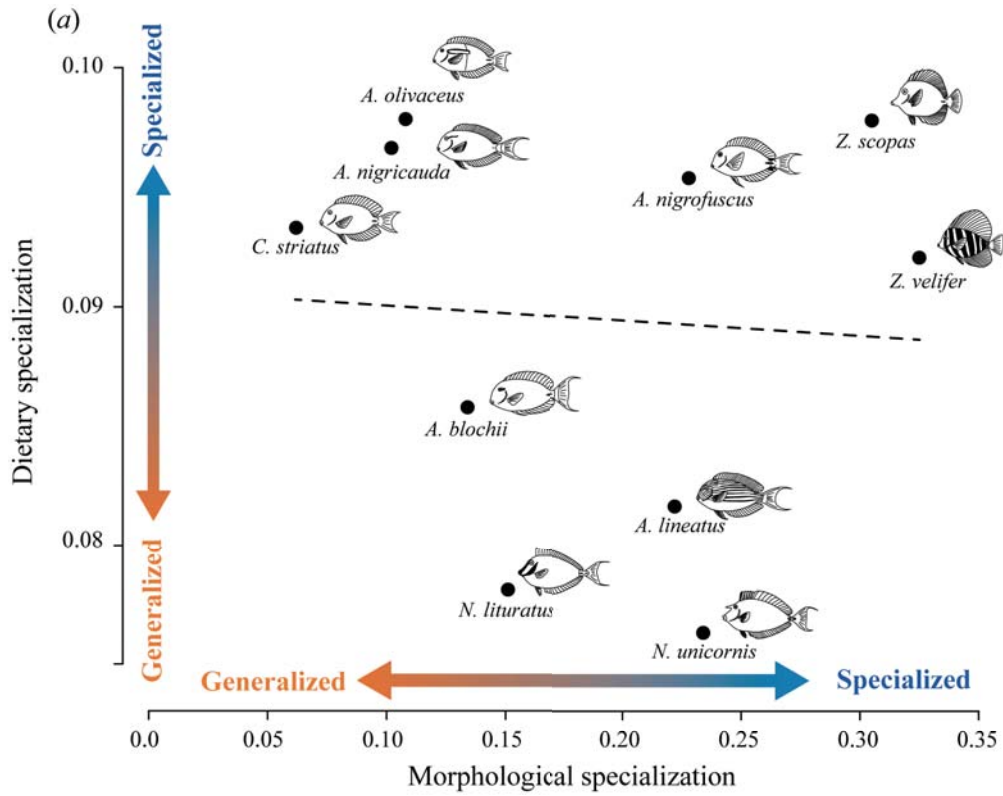


Figure 6.6: Relationships between morphological and behavioural specialization for (a) diet and (b) foraging microhabitat utilization. Morphological specialization and dietary niche space occupation showed no relationship, while there was a strong relationship between morphological specialization and microhabitat niche space occupation. For graphical purposes, behavioural specialization on the Y-axis was inverted by subtracting the occupied niche space value from 0.1.

Accounting for shared ancestry in a morphological PCA and in the phylogenetic BMMs did not change the overall result, although it slightly altered the effect sizes and CIs (**Figure 6.7**). For the relationship between morphological and dietary specialization, the mean posterior estimate for the fixed effect was centered around zero (**Figure 6.7a**) (phylogenetic BMM: $\beta = -0.020$, $CI = -0.081$ to 0.039), with both the 95% and 50% CIs intersecting zero. The relationship between morphological specialization and microhabitat specialization (based on species' niche volumes) was less diffuse (**Figure 6.7b**) with consistently negative estimates for the mean ($\beta = -0.294$, $CI = -0.527$ to -0.084) despite a slight shift towards zero compared to the uncorrected model. When regressing morphological specialization against Manly's χ^2 log-likelihood statistic of resource specialization (figure 3c), the relationship also showed a negative trend but 95% CIs did intersect zero ($\beta = -0.929$, $CI = -2.117$ to 0.326).

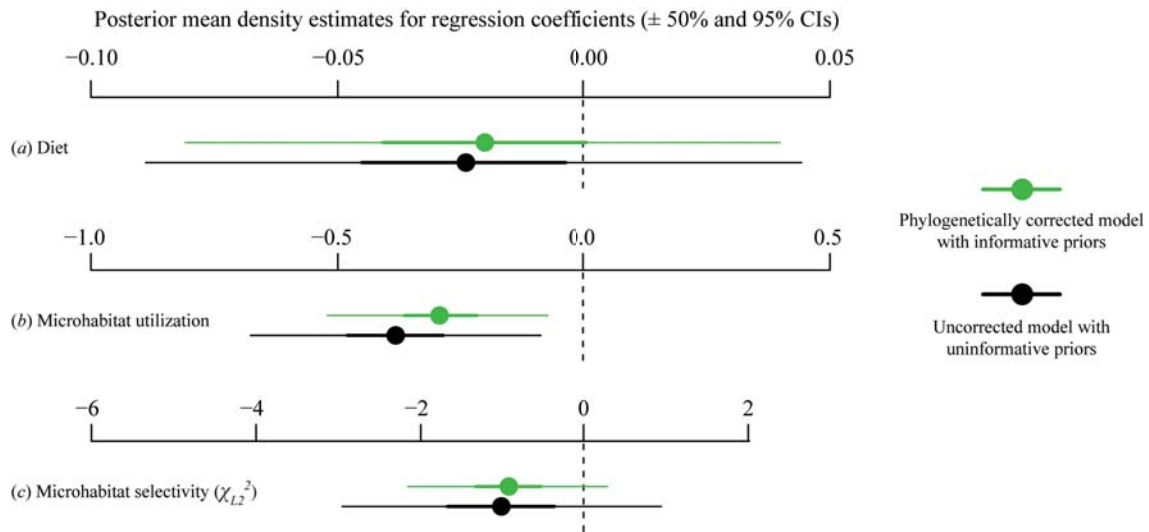


Figure 6.7: Comparisons of models estimating the relationship between morphological specialization and (a) dietary specialization, (b) foraging microhabitat specialization, and (c) microhabitat selectivity. Plots show the mean posterior density estimates of regression coefficients and their 50% and 95% CIs in phylogenetically corrected (green) and uncorrected models (black). Relationships between morphological and behavioural specialization were consistent after phylogenetic corrections, with no relationship between morphological and dietary specialization (a). In contrast, a clear negative relationship is seen between morphological and microhabitat utilization specialization regardless of phylogenetic correction (b). A possible negative relationship is seen between morphological specialization and microhabitat selectivity; however, 95% CIs extend beyond zero (c).

While the results from the phylogenetically corrected BMMs indicate that substantial information for the posterior estimates came from the specified priors, the decrease of the CIs surrounding the posterior estimates in the phylogenetic BMMs suggests that the data did support the initial findings (**Figure 6.7**). This is further supported by the results of the sensitivity analysis (**Figure 6.8, Table 6.3**). For the relationship between morphological and dietary specialization, none of the models yielded a clear trend, regardless of the specified priors. In contrast, an unambiguous negative relationship between morphological specialization and microhabitat specialization was present in the models featuring the negative priors, while there was no clear relationship when neutral or positive priors were specified. The same pattern was present for the relationship between morphological specialization and Manly's χ^2 log-likelihood statistic of resource specialization, with a clear relationship only found when a negative prior was used. Thus, although the data did not override the neutral/positive priors, for both aspects of microhabitat specialization, the models with the negative priors resulted in a clear negative relationship in the posterior, suggesting that the data supported such negative relationship. Therefore, although small sample sizes may limit the strength of the support, the main suggestion of my analyses is that morphological specialization is negatively related to microhabitat utilization specialization, while dietary specialization shows no relationship with morphology.

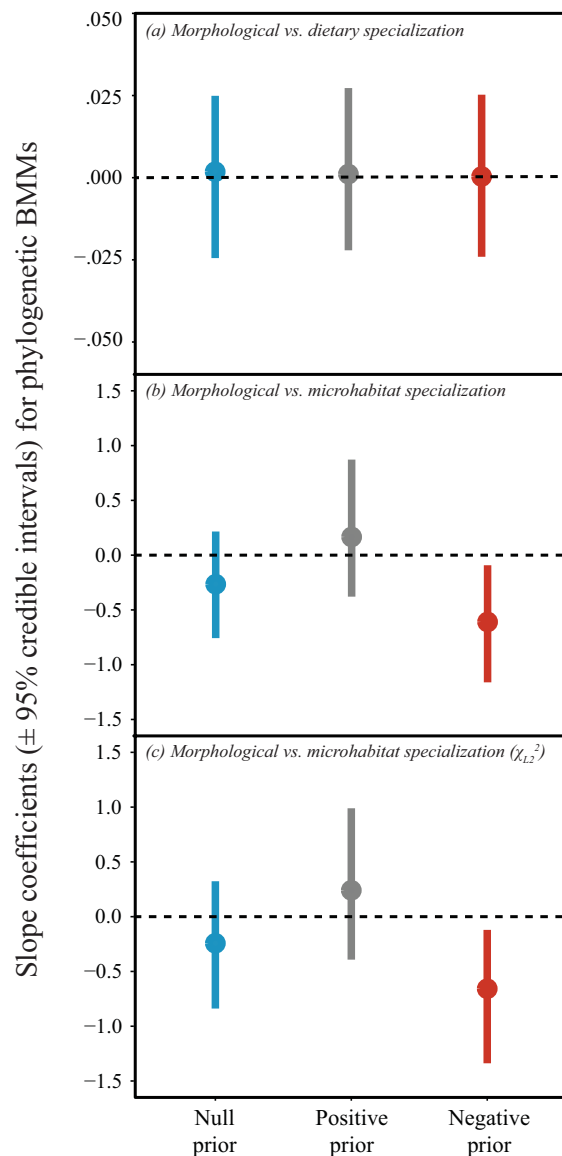


Figure 6.8: Results of the sensitivity analysis. The plot shows the mean posterior estimates of phylogenetic Bayesian Mixed Models (BMMs) testing the effect of morphological specialization on dietary and foraging microhabitat specialization (niche volumes and Manly's χ^2 log-likelihood statistic). The y-axis denotes the mean posterior estimate for the slope parameter. Three different priors were used to simulate prior beliefs about the relationship between morphological and behavioural specialization, specifically a) no relationship ("null prior"), b) a positive relationship

(i.e. morphological specialists are behavioural specialists, “positive prior”), and c) a negative relationship (i.e. morphological specialists are behavioural generalists, “negative prior”). For both metrics of foraging microhabitat specialization, only the negative prior yielded an unambiguous negative relationship. No effect was found for dietary specialization regardless of the used prior.

Table 6.3: Prior specifications and posterior parameter estimates for sensitivity analysis. Symbol indicates the nature of the prior (no relationship, positive, negative).

Relationship modeled	Direction	Fixed effects prior: intercept (α)	Fixed effects prior: slope (β)	Mean posterior slope estimate	Lower/upper 95% CIs
Morphology vs. dietary niche volume	(0)	$\text{Pr}(\mu) \sim \text{N}(0, 0.2)$	$\text{Pr}(\mu) \sim \text{N}(0, 0.2)$	0.002	-0.03 / 0.03
	(+)	$\text{Pr}(\mu) \sim \text{N}(0, 0.2)$	$\text{Pr}(\mu) \sim \text{N}(1, 0.2)$	0.001	-0.02 / 0.03
	(-)	$\text{Pr}(\mu) \sim \text{N}(1, 0.2)$	$\text{Pr}(\mu) \sim \text{N}(-1, 0.2)$	0.0004	-0.02 / 0.03
Morphology vs. microhabitat niche volume	(0)	$\text{Pr}(\mu) \sim \text{N}(0, 0.2)$	$\text{Pr}(\mu) \sim \text{N}(0, 0.2)$	-0.265	-0.76 / 0.22
	(+)	$\text{Pr}(\mu) \sim \text{N}(0, 0.2)$	$\text{Pr}(\mu) \sim \text{N}(1, 0.2)$	0.166	-0.38 / 0.87
	(-)	$\text{Pr}(\mu) \sim \text{N}(1, 0.2)$	$\text{Pr}(\mu) \sim \text{N}(-1, 0.2)$	-0.610	-1.16 / -0.09
Morphology vs. microhabitat selectivity ($\chi_{1,2}^2$)	(0)	$\text{Pr}(\mu) \sim \text{N}(0, 0.2)$	$\text{Pr}(\mu) \sim \text{N}(0, 0.2)$	-0.243	-0.84 / 0.32
	(+)	$\text{Pr}(\mu) \sim \text{N}(0, 0.2)$	$\text{Pr}(\mu) \sim \text{N}(1, 0.2)$	0.241	-0.39 / 0.99
	(-)	$\text{Pr}(\mu) \sim \text{N}(1, 0.2)$	$\text{Pr}(\mu) \sim \text{N}(-1, 0.2)$	-0.658	-1.34 / 0.12

6.4 Discussion

6.4.1 Ecological specialization in surgeonfishes

Links between morphological specialization and dietary strategies are widespread, supporting the paradigm that specialized morphological features are related to dietary specialization (Forister et al. 2012; Ferry-Graham et al. 2002a; Douglas & Matthews 1992). However, I found no such relationship in coral reef surgeonfishes, supporting a small number of similar studies in teleost fishes (Liem, 1980; Motta et al. 1995; Barnett et al. 2006; Bellwood et al. 2006; Wainwright et al. 2005). Basically, morphologically similar surgeonfish species showed marked differences in the range of food items ingested (e.g. *A. blochii* vs. *A. nigricauda*), while morphologically disparate species shared a quantitatively similar dietary range (*A. olivaceus* vs. *Z. scopas*). These results call for caution when identifying functional groups based solely on morphological similarity or phylogenetic relatedness: realized niches need to be examined carefully if we are to understand and monitor ecosystem functioning of coral reefs (Bellwood et al. 2004).

In contrast, I found an interesting, negative relationship between morphological specialization and foraging microhabitat preferences in surgeonfishes. While foraging microhabitat utilization has been identified as an important axis of niche partitioning in fishes (Brandl & Bellwood 2014a), few ecomorphological studies have considered the relationship between microhabitat specialization and morphological specialization in a broad quantitative framework (but see Snorrason et al. 1994). For surgeonfishes, it has been proposed that certain morphological characteristics such as a small and deep body, large eyes, a truncate caudal fin, or a concave forehead are linked to the exploitation of concealed resources and that these features enable species to utilize a broad range of resources (Brandl & Bellwood 2013a; Robertson & Gaines 1986). This was supported by my results: morphologically specialized species are microhabitat generalists, while morphological generalists are foraging

microhabitat specialists. Two extreme examples are *Z. scopas* and *A. nigricauda*. While *Z. scopas* exhibits a highly specialized morphology, distinguishing it from most other surgeonfish species, this species foraged over the widest range of microhabitats. In contrast, despite exhibiting the generalized *Acanthurus* morphology, *A. nigricauda* almost exclusively targeted flat and exposed rocky substrata.

Furthermore, these patterns are possibly consistent, when the availability of microhabitats is accounted for. For example, the three closely related species *A. blochii*, *A. nigricauda*, and *A. olivaceus* forage almost exclusively on flat horizontal surfaces and although these surfaces are readily available, their disproportional use of open sand, rubble, and rock microhabitats results in a high level of selectivity. In contrast, species such as *Z. velifer* and *A. nigrofuscus* appear to distribute their foraging efforts relatively evenly across available microhabitats, resulting in low selectivity scores for these species. Such patterns of specific habitat or microhabitat use by specialized species has been reported previously in parrotfishes (Bellwood & Choat 1990), but not to the extent seen in surgeonfishes.

The links between morphological and behavioural specialization in surgeonfishes are, in part, driven by phylogenetic relatedness. For example, the two species of *Zebrasoma* are closely related to each other, but phylogenetically distinct from most species within the *Acanthurus* clade (Sorenson et al. 2013). It is therefore not surprising that the two *Zebrasoma* species exhibit a similar morphology and foraging strategies but differ markedly from most of the *Acanthurus* species. Despite this clear influence of shared ancestry, the phylogenetically corrected BMMs suggest that the observed relationships are robust and still supported when phylogeny is accounted for. Thus, there appears to be no relationship between morphological and dietary specialization, while the negative relationship between morphological and microhabitat specialization is consistent for the overall niche volume measures and, to a lesser extent, possibly for microhabitats selectivity (Manly's χ^2 log-

likelihood ratio). However, given the small sample size of the phylogenetically corrected dataset and the influence of the used priors on the posterior estimates (revealed in the sensitivity analysis), the broader validity of these findings will have to be confirmed using a more extensive sampling protocol, ideally across a broader phylogenetic sample.

Interestingly, the described relationship between morphological and foraging microhabitat specialization is strikingly similar to patterns found in terrestrial plant-pollinator networks. In both arthropod and avian pollinator communities, species with “specialized” morphologies such as long probosces or bills have been identified as generalists, feeding on a wide range of flower types, while species with shorter mouthparts are restricted to flowers with shallow tubes (Borrell 2005; Temeles et al. 2009). These ecological parallels may not be coincidental. Algal and detrital resources on coral reefs and nectar in plant-pollinator communities are both replenished at high rates (Thomson et al. 1989; Bonaldo & Bellwood 2011) and form a highly diverse network of resources with varying accessibility (Jordano et al. 2003). Furthermore, similar to flowers with long tubes, which provide high rewards in terms of available nectar (Gómez et al. 2008), inaccessible microhabitats appear to harbor richer algal resources (Brandl et al. 2014), potentially offering higher rewards than flat microhabitats. Thus, although the two systems exhibit different consumer-producer relationships (mutualism vs. prey-predator relationships; Johnson & Steiner 2000), the ecological processes shaping both plant-pollinator and herbivore-benthos dynamics may be more similar than previously thought. This further suggests that the relationship between morphological and foraging microhabitat specialization present in both systems may be more broadly applicable.

6.4.2 The role of microhabitat specialization for the evolution of coral reefs

The observed pattern in surgeonfishes is superficially consistent with a phenomenon termed Liem's paradox. While the idea of Liem's paradox operating on coral reefs seems appealing in light of the extraordinary diversity present in reef fishes, there are theoretical inconsistencies in the context of my findings. The most popular evolutionary framework for Liem's paradox posits that the basic strategy for every species is to forage on an easily exploitable resource, which is subsequently followed by 'secondary' specialization on less abundant or accessible resources (Snorrason et al. 1994). In the Acanthuridae, however, specialized morphotypes similar to *Zebrasoma* or *Naso* are recorded from coral reefs in the Eocene 60 Ma (Bellwood et al. 2014), suggesting that microhabitat generalists have been present on reefs throughout most of their evolutionary history. Furthermore, although generalized morphs were also present in the Eocene (Sorenson et al. 2013), the expansion of morphological generalists specialized on flat microhabitats appears to have been a later trend (rapidly increasing species richness and relative abundance in the Miocene ca. 25 Ma), questioning the applicability of Liem's paradox in this case. Interestingly, reef parrotfishes which predominantly target open microhabitats (Price et al. 2010; Brandl & Bellwood 2014a), also appeared on scleractinian-dominated reefs around the same time as surgeonfishes of the genus *Acanthurus* (Choat et al. 2012), supporting the hypothesis that increased exploitation of open microhabitats occurred predominantly in the Miocene. It appears probable that morphological specialists added the exploitation of open microhabitats to their initial feeding repertoire and that the subsequent major radiations in surgeonfish lineages specialized on flat open microhabitats may be due to the increased availability of these foraging surfaces.

There are three non-exclusive hypotheses that may explain this trend: first, mirroring recent trends towards the loss of topographical complexity on coral reefs (Alvarez-Filip et al. 2009), coral reefs may have undergone a transition from complex, micro-topographically

intricate systems to flatter, less complex systems over the last fifty million years through increasing rates of external bioerosion (Bonaldo et al. 2014), resulting in an increase in the proportion of planar surfaces with easily accessible resources for herbivorous/detritivorous fishes. Second, biotic or abiotic processes such as sea level fluctuations or increasing calcification and accretion may have increased the extent of reef flats, a habitat that frequently supports the largest area of hard, open surfaces on modern coral reefs (Bellwood & Wainwright 2002) in the most productive zone of primary production (Steneck 1988). Finally, reef flats may have been available throughout the last 60 Ma, but the establishment of a nutritionally attractive epilithic algal matrix and its associated detrital resources may have only arisen once coral reef fishes increased grazing intensities throughout their radiation in the Miocene (Bellwood et al. 2014; Steneck 1983; Cowman & Bellwood 2013), leading to the evolution of specialized herbivores foraging exclusively on flat open surfaces. In all three cases, the expansion of specialized open-microhabitat foraging species appears to have been an essential step in the establishment of modern herbivorous coral reef fish assemblages.

6.4.3 Significance, limitations, and future perspectives

Ecomorphological assessments attempting to relate morphological specialization to dietary specialization have been invaluable for our understanding of the ecology and evolution in animals. However, there is increasing evidence that the relationship between morphological specialization and dietary preferences is multifaceted and often less clear and linear than expected. I do not suggest that the link between morphological specialization and diet in reef fishes (or, more specifically, surgeonfishes) is absent; such a conclusion would be unwarranted given the potential for more detailed assessments of this relationship (such as a higher resolution classification of morphology or dietary items or the mapping of the latter onto different microhabitats). However, my study serves as a cautionary note, suggesting that

there is a potential for more than one ecologically significant axis that one may find reflected in a species' morphology.

Our findings present an intriguing case for the exploration of foraging microhabitats as a potentially important niche axis in benthic feeding reef fishes. Although limited by the relatively small sample size of ten species with shared ancestries within a large lineage, the results suggest that the identified pattern may persist if the assessment is expanded. Furthermore, the list of factors that may influence the foraging behavior and microhabitat selection of reef fishes is long: exploring the range of dietary items present in various microhabitats and their rate of replenishment, a finer classification of dietary items, or nutritional analyses of resources in different microhabitats and their post-ingestive utilization by both specialists and generalists would offer interesting avenues for future research and may help uncover more detail about the dynamics between benthic organisms and mobile consumer communities on coral reefs. Here, I am only scraping the surface of a largely untouched ecological niche axis. Nevertheless, my findings, particularly regarding the similarities between plant-pollinator communities and coral reef herbivores and the superficial parallels to Liem's paradox, are encouraging for future investigations of foraging microhabitat utilization on coral reefs.

6.4.4 Conclusion

Our study shows that morphological specialization and its behavioural correlates are not as easily inferred as frequently assumed. I demonstrate that there is no evidence for a correlation between morphological specialization and dietary specialization, but reveal a potential link between morphological specialization and foraging microhabitat specialization. However, this relationship is inverted: species with specialized morphologies are foraging microhabitat generalists while generalized morphotypes are behaviorally specialized on exposed

microhabitats. These results suggest that we may need to look beyond the most commonly investigated links between species' morphological and behavioural niches and that on coral reefs, morphological specialization may be tightly intertwined with foraging microhabitat utilization. Consequently, the use of different microhabitats may be an important ecological axis to understand the past, present, and future of coral reefs.

Chapter 7: Micro-topography mediates interactions between corals, algae, and herbivorous reef fishes on a mid-shelf reef

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7.1 Introduction

The last two to three decades have seen the degradation of many coral reef systems around the world. Although subject to various stressors, degraded reef systems are commonly characterized by marked changes to the benthic community composition (Bellwood et al. 2004; Norström et al. 2009). One of the most prominent is a shift from scleractinian corals to macroalgae as the dominant benthic component (Hughes 1994; Cheal et al. 2010; Rasher et al. 2013). If we are to understand the basis of such transitions, it is essential to understand the population dynamics of both corals and macroalgae, and the factors that shape their interactions (McCook et al. 2001; Vermeij et al. 2009; Sotka & Hay 2009; Diaz-Pulido et al. 2010).

The life history and population dynamics of scleractinian corals have attracted much attention. Due to their planktonic larval stage, coral populations are largely shaped by the addition of new individuals from the pelagic environment (i.e., settlement) and early post-settlement mortality (Baird & Hughes 1997; Hughes et al. 1999; Mundy & Babcock 2000; Ritson-Williams et al. 2010). The settlement of corals has been studied in both the field and laboratory using experimental settlement surfaces (e.g., terracotta tiles), which are readily colonized by coral planulae (e.g., Harriott & Fisk 1987; Mundy 2000; Penin et al. 2010). These studies suggest that coral planulae actively choose microhabitats based on environmental factors, including exposure (Maida et al. 1994), depth (Baird et al. 2003) or the presence and type of crustose coralline algae (CCA) (Raimondi and Morse 2000;

Harrington et al. 2004). Similarly, by settling recruits on experimental tiles and transplanting the tiles onto the reef, many studies have investigated spatial variation in post-settlement survival of coral spat (e.g., Babcock & Mundy 1996; Ritson-Williams et al. 2010; Penin et al. 2011). The most frequently identified environmental factors influencing the post-settlement survival of juvenile or newly settled corals include sedimentation, grazing pressure by herbivorous fishes or invertebrates, and competition with other benthic organisms, particularly algae (Sammarco 1980; Hodgson 1990; Birrell et al. 2008).

Several studies have emphasized the importance of algal competition and herbivorous grazing in shaping post-settlement mortality in corals (Mundy and Babcock 2000; Arnold et al. 2010; Penin et al. 2010; Trapon et al. 2013a,b), identifying three main types of interaction. Firstly, macroalgae can be detrimental for the settlement and survival of coral recruits as they compete for space (e.g., Box & Mumby 2007). Although the mechanisms of algal interference with coral juveniles are not well resolved (Vermeij et al. 2009), and may depend on algal and coral identity (Birrell et al. 2008), mechanical and chemical/allelopathic defenses of macroalgae do appear to play an important role (Rasher et al. 2011). Secondly, coral reef herbivores control algal growth and can facilitate coral survival and replenishment by removing algal competitors and opening space for settlement (Hughes et al. 2007; Arnold et al. 2010; Rasher & Hay 2010; Rasher et al. 2012, 2013). Finally, herbivores may also increase mortality of small or recently settled corals via incidental grazing. This has been reported for both echinoid grazers (e.g., Sammarco 1980) and scraping parrotfishes (e.g., Brock 1979; Day 1983; Penin et al. 2010, 2011). Given the influence of algae and herbivores on the post-settlement survival of juvenile corals, it may be postulated that algal competitors and grazing fishes would play a key role in shaping abundance patterns of juvenile corals. However, their relative importance in shaping patterns of settlement and post-settlement mortality remains to be determined, especially for corals on natural reef substrata.

It is often assumed that many corals settle in cryptic habitats to avoid mortality due to incidental grazing, despite intense competition for space in these concealed microhabitats (Day 1983; Carlon 2001; Edmunds et al. 2004). In most experimental frameworks, coral planulae settle preferentially to cryptic habitats (i.e., the undersurfaces of tiles) as opposed to open, exposed surfaces (Maida et al. 1994; Babcock & Mundy 1996; Vermeij 2005; Arnold et al. 2010). Further, the availability of micro-crevices on settlement tiles has been shown to significantly increase the survival of corals in the first year post-settlement (Nozawa 2008). However, in the few observational field studies that have reported microhabitat associations on natural reef surfaces, juvenile corals have been found to be most abundant on open surfaces (Florida Keys: Edmunds et al. 2004; Palmyra Atoll: Roth & Knowlton 2009; Great Barrier Reef: Trapon et al. 2013a). This apparent disparity between coral settlement preferences on artificial substrata and observations of micro-habitat associations on natural reef substrata may be related to the relative size of the crevices, differential settlement preferences and/or juvenile survival patterns, or be an artefact of the experimental set-ups. For example, coral larvae may settle into micro-crevices and then outgrow the micro-crevice to become associated with more open substrata as a juvenile. Alternatively, the common use of artificial surfaces in experimental studies and the exclusion of extrinsic factors such as competition or predators under experimental conditions may have contributed to differences between experimental and observational studies. There is a clear need to understand the role of these extrinsic factors in shaping juvenile coral assemblages.

Of the observational studies assessing the microhabitat associations of juvenile coral none included distributions of potential competitors, and only one study (Trapon et al. 2013a) evaluated the intensity of herbivorous grazing on juvenile coral survival. Grazing pressure on juvenile corals is believed to be an important source of mortality and is commonly associated with echinoids and / or scraping and excavating parrotfishes (f. Labridae). The vast majority

of studies assessing the effect of herbivores on juvenile corals do not distinguish the relative impact of the various herbivorous groups (Brock 1979; Penin et al. 2010, 2011; but see Sammarco 1980; Trapon et al. 2013a). Yet herbivorous fishes differ markedly in their diet and feeding mode and can have highly dissimilar impacts on the reef substratum and its benthic community (Bellwood & Choat 1990; Burkepile & Hay 2008, 2011; Green and Bellwood 2009; Rasher et al. 2013; Hoey et al. 2013), including corals (Bonaldo and Bellwood 2011). As such, parrotfishes are more likely to cause incidental damage through non-selective scraping of the reef substratum, while the more selective removal of algal filaments by cropping and browsing species is less likely to affect coral juveniles (Trapon et al. 2013b). It thus appears necessary to evaluate the functional relationship between corals, algae and individual taxa of herbivores, rather than examining the distribution of herbivorous fishes *per se*, in order to explore their possible impact on benthic organisms (Baria et al. 2010).

The aims of the present study, therefore, were to 1) quantify the distribution patterns of juvenile corals and early life-stages of a potential macroalgal competitor, *Turbinaria ornata* (Gleason 1996; Shearer et al. 2012) with regards to microhabitats and reef zones, and 2) to examine their distribution in relation to abundance patterns of large grazing herbivorous fishes.

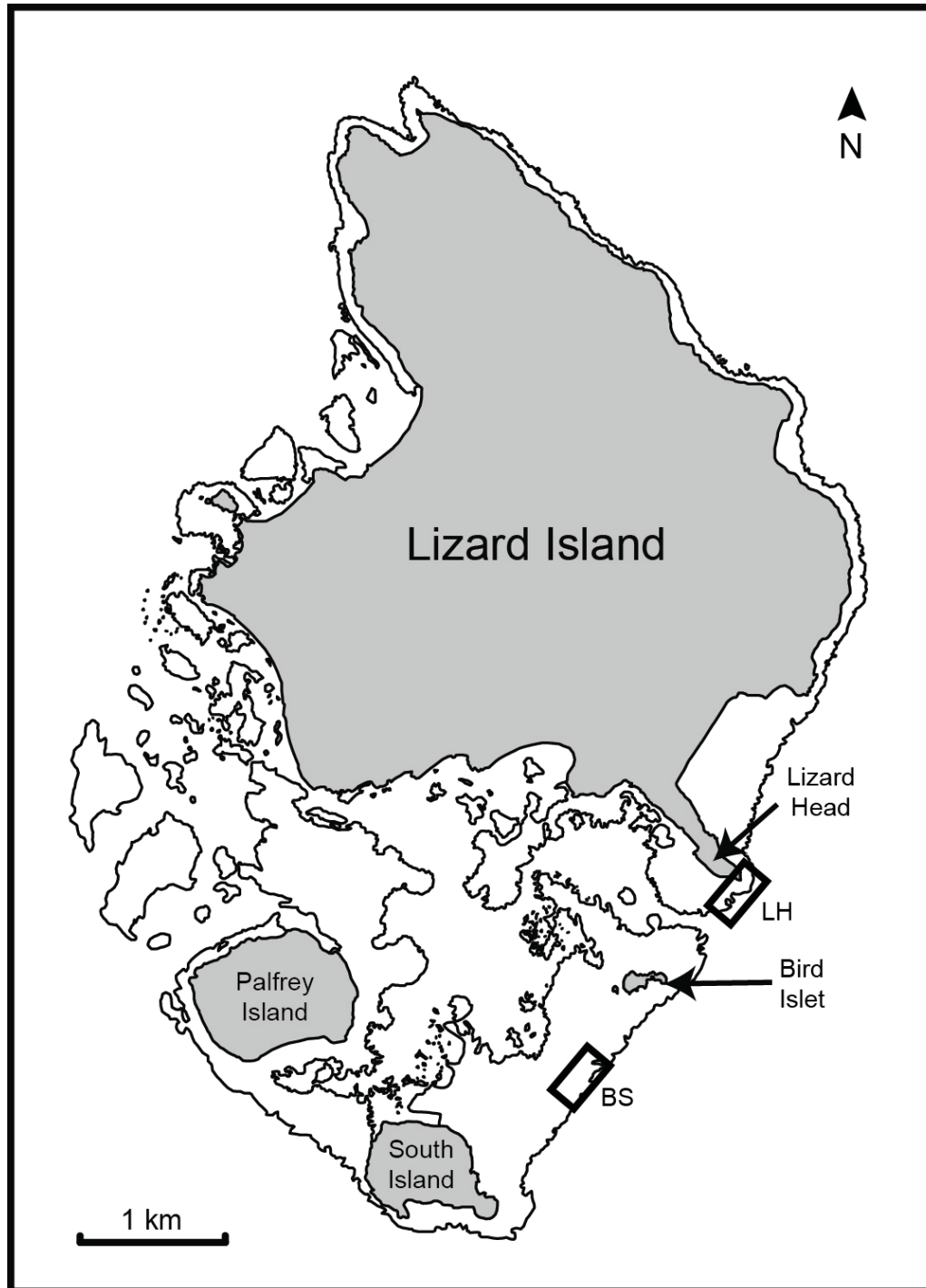


Figure 7.1: Map of Lizard Island. The two sites (BS = Bird – South; LH = Lizard Head) are marked by rectangles.

7.2 Materials and methods

The study was conducted on reefs around Lizard Island, a mid-shelf island in the Northern Great Barrier Reef, Australia (14°40'08"S 145°27'34"E). Two reef sites were examined, between Bird Islet and South Island (BS) and Lizard Head (LH) (**Figure 7.1**). Both sites are typical exposed reef systems and display distinct topographic zonation patterns. This includes a reef base, a distinct reef crest and a low complexity reef flat. The reef crest at both sites was marked by high adult coral cover and low macroalgal abundance, while the reef flat in both areas was characterized by low adult coral cover and the presence of large, erect macroalgae, particularly the fleshy brown alga *T. ornata* (Fucales, Phaeophyceae). Juvenile corals and algal thalli were quantified in two zones, the reef crest and the outer reef-flat (approximately 10 – 15 m behind the crest). These zones were chosen because they offer the highest potential for interactions between juvenile corals and macroalgae, as both adult coral colonies and macroalgal thalli were present in the immediate vicinity.

7.2.1 Benthic transects

To quantify the distribution of juvenile corals and macroalgae, five quadrats (1m²) were placed haphazardly along a 50 m belt transect, laid parallel to the reef crest. Five transects were laid in each reef zone (crest, outer flat), at each site (BS, LH) resulting in a total of 20 transects and 100 quadrats. In each quadrat, juvenile corals (< 50mm diameter; following Rylaarsdam 1983) and early life-stage macroalgal thalli (< 50mm high) were counted using SCUBA. Due to low abundances and difficulties in reliably distinguishing between juveniles and partial mortality remnants in massive corals, the juvenile coral assessment was restricted to the families Acroporidae and Pocilloporidae. For the investigation of macroalgal distribution patterns, *T. ornata* was chosen, as it was by far the most abundant macroalgal species in the area (> 95% of all fleshy macroalgae observed) and allowed for reliable

identification of early life-stages based on the height of the thalli. The microhabitat of every sampled juvenile coral colony or macroalgal thallus was characterized as concealed, semi-concealed or open. Microhabitats were considered concealed if the colony / thallus was found within a feature that was only accessible from one direction (i.e., holes or crevices). Semi-concealed habitats had access from two different directions (i.e., grooves or ledges), while in open microhabitats, the colony / thallus was fully exposed and accessible from every lateral direction within 20 cm of the colony / thallus. Grooves are defined as long concavities accessible from the top and two sides while ledges are defined as steps in the substratum with the organism growing in the angle. Concealed and semi-concealed microhabitats represented less than 10% of the total available area within all quadrats examined. All sampled colonies or thalli were measured using vernier calipers (corals: length, width and height with the 50mm cutoff based on mean colony diameter; algae: height) and photographed. Each quadrat was searched systematically for a minimum of 15 minutes and from four different angles to ensure standardized, comprehensive sampling of juvenile coral colonies and *T. ornata* thalli. Each *T. ornata* thallus recorded was examined for any evidence of feeding by macro-herbivores. Feeding events were readily identified by hemispherical bite marks and clearly severed blades.

7.2.2 Fish transects

The herbivorous fish community was assessed using underwater visual censuses along haphazardly laid 50 x 5 m belt transects in the same reef zones and sites as above (5 transects per zone at each site). All large mobile, herbivorous fishes (families Acanthuridae, Labridae, Siganidae) were counted while deploying the transect tape, to minimize disturbance (Dickens et al. 2011). Smaller, sedentary herbivores such as territorial pomacentrids were not included as they were extremely uncommon at the examined sites (cf. Ceccarelli et al. 2005; Hoey &

Bellwood 2010a). Fishes were identified to species and placed in six 5 cm size classes (10 – 15cm etc. to 35cm and > 35cm) based on visual assessments. Above 35 cm total length, a size estimate was recorded to allow for a more accurate assessment of fish biomass. All observations were performed by the same trained observer (SJB) between 1000 and 1600 hrs.

7.2.3 Statistical analysis

The distribution patterns of juvenile corals and macroalgae were analyzed using a generalized linear mixed effects model (GLMM) with a Poisson error distribution and log-link function. GLMMs were chosen as the data violated parametric assumptions even after transformation. Site, zone and microhabitat were treated as fixed effects, while transects were treated as a random factor. Deviance statistics were generated for different models and Chi-squared tests were implemented on the deviance statistics to compare the significance of explanatory variables in the models. GLMMs were applied to coral and macroalgae counts separately. Models were selected using the Akaike information criterion (AIC). In addition, the occurrence of bite marks on macroalgal thalli was compared between the crest and the outer flat using Pearson's Chi-squared test. To visualize results and obtain a distance matrix for the BEST ENV-BIO routine (see below), a non-metric multidimensional scaling analysis was performed using Euclidean distances of juvenile corals and early life-stage macroalgae in different microhabitats, grouped by transects.

Fish abundance data were converted to biomass 250 m^{-2} (using published length-weight relationships; following Hoey & Bellwood 2009), and then differences between sites and zones were analyzed using a two-way analysis of variance (ANOVA). Assumptions of the ANOVA were tested using a Shapiro-Wilk test for normality of residuals and Bartlett's test for homogeneity of variances. The species community composition within transects was

analyzed using non-metric multidimensional scaling of a Bray-Curtis distance matrix of the abundance data.

Subsequently, a BEST ENV-BIO routine was performed using the two distance matrices for coral juveniles / macroalgae and fish abundance (following Clarke and Ainsworth 1993; Clarke & Warwick 2001). This procedure allows for the exploration of relationships between two distance matrices, one of which being patterns observed in selected biota (i.e., coral and algal distribution patterns) while the other distance matrix is based on extrinsic or environmental factors (in this case abundance of herbivorous fish species). To this end, a Spearman's rank correlation coefficient (ρ) between the two matrices is calculated for every combination of variables, extrapolating the combination of extrinsic factors (i.e. a suite of herbivorous fish species) that best predicts the observed distribution patterns. The significance of explanatory variables was assessed using 999 randomized permutations under the null hypothesis that there is no correlation of variables between distance matrices (Clarke et al. 2008). The BEST ENV-BIO routine was performed using the software PRIMER. All other statistical analyses were performed using the software R and the packages *lme4* and *MuMIn* (R Development Core Team 2012).

7.3 Results

Overall, 424 juvenile coral colonies and 122 juvenile macroalgae thalli were recorded across the study area. The mean number of juvenile corals per quadrat was 4.2 ± 0.3 colonies m^{-2} (SE) and early life-stage *T. ornata* thalli was 1.2 ± 0.2 thalli m^{-2} (SE).

7.3.1 Distribution patterns of juvenile coral colonies

The majority of juvenile corals were found in concealed microhabitats and were most abundant on the reef crest (**Figures 7.2, 7.3**). Concealed microhabitats supported more than

three times more colonies (2.9 ± 0.2 colonies m^{-2} SE) than semi-concealed (0.8 ± 0.1 m^{-2} SE) or open (0.5 ± 0.1 m^{-2} SE) microhabitats. Coral juveniles were almost twice as abundant on the reef crest (5.5 ± 0.4 colonies m^{-2} SE) than on the outer flat (3.0 ± 0.3 colonies m^{-2} SE). This was reflected in the GLMM, where the outer flat ($P < 0.001$) and semi-concealed and open microhabitats ($P < 0.001$) showed a significant negative association with the abundance of juvenile corals. The model including only reef zone and microhabitat performed better than a GLMM including site as a fixed factor ($\chi^2 = 0.8$, $P = 0.4$) and/or a site \times zone interaction term ($\chi^2 = 2.5$, $P = 0.3$, **Table 7.1**), suggesting that sites had no significant effect on the observed distribution patterns. In only 1.6% of all records (6 occasions) were juvenile corals and early life-stage *T. ornata* found in the same hole or crevice (**Figure 7.3**).

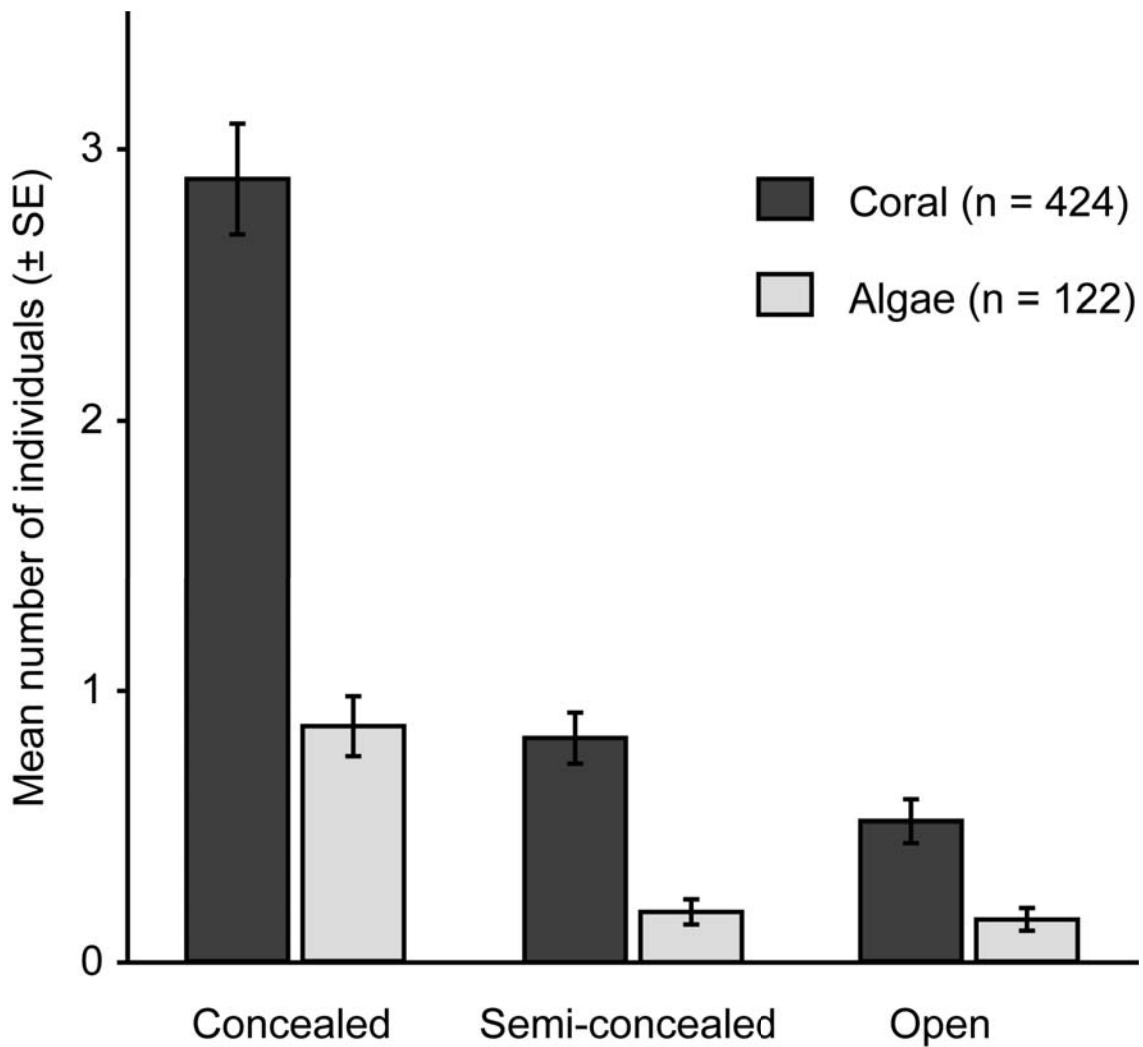


Figure 7.2: Distribution patterns of juvenile corals and early life-stage *Turbinaria ornata* thalli among microhabitats. The average number of individuals per quadrat is significantly higher in concealed microhabitats for both organisms with more than four times more individuals present in concealed microhabitats than in semi-concealed or exposed habitats.

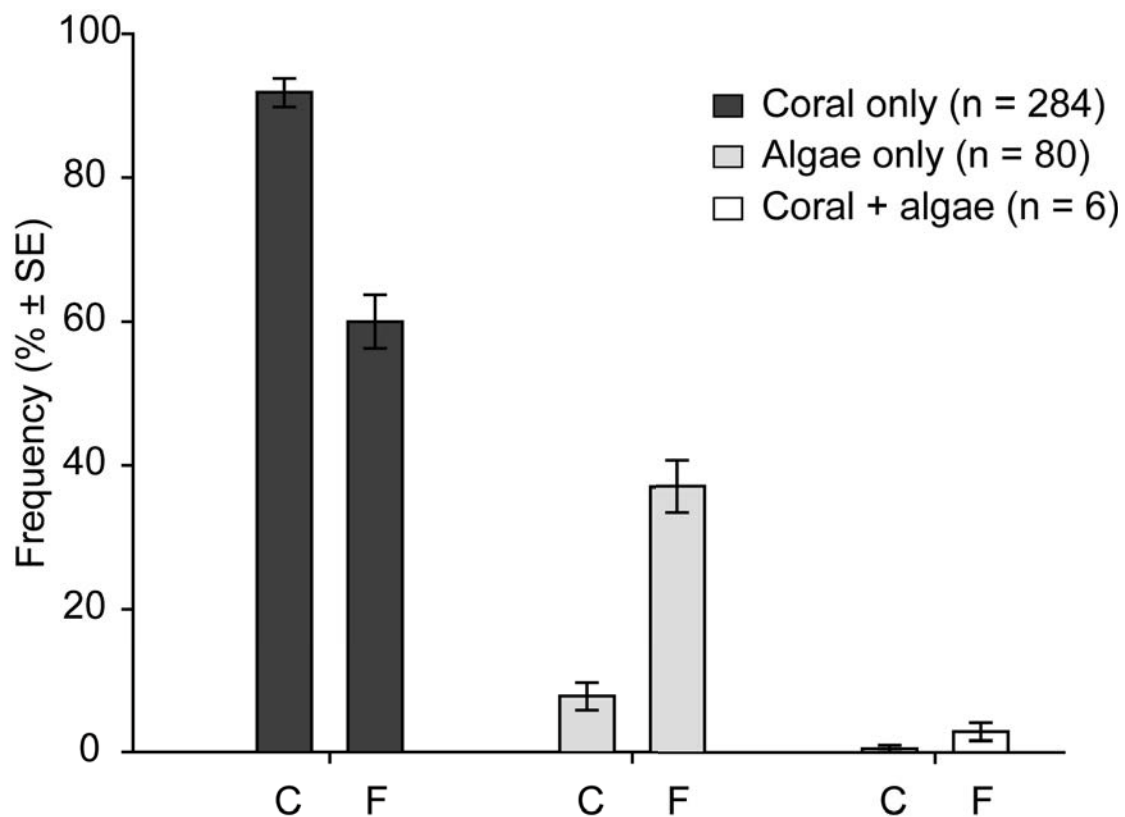


Figure 7.3: Proportion of juvenile corals, algae and both organisms in crevices in different reef zones. On the crest, 92% of records were corals only, while only 60% of records on the flat were corals only. Records of both organisms within the same crevice are rare in both reef zones ($n = 6$), amounting to only 1.6% of all records. C = Crest, F = Flat.

