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# Body size and stable isotope composition of zooplankton in the western tropical Atlantic

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#### Abstract:

Size-based approaches are paramount tools for the study of marine food webs. Here, we investigated the relationship between zooplankton body size, stable isotope composition and trophic level (TL) along a large-scale onshore-offshore gradient in the western tropical Atlantic. Samples were obtained on the Brazilian continental shelf, slope and in oceanic waters (off Fernando de Noronha archipelago and Rocas Atoll) in September and October 2015. Zooplankton was sieved into five size fractions. Zooplankton was dominated by copepods, except for the largest (>2000 µm) size fraction, that showed a high biovolume of chaetognaths, decapods, and fish larvae, Maximum zooplankton abundance and biovolume was found at the continental slope. POM showed consistently lower δ13C than zooplankton, indicating a selective use of 13C-rich primary food sources by zooplankton. Particulate organic matter (POM) was more 13Cenriched in shelf areas (average: -22.8, -23.6 and -24.3% at the shelf, slope and oceanic islands, respectively), probably due to the higher abundance of diatoms nearshore. POM had δ15N values between 2.5 and 6.9% (average: 4.0, 4.9 and 4.2% at the shelf, slope and oceanic islands, respectively). Zooplankton δ15N and TL increased with body size. The δ15N of the 200–500 μm size fraction was used as baseline for TL estimation. Oceanic areas (average baseline δ15N = 5.8% ± 0.52, n = 14) showed a higher baseline  $\delta 15N$  than the shelf (average = 3.9% ± 0.69, n = 9) and the slope areas (average =  $3.1\% \pm 0.93$ , n = 9). In spite of differing baselines, the  $\delta 15N$  data produced a consistent pattern of log-linear increase in TL with increasing size, in all areas. The choice of input trophic enrichment factor (TEF) values only slightly changed the log10 (body size) vs TL slopes, but this choice had a considerable effect on the estimates of predator/prey size ratio (PPSR) and predator/prey mass ratio (PPMR). Using a TEF above 2.3 leads to unrealistic PPSR and PPMR estimates. Overall average slope was 0.59 ± 0.08 TL  $\mu$ m-1 with TEF = 2.3 and 0.42 ± 0.07 TL  $\mu$ m-1 with TEF = 3.2.

#### Highlights

▶ Maximum zooplankton abundance and biovolume was found at the continental slope. ▶ POM showed lower  $\delta^{13}$ C than zooplankton, indicating selective feeding. ▶ Zooplankton  $\delta^{15}$ N and trophic level increased with body size. ▶ Using a  $\delta^{15}$ N trophic enrichment factor above 2.3 leads to unrealistic estimates.

Keywords: Plankton,  $\delta$ 13C,  $\delta$ 15N, Size classes, Trophic level, Predator / prey mass ratio

#### 1. Introduction

A large number of factors influence the functioning of pelagic ecosystems, including the spatio-temporal variability of physical and chemical variables, as well as dynamic interactions between species. These processes shape species distribution and trophic structure. Body size, which is easily and quickly determined, is considered one of the most important determinants of ecosystem structure (Jennings et al., 2007; Petchey et al., 2008) since metabolic requirements, food selection, growth and reproductive capacity are related to body size. Size-used analyses are therefore paramount tools for studying marine food webs (Jen.ings et al., 2001; Bănaru et al., 2014; Hunt et al., 2015; Yang et al., 2016). In pe'ag. marine food webs, predators are generally larger than their prey (Cohen et al. 1993) and trophic level (TL) is almost exclusively size-based. Thus, pelagic nar ne ecosystems can be well described by a loglinear size spectrum, where abundance continuously decreases with size, due to the energy loss in trophic transfers (1911.tt and Denman, 1977; Fry and Quinones, 1994; Jennings et al., 2002; Hunt et al., 2015). This theory was reinforced by Fry and Quinones (1994), who extinated zooplankton TLs with stable carbon and nitrogen isotopes, and deternined how TLs changed with body size. Their results supported the arguments that trophic organization and metabolism are important structuring forces. Analyzes of pelagic organisms aggregating them by size, may contribute to the understanding of the functioning of pelagic ecosystems.

