

Feeding ecology and ontogenetic dietary shift of yellowstripe goatfish *Mulloidichthys flavolineatus* (Mullidae) at Reunion Island, SW Indian Ocean

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ABSTRACT: We analyzed stomach contents and muscle isotopic composition ($\delta^{13}\text{C}$, $\delta^{15}\text{N}$) of yellowstripe goatfish *Mulloidichthys flavolineatus* from a coral reef to investigate size-related dietary changes and resource or habitat partitioning. Juveniles (<12 cm total length [TL]), young adults (12 ≤ TL < 17 cm) and adults (≥17 cm TL) showed a high diet overlap, especially between juveniles and young adults. According to stomach contents analysis, *M. flavolineatus* widens its prey spectrum with increasing size from a common prey pool that includes polychaetes, tanaids and harpacticoid copepods. We observed a significant increase in $\delta^{13}\text{C}$ values (from $-17.1 \pm 0.5\text{‰}$ for juveniles to $-10.7 \pm 0.8\text{‰}$ for adults), which were correlated to fish size. Adults ($\delta^{15}\text{N}$ mean of $11.1 \pm 1.8\text{‰}$) were one trophic level above juveniles and young adults (7.7 ± 0.5 and $7.4 \pm 0.5\text{‰}$, respectively). These patterns of isotopic changes confirmed ontogenetic dietary shifts. However, trophodynamics can be influenced by physiological factors such as growth and sexual maturity. *M. flavolineatus* shift from a pelagic to a macrobenthic diet, which is equilibrated at the adult stage. Results from combined stomach contents (prey volume) and stable isotope analyses suggested a dominant contribution of polychaetes (macrofauna), possibly through selective feeding. Conversely, on the basis of prey volume and stable isotope data, meiofauna did not feature significantly in the diet despite their high abundance in stomachs. Two adult groups were distinguished based on their $\delta^{15}\text{N}$ values (11.9 ± 0.8 and $7.8 \pm 0.6\text{‰}$), indicating possible stage-specific partitioning in habitat use inside the reef.

KEY WORDS: Yellowstripe goatfish · Ontogenetic diet shift · Stomach contents · Stable isotopes · Mixing model · Southwestern Indian Ocean

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INTRODUCTION

Goatfish (Perciformes, Mullidae) are widespread tropical and subtropical fish, and are particularly abundant in shallow-water ecosystems (Munro 1976, McCormick 1995) that serve as nurseries or habitats. Numerous fish species are found in these environments mainly due to favorable conditions both in terms of prey availability and shelter (Parrish 1989,

Robertson & Blaber 1992). Of the 10 goatfish species inhabiting Reunion Island reefs (southwestern Indian Ocean), the yellowstripe goatfish *Mulloidichthys flavolineatus* is the most common (Letourneur 1996). Despite their importance for artisanal fisheries, little is known about the feeding ecology, recruitment and habitat preferences of this species. Such data are critical for designing efficient conservation and management policies.

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Knowledge of the trophic interactions between fish and their food resources is of immediate significance to understanding processes that structure shallow-water tropical communities. A number of studies on goatfish diets are reported from the Pacific (Sorden 1982, Lukoschek & McCormick 2001, Sampey et al. 2007), the tropical West Atlantic (Munro 1976, Humann & DeLoach 2002), the Mediterranean and the Red Sea (Ben-Eliahu & Golani 1990, Golani & Galil 1991, Wahbeh 1992). Published data from the Indian Ocean are scarce and concern only the genus *Upeneichthys* (Platell et al. 1998). Sympatric mullid species can greatly differ in their depth distribution, foraging substratum selection and feeding modes (Gosline 1984, McCormick 1995, Platell et al. 1998). While main food items vary widely among species, all Mullidae are benthic carnivores, using their sensory hyoid barbels to drive out prey. *Mulloidichthys flavolineatus* is considered to be a specialist benthic predator, foraging mainly on sediments around reefs (McCormick 1995). Past studies from the Hawaiian Midway Islands (Sorden 1982) and the northern Red Sea (Ben-Eliahu & Golani 1990, Wahbeh 1992) provide general descriptions of prey items consumed by *M. flavolineatus*. While Sorden (1982) and Ben-Eliahu & Golani (1990) focused on polychaetes in stomach contents, Wahbeh (1992) investigated the overall diet of *M. flavolineatus* and reported the presence of fish eggs, nematodes, molluscs and arthropods. Wahbeh (1992) also reported differences in diet composition between adults of different age groups: while fish eggs and arthropods were dominant in younger *M. flavolineatus*, nematodes and molluscs were more commonly found in older individuals. Such ontogenetic changes in diet composition have been shown to be important for many fish species, in particular for benthic carnivores (McCormick 1998, Lukoschek & McCormick 2001). To date, ontogenetic dietary changes in early post-settlement *M. flavolineatus* have not been reported.

Stomach content analysis provides a taxonomic resolution of an individual's recent (hours to days) feeding habits. However, this approach only presents a snapshot of the food habits and includes some uncertainties when prey identification is difficult or impossible due to digestion. More recently, combined stomach content and stable isotope analyses have been used in trophic ecology (e.g. Wells et al. 2008). Stable isotopic compositions of consumer tissues can often be predictably related to stable isotopic compositions of the corresponding diet (DeNiro & Epstein 1978). This approach relies on the consistent enrichment of stable carbon and nitrogen isotopes that occurs with each successive trophic level due to isotopic fractionation during assimilation (e.g. Minagawa & Wada 1984). The mean trophic shift of consumers' tissue relative to their food

source is around 2.9 to 3.8‰ for $\delta^{15}\text{N}$ values and 0.0 to 1.5‰ for $\delta^{13}\text{C}$ values in marine systems (Michener & Schell 1994, Vander Zanden & Rasmussen 2001).

Unlike stomach content analysis, stable isotope ratios of an organism's tissue provide information on the time-integrated assimilated diet. If the isotopic composition of available prey differs with habitat, stable isotopes can provide a powerful tool to investigate animal movement (Fry & Arnold 1982, Hesslein et al. 1991). This approach depends on distinctive isotopic signatures of primary producers found in specific habitats which are propagated through a local food web (Michener & Schell 1994) and has been successfully used to trace the early life stages of fish (Herzka et al. 2002a) and migration of shrimp (Fry 1983, Fry et al. 2003). Stable isotope ratios can thus serve as natural chemical tracers for discriminating recent immigrants from those that have partially or fully equilibrated to the isotopic composition of the prey consumed in a new habitat (Fry et al. 1999, Herzka et al. 2002b).

We investigated the feeding ecology of *Mulloidichthys flavolineatus* from a coral reef at Reunion Island. A wide size range of individuals was sampled across a fringing reef in order to describe possible ontogenetic changes that occur in their diets. Using stomach content and stable isotope analyses, the present study was designed to address the following questions: (1) Are there ontogenetic variations in diet composition of *M. flavolineatus*? If so, (2) can ontogenetic change patterns be used to infer possible resource or habitat partitioning within the *M. flavolineatus* population?

MATERIALS AND METHODS

Study site and sample collection. Reunion Island (21° 07' S, 55° 32' E) is situated in the SW Indian Ocean, 800 km east of Madagascar (Fig. 1) and is characterized by a tropical oceanic climate with a wet summer season extending from January to April. The continental shelf is very narrow, and coral reefs fringe less than 10% of the coastline. The present study was carried out at the La Saline fringing reef (4 km long, 500 m maximum width), located on the west coast of the island (Fig. 1). Soft-bottom habitats are composed of coarse sand scattered with coral fragments.