Table 7.1: Summary of generalized linear mixed model (GLMM) using a Poisson error distribution. Models are compared using the Akaike information criterion (AICc). Only models that perform better than the null model and the null model itself are considered. Shown are degrees of freedom (df), model maximum log-likelihood (logLik), Akaike's information criterion (AICc), changes in the AIC (Δ) and AIC weight (wAIC).

Coral models	df	logLik	AICc	Δ	wAIC
Exposure + Zone	4	-177.73	363.6	0.00	0.507
Exposure + Site + Zone	5	-177.33	364.9	1.27	0.269
Exposure + Site + Zone + Site \times Zone	6	-176.48	365.2	1.65	0.223
Exposure	3	-185.36	376.8	13.21	0.001
Exposure + Site	4	-185.15	378.4	14.84	0.000
Zone	3	-283.85	573.8	210.18	0.000
Site + Zone	4	-283.44	575.0	211.43	0.000
Site + Zone + Site \times Zone	5	-282.59	575.4	211.80	0.000
Null	2	-291.48	587.0	223.4	0.000

Algae models	df	logLik	AICc	Δ	wAIC
Exposure + Zone	4	-108.76	225.7	0.00	0.480
Exposure + Site + Zone	5	-108.06	226.3	0.68	0.341
Exposure + Site + Zone + Site \times Zone	6	-107.67	227.6	1.97	0.179
Exposure	3	-120.98	248.0	22.40	0.000
Exposure + Site	4	-120.90	249.9	24.29	0.000
Zone	3	-142.07	290.2	64.56	0.000
Site + Zone	4	-141.37	290.0	65.23	0.000
Site + Zone + Site \times Zone	5	-140.98	292.2	66.51	0.000
Null	2	-154.21	314.5	88.85	0.000

7.3.2 Distribution patterns of juvenile algal thalli

As in corals, early life-stage *T. ornata* were more abundant in concealed microhabitats (0.9 ± 0.1 thalli m^{-2} SE) than in semi-concealed (0.2 ± 0.1 m^{-2} SE) or open (0.1 ± 0.0 m^{-2} SE) microhabitats (**Figures 7.2**). However, in contrast to coral juveniles, early life-stage *T. ornata* were more abundant on the outer flat (2.1 ± 0.2 thalli m^{-2} SE) than on the crest (0.4 ± 0.1 m^{-2} SE) (**Figure 7.3**). This was reflected in the GLMM, which identified a significant negative association of *T. ornata* thalli abundance with the reef crest ($P < 0.001$) and semi-concealed and open microhabitats ($P < 0.001$). As in the model describing the distribution of juvenile coral colonies, this model was preferred over models including site as a fixed factor ($\chi^2 = 1.4$, $P = 0.2$) and/or an interaction term between site and zone ($\chi^2 = 2.2$, $P = 0.3$, **Table 7.1**), indicating that site had again no significant effect on the observed distribution patterns. The distribution of bite marks on early life-stage *T. ornata* varied significantly between reef zones, with a higher proportion of bite marks present on thalli from the reef crest (42.1% with bite marks) than on the outer flat (21.2% with bite marks) ($\chi^2 = 4.5$, $P < 0.05$).

7.3.3 Distribution patterns of herbivorous reef fishes

A total of 1,234 herbivorous fishes, in 30 species within three families (Acanthuridae, Labridae, Siganidae) were recorded. The average biomass of herbivorous fishes displayed no consistent relationships between habitats or sites (Site \times Zone: $F_{1,20} = 9.2$, $P = 0.008$). On the reef crest the average biomass of herbivorous fish was lower at BS than LH (13.9 ± 1.9 and 21.8 ± 3.8 $\text{kg} \cdot 250\text{m}^{-2}$ SE, respectively), while on the outer reef flat herbivorous fish biomass was lower at LH than BS (10.5 ± 3.0 and 18.8 ± 1.3 $\text{kg} \cdot 250\text{m}^{-2}$ SE, respectively). Sites and reef zones alone had no significant effect on fish biomass (Site: $F_{1,20} = 0.003$, $P = 0.957$; Zone: $F_{1,20} = 1.4$, $P = 0.249$).

Non-metric multidimensional scaling of the distance matrices computed for coral juveniles / early life-stage *T. ornata* and fish data both yielded a clear separation of the reef crest from the outer flat (**Figure 7.4**). Corals, in all three microhabitats, characterized the reef crest, while the outer flat was distinguished by *T. ornata*. Among herbivorous fish species, the reef crest was characterized by the surgeonfishes *Zebrasoma scopas* and *Ctenochaetus striatus*, pairing rabbitfishes (including *Siganus doliatus*, *S. vulpinus*, *S. corallinus*, *S. punctatus*) and some scraping parrotfishes (*Scarus niger*, *S. dimidiatus*, *S. oviceps*). The outer reef flat was characterized by schooling rabbitfishes (*Siganus argenteus* and *S. canaliculatus*), several scraping parrotfishes (*Scarus psittacus*, *S. rivulatus* and *S. schlegeli*) and surgeonfishes (*Acanthurus nigrofuscus*, *A. lineatus*, *A. olivaceus*). The BEST ENV-BIO routine identified pairing rabbitfishes (*Siganus puellus*, *S. punctatus*) and the surgeonfish *Zebrasoma scopas* to be the species that best predicted the observed patterns in the abundance of coral juveniles and early life-stage *T. ornata* (Spearman rank correlation obtained under 999 permutations: $r_s = 0.72$, $P < 0.001$). All three species were completely absent from reef flat habitats and, although not numerically abundant (mean 2.0 ± 0.5 individuals 250m^{-2}), were consistently present on the reef crest, suggesting a positive relationship between the abundance of the three species and the observed distribution of corals and a negative relationship with the distribution of *T. ornata* when considering the resultant MDS plots (**Figure 7.5**).

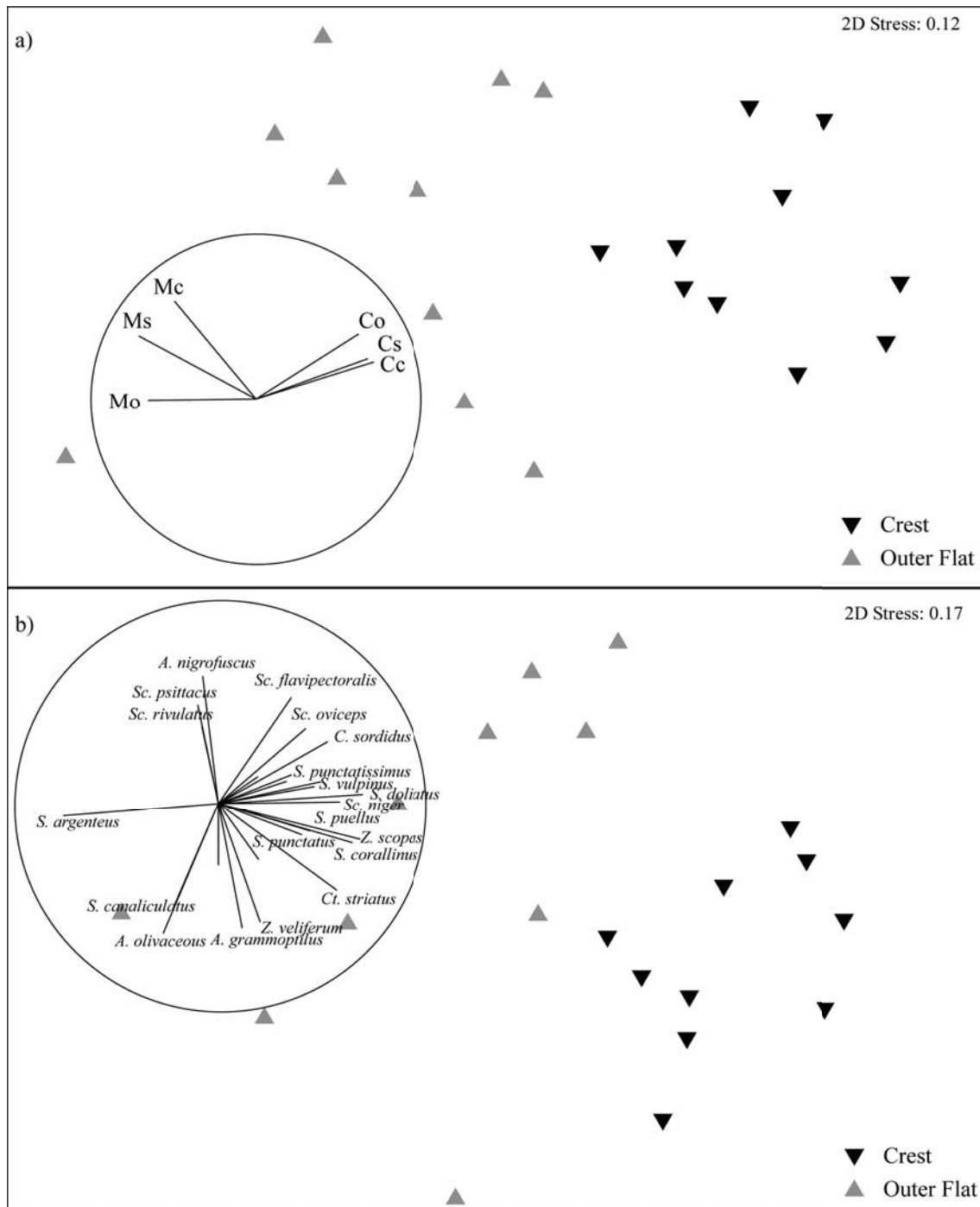


Figure 7.4: Non-metric multidimensional scaling analyses performed on distance matrices for (a) benthic patterns and (b) the distribution of herbivorous fishes. Both ordination plots show a clear segregation in transects on the crest and outer flat. (a)

The crest is characterized by corals in all microhabitats (Cc = coral concealed, Cs = coral semi-concealed, Co = coral open) while the outer flat supports macroalgal thalli in all microhabitats (Mc = macroalgae concealed, Ms = macroalgae semi-concealed, Mo = macroalgae open). (b) Fish species characteristic for the different reef zones.

The reef crest is characterized predominantly by pairing rabbitfishes, *Zebrasoma* spp. and several scraping parrotfishes while the outer flat is dominated by schooling rabbitfishes, detritivorous surgeonfishes and scraping parrotfishes.

7.4 Discussion

Cryptic microhabitats, such as holes and crevices, are often assumed to be important habitats for early life stages of benthic coral reef organisms, primarily as a spatial refuge from predation (Penin et al. 2011; Trapon et al. 2013a, b). There are, however, surprisingly few studies examining the role of these concealed microhabitats for corals and macroalgae, especially on natural reef substrata. My study provides evidence that highlights the potential importance of concealed microhabitats to juvenile corals and macroalgae on the exposed reef crest and outer reef flat zones. Despite juvenile corals and early life-stage *T. ornata* displaying different among-habitat distributions, both were 4-5 times more abundant within concealed microhabitats (i.e., holes and crevices) than in semi-concealed (i.e., cracks and ledges) or open microhabitats. This association with concealed microhabitats is even more striking given that these habitats account for less than 10 % of the available area at the study sites. Furthermore, the observed distribution patterns of juvenile corals and algae were best explained by three fish species, the rabbitfishes *S. puellus* and *S. punctatus*, and the surgeonfish *Z. scopas*. Interestingly, these three species have recently been reported to feed extensively in holes and crevices (Fox & Bellwood 2013), the microhabitat where most juvenile corals and algae were found. This provides correlative evidence, suggesting that microhabitat characteristics may play an important role in the early benthic life-stages of corals and macroalgae, which may add another facet to the complex dynamics between corals, algae and grazing fishes on coral reefs.

7.4.1 Distribution patterns of juvenile corals and macroalgal thalli

The distribution patterns observed in the present study suggest that juvenile corals survive predominantly in concealed microhabitats. This is consistent with previous experimental studies that have found a strong preference for coral planulae to settle in narrow cracks and

crevices (Harriott & Fisk 1987; Babcock & Mundy 1996; Raimondi & Morse 2000; Vermeij 2006). However, the results of the present study are in marked contrast to previous field observations (Edmunds et al. 2004; Roth & Knowlton 2009; Traçon et al. 2013a), in which the majority of coral juveniles were found in open, exposed microhabitats. The reasons for these differences are not readily apparent but may be related to variation in the reef zone examined, scale and definition of open vs. concealed microhabitats, availability and size of microhabitats, or a range of biotic (e.g., grazing rates, benthic assemblages) and abiotic (e.g., sedimentation, wave action) factors. Of these, the size of the available crevices may have a large bearing on the microhabitat associations of juvenile corals. Corals may preferentially settle to micro-crevices (<10mm; Nozawa 2008), and then outgrow the microhabitat to appear as a juvenile colony on open, or exposed, substratum. Furthermore, two of these studies were located in biogeographically distinct locations with relatively low-diversity coral communities (Palmyra Atoll: Roth & Knowlton 2009; Florida Keys: Edmunds et al. 2004) and may not be directly comparable to the present study. The third study, however, was conducted on exposed reef crests of nine reefs on the GBR, including those around Lizard Island (Traçon et al. 2013a). Traçon et al. (2013a) reported that relatively few juvenile corals (10-50mm diameter) were found within crevices, but there was significant variation among coral genera and latitudes (i.e. between the southern, central, and northern GBR). The microhabitat associations of juvenile corals may thus depend strongly on local abiotic and biotic factors.

A common explanation for the preference for cryptic habitats by coral planulae is that these habitats reduce post-settlement mortality through incidental grazing by reef herbivores, especially scraping and excavating parrotfishes (Brock 1979; Sammarco 1980; Penin et al. 2011; Evans et al. 2013). Indeed, a recent study at one of the sites used in the present study (BS) demonstrated that the exclusion of herbivorous fishes increased early post-settlement

survival of coral recruits on settlement tiles (Traçon et al. 2013b). Furthermore, the survivorship of coral recruits on tiles exposed to herbivores was negatively correlated with the density of parrotfish feeding scars on tiles.

Although there was no relationship between the distributions of juvenile corals or macroalgae and scraping and excavating parrotfishes in the present study, these fishes were present at both sites and in both reef zones, suggesting that grazing pressure on open surfaces (cf. Fox & Bellwood 2013) was high. Grazing-induced mortality as a result of feeding by parrotfishes may have contributed to the abundance of juvenile corals and algae in concealed microhabitats. However, there are many other factors that may contribute to these patterns, including increased survival of juveniles due to reduced abrasion (Hunte & Wittenberg 1992) or decreased potential of dislodgement (Madin & Connolly 2006). Alternatively, crevices may present preferred settlement locations due to, for instance, favourable light regimes (Maida et al. 1994; Babcock & Mundy 1996; Mundy & Babcock 1998; Baird et al. 2003) or the presence of specific coralline algae (Harrington et al. 2004). Given the number of processes that influence settlement choice and post-settlement survival of corals, the grazing pressure administered by herbivorous fishes is likely to be only one of many factors influencing microhabitat distribution patterns of juvenile corals.

Juvenile *T. ornata* were also most abundant in concealed microhabitats, emphasizing the possible importance of concealed microhabitats for the ecology of macroalgae (Diaz-Pulido & McCook 2004). Spatial refuges are assumed to be a beneficial factor for the settlement and survival of macroalgae (Fletcher & Callow 1992) but the effect of crevices on mortality of early life-stage macroalgae have rarely been assessed in the field (but see Diaz-Pulido & McCook 2004). As for corals, multiple factors (e.g., larval supply, water motion, sedimentation) are likely to influence the settlement preferences (Fletcher and Callow 1992), and post-settlement growth and survival (e.g., nutrients, herbivory) of algae. Among these,

herbivorous grazing is known to have a major influence on the survival and distribution of algae on coral reefs (e.g., Wright & Steinberg 2001; Hughes et al. 2007; Cheal et al. 2010).

Our two study habitats, the reef crest and outer reef flat, are commonly identified as areas of high herbivore activity (Hay 1981; Fox & Bellwood 2007). Given the detrimental effects of grazing on early life-stages of macroalgae, including *T. ornata* (Vadas et al. 1992; Lotze et al. 2001; Diaz-Pulido & McCook 2003; Stiger & Payri 2005), the high rates of herbivory within these habitats are likely to have reduced the survivorship of any *T. ornata* propagules that settled on exposed microhabitats. Moreover, variation in the consumption of adult or large fleshy fucal macroalgae (*Sargassum swartzii* and *S. cristafolium*) among seven habitats around Lizard Island, revealed that herbivory was greatest on the exposed reef crest and outer flat, including one of the sites in the present study (Hoey & Bellwood 2009, 2010b). Such high rates of grazing and browsing lends support to the hypothesis that herbivory may be contributing to the observed association of *T. ornata* with concealed microhabitats. However, the differential distribution of adult algae among zones suggests that other factors, such as wave action, sedimentation or larval supply, are also influencing reef-scale distribution patterns of *T. ornata*.

7.4.2 Coral-algae interactions

Although corals and algae are most abundant in concealed microhabitats, they rarely occupied the same hole or crevice. The majority of crevices contained either juvenile corals only, or small *T. ornata* only. In the few cases ($n = 6$), where a juvenile coral and a small *T. ornata* were found in the same crevice, the corals showed signs of bleaching and/or tissue loss that may reflect the outcomes of competitive interactions (tissue loss, bleaching; **Figure 7.5**). Mechanisms and consequences of algae-coral interactions are still not well resolved, especially between juvenile corals and early life-stage macroalgae (McCook et al. 2001;

Vermeij et al. 2009). However, it is commonly assumed that erect macroalgae can have detrimental effects on corals, especially during their early life stages (Sammarco 1980; Birrell et al. 2008; Diaz-Pulido et al. 2010). *T. ornata* has been suggested to have negative impacts on coral health, growth and recruitment, most likely through mechanical effects such as shading and abrasion (Gleason 1996; Rasher et al. 2012; Shearer et al. 2012). It is possible, therefore, that early life-stages of *T. ornata* may compete with juvenile corals for space within crevices. However, the factors influencing the potential competition between these organisms as well as the results of their competitive interactions are unclear. Thus, future experimental studies are needed to assess if juvenile corals interact with early life-stage macroalgae such as *T. ornata* and, if so, whether concealed microhabitats are a significant venue for these interactions.

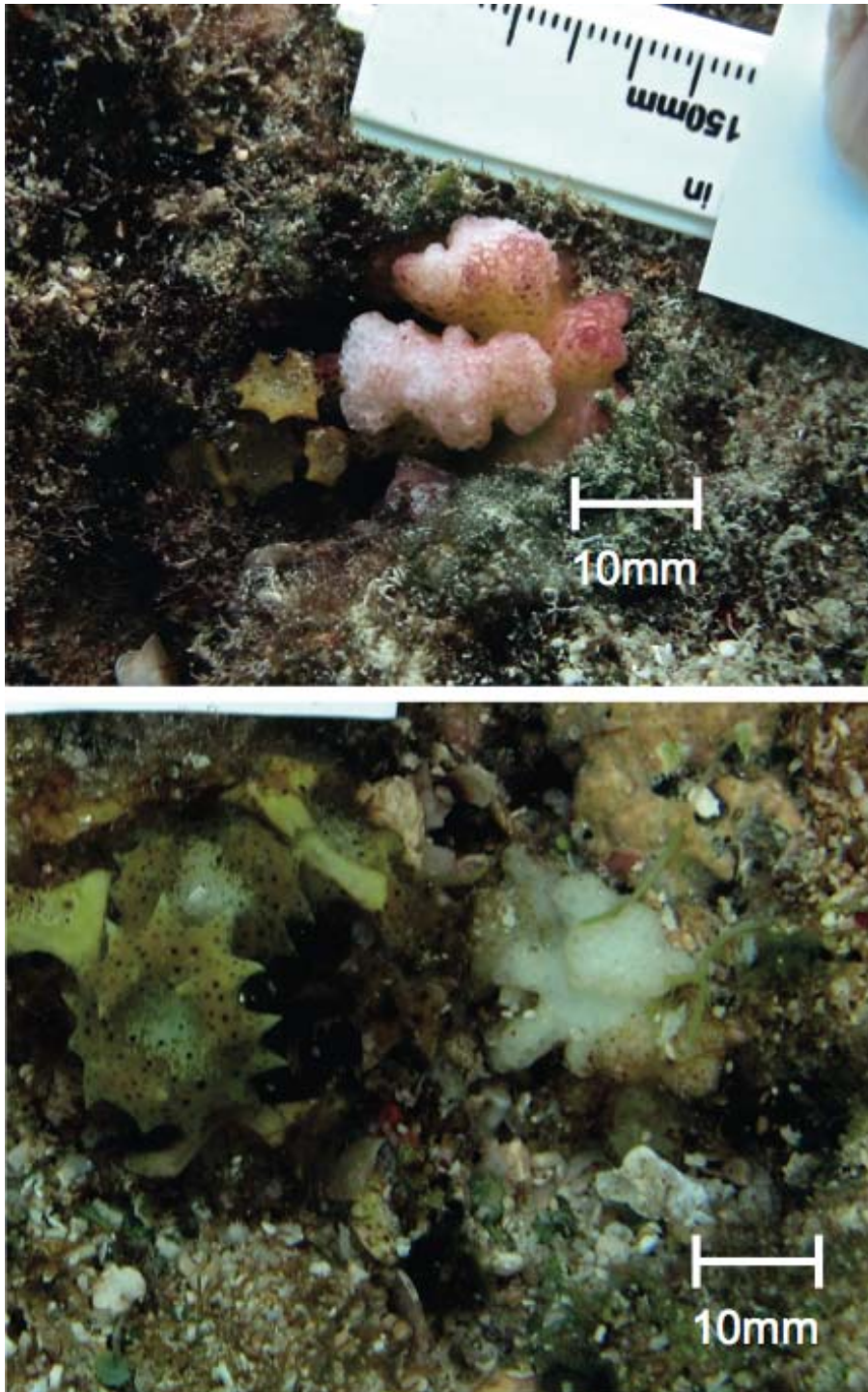


Figure 7.5: Juvenile corals and early life-stage *T. ornata* thalli in the same crevice.

a) The *T. ornata* appears to be in direct contact with the juvenile pocilloporid coral, which shows signs of tissue loss and abrasion. b) With *T. ornata* in close proximity,

the juvenile coral is fully bleached. In both cases, the coral exhibits clear signs of competitive interactions with the algae. Signs of bleaching were observed in 66.7% of cases, where corals and algae were found in the same crevice ($n = 6$), whereas only 6.7% of all other colonies ($n = 418$) showed signs of bleaching. Photographs are taken at similar magnifications (scale = 10mm).

7.4.3 The distribution of herbivorous reef fishes

Reef crests have been repeatedly reported to support higher species richness and biomass of herbivorous fishes than adjacent reef zones (e.g., Russ 1984; Fox & Bellwood 2007; Hoey & Bellwood 2008). In the present study total herbivore biomass did not differ consistently between the two reef zones, but there was marked variation in the composition of herbivorous fish assemblages between the two zones. Grazing parrotfishes and surgeonfishes were common throughout the study areas, and are known to feed heavily the epilithic algal matrix on exposed surfaces (Bellwood 1995; Fox & Bellwood 2013). Grazing by parrotfishes and surgeonfishes may therefore have contributed to the low abundances of juvenile corals and *T. ornata* on exposed surfaces, however, these fish groups avoid concave surfaces and crevices when feeding (Bellwood and Choat 1990; Bellwood 1995; Fox & Bellwood 2013), which suggests that their influence on the benthos within concealed microhabitats will be limited.

In contrast to the widespread parrotfishes, pairing rabbitfishes and surgeonfishes in the genus *Zebrasoma* were largely restricted to the reef crest in the present study. Interestingly, these species are among a limited suite of nominally herbivorous fishes that have the capacity to exploit a wide range of microhabitats, including holes and crevices, when feeding (Robertson & Gaines 1986; Brandl & Bellwood 2013a,b; Fox & Bellwood 2013). It thus appears that the reef crest harbours a significantly larger biomass of herbivorous species with the capacity to feed on organisms within coral reef crevices, and that the distribution of these species was related to the distribution of juvenile corals (positively) and early life-stage *T. ornata* (negatively) across the study sites. Although only correlative, these relationships point to a potential link between these crevice-feeding fishes and the benthic community within crevices. However, further experimental studies are required to explore the role of fishes in the dynamics of benthic organisms in concealed microhabitats.

The results of the present study provide a useful insight into the interactions between juvenile corals, macroalgae, and grazing herbivores. My data suggests that concealed microhabitats, such as holes and crevices, may play an important role in the early life-stages of corals and algae. Although only providing a preliminary snapshot, my study highlights the potential importance of micro-topography in the dynamics of reef systems, and will hopefully stimulate further research on the role of concealed microhabitats in structuring reef assemblages.

Chapter 8: Micro-topographic refuges drive consumer-producer dynamics by mediating consumer functional diversity

Currently in review as: Micro-topographic refuges drive consumer-producer dynamics by mediating consumer functional diversity (in review).

8.1 Introduction

Herbivory is universally accepted as a critical ecosystem process in vegetated environments (Gruner et al. 2008). Through the disturbance induced by feeding, herbivorous organisms affect producer communities in terms of their biomass, diversity, and productivity (Olf and Ritchie 1998; Worm and Duffy 2003). However, herbivory is multifaceted, and numerous different aspects can influence the dynamics between consumer and producer communities. One crucial factor relates to the functional identity of herbivores. In almost every ecosystem, herbivore communities comprise multiple species that vary in their functional niches, which modulates their effects on producer communities (Sommer 1999). Similarly, producer avoidance of, or tolerance to, herbivory affects dynamics between consumer and producer communities (Duffy and Hay 1990; Rasher et al. 2013). While intrinsic mechanisms to avoid being grazed, such as morphological or chemical defense mechanisms, have received considerable attention (Berenbaum 1995), extrinsic factors that permit producers to escape herbivore pressure are less well understood (Milchunas and Noy-Meir 2002).

Extrinsic factors underlying grazer-avoidance include both biological and physical protection from grazing, i.e. refuges (Duffy and Hay 1990). Biological refuges comprise positive interactions among plant-species, in which species susceptible to grazing gain associational refuge by growing close to an unpalatable or non-preferred species (Pfister and Hay 1988; Stachowicz 2001). Physical refuges usually relate to physical or geological features, which reduce the accessibility of producers to herbivorous grazers (Milchunas and

Noy-Meir 2002). Compared to biological refuges, such physical refuges have received relatively little consideration, despite their demonstrated importance in both terrestrial (Shitzer et al. 2008) and aquatic (Hay 1981; Bergey 2005) environments. Importantly, physical refuges can vary dramatically in scale, ranging from entire islands (Milchunas and Noy-Meir 2002) to micro-topographic structures on the scale of a few millimetres (Menge and Lubchenco 1981; Dudley and D'Antonio 1991; Bergey 2005). Although often inconspicuous, the latter can significantly influence consumer-producer dynamics, especially when grazing pressure is intense (Menge and Lubchenco 1981; Milchunas and Noy-Meir 2002; Bergey 2005).

Many primary producers in marine environments are subject to particularly rigorous consumer control (Gruner et al. 2008; Bennett et al. 2015) and among marine habitats, benthic communities experience the strongest level of herbivore control (Poore et al. 2012). On tropical coral-dominated reefs, grazing pressure is particularly strong and reductions of grazing pressure often lead to the rapid establishment of fleshy macroalgae (Burkepile and Hay 2008; Cheal et al. 2010; Rasher et al. 2013). As a consequence, tight links between strong consumer control by herbivorous fishes and the resilience of coral reefs have been established (Nyström et al. 2008; Graham et al. 2015).

This emphasizes the potential importance of grazing refuges in these ecosystems (Littler et al. 1986). Several cases of biological refuges, in which readily consumed algae associate with species unpalatable to most herbivores, have been reported from coral reefs (Littler et al. 1986; Pfister and Hay 1988). Likewise, physical refuges exist at several different scales, ranging from regional refuges (inner shelf vs. outer shelf on the Great Barrier Reef (GBR) (Wismer et al. 2009), to reef zones (inner flat or sandy plains vs. crest, (Hay 1981; Fox and Bellwood 2007). However, micro-topographic refuges from grazing pressure (i.e. the three-dimensional structure of the reef on the scale of a few centimetres) also

represent a widespread type of physical refuge within coral reef systems, occupying up to 25% of the available microhabitats on coral reefs (Brandl et al. 2015). While crevices appear to be critical for the recruitment and survival of scleractinian corals (Brock 1979; Nozawa 2008; Arnold et al. 2010; Brandl et al. 2014; Edmunds et al. 2014), and represent an important ecological axis for niche partitioning in herbivorous fishes (Robertson and Gaines 1986; Fox and Bellwood 2013; Brandl and Bellwood 2014; Brandl et al. 2015), no examination of the overarching effect of micro-topographic refuges on consumer-producer dynamics exists to date.

The purpose of the present study, therefore, was to provide an experimental evaluation of the effects of micro-topographic refuges on grazing dynamics on coral reefs. In quantifying grazing pressure, average turf lengths, the functional identity of grazer communities, and the community composition of the benthos in different microhabitats, we sought to provide answers to the following questions: i) Do micro-topographic refuges reduce the grazing pressure exerted by herbivorous fishes on the benthic community? ii) Is this exclusion evenly spread among fish species and functional groups? and iii) How do changes in grazing pressure and grazer identity affect the benthic community?

By answering these questions, we demonstrate that micro-topographic complexity leads to two distinct grazing regimes on coral reefs, which coexist at the scale of a few centimetres and may represent a key feature of consumer-producer dynamics in reef ecosystems.

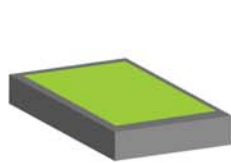
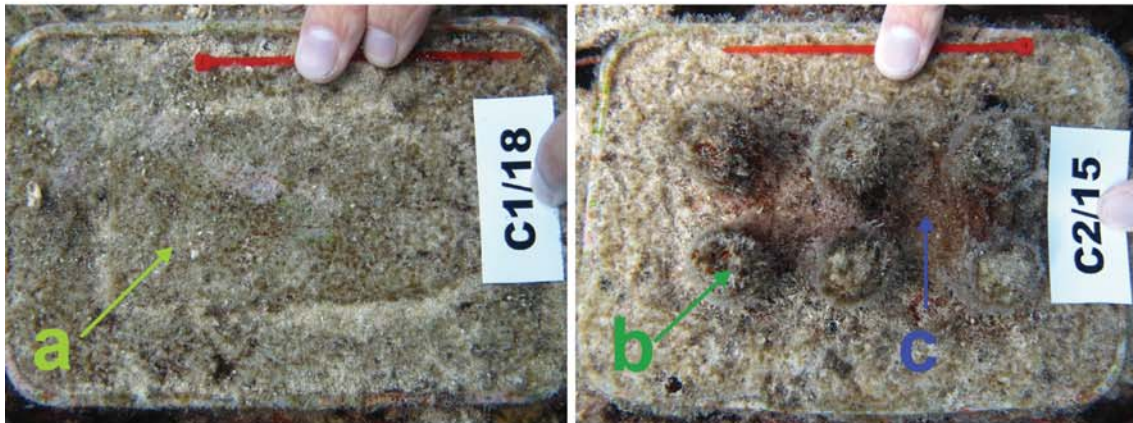
8.2 Materials and methods

8.2.1 Data collection

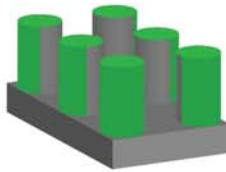
To assess the effects of micro-topographic refuges on fish grazing patterns and the benthic community, I constructed two distinct grazing surfaces. The first, flat surface, consisted of a

flat rectangle (length = 115mm, width = 65mm, height = 15mm). The second, complex surface, featured the same dimensions but included six evenly spaced vertical cylinders (radius = 10mm, h = 35mm; spaces between cylinders = 25mm) in two rows of three, to simulate micro-topographic refuges as found on coral reefs (specifically related to dead coral skeletons). On these surfaces, three distinct microhabitats were identified (**Figure 8.1**): flat and exposed (flat tiles without cylinders, henceforth 'flat'), complex and exposed (the outside and top of cylinders, henceforth 'exposed'), and complex and concealed (the inside of cylinders and the flat area between, henceforth 'concealed'). To ensure uniformity among surfaces, I used standardized moulds made from neutral cure sealant, dry cornstarch, and mineral spirits (2:2:1 by volume). Flat surfaces had only the recess (10mm) for the flat rectangular base. For complex surface moulds, holes (depth = 35mm, radius = 10mm) were cut out of the mould using a drill press with a hole-saw fitting. To cast surfaces, a mixture of powdered coral rubble (dried for 72 hours, and pulverized using a sledge hammer), river sand, and cement (1:1:1) was mixed with water and poured into the moulds. This mixture was selected following a pilot study that revealed no differences in the benthic community between natural tiles of dead coral and the rubble-cement mixture (cf. Hixon & Brostoff 1985). After setting the concrete mixture for 48 hours, surfaces were extracted from the moulds and each surface was set centrally in a 750ml disposable plastic container filled with a standard concrete mixture. After setting for 48 hours, the containers were tied together for stability, producing grazing arrays of 20 surfaces (10 of each type). Six arrays were created yielding a total of 120 surfaces with 180 grazing microhabitats (flat $n = 60$, exposed $n = 60$, concealed $n = 60$). Arrays were deployed in groups of three at two different sites along the reef crest between Bird Islets and South Island (near Lizard Island, northern GBR) in the beginning of June 2013.

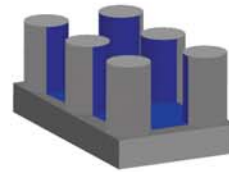
Microhabitats



(a) flat

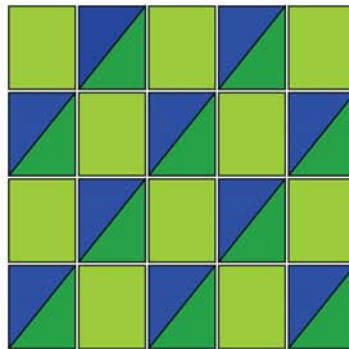


(b) exposed

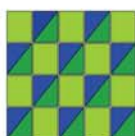
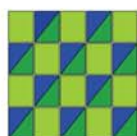
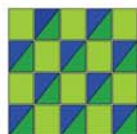


(c) concealed

Array



Site 1



Site 2

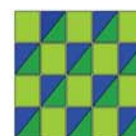
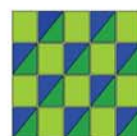
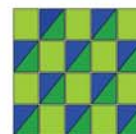


Figure 8.1: Description of the three focal microhabitats and deployment of the experiment. Flat microhabitats (a) are flat and exposed, exposed microhabitats (b) are complex and exposed on the outside or top of cylinders, and concealed microhabitats (c) are on the inside of cylinders and in the flat area between cylinders. Arrays consist of ten flat surfaces and 10 surfaces with cylinders, creating 30 non-independent microhabitats per array. Three arrays were deployed at each site.

The arrays were left on the reef for 35 weeks to establish natural benthic communities. Subsequently, the lengths of epilithic turf filaments from each microhabitat were measured using vernier callipers. To do so, the turf filament closest to a haphazardly chosen point was straightened and its vertical extension measured to the nearest millimetre. Ten measurements were taken from each microhabitat. No measurements were taken within 5mm of the edge of the rectangular base to avoid edge effects. Photographs were taken of each microhabitat from a fixed distance of approximately 10 cm to quantify the benthic community composition. Finally, the foraging activity of fish assemblages grazing on the microhabitats was monitored using remote underwater videos (GoPro Hero III) in January/February 2014. Each array was filmed for three to four hours on at least five non-consecutive days, resulting in a total observation period of 111 hours and an average of 18.52 ± 0.07 hours per microhabitat. Subsequently, both benthic photographs and grazing videos were analysed in the lab. On each photograph, I quantified the proportional cover of different benthic organisms from a planar view of the respective microhabitats (all of the flat microhabitat, upward facing circular surfaces for exposed microhabitats, horizontal surface between bases of cylinders for concealed microhabitats). Benthic organisms were assigned to a functional category (Steneck & Dethier 1994), and their proportional cover estimated using the software ImageJ. Videos were analysed by counting the number of bites taken on each microhabitat by all large (>10 cm) mobile herbivorous fish species (*sensu* Choat et al. 2002), since identification of smaller fishes was impossible in the videos.

Published data were used to characterize fish species in terms of their morphology, diet, and gut short-chain fatty acid (SCFA) profile (**Table 8.1**). For the morphology, five traits previously linked to foraging microhabitat utilization (body depth, eye diameter, snout length, snout angle, and head angle), were considered (Brandl & Bellwood 2013a). Trait measurements were regressed against standard length in a dataset comprising 260 individuals

in 99 species and residual values were averaged for each species. For the dietary data, due to differences among sources in the functional categorizations of dietary items, a broad classification scheme was created to ensure congruent dietary classifications across taxa. Categories included sediment, organic material, filamentous algae, fleshy/thallose algae, and other items such as benthic invertebrates or foraminifera. For the SCFA profiles, the overall amount of dominant SCFAs (Acetate, Butyrate, Isovalerate, and Propionate) present in the gut segment with the highest SCFA concentrations was used, as well as the proportional composition of SCFAs.

Table 8.1: Published functional traits describing the morphology, diet, and gut SCFA profiles of the herbivorous fish species present in the study. Morphological trait values are averaged individual residuals extracted from a linear model regressing the respective trait against the measured individuals' standard length (Brandl & Bellwood 2013). Diet values are averaged proportional values of each group of items in the examined individuals' guts (Choat 1969; Choat et al. 2002; Hoey et al. 2013). SCFA values represent the averaged total SCFA concentration in the guts of examined individuals and its proportional composition (Clements & Choat 1995).

Morphology					
Species	M1	M2	M3	M4	M5
<i>Acanthurus nigricauda</i>	0.018	0.568	0.031	0.021	-0.064
<i>Acanthurus nigrofuscus</i>	0.008	0.397	0.019	0.020	-0.005
<i>Chlorurus spilurus</i>	-0.069	0.379	-0.061	0.011	-0.070
<i>Ctenochaetus striatus</i>	0.043	0.555	-0.025	-0.002	0.014
<i>Scarus frenatus</i>	-0.057	0.339	-0.056	-0.001	-0.040
<i>Scarus globiceps</i>	-0.048	0.339	-0.060	0.008	-0.078
<i>Scarus niger</i>	-0.011	0.338	-0.040	0.006	-0.049
<i>Scarus psittacus</i>	-0.026	0.349	-0.065	0.002	-0.040
<i>Scarus schlegeli</i>	-0.038	0.317	-0.060	-0.013	-0.035
<i>Siganus corallinus</i>	0.009	0.414	0.008	-0.045	0.066
<i>Siganus punctatissimus</i>	0.035	0.315	0.117	0.027	0.084
<i>Siganus punctatus</i>	0.010	0.381	0.040	0.015	0.022
<i>Zebrasoma scopas</i>	0.142	0.600	-0.013	-0.088	0.080
<i>Zebrasoma velifer</i>	0.089	0.625	0.072	-0.055	0.085

Diet					
Species	D1	D2	D3	D4	D5
<i>Acanthurus nigricauda</i>	37.310	56.197	5.240	0.310	0.943
<i>Acanthurus nigrofuscus</i>	0.829	0.501	0.167	36.652	61.048
<i>Chlorurus spilurus</i>	25.000	16.400	0.000	58.600	0.200
<i>Ctenochaetus striatus</i>	72.581	18.455	1.964	2.957	3.278
<i>Scarus frenatus</i>	10.800	26.700	1.500	55.000	1.600
<i>Scarus globiceps</i>	8.200	27.200	2.300	60.600	1.700
<i>Scarus niger</i>	6.800	27.400	0.400	63.000	2.400
<i>Scarus psittacus</i>	9.700	43.700	1.800	37.700	4.100
<i>Scarus schlegeli</i>	9.700	43.700	1.800	37.700	4.100
<i>Siganus corallinus</i>	7.600	1.000	1.100	28.700	59.750

<i>Siganus punctatissimus</i>	20.500	17.500	21.350	12.200	28.250
<i>Siganus punctatus</i>	11.700	8.000	38.600	6.250	36.100
<i>Zebrasoma scopas</i>	0.000	0.045	2.365	31.368	59.065
<i>Zebrasoma velifer</i>	0.763	0.683	0.297	4.844	92.562

Gut SCFAs					
Species	S1	S2	S3	S4	S5
<i>Acanthurus nigricauda</i>	9.660	63.730	2.500	2.340	30.820
<i>Acanthurus nigrofuscus</i>	18.180	86.590	4.310	6.190	2.490
<i>Chlorurus spilurus</i>	11.480	87.000	3.070	1.230	8.020
<i>Ctenochaetus striatus</i>	9.070	63.710	3.880	7.030	25.070
<i>Scarus frenatus</i>	9.800	85.260	1.310	1.830	11.290
<i>Scarus globiceps</i>	12.070	75.430	4.340	2.640	17.610
<i>Scarus niger</i>	9.800	85.260	1.310	1.830	11.290
<i>Scarus psittacus</i>	6.170	70.830	1.860	0.910	23.900
<i>Scarus schlegeli</i>	6.170	70.830	1.860	0.910	23.900
<i>Siganus corallinus</i>	10.790	85.390	10.180	1.760	1.640
<i>Siganus punctatissimus</i>	21.860	82.510	10.600	2.240	2.770
<i>Siganus punctatus</i>	14.290	80.990	11.820	2.500	3.000
<i>Zebrasoma scopas</i>	31.940	68.100	25.080	5.240	0.780
<i>Zebrasoma velifer</i>	36.970	72.420	17.150	8.540	1.050

M1: Body depth	D1: Organic particulates	S1: Total gut SCFAs
M2: Snout length	D2: Calcareous sediment	S2: Acetate
M3: Eye diameter	D3: Others	S3: Propionate
M4: Snout angle	D4: Filamentous algae	S4: Butyrate
M5: Head angle	D5: Fleshy algae	S5: Isovalerate

8.2.2 Statistical analysis

To assess differences in the overall grazing pressure (number of bites.day⁻¹.cm⁻²) and the length of turf algal filaments (mm) on the three microhabitat types, Bayesian Mixed Models (BMMs) were performed specifying the microhabitat type (flat, exposed, concealed) as a fixed effect. To account for the spatial non-independence of microhabitats within an array, array was specified to have a random effect. In addition, site was included as a fixed effect as the estimation of random variance can be unreliable if only two levels are available. For the comparison of grazing pressure among microhabitat types (overall grazing pressure model), a Poisson error distribution with a log-link function was specified. Furthermore, bite counts for each microhabitat were modelled against an offset specifying the overall observation period (time in d) and the area available for grazing (area in cm²). To do so, the log of time and area were included as fixed effects, and informative priors were formulated for the regression coefficients of both factors, specifying a β -parameter of 1 with a variance of 0.000001. Uninformative priors were used on the other fixed effect parameters ($\mu = 0$, $V = 100,000,000$), and informative priors ($V = 0.000001$, $\nu = 100$) were used for the residual variance due to over-dispersion in the response variable. The model was run for 10,000,000 iterations, while all other models (throughout the study) were run for 3,000,000 iterations. Chain convergence was validated using trace chain plots. For the comparison of algal filament length (turf length model), a Gaussian error distribution with a link function was used, as data were approximately normally distributed. Uninformative priors were used on the fixed effects and inverse Wishart priors on the random and residual variance ($V = 1$, $\nu = 0.002$).

To investigate the consumer community feeding on different microhabitat types, as well as their benthic communities, two non-metric multidimensional scaling ordinations were performed on Manhattan distance matrices. Subsequently, permutational multivariate

analyses of variance (PERMANOVA) with 999 permutations were performed to investigate compositional differences among microhabitat types, again fitting array as a random effect. I also calculated patterns of consumer species richness interacting with the benthic community, and functional group richness for and benthic communities, respectively, and compared these patterns among microhabitat types using BMMs with a Poisson error distribution and uninformative priors.

Furthermore, I evaluated the functional composition of consumers feeding on the different microhabitat types. I combined the three trait datasets (morphology, diet, and SCFAs) and normalized data to a mean of zero and a standard deviation of 1. I then calculated pairwise distances among species using Gower's distance metric and divided the community into functional groups using a hierarchical clustering analysis with Ward's method. This resulted in two broad functional groups. I then extracted the number of bites taken on each microhabitat by the two functional groups and computed the bite-weighted trait means for the grazer community feeding on each microhabitat.

In order to compare the grazing pressure exerted on the different microhabitat types by each of the two functional groups, I used two zero-inflated Poisson BMMs (one for each group). I modelled the number of bites against two offset variables specifying the observation period and grazing area (as for the overall grazing pressure model above). Microhabitat type was formulated as a fixed effect and specified to interact with both the Poisson part of the response (counts of fish bites) and the binomial part denoting the probability of an observation being zero. Due to the presence of near complete separation in the data, I used weakly informative Cauchy distributed priors on the fixed effects (Hadfield 2010). Remaining priors were specified as in previous models, although the random (array) and residual variance were fixed to 1 in the binomial part of the model. Finally, to examine the average traits of fishes feeding on the three microhabitat types in terms of their morphology,

diet, and SCFA profiles, I compared the functional trait means for the consumer community of each microhabitat type using three multi-response BMMs to account for the non-independence of the five trait variables in each broader category. To provide comparable estimates, all response variables were normalized to a mean of zero and a standard deviation of 1. Again, the microhabitat type was specified as a fixed effect while array was specified to have a random effect. A Gaussian error distribution was specified for all response variables. Uninformative priors were specified for the fixed effects. Parameter expanded priors were used on the random variance (Hadfield 2010). Residual variance-covariance matrices were assessed to extrapolate the covariance of the five trait variables in each model.

More detail on model specifications is provided in the appendix. All analyses were performed in the software R, using the packages *vegan* (Oksanen et al. 2013), *FD* (Laliberté & Legendre 2010), and *MCMCglmm* (Hadfield 2010).

8.3 Results

8.3.1 Overall grazing pressure and turf filament length

I found marked differences in grazing pressure among the three investigated microhabitat types. The overall grazing pressure was strongest on flat microhabitats (2.86 ± 0.15 [mean \pm SE] bites.day⁻¹.cm⁻²), followed by exposed microhabitats (0.81 ± 0.05 bites.day⁻¹.cm⁻²), while the lowest grazing pressure occurred on concealed microhabitats (0.17 ± 0.02 bites.day⁻¹.cm⁻²). Concealed microhabitats had an order of magnitude fewer bites compared to flat microhabitats (**Figure 8.2a**). There was no difference among sites, and random variance was small (**Table 8.2**).