Stable isotope analysis (SIA) is widely used to elucidate trophic relationships in marine food webs (Post, 2002; Hunt et al., 2015). Nitrogen and carbon stable isotopes ( $\delta^{15}$ N and  $\delta^{13}$ C) are natural tracers that incorporate information on trophic position and food source (Fry, 2006; Wang et al., 2013). For instance,  $\delta^{15}$ N increases with trophic

position (Post, 2002), i.e., predators are generally  $^{15}$ N-enriched in relation to their prey (Minagawa and Wada, 1984). Conversely, carbon stable isotopes ( $\delta^{13}$ C) are incorporated by organisms with less modification between predators and prey (McConnaughey and McRoy, 1979). Since predator  $\delta^{13}$ C values do not differ too much from their food sources (Miller et al., 2008), they can be used as tracers of food sources or feeding areas.

Lower trophic levels, such as most plankton organisms, are important for SIA in marine environments, since they are very sensitive to physical processes that result in changes in hydrographic patterns (Bode et al., 2007; Monthéan et al., 2013; Espinasse et al., 2014). Zooplankton plays a key role in pelagic ecosystems by connecting primary producers to higher trophic levels (Saiz et al., 2007; Guidi et al., 2016; Gove et al., 2016). Understanding the structure and forcion of zooplankton communities is necessary to understand their role for marine ecosystems (Yang et al., 2017). SIA provides a tool to analyze zooplankton food webs and to measure trophic relationships between food web components Cang et al., 2016). Combined with size-based approaches, SIA can provide comprehensive understanding of zooplankton food webs (Jennings et al., 2002; Bànara et al., 2014; Espinasse et al., 2014; Yang et al., 2016).

The western tropi al Atlantic off northeastern Brazil is mostly dominated by the warm oligotrophic waters that are derived from the south equatorial current (SEC). This region comprehends the extension from the shelf, slope and slope, seamounts and oceanic islands, all with low levels of nutrient supply and productivity and high zooplankton diversity (Boltovskoy et al., 1999). Numerous studies have investigated zooplankton communities in this region, most of which focusing on community structure, abundance diversity (e.g., Neumann-Leitão et al., 1999; Diaz et al., 2009; Brandão et al., 2013; Lira et al., 2014; Santana et al., 2018), and size spectra (Marcolin

et al., 2013). In this region, the stable isotope composition ( $\delta^{13}$ C and  $\delta^{15}$ N) of zooplankton and particulate organic matter (POM) has been investigated in mangrove estuaries, on the shelf and at the adjacent slope, revealing that any measurable estuarine influence was limited to the line of intertidal coastal reefs, approximately 10 km from the coast (Schwamborn et al., 1999, 2002).

In this study, we investigate the relationship between body size and stable isotope composition of zooplankton communities in the western tropical Atlantic, to understand the relationship between trophic levels and body size, and to test the hypothesis that there are differences in zooplankton trophic structure among different environments along a large-scale onshore-offshore gradient.

#### 2. Methodology

#### 2.1 Sampling strategy

Zooplankton and particulate organic matter (POM) were sampled on the continental shelf, at the slope, and in occanic waters off northeastern Brazil (Figure 1). Two oceanic island systems were sameled: Fernando de Noronha archipelago (FN) and Rocas Atoll (RA, Figure 1). Samples were collected between September 29 and October 21, 2015, during the Acoustic along the Brazilian coast cruise (ABRACOS, Bertrand, 2015) conducted on board R/V ANTEA.

At each station, vertical profiles of conductivity, temperature and chlorophyll a fluorescence were acquired using a Seabird SBE911+ CTD probe. Zooplankton sampling was conducted by towing a regular bongo frame with four nets fitted with mesh sizes of 64 μm (30 cm mouth opening diameter), 120 μm (30 cm diam.), 300 μm (60 cm diam.) and

 $500 \mu m$  (60 cm diam.). At each station, two oblique hauls were conducted between  $200 \mu m$  meters and the surface or between  $10 \mu m$  meters from the bottom to the surface at locations with less than  $200 \mu m$  of bottom depth. A total of  $14 \mu m$  and  $18 \mu m$  samples were collected during day night, respectively.

A Hydro-Bios flowmeters were fixed inside the net opening of each net to estimate filtered volume. Seawater samples for particulate organic matter (POM) were collected at the surface (3 m) and at the depth of maximum fluorescence using a CTD/rosette equipped with Niskin bottles. A total of 8 liters of water were filtered on a 47 mm diameter Whatman GF/F filter. In the laboratory, all filter sampler were dried at 40°C during 36h.