After an offshore pelagic larval phase, *Mulloidichthys flavolineatus* cross the reef front at 10.7 ± 0.9 cm total length (TL) to recruit in back-reef and inner-reef flat habitats of Reunion Island (Durville 2002). Post-larval *M. flavolineatus* recruit to the La Saline reef during the summer season (Durville 2002). Fish were caught in March 2007 inside a 0.5 km² area (Fig. 1) using a 3 × 10 m wide barrier net with 1 cm

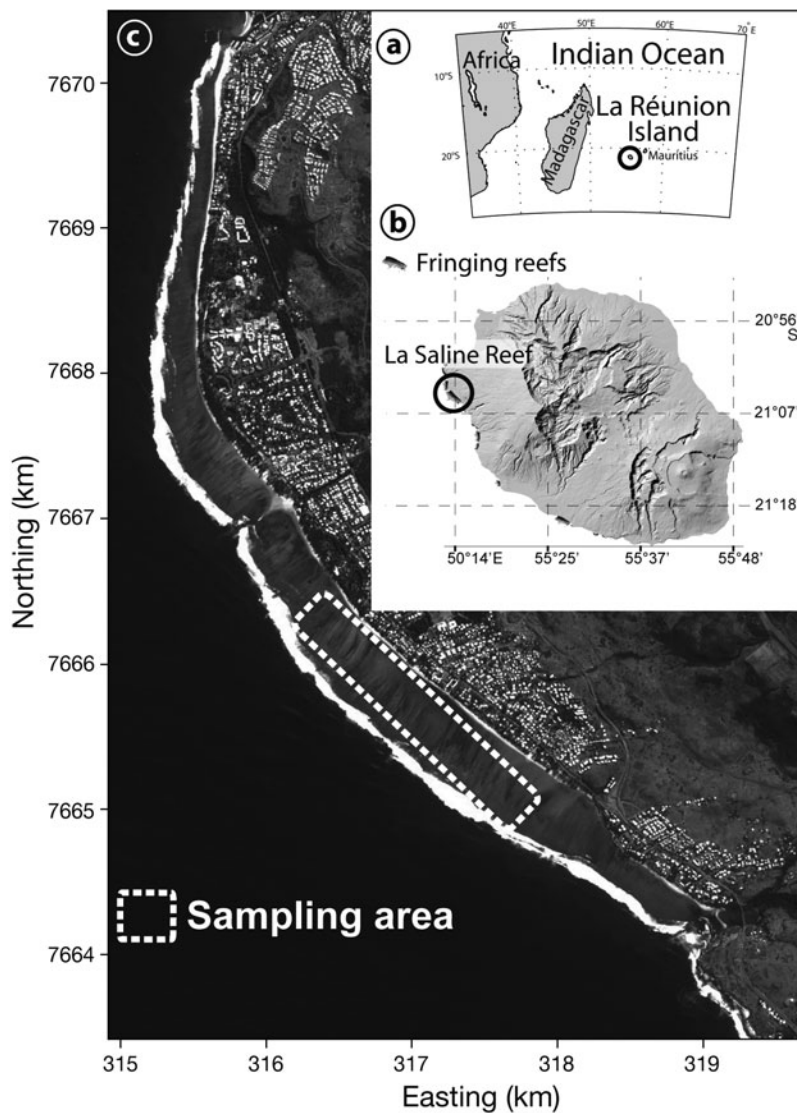


Fig. 1. La Saline reef, Réunion Island, southwestern Indian Ocean. Dashed rectangle indicates the sampling area

mesh size and spear guns. After collection, fish were placed in an ice bath, transported to the laboratory and frozen at -20°C . Size (TL), wet mass, sex and gonad weight were recorded for each individual before further analysis.

Particulate organic matter (POM) was used as a proxy for pelagic food sources and was obtained by filtering oceanic seawater (at 3 and 8 km from the coast) using 25 mm GF/C precombusted filters (4 h at 450°C). Soft-bottom macrofauna (organisms ≥ 1 mm: annelids, arthropods and molluscs) were sampled using an air-lift system and used as an end member proxy of macrobenthic food sources. Although we found meiofauna prey items (organisms $< 500\ \mu\text{m}$) in *Mulloidichthys flavolineatus* stomach contents, we could not exclude the presence of other

microbenthic organisms (organisms $< 45\ \mu\text{m}$: bacteria, ciliates and micro-phytobenthos). Hence, we used reef sediment total organic matter (STOM) as a proxy of micro/meiobenthos food sources. All associated samples were obtained at the time of the fish sampling.

Stomach content analysis. Stomachs were removed, thawed, drained and treated as follows. Total mass of the content of each stomach was determined. Relative mass was calculated as stomach content wet mass divided by body wet mass, minus stomach content wet mass. Stomach contents were sorted into main prey categories and identifiable prey items were counted (see lines in Table 1). Due to the very small size of prey, the mean volume of each prey taxa was assessed from width and length of subsampled individuals, converted into basic geometric forms. Prey were identified to the lowest taxonomic level possible and assigned to one of 4 prey types: vegetal, micro-, meio- and macrofauna (see Table 1).

To characterize the diet composition of each fish size group, 3 dietary indices have been used for each identified prey item i : frequency of occurrence in stomachs (O_i = number of stomachs including prey category i divided by total number of non-empty stomachs), numerical importance (N_i = total number of prey category i divided by total number of prey items) and volume (V_i = total volume of prey category i divided by total volume of prey items). Sediment volume was also estimated since it comprised a significant proportion of most stomach contents.

Diet overlap between the size groups was assessed using the Morisita-Horn index (see Krebs 1998). This overlap index was calculated from abundance and volume data of prey categories identified at the family level, using the following formula:

$$C_{mh} = \frac{2 \sum_{i=1}^S p_{A,i} \times p_{B,i}}{\sum_{i=1}^S p_{A,i}^2 + \sum_{i=1}^S p_{B,i}^2} \quad (1)$$

where C_{mh} = Morisita-Horn index of overlap between predator A and B; S = total number or volume of identified prey species in the feeding regime of both predators; $p_{A,i}$ = number or volume of individuals of prey species i consumed by predator A; and $p_{B,i}$ = number or

volume of individuals of prey species i consumed by predator B . C_{mh} ranges from 0 (no prey in common) to 1 (complete overlap). A significant overlap is commonly assumed for index values greater than 0.6 (Zaret & Rand 1971).

Stable isotope analyses. Dorsal white muscle tissue was sampled from individuals for carbon and nitrogen stable isotope analyses. To obtain STOM samples, detritus, macroalgae fragments and macrofauna were removed from sediment cores (top 2 cm of sediment, core $\varnothing = 3.5$ cm) and were acidified (1 N HCl) to eliminate inorganic carbon for $\delta^{13}\text{C}$ measurements. Samples were freeze-dried and ground to a homogenous powder using an automated grinder. POM filters were pre-treated and analyzed according to Lorrain et al. (2003). Macrofauna samples were analyzed following Kolasinski et al. (2008). Lipids were not removed as *Mulloidichthys flavolineatus* white muscle, macrofauna and STOM have a low lipid content (C/N ratio of all samples were <4), which does not affect $\delta^{13}\text{C}$ values of these measurements (Post et al. 2007). Fish muscle (1.5 mg), POM (whole filter), STOM (100 mg) and macrofauna (1.5 to 2.4 mg) samples were weighed into tin capsules and analyzed in duplicate.

Carbon and nitrogen content and isotopic composition were analyzed at the Stable Isotope Laboratory, GNS Science, New Zealand, using a Europa Geo 20/20 isotope ratio mass spectrometer (IRMS; PDZ Europa) interfaced to an ANCA-SL elemental analyzer (EA; PDZ Europa) in continuous flow mode (EA-IRMS). Carbon dioxide and nitrogen gases were separated on a gas chromatography (GC) column at 68°C and analyzed for isotopic abundance. The $^{13}\text{C}/^{12}\text{C}$ and $^{15}\text{N}/^{14}\text{N}$ compositions are expressed in conventional delta notation in per mille (‰) relative to the levels of $\delta^{13}\text{C}$ in Vienna Pee Dee Belemnite (VPDB) and $\delta^{15}\text{N}$ in atmospheric air, according to the following equation:

$$\delta X = [(R_{\text{sample}}/R_{\text{standard}}) - 1] \times 1000 \quad (2)$$

where X is $\delta^{13}\text{C}$ or $\delta^{15}\text{N}$ and R is the ratio of heavy to light isotope ($^{13}\text{C}/^{12}\text{C}$ or $^{15}\text{N}/^{14}\text{N}$). Repeated measurement of an internal standard exhibited a precision of $\pm 0.1\%$ for $\delta^{13}\text{C}$ and $\pm 0.3\%$ for $\delta^{15}\text{N}$.