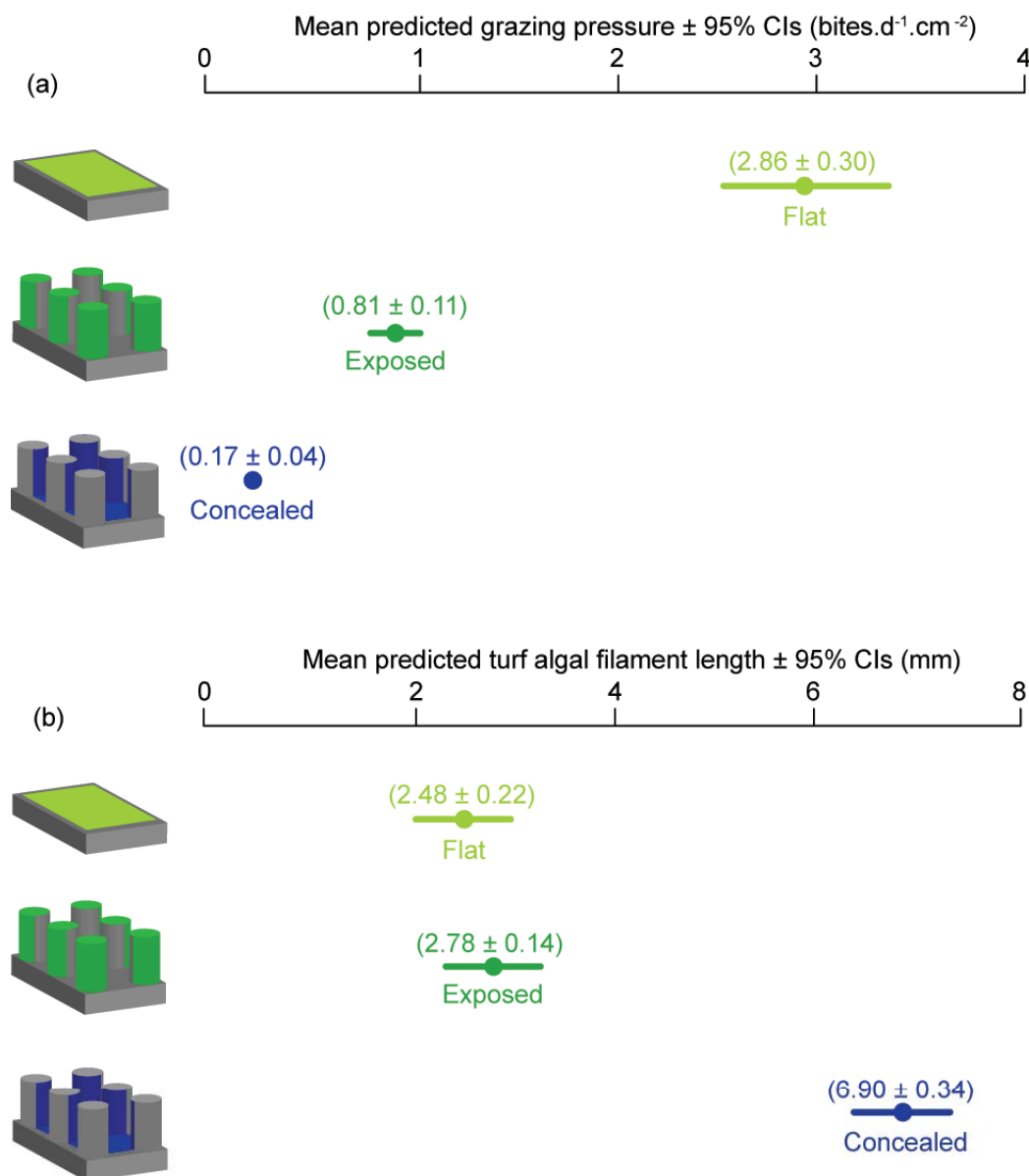


Figure 8.2: Predicted mean values from the grazing pressure model (a) and the turf length model (b), with observed means in parentheses. Flat microhabitats are subject to the highest grazing pressure, while concealed microhabitats experience the lowest grazing pressure. In contrast, turfs are longest in concealed microhabitats, while exposed and flat microhabitats both have shorter turfs. Plots show the

predicted mean \pm 95% credible intervals (CIs). Statistical significance can be assumed where CIs do not overlap with the mean of another treatment.

Table 8.2: Model output from the Poisson Bayesian Mixed Model comparing grazing pressure among microhabitat types and sites. Microhabitat type and site were specified as fixed effects with uninformative priors ($\mu = 0$, $V = 100,000,000$), while the two offset parameters, time and area, were fixed effects with informative priors ($\mu = 1$, $V = 0.000001$). Due to overdispersion, priors on the residual variance were specified as $V = 0.000001$ and $\nu = 100$, while uninformative priors were used for the random variance ($V = 1$, $\nu = 0.002$). To achieve well-mixed chains, the model was run for 10,000,000 iterations. Estimates for effect levels are based on a global intercept containing estimates for microhabitat type “flat” and site 1. Where 95% CIs do not intersect 0, statistically significant differences can be inferred.

Grazing pressure model

Iterations: 100001:9990001
 Thinning interval: 10000
 Sample size: 990

G-structure (random variance):		~array		
	Post. mean parameter est.	lower 95% CI	upper 95% CI	Eff. sample size
array	0.3384	0.02386	0.9559	990
R-structure (residual variance):		~units		
	Post. mean parameter est.	lower 95% CI	upper 95% CI	Eff. sample size
units	0.1517	0.119	0.1834	865
Fixed effects:		Number of bites ~ microhabitat type + site + log(time.d) + log(cm.sq)		
	Post. mean parameter est.	lower 95% CI	upper 95% CI	Eff. sample size
type(flat)	0.91894	0.36830	1.59062	990
type(exposed)	-1.20772	-1.35678	-1.06948	990
type(concealed)	-2.91238	-3.05042	-2.73961	990
site1	0.91894	0.36830	1.59062	990
site2	0.08132	-0.83559	0.93921	990
log(time.d)	1.00006	0.99819	1.00226	1324.1
log(cm.sq)	1.00004	0.99801	1.00196	1119.5

In contrast to grazing pressure, there were strong, but opposite, differences in the average turf filament length among the three microhabitat types. Flat microhabitats had the shortest turf filaments (2.48 ± 0.11 mm), closely followed by exposed microhabitats (2.78 ± 0.07 mm), while concealed microhabitats had by far the longest turf filaments (6.90 ± 0.17 mm) (**Figure 8.2b**). As for grazing pressure, site had no effect and random variance from the different arrays was small (**Table 8.3**).

Table 8.3: Model output from the Gaussian Bayesian Mixed Model comparing turf filament length among microhabitat types and sites. Microhabitat type and site were specified as fixed effects with uninformative priors ($\mu = 0$, $V = 100,000,000$). Uninformative priors were used for the residual and random variance parameters ($V = 1$, $\nu = 0.002$). Estimates for effect levels are based on a global intercept containing estimates for microhabitat type “flat” and site 1. Where 95% CIs do not intersect 0, statistically significant differences can be inferred.

Turf length model

Iterations: 10001:299001
 Thinning interval: 1000
 Sample size: 2990

G-structure (random variance):		~array		
	Post. mean parameter est.	lower 95% CI	upper 95% CI	Eff. sample size
array	0.1208	0.0002547	0.4463	2990
R-structure (residual variance):		~units		
	Post. mean parameter est.	lower 95% CI	upper 95% CI	Eff. sample size
units	0.9308	0.7487	1.141	2990
Fixed effects:		Turf filament length ~ microhabitat type + site - 1		
	Post. mean parameter est.	lower 95% CI	upper 95% CI	Eff. sample size
type(flat)	2.46588	1.98738	2.92477	2990
type(exposed)	0.29921	-0.06599	0.63978	2849
type(concealed)	4.41391	4.05343	4.75424	3169
site1	2.46588	1.98738	2.92477	2990
site2	0.03930	-0.62722	0.60480	2808

There was also a distinct difference in the mean number of species feeding on the three microhabitat types (species richness). While flat and exposed microhabitats were statistically indistinguishable (flat: 3.85 ± 0.16 [mean \pm SE] number of species feeding on a given microhabitat over the entire observation period; exposed: 3.18 ± 0.17 species), concealed microhabitats had substantially less grazer species richness (1.93 ± 0.14 species). Interestingly, patterns of mean functional group richness of the benthic communities showed the opposite trend, with flat microhabitats supporting a markedly lower number of functional groups (3.15 ± 0.15 [mean \pm SE] number of groups per microhabitat), compared to both exposed and concealed microhabitats (exposed: 4.333 ± 0.14 functional groups; concealed: 4.75 ± 0.15 functional groups). Again, site had no effect on the observed patterns and random variance was small (**Tables 8.4, 8.5**). Therefore, site data were pooled for subsequent analyses.

Table 8.4: Model output from the Poisson Bayesian Mixed Model comparing the number of grazer species observed to feed on a given microhabitat among microhabitat types and sites. Microhabitat type and site were specified as fixed effects with uninformative priors ($\mu = 0$, $V = 100,000,000$). Uninformative priors were used for the residual and random variance parameters ($V = 1$, $\nu = 0.002$). Estimates for effect levels are based on a global intercept containing estimates for microhabitat type “flat” and site 1. Where 95% CIs do not intersect 0, statistically significant differences can be inferred.

Grazer species richness model

Iterations: 10001:299001
 Thinning interval: 1000
 Sample size: 2990

G-structure (random variance):		~array		
	Post. mean parameter est.	lower 95% CI	upper 95% CI	Eff. sample size
array	0.04014	0.0002354	0.1269	2990
R-structure (residual variance):		~units		
	Post. mean parameter est.	lower 95% CI	upper 95% CI	Eff. sample size
units	0.003819	0.0001635	0.01174	2990
Fixed effects:		Number of grazer species ~ microhabitat type + site		
	Post. mean parameter est.	lower 95% CI	upper 95% CI	Eff. sample size
type(flat)	1.169238	0.909828	1.434785	2990
type(exposed)	-0.193356	-0.377083	0.006359	2637
type(concealed)	-0.692251	-0.925175	-0.474703	2311
site1	1.169238	0.909828	1.434785	2990
site2	0.311634	-0.022325	0.668442	3262

Table 8.5: Model output from the Poisson Bayesian Mixed Model comparing the number of benthic functional groups present on each microhabitat among microhabitat types and sites. Microhabitat type and site were specified as fixed effects with uninformative priors ($\mu = 0$, $V = 100,000,000$). Uninformative priors were used for the residual and random variance parameters ($V = 1$, $\nu = 0.002$). Estimates for effect levels are based on a global intercept containing estimates for microhabitat type “flat” and site 1. Where 95% CIs do not intersect 0, statistically significant differences can be inferred.

Benthic functional group richness model

Iterations: 10001:299001
 Thinning interval: 1000
 Sample size: 2990

G-structure (random variance):		~array		
	Post. mean parameter est.	lower 95% CI	upper 95% CI	Eff. sample size
array	0.02866	0.0001762	0.08015	2990
R-structure (residual variance):		~units		
	Post. mean parameter est.	lower 95% CI	upper 95% CI	Eff. sample size
units	0.002871	0.0001734	0.008534	2990
Fixed effects:		Number of functional groups ~ microhabitat type + site		
	Post. mean parameter est.	lower 95% CI	upper 95% CI	Eff. sample size
type(flat)	1.17987	0.95788	1.38807	2990
type(exposed)	0.31975	0.14029	0.50751	2065
type(concealed)	0.41193	0.23086	0.60103	2589
Site1	1.17987	0.95788	1.38807	2990
Site2	-0.07846	-0.21591	0.06792	2990

8.3.2 Community compositions

The three microhabitat types showed distinct differences in the grazer and benthic communities (**Figure 8.3**). The nMDS ordination for the grazer communities showed a clear separation between concealed microhabitats and the two other microhabitat types, but not between flat and exposed microhabitats (**Figure 8.3a**). The separation was driven largely by rabbitfishes (*Siganus corallinus*, *S. punctatus*, *S. punctatissimus*), and the two surgeonfish species *Zebrasoma scopas* and *Acanthurus nigrofuscus*, fishes that characteristically grazed in concealed microhabitats. In contrast, flat and exposed microhabitats were characterized by parrotfishes (*Scarus frenatus*, *S. globiceps*, *S. niger*, *S. psittacus*, *S. schlegeli*, *Chlorurus spilurus*) and the surgeonfish species *A. nigricauda* and *Ctenochaetus striatus*. This separation of microhabitat types was statistically significant in the PERMANOVA (pseudo $F_{2,174} = 37.12$, $P < 0.001$), explaining approximately one third of the variation ($R^2 = 0.30$).

Benthic communities likewise varied markedly, with the nMDS ordination again showing a clear distinction between concealed microhabitats and the two remaining microhabitats, which showed extensive overlap (**Figure 8.3b**). While concealed microhabitats were composed of a wide array of benthic organisms, including scleractinian corals and various stands of filamentous, fleshy, and foliose algae, as well as fish faeces, both flat and exposed microhabitats showed a more uniform community composition, comprising a polycultural turf algal mat, occasionally with exposed crustose coralline algae (CCA), cyanobacterial mats, and a few non-coralline sessile invertebrates (on exposed microhabitats). Again, these differences distinctly characterized the respective microhabitat types in the PERMANOVA (pseudo $F_{2,180} = 128.16$, $P < 0.001$), explaining more than half of the variance ($R^2 = 0.59$).

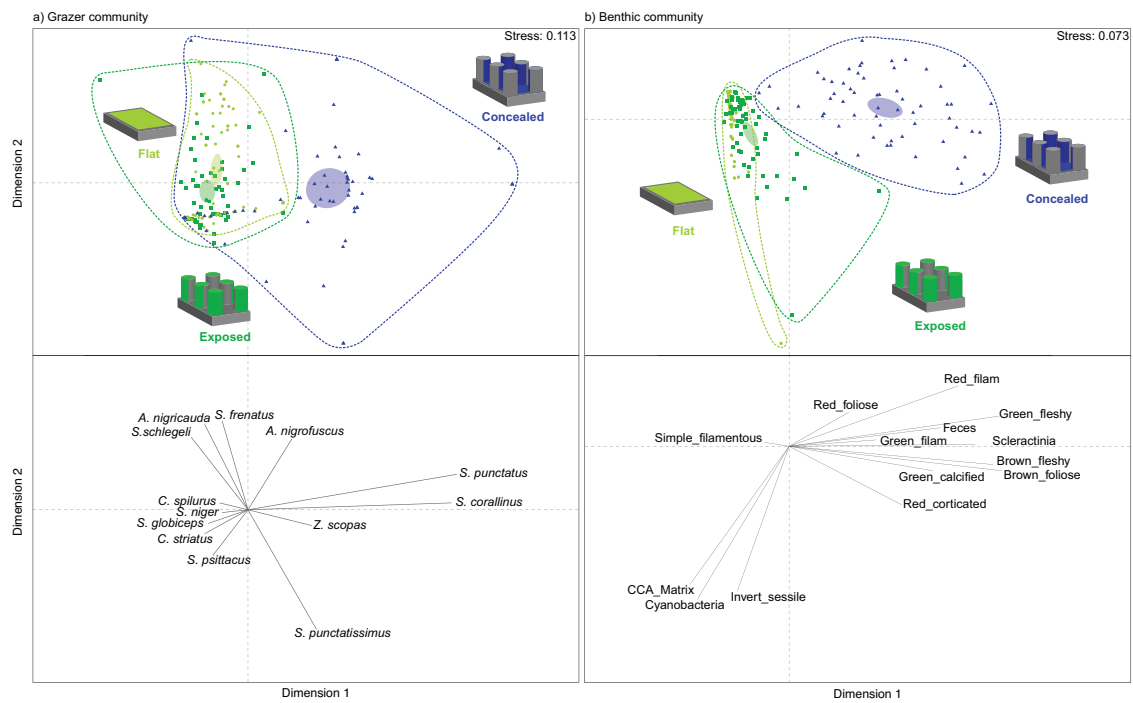
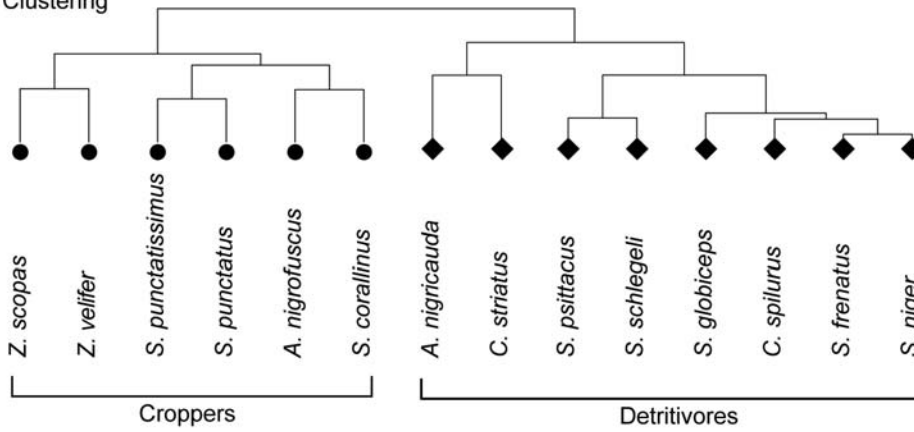


Figure 8.3: The grazer communities and benthic communities of the three microhabitat types. Ordinations represent nMDS ordinations with each symbol representing a microhabitat type. Shaded ellipses represent the 95% confidence intervals of the centroids, while coloured dashed lines comprise all samples within a given microhabitat. Bottom panels represent vectors. Concealed microhabitats are markedly different from flat and exposed habitats with regards to both their grazer communities (a), and benthic community composition (b). For the grazer community, six concealed microhabitats were omitted due to no grazing.

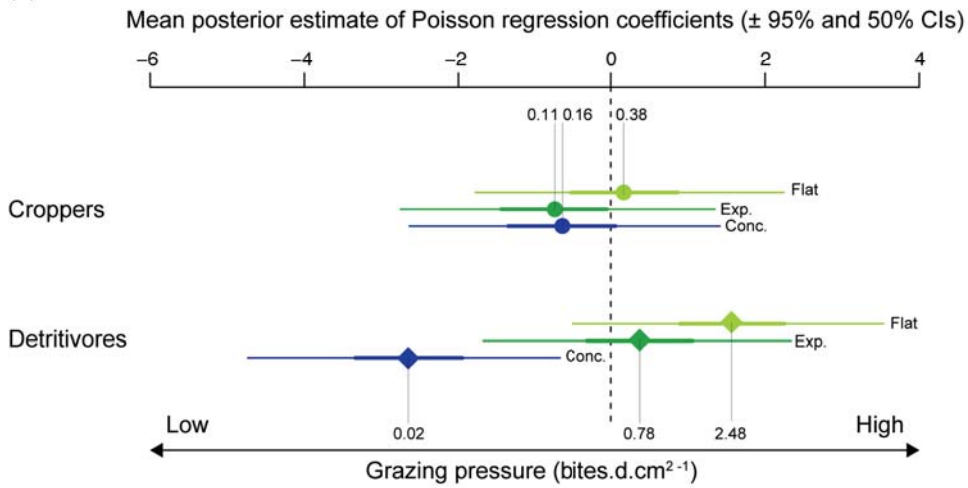
8.3.3 Functional profiles of grazer communities

When dividing grazer communities into functional groups based on their morphology, gut contents, and SCFA profiles, I found further, and consistent, differences in the grazing pressure exerted on the three microhabitat types. The first group encompassed the fish community typically feeding on concealed microhabitats (see nMDS above) and consisted of all observed rabbitfishes (*S. corallinus*, *S. punctatissimus*, *S. punctatus*), both surgeonfishes in the genus *Zebrasoma* (*Z. scopas*, *Z. velifer*), and *Acanthurus nigrofuscus*. Collectively, these species are ‘croppers’, feeding largely on the apical portions of algae. The second group encompassed the fish community typically feeding on flat and exposed microhabitats (see nMDS above), and comprised all observed parrotfishes (*C. spilurus*, *S. frenatus*, *S. globiceps*, *S. niger*, *S. psittacus*, *S. schlegeli*), and the two surgeonfish species *C. striatus* and *A. nigricauda*. Collectively, these species are ‘detritivores’, which remove both filamentous algae and particulate matter, or just the particulate portion of the epilithic algal matrix, when feeding. There were no differences in the grazing pressure exerted on the respective microhabitat types by croppers, as both bite counts and the probability of zero counts were similar among microhabitats. In strong contrast, grazing pressure by detritivores differed extensively, with concealed microhabitats being substantially less heavily-grazed and having a much higher probability of zero counts than flat or exposed microhabitats, for which zero-counts were absent (**Figure 8.4, Table 8.6**).

(a) Clustering



(b) Bite counts



(c) Zero-inflation

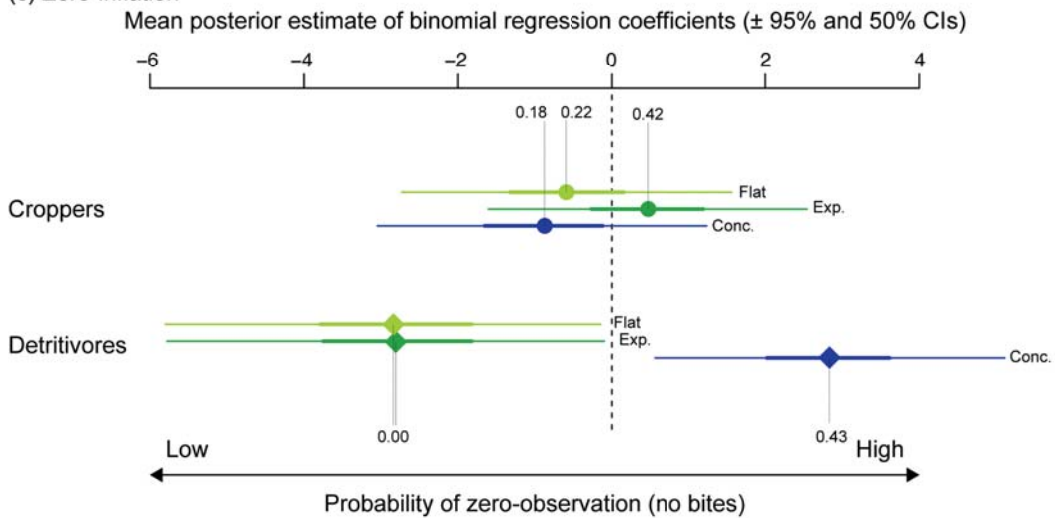


Figure 8.4: The grazing pressure of the two functional groups on the respective microhabitats. Ward's clustering analysis of the trait dataset (a) divided herbivores into two functional groups, croppers and detritivores. The model coefficients from a zero-inflated Poisson BMM indicate that there are no differences in the grazing pressure exerted on different microhabitats by croppers, with regards to both the bite rate (b) and the likelihood of no bites at all (c). In stark contrast, concealed microhabitats are significantly less grazed on by detritivores, as indicated by both the bite counts and zero-probability.

Table 8.6: Model output from the zero-inflated Poisson Bayesian Mixed Model comparing the grazing pressure of croppers and detritivores, respectively, among the three microhabitat types. Due to complete separation in the data, weakly informative priors were used on the fixed effect microhabitat type ($\mu = 0$, $V = 1 + \pi^{2/3}$), while highly informative priors were specified for the two offset parameters controlling for time and area ($\mu = 1$, $V = 0.0000001$). For the Poisson component of the model, uninformative priors were used on the residual and random variance ($V = 1$, $\nu = 0.002$), while residual and random variance were fixed at 1 and 0.000001, respectively, for the binomial zero-inflation component of the model. Microhabitat type was specified to interact with both the Poisson and the binomial part of the model to obtain estimates of the effects of microhabitat type on the grazing rates and the likelihood of complete avoidance (zero counts). To facilitate interpretation, the global intercept was suppressed.

Functional group grazing pressure model: (a) Croppers

Iterations: 10001:299001
 Thinning interval: 1000
 Sample size: 2990

G-structure (random variance):		Post. mean parameter est.	lower 95% CI	upper 95% CI	Eff. sample size
array	~array	0.055537	0.0002152	0.195891	2990
ZI-array		0.000001	0.000001	0.000001	0

R-structure (residual variance):		Post. mean parameter est.	lower 95% CI	upper 95% CI	Eff. sample size
units	~units	0.6144	0.4393	0.8112	2990
ZI-units		1.0000	1.0000	1.0000	0

Fixed effects:		Post. mean parameter est.	lower 95% CI	upper 95% CI	Eff. sample size
Number of bites ~ trait - 1 + trait(Poisson)*microhabitat type + trait(ZI)*microhabitat type + trait(Poisson)*log(time.d) + trait(Poisson)*log(cm.sq)					

P(bite)	-1.2636	-3.2340	0.8421	2800
ZI(bite)	-0.9651	-2.9121	1.1923	2990
P(bite)*type(flat)	0.1786	-1.8414	2.2165	2780
P(bite)*type(exp)	-0.7377	-2.9541	1.1963	2788
P(bite)*type(con)	-0.6485	-2.6633	1.3711	2781
ZI(bite)*type(flat)	-0.5873	-2.6187	1.5552	2990
ZI(bite)*type(exp)	0.4780	-1.5740	2.6354	2990
ZI(bite)*type(con)	-0.8578	-2.9655	1.3290	3351
P(bite)*log(time)	1.0000	0.9982	1.0019	2990
P(bite)*log(area)	1.0000	0.9981	1.0020	2990

Functional group grazing pressure model: (b) Detritivores

Iterations: 10001:299001
 Thinning interval: 1000
 Sample size: 2990

G-structure (random variance):		Post. mean parameter est.	lower 95% CI	upper 95% CI	Eff. sample size
array	~array	0.144413	0.012310	0.410385	2990
ZI-array		0.000001	0.000001	0.000001	0
R-structure (residual variance):		Post. mean parameter est.	lower 95% CI	upper 95% CI	Eff. sample size
units	~units	0.2097	0.154	0.2717	2983
ZI-units		1.0000	1.0000	1.0000	0
Fixed effects:		Post. mean parameter est.	lower 95% CI	upper 95% CI	Eff. sample size
Number of bites ~ trait + trait(Poisson)*microhabitat type + trait(ZI)*microhabitat type + trait(Poisson)*log(time.d) + trait(Poisson)*log(cm.sq)					
Poisson		-0.7581	-2.8416	1.2167	2990
ZI		-2.7932	-5.1436	-0.6116	1366.8
Poisson*type(flat)		1.5777	-0.3790	3.6576	2990
Poisson*type(exp)		0.3793	-1.5568	2.4992	2990
Poisson*type(con)		-2.6341	-4.5617	-0.4463	2990
ZI*type(flat)		-2.8630	-5.7930	-0.1750	744.2
ZI*type(exp)		-2.8041	-5.8088	-0.2844	745.4
ZI*type(con)		2.8661	0.5453	5.0674	1343.2
Poisson*log(time)		1.0000	0.9982	1.0019	2990
Poisson*log(area)		1.0000	0.9981	1.0020	2990

The fundamental differences in the feeding pressure exerted on the three microhabitat types by croppers and detritivores is closely reflected in the bite-weighted average functional profiles of fishes feeding on the respective microhabitats. Compared to fishes grazing on flat and exposed microhabitats, fishes feeding on concealed microhabitats exhibit larger head angles (i.e. a concave head), smaller snout angles (i.e. pointed snouts), larger eyes, and deeper bodies. Furthermore, fishes grazing on concealed microhabitats exhibit higher proportions of fleshy algae, filamentous algae, and other benthic material in their guts, and substantially lower amounts of calcareous sediments and organic particulates. These dietary differences are also apparent in the SCFA profiles, with fishes feeding on concealed microhabitats exhibiting high levels of SCFAs, containing high levels of acetate and propionate but low levels of isovalerate (**Figure 8.5, Table 8.7**).

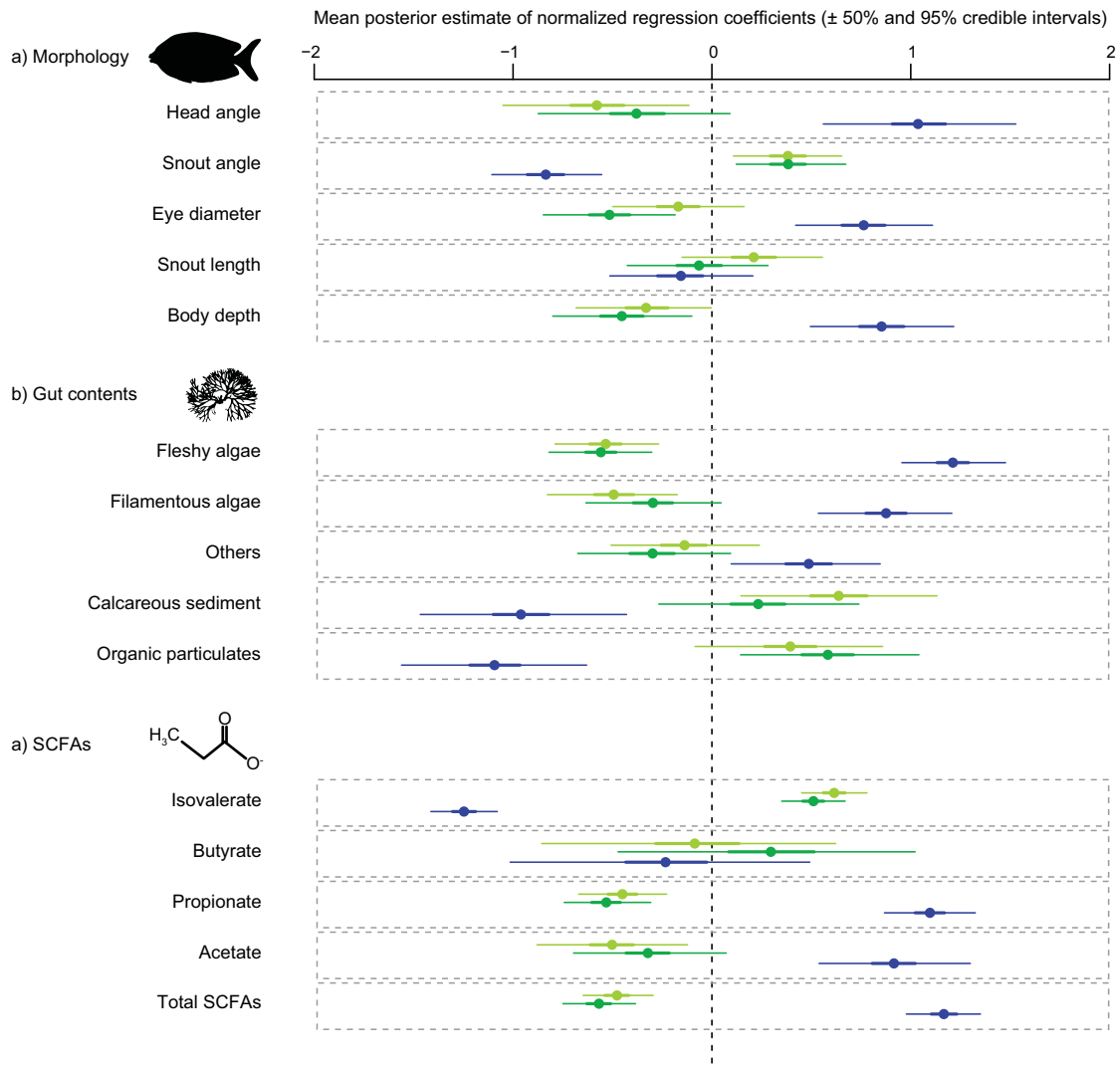


Figure 8.5: The average trait values of consumer communities feeding on the three microhabitat types. There were strong differences between concealed and exposed microhabitats in terms of morphology (a), gut contents (b), and SCFA profiles (c). Consumer communities feeding on concealed microhabitats were characterized by concave foreheads, pointed snouts, large eyes, and deep bodies. They predominantly consume fleshy and filamentous algae, while taking only little calcareous sediment and organic particulates. This is supported by the SCFA profiles, which show low levels of isovalerate, but high levels of propionate, acetate,

and total SCFA concentrations. Light green = flat microhabitats, dark green = exposed microhabitats, blue = concealed microhabitats.

Table 8.7: Model output from the three multi-response models comparing the bite-weighted average trait means of consumers in terms of morphology, diet, and SCFAs among the three microhabitat types. For all three model, uninformative priors were specified on the fixed effect (microhabitat type). For the random variance, parameter expanded priors ($V = 1$, $\nu = 5.02$, $\alpha.\mu = 0$, $\alpha.V = 1000$) were specified on an idh-structure, while residual variance was specified to co-vary among the different response variables under an us-structure (priors: $V = 1$, $\nu = 5.02$). To facilitate interpretation, the global intercept was suppressed.

(a) Morphology

Iterations: 10001:2999001
 Thinning interval: 1000
 Sample size: 2990

G-structure (random variance):		~idh(trait):array			
	Post. mean parameter est.	lower 95% CI	upper 95% CI	Eff. Sample size	
Body depth.array	0.132	0.000	0.419	2990	
Snout length.array	0.121	0.000	0.332	2592	
Eye diameter.array	0.103	0.000	0.348	2990	
Snout angle.array	0.042	0.000	0.145	2990	
Head angle.array	0.328	0.021	0.946	2990	

R-structure (residual variance-co-variance)		~us(trait):array				
		Body d	Snout l	Eye d	Snout a	Head a
Body depth	Post. Mean	0.631				
	lower 95% CI	0.498				
	upper 95% CI	0.767				
Snout length	Post. Mean	0.587	0.977			
	lower 95% CI	0.446	0.771			
	upper 95% CI	0.747	1.195			
Eye diameter	Post. Mean	-0.160	-0.312	0.719		
	lower 95% CI	-0.262	-0.458	0.550		
	upper 95% CI	-0.052	-0.180	0.866		
Snout angle	Post. Mean	-0.553	-0.506	0.367	0.714	
	lower 95% CI	-0.677	-0.656	0.243	0.564	
	upper 95% CI	-0.416	-0.356	0.491	0.868	
Head angle	Post. mean	0.329	0.154	0.026	-0.336	0.411

	lower 95% CI	0.244	0.061	-0.057	-0.428	0.329
	upper 95% CI	0.426	0.256	0.110	-0.242	0.501
<hr/>						
Fixed effects:	(body d, snout l, eye d, snout a, head a) ~ (trait:microhabitat type - 1)					
	Post. mean parameter est.	lower 95% CI	upper 95% CI	Eff. Sample size		
Body d*type(flat)	-0.331	-0.677	-0.002	2990		
Snout l*type(flat)	0.210	-0.142	0.559	2781		
Eye d*type(flat)	-0.169	-0.508	0.151	3225		
Snout a*type(flat)	0.382	0.099	0.638	3263		
Head a*type(flat)	-0.578	-1.008	-0.083	2990		
Body d*type(exp)	-0.453	-0.797	-0.098	2488		
Snout l*type(exp)	-0.065	-0.434	0.268	2990		
Eye d*type(exp)	-0.515	-0.835	-0.179	2990		
Snout a*type(exp)	0.384	0.125	0.674	2830		
Head a*type(exp)	-0.380	-0.876	0.086	2990		
Body d*type(con)	0.853	0.482	1.202	2428		
Snout l*type(con)	-0.156	-0.498	0.215	2990		
Eye d*type(con)	0.763	0.431	1.114	2990		
Snout a*type(con)	-0.835	-1.107	-0.554	2740		
Head a*type(con)	1.037	0.532	1.491	2990		

(b) Diet

Iterations: 10001:2999001
 Thinning interval: 1000
 Sample size: 2990

G-structure (random variance):	~idh(trait):array					
	Post. mean parameter est.	lower 95% CI	upper 95% CI	Eff. Sample size		
Organic_matter.array	0.341	0.017	0.880	1612		
Calc_sediment.array	0.324	0.019	0.955	2718		
Others.array	0.133	0.000	0.410	3172		
Filamentous.array	0.097	0.000	0.318	2990		
Fleshy.array	0.074	0.000	0.220	2832		

R-structure (residual variance-co- variance)	~us(trait):array					
		Organics	Sedimen t	Others	Filame nts	Fleshy
Organic matter	Post. Mean	0.353				
	lower 95% CI	0.279				
	upper 95% CI	0.431				
Calcareous sediment	Post. Mean	0.097	0.509			
	lower 95% CI	0.031	0.412			
	upper 95% CI	0.166	0.628			

Others	Post. Mean	0.016	0.225	0.893		
	lower 95% CI	-0.078	0.117	0.702		
	upper 95% CI	0.094	0.335	1.075		
Filamentous algae	Post. Mean	-0.360	-0.237	-0.247	0.640	
	lower 95% CI	-0.454	-0.331	-0.369	0.508	
	upper 95% CI	-0.272	-0.145	-0.130	0.787	
Fleshy algae	Post. mean	-0.221	-0.304	-0.152	0.258	0.338
	lower 95% CI	-0.287	-0.388	-0.247	0.179	0.268
	upper 95% CI	-0.159	-0.231	-0.072	0.342	0.414

Fixed effects: (organics, sediment, others, filamentous, fleshy) ~ (trait:microhabitat type - 1)

	Post. mean parameter est.	lower 95% CI	upper 95% CI	Eff. Sample size
Organics*type(flat)	0.394	-0.041	0.879	2990
Sediment*type(flat)	0.638	0.141	1.122	2990
Others*type(flat)	-0.138	-0.523	0.214	2990
Filamentous*type(flat)	-0.494	-0.843	-0.197	2990
Fleshy*type(flat)	-0.534	-0.775	-0.258	2990
Organics*type(exp)	0.583	0.126	1.014	2990
Sediment*type(exp)	0.233	-0.284	0.707	3411
Others*type(exp)	-0.298	-0.693	0.072	3107
Filamentous*type(exp)	-0.297	-0.650	0.019	3562
Fleshy*type(exp)	-0.558	-0.838	-0.322	2828
Organics*type(con)	-1.093	-1.546	-0.621	2990
Sediment*type(con)	-0.961	-1.521	-0.491	3486
Others*type(con)	0.487	0.110	0.856	2990
Filamentous*type(con)	0.876	0.527	1.190	2990
Fleshy*type(con)	1.212	0.953	1.471	4019

(c) SCFAs

Iterations: 10001:2999001
 Thinning interval: 1000
 Sample size: 2990

G-structure (random variance):	~idh(trait):array			
	Post. mean parameter est.	lower 95% CI	upper 95% CI	Eff. Sample size
totalscfa.array	0.009	0.000	0.034	2990
acetate.array	0.163	0.003	0.516	2990
propionate.array	0.027	0.000	0.092	2710
butyrate.array	0.896	0.045	2.273	2990
isovalerate.array	0.010	0.000	0.044	2822

R-structure (residual variance-co-variance) ~us(trait):array

		Total	Acetate	Prop	But	Iso
Total SCFA	Post. Mean	0.419				
	lower 95% CI	0.328				
	upper 95% CI	0.508				
Acetate	Post. Mean	-0.037	0.572			
	lower 95% CI	-0.110	0.455			
	upper 95% CI	0.039	0.702			
Propionate	Post. Mean	0.379	-0.128	0.485		
	lower 95% CI	0.294	-0.212	0.385		
	upper 95% CI	0.473	-0.050	0.592		
Butyrate	Post. Mean	0.074	-0.116	0.021	0.717	
	lower 95% CI	-0.007	-0.212	-0.069	0.565	
	upper 95% CI	0.163	-0.050	0.114	0.880	
Isovalerate	Post. mean	-0.247	-0.216	-0.237	-0.025	0.322
	lower 95% CI	-0.312	-0.290	-0.305	-0.098	0.254
	upper 95% CI	-0.178	-0.143	-0.166	0.049	0.389
Fixed effects:		(total scfa, acetate, propionate, butyrate, isovalerate) ~ (trait: microhabitat type - 1)				
	Post. mean parameter est.	lower 95% CI	upper 95% CI	Eff. Sample size		
TotalSCFA*type(flat)	-0.478	-0.648	-0.296	2990		
Acetate*type(flat)	-0.503	-0.855	-0.107	2990		
Propionate*type(flat)	-0.451	-0.657	-0.216	2990		
Butyrate*type(flat)	-0.087	-0.856	0.625	2990		
Isovalerate*type(flat)	0.614	0.456	0.783	3085		
TotalSCFA*type(exp)	-0.569	-0.746	-0.383	2990		
Acetate*type(exp)	-0.323	-0.728	0.036	3890		
Propionate*type(exp)	-0.532	-0.741	-0.306	2990		
Butyrate*type(exp)	0.295	-0.522	0.966	2990		
Isovalerate*type(exp)	0.510	0.358	0.676	2990		
TotalSCFA*type(con)	1.165	0.978	1.351	2990		
Acetate*type(con)	0.914	0.543	1.299	2990		
Propionate*type(con)	1.096	0.877	1.332	3742		
Butyrate*type(con)	-0.234	-0.984	0.501	2990		
Isovalerate*type(con)	-1.248	-1.410	-1.078	2810		

8.4 Discussion

My results reveal that micro-topographic refuges profoundly affect the dynamics between grazing herbivorous fishes and benthic organisms on coral reefs. Specifically, increased complexity at the scale of a few centimetres alters the grazing rates of consumer communities comparable to variation previously reported on coral reefs at local, regional, and global scales. Furthermore, micro-topographic refuges fundamentally change the functional identity of the consumer communities, essentially restricting the spectrum of herbivorous fishes that can feed on concealed microhabitats to few algae-cropping species. In contrast, detritivorous species, which incorporate large proportions of organic particulates and calcareous sediments into their diet, appear to be excluded from concealed microhabitats, although they exert intense grazing pressure on flat and exposed microhabitats. Thus, micro-topographic refuges appear to eliminate an entire ecosystem process, the removal of particulates. As a consequence, the benthic communities vary drastically between concealed and flat or exposed microhabitats. While algal turfs dominate both communities, turf filaments are markedly longer in concealed microhabitats and incorporate a wide range of algal forms and other benthic components, including juvenile scleractinian corals. I suggest that the refuge from grazing provided by micro-topographic complexity creates an environment that fosters benthic diversity through intermediate grazing pressure by a few selected cropping species.

8.4.1. Effects of micro-topographic refuges on grazing

Micro-topographic refuges clearly reduce the grazing pressure exerted by herbivorous fishes, resulting in a more than tenfold decrease in the number of bites per day per cm² within refuges. This corroborates the results of previous studies in both aquatic and terrestrial environments (Menge and Lubchenco 1981; Dudley and D'Antonio 1991; Bergey 2005; Shitzer *et al.* 2008). In the context of coral reefs, the effect of micro-topographic complexity

on grazing rates rivals well-documented differences at larger scales. The approximately ten-fold differences observed between flat and concealed microhabitats resemble the variation in grazing pressure outside and inside of damselfish territories (Hixon and Brostoff 1996), and are comparable with the variation along some major reef habitats. Specifically, Hay (1981) demonstrated a tenfold difference in the consumption of *Thalassia* blades between seaward sides of outer fore-reefs (9.0% consumed) and rubble areas of back reefs (92.8% consumed). Furthermore, if grazing pressure scales approximately linearly with herbivorous reef fish biomass, then the difference between open and concealed microhabitats exceeds both regional (0.8 to 7.2kg.100m⁻² on the inner shelf of the GBR, compared to 5.0 to 31.7kg.100m⁻² on the outer shelf, Wismer *et al.* 2009) and global disparities in grazing pressure on coral reefs (mean of 20.9g.m⁻² in Brazil compared to 83.7g.m⁻² in New Caledonia; supplementray material in Edwards *et al.* 2014). The gradient reported herein is also greater than the gradient in herbivorous fish biomass induced by fishing pressure globally (20.5g.m⁻² in fished areas compared to 56.4g.m⁻², Edwards *et al.* 2014) and exceeds herbivore biomass reductions achieved through the employment of herbivore exclusion cages (Hughes *et al.* 2007). Thus, the presence of micro-topographic complexity on the scale of a few centimetres appears to profoundly affect consumer pressure, equivalent to some of the largest gradients in grazing rates reported for coral reef ecosystems.

Besides overall grazing pressure, however, micro-topographic refuges profoundly affect the functional diversity of consumer species capable of interacting with the benthos. Concealed microhabitats are only grazed by a subset of fish species, resulting in marked reductions of consumer taxonomic and functional diversity. Microhabitat utilization has been recognized as a strong axis of niche partitioning in herbivorous fishes (Brandl and Bellwood 2014; Adam *et al.* 2015), but the effects of this environmental filter on functional processes performed by herbivorous fishes are largely unknown. Our results show that there is a clear

morphological segregation, as feeding in concealed microhabitats appears to be restricted to ‘cropping’ species, with relatively large eyes, deep bodies, concave foreheads, and pointed snouts, all of which favour the exploitation of small prey items in complex environments (Brandl and Bellwood 2013). Furthermore, micro-topographic complexity almost completely eliminates feeding by ‘detritivorous’ fishes that remove organic and inorganic particulate material. Previously reported dietary information (Choat et al. 2002) and evidence of digestive use of ingested material (SCFA profiles; Clements and Choat 1995), suggest that there is very little extraction of either the entire epilithic algal matrix (including particulates; Wilson et al. 2003), or the particulate component alone, in concealed microhabitats. Instead, grazing pressure exerted on concealed microhabitats is almost exclusively restricted to the selective cropping of the apical portions of both filamentous and foliose/fleshy algae, leaving the base layers of the turf community largely intact (Hixon and Brostoff 1996). Thus, the identity of consumer communities and the ecosystem processes they deliver differ vastly among microhabitats. This segregation may explain previously reported effects of herbivore identity and species richness on benthic community composition and succession on coral reefs (Burkepile and Hay 2008, 2010; Cheal *et al.* 2010).

Despite these striking functional differences between grazer species, many of the species observed in the present study have been consistently placed in the same functional group (grazers/detritivores) in order to assess their status or ecosystem function (e.g. Green and Bellwood 2009; Cheal et al. 2010; Doropoulos et al. 2013; Edwards et al. 2014). For example, *Acanthurus nigrofuscus* and *Ctenochaetus striatus* have been considered to be functionally equivalent in several instances (Green and Bellwood 2009; Marshall and Mumby 2011); yet, my results demonstrate that they vary drastically in their use of microhabitats and resources therein. This is likewise applicable to *Acanthurus nigrofuscus* and the rabbitfishes observed in this study (Green and Bellwood 2009). Hence, the results of the present study

emphasize the need for caution when assigning species to functional groups and assuming within-group functional equivalence. Clearly, differences in the utilization of foraging microhabitats need to be considered when characterizing species with regards to their functional impact (Adams et al. 2015) and indeed, it would not be surprising if even finer partitioning of foraging microhabitats, dietary items, or digestive use of resources would create a unique functional role for most herbivorous fish species on coral reefs.