#### 2.2 Size spectra

Oblique bongo haul samples, taken sir unameously with 120 µm, 300 µm and 500 µm meshes, were used to estimate zoo<sub>F</sub>1.nkton size spectra. They were preserved in 4% formaldehyde buffered with sodium tetraborate (0.5 g.1<sup>-1</sup>, Newell and Newell, 1963). These formaldehyde-preserved samples were used for the size spectrum analysis under a stereo microscope (120 µm) and in the ZooScan system (300 µm, 500 µm). For stereo microscope analyses (Zaiss stemi 2000 – C), the samples were split in 1/2 to 1/256 to obtain at least 300 organisms per sample. The lengths and widths of 30 specimens for each taxonomic group were measured. If there were more than 30 individuals for each taxonomic group, they were counted for abundance analyses.

For ZooScan analyses, each zooplankton sample was separated into two fractions with a 1000  $\mu$ m mesh (Gorsky et al., 2010). Each size fraction was split by a Motoda splitter into 1/2 to 1/64 subsamples to obtain up to 2,000 objects in each scan. Large, rare organisms (> 1000  $\mu$ m size fraction) were scanned from 1/2 to 1/8 subsamples. For the more plentiful size fraction of small-sized organisms (< 1000  $\mu$ m) a smaller

subsample (1/4 to 1/64) was scanned. Subsamples were digitalized by the ZooScan system and processed with the ZooProcess software, which isolates each object into one vignette and generates a range of quantitative descriptors (size, grey level distribution and shape parameters) for each vignette. A semi-automatic approach was used to classify the vignettes into pre-established taxonomic groups, using the Plankton Identifier software (Gorsky et al., 2010). After classification, all results were manually validated to correct any misclassifications.

Equivalent spherical diameter (ESD), abundance and I iove lume were calculated for each organism. Zooplankton biovolume was estimated as the ellipsoidal volume:

$$Biovolume = \frac{4\pi}{3} \left( \frac{Major\ axis}{2} \right) \left( \frac{Minor\ axis}{2} \right)^2$$

where the major and minor axis of rac's zooplankton organism were measured by the ZooScan or under the stereo microscope. Zooplankton abundance and biovolume were classified into discrete size crasses, based on their equivalent spherical diameter (ESD, calculated from the 2D area). Five ESD size classes (0-IV) were defined (0 < 200  $\mu$ m; I 200 – 500  $\mu$ m; II 500 – 1000  $\mu$ m; III 1000 – 2000  $\mu$ m and IV >2000  $\mu$ m).

#### 2.3 Stable isotopes

Stable isotope analyses were conducted on particulate organic matter (POM) and on size-fractioned zooplankton samples that were obtained with a bongo net (additional subsurface hauls with 64  $\mu$ m, 120  $\mu$ m, 300  $\mu$ m and 500  $\mu$ m mesh). For each haul, all samples were pooled and sieved into five size fractions, using a multi-mesh array (0 < 200  $\mu$ m; I: 200  $\mu$ m – 500  $\mu$ m; II: 500  $\mu$ m – 1000  $\mu$ m; III: 1000  $\mu$ m – 2000  $\mu$ m and IV >

 $2000 \mu m$ ). Each size fraction sample was packed into previously calcined aluminum envelopes, and kept frozen at  $-20^{\circ}$ C.

In the laboratory, zooplankton samples were stored in Eppendorf micro tubes, and freeze dried during at least 24 hours. Once dried, each sample was homogenized to obtain a fine powder and weighed. In order to obtain unbiased values of δ<sup>13</sup>C, part of the samples was separated to remove the carbonates (CaCO<sub>3</sub>). These samples were acidified according to Fry (1988) by adding approximately 2 ml of C.5 mol.l<sup>-1</sup> hydrochloric acid (HCl). After this procedure, samples were re-dried at c.9°C for 24 hours and homogenized. Water samples collected at subsurface (~3 n) and at the depth of the chlorophyll a fluorescence maximum (Fmax) were used to obtain POM samples by using pre-combusted GFF filters. POM samples were acidified for extracting the carbonates. For this, the filters were exposed to hydrochloric acid (HCl) vapor. After 4 hours, the filters were left in an aluminum covered box and dried at 40°C during 36h.