Data analysis. *Mulloidichthys flavolineatus* samples ranged from 9.3 to 24.9 cm TL (median = 11.7 cm). Three groups were recognized based on length frequency modes and gonad development: juveniles (<12 cm TL) newly recruited to the reef and presenting undifferentiated gonads, young adults ($12 \leq \text{TL} < 17$ cm) which are both non-sexually and sexually differentiated, and adults (≥ 17 cm TL) with sexually mature gonads. Multivariate analyses were performed on prey item volume for each size group to assess ontogenetic changes in yellowstripe goatfish diet. Data

were log-transformed to minimize effects of dominant consumed items. A similarity matrix based on Bray-Curtis coefficients was classified by hierarchical agglomerative clustering using the unweighted pair group mean arithmetic (UPGMA) linking method. All calculations and tests were performed with PRIMER 6.0 (Clarke & Gorley 2006). Differences in feeding regime were tested using non-parametric procedures when data did not fulfill homoscedasticity requirements.

Sex effects on $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values were investigated within size groups using univariate analysis of covariance (ANCOVA) with TL as a covariate to take into account size effects on stable isotope values. $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values were analyzed separately to investigate size effects using regression models. Trophic level was calculated for each size group following Hobson & Welch (1992):

$$\text{Trophic level} = 2.5 + (\delta^{15}\text{N}_{\text{observed}} - 7.5)/2.3 \quad (3)$$

where the average macrofauna trophic level is 2.5, $\delta^{15}\text{N}_{\text{observed}}$ is the $\delta^{15}\text{N}$ value of the yellowstripe goatfish, the average $\delta^{15}\text{N}$ value of the end member macrofauna is 7.5‰ and the $\delta^{15}\text{N}$ enrichment value per trophic level is 2.3‰. Post (2002) reported that $\delta^{15}\text{N}$ enrichment can range from 2 to 5‰ in aquatic ecosystems. Since fractionation values are undetermined for *Mulloidichthys flavolineatus*, the trophic fractionation of 2.3‰ (McCutchan et al. 2003) represents a mean value for aquatic organisms feeding on either high (animal, microbial) or low (algal) protein content diets.

A 3-source isotope mixing model (Phillips & Gregg 2003) was used to estimate the proportional contributions of pelagic, micro/meiobenthic and macrobenthic food sources to the diets of juveniles and young adults. Pelagic, micro/meiobenthic and macrobenthic sources are the average isotopic signatures of POM, STOM and macrofauna, respectively. Trophic enrichment factors of 2.3 and 0.5‰ were used for $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$, respectively. A von Bertalanffy model for the growth rates of *Mulloidichthys flavolineatus* (Holland et al. 1993) indicated that minimum size of juveniles (8.7 cm TL), young adults (12 cm TL) and adults (17 cm TL) corresponded to ages of around 10, 12 and 18 mo, respectively. Hence, a 2-source mixing model was used to investigate the respective contribution of micro/meiobenthic and macrobenthic food sources to the diet of adults since we assume that pelagic carbon and nitrogen have been totally replaced by a corresponding benthic signature in muscle tissue around 8 mo after settlement inside the reef environment. The 2-source isotope mixing model (Fry 2006) was calculated with a trophic enrichment factor (I) of 0.5‰:

$$\% \text{macrobenthic} = 100 \times \frac{(\delta^{13}\text{C}_{\text{observed}} - \delta^{13}\text{C}_{\text{micro/meiobenthic}} - D)}{(\delta^{13}\text{C}_{\text{macrobenthic}} - \delta^{13}\text{C}_{\text{micro/meiobenthic}})} \quad (4)$$

and $\% \text{micro/meiobenthic} = 100 - \% \text{macrobenthic}$. The mixing model calculates all feasible combinations of sources that could explain the consumer isotope value, thereby placing limits on the dietary contributions of each source. Various diet contributions were tested using ANCOVA with TL as a covariate.

RESULTS

Stomach content analysis

Overall diet

We analyzed stomach contents of 110 yellowstripe goatfish. No empty stomachs were found. Average volume of the juvenile's stomach content ($n = 64$) was 55.7 mm^3 (ranging from <0.1 to 207.1 mm^3). The diet of juveniles was dominated by annelids and arthropods (56.0 and 36.6% of volume, respectively), while nematodes contributed only a minor part (5.7% of volume) (Table 1). Stomach content volume in young adults ($n = 26$) was on average 154.0 mm^3 (ranging from 10.9 to 316.2 mm^3). Annelids were the most important food source for this group (70.9% of volume), while arthropods ranked second (22.5%) and echinoderms and nematodes represented 2.4 and 2.3%, respectively, of total volume. The average stomach content volume of the 20 adults studied was 146.8 mm^3 (ranging from 0.5 to 332.5 mm^3). Annelids were the most important food source for adults (annelids: 52.1%; arthropods: 38.7%; echinoderms: 6.5% of volume). On the basis of prey number, the trend differed since arthropods were the most common food source in adults (55.8%) and juveniles (47.5%) diets, but ranked second for young adults (44.7%). Nematodes were most common for juveniles (46.7% by number) and young adults (47.4%), and decreased in adults (31.0%). The contribution of polychaetes to the *Mulloidichthys flavolineatus* diet was low for all size groups, increasing from 1.7% for juveniles to 3.1 and 3.6% for adults and young adults, respectively.

Sediment occurred in stomachs more frequently in young adults and adults (88 and 90% of individuals, respectively) than in juveniles (11%). Differences in sediment occurrence in stomach contents was significant between juveniles and adults ($\chi^2 = 41.86$, $p < 0.01$) and between juveniles and young adults ($\chi^2 = 46.58$, $p < 0.01$), but not between adults and young adults ($\chi^2 = 0.10$, $p = 0.76$). Sediment proportion in each stomach content's total volume decreased from young

adults (mean of 19.0%) to adults (11.8%) and juveniles (0.5%). Differences in ingested sediment volume was significant between juveniles and both young adults (Mann-Whitney test: $Z = 7.59$, $p < 0.01$) and adults ($Z = 7.12$, $p < 0.01$); no difference was observed between adults and young adults ($Z = 1.73$, $p = 0.083$).

Prey taxonomic analysis

Juveniles. Ten prey families and 8994 prey items were found in stomachs of the 64 juveniles sampled (Table 1), with a mean (\pm SE) of 140.5 ± 31.8 prey items per stomach. Malacostracans were the dominant prey, followed by Adenophora nematodes and polychaetes. Major food items were harpacticoid copepods (37.1% by number, 27.3% by volume), tanaids of the Leptocheliidae family (10.4% by number, 9.3% by volume) and nematodes of the Chromadoridae family (46.7% by number, 5.7% by volume). In total, these prey contributed 94.2 and 42.3% by number and volume, respectively. Despite the low number of polychaetes (1.7%), they were the dominant prey by volume (56.0%). We note that foraminifera of the Miliolidae and Amphisteginidae families occurred frequently in stomach contents (48.4 and 64.1%, respectively) although their abundance and volume were low. Naticidae occurred in 26.6% of stomach contents and was the unique gastropod family consumed by juveniles.

Young adults. Seventeen prey families and 4040 prey items were found in the stomachs of the 26 young adults sampled (Table 1), with a mean (\pm SE) of 155.4 ± 37.6 prey items per stomach. Malacostracans occurred in 25 samples (96.2%) and contributed 44.7% by number and 22.5% by volume. Harpacticoid copepods and tanaids of the Leptocheliidae family were the main malacostracan components; together they represented 43.2% of the diet by number and 21.3% by volume. Other malacostracan prey included amphipods of the Melitidae family (observed in 46.2% of the stomachs) and some undetermined decapods and isopods. Nematodes of the Chromadoridae family dominated the young adult's diet (47.4% by number) but, owing to their small size, only represented 2.3% of the diet by volume. Polychaetes did not contribute significantly to the total number of prey (3.6%) but they constituted the main prey item by volume (70.9%). Gastropods (Nassariidae, Cypraeidae and Naticidae) were also present in the diet; Naticidae were found in 42.3% of the stomachs.