8.4.2. Effects of micro-topographic refuges on benthic communities

As expected, given the differences in both grazing pressure and the functional identity of grazers, the benthic communities varied strongly among microhabitat types. In close agreement with the reduced grazing pressure, we found an almost three-fold increase in the length of turf algal filaments in concealed microhabitats. This closely matches situations in which herbivory is naturally or experimentally suppressed (Arnold et al. 2010; Bonaldo and Bellwood 2011; Rasher et al. 2012). The lack of a difference between turf filaments growing on flat and exposed microhabitats suggests that there is a grazing threshold above which short turf filaments are maintained. Specifically, lower grazing pressure on exposed microhabitats (0.81 ± 0.05 bites.day⁻¹.cm⁻²) appears to be sufficient to maintain a closely cropped turf community, while the even lower grazing pressure on concealed microhabitats (0.17 ± 0.02 bites.day⁻¹.cm⁻²) clearly allows for longer turf filaments, suggesting a threshold between 0.17 and 0.81 bites.day⁻¹.cm⁻² (**Figure 8.2**).

While algal turfs dominated all microhabitats, we found a marked difference between the benthic communities of open and concealed microhabitats. This highlights a fundamental problem when describing benthic communities on coral reefs merely as algal turfs (or the epilithic algal matrix), as these communities are far from uniform (Connell et al. 2014; Harris et al. 2015). The benthic community on flat and exposed microhabitats appears to closely

resemble a state of ‘successional deflection’ (*sensu* Hixon and Brostoff 1996), where extreme and destructive grazing by a wide range of fish species only permits the existence of short, low-complexity algal filaments, CCA, and cyanobacterial mats (Hixon and Brostoff 1996; Wismer et al. 2009). In contrast, the benthic community in concealed microhabitats was far more diverse, including a wide array of algal growth forms as well as juvenile scleractinian corals. This not only corroborates previous findings, which suggest that micro-topographic complexity can be beneficial to benthic organisms by providing a refuge from grazing pressure (Brandl et al. 2014; Edmunds et al. 2014; Franco et al. 2015), but it also supports the role of micro-topographic complexity, or any refuge from intense grazing, in fostering benthic diversity across ecosystems (Menge and Lubchenco 1981; Hixon and Brostoff 1985, 1996; Casey et al. 2014). Indeed, the communities present in concealed microhabitats appear analogous to a successional state of peak diversity (*sensu* Hixon and Brostoff 1996), which is commonly maintained by intermediate levels of consumer-mediated disturbance (Worm et al. 2002). This may apply herein, as detritivorous parrotfish species that scrape the entire turf community off the substratum, therefore representing a severe disturbance, are excluded from concealed microhabitats. In contrast, the selective cropping of algae by fishes feeding in concealed microhabitats may represent a partial and therefore intermediate level of disturbance.

Thus, micro-topographic refuges on coral reefs create two distinct regimes, which are maintained by two separate groups of grazers: detritivorous fishes, which feed on the entire epilithic algal matrix (EAM) or its particulate component, maintain a short, intensively grazed, low-diversity epilithic turf community, while cropping herbivores sustain a longer turf community with diverse algal growth forms and juvenile corals.

8.4.3. *Effects of micro-topographic refuges on coral reef ecosystems*

Our results suggest that the relationship between grazing herbivorous fishes and benthic communities is overwhelmingly influenced by micro-topographic refuges, and as such, far more complex than previously assumed. Since the identification of herbivory as a critical process for coral reef ecosystems, many studies have identified the direct positive effect of grazing herbivores on reef resilience (reviewed by Bellwood *et al.* 2004; Nyström *et al.* 2008; Roff and Mumby 2012; Graham *et al.* 2015). While there is clear evidence that it is the failure of coral recruitment, which causes the large-scale and long-term decline of coral-dominated reefs (Hughes and Tanner 2000; Hughes *et al.* 2007), our results suggest that micro-topographic refuges may play a critical role in the dynamics between herbivorous grazing and coral replenishment (**Figure 8.6**). Coral replenishment is tightly linked to the survival of juvenile corals, which depends on the avoidance of mortality induced by both intense grazing pressure and the smothering by algal turfs (Arnold *et al.* 2010; Steneck *et al.* 2014). My results suggest that by creating an interface between open and concealed microhabitats, micro-topographic refuges represent an environment that permits corals to avoid both threats, provided that the herbivore community comprises fishes capable of cropping algae within refuges, and detritivorous fishes that maintain a shortly grazed EAM outside refuges. In essence, by selectively trimming algae within refuges, cropping grazers prevent the smothering of coral recruits from within refuges. Simultaneously, detritivorous grazers maintain a short EAM outside refuges, enabling corals to access light whilst being sheltered from grazing damage within refuges. Thus, although easily overlooked in a world of global syntheses, micro-topographic complexity on the scale of a few centimetres, and the functional roles of herbivorous fishes grazing within and outside of micro-topographic refuges, may be a critical aspect of the dynamics between herbivorous fishes and the benthic community in coral reef ecosystems.

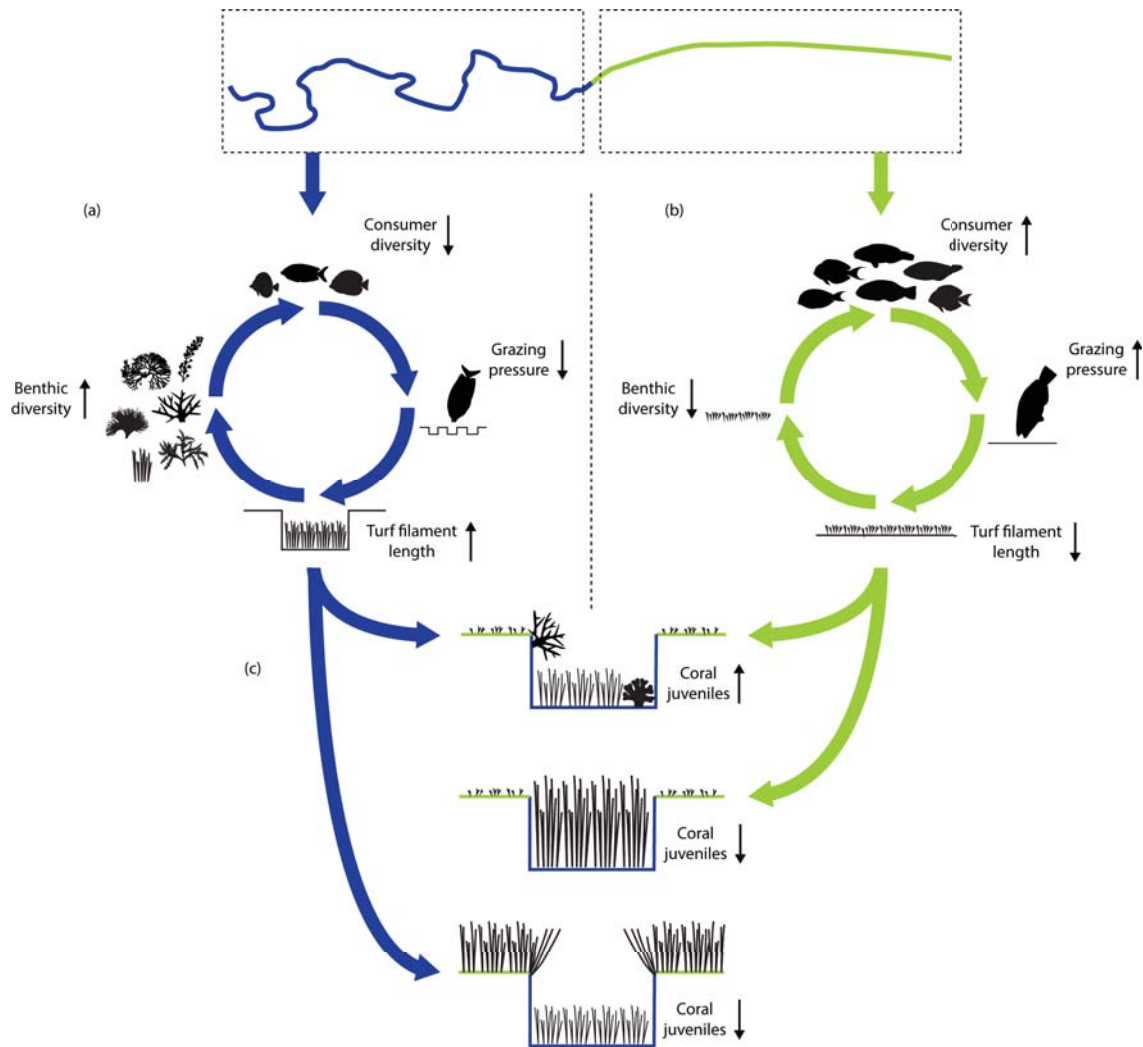


Figure 8.6: Conceptual synthesis of the effects of refuges. Refuges restrict the grazer community to a few cropping species, which exert moderate grazing pressure. Consequently, turf algal filaments grow longer and a diverse benthic community can establish (a). In contrast, a diverse assemblage (incl. croppers and detritivores), grazes on flat microhabitats, leading to intense overall grazing pressure. Algal filaments are tightly grazed and a low-diversity epilithic algal matrix is established (b). Thus, three possible scenarios emerge (c): if both regimes coexist, corals can settle in refuges to avoid grazing pressure, and can outgrow these microhabitats, as turfs are controlled within and outside refuges. If no croppers are

present, corals become smothered by algal turfs in refuges. If detritivores are absent, grazing pressure is reduced, but corals are smothered by algae expanding from flat microhabitats.

In summary, I demonstrate that micro-topographic complexity can act as an environmental refuge for benthic organisms by means of excluding a suite of high-intensity grazers, while permitting the grazing activity of low-intensity grazers that crop algal filaments. As a consequence, two fundamentally different benthic regimes can coexist at the scale of a few centimetres. I suggest that a balance between these two regimes underpins processes relating to coral replenishment on coral reefs and propose that the role of micro-topographic complexity and functional differences among herbivorous fishes be considered in future assessments of grazing dynamics on coral reefs.

Chapter 9: General Discussion

Although the interest in species diversity *per se* continues (Mora et al. 2011; Fisher et al. 2015), the question of what species are doing within a given ecosystem has established itself as one of the most urgent issues in ecology (Diaz & Cabido 2001; Cardinale et al. 2002; Naeem et al. 2012; Cardinale et al. 2012). Yet, as more and more information is compiled into global databases of species' functional traits (Kattge et al. 2011), the call for the consideration of realized niches, which include individual variability in environments governed by countless, and often unquantified, biotic and abiotic interactions, is intensifying (Schleuter et al. 2010; Violle et al. 2012; Albert et al. 2012; Brandl & Bellwood 2014a).

The importance of this approach is demonstrated in this thesis. In Chapter 2 and 3, I extracted social and morphological traits that can serve as predictors for species' fundamental niches, suggesting that pairing species with large eyes, concave foreheads, pointed snouts, and deep bodies may be able to feed in more topographically complex microhabitats. While Chapter 4 presented evidence for one of the mechanisms driving this relationship, it also shows that intraspecific plasticity in the social system can have significant implications for a species' realized functional niche. Chapter 5 then demonstrated, how species that have been consistently placed in the same functional group by previous work (Green et al. 2009; Cheal et al. 2012; Doropoulos et al. 2013), thus implying that their impact on reefs is equivalent, can vary dramatically in their functional role when realized niches are considered. This realization was facilitated by the introduction of a new framework for the community-wide analysis of realized niche overlap. Chapter 6 further supported the necessity of considering realized niches: while it is intuitive to infer that morphologically specialized species will have specialized realized niches, the opposite can be the case, as found in coral reef surgeonfishes. Finally, Chapter 7 and 8 provided observational and experimental evidence for how small

variation in the realized niches of species can markedly modify their functional role within ecosystems and how, in the case of herbivorous fishes, this may lead to significant changes in benthic communities. In disentangling the relationship between pair-formation, herbivorous fishes, micro-topographic complexity, and the benthic community on coral reefs, this thesis contributes to our knowledge of the effects of *realized* functional diversity for coral reef ecosystems. The information provided in this thesis, therefore, may be considered in future assessments of functional diversity on coral reefs, in order to improve the accuracy with which we can predict the contribution of biodiversity to ecosystem functioning in one of the world's most diverse ecosystems.

Consumer control of primary producers is extraordinarily strong on coral reefs and many studies have emphasized the importance of herbivorous fishes in preventing the proliferation of macroalgae (e.g. Hughes 1994; Bellwood et al. 2004; Hughes et al. 2007; Burkepille & Hay 2006; Graham et al. 2015). The investigation of herbivorous fishes and their functional roles on coral reefs has flourished throughout the last two decades of the 20th century (e.g. Robertson et al. 1979; Robertson & Gaines 1986; Choat & Bellwood 1985; Hay 1991; Hay & Fenical 1988; Choat & Clements 1993; Bellwood & Choat 1990; Bellwood 1995; Clements & Choat 1995), laying the foundation for subsequent functional classifications. Since then, many researchers have engaged in studies that monitor herbivore abundances (Floeter et al. 2005; Wismer et al. 2009; Cheal et al. 2012), assess herbivore biomass and diversity on large spatial scales (Edwards et al. 2014), investigate the effects of herbivore extraction (McClanahan et al. 2001; Bellwood et al. 2012), or examine the socio-ecological factors that can lead to over-exploitation of herbivorous fishes (Cinner et al. 2009; Cinner et al. 2012). While the advances made in these fields are arguably indispensable for our overarching goal to manage and conserve coral reefs, the vast majority of studies incorporating herbivore functional diversity into their framework use classification schemes

that are, at best, coarse representations of species' fundamental niches, suggesting extensive functional overlap between species within groups.

This contrasts strongly with the burgeoning evidence of differences in the realized niches of herbivorous fishes. Over the last five years, research has revealed marked differences among herbivorous fish species commonly considered to be functionally equal. These differences relate to their diet (Fox et al. 2009; Mantyka & Bellwood 2007a,b; Hoey & Bellwood 2011; Hoey et al. 2013; Rasher et al. 2013), spatial ecology (Marshall et al. 2011; Claisse et al. 2011; Fox & Bellwood 2011; Welsh & Bellwood 2012; Brandl & Bellwood 2013b), foraging microhabitat utilization (Cardoso et al. 2006; Fox & Bellwood 2013; Brandl & Bellwood 2014a [Chapter 5]), or their overall impact on the benthic community (Burkepile & Hay 2008; 2010; 2011; [Chapter 8]). Yet, unfortunately, little of the information provided by these studies has since been incorporated into studies attempting to extrapolate the functional structure of herbivore communities and the effects of human-mediated pressure on ecosystem function. This is concerning as it suggests that many studies underestimate the functional diversity of herbivorous fishes, and thus overestimate the extent of functional redundancy (Bellwood et al. 2003; Johannsson et al. 2013; Micheli et al. 2013). Thus, I posit that more dedicated efforts need to be made to incorporate current findings with regards to functional differences among herbivorous reef fishes, as provided in this thesis, into evaluations of herbivore diversity and ecosystem functioning. At present, the continued use of functional classification schemes that have been shown to be of limited resolution impedes our attempts to fully understand herbivory as an ecosystem process and the effect of current environmental changes on herbivorous grazing on coral reefs.

Several of the results presented in this thesis may provide an impetus for future research. Notably, the results of Chapters 2, 3, and 4 present pair-formation in fishes in a new light. While fishes are often assumed to associate predominantly for the purpose of

reproduction (Pratchett et al. 2006), I demonstrate that direct ecological benefits of associating with a partner may override potential reproductive advantages. Given the frequency of pair-formation in reef fishes (Brandl & Bellwood 2014b [Chapter 2]), the potential for case studies examining pairing behaviour from an ecological perspective is vast, and likely to yield other cases in which both reproductive and direct ecological benefits underpin pairing behaviour in fishes. Chapter 4 builds on the social-ecological aspect of pair-formation in fishes and, as such, presents rabbitfishes as a model organism for the study of the cognitive and social capacities that allow fishes to engage in cooperative relationships. While the last decade has seen the establishment of fishes as a group of organisms that can cope with the complex cognitive and social demands of cooperative interactions (Bshary et al. 2002; Bshary & Grutter 2006; Brown 2015), the cooperative vigilance system observed in rabbitfishes opens several avenues for research that could be considered. First, genetic parentage analyses would provide a useful tool in order to examine whether kinship may play a role in the maintenance of cooperation among pair members. Kinship has long been argued to be a crucial factor for the evolution of cooperation in animals (Axelrod & Hamilton 1981; Dugatkin 1997; Clutton-Brock et al. 2001; Clutton-Brock 2009), but its effect on cooperation between rabbitfishes has yet to be determined. While the pelagic larval phase of rabbitfishes (Thresher 1984) makes close kinship between pair members rather unlikely, it appears possible that kin selection plays a factor in maintaining high levels of cooperation. Second, further exploration of the factors that stabilize cooperation in pairs of coral reef rabbitfishes could be performed, such as experimental manipulations of the partner's cooperativeness or the ambient predation pressure. In addition, the potential role of fin-flicking behaviour as a communicative signal could be examined using experimental setups. Third, the cognitive demands of cooperation between rabbitfishes could be assessed by confronting rabbitfishes with challenges such as individual recognition or memorization of individual's previous

actions. Along with these assessments, a closer examination of life-history parameters in rabbitfish pairs would be helpful to determine the fishes' sex, age, and respective body condition. Finally, the reproductive strategies of rabbitfishes are still largely unexplored. Aside from anecdotal information, which suggests that pairing rabbitfishes spawn in large aggregations (Johannes 1981), no spawning observations have been made to date.

The results from the remaining chapters suggest that a closer examination of the interactions between micro-topographic complexity and the foraging behaviour of fishes may provide interesting information on consumer-producer dynamics on coral reefs. First, a detailed account of the benthic communities within crevices *in situ* could be performed, including assessments of particulate components. The dynamics of sediment and organic particulates, in particular, may provide interesting insights. In addition, a species-level assessment of the benthic communities in different types of concealed microhabitats (e.g. scaling in size) along with high-resolution examinations of the dietary preferences of herbivorous fishes, potentially making use of stable isotope analyses, may reveal interesting aspects of grazing dynamics on coral reefs and enable us to further unravel niche overlap among herbivorous fishes and its effect on the benthic community. Finally, a replication of the experiment presented in Chapter 8 in different locations would be desirable. It is striking that none of the genera identified to feed in concealed environments on the Great Barrier Reef are present in the Caribbean. If herbivory in concealed environments represent a critical ecosystem process on Indo-Pacific coral reefs, the question arises whether this process is equally important in the Caribbean, and if so, which species are responsible for performing the functional role. Given the comparably low diversity of Caribbean coral reefs, such approach would yield interesting details and may reveal one pathway that has led to the large-scale degradation of Caribbean coral reefs.

Finally, I suggest that the findings of this thesis be incorporated into larger-scale assessments of the functional diversity of herbivorous fishes. This could make use of either the morphological information provided in Brandl & Bellwood (2013a [Chapter 3]), or classify species based on their realized microhabitat realization in order to extract their contribution to overall ecosystem functioning. In this context, the assembly of a catalogue of morphological and categorical ecological traits for coral reef organisms may prove to be a valuable endeavour. In such a catalogue, current information about the fundamental and realized niches of coral reef organisms, such as fishes, could be regularly updated and made accessible to researchers, while encouraging authors to provide information about the functional niches of coral reef organisms in a standardized format. For instance, a similar framework to the one suggested in Brandl & Bellwood (2014a [Chapter 5]) may be applied in order to examine functional niche partitioning in consumer assemblages and standardized estimates of niche overlap could be entered in the database. If used continuously in various locations around the world, larger-scale evaluations of functional diversity and ecosystem processes will enable researchers to consider information on the realized niche of fishes, rather than their potential function within the ecosystem. Given the downward trajectory of coral reefs and the worldwide losses of diversity (Knowlton 2009), such approaches will become more and more critical in order to predict the response of reef systems to anthropogenic disturbances or the implementation of conservation measures to counteract these disturbances.

In summary, this thesis presents evidence for fine-scale functional niche partitioning among herbivorous coral reef fishes. Herbivorous fish species differ markedly in their use of different benthic foraging microhabitats, and ecological segregation that is facilitated by the differences in morphological features, social preferences, and environmental variation among

microhabitats. While this thesis is restricted to coral reef environments, processes similar to the ones described herein may also be operating in other ecosystems.

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Appendix A

Publications arising from this thesis:

1. Brandl SJ, Bellwood DR (2014). Pair-formation in coral reef fishes: an ecological perspective. *Oceanogr Mar Biol Annu Rev* 52, 1-80 (doi: 10.1201/b17143-2).
2. Brandl SJ, Bellwood DR (2013). Morphology, ecology, and sociality: can morphology predict pair-forming behaviour in coral reef fishes. *Coral Reefs* 32, 835-846 (doi: 10.1007/s00338-013-1042-0).
3. Brandl SJ, Bellwood DR. Coordinated, cooperative vigilance in pairing coral reef fishes permits the exploitation of cryptic resources. *Sci Rep* 5, 14556 (doi:10.1038/srep14556).
4. Brandl SJ, Bellwood DR (2014). Individual-based analyses reveal limited functional overlap in a coral reef fish community. *J Anim Ecol* 83, 661-670 (doi: 10.1111/1365-2656.12171).
5. Brandl SJ, Robbins WR, Bellwood DR. Exploring the nature of ecological specialization in a coral reef fish community: morphology, diet, and foraging microhabitat use. *Proc R Soc B* 20151147 (doi: 10.1098/rspb.2015.1147).
6. Brandl SJ, Hoey AS, Bellwood DR. (2014) Micro-topography mediates interactions between corals, algae, and herbivorous fishes on coral reefs. *Coral Reefs* 33, 421-430 (doi: 10.1007/s00338-013-1110-5).
7. Brandl SJ, Bellwood DR. Micro-topographic refuges drive consumer-producer dynamics by mediating consumer functional diversity (under review).

Appendix B

Publications during candidature not arising from this thesis:

1. Hoey AS, Brandl SJ, Bellwood DR (2013). Diet and cross-shelf distribution of rabbitfishes (f. Siganidae) on the northern Great Barrier Reef: implications for ecosystem function. *Coral Reefs* 32, 973-982 (doi: 10.1007/s00338-013-1043-z).
2. Bellwood DR, Goatley CHR, Brandl SJ, Bellwood O (2014). Fifty million years of herbivory: fossils, fishes and functional innovations. *Proc R Soc B* 281, 20133046 (doi: 10.1098/rspb.2013.3046).
3. Brooker RM, Munday PL, Brandl SJ, Jones GP (2014). Local extinction of a coral reef fish explained by inflexible prey choice. *Coral Reefs* 33, 891-896 (doi: 10.1007/s00338-014-1197-3).
4. Rocker MM, Brandl SJ (2015). Transplantation of corals into a new environment results in substantial skeletal loss in *Acropora tenuis*. *Mar Biodiv* (doi: 10.1007/s12526-014-0239-y).
5. Heinrich DDU, Watson SA, Rummer JL, Brandl SJ, Simpfendorfer CA, Heupel MR, Munday PL (2015) Foraging behaviour of the epaulette shark *Hemiscyllium ocellatum* is not affected by elevated CO₂. ICES (in press, doi: 10.1093/icesjms/fsv085).
6. Brooker RM, Brandl SJ, Dixson DL. Cryptic effects of habitat declines: coral-associated fishes avoid coral-seaweed interactions due to visual and chemical cues. *Sci Rep* 6, 18842 (doi:10.1038/srep18842).
7. Brandl SJ, Emslie MJ, Ceccarelli DM. Functional redundancy buffers trait-specific susceptibility to disturbance in a high-diversity system (under review).

8. Mirbach CE, Brandl SJ. The ontogeny of pairing in coral reef rabbitfishes (f. Siganidae) (under review).
9. Casey JM, Baird AH, Brandl SJ, Hoogenboom MO, Rizzari JR, Mirbach CE, Frisch AJ, Connolly SR. A test of trophic cascade theory: fish and benthic assemblages across a predator density gradient on coral reefs (under review).

Appendix C

Supplemental table for **Chapter 2: Pair-formation in coral reef fishes: an ecological perspective**

List of Indo-Pacific reef fishes and their social system, trophic affiliation, burrowing behaviour, activity patterns, size and reproductive traits. Troph. = trophic group; Ca = carnivore; Co = corallivore; De = detritivore; He = herbivore; Ma = macroinvertevore; Mi = microinvertevore; Om = omnivore; Pl = planktivore; Pi = Piscivore; Sp = spongivore; Burr. = burrowing; Act. = activity period; D = diurnal; N = nocturnal; Spa. = spawning mode; G = group; P = pair; B = both; Gam. = gamete release; Br = broadcast; Dm = demersal; Gel = gelatinous egg mass; Es = egg-scattering; Mo = mouthbrooding; Po = pouchbrooding; Mon. = monogamy; Refs. = references. N/A = no available information. *Asterisks* mark burrowing species that associate with alpheid shrimps. References are listed numerically at the end of the table.

Family	Species	Pairs	Troph.	Burr.	Act.	Size (mm)	Spa.	Gam.	Mon.	Refs.
Acanthuridae	<i>Acanthurus achilles</i>	Yes	He	No	D	200	G	Br	Yes	5
	<i>Acanthurus albipectoralis</i>	No	Pl	No	D	330	G	Br	No	6
	<i>Acanthurus auranticavus</i>	No	De	No	D	350	G	Br	No	6
	<i>Acanthurus bariene</i>	Yes	De	No	D	500	G	Br	No	6
	<i>Acanthurus blochii</i>	No	De	No	D	420	G	Br	No	6
	<i>Acanthurus dussumieri</i>	No	De	No	D	500	G	Br	No	7
	<i>Acanthurus fowleri</i>	Yes	He	No	D	450	G	Br	No	2
	<i>Acanthurus grammoptilus</i>	No	De	No	D	350	G	Br	No	6
	<i>Acanthurus guttatus</i>	No	He	No	D	280	B	Br	No	1;3
	<i>Acanthurus japonicus</i>	No	He	No	D	210	G	Br	No	2
	<i>Acanthurus leucocheilus</i>	No	De	No	D	200	G	Br	No	2
	<i>Acanthurus leucopareius</i>	No	He	No	D	200	G	Br	No	1;3
	<i>Acanthurus leucosternon</i>	Yes	He	No	D	380	G	Br	Yes	8
	<i>Acanthurus lineatus</i>	No	He	No	D	380	G	Br	No	6
	<i>Acanthurus maculiceps</i>	No	He	No	D	200	G	Br	No	6
	<i>Acanthurus mata</i>	No	Pl	No	D	500	G	Br	No	1;2;3
	<i>Acanthurus nigricans</i>	Yes	He	No	D	210	G	Br	Yes	5

<i>Acanthurus nigricauda</i>	No	De	No	D	400	G	Br	No	7;6	
<i>Acanthurus nigrofuscus</i>	No	He	No	D	210	B	Br	No	8;9	
<i>Acanthurus nigroris</i>	No	He	No	D	250	G	Br	No	1;2;3	
<i>Acanthurus nubilus</i>	No	Pl	No	D	450	G	Br	No	2;3;4	
<i>Acanthurus olivaceus</i>	No	De	No	D	350	G	Br	No	7	
<i>Acanthurus pyroferus</i>	No	He	No	D	250	G	Br	No	6	
<i>Acanthurus reversus</i>	No	De	No	D	340	G	Br	No	3	
<i>Acanthurus tennenti</i>	Yes	He	No	D	310	G	Br	No	8	
<i>Acanthurus thompsoni</i>	No	Pl	No	D	270	G	Br	No	1;2;3	
<i>Acanthurus triostegus</i>	Yes	He	No	D	260	G	Br	No	8;6	
<i>Acanthurus tristis</i>	No	He	No	D	250	G	Br	No	2	
<i>Acanthurus xanthopterus</i>	No	De	No	D	560	G	Br	No	1;2;3	
<i>Ctenochaetus binotatus</i>	No	De	No	D	220	G	Br	No	2;3;4	
<i>Ctenochaetus cyanocheilus</i>	No	He	No	D	200	G	Br	No	2;3	
<i>Ctenochaetus flavicauda</i>	No	De	No	D	130	G	Br	No	2	
<i>Ctenochaetus hawaiiensis</i>	No	De	No	D	250	G	Br	No	2;3	
<i>Ctenochaetus marginatus</i>	No	De	No	D	220	G	Br	No	2;3	
<i>Ctenochaetus striatus</i>	No	De	No	D	260	B	Br	No	8;6	
<i>Ctenochaetus strigosus</i>	No	De	No	D	180	P	Br	No	8;1	
<i>Ctenochaetus tomiensis</i>	No	De	No	D	150	G	Br	No	2;3	
<i>Naso annulatus</i>	No	Pl	No	D	1000	G	Br	No	1;2;3	
<i>Naso brachycentron</i>	No	He	No	D	600	G	Br	No	1;2;3;4	
<i>Naso brevirostris</i>	No	Pl	No	D	500	G	Br	No	1;2;3	
<i>Naso caeruleacauda</i>	No	Pl	No	D	300	G	Br	No	2;3	
<i>Naso caesioides</i>	No	Pl	No	D	600	G	Br	No	1;2;3	
<i>Naso hexacanthus</i>	No	Pl	No	D	750	G	Br	No	1;2;3	
<i>Naso lituratus</i>	Yes	He	No	D	300	G	Br	No	8;6;9	
<i>Naso lopezi</i>	No	Pl	No	D	650	G	Br	No	1;2;3	
<i>Naso maculatus</i>	No	Pl	No	D	600	G	Br	No	3;10	
<i>Naso minor</i>	No	Pl	No	D	190	G	Br	No	2;4	
<i>Naso thynnoides</i>	No	Pl	No	D	300	G	Br	No	1;2;3	
<i>Naso tonganus</i>	No	He	No	D	600	G	Br	No	2;3;4	
<i>Naso tuberosus</i>	No	He	No	D	600	G	Br	No	1;2;3;6	
<i>Naso unicornis</i>	No	He	No	D	700	G	Br	No	8;1;2;3;6	
<i>Naso vlamingii</i>	No	Pl	No	D	500	G	Br	No	1;2;3	
<i>Paracanthurus hepatus</i>	No	Pl	No	D	310	G	Br	No	1;2;3;6	
<i>Prionurus maculatus</i>	No	He	No	D	435	G	Br	No	1;3	
<i>Prionurus microlepidotus</i>	No	He	No	D	700	P	Br	No	1	
<i>Zebrasoma desjardini</i>	Yes	He	No	D	400	P	Br	No	2;4	
<i>Zebrasoma flavescens</i>	No	He	No	D	200	B	Br	No	2;4	
<i>Zebrasoma rostratum</i>	No	He	No	D	210	P	Br	No	2;3	
<i>Zebrasoma scopas</i>	Yes	He	No	D	200	B	Br	Yes	8;6;9	
<i>Zebrasoma veliferum</i>	Yes	He	No	D	400	P	Br	No	8;6;9	
Anomalopidae	<i>Anomalops katoptron</i>	No	Pl	No	N	350	P	Br	No	4;11
	<i>Photoblepharon palpebratum</i>	No	Pl	No	N	120	P	Br	No	4
Antennariidae	<i>Antennarius biocellatus</i>	No	Pi	No	D	150	P	Gel	No	2;4
	<i>Antennarius coccineus</i>	No	Pi	No	D	120	P	Gel	No	1;2;3;12
	<i>Antennarius commersoni</i>	No	Pi	No	D	300	P	Gel	No	1;2;3;4
	<i>Antennarius dorehensis</i>	No	Pi	No	D	50	P	Gel	No	2
	<i>Antennarius maculatus</i>	No	Pi	No	D	90	P	Gel	No	1;2;3;13
	<i>Antennarius nummifer</i>	No	Pi	No	D	100	P	Gel	No	1;2;3
	<i>Antennarius pictus</i>	No	Pi	No	D	160	P	Gel	No	1;2;3
	<i>Antennarius randalli</i>	No	Pi	No	D	45	P	Gel	No	1;2;3

	<i>Antennarius striatus</i>	No	Pi	No	D	220	P	Gel	No	1;2;3
	<i>Antennatus rosaceus</i>	No	Pi	No	D	58	P	Gel	No	3
	<i>Antennatus tuberosus</i>	No	Pi	No	D	80	P	Gel	No	2
	<i>Histiophryne cryptacanthus</i>	No	Pi	No	D	85	P	Gel	No	1;2;3
	<i>Histrio histrio</i>	No	Ca	No	D	140	P	Gel	No	1;2;3;4
	<i>Lophiocharon trisignatus</i>	No	Pi	No	D	180	P	Gel	No	1;2;4
	<i>Tathicarpus butleri</i>	No	Pi	No	D	120	P	Gel	No	2
Apistidae	<i>Apistus carinatus</i>	No	Pl	No	D	88	N/A	N/A	No	2;14
Aploactinidae	<i>Paraploactis kagoshimensis</i>	No	N/A	No	D	120	N/A	N/A	No	2
Apogonidae	<i>Apogon amboinensis</i>	No	Ca	No	D	100	P	Mo	No	2;15
	<i>Apogon exostigma</i>	No	Ca	No	N	110	P	Mo	No	1;2;16
	<i>Apogon guamensis</i>	Yes	Pl	No	N	110	P	Mo	No	1;2;3;16
	<i>Apogon hyalosoma</i>	No	Ca	No	N	200	P	Mo	No	2;17
	<i>Apogon norfolcensis</i>	No	Ca	No	N	100	P	Mo	No	2;3;4
	<i>Apogon sangiensis</i>	No	Ca	No	N	80	P	Mo	No	1;4
	<i>Apogon semiornatus</i>	No	Ca	No	N	70	P	Mo	No	1;3;4
	<i>Apogonichthyoides melas</i>	No	Ca	No	D	130	P	Mo	No	2;4
	<i>Apogonichthyoides timorensis</i>	No	Ca	No	N	80	P	Mo	No	2;4
	<i>Apogonichthyoides unnotatus</i>	No	Ca	No	D	60	P	Mo	No	2
	<i>Apogonichthys ocellatus</i>	No	Ca	No	N	35	P	Mo	No	2;4
	<i>Archamia biguttata</i>	No	Pl	No	N	90	P	Mo	No	1;2;3;12
	<i>Archamia bleekeri</i>	No	Pl	No	N	90	P	Mo	No	2;4
	<i>Archamia fucata</i>	No	Pl	No	N	90	P	Mo	No	1;2;3;18
	<i>Archamia leai</i>	No	Pl	No	N	80	P	Mo	No	1;2;3;16
	<i>Archamia macroptera</i>	No	Pl	No	D	90	P	Mo	No	2;3;
	<i>Archamia zosterophora</i>	No	Pl	No	N	80	P	Mo	No	1;2;4
	<i>Cercamia cladara</i>	No	N/A	No	N	40	P	Mo	No	3;4
	<i>Cercamia eremia</i>	No	N/A	No	N	40	P	Mo	No	2
	<i>Cheilodipterus alleni</i>	No	N/A	No	N	40	P	Mo	No	2;4
	<i>Cheilodipterus artus</i>	No	Ca	No	D	120	P	Mo	No	1;2;3;18
	<i>Cheilodipterus intermedius</i>	No	N/A	No	D	110	P	Mo	No	2;4
	<i>Cheilodipterus isostigmus</i>	No	Pi	No	D	100	P	Mo	No	2;3;4
	<i>Cheilodipterus macrodon</i>	No	Pi	No	D	200	P	Mo	Yes	1;2;3;18
	<i>Cheilodipterus nigrotaeniatus</i>	Yes	Ma	No	D	80	P	Mo	No	2;12
	<i>Cheilodipterus parazonatus</i>	Yes	Ma	No	D	70	P	Mo	No	1;2;12
	<i>Cheilodipterus quinquelineatus</i>	No	Ca	No	N	120	P	Mo	No	1;2;3;18
	<i>Cheilodipterus singapurensis</i>	No	N/A	No	N	175	P	Mo	No	2
	<i>Cheilodipterus zonatus</i>	Yes	Ma	No	D	70	P	Mo	No	2;12
	<i>Foa brachygramma</i>	No	Pl	No	D	60	P	Mo	No	1;4
	<i>Foa fo</i>	No	Pl	No	D	35	P	Mo	No	1;2;3
	<i>Foa hyalina</i>	No	Pl	No	D	50	P	Mo	No	1;2;3
	<i>Fowleria marmorata</i>	No	N/A	No	D	45	P	Mo	No	1;2;3
	<i>Fowleria punctulata</i>	No	N/A	No	N	60	P	Mo	No	1;4
	<i>Fowleria vaiulae</i>	No	N/A	No	D	50	P	Mo	No	1;2;3
	<i>Fowleria variegata</i>	No	N/A	No	D	50	P	Mo	No	1;2;3
	<i>Nectamia bandanensis</i>	No	Pl	No	N	90	P	Mo	No	2;3;4
	<i>Nectamia fusca</i>	No	Pl	No	N	100	P	Mo	No	2;3;4
	<i>Nectamia octospina</i>	No	Pl	No	D	50	P	Mo	No	1;2;3
	<i>Nectamia savayensis</i>	No	Ma	No	D	110	P	Mo	No	1;4
	<i>Ostorhinchus angustatus</i>	No	Mi	No	N	90	P	Mo	No	1;2;3;4
	<i>Ostorhinchus apogonides</i>	No	Pl	No	D	100	P	Mo	No	1;2;3;12
	<i>Ostorhinchus aureus</i>	No	N/A	No	D	120	P	Mo	No	1;2;3
	<i>Ostorhinchus capricornis</i>	Yes	N/A	No	D	100	P	Mo	No	2

<i>Ostorhinchus cavitiensis</i>	No	N/A	No	D	80	P	Mo	No	2
<i>Ostorhinchus chrysopomus</i>	No	N/A	No	D	90	P	Mo	No	2
<i>Ostorhinchus chrysotaenia</i>	No	N/A	No	N	100	P	Mo	No	2;4
<i>Ostorhinchus compressus</i>	No	Pl	No	N	120	P	Mo	No	1;2;3;19
<i>Ostorhinchus cookii</i>	No	N/A	No	N	100	P	Mo	No	1;2;3;4
<i>Ostorhinchus crassiceps</i>	No	Ma	No	N	60	P	Mo	No	1;2;3;4
<i>Ostorhinchus cyanosoma</i>	Yes	Ma	No	D	80	P	Mo	No	1;2;3;16
<i>Ostorhinchus dispar</i>	No	N/A	No	N	50	P	Mo	No	2;4
<i>Ostorhinchus doederleini</i>	Yes	Ca	No	N	120	P	Mo	No	1;2;3;16
<i>Ostorhinchus endekataenia</i>	No	N/A	No	N	140	P	Mo	No	2;4
<i>Ostorhinchus evermanni</i>	Yes	N/A	No	N	120	P	Mo	No	2;4
<i>Ostorhinchus fleurieu</i>	No	N/A	No	D	110	P	Mo	No	2
<i>Ostorhinchus franssedai</i>	No	N/A	No	D	75	P	Mo	No	2
<i>Ostorhinchus griffini</i>	Yes	N/A	No	D	140	P	Mo	No	2;4
<i>Ostorhinchus hartzfeldii</i>	No	Ma	No	N	120	P	Mo	No	2;20
<i>Ostorhinchus hoevenii</i>	No	Ma	No	D	50	P	Mo	No	2;20
<i>Ostorhinchus holotaenia</i>	No	N/A	No	D	80	P	Mo	No	1;2;3
<i>Ostorhinchus kiensis</i>	No	N/A	No	D	90	P	Mo	No	2;3
<i>Ostorhinchus komodoensis</i>	No	Pl	No	D	70	P	Mo	No	2;4
<i>Ostorhinchus lateralis</i>	No	Ma	No	N	80	P	Mo	No	2;21
<i>Ostorhinchus lineomaculatus</i>	No	N/A	No	N	65	P	Mo	No	2
<i>Ostorhinchus luteus</i>	No	Pl	No	D	50	P	Mo	No	1;2;3;12
<i>Ostorhinchus margaritophorus</i>	No	N/A	No	N	55	P	Mo	No	2;22
<i>Ostorhinchus moluccensis</i>	No	N/A	No	N	90	P	Mo	No	2
<i>Ostorhinchus monospilus</i>	Yes	N/A	No	N	80	P	Mo	No	2;4
<i>Ostorhinchus multilineatus</i>	No	N/A	No	D	100	P	Mo	No	2;4
<i>Ostorhinchus nanus</i>	No	N/A	No	D	35	P	Mo	No	2;4
<i>Ostorhinchus neotes</i>	No	N/A	No	D	30	P	Mo	No	2
<i>Ostorhinchus nigrofasciatus</i>	Yes	Mi	No	N	80	P	Mo	No	1;2;3;4
<i>Ostorhinchus notatus</i>	No	Pl	No	N	100	P	Mo	Yes	1;2;23
<i>Ostorhinchus novemfasciatus</i>	Yes	Ca	No	N	90	P	Mo	No	1;2;3;4
<i>Ostorhinchus ocellicaudus</i>	Yes	N/A	No	D	60	P	Mo	No	2
<i>Ostorhinchus parvulus</i>	No	N/A	No	D	80	P	Mo	No	2;4
<i>Ostorhinchus properupta</i>	Yes	N/A	No	D	75	P	Mo	No	2
<i>Ostorhinchus quadrifasciatus</i>	No	Ma	No	N	100	P	Mo	No	2;24
<i>Ostorhinchus rubrimacula</i>	No	Ma	No	D	80	P	Mo	No	2;3;25
<i>Ostorhinchus ruepellii</i>	No	N/A	No	N	120	P	Mo	No	2;4
<i>Ostorhinchus sealei</i>	No	N/A	No	D	90	P	Mo	No	2
<i>Ostorhinchus selas</i>	No	N/A	No	D	55	P	Mo	No	2
<i>Ostorhinchus semiornatus</i>	No	N/A	No	N	70	P	Mo	No	2;3
<i>Ostorhinchus taeniophorus</i>	N/A	N/A	No	N	100	P	Mo	No	1;2;3;4
<i>Ostorhinchus talboti</i>	No	N/A	No	N	100	P	Mo	No	1;2;3
<i>Ostorhinchus thermalis</i>	No	N/A	No	D	80	P	Mo	No	2
<i>Ostorhinchus wassinki</i>	Yes	N/A	No	D	70	P	Mo	No	2
<i>Pristiapogon fraenatus</i>	No	N/A	No	N	110	P	Mo	No	1;2;4
<i>Pristiapogon kallopterus</i>	No	Pl	No	N	150	P	Mo	No	2;12
<i>Pristicon rhodopterus</i>	No	N/A	No	N	150	P	Mo	No	2
<i>Pristicon trimaculatus</i>	Yes	Pl	No	N	150	P	Mo	No	2;3;12
<i>Pseudamia gelatinosa</i>	No	N/A	No	N	100	P	Mo	No	1;2;3
<i>Pseudamia hayashii</i>	No	N/A	No	N	75	P	Mo	No	2
<i>Pseudamia zonata</i>	No	N/A	No	N	90	P	Mo	No	2;3
<i>Pseudamiops gracilicauda</i>	No	Pl	No	N	50	P	Mo	No	3
<i>Pseudamiops phasma</i>	N/A	N/A	No	D	47	P	Mo	No	3

	<i>Pterapogon kauderni</i>	No	Pl	No	N	65	P	Mo	No	2;4
	<i>Pterapogon mirifica</i>	No	N/A	No	N	140	P	Mo	No	2;4
	<i>Rhabdamia cypselurus</i>	No	Pl	No	N	60	P	Mo	No	1;2;3;4
	<i>Rhabdamia gracilis</i>	No	Pl	No	N	60	P	Mo	No	1;2;3;4
	<i>Rhabdamia spilota</i>	No	Pl	No	D	60	P	Mo	No	2;4
	<i>Siphamia corallicola</i>	No	N/A	No	D	38	P	Mo	No	2;4
	<i>Siphamia elongata</i>	No	N/A	No	D	38	P	Mo	No	2
	<i>Siphamia fuscolineata</i>	No	N/A	No	D	35	P	Mo	No	2
	<i>Siphamia majimai</i>	N/A	N/A	No	N	35	P	Mo	No	1;4
	<i>Siphamia tubifer</i>	No	N/A	No	D	40	P	Mo	No	2
	<i>Siphamia versicolor</i>	No	N/A	No	D	40	P	Mo	No	2;3
	<i>Sphaeramia nematoptera</i>	No	Pl	No	N	80	P	Mo	No	1;2;3;26
	<i>Sphaeramia orbicularis</i>	No	Pl	No	N	115	P	Mo	No	2;3;4
	<i>Zoramia fragilis</i>	Yes	Mi	No	N	55	P	Mo	No	1;2;3;18
	<i>Zoramia gilberti</i>	No	N/A	No	D	55	P	Mo	No	2;4
	<i>Zoramia leptacanthus</i>	No	N/A	No	N	60	P	Mo	No	2;3;4
	<i>Zoramia perlita</i>	No	N/A	No	N	55	P	Mo	No	2;4
Aulostomidae	<i>Aulostomus chinensis</i>	No	Ca	No	D	800	P	Br	No	4
Balistidae	<i>Abalistes stellatus</i>	No	N/A	No	D	600	P	Dm	No	1;2;3
	<i>Balistapus undulatus</i>	No	Om	No	D	300	P	Dm	No	1;2;3
	<i>Balistes polylepis</i>	No	Ma	No	D	760	G	Dm	No	4
	<i>Balistoides conspicillum</i>	No	Ma	No	D	500	G	Dm	No	4
	<i>Balistoides viridescens</i>	No	Ma	No	D	750	G	Dm	No	1;2;3
	<i>Canthidermis maculata</i>	No	Pl	No	D	300	P	Dm	No	12
	<i>Melichthys indicus</i>	No	Om	No	D	240	P	Dm	No	4
	<i>Melichthys niger</i>	No	Om	No	D	350	P	Dm	No	1;2;3
	<i>Melichthys vidua</i>	No	Om	No	D	300	P	Dm	No	1;2;3
	<i>Odonus niger</i>	No	Pl	No	D	400	P	Dm	No	1;2;3
	<i>Pseudobalistes flavimarginatus</i>	No	Ma	No	D	600	G	Dm	No	4
	<i>Pseudobalistes fuscus</i>	No	Ma	No	D	550	G	Dm	Yes	1;2;3
	<i>Rhinecanthus aculeatus</i>	No	Om	No	D	250	P	Dm	No	27
	<i>Rhinecanthus lunula</i>	No	Om	No	D	280	P	Dm	No	1;2;3
	<i>Rhinecanthus rectangulus</i>	No	Om	No	D	250	P	Dm	No	4
	<i>Rhinecanthus verrucosus</i>	No	Om	No	D	230	P	Dm	No	28
	<i>Sufflamen bursa</i>	No	Om	No	D	240	P	Dm	No	4
	<i>Sufflamen chrysopterum</i>	No	Ma	No	D	220	P	Dm	Yes	4
	<i>Sufflamen fraenatum</i>	No	Om	No	D	380	P	Dm	No	4
	<i>Xanthichthys auromarginatus</i>	No	Pl	No	D	220	P	Dm	No	12
	<i>Xanthichthys caeruleolineatus</i>	No	Pl	No	D	350	P	Dm	No	12
	<i>Xanthichthys mento</i>	No	Pl	No	D	220	P	Dm	No	12
Batrachoididae	<i>Halophyrne diemensis</i>	No	Ca	No	D	260	N/A	N/A	No	1;2;3;4
Belonidae	<i>Platybelone argalus</i>	No	Pi	No	D	370	N/A	Es	No	1;2;3
	<i>Strongylura incisa</i>	No	Pi	No	D	700	N/A	Es	No	1;2;3
	<i>Tylosurus acus</i>	No	Pi	No	D	1000	N/A	Es	No	3
	<i>Tylosurus crocodilus</i>	No	Pi	No	D	1500	N/A	Es	No	1;2;4
Blenniidae	<i>Andamia tetradactylus</i>	N/A	N/A	No	D	65	P	Dm	No	2
	<i>Aspidontus dussumieri</i>	No	Om	No	D	120	P	Dm	No	3
	<i>Aspidontus taeniatus</i>	No	Ca	No	D	115	P	Dm	No	1;2;3;4
	<i>Atrosalaria fuscus</i>	No	De	No	D	145	P	Dm	No	1;2;3;29
	<i>Blenniella caudolineata</i>	N/A	N/A	No	D	100	P	Dm	No	3
	<i>Blenniella chrysoopilus</i>	No	Om	No	D	140	P	Dm	No	1;2;3;4
	<i>Blenniella gibbifrons</i>	No	N/A	No	D	100	P	Dm	No	2;3
	<i>Blenniella interrupta</i>	No	N/A	No	D	80	P	Dm	No	2