Samples were analyzed using an elemental analyzer (Thermo Scientific Flash EA 2000) coupled to an Isotope Kerio Mass Spectrometer (Delta V+ mass) at the Pôle de Spectrométrie Océan (P!nuzané, France). Results were expressed in standard  $\delta$  notation based on international standards (Vienna Pee Dee Belemnite for  $\delta^{13}C$  and  $N_2$  in air for  $\delta^{15}N$ ) following the equation:  $\delta X = [(R_{sample}/R_{standard}) -1] \times 10^3$  (in %), where R is  $^{13}C/^{12}C$  or  $^{15}N/^{14}N$ . The standard values were reproduced into a confidence limit: IAEA-600 (certified values:  $-27.77 \pm 0.04\%$  for  $\delta^{13}C$  and  $1.00 \pm 0.20\%$  for  $\delta^{15}N$ ; measured value:  $-27.79 \pm 0.10\%$  for  $\delta^{13}C$  and  $1.09 \pm 0.06\%$  for  $\delta^{15}N$ ), IAEACH-6 (certified values:  $-10.45 \pm 0.08\%$  for  $\delta^{13}C$ ; measured value:  $-10.43 \pm 0.08\%$  for  $\delta^{13}C$ ), IAEA-N-1 (certified values:  $0.4 \pm 0.2\%$  for  $\delta^{15}N$ ; measured value:  $0.45 \pm 0.10\%$  for  $\delta^{15}N$ ) and IAEA-N-2 (certified values:  $20.3 \pm 0.2\%$  for  $\delta^{15}N$ ; measured value:  $20.24 \pm 0.12\%$  for

 $\delta^{15}$ N). One sample of a home standard (Thermo acetanilide) was analyzed for experimental precision, after every six samples.

#### 2.4 Trophic levels

Trophic level (TL) was calculated based on  $\delta^{15}N$  (Vander Zanden and Fetzer, 2007):

$$TL = \frac{(\delta^{15} N_{consumer} - \delta^{15} N_{baselin})}{TEF} \Gamma \lambda$$

where TEF is the trophic enrichment factor and λ is the trophic level of the baseline. Since isotopic values of phytoplanl to a (TL1) are based on POM, which may be influenced by the co-occurrence of detritus (Montoya et al., 2002) and microzooplankton in the water column (Post, 2002; Hunt et al., 2015; Yang et al., 2017), primary consumers were used to estimate the trophic level, once the they are less influenced by small-scale spatial and temporal variation (Hunt et. al. 2015). Small-sized zooplankton is classically used as a baseline (TL 2; e.g., Fry and Quinones, 1994; Kline and Pauly 1998, Montovo et al., 2002; Hauss et al., 2013, Hunt et al., 2015). In this study, we assume the size-class I (200 – 500 μm), to be closest to the primary consumer with TL 2. A TEF value of 3.2‰ per TL was applied to estimate relative TLs for each zooplankton size class and sampling area (Post, 2002; Ménard et al., 2014). For comparison, TLs were also estimated using a TEF of 2.3‰ per TL, as given by Schwamborn and Giarrizzo (2015).

Differences in trophic level, stable isotopes composition, abundance and biovolume of zooplankton between areas (shelf, slope and oceanic islands) and between size

classes were tested by non-parametric Kruskal-Wallis ANOVA ( $p_{crit} = 0.05$ ), since these data displayed non-normality and heteroscedasticity. Post-hoc comparisons between pairs of samples were conducted with Dunn's test ( $p_{crit} = 0.05$ ) (Zar, 1996).

Ordinary least squares linear regression was used to investigate the linear relationship between log<sub>10</sub>-transformed average body size (ESD, μm) and trophic level (TL, estimated with TEF = 3.2 and 2.3). Body size (ESD) was log10-transformed to obtain linear relationships for analysis and to improve hor escedasticity. The slope of this linear regression model was used to estimate the ave age predator/ prey size ratio (PPSR) and predator/ prey mass ratio (PPMR), using the fellowing equations: PPSR =  $10^{(1/\text{slope})}$ , if  $\log_{10}$  is being used in the linear movel (Hunt et al., 2015), and PPMR = PPSR<sup>3</sup>, assuming isometry and size-invariant density (Lins et al., 2019). PPSR and PPMR estimates obtained with TEF = 2 2 and 2.3 were then compared to previously published estimates. All available data at: https://figshare.com/articles/datas.zu?ody\_Size\_Stable\_Isotope\_Figueiredo\_et\_al\_Meta data\_csv/12620807.

#### 3. Results

#### 3.1 Hydrography

Sea surface temperature was nearly homogeneous (median: 26.6°C) over the whole study area. Surface salinity ranged from 36.1 to 36.6, with higher values along the continental slope. The thermocline ranged between ~80 and ~180 m in the slope area

and ~90 and ~130 m around oceanic islands (Assunção et al., in press). The chlorophyll *a* fluorescence maximum was generally located at the upper limit of the thermocline.