Adults. A total of 2706 prey items belonging to 16 families were recorded in the stomachs of the 20 adults sampled (Table 1), with a mean (\pm SE) of 135.3 ± 49.5 prey items per stomach. Malacostracans occurred in

Table 1. *Mulloidichthys flavolineatus*. Number (N, %), volume (V, %) and occurrence (O, %) of prey taxa or categories recovered from juvenile (n = 64), young adult (n = 26) and adult (n = 20) stomach contents. Un.: unidentified; Micro: microfauna; Meio: meiofauna; Macro: macrofauna

Phylum Class	Order	Family	Type	Juveniles			Young adults			Adults												
				N	%	V	N	%	V	N	%	V	N	%	O							
Sarcomastigophora																						
Granuloreticulosea	Foraminiferida	Miliolidae	Micro	176	2.0	2.8	0.1	31	48.4	23	0.6	0.3	<0.1	10	38.5	19	0.7	0.1	<0.1	9	45.0	
		Amphisteginidae	Micro	125	1.4	41.6	1.2	41	64.1	77	1.9	0.7	<0.1	20	76.9	103	3.8	15.9	0.4	17	85.0	
Total Sarcomastigophora				301	3.3	44.4	1.2	51	79.7	100	2.5	1.0	<0.1	24	92.3	122	4.5	16.0	0.4	17	85.0	
Nematoda																						
Adenophorea	Desmodorida	Chromadoridae	Meio	4203	46.7	202.1	5.7	56	87.5	1915	47.4	92.1	2.3	21	80.8	838	31.0	42.5	1.2	17	85.0	
Annelida																						
Polychaeta			Macro	157	1.7	1997.6	56.0	44	68.8	146	3.6	2838.0	70.9	23	88.5	77	2.8	1899.7	52.1	16	80.0	
Echinodermata																						
Ophiuroidea			Macro							20	0.5	94.2	2.4	6	23.1	50	1.8	235.6	6.5	8	40.0	
Mollusca																						
Gastropoda	Neogastropoda	Nassaridae	Macro							2	<0.1	1.0	<0.1	2	7.7	11	0.4	5.8	0.2	6	30.0	
	Neotaenioglossa	Cypraeidae	Macro							3	0.1	<0.1	<0.1	3	11.5						<0.1	
		Naticidae	Macro	27	0.3	1.8	<0.1	17	26.6	14	0.3	1.2	<0.1	11	42.3	45	1.7	7.4	0.2	13	65.0	
Total Mollusca				27	0.3	1.8	<0.1	17	26.6	19	0.5	2.4	0.1	12	46.1	56	2.1	13.2	0.4	14	70.0	
Arthropoda																						
Malacostraca	Harpacticoida		Meio	3338	37.1	972.5	27.3	57	89.1	1254	31.0	415.9	10.4	25	96.2	319	11.8	163.0	4.5	14	70.0	
	Tanaidacea		Meio	934	10.4	332.4	9.3	46	71.9	494	12.2	436.8	10.9	25	96.2	314	11.6	394.5	10.8	15	75.0	
	Decapoda		Macro							5	0.1	19.2	0.5	3	11.5	2	0.1	58.1	1.6	2	10.0	
	Amphipoda	Melitidae	Meio	3	<0.1	1.6	<0.1	2	3.1	51	1.3	27.5	0.7	12	46.2	40	1.5	76.3	2.1	10	50.0	
		Un. amphipods	Macro												4	0.1	7.9	0.2	2	10.0		
	Isopoda	Un. isopods	Macro	4275	47.5	1306.5	36.6	57	89.1	1807	44.7	902.0	22.5	25	96.2	1511	55.8	1409.2	38.7	19	95.0	
Total Arthropoda										2	<0.1	3.0	0.1	2	7.7	1	<0.1	1.5	<0.1	1	5.0	
Total unidentified animals																						
Heterokontophyta																						
Phaeophyceae										17	0.4	71.2	1.8	10	38.5	6	0.2	25.1	0.7	6	30.0	
Rhodophyta																						
Rhodophyceae										2	<0.1	0.1	<0.1	1	3.8	39	1.4	1.2	<0.1	6	30.0	
Chlorophyta										1	<0.1	0.5	<0.1	1	3.8	6	0.2	1.1	<0.1	1	5.0	
Unidentified plants										11	0.3	0.5	<0.1	2	7.7							
Grand total				8992		3565.6	64	64		4040		4005		26		2706		3645.1		20		

95% of stomachs; they dominated the diet by number (55.8%) and ranked second by volume (38.7%). Isopods (30.5% by number, 19.5% by volume) and tanaids of the Leptocheliidae family (11.6% by number, 10.8% by volume) were the dominant malacostracan prey. Among other malacostracan families, harpacticoid copepods and Melitidae had the highest occurrence (70 and 50%, respectively), and polychaetes ranked first by volume (52.1%). Nematodes of the Chromadoridae family occurred in 85% of the stomachs, and they contributed 31.0% to diet by number but only 1.2% by volume. Amphisteginidae foraminifers were observed in 85% of stomachs but their contribution to diet by number (3.8%) and volume (0.4%) were low. Gastropod prey were represented by Nassariidae (40% occurrence) and Naticidae (60% occurrence), but their diet contribution was negligible (0.4 and 1.7% by number, respectively, 0.2% each by volume).

Comparison of feeding regimes

Differences in prey number were significant between juveniles (mean of 4.9 ± 1.9) and both young adults (6.9 ± 2.0) and adults (7.2 ± 2.7 ; Mann-Whitney: $Z = 3.98$, $p < 0.01$ and $Z = 3.48$, $p < 0.01$, respectively). No significant differences were observed between young adults and adults ($Z = 0.81$, $p = 0.426$). The effect of fish size on prey number was not significant (Kruskal-Wallis: $H = 1.58$, $p = 0.451$); however, a significant size effect was observed for both total volume ($H = 26.90$, $p < 0.01$) and relative mass ($H = 20.14$, $p < 0.01$). For each pairing of size groups, Morisita-Horn index values calculated both on number and volume of prey items were >0.6 (Table 2), indicating a strong overlap between the different size groups. Juveniles and young adults had the highest overlap indices among all size-group pairs (0.993 based on prey number and 0.943 based on prey volume). Fish pairs containing the young adult group presented the highest index values. Diet analysis based on volume (Table 1) indicated that *Mulloid-*

ichthys flavolineatus gradually shifted to a mixed meio-/macrofauna diet in juveniles (43.5 and 56.1% of prey total volume, respectively) to a macrofauna-dominated diet in young adults (24.3% for meiofauna, 74.2% for macrofauna) and adults (19 and 80.4%). Moreover, clusters calculated from prey volumes showed that all groups shared 3 common prey taxa: polychaetes (macrofauna), and Leptocheliidae tanaids and harpacticoid copepods (meiofauna) (Fig. 2). Together, these prey dominated the diet by volume (92.6% of total prey volume in juveniles, 95.8% in young adults and 67.4% in adults) (Table 1). In juvenile and young adult diets, meiofauna taxa were added to the common prey pool: Chromadoridae nematodes in both groups, Amphisteginidae foraminifers in juveniles (Fig. 2a) and Melitidae amphipods in young adults (Fig. 2b). Melitidae amphipods were maintained into the common prey pool in adult group and macrofauna taxa (ophiuroids) were added (Fig. 2c). Surprisingly, phaeophyta had a noticeable presence in stomach contents (from 30.0 to 38.5% of occurrence) (Table 1).

Stable isotope analysis

A total of 103 muscle samples of *Mulloidichthys flavolineatus* were analyzed for stable isotopes ($\delta^{13}\text{C}$ and $\delta^{15}\text{N}$). Mean values for each size group are presented in Table 3. There were significant variations in $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values among size groups (ANCOVA: $\delta^{13}\text{C}$, $F = 12.81$, $p < 0.001$; $\delta^{15}\text{N}$, $F = 9.14$, $p < 0.001$). Newman-Keuls *post hoc* tests revealed differences in $\delta^{13}\text{C}$ values between juveniles (mean $\delta^{13}\text{C}$ of $-17.1 \pm 0.5\text{‰}$), young adults ($-14.1 \pm 2.3\text{‰}$) and adults ($-10.7 \pm 0.8\text{‰}$). For $\delta^{15}\text{N}$, significant differences were observed between adults (mean $\delta^{15}\text{N}$: $11.1 \pm 1.8\text{‰}$) and the 2 smaller size groups (juveniles and young adults, mean $\delta^{15}\text{N}$ of 7.7 ± 0.5 and $7.4 \pm 0.5\text{‰}$, respectively). There were significant differences in the $\delta^{13}\text{C}$ values of young adults with undifferentiated gonads (mean of $-15.2 \pm 2.2\text{‰}$) and sexually differentiated young adults ($-13.8 \pm 0.8\text{‰}$; $F = 21.32$, $p < 0.001$). There were no significant differences in $\delta^{13}\text{C}$ values between male and female young adults (Kruskal-Wallis: $H = 0.22$, $p = 0.638$). No significant sex effects were observed in $\delta^{13}\text{C}$ values of adults ($F = 2.32$, $p = 0.115$), nor in $\delta^{15}\text{N}$ values of both young adults and adults ($F = 1.93$, $p = 0.155$ and $F = 1.08$, $p = 0.385$, respectively).