<i>Blenniella paula</i>	No	Om	No	D	130	P	Dm	No	1;2;3;4
<i>Cirripectes auritus</i>	No	Om	No	D	90	P	Dm	No	2;3
<i>Cirripectes chelomatus</i>	No	De	No	D	120	P	Dm	No	1;2;3;29
<i>Cirripectes filamentosus</i>	No	De	No	D	90	P	Dm	No	1;2;3
<i>Cirripectes polyzona</i>	No	De	No	D	85	P	Dm	No	1;2;3;4
<i>Cirripectes springeri</i>	No	Om	No	D	80	P	Dm	No	2
<i>Cirripectes stigmaticus</i>	No	De	No	D	130	P	Dm	No	2
<i>Cirripectes variolosus</i>	No	De	No	D	80	P	Dm	No	2
<i>Cirripectus castaneus</i>	Yes	Om	No	D	125	P	Dm	No	1;2;3;4
<i>Crossosalarias macrospilus</i>	No	N/A	No	D	85	P	Dm	No	1;2;3
<i>Ecsenius alleni</i>	No	De	No	D	85	P	Dm	No	2
<i>Ecsenius australianus</i>	No	De	No	D	50	P	Dm	No	1;2
<i>Ecsenius axelrodi</i>	No	De	No	D	50	P	Dm	No	2
<i>Ecsenius bathi</i>	No	De	No	D	40	P	Dm	No	2
<i>Ecsenius bicolor</i>	No	De	No	D	100	P	Dm	No	1;2;3;29
<i>Ecsenius bimaculatus</i>	No	De	No	D	50	P	Dm	No	2;3
<i>Ecsenius caeruliventris</i>	No	De	No	D	30	P	Dm	No	2
<i>Ecsenius collettei</i>	No	De	No	D	40	P	Dm	No	2
<i>Ecsenius dilemma</i>	No	De	No	D	50	P	Dm	No	2
<i>Ecsenius fijiensis</i>	No	De	No	D	40	P	Dm	No	2;3
<i>Ecsenius fourmanoiri</i>	N/A	De	No	D	62	P	Dm	No	3
<i>Ecsenius isos</i>	N/A	De	No	D	40	P	Dm	No	3
<i>Ecsenius kurti</i>	No	De	No	D	45	P	Dm	No	2
<i>Ecsenius lineatus</i>	No	De	No	D	70	P	Dm	No	2;30
<i>Ecsenius lividanalisis</i>	No	De	No	D	50	P	Dm	No	2
<i>Ecsenius lubbocki</i>	Yes	De	No	D	40	P	Dm	No	2
<i>Ecsenius mandibularis</i>	No	De	No	D	72	P	Dm	No	29;4
<i>Ecsenius melarchus</i>	No	De	No	D	60	P	Dm	No	2;3
<i>Ecsenius midas</i>	No	Pl	No	D	130	P	Dm	No	1;2;3;4
<i>Ecsenius monoculus</i>	No	De	No	D	60	P	Dm	No	2
<i>Ecsenius namiyei</i>	No	De	No	D	100	P	Dm	No	2;30
<i>Ecsenius oculus</i>	No	De	No	D	60	P	Dm	No	2
<i>Ecsenius ops</i>	No	De	No	D	55	P	Dm	No	2
<i>Ecsenius pardus</i>	N/A	De	No	D	62	P	Dm	No	3
<i>Ecsenius pictus</i>	Yes	De	No	D	50	P	Dm	No	2
<i>Ecsenius portenoyi</i>	N/A	De	No	D	58	P	Dm	No	3
<i>Ecsenius prooculis</i>	No	De	No	D	50	P	Dm	No	2
<i>Ecsenius schroederi</i>	No	De	No	D	50	P	Dm	No	2
<i>Ecsenius sellifer</i>	No	De	No	D	45	P	Dm	No	2
<i>Ecsenius shirleyae</i>	No	De	No	D	40	P	Dm	No	2
<i>Ecsenius stictus</i>	No	De	No	D	55	P	Dm	No	2;29
<i>Ecsenius stigmatura</i>	No	De	No	D	55	P	Dm	No	2
<i>Ecsenius taeniatus</i>	No	De	No	D	40	P	Dm	No	2
<i>Ecsenius tessera</i>	N/A	De	No	D	40	P	Dm	No	3
<i>Ecsenius tigris</i>	No	De	No	D	40	P	Dm	No	1;2
<i>Ecsenius tricolor</i>	No	De	No	D	50	P	Dm	No	2;3
<i>Ecsenius trilineatus</i>	No	De	No	D	35	P	Dm	No	2;3
<i>Ecsenius yaeyamaensis</i>	No	De	No	D	60	P	Dm	No	2
<i>Enchelyurus ater</i>	N/A	N/A	No	D	53	P	Dm	No	3
<i>Enchelyurus kraussii</i>	N/A	N/A	No	D	50	P	Dm	No	3
<i>Entomacrodus caudofasciatus</i>	N/A	N/A	No	D	66	P	Dm	No	3
<i>Entomacrodus corneliae</i>	N/A	N/A	No	D	50	P	Dm	No	3
<i>Entomacrodus decussatus</i>	N/A	N/A	No	D	190	P	Dm	No	3

<i>Entomacrodus epalzeocheilos</i>	N/A	N/A	No	D	128	P	Dm	No	3
<i>Entomacrodus macrospilus</i>	N/A	N/A	No	D	42	P	Dm	No	3
<i>Entomacrodus randalli</i>	N/A	N/A	No	D	109	P	Dm	No	3
<i>Entomacrodus sealei</i>	N/A	N/A	No	D	109	P	Dm	No	3
<i>Entomacrodus striatus</i>	N/A	N/A	No	D	118	P	Dm	No	3
<i>Entomacrodus thalassinus</i>	N/A	N/A	No	D	60	P	Dm	No	3
<i>Exallias brevis</i>	No	Co	No	D	145	P	Dm	No	1;2;3;31
<i>Glyptoparus delicatulus</i>	Yes	Om	No	D	50	P	Dm	No	1;2;3;29
<i>Istiblennius bellus</i>	No	N/A	No	D	150	P	Dm	No	2
<i>Istiblennius dussumieri</i>	No	N/A	No	D	125	P	Dm	No	2
<i>Istiblennius edentulus</i>	No	He	No	D	170	P	Dm	No	1;2;3;32
<i>Istiblennius lineatus</i>	No	He	No	D	140	P	Dm	No	2
<i>Meiacanthus abditus</i>	No	Pl	No	D	65	P	Dm	No	2
<i>Meiacanthus anema</i>	Yes	N/A	No	D	65	P	Dm	No	2;4
<i>Meiacanthus atrodorsalis</i>	Yes	Mi	No	D	110	P	Dm	No	1;2;3
<i>Meiacanthus bundoon</i>	Yes	Pl	No	D	80	P	Dm	No	2;3
<i>Meiacanthus crinitus</i>	Yes	N/A	No	D	65	P	Dm	No	2
<i>Meiacanthus ditrema</i>	No	Pl	No	D	65	P	Dm	No	1;2;3;4
<i>Meiacanthus geminatus</i>	No	N/A	No	D	65	P	Dm	No	2
<i>Meiacanthus grammistes</i>	Yes	N/A	No	D	100	P	Dm	No	1;2;3
<i>Meiacanthus kamoharai</i>	Yes	Om	No	D	85	P	Dm	No	2;4
<i>Meiacanthus lineatus</i>	Yes	N/A	No	D	95	P	Dm	No	1;2
<i>Meiacanthus oualanensis</i>	No	N/A	No	D	100	P	Dm	No	2
<i>Meiacanthus smithi</i>	Yes	Mi	No	D	80	P	Dm	No	2;4
<i>Meiacanthus urostigma</i>	No	N/A	No	D	55	P	Dm	No	2
<i>Meiacanthus vicinus</i>	Yes	N/A	No	D	65	P	Dm	No	2
<i>Meiacanthus vittatus</i>	No	N/A	No	D	65	P	Dm	No	2
<i>Omobranchus elongatus</i>	No	N/A	No	D	55	P	Dm	No	2
<i>Omobranchus germaini</i>	No	Om	No	D	78	P	Dm	No	3
<i>Omobranchus obliquus</i>	No	N/A	No	D	70	P	Dm	No	2
<i>Paralticus amboinensis</i>	No	N/A	No	D	150	P	Dm	No	2
<i>Petroscirtes breviceps</i>	Yes	Om	No	D	130	P	Dm	No	1;2;33
<i>Petroscirtes fallax</i>	Yes	Om	No	D	95	P	Dm	No	1;2
<i>Petroscirtes lupus</i>	No	He	No	D	130	P	Dm	No	3;34
<i>Petroscirtes mitratus</i>	No	Om	No	D	150	P	Dm	No	1;2;3;26
<i>Petroscirtes variabilis</i>	No	Om	No	D	75	P	Dm	No	1;2;3;26
<i>Petroscirtes xestus</i>	No	N/A	No	D	95	P	Dm	No	2
<i>Plagiotremus flavus</i>	No	Ca	No	D	70	P	Dm	No	2
<i>Plagiotremus laudandus</i>	No	Ca	No	D	130	P	Dm	No	1;2;3;4
<i>Plagiotremus rhinorhynchus</i>	No	Ca	No	D	130	P	Dm	No	1;2;3;4
<i>Plagiotremus tapeinosoma</i>	No	Ca	No	D	130	P	Dm	No	1;2;3;4
<i>Salarias alboguttatus</i>	No	N/A	No	D	65	P	Dm	No	1;2;3
<i>Salarias ceramensis</i>	No	He	No	D	140	P	Dm	No	2;34
<i>Salarias fasciatus</i>	No	De	No	D	140	P	Dm	No	1;2;3;29
<i>Salarias guttatus</i>	No	De	No	D	140	P	Dm	No	2;29
<i>Salarias nigrocinctus</i>	No	De	No	D	53	P	Dm	No	3
<i>Salarias obscurus</i>	No	De	No	D	130	P	Dm	No	2
<i>Salarias patzneri</i>	No	De	No	D	50	P	Dm	No	2;29
<i>Salarias ramosus</i>	No	De	No	D	50	P	Dm	No	2
<i>Salarias segmentatus</i>	No	He	No	D	75	P	Dm	No	2;4
<i>Salarias sinuosus</i>	No	De	No	D	60	P	Dm	No	1;2;3
<i>Stanulus seychellensis</i>	Yes	N/A	No	D	33	P	Dm	No	2
<i>Stanulus talboti</i>	No	N/A	No	D	48	P	Dm	No	3

Appendix C

	<i>Xiphasia matsubarai</i>	No	Ca	No	N	300	P	Dm	No	3;4
	<i>Xiphasia setifer</i>	No	Ca	No	N	530	P	Dm	No	2;3;4
Bothidae	<i>Asterorhombus fijiensis</i>	No	Ca	No	D	150	P	Br	No	2
	<i>Asterorhombus filifer</i>	No	Ca	No	D	130	P	Br	No	3
	<i>Asterorhombus intermedius</i>	No	Ca	No	D	160	P	Br	No	4
	<i>Bothus mancus</i>	No	Pi	No	N	420	P	Br	No	1;2;3;4
	<i>Bothus pantherinus</i>	No	Ca	No	D	390	P	Br	No	4
	<i>Engyprosopon grandisquama</i>	No	Ma	No	D	110	P	Br	No	4
Caesionidae	<i>Caesio caeruleaurea</i>	No	Pl	No	D	400	G	Br	No	2;4
	<i>Caesio cuning</i>	No	Pl	No	D	500	G	Br	No	1;2;3;35;12
	<i>Caesio lunaris</i>	No	Pl	No	D	400	G	Br	No	2;4
	<i>Caesio teres</i>	No	Pl	No	D	400	G	Br	No	1;2;3;12
	<i>Caesio varilineata</i>	No	Pl	No	D	400	G	Br	No	2;4
	<i>Caesio xanthonota</i>	No	Pl	No	D	400	G	Br	No	2;4
	<i>Dipterygnotus balteatus</i>	No	Pl	No	D	140	G	Br	No	2;3;4
	<i>Gymnocaesio gymnoptera</i>	No	Pl	No	D	180	G	Br	No	2;3;4
	<i>Pterocaesio chrysozona</i>	No	Pl	No	D	210	G	Br	No	2;4
	<i>Pterocaesio digramma</i>	No	Pl	No	D	210	G	Br	No	2;3;35
	<i>Pterocaesio lativittata</i>	No	Pl	No	D	200	G	Br	No	2;4
	<i>Pterocaesio marri</i>	No	Pl	No	D	350	G	Br	No	1;2;3;12
	<i>Pterocaesio pisang</i>	No	Pl	No	D	210	G	Br	No	2;3;4
	<i>Pterocaesio randalli</i>	No	Pl	No	D	250	G	Br	No	2;4
	<i>Pterocaesio tessellata</i>	No	Pl	No	D	250	G	Br	No	2;4
	<i>Pterocaesio tile</i>	No	Pl	No	D	250	G	Br	No	1;2;3;12
	<i>Pterocaesio trilineata</i>	No	Pl	No	D	200	G	Br	No	1;2;3;12
Callionymidae	<i>Anaora tentaculata</i>	No	Mi	No	D	45	P	Br	No	2;26
	<i>Callionymus enneactis</i>	Yes	Mi	No	D	45	P	Br	No	2;12
	<i>Callionymus filamentosus</i>	No	Mi	No	D	165	P	Br	No	3;4
	<i>Callionymus keeleyi</i>	No	Mi	No	D	60	P	Br	No	2;12
	<i>Callionymus marquesensis</i>	No	Mi	No	D	55	P	Br	No	3
	<i>Callionymus simplicicornis</i>	No	Mi	No	D	60	P	Br	No	3
	<i>Callionymus superbus</i>	Yes	Mi	No	D	150	P	Br	No	2;12
	<i>Dactylopus dactylopus</i>	Yes	Mi	No	D	150	P	Br	No	2;24
	<i>Dactylopus kuiteri</i>	Yes	Mi	No	D	150	P	Br	No	2
	<i>Diplogrammus goramensis</i>	No	Mi	No	D	80	P	Br	No	1;2;4
	<i>Diplogrammus xenicus</i>	Yes	Mi	No	D	70	P	Br	No	2;36
	<i>Synchiropus bartelsi</i>	Yes	Mi	No	D	45	P	Br	No	2
	<i>Synchiropus morrisoni</i>	Yes	Mi	No	D	45	P	Br	No	1;2;3;4
	<i>Synchiropus moyeri</i>	No	Pl	No	D	75	P	Br	No	2;12
	<i>Synchiropus ocellatus</i>	No	Mi	No	D	70	P	Br	No	1;2;3;12
	<i>Synchiropus picturatus</i>	No	Mi	No	D	60	P	Br	No	2;4
	<i>Synchiropus splendidus</i>	No	Mi	No	D	60	P	Br	No	1;2;3;4
<i>Synchiropus stellatus</i>	No	Mi	No	D	60	P	Br	No	2;12	
Caracanthidae	<i>Caracanthus maculatus</i>	No	Ma	No	D	50	P	Gel	No	2;12
Carangidae	<i>Alectis ciliaris</i>	No	Ca	No	D	1300	G	Br	No	1;2;3
	<i>Carangoides bajad</i>	No	Ca	No	D	610	G	Br	No	2;37
	<i>Carangoides chrysophrys</i>	No	Pi	No	D	600	G	Br	No	38;4
	<i>Carangoides coeruleopinnatus</i>	No	N/A	No	D	400	G	Br	No	1;3
	<i>Carangoides dinema</i>	No	N/A	No	D	580	G	Br	No	3
	<i>Carangoides equula</i>	No	N/A	No	D	370	G	Br	No	3
	<i>Carangoides ferdau</i>	No	Ca	No	D	700	G	Br	No	1;2;3
	<i>Carangoides fulvoguttatus</i>	No	Pi	No	D	1300	G	Br	No	1;2;3;39
	<i>Carangoides gymnostethus</i>	No	Ca	No	D	900	G	Br	No	1;2;3;4

<i>Carangoides hedlandensis</i>	No	Pi	No	D	320	G	Br	No	1;3
<i>Carangoides humerosus</i>	No	Ca	No	D	250	G	Br	No	1;38
<i>Carangoides malabricus</i>	No	Ca	No	D	280	G	Br	No	1;38
<i>Carangoides oblongus</i>	No	Ca	No	D	460	G	Br	No	2;3;40
<i>Carangoides orthogrammus</i>	No	Ca	No	D	700	G	Br	No	2;3;41
<i>Carangoides plagiotaenia</i>	No	N/A	No	D	420	G	Br	No	2;3
<i>Carangoides talamparoides</i>	No	Pi	No	D	320	G	Br	No	1;38
<i>Caranx bucculentus</i>	No	Ca	No	D	660	G	Br	No	42;3
<i>Caranx ignobilis</i>	No	Ca	No	D	1650	G	Br	No	1;2;3;43
<i>Caranx lugubris</i>	No	Pi	No	N	740	G	Br	No	2;4
<i>Caranx melampygyus</i>	No	Pi	No	D	1000	G	Br	No	44;1;2;3;41
<i>Caranx papuensis</i>	No	Pi	No	D	800	G	Br	No	45;1;2;3
<i>Caranx sexfasciatus</i>	No	Ca	No	N	1000	G	Br	No	2;3
<i>Caranx tille</i>	No	Ca	No	D	690	G	Br	No	3;4
<i>Decapterus kurroides</i>	No	Pl	No	D	500	G	Br	No	1;4
<i>Decapterus macarellus</i>	No	Pl	No	D	350	G	Br	No	1;2;3;4
<i>Decapterus macrosoma</i>	No	Pl	No	D	320	G	Br	No	3;4
<i>Decapterus muroadsi</i>	No	Pl	No	D	450	G	Br	No	3
<i>Decapterus russelli</i>	No	Pl	No	D	380	G	Br	No	1;4
<i>Decapterus tabl</i>	No	Pl	No	D	500	G	Br	No	1;4
<i>Elagatis bipinnulatus</i>	No	Ca	No	D	1200	G	Br	No	1;2;4
<i>Gnathodon speciosus</i>	No	Ca	No	D	1400	G	Br	No	1;2;3
<i>Megalaspis cordyla</i>	No	Ca	No	D	800	G	Br	No	3;46
<i>Naucrates ductor</i>	No	Ca	No	D	750	G	Br	No	2;4
<i>Pseudocaranx dentex</i>	No	Ca	No	D	940	G	Br	No	1;2;3
<i>Scomberoides commersonianus</i>	No	Pi	No	D	1200	G	Br	No	1;2;3;39
<i>Scomberoides lysan</i>	No	Ca	No	D	700	G	Br	No	1;2;3;4
<i>Scomberoides tol</i>	No	Ca	No	D	510	G	Br	No	1;3;4
<i>Selar boops</i>	No	Ca	No	N	220	G	Br	No	1;2;3;4
<i>Selar crumenophthalmus</i>	No	Ca	No	N	300	G	Br	No	1;2;3;4
<i>Selaroides leptolepis</i>	No	Ca	No	D	220	G	Br	No	2;47
<i>Seriola dumerili</i>	No	Pi	No	D	1880	G	Br	No	1;2;3
<i>Seriola lalandi</i>	No	Ca	No	D	1930	G	Br	No	1;2;3
<i>Seriola rivoliana</i>	No	Pi	No	D	900	G	Br	No	1;2;3;48
<i>Trachinotus bailloni</i>	No	Pi	No	D	540	G	Br	No	1;2;3;4
<i>Trachinotus blochii</i>	No	Ma	No	D	650	G	Br	No	1;2;3;4
<i>Trachinotus botla</i>	No	Ma	No	D	610	G	Br	No	1;4
<i>Uraspis helvola</i>	No	N/A	No	N	500	G	Br	No	2;4
Centriscidae									
<i>Aeoliscus strigatus</i>	No	Pl	No	D	150	N/A	N/A	No	4
<i>Centriscus scutatus</i>	No	Pl	No	D	140	N/A	N/A	No	4
Chaetodontidae									
<i>Amphichaetodon howensis</i>	No	Ma	No	D	180	P	Br	No	2;4
<i>Chaetodon adiergastos</i>	Yes	N/A	No	D	160	P	Br	No	2
<i>Chaetodon andamanensis</i>	Yes	Co	No	D	150	P	Br	No	49;1;2;3
<i>Chaetodon argentatus</i>	Yes	Om	No	D	200	P	Br	Yes	49;2
<i>Chaetodon assarius</i>	No	Pl	No	D	130	P	Br	No	2;49
<i>Chaetodon aureofasciatus</i>	Yes	Co	No	D	125	P	Br	Yes	49;2;50
<i>Chaetodon auriga</i>	Yes	Om	No	D	230	G	Br	Yes	51;49
<i>Chaetodon auripes</i>	Yes	Om	No	D	200	P	Br	Yes	52;2
<i>Chaetodon baronessa</i>	Yes	Co	No	D	150	P	Br	Yes	1;2;49
<i>Chaetodon bennetti</i>	Yes	Co	No	D	180	P	Br	No	52;1;2
<i>Chaetodon burgessi</i>	Yes	N/A	No	D	140	P	Br	No	2
<i>Chaetodon citrinellus</i>	Yes	Om	No	D	130	P	Br	Yes	52;2;3
<i>Chaetodon collare</i>	Yes	Om	No	D	160	P	Br	Yes	53;2

<i>Chaetodon daedalma</i>	No	Om	No	D	150	P	Br	No	52;2
<i>Chaetodon declivis</i>	No	N/A	No	D	150	P	Br	No	4
<i>Chaetodon decussatus</i>	Yes	Om	No	D	200	P	Br	No	2
<i>Chaetodon ephippium</i>	Yes	Om	No	D	230	G	Br	Yes	49;1;2;3
<i>Chaetodon falcula</i>	Yes	Mi	No	D	200	P	Br	Yes	2;4
<i>Chaetodon flavirostris</i>	Yes	Om	No	D	200	P	Br	Yes	49;2
<i>Chaetodon flavocoronatus</i>	Yes	N/A	No	D	120	P	Br	No	2
<i>Chaetodon guentheri</i>	No	N/A	No	D	130	P	Br	No	2;4
<i>Chaetodon guttatissimus</i>	Yes	N/A	No	D	120	P	Br	No	2
<i>Chaetodon interruptus</i>	No	N/A	No	D	200	P	Br	No	2
<i>Chaetodon kleinii</i>	Yes	Om	No	D	140	G	Br	Yes	49;1;2;3
<i>Chaetodon lineolatus</i>	Yes	Om	No	D	300	G	Br	Yes	49;1;2;3
<i>Chaetodon litus</i>	No	N/A	No	D	150	P	Br	No	2;
<i>Chaetodon lunula</i>	Yes	Om	No	D	210	G	Br	Yes	49;1;2;3
<i>Chaetodon lunulatus</i>	Yes	Co	No	D	150	P	Br	Yes	49;1;2;3
<i>Chaetodon melanotus</i>	Yes	Co	No	D	150	G	Br	Yes	2;50;54
<i>Chaetodon mertensii</i>	Yes	Om	No	D	125	P	Br	No	1;2
<i>Chaetodon meyeri</i>	Yes	Co	No	D	180	P	Br	Yes	2;3
<i>Chaetodon modestus</i>	No	N/A	No	D	170	P	Br	No	4
<i>Chaetodon nippon</i>	Yes	Om	No	D	150	P	Br	No	52;2
<i>Chaetodon ocellicaudus</i>	Yes	Co	No	D	140	P	Br	No	2;4
<i>Chaetodon octofasciatus</i>	Yes	Co	No	D	120	P	Br	Yes	4
<i>Chaetodon ornatissimus</i>	Yes	Co	No	D	180	P	Br	Yes	49;2
<i>Chaetodon oxycephalus</i>	Yes	Co	No	D	250	P	Br	No	1;2
<i>Chaetodon pelewensis</i>	Yes	Mi	No	D	125	P	Br	Yes	49;1;2;3
<i>Chaetodon plebeius</i>	Yes	Co	No	D	150	P	Br	Yes	49;1;2;3
<i>Chaetodon punctofasciatus</i>	Yes	Om	No	D	120	P	Br	Yes	49;1;2;3
<i>Chaetodon quadrimaculatus</i>	Yes	Om	No	D	160	P	Br	Yes	49;2
<i>Chaetodon rafflesi</i>	Yes	Mi	No	D	150	G	Br	No	2;55;3
<i>Chaetodon rainfordi</i>	Yes	Om	No	D	150	P	Br	Yes	49;1;2;55
<i>Chaetodon reticulatus</i>	Yes	Om	No	D	160	P	Br	Yes	2;3
<i>Chaetodon selene</i>	Yes	Mi	No	D	160	P	Br	No	2;4
<i>Chaetodon semeion</i>	Yes	Mi	No	D	240	G	Br	No	2;55
<i>Chaetodon smithi</i>	No	Pl	No	D	170	P	Br	No	53;2
<i>Chaetodon speculum</i>	Yes	Om	No	D	180	P	Br	Yes	49;2;3
<i>Chaetodon tinkeri</i>	Yes	Om	No	D	150	P	Br	No	53;2
<i>Chaetodon triangulum</i>	Yes	Co	No	D	150	P	Br	Yes	51;2
<i>Chaetodon trichrous</i>	Yes	Om	No	D	120	P	Br	Yes	2;56
<i>Chaetodon tricinctus</i>	Yes	N/A	No	D	150	P	Br	Yes	2
<i>Chaetodon trifascialis</i>	No	Co	No	D	180	P	Br	No	51;57
<i>Chaetodon ulietensis</i>	Yes	Om	No	D	150	P	Br	Yes	49;2;3
<i>Chaetodon unimaculatus</i>	Yes	Om	No	D	200	P	Br	Yes	49;1;2;3
<i>Chaetodon vagabundus</i>	Yes	Om	No	D	230	P	Br	Yes	51;2;54
<i>Chaetodon wiebeli</i>	Yes	He	No	D	180	P	Br	Yes	2;4
<i>Chaetodon xanthurus</i>	Yes	Om	No	D	140	P	Br	Yes	2;3
<i>Chelmon marginalis</i>	Yes	Mi	No	D	180	P	Br	No	58;2
<i>Chelmon muelleri</i>	Yes	Mi	No	D	180	P	Br	No	1;58;2
<i>Chelmon rostratus</i>	Yes	Mi	No	D	200	P	Br	Yes	49;58;2
<i>Coradion altivelis</i>	No	N/A	No	D	200	P	Br	No	2;4
<i>Coradion chrysozonus</i>	Yes	Sp	No	D	150	P	Br	No	2;4
<i>Coradion melanopus</i>	Yes	Sp	No	D	150	P	Br	No	2;4
<i>Forcipiger flavissimus</i>	Yes	Mi	No	D	220	P	Br	Yes	1;2;3
<i>Forcipiger longirostris</i>	Yes	Mi	No	D	220	P	Br	Yes	49;1;2;3

Appendix C

	<i>Hemitaurichthys multispinosus</i>	No	Pl	No	D	208	P	Br	No	3;4
	<i>Hemitaurichthys polylepis</i>	Yes	Pl	No	D	180	P	Br	No	49;1;2;3
	<i>Hemitaurichthys thompsoni</i>	Yes	Pl	No	D	180	P	Br	No	49;1;2;3
	<i>Hemitaurichthys zoster</i>	Yes	Pl	No	D	160	P	Br	No	49;1;2;3
	<i>Heniochus acuminatus</i>	Yes	Om	No	D	250	G	Br	No	49;1;2;3
	<i>Heniochus chrysostomus</i>	Yes	Co	No	D	180	G	Br	No	1;2;3
	<i>Heniochus diphreutis</i>	No	Pl	No	D	210	G	Br	No	1;2;3
	<i>Heniochus monocerus</i>	Yes	Om	No	D	230	G	Br	No	1;2;3;4
	<i>Heniochus pleurotaenia</i>	Yes	N/A	No	D	170	G	Br	No	2
	<i>Heniochus singularius</i>	Yes	Om	No	D	230	G	Br	No	1;2;3
	<i>Heniochus varius</i>	Yes	Mi	No	D	190	G	Br	No	2;3
	<i>Parachaetodon ocellatus</i>	Yes	Mi	No	D	180	P	Br	No	1;2;59
Chanidae	<i>Chanos chanos</i>	No	Om	No	D	1800	G	Br	No	1;2;3;4
Cirrhitidae	<i>Amblycirrhitus bimacula</i>	Yes	Ma	No	D	85	P	Br	No	1;2;3;60;12
	<i>Amblycirrhitus unimacula</i>	N/A	N/A	No	D	110	P	Br	No	3
	<i>Cirrhitichthys aprinus</i>	No	Ca	No	D	100	P	Br	No	1;2;3;61
	<i>Cirrhitichthys aureus</i>	No	Ca	No	D	150	P	Br	No	1;2;3
	<i>Cirrhitichthys falco</i>	Yes	Mi	No	D	70	P	Br	No	1;2;3;60;12
	<i>Cirrhitichthys oxycephalus</i>	No	Ca	No	D	95	P	Br	No	1;2;3;60;4
	<i>Cirrhitichthys splendens</i>	No	Ca	No	D	230	P	Br	No	2
	<i>Cirrhitops hubbardi</i>	N/A	Ca	No	D	75	P	Br	No	3
	<i>Cirrhitus pinnulatus</i>	No	Ca	No	D	280	P	Br	No	1;2;3;4
	<i>Cyprinocirrhites polyactis</i>	No	Pl	No	D	150	P	Br	No	1;2;3
	<i>Isocirrhites sexfasciatus</i>	N/A	N/A	No	D	75	P	Br	No	3
	<i>Itycirrhites wilhelmi</i>	N/A	N/A	No	D	75	P	Br	No	3
	<i>Neocirrhites armatus</i>	Yes	Ma	No	D	90	P	Br	Yes	1;2;3;62
	<i>Oxycirrhites typus</i>	Yes	Mi	No	D	130	P	Br	Yes	1;2;3;62
	<i>Paracirrhites arcatus</i>	No	Ca	No	D	130	P	Br	No	1;2;3;63
	<i>Paracirrhites forsteri</i>	Yes	Ca	No	D	220	P	Br	No	1;2;3;62
	<i>Paracirrhites hemistictus</i>	No	Ma	No	D	280	P	Br	No	1;2;3;62
	<i>Paracirrhites nisus</i>	Yes	Ma	No	D	100	P	Br	No	2;3;4
	<i>Paracirrhites xanthus</i>	No	Ma	No	D	110	P	Br	No	1;2;3;4;12
Congridae	<i>Ariosoma anagoides</i>	No	N/A	No	N	400	G	Br	No	1;2;3
	<i>Ariosoma scheelei</i>	No	N/A	No	N	200	G	Br	No	1;2;3
	<i>Conger cinereus</i>	No	Ca	No	N	1300	G	Br	No	4
	<i>Gorgasia maculata</i>	No	Pl	No	D	550	P	Br	No	1;2;3
	<i>Gorgasia preclara</i>	No	Pl	No	D	400	P	Br	No	4
	<i>Heteroconger enigmaticus</i>	No	N/A	No	D	450	G	Br	No	1;2;3
	<i>Heteroconger perissodon</i>	No	N/A	No	D	600	G	Br	No	1;2;3
	<i>Heteroconger polyzona</i>	Yes	N/A	No	D	700	G	Br	No	4
	<i>Heteroconger taylori</i>	No	N/A	No	D	400	G	Br	No	1;2;3
	<i>Poecilconger fasciatus</i>	No	N/A	No	D	600	G	Br	No	1;2;3
Coryphaenidae	<i>Coryphaena hippurus</i>	No	Ca	No	D	1620	G	Br	No	4
Dactylopteridae	<i>Dactyloptena orientalis</i>	No	Ca	No	D	380	N/A	Br	No	1;2;3;4
Diodontidae	<i>Chilomycterus reticularis</i>	No	Ma	No	D	550	P	Br	No	4;64
	<i>Cyclichthys orbicularis</i>	No	Ma	No	N	150	P	Br	No	4
	<i>Cyclichthys spilostylus</i>	No	Ma	No	N	340	P	Br	No	4
	<i>Diodon holocanthus</i>	No	Ma	No	N	290	P	Br	No	4
	<i>Diodon hystrix</i>	No	Ma	No	N	710	P	Br	No	4
	<i>Diodon liturosus</i>	No	Ma	No	N	500	P	Br	No	4
	<i>Lophodiodon calori</i>	No	Ma	No	N	300	P	Br	No	4
Echeneidae	<i>Echeneis naucrates</i>	No	Ca	No	D	1000	N/A	N/A	No	4
Ephippidae	<i>Platax batavianus</i>	Yes	Om	No	D	500	G	Br	No	2;4

	<i>Platax boersii</i>	No	Om	No	D	470	G	Br	No	2
	<i>Platax orbicularis</i>	Yes	Om	No	D	280	G	Br	No	2;3
	<i>Platax pinnatus</i>	No	Om	No	D	370	G	Br	No	2;65
	<i>Platax teira</i>	No	Om	No	D	410	G	Br	No	1;2;3;66
	<i>Zabidius novemaculeatus</i>	No	N/A	No	D	450	G	Br	No	2
Fistulariidae	<i>Fistularia commersonii</i>	No	Ca	No	D	1500	P	Br	No	4
Gobiesocidae	<i>Diademichthys lineatus</i>	No	Mi	No	D	50	P	Dm	No	1;2;3;67
	<i>Lepadichthys crinophilum</i>	Yes	Mi	No	D	30	P	Dm	No	1;2;3
	<i>Lepadichthys lineatus</i>	Yes	Mi	No	D	30	P	Dm	No	1;2;3;4
Gobiidae	<i>Acentrogobius janthinopterus</i>	No	N/A	No	D	70	P	Dm	No	2
	<i>Amblyeleotris arcupinna</i>	No	Pl	Yes*	D	85	P	Dm	No	2
	<i>Amblyeleotris aurora</i>	No	Pl	Yes*	D	90	P	Dm	No	2
	<i>Amblyeleotris bellicauda</i>	No	Pl	Yes*	D	70	P	Dm	No	3;12
	<i>Amblyeleotris biguttata</i>	No	Pl	Yes*	D	104	P	Dm	No	3;12
	<i>Amblyeleotris diagonalis</i>	No	Pl	Yes*	D	90	P	Dm	No	2;3
	<i>Amblyeleotris ellipse</i>	No	Pl	Yes*	D	73	P	Dm	No	3;12
	<i>Amblyeleotris fasciata</i>	Yes	Pl	Yes*	D	80	P	Dm	No	1;2;3;4
	<i>Amblyeleotris fontanesii</i>	No	Pl	Yes*	D	170	P	Dm	No	2;3
	<i>Amblyeleotris guttata</i>	No	Pl	Yes*	D	90	P	Dm	No	1;2;3;12
	<i>Amblyeleotris gymnocephala</i>	No	Pl	Yes*	D	100	P	Dm	No	2
	<i>Amblyeleotris katherine</i>	Yes	Pl	Yes*	D	60	P	Dm	No	3;12;4
	<i>Amblyeleotris latifasciata</i>	No	Pl	Yes*	D	100	P	Dm	No	2
	<i>Amblyeleotris marquesas</i>	No	Pl	Yes*	D	85	P	Dm	No	3;12
	<i>Amblyeleotris novaecaledoniae</i>	No	Pl	Yes*	D	100	P	Dm	No	3;12
	<i>Amblyeleotris ogasawarensis</i>	No	Pl	Yes*	D	110	P	Dm	No	3;12
	<i>Amblyeleotris periophthalma</i>	No	Pl	Yes*	D	80	P	Dm	No	1;2;3
	<i>Amblyeleotris randalli</i>	No	Pl	Yes*	D	90	P	Dm	No	2;3
	<i>Amblyeleotris rhyax</i>	No	Pl	Yes*	D	80	P	Dm	No	1;2
	<i>Amblyeleotris rubrimarginata</i>	No	Pl	Yes*	D	90	P	Dm	No	2;3
	<i>Amblyeleotris steinitzi</i>	No	Pl	Yes*	D	80	P	Dm	No	1;2;3;12
	<i>Amblyeleotris stenoaeniata</i>	No	Pl	Yes*	D	95	P	Dm	No	3;12
	<i>Amblyeleotris wheeleri</i>	No	Pl	Yes*	D	80	P	Dm	No	1;2
	<i>Amblyeleotris yanoi</i>	No	Pl	Yes*	D	120	P	Dm	No	2
	<i>Amblygobius buanensis</i>	Yes	Om	No	D	60	P	Dm	No	2;4
	<i>Amblygobius bynoensis</i>	Yes	He	Yes	D	80	P	Dm	No	1;2;3;68
	<i>Amblygobius decussatus</i>	No	Om	No	D	60	P	Dm	No	1;2;3;69
	<i>Amblygobius esakiae</i>	Yes	He	No	D	65	P	Dm	No	2
	<i>Amblygobius nocturnis</i>	Yes	De	Yes	D	50	P	Dm	Yes	1;2;3;70
	<i>Amblygobius phalaena</i>	Yes	He	Yes	D	135	P	Dm	Yes	1;2;3;68
	<i>Amblygobius semicinctus</i>	Yes	Om	Yes	D	140	P	Dm	Yes	2;4
	<i>Amblygobius sphynx</i>	Yes	Mi	No	D	165	P	Dm	No	1;2;3
	<i>Asterropteryx bipunctatus</i>	No	De	Yes	D	40	P	Dm	No	2;71;4
	<i>Asterropteryx ensifera</i>	No	Pl	No	D	35	P	Dm	No	1;2;3;4
	<i>Asterropteryx semipunctatus</i>	No	De	Yes	D	60	P	Dm	No	1;2;3;70
	<i>Asterropteryx spinosa</i>	No	De	No	D	35	P	Dm	No	3;71
	<i>Asterropteryx striata</i>	No	Pl	No	D	35	P	Dm	No	2;72
	<i>Bathygobius coalitus</i>	No	De	No	D	120	P	Dm	No	1;3;71
	<i>Bathygobius cocosensis</i>	Yes	Mi	No	D	50	P	Dm	No	1;2;3;73
	<i>Bathygobius cotticeps</i>	No	De	No	D	110	P	Dm	No	3;71
	<i>Bathygobius cyclopterus</i>	No	De	No	D	70	P	Dm	No	3;71
	<i>Bathygobius fuscus</i>	No	De	No	D	80	P	Dm	No	1;3;70
	<i>Bathygobius laddi</i>	No	De	No	D	50	P	Dm	No	1;3;71
	<i>Bathygobius padangensis</i>	No	De	No	D	50	P	Dm	No	2;71

<i>Bryaninops amplus</i>	N/A	N/A	No	D	50	P	Dm	No	2
<i>Bryaninops dianneae</i>	N/A	N/A	No	D	24	P	Dm	No	3
<i>Bryaninops erythroptus</i>	N/A	N/A	No	D	20	P	Dm	No	2
<i>Bryaninops loki</i>	N/A	N/A	No	D	30	P	Dm	No	1;2;3
<i>Bryaninops natans</i>	No	N/A	No	D	25	P	Dm	No	1;2;3
<i>Bryaninops tigris</i>	No	Pl	No	D	55	P	Dm	No	2;4
<i>Bryaninops yongei</i>	Yes	Pl	No	D	35	P	Dm	No	2;74
<i>Callogobius clitellus</i>	No	N/A	No	D	45	P	Dm	No	2
<i>Callogobius hasseltii</i>	No	N/A	No	D	60	P	Dm	No	1;2;3
<i>Callogobius maculipinnis</i>	No	N/A	No	D	90	P	Dm	No	1;3
<i>Callogobius sclateri</i>	No	N/A	No	D	45	P	Dm	No	3
<i>Calumia profunda</i>	No	N/A	No	D	23	P	Dm	No	2
<i>Cristatogobius lophius</i>	No	N/A	Yes*	D	75	P	Dm	No	2;4
<i>Cryptocentrus caeruleopunctatus</i>	No	Mi	Yes*	D	80	P	Dm	No	1;2;75
<i>Cryptocentrus cinctus</i>	Yes	Mi	Yes*	D	70	P	Dm	No	1;2;75
<i>Cryptocentrus cyanotaenia</i>	Yes	Mi	Yes*	D	70	P	Dm	No	1;2;75;4
<i>Cryptocentrus fasciatus</i>	Yes	Mi	Yes*	D	80	P	Dm	No	1;2;3;75;4
<i>Cryptocentrus inexplicatus</i>	No	Mi	Yes*	D	75	P	Dm	No	2;75
<i>Cryptocentrus insignitus</i>	No	Mi	Yes*	D	80	P	Dm	No	1;2;75
<i>Cryptocentrus leptocephalus</i>	No	Mi	Yes*	D	100	P	Dm	No	2;75
<i>Cryptocentrus leucostictus</i>	No	Mi	Yes*	D	90	P	Dm	No	1;2;3;75
<i>Cryptocentrus pavaninoides</i>	No	Mi	Yes*	D	140	P	Dm	No	2;75
<i>Cryptocentrus polyophthalmus</i>	No	Mi	Yes*	D	90	P	Dm	No	2;75
<i>Cryptocentrus strigilliceptus</i>	No	Mi	Yes*	D	60	P	Dm	No	2;75
<i>Ctenogobiops aurocingulus</i>	No	N/A	Yes*	D	55	P	Dm	No	2;3
<i>Ctenogobiops crocineus</i>	No	N/A	Yes*	D	50	P	Dm	No	2
<i>Ctenogobiops feroculus</i>	Yes	N/A	Yes*	D	55	P	Dm	No	2;3;4
<i>Ctenogobiops pomasticus</i>	No	N/A	Yes*	D	50	P	Dm	No	1;2;3
<i>Ctenogobiops tangaroai</i>	No	N/A	Yes*	D	50	P	Dm	No	1;2;3
<i>Ctenogobiops tongaensis</i>	No	N/A	Yes*	D	55	P	Dm	No	3
<i>Discordipinna griessingeri</i>	No	N/A	No	D	25	P	Dm	No	1;2;3
<i>Echinogobius hayashi</i>	No	N/A	Yes	D	120	P	Dm	No	2;4
<i>Eviota afelei</i>	No	N/A	No	D	23	P	Dm	No	1;3
<i>Eviota albolineata</i>	No	Mi	No	D	35	P	Dm	No	1;2;3;12
<i>Eviota bifasciata</i>	No	Pl	No	D	30	P	Dm	No	1;2;4
<i>Eviota cometa</i>	No	N/A	No	D	24	P	Dm	No	3
<i>Eviota distigma</i>	No	N/A	No	D	27	P	Dm	No	3
<i>Eviota fasciola</i>	No	N/A	No	D	24	P	Dm	No	3
<i>Eviota guttata</i>	No	Mi	No	D	25	P	Dm	No	1;2;3;12
<i>Eviota infulata</i>	No	N/A	No	D	24	P	Dm	No	3
<i>Eviota lachdeberi</i>	No	N/A	No	D	25	P	Dm	No	2
<i>Eviota latifasciata</i>	No	N/A	No	D	20	P	Dm	No	1;2;3
<i>Eviota melasma</i>	No	N/A	No	D	32	P	Dm	No	1;2;3
<i>Eviota mikiae</i>	No	N/A	No	D	25	P	Dm	No	2
<i>Eviota monostigma</i>	No	N/A	No	D	33	P	Dm	No	3
<i>Eviota nebulosa</i>	No	N/A	No	D	24	P	Dm	No	3
<i>Eviota nigriventris</i>	No	N/A	No	D	25	P	Dm	No	1;2;3
<i>Eviota pellucida</i>	No	N/A	No	D	25	P	Dm	No	1;2;3
<i>Eviota prasina</i>	No	N/A	No	D	30	P	Dm	No	1;2
<i>Eviota prasites</i>	No	Mi	No	D	25	P	Dm	No	1;2;3;12
<i>Eviota punctulata</i>	No	N/A	No	D	20	P	Dm	No	1;2;3
<i>Eviota queenslandica</i>	No	Om	No	D	30	P	Dm	No	1;2;70
<i>Eviota sebreei</i>	No	N/A	No	D	25	P	Dm	No	1;2;3