Table 2. *Mulloidichthys flavolineatus*. Morisita-Horn overlap index of dietary composition based upon number of prey items (values in *italics*) and volume (values in **bold**) for each size group

Size group	Juveniles	Young adults	Adults
Juveniles		<i>0.993</i>	<i>0.685</i>
Young adults	0.943		<i>0.715</i>
Adults	0.865	0.908	

$\delta^{13}\text{C}$ isotopes

$\delta^{13}\text{C}$ values of juveniles ranged from -17.6 to -14.3‰ (Fig. 3a). A slight increase in $\delta^{13}\text{C}$ values with TL was

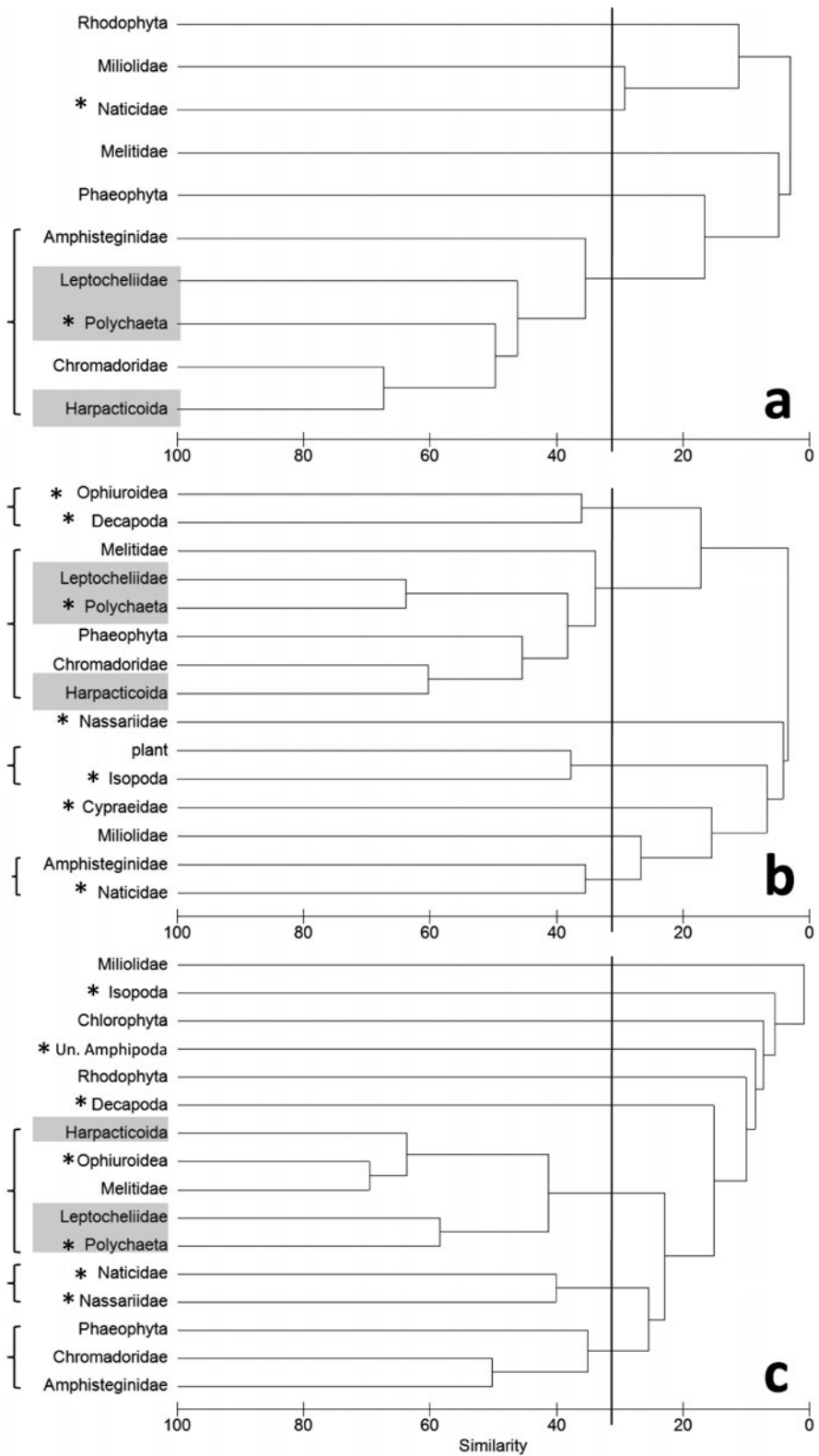


Fig. 2. *Mulloidichthys flavolineatus*. Hierarchical cluster analysis of stomach contents based on a Bray-Curtis similarity matrix of prey taxa volume, performed for each size group: (a) juveniles, (b) young adults and (c) adults. Vertical bars indicate the similarity level (%) selected to define groups. Grey areas illustrate the common pool of prey taxa which dominated the diet by volume. Asterisks distinguish macrofauna taxa from micro- and meiofauna taxa (no asterisks). Un.: unidentified

Table 3. *Mulloidichthys flavolineatus*. Mean (\pm SD) and range of carbon and nitrogen stable isotope values and trophic level estimation of each size group. TL = total length, n = number of samples

Size group	Stable isotope	Mean (‰)	Range (‰)	Trophic level
Juveniles <12 cm TL (n = 59)	$\delta^{13}\text{C}$	-17.1 ± 0.5	-17.6: -14.3	2.6 ± 0.2
	$\delta^{15}\text{N}$	7.7 ± 0.5	6.7: 8.7	
Young adults 12 \leq TL < 17 cm (n = 26)	$\delta^{13}\text{C}$	-14.1 ± 2.3	-17.6: -11.2	2.5 ± 0.1
	$\delta^{15}\text{N}$	7.4 ± 0.5	7.1: 8.8	
Adults ≥ 17 cm TL (n = 18)	$\delta^{13}\text{C}$	-10.7 ± 0.8	-12.3: -9.5	3.8 ± 0.7
	$\delta^{15}\text{N}$	11.1 ± 1.8	7.3: 12.9	

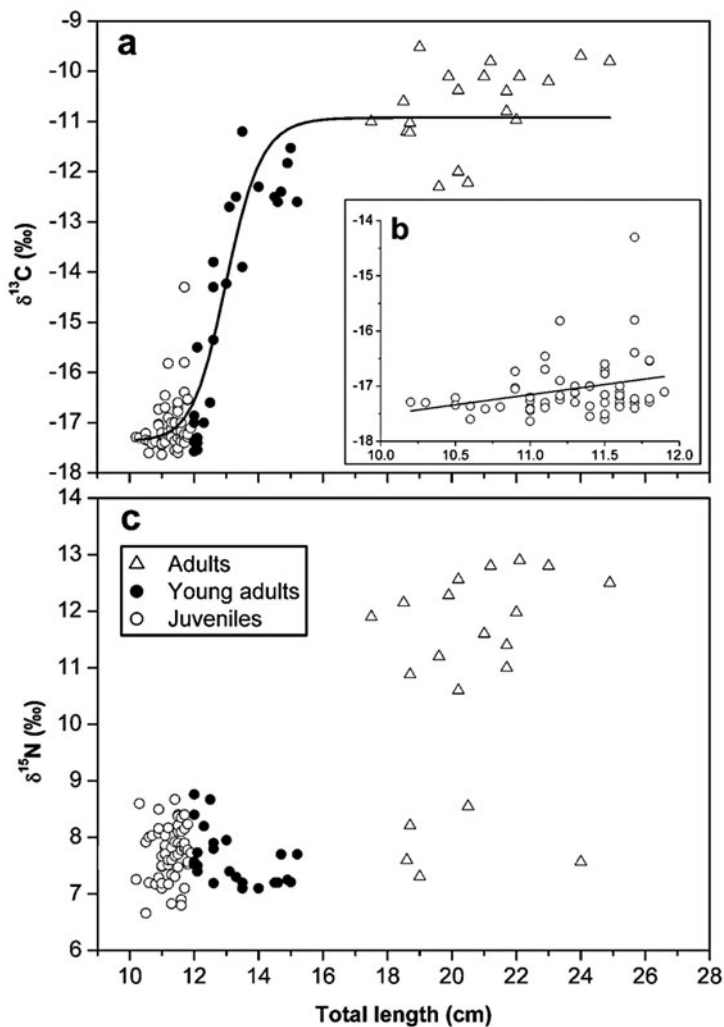


Fig. 3. *Mulloidichthys flavolineatus*. (a) $\delta^{13}\text{C}$ and (c) $\delta^{15}\text{N}$ values of *M. flavolineatus* as a function of total length (TL; n = 103). Symbols represent size groups: juveniles (n = 59), young adults (n = 26) and adults (n = 18). (a) A logistic model was fitted to $\delta^{13}\text{C}$ data (solid line; $\delta^{13}\text{C} = -6.32/[1 + (\text{TL}/12.98)^{26.27}] - 11.00$, $R^2 = 0.928$, $p < 0.001$). (b) A linear regression model was fitted to juvenile $\delta^{13}\text{C}$ values (solid line; $\delta^{13}\text{C} = 0.37\text{TL} - 21.20$; $R^2 = 0.064$, $p = 0.030$)

observed (Fig. 3b) for juveniles ($\delta^{13}\text{C}_{\text{juveniles}} = 0.37\text{TL} - 21.20$; $R^2 = 0.064$, $n = 59$, $p = 0.030$). The range of $\delta^{13}\text{C}$ values was greater for young adults (-17.6 to -11.2‰) than for juveniles or adults, and increased rapidly as a function of TL ($\delta^{13}\text{C}_{\text{young-adults}} = 1.40\text{TL} - 32.99$; $R^2 = 0.736$, $n = 26$, $p < 0.001$). $\delta^{13}\text{C}$ values of adults ranged from -12.3 to -9.5‰, but were not significantly correlated with TL ($\delta^{13}\text{C}_{\text{adults}} = 0.19\text{TL} - 14.71$; $R^2 = 0.079$, $n = 18$, $p = 0.128$). Hence, a logistic regression between $\delta^{13}\text{C}$ values and TL was modelled for *Mulloidichthys flavo-*

lineatus (Fig. 3a): $\delta^{13}\text{C}_{\text{yellowstripe goatfish}} = -6.32/[1 + (\text{TL}/12.98)^{26.27}] - 11.00$; $R^2 = 0.928$, $n = 103$, $p < 0.001$.

$\delta^{15}\text{N}$ isotopes

$\delta^{15}\text{N}$ values of juvenile fish ranged from 6.7 to 8.7‰, and were not correlated with TL ($\delta^{15}\text{N}_{\text{juveniles}} = 0.16\text{TL} + 5.94$; $R^2 = 0.002$, $n = 59$, $p = 0.299$). For young adults, $\delta^{15}\text{N}$ values ranged from 7.1 to 8.8‰, decreasing slightly with size ($\delta^{15}\text{N}_{\text{young-adults}} = -0.20\text{TL} + 10.21$; $R^2 = 0.181$, $n = 26$, $p = 0.019$) (Fig. 3c). $\delta^{15}\text{N}$ values of adults varied highly among individuals and ranged from 7.3 to 12.9‰. Two groups of individuals were identified (Fig. 3) with mean $\delta^{15}\text{N}$ values of 11.9 ± 0.8 and 7.8 ± 0.6 ‰ (Mann-Whitney: $Z = 3.27$, $p < 0.01$). $\delta^{15}\text{N}$ values of adults were not correlated with TL ($\delta^{15}\text{N}_{\text{adults}} = 0.23\text{TL} + 6.27$; $R^2 = -0.003$, $n = 18$, $p = 0.344$). There was a difference of one trophic level between adults (mean trophic level of 3.8 ± 0.7) and young adults and juveniles (2.5 ± 0.1 and 2.6 ± 0.2 , respectively; Kruskal-Wallis: $H = 30.44$, $p < 0.001$) (Table 3). As $\delta^{15}\text{N}$ values greatly varied within each group (up to 1 trophic level in adult group), data were not fitted using a model approach.

Food source contribution to diet

The average $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values of pelagic POM were -22.5 ± 0.5 and 4.4 ± 0.2 ‰, respectively. STOM, the proxy for micro-/meiobenthos food sources, had average $\delta^{13}\text{C}$ values of -16.0 ± 0.9 ‰ and $\delta^{15}\text{N}$ values of 3.3 ± 0.3 ‰. Macrofauna was the most enriched food source with mean $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values of -11.9 ± 1.3 and 7.5 ± 1.4 ‰, respectively (Fig. 4). Results of isotope mixing models are presented in Fig. 5. Contribution of pelagic food sources to overall diet decreased significantly (ANCOVA: $F = 84.3$, $p < 0.01$) from juveniles (mean of 47.7%) to young adults (17.0%), concurrent

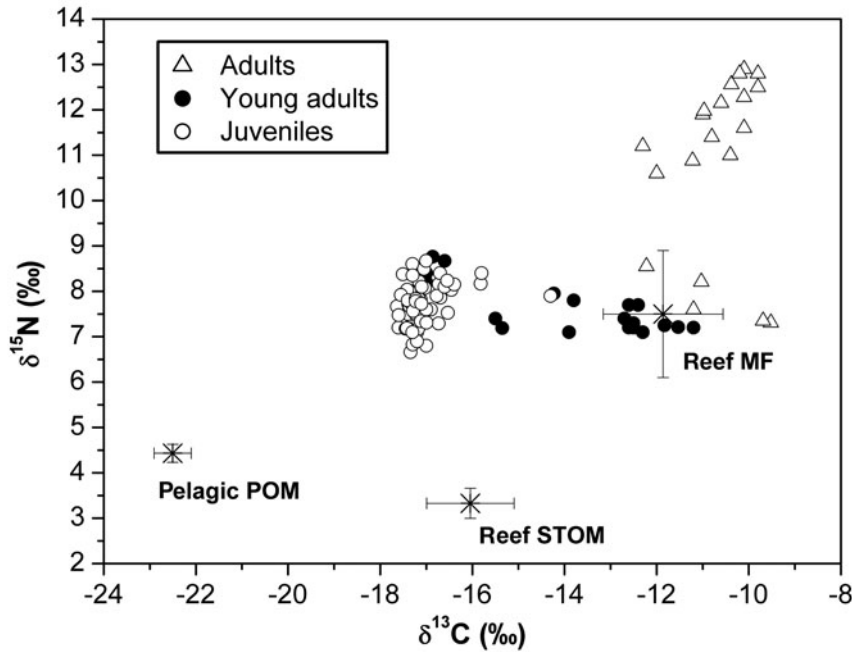


Fig. 4. *Mulloidichthys flavolineatus*. Mean (\pm SD) $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values of *M. flavolineatus* and potential food sources. Symbols represent size groups: juveniles (n = 59), young adults (n = 26) and adults (n = 18). $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ signatures of pelagic particulate organic matter (POM; n = 9), reef sediment total organic matter (STOM; n = 15) and reef macrofauna (MF; annelids, n = 31; arthropods, n = 24; molluscs, n = 27) are also shown