<i>Eviota sigillata</i>	No	N/A	No	D	25	P	Dm	No	1;2
<i>Eviota smaragdus</i>	No	N/A	No	D	24	P	Dm	No	2;3
<i>Eviota storthynx</i>	No	N/A	No	D	25	P	Dm	No	2
<i>Eviota zebrina</i>	No	Om	No	D	24	P	Dm	No	1;3;69
<i>Exyrias bellissimus</i>	No	Om	No	D	130	P	Dm	No	1;2;3
<i>Exyrias ferrarisi</i>	Yes	N/A	No	D	95	P	Dm	No	2
<i>Exyrias puntang</i>	No	Om	No	D	135	P	Dm	No	1;2;3;15
<i>Favonigobius reichei</i>	No	Ma	No	D	45	P	Dm	No	2;15
<i>Fusigobius aureus</i>	No	N/A	No	D	75	P	Dm	No	2
<i>Fusigobius duospilus</i>	No	Ma	No	D	55	P	Dm	No	1;2;3;12
<i>Fusigobius inframaculatus</i>	No	N/A	No	D	75	P	Dm	No	2
<i>Fusigobius neophytus</i>	No	Ma	No	D	70	P	Dm	No	1;2;3;12
<i>Fusigobius signipinnis</i>	No	N/A	No	D	30	P	Dm	No	2
<i>Gnatholepis anjerensis</i>	No	N/A	No	D	55	P	Dm	No	1;2;3
<i>Gnatholepis cauerensis</i>	No	N/A	No	D	45	P	Dm	No	1;2;3
<i>Gobiodon acicularis</i>	No	Om	No	D	38	P	Dm	No	2;71
<i>Gobiodon axillaris</i>	N/A	N/A	No	D	40	P	Dm	No	3
<i>Gobiodon brochus</i>	Yes	Om	No	D	34	P	Dm	No	1;3;76
<i>Gobiodon ceramensis</i>	Yes	N/A	No	D	35	P	Dm	No	2
<i>Gobiodon citrinus</i>	No	Co	No	D	55	P	Dm	No	1;2;3;77
<i>Gobiodon histrio</i>	No	Om	No	D	35	P	Dm	No	1;3;76
<i>Gobiodon okinawae</i>	No	Om	No	D	35	P	Dm	No	1;2;3;76
<i>Gobiodon quinquestrigatus</i>	Yes	Om	No	D	35	P	Dm	Yes	1;2;3;76
<i>Gobiodon spilophthalmus</i>	No	N/A	No	D	35	P	Dm	No	2
<i>Gobiopsis exigua</i>	No	N/A	No	D	50	P	Dm	No	2
<i>Heteroplopomus barbatus</i>	No	N/A	No	D	25	P	Dm	No	2
<i>Istigobius decoratus</i>	No	De	No	D	90	P	Dm	No	1;2;3;70
<i>Istigobius goldmanni</i>	No	De	No	D	62	P	Dm	No	3;70
<i>Istigobius ornatus</i>	No	Mi	No	D	90	P	Dm	No	1;2;3;4
<i>Istigobius rigilius</i>	No	Om	No	D	90	P	Dm	No	1;2;3;69
<i>Koumansetta hectori</i>	No	Om	No	D	50	P	Dm	No	2;78
<i>Koumansetta rainfordi</i>	No	He	No	D	55	P	Dm	No	1;2;3;70
<i>Lotilia graciliosa</i>	No	N/A	Yes*	D	45	P	Dm	No	1;2;3
<i>Luposicya lupus</i>	No	N/A	No	D	35	P	Dm	No	2
<i>Macrodontogobius wilburi</i>	No	Om	No	D	65	P	Dm	No	1;2;3;26
<i>Mahidolia mystacina</i>	No	Mi	Yes*	D	70	P	Dm	No	1;2;3;79
<i>Myersina lachneri</i>	No	N/A	Yes*	D	50	P	Dm	No	2
<i>Myersina nigrivirgata</i>	No	N/A	Yes*	D	100	P	Dm	No	2
<i>Oplopomus caninoides</i>	No	N/A	No	D	60	P	Dm	No	2
<i>Oplopomus oplopomus</i>	Yes	Mi	No	D	75	P	Dm	No	1;2;3;80
<i>Oxyurichthys notonema</i>	N/A	N/A	Yes	D	160	P	Dm	No	3
<i>Oxyurichthys papuensis</i>	No	N/A	Yes	D	200	P	Dm	No	1;2;3
<i>Pandaka pusilila</i>	No	N/A	No	D	12	P	Dm	No	2
<i>Paragobiodon echinocephalus</i>	No	N/A	No	D	35	P	Dm	Yes	1;2;3
<i>Paragobiodon lacunicolus</i>	No	N/A	No	D	35	P	Dm	No	1;2;3
<i>Paragobiodon modestus</i>	No	N/A	No	D	35	P	Dm	No	1;2;3
<i>Paragobiodon xanthosomus</i>	No	N/A	No	D	40	P	Dm	Yes	1;2;3
<i>Phyllogobius platycephalops</i>	No	N/A	No	D	25	P	Dm	No	2
<i>Pleurosicya bilobata</i>	No	Mi	No	D	28	P	Dm	No	3;4
<i>Pleurosicya boldinghi</i>	Yes	N/A	No	D	35	P	Dm	No	2
<i>Pleurosicya coerulea</i>	N/A	N/A	No	D	22	P	Dm	No	3
<i>Pleurosicya elongata</i>	N/A	N/A	No	D	35	P	Dm	No	2
<i>Pleurosicya fringilla</i>	N/A	N/A	No	D	22	P	Dm	No	3

<i>Pleurosicya labiata</i>	No	N/A	No	D	35	P	Dm	No	2
<i>Pleurosicya micheli</i>	No	N/A	No	D	25	P	Dm	No	2;3
<i>Pleurosicya mossambica</i>	No	N/A	No	D	35	P	Dm	No	1;2;3
<i>Priolepis aureoviridis</i>	No	N/A	No	D	65	P	Dm	No	2;
<i>Priolepis cincta</i>	No	N/A	No	D	50	P	Dm	Yes	1;2;3
<i>Priolepis compita</i>	No	Mi	No	D	17	P	Dm	No	3
<i>Priolepis fallacincta</i>	N/A	N/A	No	D	32	P	Dm	No	3
<i>Priolepis inhaca</i>	No	N/A	No	D	35	P	Dm	No	1;2;3
<i>Priolepis kappa</i>	N/A	N/A	No	D	26	P	Dm	No	3
<i>Priolepis nocturna</i>	No	N/A	No	D	35	P	Dm	No	1;2;3;
<i>Priolepis nuchifasciata</i>	Yes	Mi	No	D	40	P	Dm	No	1;70
<i>Priolepis pallidicincta</i>	N/A	N/A	No	D	36	P	Dm	No	3
<i>Priolepis semidoliata</i>	N/A	N/A	No	D	36	P	Dm	No	3
<i>Priolepis squamogena</i>	N/A	N/A	No	D	54	P	Dm	No	3
<i>Priolepis triops</i>	N/A	N/A	No	D	26	P	Dm	No	3
<i>Signigobius biocellatus</i>	Yes	Mi	Yes	D	65	P	Dm	Yes	1;2;4
<i>Stonogobiops larsonae</i>	No	N/A	Yes*	D	60	P	Dm	No	2
<i>Stonogobiops medon</i>	No	N/A	Yes*	D	60	P	Dm	No	2
<i>Stonogobiops nematodes</i>	Yes	N/A	Yes*	D	60	P	Dm	No	1;2 4
<i>Stonogobiops xanthorhinica</i>	Yes	N/A	Yes*	D	60	P	Dm	No	1;2;3 4
<i>Stonogobiops yasha</i>	Yes	N/A	Yes*	D	60	P	Dm	No	1;2;3;4
<i>Tomiyamichthys lanceolata</i>	No	N/A	Yes*	D	50	P	Dm	No	2;4
<i>Tomiyamichthys oni</i>	No	N/A	Yes*	D	100	P	Dm	No	2
<i>Trimma anaima</i>	No	N/A	No	D	20	P	Dm	No	2
<i>Trimma annosum</i>	No	N/A	No	D	25	P	Dm	No	2
<i>Trimma benjamini</i>	No	Pl	No	D	25	P	Dm	No	1;2;3;4
<i>Trimma caesiura</i>	No	Om	No	D	25	P	Dm	No	1;2;3;70
<i>Trimma cana</i>	No	N/A	No	D	25	P	Dm	No	2
<i>Trimma emeryi</i>	N/A	N/A	No	D	24	P	Dm	No	3
<i>Trimma griffithsi</i>	No	Pl	No	D	23	P	Dm	No	2;81
<i>Trimma halonevum</i>	No	N/A	No	D	28	P	Dm	No	2
<i>Trimma hoesei</i>	Yes	N/A	No	D	25	P	Dm	No	25
<i>Trimma macrophthalma</i>	No	N/A	No	D	25	P	Dm	No	1;2;3
<i>Trimma milta</i>	N/A	N/A	No	D	30	P	Dm	No	3
<i>Trimma naudei</i>	No	Mi	No	D	30	P	Dm	No	2;4
<i>Trimma okinawae</i>	No	N/A	No	D	28	P	Dm	No	1;2;3
<i>Trimma rubromaculatus</i>	No	N/A	No	D	22	P	Dm	No	2
<i>Trimma stobbsi</i>	No	N/A	No	D	20	P	Dm	No	2
<i>Trimma striata</i>	No	Mi	No	D	35	P	Dm	No	1;2;3;70
<i>Trimma taylori</i>	No	Pl	No	D	25	P	Dm	No	1;2;3;82
<i>Trimma tevegae</i>	No	Pl	No	D	40	P	Dm	No	1;2;3;4
<i>Trysogobius colini</i>	No	N/A	No	D	50	P	Dm	No	2
<i>Valenciennea alleni</i>	Yes	Mi	Yes	D	65	P	Dm	No	2;4
<i>Valenciennea bella</i>	Yes	Mi	Yes	D	90	P	Dm	No	2;4
<i>Valenciennea decora</i>	Yes	Mi	Yes	D	140	P	Dm	No	1;2;3
<i>Valenciennea helsdingeni</i>	Yes	Mi	Yes	D	145	P	Dm	Yes	1;2;3;83
<i>Valenciennea immaculata</i>	Yes	Mi	Yes	D	105	P	Dm	No	1;2;3;4
<i>Valenciennea limicola</i>	Yes	Mi	Yes	D	80	P	Dm	No	2
<i>Valenciennea longipinnis</i>	Yes	Mi	Yes	D	150	P	Dm	Yes	1;2;3;4
<i>Valenciennea muralis</i>	Yes	Mi	Yes	D	115	P	Dm	Yes	1;2;3;50;4
<i>Valenciennea parva</i>	Yes	Mi	Yes	D	65	P	Dm	Yes	1;2;3
<i>Valenciennea puellaris</i>	Yes	Mi	Yes	D	155	P	Dm	Yes	1;2;3;4
<i>Valenciennea randalli</i>	No	Mi	Yes	D	90	P	Dm	No	1;2;3

	<i>Valenciennea sexguttata</i>	Yes	Mi	Yes	D	115	P	Dm	Yes	1;2;3
	<i>Valenciennea strigata</i>	Yes	Mi	Yes	D	155	P	Dm	Yes	1;2;3;84
	<i>Valenciennea wardii</i>	Yes	Mi	Yes	D	100	P	Dm	No	1;2;3
	<i>Vanderhorstia ambanoro</i>	No	N/A	Yes*	D	120	P	Dm	No	2;3
	<i>Vanderhorstia flavilineata</i>	No	N/A	Yes*	D	40	P	Dm	No	2;4
	<i>Vanderhorstia macropteryx</i>	No	N/A	Yes*	D	60	P	Dm	No	2
	<i>Vanderhorstia ornatissima</i>	No	N/A	Yes*	D	60	P	Dm	No	2;4
	<i>Yongeichthys nebulosus</i>	No	Pl	No	D	160	P	Dm	No	2;17
Haemulidae	<i>Diagramma melanacrum</i>	No	Ma	No	N	500	G	Br	No	2
	<i>Diagramma pictum</i>	No	Ca	No	N	940	G	Br	No	1;2;3;4
	<i>Plectorhinchus albovittatus</i>	No	Ma	No	N	1000	G	Br	No	1;2;3
	<i>Plectorhinchus chaetodonoides</i>	No	Ca	No	N	720	G	Br	No	1;2;3;4
	<i>Plectorhinchus chrysotaenia</i>	No	Ma	No	N	500	G	Br	No	2;3
	<i>Plectorhinchus flavomaculatus</i>	No	Ca	No	N	600	G	Br	No	1;2;3
	<i>Plectorhinchus gibbosus</i>	No	Ma	No	N	600	G	Br	No	1;2;3
	<i>Plectorhinchus lessonii</i>	No	Ma	No	N	480	G	Br	No	1;2;3
	<i>Plectorhinchus lineatus</i>	No	Ma	No	N	480	G	Br	No	1;2;3;4
	<i>Plectorhinchus multivittatum</i>	No	Ma	No	N	500	G	Br	No	2;3
	<i>Plectorhinchus picus</i>	No	Ma	No	N	850	G	Br	No	1;2;3;4
	<i>Plectorhinchus polytaenia</i>	No	Mi	No	N	400	G	Br	No	2
	<i>Plectorhinchus unicolor</i>	No	Ma	No	N	800	G	Br	No	1;2;3
	<i>Plectorhinchus vittatus</i>	No	Ma	No	N	850	G	Br	No	1;2;3
	<i>Pomadasyd argenteus</i>	N/A	N/A	No	D	520	G	Br	No	1
Holocentridae	<i>Myripristis adusta</i>	No	Pl	No	N	320	G	Br	No	4
	<i>Myripristis amaena</i>	No	Pl	No	N	320	G	Br	No	4
	<i>Myripristis berndti</i>	No	Pl	No	N	300	G	Br	No	4
	<i>Myripristis botche</i>	Yes	Pl	No	N	300	G	Br	No	4
	<i>Myripristis chryseres</i>	No	Pl	No	N	250	G	Br	No	4
	<i>Myripristis earlei</i>	No	Ma	No	N	300	G	Br	No	12
	<i>Myripristis hexagona</i>	No	Pl	No	N	200	G	Br	No	4
	<i>Myripristis kuntee</i>	No	Pl	No	N	200	G	Br	No	4
	<i>Myripristis murdjan</i>	No	Pl	No	N	270	G	Br	No	4
	<i>Myripristis pralinia</i>	No	Pl	No	N	200	G	Br	No	4
	<i>Myripristis trachyacron</i>	No	Pl	No	N	150	G	Br	No	12
	<i>Myripristis violacea</i>	No	Pl	No	N	200	G	Br	No	4
	<i>Myripristis vittata</i>	No	Pl	No	N	200	G	Br	No	4
	<i>Myripristis woodsi</i>	No	Pl	No	N	210	G	Br	No	4
	<i>Neoniphon argenteus</i>	No	Ma	No	N	190	G	Br	No	4
	<i>Neoniphon aurolineatus</i>	No	Ma	No	N	250	G	Br	No	12
	<i>Neoniphon opercularis</i>	No	Ma	No	N	350	G	Br	No	4
	<i>Neoniphon sammara</i>	No	Ca	No	N	320	G	Br	No	4
	<i>Plectrypops lima</i>	No	Ca	No	N	160	G	Br	No	4
	<i>Sargocentron caudimaculatum</i>	No	Ma	No	N	250	G	Br	No	4
	<i>Sargocentron cornutum</i>	No	Ma	No	N	180	G	Br	No	4
	<i>Sargocentron diadema</i>	No	Ma	No	N	170	G	Br	No	85
	<i>Sargocentron ensiferum</i>	No	Ma	No	N	250	G	Br	No	1;2;3
	<i>Sargocentron iota</i>	No	Ma	No	N	80	G	Br	No	1;2;3
	<i>Sargocentron itodai</i>	No	Ma	No	N	170	G	Br	No	85
	<i>Sargocentron melanospilos</i>	No	Ma	No	N	250	G	Br	No	4
	<i>Sargocentron microstoma</i>	No	Ma	No	N	190	G	Br	No	3
	<i>Sargocentron punctatissimum</i>	No	Ma	No	N	200	G	Br	No	4
	<i>Sargocentron rubrum</i>	No	Ma	No	N	270	G	Br	No	4
	<i>Sargocentron spiniferum</i>	Yes	Ca	No	N	450	G	Br	No	4

	<i>Sargocentron tiere</i>	No	Ca	No	N	330	G	Br	No	4
	<i>Sargocentron tiereoides</i>	No	Ma	No	N	160	G	Br	No	4
	<i>Sargocentron violaceum</i>	No	Ma	No	N	250	G	Br	No	4
Kyphosidae	<i>Kyphosus bigibbus</i>	No	He	No	D	700	G	Br	No	1;2;3;86
	<i>Kyphosus cinerascens</i>	No	He	No	D	450	G	Br	No	1;2;3;86
	<i>Kyphosus cornelii</i>	No	He	No	D	600	G	Br	No	2;3;87
	<i>Kyphosus pacificus</i>	No	He	No	D	650	G	Br	No	3
	<i>Kyphosus sydneyanus</i>	No	He	No	D	600	G	Br	No	3;86
	<i>Kyphosus vaigiensis</i>	No	He	No	D	450	G	Br	No	1;2;3;86
	<i>Sectator ocyurus</i>	No	He	No	D	380	G	Br	No	3;12
Labridae	<i>Anampses caeruleopunctatus</i>	Yes	Ma	No	D	420	G	Br	No	1;2;3;4
	<i>Anampses elegans</i>	No	Ma	No	D	300	P	Br	No	2;4
	<i>Anampses femininus</i>	No	Mi	No	D	240	P	Br	No	1;2;3;87
	<i>Anampses geographicus</i>	No	Pl	No	D	200	P	Br	No	1;2;3;4
	<i>Anampses lennardi</i>	Yes	Mi	No	D	280	P	Br	No	2;4
	<i>Anampses lineatus</i>	No	Ma	No	D	120	P	Br	No	2;12
	<i>Anampses melanurus</i>	Yes	Mi	No	D	120	P	Br	No	1;2;3
	<i>Anampses meleagrides</i>	No	Ma	No	D	220	P	Br	No	1;2;3;12
	<i>Anampses neoguinaicus</i>	No	Ma	No	D	170	P	Br	No	2;4
	<i>Anampses twistii</i>	Yes	Ma	No	D	180	P	Br	No	1;2;3;4
	<i>Bodianus anthioides</i>	No	Ma	No	D	210	P	Br	No	1;2;3
	<i>Bodianus axillaris</i>	No	Ma	No	D	200	P	Br	No	1;2;3;88
	<i>Bodianus bilunulatus</i>	No	Ma	No	D	550	P	Br	No	2;3
	<i>Bodianus bimaculatus</i>	No	Ma	No	D	100	P	Br	No	1;2;12
	<i>Bodianus diana</i>	Yes	Ma	No	D	250	P	Br	No	1;2;12
	<i>Bodianus loxozonus</i>	No	Ma	No	D	400	P	Br	No	1;2;3;4
	<i>Bodianus mesothorax</i>	No	Ma	No	D	190	P	Br	No	1;2;3;4;12
	<i>Bodianus neilli</i>	No	Ma	No	D	200	P	Br	No	2;4
	<i>Bodianus opercularis</i>	No	Ma	No	D	150	P	Br	No	2
	<i>Bodianus perditio</i>	No	Ma	No	D	150	P	Br	No	1;2;3
	<i>Bodianus prognathus</i>	No	Ma	No	D	200	P	Br	No	1;2;12
	<i>Bolbometopon muricatum</i>	No	Co	No	D	1260	B	Br	No	1;2;3;89
	<i>Calotomus carolinus</i>	No	He	No	D	500	B	Br	No	1;2;3;4
	<i>Calotomus spinidens</i>	No	He	No	D	190	B	Br	No	1;2;3;4
	<i>Cetoscarus bicolor</i>	No	De	No	D	800	B	Br	No	1;2;3
	<i>Cheilinus chlorourus</i>	No	Ma	No	D	360	P	Br	No	2;90
	<i>Cheilinus fasciatus</i>	No	Ma	No	D	360	P	Br	No	1;2;3;90
	<i>Cheilinus oxycephalus</i>	Yes	Mi	No	D	170	P	Br	No	1;2;3;91
	<i>Cheilinus trilobatus</i>	No	Ma	No	D	450	P	Br	No	1;2;3;90
	<i>Cheilinus undulatus</i>	No	Ca	No	D	2290	G	Br	No	1;2;3
	<i>Cheilio inermis</i>	No	Ca	No	D	500	P	Br	No	1;2;3
	<i>Chlorurus bleekeri</i>	No	De	No	D	480	B	Br	No	1;2;3;92
	<i>Chlorurus bowersi</i>	No	De	No	D	300	B	Br	No	2
	<i>Chlorurus capistratoides</i>	No	De	No	D	550	B	Br	No	2
	<i>Chlorurus frontalis</i>	No	De	No	D	500	B	Br	No	1;2;3
	<i>Chlorurus japanensis</i>	No	De	No	D	300	B	Br	No	2;3
	<i>Chlorurus microrhinos</i>	No	De	No	D	800	B	Br	No	1;2;3;92
	<i>Chlorurus sordidus</i>	No	De	No	D	400	B	Br	No	1;2;3
	<i>Chlorurus strongycephalus</i>	No	De	No	D	700	B	Br	No	2
	<i>Chlorurus troschelii</i>	No	De	No	D	350	B	Br	No	2
	<i>Choerodon anchorago</i>	No	Ma	No	D	380	P	Br	No	1;2;3;26
	<i>Choerodon cephalotes</i>	No	Ma	No	D	380	P	Br	No	1;4
	<i>Choerodon cyanodus</i>	No	Ma	No	D	700	P	Br	No	1;2

<i>Choerodon fasciatus</i>	No	Ma	No	D	300	P	Br	Yes	1;2;3
<i>Choerodon graphicus</i>	No	Ma	No	D	460	P	Br	No	1;2;3;4
<i>Choerodon jordani</i>	No	Ma	No	D	170	P	Br	No	2;3
<i>Choerodon monostigma</i>	No	Ma	No	D	250	P	Br	No	2
<i>Choerodon oligacanthus</i>	Yes	Ma	No	D	350	P	Br	No	2
<i>Choerodon rubescens</i>	No	Ma	No	D	900	P	Br	No	2;39
<i>Choerodon schoenleinii</i>	No	Ma	No	D	900	P	Br	No	1;2
<i>Choerodon vitta</i>	No	Ma	No	D	200	P	Br	No	2
<i>Choerodon zosterophorus</i>	No	Ma	No	D	250	P	Br	No	2
<i>Cirrhilabrus aurantidorsalis</i>	No	Pl	No	D	100	G	Br	No	2;12
<i>Cirrhilabrus bathyphilus</i>	No	Pl	No	D	110	G	Br	No	2;12
<i>Cirrhilabrus condei</i>	No	Pl	No	D	100	G	Br	No	2;12
<i>Cirrhilabrus cyanopleura</i>	No	Pl	No	D	100	G	Br	No	2;4
<i>Cirrhilabrus exquisitus</i>	No	Pl	No	D	120	G	Br	No	1;2;3;12
<i>Cirrhilabrus filamentosus</i>	No	Pl	No	D	120	G	Br	No	2;4
<i>Cirrhilabrus flavidorsalis</i>	No	Pl	No	D	65	G	Br	No	2;12
<i>Cirrhilabrus joanalleneae</i>	No	Pl	No	D	60	G	Br	No	2;12
<i>Cirrhilabrus katherinae</i>	No	Pl	No	D	90	G	Br	No	2;12
<i>Cirrhilabrus laboutei</i>	No	Pl	No	D	110	G	Br	No	1;2;3;12
<i>Cirrhilabrus lineatus</i>	No	Pl	No	D	120	G	Br	No	1;2;3;4
<i>Cirrhilabrus lubbocki</i>	No	Pl	No	D	80	G	Br	No	2;12;4
<i>Cirrhilabrus luteovittatus</i>	No	Pl	No	D	120	G	Br	No	2;12
<i>Cirrhilabrus marjorie</i>	No	Pl	No	D	70	G	Br	No	2;12
<i>Cirrhilabrus morrisoni</i>	No	Pl	No	D	80	G	Br	No	2;12
<i>Cirrhilabrus punctatus</i>	No	Ca	No	D	130	G	Br	No	1;2;3;12
<i>Cirrhilabrus randalli</i>	No	Pl	No	D	85	G	Br	No	2;12
<i>Cirrhilabrus rhomboidalis</i>	No	Pl	No	D	85	G	Br	No	2;12
<i>Cirrhilabrus rubripinnis</i>	No	Pl	No	D	80	G	Br	No	2;12
<i>Cirrhilabrus rubrimarginatus</i>	No	Pl	No	D	150	G	Br	No	1;2;3;12
<i>Cirrhilabrus scottorum</i>	No	Pl	No	D	120	G	Br	No	1;2;3;4
<i>Cirrhilabrus solorensis</i>	No	Pl	No	D	120	G	Br	No	2
<i>Cirrhilabrus temminckii</i>	No	Pl	No	D	100	G	Br	No	2;12;4
<i>Cirrhilabrus tonozukai</i>	No	Pl	No	D	70	G	Br	No	2;4;12
<i>Cirrhilabrus walindi</i>	No	Pl	No	D	70	G	Br	No	2;12
<i>Cirrhilabrus walshi</i>	No	Pl	No	D	100	G	Br	No	2;12
<i>Coris auricularis</i>	No	Ma	No	D	400	P	Br	No	2;93
<i>Coris aurilineata</i>	No	Om	No	D	115	P	Br	No	2;4;94
<i>Coris aygula</i>	No	Ma	No	D	400	P	Br	No	1;2;3
<i>Coris batuensis</i>	No	Ma	No	D	170	P	Br	No	2;3;4;12
<i>Coris bulbifrons</i>	No	Ma	No	D	1400	P	Br	No	2;4
<i>Coris caudimacula</i>	No	Ma	No	D	200	P	Br	No	2;3
<i>Coris centralis</i>	No	Ma	No	D	300	P	Br	No	2;3;4;12
<i>Coris dorsomacula</i>	No	Ma	No	D	200	P	Br	No	1;2;3;12
<i>Coris gaimard</i>	No	Ma	No	D	380	P	Br	No	1;2;3
<i>Coris pictoides</i>	No	Ma	No	D	120	P	Br	No	1;2;12
<i>Cymolutes praetextatus</i>	No	Ma	No	D	200	P	Br	No	1;2;3;12
<i>Cymolutes torquatus</i>	No	N/A	No	D	120	P	Br	No	1;2;3
<i>Diprocatacanthus xanthurus</i>	No	Mi	No	D	80	P	Br	No	1;2;12
<i>Epibulus insidiator</i>	No	Ca	No	D	350	P	Br	No	1;2;3
<i>Gomphosus varius</i>	No	Mi	No	D	280	P	Br	No	1;2;3
<i>Halichoeres argus</i>	No	N/A	No	D	110	P	Br	No	1;2
<i>Halichoeres binotopsis</i>	No	N/A	No	D	120	P	Br	No	2;4
<i>Halichoeres biocellatus</i>	No	N/A	No	D	120	P	Br	No	1;2;3

<i>Halichoeres chlorocephalus</i>	No	N/A	No	D	90	P	Br	No	2
<i>Halichoeres chloropterus</i>	No	Ma	No	D	190	P	Br	No	1;2;4
<i>Halichoeres chrysus</i>	No	Mi	No	D	120	P	Br	No	1;2;3;12
<i>Halichoeres cosmetus</i>	No	N/A	No	D	130	P	Br	No	2
<i>Halichoeres hortulanus</i>	No	Ma	No	D	270	P	Br	No	1;2;3;4
<i>Halichoeres leucoxanthus</i>	No	N/A	No	D	110	P	Br	No	2
<i>Halichoeres leucurus</i>	Yes	N/A	No	D	120	P	Br	No	2
<i>Halichoeres margaritaceus</i>	No	Ma	No	D	130	P	Br	No	1;2;3;4
<i>Halichoeres marginatus</i>	No	Ma	No	D	170	P	Br	No	1;2;3
<i>Halichoeres melanochir</i>	Yes	N/A	No	D	100	P	Br	No	2
<i>Halichoeres melanurus</i>	No	Mi	No	D	120	P	Br	No	1;2;3 4
<i>Halichoeres melasmapomus</i>	No	Mi	No	D	140	P	Br	No	1;2;3
<i>Halichoeres nebulosus</i>	No	N/A	No	D	120	P	Br	No	2;3
<i>Halichoeres nigrescens</i>	No	N/A	No	D	140	P	Br	No	2
<i>Halichoeres ornatissimus</i>	No	Mi	No	D	150	P	Br	No	1;2;3;4
<i>Halichoeres pallidus</i>	No	Mi	No	D	80	P	Br	No	1;2;12
<i>Halichoeres papilionaceus</i>	No	N/A	No	D	100	P	Br	No	2
<i>Halichoeres podostigma</i>	No	N/A	No	D	190	P	Br	No	2
<i>Halichoeres prosopeton</i>	No	N/A	No	D	130	P	Br	No	1;2;3
<i>Halichoeres richmondi</i>	No	Mi	No	D	190	P	Br	No	2
<i>Halichoeres rubricephalus</i>	No	N/A	No	D	100	P	Br	No	2
<i>Halichoeres scapularis</i>	No	Mi	No	D	200	P	Br	No	1;2;4
<i>Halichoeres solorensis</i>	No	Ma	No	D	180	P	Br	No	2;4
<i>Halichoeres trimaculatus</i>	No	Ma	No	D	270	P	Br	No	1;2;3 4
<i>Halichoeres trispilus</i>	No	N/A	No	D	90	P	Br	No	2
<i>Halichoeres vrolikii</i>	No	N/A	No	D	130	P	Br	No	2
<i>Halichoeres zeylonicus</i>	No	N/A	No	D	150	P	Br	No	2
<i>Hemigymnus fasciatus</i>	No	Ma	No	D	500	P	Br	No	1;2;3
<i>Hemigymnus melanopterus</i>	No	Ma	No	D	600	P	Br	No	1;2;3
<i>Hipposcarus harid</i>	No	De	No	D	750	B	Br	No	2
<i>Hipposcarus longiceps</i>	No	De	No	D	600	B	Br	No	1;2;3
<i>Hologymnosus annulatus</i>	No	Pi	No	D	400	P	Br	No	1;2;3
<i>Hologymnosus doliatus</i>	No	Ca	No	D	400	P	Br	No	1;2;3
<i>Hologymnosus longipes</i>	No	N/A	No	D	400	P	Br	No	1;3
<i>Hologymnosus rhodonotus</i>	No	N/A	No	D	320	P	Br	No	2
<i>Iniistius aneitensis</i>	No	Ma	No	D	240	P	Br	No	2;3;12
<i>Iniistius celebicus</i>	No	Ma	No	D	250	P	Br	No	2;3;12
<i>Iniistius pavo</i>	No	Ma	No	D	350	P	Br	No	2;3;4
<i>Iniistius pentadactylus</i>	No	Ma	No	D	250	P	Br	No	2;4
<i>Iniistius tetrazona</i>	No	Ma	No	D	250	P	Br	No	2
<i>Labrichthys unilineatus</i>	No	Co	No	D	160	P	Br	No	77
<i>Labroides bicolor</i>	Yes	Mi	No	D	140	P	Br	No	1;2;3;4
<i>Labroides dimidiatus</i>	Yes	Mi	No	D	115	P	Br	Yes	1;2;3;4
<i>Labroides pectoralis</i>	No	Mi	No	D	80	P	Br	No	1;2;3;4
<i>Labroides rubrolabiatus</i>	Yes	Mi	No	D	90	P	Br	No	2;3;12
<i>Labropsis alleni</i>	Yes	Co	No	D	100	P	Br	No	2;4
<i>Labropsis manabei</i>	No	Co	No	D	130	P	Br	No	77
<i>Labropsis micronesica</i>	No	Mi	No	D	130	P	Br	No	2;4;12
<i>Labropsis xanthonota</i>	No	Co	No	D	140	P	Br	No	1;2;3;12
<i>Leptojulius cyanopleura</i>	No	Pl	No	D	130	P	Br	No	2
<i>Leptoscarus vaigiensis</i>	No	He	No	D	350	P	Br	No	1;2;3;4
<i>Macropharyngodon choati</i>	No	Ma	No	D	100	P	Br	No	2
<i>Macropharyngodon kùteri</i>	No	Ma	No	D	100	P	Br	No	1;2;3

<i>Macropharyngodon meleagris</i>	No	Ma	No	D	150	P	Br	No	1;2;3;4
<i>Macropharyngodon negrosensis</i>	No	Ma	No	D	150	P	Br	No	1;2;3;4
<i>Macropharyngodon ornatus</i>	No	Ma	No	D	120	P	Br	No	2
<i>Novaculichthys macrolepidotus</i>	No	Ma	No	D	150	P	Br	No	2;3
<i>Novaculichthys taeniourus</i>	No	Ma	No	D	270	P	Br	No	1;2;3
<i>Oxycheilinus arenatus</i>	No	Ca	No	D	190	P	Br	No	2;1;4
<i>Oxycheilinus bimaculatus</i>	No	Ca	No	D	150	P	Br	No	2;1;12
<i>Oxycheilinus celebicus</i>	No	N/A	No	D	240	P	Br	No	2;1
<i>Oxycheilinus digrammus</i>	No	Pi	No	D	300	P	Br	No	2;1;12
<i>Oxycheilinus orientalis</i>	No	Ca	No	D	170	P	Br	No	2;4
<i>Oxycheilinus rhodochrous</i>	No	Ca	No	D	200	P	Br	No	2;4
<i>Oxycheilinus sp.</i>	No	Ca	No	D	140	P	Br	No	2;12
<i>Oxycheilinus unifasciatus</i>	No	Ca	No	D	460	P	Br	No	2;1;4
<i>Paracheilinus angulatus</i>	No	Pl	No	D	70	P	Br	No	2
<i>Paracheilinus carpenteri</i>	No	Pl	No	D	80	P	Br	No	2
<i>Paracheilinus filamentosus</i>	No	Pl	No	D	80	P	Br	No	2;4
<i>Paracheilinus flavianalis</i>	No	Pl	No	D	70	P	Br	No	2
<i>Paracheilinus mccoskeri</i>	No	Pl	No	D	70	P	Br	No	2
<i>Paracheilinus rubricaudalis</i>	No	Pl	No	D	80	P	Br	No	2;3
<i>Paracheilinus togeanensis</i>	No	Pl	No	D	80	P	Br	No	2
<i>Pseudocheilinus ataenia</i>	No	Mi	No	D	50	P	Br	No	2;4
<i>Pseudocheilinus evanidus</i>	No	Mi	No	D	80	P	Br	No	1;2;3;4
<i>Pseudocheilinus hexataenia</i>	No	Mi	No	D	75	P	Br	No	1;2;3;4
<i>Pseudocheilinus ocellatus</i>	No	Mi	No	D	85	P	Br	No	2;1
<i>Pseudocheilinus octotaenia</i>	No	Mi	No	D	135	P	Br	No	1;2;3;4
<i>Pseudocheilinus tetrataenia</i>	No	Mi	No	D	75	P	Br	No	2;3;4
<i>Pseudocoris aurantiofasciatus</i>	No	Pl	No	D	200	G	Br	No	2;3
<i>Pseudocoris bleekeri</i>	No	Pl	No	D	150	G	Br	No	2;4
<i>Pseudocoris heteroptera</i>	No	Pl	No	D	200	G	Br	No	2;4
<i>Pseudocoris yamashiroi</i>	No	Pl	No	D	150	G	Br	No	2;3
<i>Pseudodax moluccanus</i>	No	Ma	No	D	250	P	Br	No	1;2;3;12
<i>Pseudojuloides atavai</i>	No	Ma	No	D	130	P	Br	No	2;3;12
<i>Pseudojuloides cerasinus</i>	No	Ma	No	D	120	P	Br	No	1;2;3;12
<i>Pseudojuloides kaleidios</i>	No	N/A	No	D	100	P	Br	No	2
<i>Pseudojuloides severnsi</i>	No	Om	No	D	100	P	Br	No	2;4
<i>Pteragogus cryptus</i>	No	Mi	No	D	95	P	Br	Yes	1;2;3;4
<i>Pteragogus enneacanthus</i>	No	Ma	No	D	120	P	Br	No	2;3;12
<i>Pteragogus flagellifer</i>	No	N/A	No	D	200	P	Br	No	2
<i>Scarus altipinnis</i>	No	De	No	D	600	B	Br	No	1;2;3
<i>Scarus chameleon</i>	No	De	No	D	310	B	Br	No	1;2;3
<i>Scarus dimidiatus</i>	No	De	No	D	300	B	Br	No	1;2;3
<i>Scarus festivus</i>	No	De	No	D	430	B	Br	No	2;3
<i>Scarus flavipectoralis</i>	No	De	No	D	410	B	Br	No	1;2;3
<i>Scarus forsteni</i>	No	De	No	D	550	B	Br	No	1;2;3
<i>Scarus frenatus</i>	No	De	No	D	470	B	Br	No	1;2;3
<i>Scarus ghobban</i>	No	De	No	D	750	B	Br	No	1;2;3
<i>Scarus globiceps</i>	No	De	No	D	270	B	Br	No	1;2;3
<i>Scarus hypselopterus</i>	No	De	No	D	310	B	Br	No	2;3
<i>Scarus koputea</i>	N/A	De	No	D	310	B	Br	No	2;3
<i>Scarus longipinnis</i>	No	De	No	D	400	B	Br	No	1;2;3
<i>Scarus niger</i>	No	De	No	D	350	B	Br	No	1;2;3
<i>Scarus oviceps</i>	No	De	No	D	310	B	Br	No	1;2;3
<i>Scarus prasiognathus</i>	No	De	No	D	700	B	Br	No	2

	<i>Scarus psittacus</i>	No	De	No	D	300	B	Br	No	1;2;3
	<i>Scarus quoyi</i>	No	De	No	D	210	B	Br	No	2
	<i>Scarus rivulatus</i>	No	De	No	D	400	B	Br	No	1;2;3
	<i>Scarus rubroviolaceus</i>	Yes	De	No	D	700	B	Br	No	1;2;3
	<i>Scarus russellii</i>	No	De	No	D	510	B	Br	No	2
	<i>Scarus schlegeli</i>	No	De	No	D	380	B	Br	No	1;2;3
	<i>Scarus spinus</i>	No	De	No	D	300	B	Br	No	1;2;3
	<i>Scarus tricolor</i>	No	De	No	D	550	B	Br	No	1;2;3
	<i>Scarus viridifucatus</i>	No	De	No	D	320	B	Br	No	2
	<i>Scarus xanthopleura</i>	No	De	No	D	550	B	Br	No	2;3
	<i>Stethojulis bandanensis</i>	No	Mi	No	D	160	G	Br	No	1;2;3;4
	<i>Stethojulis interrupta</i>	No	Mi	No	D	130	G	Br	No	2;3;4
	<i>Stethojulis notialis</i>	No	Mi	No	D	100	G	Br	No	2;1
	<i>Stethojulis strigiventer</i>	No	Mi	No	D	140	G	Br	No	1;2;3
	<i>Stethojulis trilineata</i>	No	Mi	No	D	140	G	Br	No	2;3
	<i>Thalassoma amblycephalum</i>	No	Pl	No	D	140	G	Br	No	1;2;3;4
	<i>Thalassoma hardwicke</i>	No	Om	No	D	200	G	Br	No	1;2;3;4
	<i>Thalassoma janseni</i>	No	N/A	No	D	200	G	Br	No	2;3
	<i>Thalassoma lunare</i>	No	Pi	No	D	250	G	Br	No	1;2;3
	<i>Thalassoma lutescens</i>	No	Ma	No	D	250	G	Br	No	1;2;3;4
	<i>Thalassoma purpureum</i>	No	Ma	No	D	430	G	Br	No	1;2;3;4
	<i>Thalassoma quinquevittatum</i>	No	Ma	No	D	250	G	Br	No	1;2;3;4
	<i>Thalassoma trilobatum</i>	No	Ma	No	D	300	G	Br	No	1;2;3;4
	<i>Wetmorella albofasciata</i>	No	Ma	No	D	60	P	Br	No	1;2;3
	<i>Wetmorella nigropinnata</i>	No	Ma	No	D	80	P	Br	No	1;2;3;4
Lethrinidae	<i>Gnathodentex aureolineatus</i>	No	Ma	No	N	300	G	Br	No	2;3;4
	<i>Gymnocranius euanus</i>	No	Ma	No	D	450	G	Br	No	2;3
	<i>Gymnocranius frenatus</i>	No	Ma	No	D	350	G	Br	No	2;4
	<i>Gymnocranius grandoculis</i>	No	Ca	No	D	800	G	Br	No	1;2;3;4
	<i>Gymnocranius griseus</i>	No	Ma	No	D	350	G	Br	No	2;4
	<i>Gymnocranius microdon</i>	No	Ma	No	D	410	G	Br	No	2;4
	<i>Lethrinus amboinensis</i>	No	Ca	No	D	570	G	Br	No	2;4
	<i>Lethrinus atkinsoni</i>	No	Ca	No	D	410	G	Br	No	1;2;3
	<i>Lethrinus erythracanthus</i>	No	Ma	No	D	700	G	Br	No	44;1;2;3
	<i>Lethrinus erythropterus</i>	No	Ca	No	D	500	G	Br	No	2;4
	<i>Lethrinus genivittatus</i>	No	Ma	No	D	200	G	Br	No	1;2;3;95
	<i>Lethrinus harak</i>	No	Ca	No	D	500	G	Br	No	1;2;3;4
	<i>Lethrinus laticaudis</i>	No	Ca	No	D	560	G	Br	No	1;2;3;4
	<i>Lethrinus lentjan</i>	No	Ma	No	D	500	G	Br	No	1;2;3
	<i>Lethrinus microdon</i>	No	Ca	No	D	700	G	Br	No	2;4
	<i>Lethrinus miniatus</i>	No	Ca	No	D	900	G	Br	No	1;2;3
	<i>Lethrinus nebulosus</i>	No	Ca	No	D	800	G	Br	No	1;2;3
	<i>Lethrinus obsoletus</i>	No	Ma	No	D	500	G	Br	No	1;2;3
	<i>Lethrinus olivaceus</i>	No	Ca	No	D	1000	G	Br	No	44;1;2;3
	<i>Lethrinus ornatius</i>	No	Ca	No	D	400	G	Br	No	1;2;4
	<i>Lethrinus ravis</i>	No	N/A	No	D	320	G	Br	No	3
	<i>Lethrinus rubrioperculatus</i>	No	Ca	No	D	500	G	Br	No	1;2;3
	<i>Lethrinus semicinctus</i>	No	Ca	No	D	290	G	Br	No	2;4
	<i>Lethrinus variegatus</i>	No	Ma	No	D	200	G	Br	No	2;3
	<i>Lethrinus xanthochilus</i>	No	Ca	No	D	600	G	Br	No	44;1;2;3
	<i>Monotaxis grandoculis</i>	No	Ma	No	N	600	G	Br	No	44;1;2;3
Lutjanidae	<i>Aphareus furca</i>	No	Ca	No	D	400	G	Br	No	1;2;3
	<i>Aphareus rutilans</i>	No	Ca	No	D	1100	G	Br	No	3;4

<i>Aprion virescens</i>	No	Ca	No	D	1000	G	Br	No	44;1;2;3
<i>Lutjanus adetii</i>	No	Ca	No	N	500	G	Br	No	3;4
<i>Lutjanus argentimaculatus</i>	No	Ca	No	D	1200	G	Br	No	1;2;3
<i>Lutjanus bengalensis</i>	No	Ca	No	D	300	G	Br	No	2
<i>Lutjanus biguttatus</i>	No	Ca	No	D	200	G	Br	No	2;3;4
<i>Lutjanus bohar</i>	No	Pi	No	D	750	G	Br	No	44;1;2;3
<i>Lutjanus bouton</i>	No	Ca	No	D	280	G	Br	No	2;4
<i>Lutjanus carponotatus</i>	No	Ca	No	D	400	G	Br	No	1;96;2;3
<i>Lutjanus decussatus</i>	No	Ca	No	D	300	G	Br	No	1;2;97
<i>Lutjanus ehrenbergii</i>	No	Ca	No	D	350	G	Br	No	1;2;3
<i>Lutjanus fulviflamma</i>	No	Ca	No	N	350	G	Br	No	1;96;2;3;
<i>Lutjanus fulvus</i>	No	Ca	No	D	400	G	Br	No	44;1;2;3
<i>Lutjanus gibbus</i>	No	Ca	No	N	500	G	Br	No	1;2;3
<i>Lutjanus johnii</i>	No	Ca	No	D	700	G	Br	No	98;3;4
<i>Lutjanus kasmira</i>	No	Ca	No	D	350	G	Br	No	1;2;3
<i>Lutjanus lemniscatus</i>	No	Ca	No	D	650	G	Br	No	1;2;4
<i>Lutjanus lunulatus</i>	No	Ca	No	D	350	G	Br	No	2
<i>Lutjanus lutjanus</i>	No	Ca	No	D	300	G	Br	No	1;2;3;4
<i>Lutjanus madras</i>	No	Ca	No	D	300	G	Br	No	2;24
<i>Lutjanus malabaricus</i>	No	Ca	No	N	1000	G	Br	No	2;4
<i>Lutjanus maxweberi</i>	No	Ca	No	D	800	G	Br	No	2
<i>Lutjanus monostigma</i>	No	Pi	No	D	600	G	Br	No	44;1;2;3
<i>Lutjanus quinquelineatus</i>	No	Ca	No	D	390	G	Br	No	99;1;2;3
<i>Lutjanus rivulatus</i>	No	Pi	No	D	800	G	Br	No	3
<i>Lutjanus rufolineatus</i>	No	Ca	No	D	280	G	Br	No	2;3;4
<i>Lutjanus russelli</i>	No	Ca	No	D	280	G	Br	No	100;1;2;3
<i>Lutjanus sebae</i>	No	Ca	No	D	600	G	Br	No	1;2;3;4
<i>Lutjanus semicinctus</i>	No	Pi	No	D	350	G	Br	No	1;2;4
<i>Lutjanus timorensis</i>	No	Pi	No	D	500	G	Br	No	2;101;4
<i>Lutjanus vitta</i>	No	Ca	No	D	400	G	Br	No	1;2;3;39
<i>Macolor macularis</i>	No	Pl	No	N	600	G	Br	No	1;2;3;4
<i>Macolor niger</i>	No	Ca	No	D	600	G	Br	No	1;2;3;4
<i>Paracaesio sordida</i>	No	Pl	No	D	400	G	Br	No	2;3;4
<i>Paracaesio xanthura</i>	No	Pl	No	D	400	G	Br	No	2;3
<i>Pinjalo lewisi</i>	No	Ma	No	D	500	G	Br	No	2;3;4
<i>Pinjalo pinjalo</i>	No	Ma	No	D	500	G	Br	No	2;3;4
<i>Symphoricthys spilurus</i>	No	Ma	No	D	600	G	Br	No	1;2;3
<i>Symphorus nematophorus</i>	No	Pi	No	D	800	G	Br	No	1;2;3
Malacanthidae									
<i>Hoplolatilus chlupaty</i>	Yes	Pl	Yes	D	150	P	Br	Yes	2;102
<i>Hoplolatilus cuniculus</i>	Yes	Pl	Yes	D	150	P	Br	Yes	1;2;3;102
<i>Hoplolatilus fronticinctus</i>	Yes	Pl	Yes	D	170	P	Br	Yes	2;102
<i>Hoplolatilus luteus</i>	Yes	Pl	Yes	D	110	P	Br	Yes	2;102
<i>Hoplolatilus marcosi</i>	Yes	Pl	Yes	D	150	P	Br	Yes	2;102
<i>Hoplolatilus purpureus</i>	Yes	Pl	Yes	D	120	P	Br	Yes	2;102
<i>Hoplolatilus starcki</i>	Yes	Pl	Yes	D	150	P	Br	Yes	1;2;102
<i>Malacanthus brevirostris</i>	Yes	Ma	Yes	D	300	P	Br	No	1;2;3;103
<i>Malacanthus latovittatus</i>	Yes	Ma	Yes	D	350	P	Br	Yes	1;2;3;103
Microcanthidae									
<i>Microcanthus strigatus</i>	No	Om	No	D	160	N/A	Br	No	1;2;3
<i>Gunnelichthys curiosus</i>	Yes	Mi	Yes	D	120	N/A	Br	No	1;2;3;4
<i>Gunnelichthys monostigma</i>	Yes	Mi	Yes	D	110	N/A	Br	No	1;2;3;4
<i>Gunnelichthys pleurotaenia</i>	Yes	Mi	Yes	D	90	N/A	Br	No	1;2;3;4
<i>Gunnelichthys viridescens</i>	Yes	Mi	Yes	D	100	N/A	Br	No	1;2;3;4
Molidae									
<i>Mola mola</i>	No	Ca	No	D	3080	N/A	Br	No	4