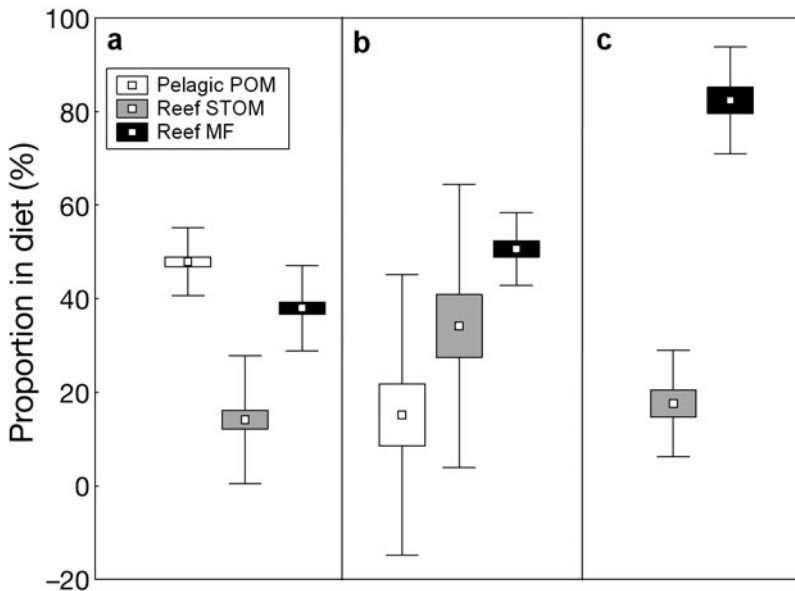


Fig. 5. Relative contribution (%) of pelagic (pelagic POM), micro-/meiobenthic (reef STOM) and macrobenthic (reef MF) food sources to (a) juvenile, (b) young adult and (c) adult *Mulloidichthys flavolineatus* diet estimated by 3- and 2-source isotopic mixing models. Boxes are means \pm SE, errorbars are \pm SD

to an increasing contribution of benthic sources. Micro-/meiobenthos food source contribution significantly increased from juveniles (mean of 14.6%) to young adults (32.3%), and decreased from young adults to adults (17.6%, $F = 24.0$, $p < 0.01$). Macrobenthic food source contribution increased as a function of fish size ($F = 16.2$, $p < 0.001$) from juveniles (37.7%) to young adults (50.6%) and adults (82.4%). Finally, young adults displayed a larger individual variability (see SD bars in Fig. 5) than adults and juveniles.

DISCUSSION

Stomach content analysis

Stomach content analysis reveals that the diet of *Mulloidichthys flavolineatus* from Reunion Island is predominantly composed of arthropods, nematodes and polychaetes. This result supports others findings for mullid fish (Ben-Eli-ahu & Golani 1990, Golani & Galil 1991, Wahbeh 1992, Platell et al. 1998, Lukoschek & McCormick 2001) and more generally reflects the diet of most coral reef benthic carnivorous fish (Kulbicki et al. 2005). Arthropods and nematodes dominated the yellowstripe goatfish diet in La Saline reef in terms of frequency of occurrence and numerical abundance, but constituted a low biomass proportion of the diet due to the small sizes of these prey types. The opposite pattern was observed for polychaetes, which constituted the most important part of stomach content volume. Their low abundance in stomachs contrasts with their high densities in La Saline reef flat sediments compared to other taxa (Taddei 2006). This suggests that *M. flavolineatus* selects specific polychaetes for their diet. Such selectivity has been already highlighted by Sorden (1982), who observed *M. flavolineatus* to select against small, abundant polychaete species in the Midway Islands (Hawaii).

Except for nematodes, our results differ from a study of *Mulloidichthys flavolineatus* diet in the Gulf of Aqaba

(Wahbeh 1992), where fish eggs, nematodes and molluscs dominated based on abundance, occurrence and gravimetric indices. In the present study, fish eggs were not observed and molluscs did not feature significantly in the feeding regime. However, specimens of *M. flavolineatus* from the Gulf of Aqaba were sampled in seagrass-rich areas, well known to host spawning aggregations (Moltschaniwskyj & Pecl 2003) and rich mollusc communities (Baron et al. 1993), while seagrass beds at La Saline reef are small and sparse. Considerable variation in diet composition has been reported between mullid species (Golani & Galil 1991, Platell et al. 1998), as well as within the same species from different locations (Lukoschek & McCormick 2001). These differences in prey selection may be related to differences in availability of prey taxa at these locations and seasonal variability.

The diet of juvenile yellowstripe goatfish was numerically dominated by nematodes and harpacticoid copepods. Several demersal juvenile fish species have been found to feed primarily on these 2 taxa, providing evidence of the crucial role of meiofauna in supporting juvenile growth (Coull 1999). Ontogenetic diet changes in yellowstripe goatfish were demonstrated from diet analyses across the 3 size groups investigated and correspond to a widening of the prey spectrum with increasing size. Differences in diet composition observed between juveniles and adults consisted of a decrease in meiofauna (nematodes, tanaids and harpacticoid copepods) and microfauna (foraminifera) prey, concurrent with an increase in macrofauna prey (gastropods, echinoderms, isopods and decapods) from a common prey pool (polychaetes, tanaids [Leptochelidae] and harpacticoid copepods). This is consistent with the conceptual model of a relationship between fish size and the size, biomass and diversity of prey (Scotto di Carlo et al. 1982, Harmelin-Vivien et al. 1989), limiting intraspecific competition for similarly sized prey (Huss et al. 2008). Intraspecific competition within fish cohorts, in particular following recruitment when juvenile density can affect resource density (Byström & Garcia-Berthou 1999), was found to have a large influence on the inter-individual variation in resource use (Svanbäck & Persson 2004). Size of the feeding apparatus, in particular the pharyngeal gape, has been associated with the size range of exploitable prey items (McCormick 1998). However, pharyngeal gape size was not the dominant factor in prey choice of *Parupeneus barberinus* (Lukoschek & McCormick 2001). Changes in diet composition associated with increasing size of *Mulloidichthys flavolineatus* from La Saline reef concur with an increase in sediment ingestion. Sediment was also found to be a significant component of stomach contents of the same species from the Red Sea (Wahbeh 1992). In the present study,

young adults and adults ingested sediment more frequently and in larger proportions than juveniles. These observations may be related to an ontogenetic change in foraging behaviour or change in microhabitat preference. Even though there is no data on foraging substrata used by *M. flavolineatus*, it is likely that juveniles concentrate their feeding efforts on prey associated with surface sediments or algal turfs, as stomachs contain a noticeable presence of fragments from algal mats, known to host a high abundance of invertebrates (Vetter 1994), while larger individuals forage deeper into the soft substrates. This has also been observed for some other mullid species (Eggold & Motta 1992, Lukoschek & McCormick 2001), associated with morphological modifications of specialized organs such as barbel development or elongation of the preorbital region of the head (Gosline 1984). The high values of the Morisita-Horn index, in particular between juveniles and young adults, attest to a high dietary overlap between successive size groups. This is due to the restricted group of prey they share which dominates the diet of each size group. The highest values of dietary overlap were observed for juvenile–young adult and young adult–adult pairs, indicating that young adults are a transient group. Such findings reflect a gradual expansion of the trophic niche of the *M. flavolineatus* with increasing size. However, stomach content analysis cannot resolve questions pertaining to the temporal dynamics of such dietary changes.

Stable isotope analyses

Interpreting intraspecific stable isotope variations with body size as evidence of ontogenetic diet changes requires underlying assumptions, which can result in ambiguous conclusions if ignored (Gannes et al. 1997). In particular, isotopic fractionation may be the result of both diet shifts as well as changes in tissue turnover rates during ontogeny. Tissue turnover rate is a function of growth and metabolism (Fry & Arnold 1982); fast-growing organisms have more rapid turnover rates and therefore more rapid isotope value changes (Hesslein et al. 1993). Despite the lack of information on fractionation rates in *Mulloidichthys flavolineatus*, we did not find any important changes in carbon isotopic composition according to size within the juvenile group, which experiences the greatest growth rate (Holland et al. 1993). Evidence from other teleosts (Fry 1983, Herzka & Holt 2000) indicate that metabolic processes have little or no effect on isotopic composition. This implies that patterns of isotopic changes are mostly the result of a dilution of the initial isotopic pool by addition of newly deposited biomass. Hence, major observed modifications in tissue carbon isotopic com-

position can be attributed to diet changes. The slight increase in $\delta^{13}\text{C}$ values with size observed in juveniles could be attributed to a group of individuals that have settled earlier in the shallow reef flat, thus reflecting the effects of a benthic signal integrated over a longer time period.

Carbon isotopes are often used to trace energy pathways within a system because pelagic carbon is depleted in ^{13}C relative to carbon derived from benthic-littoral sources (France 1995). Correlations between $\delta^{13}\text{C}$ values and *Mulloidichthys flavolineatus* TL clearly demonstrate important intraspecific $\delta^{13}\text{C}$ changes during ontogeny, consistent with other studies that have investigated ontogenetic diet shifts from juvenile to adult fish (Cocheret de la Morinière et al. 2003, Wells et al. 2008). In the La Saline reef, new recruits retained an offshore pelagic isotopic signal of -17.1‰ , while the carbon isotopic composition changed to a more enriched benthic signal (-10.7‰) in adults, indicating a transition from larval pelagic life to a benthic habitat. Tissue isotopic composition approaches the end member macrobenthic food signatures asymptotically over time, with a time lag of around 8 mo (minimum juvenile age at settlement to minimum adult age, see 'Data analysis'), corresponding to the dilution of the pelagic carbon signature. A limited number of studies have estimated isotopic turnover rates for fish, particularly in natural conditions. *M. flavolineatus* time to equilibrium is in the range found for various fish species (see Herzka 2005), and is close to the value determined for populations of the migratory goby *Rhinogobius* spp. (Maruyama et al. 2001). Most of the size-dependent changes in $\delta^{13}\text{C}$ occur at the young adult stage, for individuals with TL < 13 cm (breakpoint of the non-linear curve fit), that are expected to have a reef flat residence time of around 3.5 mo based on a von Bertalanffy growth curve (Holland et al. 1993). This provides evidence for relatively slow carbon turnover rate in *M. flavolineatus*, which is in agreement with the isotope mixing model where pelagic POM was still found to significantly contribute to the diet of young adults (mean pelagic POM diet contribution for juveniles and young adults of 47.7 and 17.0%, respectively). While isotopic changes occur rapidly from juveniles to young adults with TL < 13 cm, larger individuals (≥ 13 cm TL) took longer to reach the end member benthic carbon signature (5.5 mo). Such $\delta^{13}\text{C}$ patterns may be due to a decrease in *M. flavolineatus* growth rates with increasing size (Holland et al. 1993) or energy allocation for reproduction, as we found that 50% of young adults at 13 cm TL have already differentiated gonads. This was supported by the significant differences observed in $\delta^{13}\text{C}$ values between non-sexually differentiated (-15.2‰) and sexually differentiated (-13.8‰) young

adults. Even less is known about such physiological changes; this variation may be related to the equilibrium between assimilation and excretion, which is influenced by growth rate or metabolism (Olive et al. 2003). Physiological changes may also influence tissue $^{13}\text{C}/^{12}\text{C}$ isotopic ratios through variation in ^{13}C -depleted lipid content relative to other biochemical components (DeNiro & Epstein 1978, Pinnegar & Polunin 1999), such as those implied in reproductive organs. Such findings highlight that ontogenetic dietary changes related to settlement occur at different rates within the *M. flavolineatus* population at La Saline reef. These rates appear to be highly dependent on physiological factors, in particular for the medium-sized individuals.

Overall positive relationships between $\delta^{15}\text{N}$ values and size have been found for *Mulloidichthys flavolineatus*. The $\delta^{15}\text{N}$ values of this fish typically increased with body size and such patterns have been attributed almost exclusively to ontogenetic changes in diet (e.g. Vander Zanden & Rasmussen 2001, Badalamenti et al. 2002). Although some studies have pointed at a possible ^{15}N accumulation with age (Beaudoin et al. 1999, Overman & Parrish 2001), we did not find significant enrichment within each size group. There were no significant differences in $\delta^{15}\text{N}$ values between juveniles and young adults, and these groups thus had similar trophic level positions (mean trophic levels of 2.6 and 2.5, respectively), consistent with their high diet overlap (0.993 by number, 0.943 by volume) revealed by prey taxonomic analysis. The slight decrease in $\delta^{15}\text{N}$ values during the transition from juvenile to young adult stages reflects the shift from an offshore pelagic food source to an increasing reliance on benthic primary consumers (micro-/meiofauna), which generally have lower $\delta^{15}\text{N}$ values than their pelagic counterparts ($\delta^{15}\text{N}$ mean of 3.3 and 4.4‰ for micro-/meiofauna proxy and pelagic POM, respectively; Vander Zanden & Rasmussen 1999). Since a single trophic level is represented by a shift from 2 to 5‰ between predator and prey (Post 2002), the average shift of 3.4‰ in $\delta^{15}\text{N}$ values observed between the smallest size groups and adults (mean trophic level of 3.8) is about one trophic level and would represent the trophic status of the prey they consumed. While total prey number decreases by up to 50% between young adults and adults, the number of macrofauna prey items (mainly isopods, gastropods and ophiroids) increases to a factor of 5.3 (1.0 between juveniles and young adults). Isotopic mixing models reveal that macrofauna food sources contribute significantly to the tissue isotopic composition of each size group and attain a contribution of up to 82.4% in the adult's diet. This is consistent with the commonly observed trend that the trophic level of most aquatic species

increases with size (Jennings et al. 2002). These results suggest that *M. flavolineatus* gradually shifts to a macrofaunal diet that is totally equilibrated at the adult stage and reveal a major nutritional role of macrofauna in all post-settlement stages. This concurs with conclusions from stomach content analysis based on volume, but contradicts numerical analysis. While nematodes, tanaids and harpacticoid copepods (meiofauna) numerically dominated the diet, polychaetes (macrofauna) were dominant in volume, which was reflected in the fish isotopic signature. Polychaetes likely drive the juvenile yellowstripe goatfish isotopic signatures, since there were very few other macrofauna taxa present in their stomachs (only Naticidae, with low volumetric and numeric importance). In accordance with Sorden (1982), we suggest that *M. flavolineatus* preferentially select polychaetes. A possible preference could be based on the specific micronutrient fatty acids value necessary to support their growth (Watanabe et al. 1983).

Two distinct adult groups were identified based on their nitrogen stable isotope values (11.9 ± 0.8 and $7.8 \pm 0.6\text{‰}$). They displayed a range of sizes that excludes an age-related phenomenon, and we did not find any other discriminating factors (stomach contents, sex effect or time and place of capture) to explain such a difference. $\delta^{15}\text{N}$ isotope variability within the adult group could be explained by the migration of some individuals (individuals with $\delta^{15}\text{N}$ values of about 7.8‰) to outer reef soft-bottom areas, which have a more depleted $\delta^{15}\text{N}$ benthic food web baseline due to the increased distance from productive inner reef zones and anthropogenic runoffs (Heikoop et al. 2000). Holland et al. (1993) demonstrated that *Mulloidichthys flavolineatus* specimens presented a high degree of site fidelity, a relatively high feeding excursion distance range of 600 m across sand flats and very little post-recruitment movement of adults at Coconut Island reef, Hawaii. Hence, we hypothesize that fish in the present study with lower $\delta^{15}\text{N}$ values consisted of individuals primarily established in the outer reef zone that were caught during their daily feeding excursions in the shallow-water reef. In addition to ontogenetic dietary shifts, stable isotope analysis highlighted possible stage-specific partitioning in habitat use inside the La Saline reef that should be further investigated with tracking experiments.

CONCLUSIONS

The common use of prey numbers in stomach content analysis reveals strong limitations to the understanding of the feeding regime of *Mulloidichthys flavolineatus* and is at odds with conclusions from stable

isotope analysis in giving too much importance to meiofauna. Volumetric analysis of diet composition appears to be more adequate, highlighting a major role of polychaetes in the diet of *M. flavolineatus*. Stable isotope analyses revealed that *M. flavolineatus* shift from a pelagic to a macrobenthic diet, which is equilibrated at the adult stage. Combined $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ analysis is a relevant tool to depict dietary shifts occurring during ontogeny and migration onto the reef flats. We thus suggest that patterns of isotopic changes in response to dietary shifts of demersal reef fish can be adequately predicted based on size cohorts. We suggest that metabolism and sexual maturity can influence the rate at which isotopic equilibrium is reached following a shift to isotopically distinct food sources and should be taken into account in further trophodynamic studies. Such dietary shifts following habitat change reflects a variety of factors, including variation in taxonomic composition and trophic level of prey, foraging behaviour, pelagic–benthic differences in the dominant sources of primary production and isotopic composition of the inorganic nutrient pools. Stable isotope analysis appears to be a useful tool to investigate habitat and resource partitioning, in particular to describe patterns of transition from offshore pelagic to benthic habitats and to evaluate the contribution of benthic production that supports fish biomass.

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