Monacanthidae	<i>Acreichthys radiatus</i>	No	Co	No	D	70	P	Dm	No	4
	<i>Acreichthys tomentosus</i>	No	Ma	No	D	100	P	Dm	No	4
	<i>Aluterus monoceros</i>	No	Om	No	D	750	P	Dm	No	104
	<i>Aluterus scriptus</i>	No	Om	No	D	750	P	Dm	No	1;2;3
	<i>Amanses scopas</i>	Yes	Co	No	D	200	G	Br	No	105
	<i>Anacanthus barbatus</i>	No	N/A	No	D	350	P	Dm	No	1;2;3
	<i>Cantherhines dumerili</i>	Yes	Co	No	D	350	P	Dm	No	77
	<i>Cantherhines fronticinctus</i>	No	Ma	No	D	230	P	Dm	No	4
	<i>Cantherhines pardalis</i>	No	Ma	No	D	250	P	Dm	No	4
	<i>Cantherhines sandwichiensis</i>	No	Om	No	D	194	P	Dm	No	1;2;3
	<i>Chaetodermis penicilligera</i>	No	N/A	No	D	310	P	Dm	No	1;2;3
	<i>Monacanthus chinensis</i>	No	Om	No	D	380	P	Dm	No	4
	<i>Oxymonacanthus longirostris</i>	Yes	Co	No	D	90	P	Dm	Yes	77
	<i>Paraluteres arqat</i>	No	N/A	No	D	80	P	Dm	No	1;2;3
	<i>Paraluteres prionurus</i>	Yes	Om	No	D	100	P	Dm	No	4
	<i>Paramonacanthus choirocephalus</i>	No	Om	No	D	130	P	Dm	No	24
	<i>Paramonacanthus japonicus</i>	Yes	Mi	No	D	100	P	Dm	Yes	106
	<i>Pervagor alternans</i>	No	Mi	No	D	160	P	Dm	No	12
	<i>Pervagor aspricaudus</i>	No	Mi	No	D	120	P	Dm	No	12
	<i>Pervagor janthinosoma</i>	Yes	Mi	No	D	140	P	Dm	No	4
	<i>Pervagor melanocephalus</i>	Yes	Mi	No	D	100	P	Dm	No	12
	<i>Pervagor nigrolineatus</i>	No	Mi	No	D	100	P	Dm	No	12
	<i>Pseudalutarius nascicornis</i>	Yes	Mi	No	D	180	P	Dm	No	107
<i>Pseudomonacanthus macrurus</i>	Yes	Om	No	D	240	P	Dm	No	21	
<i>Rudarias excelsus</i>	N/A	N/A	No	D	25	G	Dm	No	1;2;3	
<i>Rudarias minutus</i>	No	N/A	No	D	30	G	Dm	No	1;2;3	
Monocentridae	<i>Cleidopus gloriamaris</i>	Yes	N/A	No	D	220	N/A	Br	No	1;2;3
	<i>Monocentris japonica</i>	Yes	N/A	No	D	170	N/A	Br	No	1;2;3
Mugilidae	<i>Chelon macrolepis</i>	No	De	No	D	400	G	Br	No	3;15
	<i>Chelon melinopterus</i>	No	De	No	D	300	G	Br	No	3;15
	<i>Chelon subviridis</i>	No	De	No	D	400	G	Br	No	3;4
	<i>Crenimugil crenilabis</i>	No	De	No	D	400	G	Br	No	2;3;108
	<i>Ellochelon vaigiensis</i>	No	Om	No	D	520	G	Br	No	2;3;15
	<i>Moolgarda engeli</i>	No	De	No	D	150	G	Br	No	2;108
	<i>Moolgarda seheli</i>	No	De	No	D	500	G	Br	No	3;4
	<i>Mugil cephalus</i>	No	De	No	D	300	G	Br	No	3;15
	<i>Neomyxus leuciscus</i>	No	Om	No	D	460	G	Br	No	2;3;4
Mullidae	<i>Mulloidichthys flavolineatus</i>	No	Ma	No	N	400	G	Br	No	1;2;3;4
	<i>Mulloidichthys mimicus</i>	No	Ma	No	N	300	G	Br	No	1;2;3
	<i>Mulloidichthys pflugeri</i>	Yes	Ma	No	D	480	G	Br	No	1;2;3;
	<i>Mulloidichthys vanicolensis</i>	No	Ma	No	N	380	G	Br	No	1;2;3;4
	<i>Parupeneus barberinoides</i>	No	Ma	No	D	250	G	Br	No	1;2;3;4
	<i>Parupeneus barberinus</i>	No	Ma	No	D	500	G	Br	No	1;2;3;4
	<i>Parupeneus ciliatus</i>	No	Ma	No	N	380	G	Br	No	1;2;3;4
	<i>Parupeneus crassilabris</i>	No	Ma	No	D	350	G	Br	No	1;2;3
	<i>Parupeneus cyclostomus</i>	Yes	Ca	No	D	500	P	Br	No	1;2;3;4
	<i>Parupeneus heptacanthus</i>	No	Ma	No	D	360	P	Br	No	1;2;3;109
	<i>Parupeneus indicus</i>	No	Ma	No	N	350	P	Br	No	1;2;3;4
	<i>Parupeneus insularis</i>	No	Ma	No	D	330	P	Br	No	1;2;3;110
	<i>Parupeneus macronemus</i>	No	Ma	No	D	320	P	Br	No	1;2;3;111
	<i>Parupeneus multifasciatus</i>	No	Ma	No	D	300	P	Br	No	1;2;3;109
	<i>Parupeneus pleurostigma</i>	No	Ma	No	D	330	P	Br	No	1;2;3;109
	<i>Parupeneus rubescens</i>	No	Ma	No	D	430	P	Br	No	1;2;3;112

	<i>Parupeneus spilurus</i>	No	Ma	No	N	350	P	Br	No	1;2;3
	<i>Parupeneus trifasciatus</i>	No	Ca	No	N	350	P	Br	No	1;2;3;4
	<i>Upeneus moluccensis</i>	No	Ca	No	D	200	P	Br	No	113
	<i>Upeneus taeniopterus</i>	No	Ca	No	D	360	P	Br	No	2
	<i>Upeneus tragula</i>	No	Ca	No	D	300	P	Br	No	114
	<i>Upeneus vittatus</i>	No	Ca	No	D	280	P	Br	No	114
Muraenidae	<i>Channomuraena vittata</i>	No	N/A	No	N	1500	P	Br	No	4
	<i>Echidna delicatula</i>	No	Ma	No	N	650	P	Br	No	115
	<i>Echidna nebulosa</i>	No	Ma	No	N	750	P	Br	No	115
	<i>Echidna polyzona</i>	No	Ma	No	N	600	P	Br	No	115
	<i>Enchelycore bayeri</i>	No	Pi	No	N	600	P	Br	No	115
	<i>Enchelycore lichenosa</i>	No	Pi	No	N	925	P	Br	No	4
	<i>Enchelycore pardalis</i>	No	Pi	No	N	800	P	Br	No	12
	<i>Enchelycore schismatorhynchus</i>	No	Pi	No	N	1200	P	Br	No	4
	<i>Enchelynassa canina</i>	No	Ca	No	N	1540	P	Br	No	4
	<i>Gymnomuraena zebra</i>	No	Ma	No	D	1540	P	Br	No	116
	<i>Gymnothorax albimarginatus</i>	No	N/A	No	N	1000	P	Br	No	4
	<i>Gymnothorax breedeni</i>	No	Pi	No	N	750	P	Br	No	115
	<i>Gymnothorax buroensis</i>	No	N/A	No	N	330	P	Br	No	1;2;3
	<i>Gymnothorax chilospilus</i>	No	N/A	No	D	500	P	Br	No	1;2;3
	<i>Gymnothorax chlamydatum</i>	No	N/A	No	D	800	P	Br	No	1;2;3
	<i>Gymnothorax cribroris</i>	No	N/A	No	N	470	P	Br	No	1;2;3
	<i>Gymnothorax enigmaticus</i>	No	N/A	No	N	580	P	Br	No	1;2;3
	<i>Gymnothorax eurostus</i>	No	Ca	No	D	650	P	Br	No	3
	<i>Gymnothorax favagineus</i>	No	Ca	No	D	1800	P	Br	No	4
	<i>Gymnothorax fimbriatus</i>	No	Ca	No	N	800	P	Br	No	4
	<i>Gymnothorax flavimarginatus</i>	No	Ca	No	N	1500	P	Br	No	3
	<i>Gymnothorax gracilicauda</i>	No	N/A	No	N	320	P	Br	No	1;2;3
	<i>Gymnothorax herrei</i>	No	N/A	No	D	300	G	Br	No	1;2;3
	<i>Gymnothorax isingteena</i>	No	N/A	No	D	1800	P	Br	No	1;2;3
	<i>Gymnothorax javanicus</i>	No	Pi	No	N	2390	P	Br	No	3
	<i>Gymnothorax melatremus</i>	No	Ma	No	N	200	P	Br	No	12
	<i>Gymnothorax meleagris</i>	No	Pi	No	N	1200	P	Br	No	3
	<i>Gymnothorax minor</i>	No	N/A	No	D	600	P	Br	No	1;2;3
	<i>Gymnothorax nudivomer</i>	No	N/A	No	D	1800	P	Br	No	1;2;3
	<i>Gymnothorax picta</i>	No	Ca	No	D	1200	P	Br	No	4
	<i>Gymnothorax richardsoni</i>	No	N/A	No	D	320	P	Br	No	1;2;3
	<i>Gymnothorax rueppelliae</i>	No	Ca	No	N	800	P	Br	No	12
	<i>Gymnothorax tile</i>	No	N/A	No	D	530	P	Br	No	1;2;3
	<i>Gymnothorax undulatus</i>	No	Ca	No	N	1500	P	Br	No	4
<i>Gymnothorax zonipectis</i>	No	N/A	No	N	800	P	Br	No	1;2;3	
<i>Pseudechidna brummeri</i>	No	N/A	No	N	1030	P	Br	No	1;2;3	
<i>Rhinomuraena quaesita</i>	Yes	Pi	No	D	850	P	Br	No	115	
<i>Scuticaria okinawae</i>	No	N/A	No	D	930	P	Br	No	1;2;3	
<i>Scuticaria tigrina</i>	No	N/A	No	N	1200	P	Br	No	1;2;3	
<i>Siderea thysioidea</i>	Yes	N/A	No	D	650	P	Br	No	4	
<i>Strophidon sathete</i>	No	Ca	No	D	3750	P	Br	No	4	
<i>Uropterygius fasciolatus</i>	No	Pi	No	D	530	P	Br	No	116	
<i>Uropterygius macrocephalus</i>	No	Pi	No	D	400	P	Br	No	116	
Nemipteridae	<i>Nemipterus furcosus</i>	No	Ca	No	D	300	G	Br	No	100;2;3
	<i>Nemipterus peronii</i>	No	Ca	No	D	280	G	Br	No	3;4
	<i>Nemipterus zysron</i>	No	Ca	No	D	250	G	Br	No	3;4
	<i>Pentapodus aureofasciatus</i>	Yes	Ca	No	D	250	G	Br	No	2;3

	<i>Pentapodus bifasciatus</i>	No	Ca	No	D	200	G	Br	No	2;4
	<i>Pentapodus caninus</i>	No	Ca	No	D	250	G	Br	No	2;3
	<i>Pentapodus emeryii</i>	No	Ca	No	D	350	G	Br	No	2;4
	<i>Pentapodus nagasakiensis</i>	No	Ma	No	D	250	G	Br	No	2;4
	<i>Pentapodus paradiseus</i>	No	Ca	No	D	350	G	Br	No	1;2;3
	<i>Pentapodus porosus</i>	No	Ca	No	D	300	G	Br	No	2
	<i>Pentapodus setosus</i>	No	Ma	No	D	250	G	Br	No	2;4
	<i>Pentapodus trivittatus</i>	Yes	Ca	No	D	300	G	Br	No	2;4
	<i>Scaevius milii</i>	No	Ca	No	D	300	G	Br	No	2;4
	<i>Scaevius vitta</i>	No	Ca	No	D	400	G	Br	No	2;3
	<i>Scolopsis affinis</i>	No	Ma	No	D	300	G	Br	No	2;117
	<i>Scolopsis auratus</i>	No	N/A	No	D	300	G	Br	No	2
	<i>Scolopsis bilineatus</i>	No	Ma	No	N	250	G	Br	No	1;2;3;118
	<i>Scolopsis ciliatus</i>	No	Ca	No	D	250	G	Br	No	2;3;4
	<i>Scolopsis ghanam</i>	No	Ca	No	D	250	G	Br	No	2;4
	<i>Scolopsis lineatus</i>	No	Ca	No	D	250	G	Br	No	1;2;3;4
	<i>Scolopsis margaritifera</i>	No	Ca	No	D	250	G	Br	No	1;119;2
	<i>Scolopsis monogramma</i>	No	Ca	No	D	380	G	Br	No	2;4
	<i>Scolopsis taeniopterus</i>	No	Ca	No	D	300	G	Br	No	2;4
	<i>Scolopsis trilineatus</i>	No	Ma	No	D	250	G	Br	No	1;2;3;20
	<i>Scolopsis vosmeri</i>	No	Ma	No	D	250	G	Br	No	2;4
	<i>Scolopsis xenochrous</i>	No	Ma	No	D	250	G	Br	No	2;4
Ophichthidae	<i>Apterichtus klazingai</i>	No	Ca	No	D	400	G	Br	No	4
	<i>Brachysomophis cirrocheilos</i>	No	Ca	No	D	1250	G	Br	No	4
	<i>Brachysomophis crocodilinus</i>	No	Ca	No	N	820	G	Br	No	120
	<i>Brachysomophis henshawi</i>	No	Ca	No	N	1060	G	Br	No	4
	<i>Callechelys catostoma</i>	No	Ca	No	D	850	G	Br	No	1;2;3
	<i>Callechelys marmorata</i>	No	Ca	No	D	900	G	Br	No	1;2;3
	<i>Leiuranus semicinctus</i>	No	Ca	No	N	600	G	Br	No	4
	<i>Leiuranus versicolor</i>	No	N/A	No	D	520	G	Br	No	1;2;3
	<i>Myrichthys colubrinus</i>	No	N/A	No	D	900	G	Br	No	1;2;3
	<i>Myrichthys maculosus</i>	No	N/A	No	N	1000	G	Br	No	4
	<i>Ophichthus bonaparti</i>	No	N/A	No	N	750	G	Br	No	4
	<i>Ophichthus cephalozona</i>	No	N/A	No	N	1080	G	Br	No	1;2;3
	<i>Ophichthus melanochir</i>	No	N/A	No	D	800	G	Br	No	4
	<i>Ophichthus polyophthalmus</i>	No	N/A	No	D	350	G	Br	No	4
	<i>Pisodonophis cancrivorus</i>	Yes	Ma	No	D	750	G	Br	No	121
Opistognathidae	<i>Opistognathus darwiniensis</i>	No	Pl	No	D	450	P	Mo	No	2
	<i>Opistognathus dendriticus</i>	No	Pl	No	D	450	P	Mo	No	1;2
	<i>Opistognathus papuensis</i>	No	Pl	No	D	450	P	Mo	No	1;2
	<i>Opistognathus solorensis</i>	No	Pl	No	D	450	P	Mo	No	1;2
Ostraciidae	<i>Lactoria cornuta</i>	No	Ma	No	D	460	P	Br	No	1;2;3;4
	<i>Lactoria diaphana</i>	No	Ma	No	D	250	P	Br	No	2;122
	<i>Lactoria formasini</i>	No	Ma	No	D	150	P	Br	No	1;2;3;122
	<i>Ostracion cubicus</i>	No	Om	No	D	450	P	Br	No	1;2;3;4
	<i>Ostracion meleagris</i>	Yes	Om	No	D	180	P	Br	No	1;2;3;4
	<i>Ostracion rhinorhynchus</i>	Yes	Ma	No	D	300	P	Br	No	2;4
	<i>Ostracion solorensis</i>	Yes	N/A	No	D	110	P	Br	No	1;2;3
	<i>Ostracion whitleyi</i>	Yes	Ma	No	D	155	P	Br	No	1;2;3;12
	<i>Rhynchostracion nasus</i>	No	N/A	No	D	300	P	Br	No	2
	<i>Tetrosomus concatenatus</i>	No	N/A	No	D	300	P	Br	No	2
	<i>Tetrosomus gibbosus</i>	No	Ma	No	D	300	P	Br	No	1;2;3;4
Pegasidae	<i>Eurypegasus draconis</i>	Yes	Mi	No	D	70	P	Br	Yes	1;2;3;4

Appendix C

	<i>Pegasus volitans</i>	Yes	Mi	No	D	110	P	Br	No	1;2;3;4
Pempheridae	<i>Parapriacanthus ransonneti</i>	No	Pl	No	N	100	G	Br	No	4
	<i>Pempheris adusta</i>	No	Pl	No	N	170	G	Br	No	4
	<i>Pempheris oualensis</i>	No	Pl	No	N	220	G	Br	No	4
	<i>Pempheris schwenkii</i>	No	Pl	No	N	150	G	Br	No	4
	<i>Pempheris vanicolensis</i>	No	Pl	No	N	200	G	Br	No	4
Pinguipedidae	<i>Parapercis clathrata</i>	No	Ma	No	D	175	P	Br	No	1;2;3;12
	<i>Parapercis cylindrica</i>	No	Ma	No	D	230	P	Br	No	2;88
	<i>Parapercis hexophthalma</i>	No	Ma	No	D	280	P	Br	No	1;2;3
	<i>Parapercis lata</i>	No	Ma	No	D	260	P	Br	No	2;3;
	<i>Parapercis lineopunctata</i>	No	Ma	No	D	120	P	Br	No	2
	<i>Parapercis maculatus</i>	No	Ma	No	D	200	P	Br	No	2
	<i>Parapercis millipunctata</i>	No	Ma	No	D	260	P	Br	No	1;2;3
	<i>Parapercis multiplicata</i>	No	Ma	No	D	120	P	Br	No	1;2;3
	<i>Parapercis schauinslandi</i>	No	Pl	No	D	130	P	Br	No	1;2;3;4
	<i>Parapercis snyderi</i>	No	Ma	No	D	110	P	Br	No	1;2;3
	<i>Parapercis tetracantha</i>	No	Ma	No	D	260	P	Br	No	2
	<i>Parapercis xanthozona</i>	No	Ma	No	D	230	P	Br	No	1;2;3
Platycephalidae	<i>Cociella punctata</i>	No	Pl	No	D	350	P	Br	No	2;14
	<i>Cymbacephalus beauforti</i>	No	Pi	No	D	470	P	Br	No	2;123
	<i>Inegocia japonica</i>	No	N/A	No	D	250	P	Br	No	2
	<i>Onigocia spinosa</i>	No	N/A	No	N	130	P	Br	No	2
	<i>Rogadius patriciae</i>	No	N/A	No	D	270	P	Br	No	2
	<i>Rogadius pristiger</i>	No	N/A	No	D	210	P	Br	No	2
	<i>Rogadius welanderi</i>	No	N/A	No	D	130	P	Br	No	2
	<i>Thysanophrys arenicola</i>	No	Ca	No	D	370	P	Br	No	1;2;3;4
	<i>Thysanophrys carbunculus</i>	No	Pi	No	D	400	P	Br	No	2;124
	<i>Thysanophrys chiltonae</i>	No	Ca	No	D	220	P	Br	No	1;2;3;4
	<i>Thysanophrys otaitensis</i>	No	Ca	No	D	250	P	Br	No	1;2;3
Plesiopidae	<i>Assessor flavissimus</i>	No	Pl	No	D	55	G	Mo	No	1;2
	<i>Assessor macneilli</i>	No	Om	No	D	60	G	Mo	No	1;2;3;70
	<i>Assessor randalli</i>	No	Pl	No	D	60	G	Mo	No	2
	<i>Belonopterygium fasciolatum</i>	No	N/A	No	D	50	G	Dm	No	2
	<i>Calloplesiops ativelis</i>	No	N/A	No	D	130	G	Dm	No	1;2;3
	<i>Paraplesiops poweri</i>	No	Ma	No	D	149	G	Dm	No	1;125
	<i>Plesiops coeruleolineatus</i>	No	Ca	No	N	80	G	Dm	No	1;2;3;4
	<i>Plesiops corallicola</i>	No	Ma	No	N	160	G	Dm	No	1;2;3;125
	<i>Plesiops insularis</i>	No	Ma	No	D	149	G	Dm	No	1;125
		<i>Plesiops verecundus</i>	No	Ma	No	D	149	G	Dm	No
Pleuronectidae	<i>Pseudorhombus duplioniocellatus</i>	No	Ca	No	D	400	P	Br	No	2
	<i>Samaris cristatus</i>	No	Ma	No	D	220	P	Br	No	1;2;3;4
	<i>Samariscus triocellatus</i>	No	Ma	No	N	90	P	Br	No	1;2;3;4
Plotosidae	<i>Paraplotosus albilabris</i>	No	Ma	No	N	1300	P	Dm	No	4
	<i>Plotosus lineatus</i>	No	Ca	No	N	320	P	Dm	No	4
Pomacanthidae	<i>Apolemichthys griffisi</i>	Yes	Sp	No	D	250	P	Dm	No	2;4
	<i>Apolemichthys trimaculatus</i>	Yes	Sp	No	D	250	P	Br	No	1;2;3
	<i>Apolemichthys xanthopunctatus</i>	No	Sp	No	D	250	P	Br	No	2;4
	<i>Centropyge aurantius</i>	No	N/A	No	D	100	P	Br	No	2
	<i>Centropyge bicolor</i>	Yes	Om	No	D	150	P	Br	No	2;126
	<i>Centropyge bispinosus</i>	No	He	No	D	100	P	Br	No	53;2;4
	<i>Centropyge boylei</i>	N/A	He	No	D	70	P	Br	No	3;4
	<i>Centropyge colini</i>	No	He	No	D	90	P	Br	No	1;2;3
	<i>Centropyge eibli</i>	Yes	He	No	D	110	P	Br	No	2;4

<i>Centropyge ferrugatus</i>	No	He	No	D	100	P	Br	No	53;2
<i>Centropyge fisheri</i>	No	He	No	D	76	P	Br	No	2
<i>Centropyge flavipectoralis</i>	No	He	No	D	100	P	Br	No	2
<i>Centropyge flavissimus</i>	No	He	No	D	100	P	Br	No	127;2
<i>Centropyge heraldi</i>	No	He	No	D	100	P	Br	No	127;2
<i>Centropyge hotumatua</i>	No	He	No	D	80	P	Br	No	2;3
<i>Centropyge interruptus</i>	No	He	No	D	150	P	Br	Yes	127
<i>Centropyge loriculus</i>	No	He	No	D	100	P	Br	No	2;3;4
<i>Centropyge multicolor</i>	No	He	No	D	90	P	Br	No	2
<i>Centropyge multifasciata</i>	No	He	No	D	100	P	Br	No	1;2;3
<i>Centropyge multispinis</i>	No	He	No	D	100	P	Br	No	2
<i>Centropyge narcosis</i>	Yes	He	No	D	72	P	Br	No	3;4
<i>Centropyge nigriocellus</i>	No	He	No	D	60	P	Br	No	2;4
<i>Centropyge nox</i>	Yes	He	No	D	90	P	Br	No	2;4
<i>Centropyge shepardi</i>	No	He	No	D	120	P	Br	No	53;2;4
<i>Centropyge tibicen</i>	No	He	No	D	180	P	Br	No	127;2
<i>Centropyge venusta</i>	Yes	He	No	D	120	P	Br	No	128
<i>Centropyge vrolikii</i>	No	Om	No	D	120	P	Br	Yes	53;2
<i>Chaetodontoplus ballinae</i>	No	Sp	No	D	200	P	Br	No	53;2
<i>Chaetodontoplus chrysocephalus</i>	No	Sp	No	D	220	P	Br	No	53;2;4
<i>Chaetodontoplus conspicillatus</i>	Yes	Sp	No	D	250	P	Br	No	1;53;2;3
<i>Chaetodontoplus dimidiatus</i>	Yes	Sp	No	D	200	P	Br	No	2
<i>Chaetodontoplus duboulayi</i>	Yes	Sp	No	D	250	P	Br	No	53;2
<i>Chaetodontoplus melanosoma</i>	Yes	Sp	No	D	200	P	Br	No	53;2;4
<i>Chaetodontoplus meredithi</i>	Yes	Sp	No	D	250	P	Br	No	53;2
<i>Chaetodontoplus mesoleucus</i>	Yes	Om	No	D	180	P	Br	No	53;2
<i>Chaetodontoplus personifer</i>	N/A	N/A	No	D	350	P	Br	No	2
<i>Chaetodontoplus septentrionalis</i>	No	Sp	No	D	200	P	Br	No	53;2;4
<i>Genicanthus bellus</i>	No	Pl	No	D	180	P	Br	No	1;53;2;3
<i>Genicanthus caudovittatus</i>	No	Pl	No	D	200	P	Br	No	53;2;3
<i>Genicanthus lamarck</i>	No	Pl	No	D	230	P	Br	No	53;2
<i>Genicanthus melanospilos</i>	Yes	Pl	No	D	180	P	Br	No	53;2;3
<i>Genicanthus semicinctus</i>	No	Pl	No	D	250	P	Br	No	53;2
<i>Genicanthus semifasciatus</i>	No	Pl	No	D	210	P	Br	No	53;2
<i>Genicanthus spinus</i>	No	Pl	No	D	375	P	Br	No	53;3
<i>Genicanthus watanabei</i>	No	Pl	No	D	150	P	Br	No	53;2
<i>Pomacanthus annularis</i>	Yes	Sp	No	D	450	P	Br	No	53;128;2
<i>Pomacanthus imperator</i>	Yes	Sp	No	D	380	P	Br	No	53;2;3
<i>Pomacanthus navarchus</i>	No	Sp	No	D	380	P	Br	No	53;2
<i>Pomacanthus semicirculatus</i>	No	Om	No	D	350	P	Br	No	53;2;3
<i>Pomacanthus sexstriatus</i>	Yes	Om	No	D	460	P	Br	No	53;2;129
<i>Pomacanthus xanthometopon</i>	No	Sp	No	D	380	P	Br	No	53;2
<i>Pygoplites diacanthus</i>	Yes	Sp	No	D	250	P	Br	No	53;2;3
Pomacentridae									
<i>Abudefduf bengalensis</i>	No	Om	No	D	170	G	Dm	No	1;2;4
<i>Abudefduf conformis</i>	N/A	N/A	No	D	170	G	Dm	No	3
<i>Abudefduf lorentzi</i>	No	He	No	D	150	G	Dm	No	130;2
<i>Abudefduf notatus</i>	No	N/A	No	D	150	G	Dm	No	2;4
<i>Abudefduf septemfasciatus</i>	No	Om	No	D	190	G	Dm	No	1;2;3
<i>Abudefduf sexfasciatus</i>	No	Pl	No	D	150	G	Dm	No	1;2;3;131
<i>Abudefduf sordidus</i>	No	Om	No	D	190	G	Dm	No	1;2;3
<i>Abudefduf vaigiensis</i>	No	Pl	No	D	190	G	Dm	No	1;2;3;131
<i>Abudefduf whitleyi</i>	No	Pl	No	D	150	G	Dm	No	132;1;2;3
<i>Acanthochromis polyacanthus</i>	Yes	Pl	No	D	130	P	Dm	Yes	1;2;137

<i>Altrichthys azurelineatus</i>	Yes	Pl	No	D	60	P	Dm	No	2;4
<i>Altrichthys curatus</i>	Yes	Pl	No	D	60	P	Dm	No	2;4
<i>Amblyglyphidodon aureus</i>	Yes	Pl	No	D	150	P	Dm	No	1;2;3
<i>Amblyglyphidodon batunai</i>	Yes	Om	No	D	100	P	Dm	No	2;4
<i>Amblyglyphidodon curacao</i>	No	Pl	No	D	115	P	Dm	No	132;1;2;3
<i>Amblyglyphidodon leucogaster</i>	No	Om	No	D	130	P	Dm	No	1;2;4
<i>Amblyglyphidodon melanopterus</i>	No	Om	No	D	140	P	Dm	No	3
<i>Amblyglyphidodon orbicularis</i>	Yes	Om	No	D	130	P	Dm	No	2;3
<i>Amblyglyphidodon ternatensis</i>	No	Om	No	D	120	P	Dm	No	2
<i>Amblypomacentrus breviceps</i>	No	N/A	No	D	80	P	Dm	No	1;2;4
<i>Amphiprion akallopisos</i>	No	Pl	No	D	100	P	Dm	No	133;2;131;4
<i>Amphiprion akindynos</i>	No	Pl	No	D	120	P	Dm	No	133;3
<i>Amphiprion chrysopterus</i>	No	Om	No	D	170	P	Dm	No	1;2;3;4
<i>Amphiprion clarkii</i>	No	Om	No	D	120	P	Dm	Yes	1;2;3;4
<i>Amphiprion ephippium</i>	Yes	Pl	No	D	120	P	Dm	No	2;4
<i>Amphiprion frenatus</i>	No	Pl	No	D	60	P	Dm	Yes	133;2;4
<i>Amphiprion latezonatus</i>	No	Pl	No	D	140	P	Dm	No	133;2;4
<i>Amphiprion leucokranos</i>	No	Pl	No	D	130	P	Dm	No	133;2;4
<i>Amphiprion mccullochi</i>	No	Pl	No	D	120	P	Dm	No	133;2;4
<i>Amphiprion melanopus</i>	Yes	Om	No	D	130	P	Dm	No	1;2;3;4
<i>Amphiprion ocellaris</i>	No	Pl	No	D	90	P	Dm	No	134;2
<i>Amphiprion percula</i>	No	Pl	No	D	90	P	Dm	No	1;2;4
<i>Amphiprion perideraion</i>	No	Pl	No	D	100	P	Dm	Yes	1;133;2;3;4
<i>Amphiprion polymnus</i>	No	Pl	No	D	120	P	Dm	No	133;2;4
<i>Amphiprion rubrocinctus</i>	No	Pl	No	D	120	P	Dm	No	2;4
<i>Amphiprion sandaracinos</i>	Yes	Pl	No	D	130	P	Dm	No	133;2;4
<i>Amphiprion sebae</i>	No	Pl	No	D	140	P	Dm	No	133;2;4
<i>Amphiprion tricinctus</i>	No	Om	No	D	130	P	Dm	No	2;4
<i>Cheiloprion labiatus</i>	No	Co	No	D	80	P	Dm	No	1;2
<i>Chromis abrupta</i>	No	Pl	No	D	80	P	Dm	No	3
<i>Chromis acares</i>	No	Pl	No	D	55	P	Dm	No	1;2;3
<i>Chromis agilis</i>	No	Pl	No	D	90	P	Dm	No	135;1;2;3
<i>Chromis albomaculata</i>	No	Pl	No	D	180	P	Dm	No	2
<i>Chromis alleni</i>	No	Pl	No	D	80	P	Dm	No	2
<i>Chromis alpha</i>	No	Pl	No	D	100	P	Dm	No	2
<i>Chromis amboinensis</i>	No	Pl	No	D	75	P	Dm	No	1;2;3
<i>Chromis analis</i>	No	Pl	No	D	150	P	Dm	No	1;2;3
<i>Chromis atripectoralis</i>	No	Pl	No	D	100	P	Dm	No	1;2;3
<i>Chromis atripes</i>	No	Pl	No	D	70	G	Dm	No	1;2
<i>Chromis bami</i>	No	Pl	No	D	85	P	Dm	No	3
<i>Chromis caudalis</i>	No	Pl	No	D	90	P	Dm	No	2
<i>Chromis chrysur</i>	No	Pl	No	D	160	P	Dm	No	1;2;3
<i>Chromis cinerascens</i>	No	Pl	No	D	130	P	Dm	No	2
<i>Chromis delta</i>	No	Pl	No	D	65	P	Dm	No	1;2;3
<i>Chromis dimidiatus</i>	No	Pl	No	D	70	P	Dm	No	2
<i>Chromis elerae</i>	No	Pl	No	D	70	P	Dm	No	1;2;3
<i>Chromis fatuhivae</i>	No	Pl	No	D	67	P	Dm	No	3
<i>Chromis flavapicis</i>	No	Pl	No	D	135	P	Dm	No	3
<i>Chromis flavipectoralis</i>	No	Pl	No	D	70	P	Dm	No	2
<i>Chromis flavomaculata</i>	No	Pl	No	D	150	P	Dm	No	136;2
<i>Chromis fumea</i>	No	Pl	No	D	130	P	Dm	No	1;2;3
<i>Chromis iomelas</i>	No	Pl	No	D	70	P	Dm	No	1;2;3
<i>Chromis lepidolepis</i>	No	Pl	No	D	80	P	Dm	No	138;1;2;3

<i>Chromis leucura</i>	No	Pl	No	D	55	P	Dm	No	2;3
<i>Chromis lineata</i>	No	Pl	No	D	50	P	Dm	No	2
<i>Chromis margaritifer</i>	No	Pl	No	D	80	P	Dm	No	2;4
<i>Chromis nitida</i>	No	Pl	No	D	90	P	Dm	No	139;1;2;3
<i>Chromis notata</i>	No	Pl	No	D	160	P	Dm	No	2
<i>Chromis opercularis</i>	No	Pl	No	D	160	P	Dm	No	2
<i>Chromis ovatifformes</i>	No	Pl	No	D	90	P	Dm	No	2
<i>Chromis pamae</i>	No	Pl	No	D	137	P	Dm	No	3
<i>Chromis retrofasciata</i>	No	Pl	No	D	55	P	Dm	No	1;140;2;3
<i>Chromis scotochloptera</i>	No	Pl	No	D	150	P	Dm	No	2;4
<i>Chromis ternatensis</i>	No	Pl	No	D	70	P	Dm	No	1;2;3;131
<i>Chromis vanderbilti</i>	No	Pl	No	D	60	P	Dm	No	1;2;3
<i>Chromis viridis</i>	No	Pl	No	D	80	P	Dm	No	1;2;3
<i>Chromis weberi</i>	No	Pl	No	D	120	P	Dm	No	1;2;3
<i>Chromis westaustralis</i>	No	Pl	No	D	100	P	Dm	No	2
<i>Chromis xanthochira</i>	No	Pl	No	D	140	P	Dm	No	138;1;2
<i>Chromis xanthura</i>	No	Pl	No	D	150	P	Dm	No	1;2;3
<i>Chrysiptera albata</i>	No	N/A	No	D	46	P	Dm	No	3
<i>Chrysiptera bioellata</i>	No	Om	No	D	70	P	Dm	No	1;2;3
<i>Chrysiptera bleekeri</i>	Yes	N/A	Yes	D	80	P	Dm	No	2;4
<i>Chrysiptera brownriggii</i>	No	Om	No	D	85	P	Dm	No	2;3;4
<i>Chrysiptera caeruleolineata</i>	No	Pl	No	D	55	P	Dm	No	1;2;3
<i>Chrysiptera cyanea</i>	No	Pl	No	D	80	P	Dm	No	1;2;4
<i>Chrysiptera cymatilis</i>	No	Pl	No	D	60	P	Dm	No	2
<i>Chrysiptera flavipinnis</i>	No	N/A	No	D	80	P	Dm	No	1;2
<i>Chrysiptera galba</i>	No	N/A	No	D	95	P	Dm	No	3
<i>Chrysiptera glauca</i>	No	He	No	D	80	P	Dm	No	1;2;3
<i>Chrysiptera hemicyanea</i>	No	Pl	No	D	60	P	Dm	No	2;4
<i>Chrysiptera kuiteri</i>	Yes	Pl	Yes	D	60	P	Dm	No	141;2
<i>Chrysiptera oxycephala</i>	Yes	Pl	No	D	80	P	Dm	No	2;4
<i>Chrysiptera parasema</i>	No	N/A	No	D	60	P	Dm	No	2;4
<i>Chrysiptera rex</i>	No	He	No	D	80	P	Dm	No	1;2;3;4
<i>Chrysiptera rollandi</i>	No	Pl	No	D	50	P	Dm	No	1;2;3;4
<i>Chrysiptera sinclairi</i>	No	N/A	No	D	60	P	Dm	No	2
<i>Chrysiptera springeri</i>	No	Om	No	D	60	P	Dm	No	138;2
<i>Chrysiptera starcki</i>	No	N/A	No	D	90	P	Dm	No	1;2;3
<i>Chrysiptera talboti</i>	No	Pl	No	D	60	P	Dm	No	1;2;3;4
<i>Chrysiptera taupou</i>	No	N/A	No	D	80	P	Dm	No	1;2;3
<i>Chrysiptera traceyi</i>	No	Om	No	D	60	P	Dm	No	2;4
<i>Chrysiptera tricincta</i>	Yes	N/A	No	D	60	P	Dm	No	141;2
<i>Chrysiptera unimaculata</i>	No	He	No	D	80	P	Dm	No	1;2;3;131
<i>Dascyllus aruanus</i>	No	Pl	No	D	80	P	Dm	No	1;2;3
<i>Dascyllus auripinnis</i>	No	Pl	No	D	140	P	Dm	No	138;2;3
<i>Dascyllus carneus</i>	No	Pl	No	D	70	P	Dm	No	2;4
<i>Dascyllus flavicaudus</i>	No	Pl	No	D	110	P	Dm	No	138;2;3
<i>Dascyllus melanurus</i>	No	Pl	No	D	80	P	Dm	No	142;1;2;3
<i>Dascyllus reticulatus</i>	No	Pl	No	D	80	P	Dm	No	142;1;2;3
<i>Dascyllus strasburgi</i>	No	Om	No	D	105	P	Dm	No	142;3
<i>Dascyllus trimaculatus</i>	No	Om	No	D	140	P	Dm	No	142;1;2;3;4
<i>Dischistodus chrysopoecilus</i>	No	De	No	D	150	P	Dm	No	1;2
<i>Dischistodus darwiniensis</i>	No	De	No	D	130	P	Dm	No	2
<i>Dischistodus fasciatus</i>	No	De	No	D	130	P	Dm	No	2
<i>Dischistodus melanotus</i>	No	De	No	D	150	P	Dm	No	1;2;143

<i>Dischistodus perspicillatus</i>	No	De	No	D	190	P	Dm	No	1;2
<i>Dischistodus prosopotaenia</i>	No	De	No	D	180	P	Dm	No	1;2
<i>Dischistodus pseudochrysopoecilus</i>	No	De	No	D	160	P	Dm	No	1;2
<i>Hemiglyphidodon plagiometopon</i>	No	De	No	D	180	P	Dm	No	1;2;144
<i>Lepidozygus tapeinosoma</i>	No	Pl	No	D	90	P	Dm	No	1;2;3
<i>Neoglyphidodon bonang</i>	No	N/A	No	D	130	P	Dm	No	2;4
<i>Neoglyphidodon carlsoni</i>	No	N/A	No	D	120	P	Dm	No	2;3
<i>Neoglyphidodon crossi</i>	No	N/A	No	D	130	P	Dm	No	2;4
<i>Neoglyphidodon melas</i>	Yes	Co	No	D	150	P	Dm	No	1;2;4;143
<i>Neoglyphidodon nigroris</i>	No	Pl	No	D	110	P	Dm	No	1;2;3;4
<i>Neoglyphidodon oxyodon</i>	No	N/A	No	D	140	P	Dm	No	2;4
<i>Neoglyphidodon polyacanthus</i>	No	N/A	No	D	140	P	Dm	No	1;2;3
<i>Neoglyphidodon thoracotaenatus</i>	No	Pl	No	D	100	P	Dm	No	2;4
<i>Neopomacentrus aquadulcis</i>	No	Pl	No	D	100	P	Dm	No	2
<i>Neopomacentrus azysron</i>	No	Pl	No	D	80	G	Dm	No	35;1;2
<i>Neopomacentrus bankieri</i>	No	Pl	No	D	70	G	Dm	No	1;2;145
<i>Neopomacentrus cyanomos</i>	No	Pl	No	D	90	G	Dm	No	132;1;2
<i>Neopomacentrus filamentosus</i>	No	Pl	No	D	80	G	Dm	No	2;3;117
<i>Neopomacentrus metallicus</i>	No	Pl	No	D	80	G	Dm	No	3;4
<i>Neopomacentrus nemurus</i>	No	Pl	No	D	80	G	Dm	No	2;3;4
<i>Neopomacentrus taeniurus</i>	No	Pl	No	D	100	G	Dm	No	2
<i>Neopomacentrus violascens</i>	No	Pl	No	D	70	G	Dm	No	2;4
<i>Parma oligolepis</i>	No	N/A	No	D	200	P	Dm	No	1
<i>Parma polylepis</i>	No	De	No	D	220	P	Dm	No	146;1
<i>Plectroglyphidodon dickii</i>	No	Om	No	D	110	P	Dm	No	1;2;3;4
<i>Plectroglyphidodon flaviventris</i>	No	N/A	No	D	80	P	Dm	No	3;4
<i>Plectroglyphidodon imparipennis</i>	No	Om	No	D	60	P	Dm	No	1;2;3;4
<i>Plectroglyphidodon johnstonianus</i>	No	Co	No	D	90	P	Dm	No	1;2;3;4
<i>Plectroglyphidodon lacrymatus</i>	No	Om	No	D	100	P	Dm	No	1;2;3;144
<i>Plectroglyphidodon leucozonus</i>	No	He	No	D	100	P	Dm	No	1;2;3;4
<i>Plectroglyphidodon phoenixensis</i>	No	De	No	D	90	P	Dm	No	1;2;3;4
<i>Plectroglyphidodon sagmarius</i>	No	N/A	No	D	64	P	Dm	No	3
<i>Pomacentrus adelus</i>	No	De	No	D	85	P	Dm	No	2;3;144
<i>Pomacentrus albimaculus</i>	No	He	No	D	90	P	Dm	No	2;4
<i>Pomacentrus alexanderae</i>	No	Om	No	D	100	P	Dm	No	2;4
<i>Pomacentrus alleni</i>	No	N/A	No	D	65	P	Dm	No	2
<i>Pomacentrus amboinensis</i>	No	Om	No	D	100	P	Dm	No	1;147;2;3
<i>Pomacentrus armillatus</i>	No	N/A	No	D	70	P	Dm	No	2
<i>Pomacentrus aurifrons</i>	No	Pl	No	D	70	P	Dm	No	2;3
<i>Pomacentrus auriventris</i>	No	N/A	No	D	70	P	Dm	No	2
<i>Pomacentrus australis</i>	No	He	No	D	80	P	Dm	No	138;1;2
<i>Pomacentrus azuremaculatus</i>	No	N/A	No	D	100	P	Dm	No	2
<i>Pomacentrus bankanensis</i>	No	De	No	D	100	P	Dm	No	138;1;2;3;144
<i>Pomacentrus bipunctatus</i>	No	N/A	No	D	100	P	Dm	No	2
<i>Pomacentrus brachialis</i>	No	Om	No	D	100	P	Dm	No	138;2;3
<i>Pomacentrus burroughi</i>	No	De	No	D	80	P	Dm	No	2;144
<i>Pomacentrus callainus</i>	No	N/A	No	D	95	P	Dm	No	3
<i>Pomacentrus chrysurus</i>	No	De	No	D	90	P	Dm	No	1;2;3;144
<i>Pomacentrus coelestis</i>	No	Pl	No	D	70	P	Dm	No	1;2;3;4
<i>Pomacentrus colini</i>	No	N/A	No	D	90	P	Dm	No	2
<i>Pomacentrus cuneatus</i>	No	N/A	No	D	90	P	Dm	No	2
<i>Pomacentrus geminospilos</i>	No	N/A	No	D	75	P	Dm	No	2
<i>Pomacentrus grammarhynchus</i>	No	De	No	D	120	P	Dm	No	1;148;2

	<i>Pomacentrus imitator</i>	No	Pl	No	D	100	P	Dm	No	1;2;3
	<i>Pomacentrus javanicus</i>	No	N/A	No	D	80	P	Dm	No	2
	<i>Pomacentrus komodoensis</i>	No	N/A	No	D	80	P	Dm	No	2
	<i>Pomacentrus lepidogenys</i>	No	Pl	No	D	90	P	Dm	No	138;1;2
	<i>Pomacentrus limosus</i>	No	N/A	No	D	70	P	Dm	No	2
	<i>Pomacentrus littoralis</i>	No	N/A	No	D	100	P	Dm	No	2
	<i>Pomacentrus melanochir</i>	No	Pl	No	D	70	P	Dm	No	2;4
	<i>Pomacentrus milleri</i>	No	De	No	D	90	P	Dm	No	2;149
	<i>Pomacentrus moluccensis</i>	No	Om	No	D	70	P	Dm	No	138;1;2;3;69
	<i>Pomacentrus nagasakiensis</i>	No	Pl	No	D	120	P	Dm	No	1;2;3;4
	<i>Pomacentrus nigromanus</i>	No	Om	No	D	90	P	Dm	No	2;4
	<i>Pomacentrus nigromarginatus</i>	No	Om	No	D	90	P	Dm	No	138;1;2;3
	<i>Pomacentrus optisthostigma</i>	No	N/A	No	D	80	P	Dm	No	2
	<i>Pomacentrus pavo</i>	No	Pl	No	D	110	P	Dm	No	1;2;3
	<i>Pomacentrus philippinus</i>	No	Pl	No	D	100	P	Dm	No	1;2;3
	<i>Pomacentrus polyspinus</i>	No	N/A	No	D	100	P	Dm	No	2
	<i>Pomacentrus proteus</i>	No	N/A	No	D	100	P	Dm	No	2
	<i>Pomacentrus reidi</i>	No	N/A	No	D	110	P	Dm	No	2
	<i>Pomacentrus saksonoi</i>	No	N/A	No	D	90	P	Dm	No	2
	<i>Pomacentrus similis</i>	No	Om	No	D	70	P	Dm	No	147;2
	<i>Pomacentrus simsiang</i>	No	De	No	D	90	P	Dm	No	138;2
	<i>Pomacentrus smithi</i>	No	Pl	No	D	70	P	Dm	No	138;2
	<i>Pomacentrus spilotoceps</i>	No	N/A	No	D	80	P	Dm	No	2
	<i>Pomacentrus stigma</i>	No	N/A	No	D	120	P	Dm	No	2
	<i>Pomacentrus taeniometopon</i>	No	De	No	D	100	P	Dm	No	2;4
	<i>Pomacentrus tripunctatus</i>	No	De	No	D	100	P	Dm	No	1;2;144
	<i>Pomacentrus vaiuli</i>	No	Om	No	D	100	P	Dm	No	2;3
	<i>Pomacentrus wardi</i>	No	De	No	D	100	P	Dm	No	138;1;2;144
	<i>Pomacentrus yoshii</i>	No	N/A	No	D	80	P	Dm	No	2;3
	<i>Pomachromis fuscidorsalis</i>	No	Pl	No	D	85	P	Dm	No	3
	<i>Pomachromis guamensis</i>	No	Pl	No	D	55	P	Dm	No	2
	<i>Pomachromis richardsoni</i>	No	N/A	No	D	55	P	Dm	No	1;2;3
	<i>Premnas biaculeatus</i>	Yes	Om	No	D	80	P	Dm	No	2;4
	<i>Pristotis obtusirostris</i>	No	Pl	No	D	100	P	Dm	No	2;3
	<i>Stegastes albifasciatus</i>	No	De	No	D	110	P	Dm	No	1;2;3
	<i>Stegastes altus</i>	No	De	No	D	150	P	Dm	No	150;2
	<i>Stegastes apicalis</i>	No	De	No	D	130	P	Dm	No	1;2;144
	<i>Stegastes aureus</i>	No	De	No	D	110	P	Dm	No	1;2;3;4
	<i>Stegastes emeryi</i>	No	N/A	No	D	90	P	Dm	No	3
	<i>Stegastes fasciolatus</i>	No	De	No	D	110	P	Dm	Yes	151;1;2;3
	<i>Stegastes gascoynei</i>	No	De	No	D	150	P	Dm	No	1;2;3;4
	<i>Stegastes lividus</i>	No	De	No	D	150	P	Dm	No	1;2;3
	<i>Stegastes nigricans</i>	No	De	No	D	140	P	Dm	No	1;2;3
	<i>Stegastes obreptus</i>	No	De	No	D	150	P	Dm	No	2;149
	<i>Stegastes punctatus</i>	No	De	No	D	150	P	Dm	No	3
Priacanthidae	<i>Heteropriacanthus cruentatus</i>	No	Ca	No	N	320	G	Br	No	4
	<i>Priacanthus blochii</i>	No	Pl	No	N	350	G	Br	No	12
	<i>Priacanthus hamrur</i>	No	Ca	No	N	400	G	Br	No	4
	<i>Pristigenys nipponia</i>	No	N/A	No	N	350	G	Br	No	1;2;3
Pseudochromidae	<i>Amsichthys knighti</i>	No	N/A	No	D	45	P	Dm	No	1
	<i>Congrogadus subducens</i>	No	Ca	No	D	450	P	Dm	No	1;2;3;125
	<i>Cypho purpurascens</i>	Yes	N/A	No	D	75	P	Dm	No	1;2
	<i>Labracinus cyclyphthalmus</i>	Yes	Ca	No	D	200	P	Dm	No	1;2;88

	<i>Labracinus lineatus</i>	No	N/A	No	D	250	P	Dm	No	2
	<i>Lubbockichthys multisquamatus</i>	No	Mi	No	D	75	P	Dm	No	1;2;152
	<i>Manonichthys polynemus</i>	No	N/A	No	D	120	P	Dm	No	2;4
	<i>Manonichthys splendens</i>	Yes	N/A	No	D	130	P	Dm	No	2;4
	<i>Ogilbyina novaehollandiae</i>	Yes	N/A	No	D	100	P	Dm	No	1;2
	<i>Ogilbyina queenslandiae</i>	No	N/A	No	D	150	P	Dm	No	1;2
	<i>Ogilbyina salvati</i>	N/A	N/A	No	D	80	P	Dm	No	3
	<i>Ogilbyina velifera</i>	No	N/A	No	D	120	P	Dm	No	1;2
	<i>Pictichromis coralensis</i>	No	N/A	No	D	70	P	Dm	No	3
	<i>Pictichromis paccagnellae</i>	No	N/A	No	D	70	P	Dm	No	1;2
	<i>Pictichromis porphyrea</i>	No	Ma	No	D	70	P	Dm	No	2;3
	<i>Pseudochromis andamanensis</i>	No	N/A	No	D	70	P	Dm	No	2
	<i>Pseudochromis bitaeniatus</i>	No	Ma	No	D	70	P	Dm	No	1;2
	<i>Pseudochromis cyanotaenia</i>	Yes	Mi	No	D	60	P	Dm	No	1;2;4
	<i>Pseudochromis diadema</i>	No	N/A	No	D	60	P	Dm	No	2
	<i>Pseudochromis elongatus</i>	No	Ma	No	D	35	P	Dm	No	2
	<i>Pseudochromis flammicauda</i>	Yes	Ma	No	D	55	P	Dm	No	1;2
	<i>Pseudochromis fuscus</i>	No	Ca	No	D	90	P	Dm	No	1;2;3;153
	<i>Pseudochromis jamesi</i>	No	Ma	No	D	55	P	Dm	No	1;3
	<i>Pseudochromis marshallensis</i>	No	Ma	No	D	70	P	Dm	No	2
	<i>Pseudochromis moorei</i>	Yes	Ma	No	D	100	P	Dm	No	2;4
	<i>Pseudochromis paranox</i>	No	Ma	No	D	70	P	Dm	No	1;2
	<i>Pseudochromis perspicillatus</i>	No	Ma	No	D	120	P	Dm	No	1;2
	<i>Pseudochromis quinqueidentatus</i>	No	Ma	No	D	95	P	Dm	No	1
	<i>Pseudochromis ransonneti</i>	No	Ma	No	D	120	P	Dm	No	2
	<i>Pseudochromis steenei</i>	Yes	Ma	No	D	120	P	Dm	No	2
	<i>Pseudochromis wilsoni</i>	No	Ma	No	D	80	P	Dm	No	1;2
	<i>Pseudoplesiops immaculatus</i>	No	N/A	No	D	50	P	Dm	No	1;2
	<i>Pseudoplesiops rosae</i>	No	Ma	No	D	23	P	Dm	No	1;4
	<i>Pseudoplesiops typus</i>	No	N/A	No	D	70	P	Dm	No	1
Ptereleotridae	<i>Aioliops megastigma</i>	No	N/A	No	D	30	P	Dm	No	2
	<i>Nemateleotris decora</i>	Yes	Pl	Yes	D	85	P	Dm	Yes	1;2;3;4
	<i>Nemateleotris helfrichi</i>	Yes	Pl	Yes	D	65	P	Dm	Yes	2;3
	<i>Nemateleotris magnifica</i>	Yes	Pl	Yes	D	80	P	Dm	Yes	1;2;3;4
	<i>Oxymetopon compressus</i>	No	N/A	No	D	200	P	Dm	No	2
	<i>Oxymetopon cyanoctenosum</i>	Yes	N/A	Yes	D	200	P	Dm	No	2;4
	<i>Parioglossus formosus</i>	No	N/A	No	D	40	P	Dm	No	2
	<i>Parioglossus interruptus</i>	No	N/A	No	D	30	P	Dm	No	2
	<i>Parioglossus nudus</i>	No	N/A	No	D	20	P	Dm	No	2
	<i>Parioglossus raoi</i>	No	N/A	No	D	30	P	Dm	No	2
	<i>Ptereleotris evides</i>	Yes	Pl	Yes	D	135	P	Dm	Yes	1;2;3;12
	<i>Ptereleotris grammica</i>	Yes	Pl	Yes	D	100	P	Dm	No	1;2;12
	<i>Ptereleotris hanae</i>	Yes	Pl	Yes	D	120	P	Dm	No	1;2;3;12
	<i>Ptereleotris heteroptera</i>	Yes	Pl	Yes	D	120	P	Dm	Yes	1;2;3;4
	<i>Ptereleotris microlepis</i>	Yes	Pl	Yes	D	150	P	Dm	Yes	1;2;3;4
	<i>Ptereleotris monoptera</i>	Yes	Pl	Yes	D	150	P	Dm	Yes	1;2;3;4
	<i>Ptereleotris uroditaenia</i>	No	Pl	No	D	100	P	Dm	No	1;2
	<i>Ptereleotris zebra</i>	No	Pl	No	D	120	P	Dm	No	1;2;3;4
Rachycentridae	<i>Rachycentron canadum</i>	No	Ca	No	D	2000	G	Br	No	4
Scatophagidae	<i>Scatophagus argus</i>	No	Om	No	D	300	N/A	Dm	No	2;3
	<i>Selenotoca multifasciata</i>	No	N/A	No	D	280	N/A	Dm	No	2
Scombridae	<i>Acanthocybium solandri</i>	No	Pi	No	D	2100	G	Br	No	2
	<i>Grammatorcynus bilineatus</i>	No	Ca	No	D	1000	G	Br	No	1;2;3

	<i>Gymnosarda unicolor</i>	No	Pi	No	D	1800	G	Br	No	44;2;3
	<i>Rastrelliger kanagurta</i>	No	Pl	No	D	380	G	Br	No	3
	<i>Sarda orientalis</i>	No	Ca	No	D	1020	G	Br	No	2;4
	<i>Scomberomorus commerson</i>	No	Pi	No	D	2350	G	Br	No	1;2;3
	<i>Thunnus albacares</i>	No	Ca	No	D	2100	G	Br	No	2;4
Scorpaenidae	<i>Brachypterois serrulata</i>	No	N/A	No	N	120	P	Gel	No	2
	<i>Dendrochirus biocellatus</i>	No	Pi	No	N	100	P	Gel	No	2;12
	<i>Dendrochirus brachypterus</i>	No	Ma	No	N	170	P	Gel	No	1;2;3;154
	<i>Dendrochirus zebra</i>	No	Ma	No	N	180	P	Gel	No	1;2;3;154
	<i>Ebosia bleekeri</i>	No	N/A	No	N	220	P	Gel	No	2
	<i>Iracundus signifer</i>	No	Pi	No	D	130	P	Gel	No	2;4;155
	<i>Parapterois heterura</i>	No	N/A	No	N	230	P	Gel	No	2
	<i>Parascorpaena mossambica</i>	No	Ma	No	N	100	P	Gel	No	1;2;3;156
	<i>Parascorpaena picta</i>	No	Ma	No	N	150	P	Gel	No	2;154
	<i>Pteroidichthys amboinensis</i>	No	N/A	No	N	120	P	Gel	No	2
	<i>Pterois antennata</i>	No	Ma	No	N	200	P	Gel	No	1;2;3;4
	<i>Pterois kodipungi</i>	No	N/A	No	N	350	P	Gel	No	2
	<i>Pterois miles</i>	No	Ca	No	N	380	P	Gel	No	1;2;3;157
	<i>Pterois mombasae</i>	No	Ma	No	N	160	P	Gel	No	2;154
	<i>Pterois radiata</i>	No	Ma	No	N	240	P	Gel	No	2;154
	<i>Pterois volitans</i>	No	Ca	No	N	380	P	Gel	No	1;2;3;157
	<i>Rhinopias aphanes</i>	No	N/A	No	D	240	P	Gel	No	2
	<i>Rhinopias eschmeyeri</i>	No	N/A	No	D	190	P	Gel	No	2
	<i>Rhinopias frondosa</i>	No	Ca	No	N	230	P	Gel	No	2;4
	<i>Scorpaenodes guamensis</i>	No	Ma	No	N	140	P	Gel	No	1;2;3;154
	<i>Scorpaenodes hirsutus</i>	No	Ma	No	N	70	P	Gel	No	1;2;3;154
	<i>Scorpaenodes kelloggi</i>	No	Ma	No	D	48	P	Gel	No	3;158
	<i>Scorpaenodes littoralis</i>	No	Ma	No	N	80	P	Gel	No	2
	<i>Scorpaenodes minor</i>	N/A	Ma	No	D	52	P	Gel	No	3
	<i>Scorpaenodes parvipinnis</i>	No	Ma	No	N	130	P	Gel	No	1;2;3;154
	<i>Scorpaenodes quadrispinosus</i>	N/A	Ma	No	D	97	P	Gel	No	3
	<i>Scorpaenodes varipinnis</i>	No	Ma	No	N	70	P	Gel	No	2;154
	<i>Scorpaenopsis diabolus</i>	Yes	Pi	No	D	280	P	Gel	No	1;2;3;12
	<i>Scorpaenopsis macrochir</i>	Yes	N/A	No	D	150	P	Gel	No	2;3;4
	<i>Scorpaenopsis neglecta</i>	No	Pi	No	D	150	P	Gel	No	2;159
	<i>Scorpaenopsis oxycephala</i>	No	Ma	No	D	360	P	Gel	No	2;12
	<i>Scorpaenopsis papuensis</i>	No	N/A	No	D	220	P	Gel	No	2
	<i>Scorpaenopsis possi</i>	No	N/A	No	D	220	P	Gel	No	2;3
	<i>Scorpaenopsis venosa</i>	No	N/A	No	D	200	P	Gel	No	2
	<i>Sebastapistes cyanostigma</i>	No	Ma	No	N	80	P	Gel	No	1;2;3;4
	<i>Sebastapistes mauritiana</i>	No	Pi	No	N	80	P	Gel	No	2
	<i>Sebastapistes strongia</i>	No	Pi	No	N	95	P	Gel	No	3;4
	<i>Taenianotus triacanthus</i>	Yes	Ca	No	D	100	P	Gel	No	1;2;3;4
Serranidae	<i>Aethaloperca rogae</i>	No	Pi	No	D	600	P	Br	No	1;3;154
	<i>Anypserdon leucogrammicus</i>	No	Pi	No	D	520	P	Br	No	1;2;3
	<i>Aporops bilinearis</i>	N/A	N/A	No	D	115	P	Br	No	1;3
	<i>Belonoperca chabanaudi</i>	No	Ma	No	D	150	P	Br	No	1;2;3;12
	<i>Belonoperca pylei</i>	N/A	N/A	No	D	80	P	Br	No	1;3
	<i>Cephalopholis argus</i>	No	Pi	No	D	550	G	Br	No	1;2;3;154
	<i>Cephalopholis boenak</i>	No	Ca	No	D	260	P	Br	No	1;2;3;4
	<i>Cephalopholis cyanostigma</i>	No	Ca	No	D	300	P	Br	No	1;2;4
	<i>Cephalopholis formosa</i>	No	N/A	No	D	340	P	Br	No	1;2
	<i>Cephalopholis leopardus</i>	No	Pi	No	D	200	P	Br	No	1;2;3;12

<i>Cephalopholis microprion</i>	No	Ca	No	D	240	P	Br	No	1;2;3;40;160
<i>Cephalopholis miniata</i>	No	Pi	No	D	410	G	Br	No	1;2;3;12
<i>Cephalopholis polleni</i>	No	N/A	No	D	430	P	Br	No	2;3
<i>Cephalopholis sexmaculata</i>	No	Pi	No	D	480	G	Br	No	1;2;3;12
<i>Cephalopholis sonnerati</i>	No	Ma	No	D	570	G	Br	No	1;2;3;154
<i>Cephalopholis spiloparaea</i>	No	Pi	No	D	220	P	Br	No	1;2;3;12
<i>Cephalopholis urodeta</i>	No	Pi	No	D	270	G	Br	No	1;2;3;12
<i>Cromileptes altivelis</i>	No	Ca	No	D	700	P	Br	No	1;2;3;4
<i>Diploprion bifasciatum</i>	No	Ca	No	D	250	P	Br	No	1;2;3;88;161
<i>Epinephelus areolatus</i>	No	Ca	No	D	400	G	Br	No	1;2;3;4
<i>Epinephelus bilobatus</i>	No	N/A	No	D	330	G	Br	No	2
<i>Epinephelus bleekeri</i>	No	Ca	No	D	750	G	Br	No	2;24
<i>Epinephelus bontoides</i>	No	N/A	No	D	300	G	Br	No	2
<i>Epinephelus caeruleopunctatus</i>	No	Ca	No	D	600	G	Br	No	1;2;3;4
<i>Epinephelus chlorostigma</i>	No	Ca	No	D	750	G	Br	No	2;4
<i>Epinephelus coioides</i>	No	Ca	No	D	950	G	Br	No	1;2;3;4
<i>Epinephelus corallicola</i>	No	N/A	No	D	490	G	Br	No	1;2
<i>Epinephelus cyanopodus</i>	No	Pi	No	D	120	G	Br	No	1;2;3;12
<i>Epinephelus erythrurus</i>	No	N/A	No	D	430	G	Br	No	2
<i>Epinephelus fasciatus</i>	No	Ca	No	D	400	G	Br	No	1;2;3;154
<i>Epinephelus fuscoguttatus</i>	No	Ca	No	D	1000	G	Br	No	1;2;3;154;44
<i>Epinephelus hexagonatus</i>	No	Ca	No	D	260	G	Br	No	1;2;3;154
<i>Epinephelus howlandi</i>	No	Pi	No	D	440	G	Br	No	1;2;3;12
<i>Epinephelus irroratus</i>	No	N/A	No	D	340	G	Br	No	3
<i>Epinephelus lanceolatus</i>	No	Ca	No	D	2340	G	Br	No	1;2;3;4
<i>Epinephelus longispinis</i>	No	Ma	No	D	500	G	Br	No	2;4
<i>Epinephelus macrospilos</i>	No	Ca	No	D	430	G	Br	No	1;2;3;154
<i>Epinephelus maculatus</i>	No	Ca	No	D	600	G	Br	No	1;2;3;4
<i>Epinephelus malabricus</i>	No	Ca	No	D	2340	G	Br	No	1;2;3;4
<i>Epinephelus melanostigma</i>	No	Pi	No	D	330	G	Br	No	2;3;12
<i>Epinephelus merra</i>	No	Ca	No	D	320	G	Br	No	1;2;3
<i>Epinephelus miliaris</i>	No	Ma	No	D	530	G	Br	No	2;3;4
<i>Epinephelus multinotatus</i>	No	Ca	No	D	1000	G	Br	No	2;4
<i>Epinephelus ongus</i>	No	Ca	No	D	350	G	Br	No	1;2;3;162
<i>Epinephelus polyphkadion</i>	No	Ca	No	D	610	G	Br	No	1;2;3
<i>Epinephelus quoyanus</i>	No	Ca	No	D	380	G	Br	No	1;2;4
<i>Epinephelus retouti</i>	No	N/A	No	D	470	G	Br	No	3
<i>Epinephelus rivulatus</i>	No	Ca	No	D	450	G	Br	No	1;2;3;4
<i>Epinephelus socialis</i>	No	Pi	No	D	420	G	Br	No	2;3;12
<i>Epinephelus spilotoceps</i>	No	Ca	No	D	310	G	Br	No	2;4
<i>Epinephelus tauvina</i>	No	Pi	No	D	330	G	Br	No	1;2;3;12
<i>Epinephelus tukula</i>	No	Ca	No	D	2000	G	Br	No	1;2;3;4
<i>Epinephelus undulosus</i>	No	Ca	No	D	500	G	Br	No	2;4
<i>Epinephelus waandersi</i>	No	N/A	No	D	600	G	Br	No	2
<i>Gracilo albomarginata</i>	No	Pi	No	D	500	N/A	Br	No	1;2;3;12
<i>Grammistes sexlineatus</i>	No	Pi	No	D	270	N/A	Br	No	1;2;3;4
<i>Grammistops ocellatus</i>	Yes	N/A	No	D	130	N/A	Gel	No	1;2;3
<i>Liopropoma flavidum</i>	N/A	N/A	No	D	61	N/A	Br	No	3
<i>Liopropoma latifasciatum</i>	N/A	N/A	No	D	160	N/A	Br	No	2
<i>Liopropoma mitratum</i>	N/A	N/A	No	D	82	N/A	Br	No	3
<i>Liopropoma multilineatum</i>	N/A	N/A	No	D	77	N/A	Br	No	3
<i>Liopropoma pallidum</i>	N/A	N/A	No	D	78	N/A	Br	No	3
<i>Liopropoma susumi</i>	N/A	N/A	No	D	91	N/A	Br	No	3

<i>Liopropoma tonstrinum</i>	N/A	N/A	No	D	80	N/A	Br	No	3
<i>Luzonichthys earlei</i>	No	Pl	No	D	44	N/A	Br	No	2
<i>Luzonichthys waitei</i>	No	Pl	No	D	70	N/A	Br	No	1;2;3;12
<i>Luzonichthys whitleyi</i>	No	Pl	No	D	60	N/A	Br	No	2;3
<i>Luzonichthys williamsi</i>	No	N/A	No	D	58	N/A	Br	No	3
<i>Plectranthias inermis</i>	No	N/A	No	D	350	N/A	Br	No	2;4
<i>Plectranthias longimanus</i>	No	N/A	No	D	350	N/A	Br	No	1;2;3
<i>Plectranthias nanus</i>	No	N/A	No	D	50	N/A	Br	No	1;2;3
<i>Plectranthias winniensis</i>	No	N/A	No	D	350	N/A	Br	No	1;2;3
<i>Plectropomus areolatus</i>	No	Pi	No	D	800	G	Br	No	1;2;3;12
<i>Plectropomus laevis</i>	No	Pi	No	D	1250	G	Br	No	1;2;3;12
<i>Plectropomus leopardus</i>	No	Pi	No	D	700	G	Br	No	1;2;3;12
<i>Plectropomus maculatus</i>	No	Pi	No	D	1250	G	Br	No	1;2;3
<i>Plectropomus oligacanthus</i>	No	Ca	No	D	750	G	Br	No	1;2;3;4
<i>Plectropomus pessuliferus</i>	No	Pi	No	D	1200	G	Br	No	1;2;3;12
<i>Pogonoperca punctata</i>	Yes	Pi	No	D	350	N/A	Gel	No	1;2;3;12
<i>Pseudanthias aurulentus</i>	No	Pl	No	D	60	P	Br	No	2
<i>Pseudanthias bartlettorum</i>	No	Pl	No	D	90	P	Br	No	2;3
<i>Pseudanthias bicolor</i>	No	Pl	No	D	130	P	Br	No	1;2;3;12
<i>Pseudanthias bimaculatus</i>	No	Pl	No	D	90	P	Br	No	2
<i>Pseudanthias carlsoni</i>	No	Pl	No	D	100	P	Br	No	3
<i>Pseudanthias cooperi</i>	No	Pl	No	D	140	P	Br	No	1;2;3
<i>Pseudanthias dispar</i>	No	Pl	No	D	95	P	Br	No	1;2;3
<i>Pseudanthias engelhardi</i>	No	Pl	No	D	100	P	Br	No	1;2;3
<i>Pseudanthias evansi</i>	No	Pl	No	D	100	P	Br	No	2
<i>Pseudanthias fasciatus</i>	No	Pl	No	D	210	P	Br	No	1;2
<i>Pseudanthias flavoguttatus</i>	No	Pl	No	D	110	P	Br	No	2
<i>Pseudanthias hiva</i>	No	Pl	No	D	140	P	Br	No	3
<i>Pseudanthias huchti</i>	No	Pl	No	D	120	P	Br	No	1;2
<i>Pseudanthias hutomoi</i>	No	Pl	No	D	120	P	Br	No	2
<i>Pseudanthias hypselosoma</i>	No	Pl	No	D	70	P	Br	No	1;2
<i>Pseudanthias ignitus</i>	No	Pl	No	D	80	P	Br	No	2;4
<i>Pseudanthias lori</i>	No	Pl	No	D	120	P	Br	No	1;2;3
<i>Pseudanthias luzonensis</i>	No	Pl	No	D	145	P	Br	No	1;2
<i>Pseudanthias mooreanus</i>	N/A	Pl	No	D	72	P	Br	No	3
<i>Pseudanthias olivaceus</i>	No	Pl	No	D	120	P	Br	No	2;3
<i>Pseudanthias parvirostris</i>	No	Pl	No	D	100	P	Br	No	2;4
<i>Pseudanthias pascalus</i>	No	Pl	No	D	170	P	Br	No	2;3
<i>Pseudanthias pictilis</i>	No	Pl	No	D	135	P	Br	No	2
<i>Pseudanthias pleurotaenia</i>	No	Pl	No	D	200	P	Br	No	1;2;3;12
<i>Pseudanthias randalli</i>	No	Pl	No	D	70	P	Br	No	2
<i>Pseudanthias regalis</i>	No	Pl	No	D	62	P	Br	No	3
<i>Pseudanthias rubrizonatus</i>	No	Pl	No	D	100	P	Br	No	2;4
<i>Pseudanthias sheni</i>	No	Pl	No	D	200	P	Br	No	2
<i>Pseudanthias smithvanzii</i>	No	Pl	No	D	95	P	Br	No	1;2
<i>Pseudanthias squamipinnis</i>	No	Pl	No	D	150	P	Br	No	1;2;3
<i>Pseudanthias tuka</i>	No	Pl	No	D	120	P	Br	No	1;2
<i>Pseudanthias venator</i>	No	Pl	No	D	70	P	Br	No	2
<i>Pseudanthias ventralis</i>	No	Pl	No	D	70	P	Br	No	1;2;3;12
<i>Pseudogramma polyacanthum</i>	No	Mi	No	D	75	N/A	Br	No	1;2;3
<i>Serranocirrhites latus</i>	No	N/A	No	D	350	N/A	Br	No	1;2;3
<i>Suttonia lineata</i>	N/A	N/A	No	D	96	N/A	Br	No	3
<i>Variola albimarginata</i>	No	Pi	No	D	550	G	Br	No	1;2;3

Appendix C

	<i>Variola louti</i>	No	Pi	No	D	550	G	Br	No	1;2;3;44
Siganidae	<i>Siganus argenteus</i>	No	He	No	D	420	G	Es	No	163;1;2;3;6
	<i>Siganus canaliculatus</i>	No	He	No	D	290	G	Es	No	163;1;2;3;164
	<i>Siganus corallinus</i>	Yes	He	No	D	250	G	Es	No	163;1;2;3;6
	<i>Siganus doliatus</i>	Yes	He	No	D	250	G	Es	No	1;2;3
	<i>Siganus guttatus</i>	No	He	No	D	350	G	Es	No	163;1;2;3
	<i>Siganus javus</i>	Yes	He	No	D	530	G	Es	No	163;1;2;3;165
	<i>Siganus lineatus</i>	No	De	No	D	350	G	Es	No	163;1;2;3;166
	<i>Siganus magnificus</i>	Yes	He	No	D	230	G	Es	No	163;2;4
	<i>Siganus puelloides</i>	Yes	He	No	D	310	G	Es	No	163;2
	<i>Siganus puellus</i>	Yes	Sp	No	D	380	G	Es	No	163;167
	<i>Siganus punctatissimus</i>	Yes	He	No	D	280	G	Es	No	163;2;6
	<i>Siganus punctatus</i>	Yes	He	No	D	300	G	Es	No	163;1;2;3;6
	<i>Siganus randalli</i>	No	He	No	D	250	G	Es	No	163;2
	<i>Siganus spinus</i>	No	He	No	D	200	G	Es	No	163;1;2;3
	<i>Siganus stellatus</i>	Yes	He	No	D	350	G	Es	No	163;2
	<i>Siganus unimaculatus</i>	Yes	He	No	D	240	G	Es	No	163;2;4
	<i>Siganus uspi</i>	Yes	He	No	D	220	G	Es	No	163;2;4
	<i>Siganus vermiculatus</i>	No	He	No	D	370	G	Es	No	163;1;2;3
<i>Siganus virgatus</i>	No	He	No	D	300	G	Es	No	163;2	
<i>Siganus vulpinus</i>	Yes	He	No	D	240	G	Es	No	163;1;2;3	
Soleidae	<i>Aseraggodes kaianus</i>	No	Ma	No	D	140	N/A	Br	No	1;2;3;4
	<i>Aseraggodes melanostictus</i>	No	Ma	No	D	40	N/A	Br	No	2
	<i>Liachirus melanospilus</i>	No	Ma	No	D	150	N/A	Br	No	2
	<i>Pardachirus pavoninus</i>	No	Ma	No	N	220	N/A	Br	No	2
	<i>Soleichthys heterorhinos</i>	No	Ma	No	N	150	N/A	Br	No	2
	<i>Synaptura marginata</i>	No	Ma	No	N	300	N/A	Br	No	1;2;3;4
	<i>Zebrias fasciatus</i>	No	Ma	No	N	250	N/A	Br	No	2
Solenostomidae	<i>Solenostomus cyanopterus</i>	Yes	Mi	No	D	160	P	Po	Yes	4
	<i>Solenostomus halimeda</i>	Yes	Mi	No	D	70	P	Po	No	1;2;3
	<i>Solenostomus paegnius</i>	Yes	Mi	No	D	120	P	Po	No	4
	<i>Solenostomus paradoxus</i>	Yes	Mi	No	D	110	P	Po	No	1;2;3
Sparidae	<i>Acanthopagrus berda</i>	No	Ca	No	D	500	G	Br	No	2;4
Sphyracnidae	<i>Sphyracna barracuda</i>	No	Pi	No	D	1800	G	Br	No	1;2;3;168
	<i>Sphyracna flavicauda</i>	No	N/A	No	N	500	G	Br	No	1;2;3;4
	<i>Sphyracna forsteri</i>	No	Ca	No	N	650	G	Br	No	1;2;3;4
	<i>Sphyracna helleri</i>	No	N/A	No	N	850	G	Br	No	1;2;3;4
	<i>Sphyracna jello</i>	No	Ca	No	D	1500	G	Br	No	1;2;3;169
	<i>Sphyracna genie</i>	No	Pi	No	N	1000	G	Br	No	1;2;3;40
Synanceiidae	<i>Choridactylus multibarbus</i>	No	N/A	No	D	120	P	Gel	No	2
	<i>Dampierosa daruma</i>	No	N/A	No	D	130	P	Gel	No	2
	<i>Inimicus caledonicus</i>	No	N/A	No	D	250	P	Gel	No	2;3
	<i>Inimicus didactylus</i>	Yes	Ca	No	D	180	P	Gel	No	1;2;3;4
	<i>Inimicus sinensis</i>	No	N/A	No	D	180	P	Gel	No	2
	<i>Minous trachycephalus</i>	No	N/A	No	N	90	P	Gel	No	2;4
	<i>Synanceia horridas</i>	No	Ca	No	N	300	P	Gel	No	1;2;3
	<i>Synanceia verrucosa</i>	No	Ca	No	D	350	P	Gel	No	2;4
Syngnathidae	<i>Acentromura breviperula</i>	Yes	Mi	No	D	50	P	Po	No	1;2;3
	<i>Choeroichthys brachysoma</i>	No	Mi	No	D	65	P	Po	No	1;2;3
	<i>Choeroichthys cinctus</i>	No	Mi	No	D	80	P	Po	No	1;2;3
	<i>Corythoichthys amplexus</i>	Yes	Mi	No	D	95	P	Po	Yes	170
	<i>Corythoichthys flavofasciatus</i>	Yes	Mi	No	D	180	P	Po	Yes	171
	<i>Corythoichthys haematopterus</i>	Yes	Mi	No	D	180	P	Po	Yes	1;2;3

	<i>Corythoichthys intestinalis</i>	Yes	Mi	No	D	180	P	Po	Yes	170
	<i>Corythoichthys nigripectus</i>	Yes	Mi	No	D	110	P	Po	Yes	1;2;3
	<i>Corythoichthys ocellatus</i>	Yes	Mi	No	D	110	P	Po	No	1;2;3
	<i>Corythoichthys polynotatus</i>	Yes	Mi	No	D	160	P	Po	No	1;2;3
	<i>Corythoichthys schultzi</i>	Yes	Mi	No	D	160	P	Po	Yes	1;2;3
	<i>Doryrhamphus excisus</i>	Yes	Mi	No	D	70	P	Po	Yes	4
	<i>Doryrhamphus janssi</i>	Yes	Mi	No	D	130	P	Po	No	4
	<i>Doryrhamphus japonicus</i>	Yes	Mi	No	D	70	P	Po	Yes	4
	<i>Doryrhamphus negrosensis</i>	Yes	Mi	No	D	55	P	Po	No	1;2;3
	<i>Dunckerocampus boylei</i>	Yes	Mi	No	D	160	P	Po	No	1;2;3
	<i>Dunckerocampus dactyliophorus</i>	Yes	Mi	No	D	180	P	Po	No	172
	<i>Dunckerocampus multiannulatus</i>	Yes	Mi	No	D	180	P	Po	No	1;2;3
	<i>Dunckerocampus pessuliferus</i>	Yes	Mi	No	D	160	P	Po	No	172
	<i>Halicampus brocki</i>	No	N/A	No	D	115	P	Po	No	1;2;3
	<i>Halicampus macrorhynchus</i>	Yes	N/A	No	D	160	P	Po	No	4
	<i>Halicampus mataafae</i>	No	N/A	No	D	150	P	Po	No	1;2;3
	<i>Halicampus nitidus</i>	No	N/A	No	D	75	P	Po	No	1;2;3
	<i>Hippichthys cyanospilos</i>	No	N/A	No	D	160	P	Po	No	1;2;3
	<i>Hippocampus barbouri</i>	No	N/A	No	D	150	P	Po	No	1;2;3
	<i>Hippocampus bargibanti</i>	Yes	Mi	No	D	20	P	Po	No	173;174
	<i>Hippocampus colemani</i>	No	Mi	No	D	10	P	Po	No	1;2;3
	<i>Hippocampus comes</i>	Yes	Mi	No	N	160	P	Po	Yes	4
	<i>Hippocampus denise</i>	No	Mi	No	D	15	P	Po	No	174
	<i>Hippocampus hystrix</i>	No	Mi	No	D	150	P	Po	Yes	174
	<i>Hippocampus kuda</i>	Yes	Mi	No	D	150	P	Po	No	4
	<i>Hippocampus trimaculatus</i>	No	Mi	No	D	220	P	Po	No	174
	<i>Micrognathus andersonii</i>	No	N/A	No	D	60	P	Po	No	1;2;3
	<i>Micrognathus pygmaeus</i>	No	N/A	No	D	60	P	Po	No	1;2;3
	<i>Phoxocampus tetrophthalmus</i>	No	N/A	No	D	80	P	Po	No	1;2;3
	<i>Siokunichthys nigrolineatus</i>	No	N/A	No	D	400	P	Po	No	1;2;3
	<i>Syngnathoides biaculeatus</i>	Yes	Mi	No	D	280	P	Po	No	175
	<i>Trachyrhamphus bicoarctatus</i>	No	N/A	No	D	400	P	Po	No	1;2;3
	<i>Trachyrhamphus longirostris</i>	No	N/A	No	D	400	P	Po	No	1;2;3
Synodontidae	<i>Saurida gracilis</i>	No	Pi	No	N	280	P	Br	No	1;2;3;4
	<i>Saurida nebulosa</i>	No	Pi	No	D	200	P	Br	No	1;2;3
	<i>Synodus binotatus</i>	Yes	Pi	No	D	170	P	Br	No	1;2;3;12
	<i>Synodus dermatogenys</i>	Yes	Pi	No	D	200	P	Br	No	1;2;3;160
	<i>Synodus jaculum</i>	Yes	Pi	No	D	140	P	Br	No	1;2;3;12
	<i>Synodus rubromarmoratus</i>	No	Pi	No	D	85	P	Br	No	1;2;3
	<i>Synodus variegatus</i>	Yes	Pi	No	D	240	P	Br	No	1;2;3;12
	<i>Trachinocephalus myops</i>	No	Ca	No	D	250	P	Br	No	1;2;3;176
Tetraodontidae	<i>Arothron caeruleopunctatus</i>	No	Om	No	D	700	G	Es	No	1;2;3
	<i>Arothron hispidus</i>	No	Om	No	D	480	G	Es	No	4
	<i>Arothron immaculatus</i>	No	Om	No	D	280	G	Es	No	175
	<i>Arothron manilensis</i>	No	Om	No	D	310	G	Es	No	15
	<i>Arothron mappa</i>	No	Om	No	D	600	G	Es	No	1;2;3
	<i>Arothron meleagris</i>	No	Co	No	D	500	G	Es	No	77
	<i>Arothron nigropunctatus</i>	Yes	Co	No	D	330	G	Es	No	77
	<i>Arothron reticularis</i>	No	Ma	No	D	300	G	Es	No	4
	<i>Arothron stellatus</i>	No	Om	No	D	900	G	Es	No	1;2;3
	<i>Canthigaster amboinensis</i>	No	Om	No	D	140	P	Dm	No	177
	<i>Canthigaster bennetti</i>	Yes	He	No	D	100	P	Dm	Yes	4;178
	<i>Canthigaster callisterna</i>	Yes	N/A	No	D	240	P	Dm	No	1;2;3

	<i>Canthigaster compressa</i>	Yes	Mi	No	D	100	P	Dm	No	107;4
	<i>Canthigaster coronata</i>	No	Om	No	D	135	P	Dm	No	4
	<i>Canthigaster epilampra</i>	Yes	Om	No	D	110	P	Dm	No	4
	<i>Canthigaster janthinoptera</i>	Yes	Om	No	D	90	P	Dm	No	4
	<i>Canthigaster leoparda</i>	Yes	N/A	No	D	70	P	Dm	No	1;2;3
	<i>Canthigaster ocellicincta</i>	Yes	N/A	No	D	65	P	Dm	No	4
	<i>Canthigaster papua</i>	Yes	Mi	No	D	90	P	Dm	No	107
	<i>Canthigaster rivulata</i>	No	N/A	No	D	180	P	Dm	No	1;2;3
	<i>Canthigaster smithae</i>	No	N/A	No	D	130	P	Dm	No	1;2;3
	<i>Canthigaster solandri</i>	Yes	Om	No	D	105	P	Dm	No	4;178
	<i>Canthigaster tyleri</i>	No	Om	No	D	80	P	Dm	No	4
	<i>Canthigaster valentini</i>	Yes	Om	No	D	90	P	Dm	Yes	4;178
	<i>Chelonodon patoca</i>	No	Om	No	D	330	P	Dm	No	179
	<i>Lagocephalus scleratus</i>	No	Ca	No	D	850	P	Dm	No	180;181
	<i>Torquigener brevipinnis</i>	No	N/A	No	D	140	P	Dm	No	1;2;3
Tetrarogidae	<i>Ablabys macracanthus</i>	Yes	N/A	No	D	150	P	Br	No	2
	<i>Ablabys taenianotus</i>	Yes	Ca	No	N	150	P	Br	No	1;2;3;20
	<i>Paracentropogon longispinus</i>	No	N/A	No	D	120	P	Br	No	2
	<i>Richardsonichthys leucogaster</i>	No	N/A	No	N	100	P	Br	No	2;4
	<i>Tetraroge barbata</i>	Yes	Ca	No	D	110	P	Br	No	2;4
	<i>Tetraroge niger</i>	Yes	N/A	No	D	135	P	Br	No	2
Trichonotidae	<i>Trichonotus elegans</i>	No	Ma	No	D	180	N/A	N/A	No	1;2;3;182
	<i>Trichonotus halsteadii</i>	No	Ma	No	D	150	N/A	N/A	No	2;182
	<i>Trichonotus setiger</i>	No	Ma	No	D	180	N/A	N/A	No	1;2;3;182
Tripterygiidae	<i>Ceratobregma helenae</i>	No	N/A	No	D	45	P	Dm	No	1;2;3
	<i>Enneapterygius atrogulare</i>	Yes	Om	No	D	53	P	Dm	No	3;73
	<i>Enneapterygius elegans</i>	N/A	N/A	No	D	35	P	Dm	No	3
	<i>Enneapterygius flavoccipitis</i>	Yes	N/A	No	D	35	P	Dm	No	2
	<i>Enneapterygius hemimelas</i>	N/A	N/A	No	D	48	P	Dm	No	3
	<i>Enneapterygius mirabilis</i>	No	N/A	No	D	35	P	Dm	No	1;2
	<i>Enneapterygius nanus</i>	N/A	N/A	No	D	28	P	Dm	No	3
	<i>Enneapterygius niger</i>	N/A	N/A	No	D	35	P	Dm	No	3
	<i>Enneapterygius nigricauda</i>	N/A	N/A	No	D	35	P	Dm	No	3
	<i>Enneapterygius pallidoserialis</i>	Yes	N/A	No	D	35	P	Dm	No	2
	<i>Enneapterygius paucifasciatus</i>	N/A	N/A	No	D	35	P	Dm	No	3
	<i>Enneapterygius philippinus</i>	Yes	N/A	No	D	35	P	Dm	No	2
	<i>Enneapterygius pyramis</i>	N/A	N/A	No	D	34	P	Dm	No	3
	<i>Enneapterygius randalli</i>	N/A	N/A	No	D	34	P	Dm	No	3
	<i>Enneapterygius rhabdotus</i>	N/A	N/A	No	D	32	P	Dm	No	3
	<i>Enneapterygius rhotion</i>	N/A	N/A	No	D	37	P	Dm	No	3
	<i>Enneapterygius rufopileus</i>	No	Mi	No	D	45	P	Dm	No	3;183
	<i>Enneapterygius similis</i>	N/A	N/A	No	D	39	P	Dm	No	3
	<i>Enneapterygius triserialis</i>	N/A	N/A	No	D	45	P	Dm	No	3
	<i>Enneapterygius williamsi</i>	N/A	N/A	No	D	33	P	Dm	No	3
	<i>Helcogramma capidatum</i>	No	N/A	No	D	41	P	Dm	No	3
	<i>Helcogramma chica</i>	No	Om	No	D	40	P	Dm	No	3;184
	<i>Helcogramma rhinoceros</i>	Yes	N/A	No	D	40	P	Dm	No	2
	<i>Helcogramma striatum</i>	No	Pl	No	D	50	P	Dm	No	1;2;3;4
	<i>Helcogramma vulcanum</i>	Yes	N/A	No	D	40	P	Dm	No	2
	<i>Norfolkia brachylepis</i>	N/A	N/A	No	D	73	P	Dm	No	3
	<i>Norfolkia squamiceps</i>	N/A	N/A	No	D	66	P	Dm	No	3
	<i>Norfolkia thomasi</i>	N/A	N/A	No	D	50	P	Dm	No	3
	<i>Springerichthys kulbickii</i>	N/A	N/A	No	D	35	P	Dm	No	3

	<i>Ucla xenogrammus</i>	No	N/A	No	D	45	P	Dm	No	1;2;3
Uranoscopidae	<i>Uranoscopus sulphureus</i>	No	N/A	No	D	350	N/A	N/A	No	2
Zanclidae	<i>Zanclus cornutus</i>	Yes	Om	No	D	160	G	Br	No	1;2;3

1: Randall et al. 1997; 2: Allen et al. 2003; 3: Randall 2005; 4: Froese & Pauly 2012; 5: Barlow 1974; 6: Cheal et al. 2012; 7: Russ 1984; 8: Robertson et al. 1979; 9: Brandl & Bellwood 2013b; 10: Randall & Struhsacker 1981; 11: Meyer-Rochow 1976; 12: Sandin & Williams 2010; 13: Pietsch & Grobecker 1987; 14: Nakane et al. 2011; 15: Nanjo et al. 2008; 16: Marnane & Bellwood 2002; 17: Zagars et al. 2013; 18: Barnett & Bellwood 2006; 19: Job & Shand 2001; 20: Unsworth et al. 2007; 21: Unsworth et al. 2009; 22: Schmitz & Wainwright 2011; 23: Fukumori et al. 2008; 24: Hajisamae 2009; 25: Longenecker & Langston 2006; 26: Nakamura et al. 2006; 27: Kuwamura 1991; 28: Chen et al. 2005; 29: Wilson 2000; 30: Ho et al. 2007; 31: Carlson 2012; 32: Roberts 1987; 33: Kwak et al. 2004; 34: Townsend & Tibbetts 2000; 35: Hamner et al. 1988; 36: Fricke & Zaiser 1982; 37: Grandcourt et al. 2004; 38: Salini et al. 1994; 39: Farmer & Wilson 2011; 40: Blaber et al. 1990; 41: Meyer et al. 2001; 42: Brewer et al. 1989; 43: Smith & Parrish 2002; 44: Randall 1980; 45: Blaber & Cyrus 1983; 46: Sreenivasan 1974; 47: Tandon 1960; 48: Barreiros et al. 2003; 49: Roberts & Ormond 1992; 50: Pratchett et al. 2006; 51: Reese 1975; 52: Sano 1989; 53: Allen et al. 1998; 54: Claydon 2004; 55: Pratchett 2005; 56: Reavis & Copus 2011; 57: Yabuta & Kawashima 1997; 58: Ferry-Graham et al. 2001; 59: Bellwood et al. 2010; 60: Donaldson 1990; 61: Kadota et al. 2011; 62: Donaldson 1989; 63: Kane et al. 2009; 64: Clemente et al. 2010; 65: Bellwood et al. 2006a; 66: Taquet et al. 2007; 67: Sakashita 1992; 68: Hernaman et al. 2009; 69: Kramer et al. 2012b; 70: Depczynski & Bellwood 2003; 71: Herler et al. 2011; 72: Allen & Munday 1995; 73: White & Brown 2013; 74: Munday et al. 2002; 75: Karplus 1979; 76: Brooker et al. 2010; 77: Cole et al. 2008; 78: Randall & Goren 1993; 79: Daroonchoo 1991; 80: Sano et al. 1984; 81: Hagiwara & Winterbottom 2007; 82: Winterbottom 1984; 83: Clark et al. 2000; 84: Reavis & Barlow 1998; 85: Randall 1998; 86: Clements & Choat 1997; 86: Rimmer & Wiebe 1987; 87: DiSalvo et al. 2007; 88: Williams & Williams 1986; 89: Bellwood et al. 2003; 90: Muñoz et al. 2006; 91: Ferry-Graham et al. 2002; 92: Choat et al. 2004; 93: Lek et al. 2011; 94: Pratchett et al. 2001; 95: Debenay et al. 2011; 96: Connell 1998; 97: Nanami & Yamada 2008; 98: Kiso & Mahyam 2003; 99: Sweatman 1992; 100: Salini et al. 1990; 101: Bonin et al. 2011; 102: Clark et al. 1998; 103: Clark & Pohle 1992; 104: López-Peralta & Arcilla 2002; 105: Harmelin-Vivien 1979; 106: Kokita & Mizota 2002; 107: Wantiez & Kulbicki 1995; 108: Blaber 1977; 109: McCormick

1995; 110: Randall & Myers 2002; 111: Randall 2004; 112: Letourneur 1996; 113: Golani & Galil 1991; 114: Sabrah & El-Ganainy 2009; 115: Mehta 2008; 116: Reece et al. 2010; 117: Mequila & Campos 2007; 118: Boaden & Kingsford 2012; 119: Russell 1997; 120: Hiatt & Strasburg 1960; 121: Tamaki et al. 1992; 122: Moyer & Sano 1987; 123: Metian et al. 2013; 124: Sivakumar & Ramaiyan 1987; 125: Mooi & Gill 2004; 126: Ang & Manica 2010; 127: Moyer & Nakazono 1978; 128: Pyle 2003; 129: Mantyka & Bellwood 2007; 130: Hensley & Allen 1977; 131: Frédéricich et al. 2009; 132: Williams & Hatcher 1983; 133: Fautin & Allen 1992; 134: Fricke & Fricke 1977; 135: Hobson 1991; 136: Tribble & Nishikawa 1982; 137: Wenger et al. 2012; 138: Allen 1991; 139: Sackley & Kaufman 1996; 140: Gluckmann & Vandewalle 1998; 141: Allen & Rajasuriya 1995; 142: Randall & Allen 1977; 143: Elliot & Bellwood 2003; 144: Ceccarelli 2007; 145: Bellwood et al. 2006b; 146: Horn 1989; 147: Öhmann et al. 1998; 148: Lewis 1998; 149: Hamilton & Dill 2003; 150: Kohda 1981; 151: Russ 1987; 152: Allen 1987; 153: Feeney et al. 2012; 154: Harmelin-Vivien & Bouchon 1976; 155: Shallenberger & Madden 1973; 156: Marguillier et al. 1997; 157: Muñoz et al. 2011; 158: Longenecker 2007; 159: Barros et al. 2008; 160: Holmes & McCormick 2011; 161: Stewart & Jones 2001; 162: Craig 2007; 163: Woodland 1990; 164: Fox & Bellwood 2008; 165: Cvitanovic & Bellwood 2009; 166: Fox et al. 2009; 167: Hoey et al. 2013; 168: Grubich et al. 2008; 169: Hajisamae et al. 2003; 170: Gronell 1984; 171: Iyer 2012; 172: Leysen et al. 2011; 173: Lourie & Randall 2003; 174: Foster & Vincent 2004; 175: Horinouchi et al. 2012; 176: Kizhakudan & Gomathy 2007; 177: Sikkell & Sikkell 2012; 178: Sikkell 1990; 179: Beumer 1978; 180: Sabrah et al. 2006; 181: Aydin 2011; 182: Clark & Pohle 1996; 183: Silberschneider & Booth 2001; 184: Hadley Hansen 1986