#### 3.2 Zooplankton abundance and biovolume size structure composition

A total of 15 taxonomic categories were identified: jellyfish, polychaetes, bivalves, gastropods, copepods, nauplii, euphausiids, mysids, decapods, salps, chaetognaths, appendicularians, fish eggs, fish larvae and "other zoop!ankton" (composed by organisms with less than 3% abundance).

No significant differences were detected between zooplankton abundance, biovolume, and stable isotope composition at both oceanic island areas, Fernando de Noronha and Rocas Atoll (K-W-ANOVA, n > 0.05). Therefore, data from these two areas were pooled together as "ocean o islands" for subsequent analyses. Samples obtained from the shelf and from the clope were analyzed separately, since they were different regarding several par interes, especially stable isotope composition (Fig. 5).

Abundance and biovol. The differed significantly between areas (K-W ANOVA, p < 0.05; Table 1). Samp'es 'aken off oceanic islands had significantly lower abundances (mean: 6.0 ind.m<sup>-3</sup>  $\pm$  3.1; Table 2) than those obtained from the shelf and from the slope (means: 10.3 ind.m<sup>-3</sup>  $\pm$  6.5 and 14.1 ind.m<sup>-3</sup>  $\pm$  16.8, respectively). The slope presented significantly higher biovolume (mean:  $11.0 \pm 15.4 \text{ mm}^3 \text{.m}^{-3}$ ) than oceanic islands (mean  $3.3 \pm 1.5 \text{ mm}^3 \text{.m}^{-3}$ ) and the shelf (mean:  $3.7 \pm 3.3 \text{ mm}^3 \text{.m}^{-3}$ , Figure 2).

Zooplankton composition varied considerably among size classes (Figure 3). Copepoda was the most abundant group for the three size fractions ranging between 200  $\mu$ m and 2000  $\mu$ m for all environments, and presented the largest contribution to biovolume. The taxonomic composition for the > 2000  $\mu$ m class showed a higher

contribution of larger zooplankton organisms, such as Chaetognatha, Decapoda, and fish larvae, especially off oceanic islands (Figure 3).

#### 3.3 Stable isotope composition of zooplankton and POM

Spatial patterns of stable isotope composition differed between POM and zooplankton. POM displayed consistently lower  $\delta^{13}$ C than zooplankton (Figure 4). Also,  $\delta^{13}$ C of POM showed a decreasing trend from the shelf towards offshore areas.  $\delta^{13}$ C of POM differed significantly between areas (Figure 5, Table 1, K-W ANOVA, p < 0.001), with higher  $\delta^{13}$ C values in shelf areas, followed by the slope and oceanic islands (Figure 5). Conversely, zooplankton displayed are apposite pattern, with increasing values towards offshore areas (Figure 5, Table 2).  $\delta^{15}$ N and  $\delta^{13}$ C values of all zooplankton size classes differed significantly between areas (Table 1, K-W ANOVA, p < 0.05), with higher values off oceanic islands (Figure 5; Table SM 1) varying from ~22 to ~19% for  $\delta^{13}$ C and from ~5 to ~6% for  $\delta^{15}$ N.

POM had  $\delta^{15}N$  values between 2.54 and 6.89‰ (mean at oceanic islands: 4.15‰, st. dev.: 1.2‰, mean at the shelf: 4.00‰, st. dev.: 0.59‰, mean at the slope: 4.86‰, st. dev.: 1.24‰). Within the meso- and macrozooplankton,  $\delta^{15}N$  generally increased with size, from the size fraction < 200 to the size fraction > 2000  $\mu$ m (Figure 4). At the shelf, small-sized mesozooplankton (size fraction 200 – 500  $\mu$ m) presented significantly more depleted values than the largest size (Table 2, 1000 – > 2000  $\mu$ m, p < 0.05, K-W ANOVA). At the slope, the 200 – 500  $\mu$ m size fraction showed significantly lower  $\delta^{15}N$  values than the largest (> 2000 $\mu$ m) size fraction (Table 2, p < 0.05, K-W ANOVA).

 $\delta^{13}$ C values were significantly different between > 2000 µm and 500 – 1000 µm at the shelf (Figure 5, p < 0.01, non-parametric ANOVA) and between > 2000 µm and the other size fractions at oceanic islands (Table 2, p < 0.05, non-parametric ANOVA).

#### 3.4 Trophic levels

To estimate trophic levels (TL), according to other studies (e.g., Fry and Quinones, 1994; Kline and Pauly, 1998, Montoya et al., 2002; Haus, et al., 2013, Hunt el al., 2015) we used the mean  $\delta^{15}N$  of the zooplankton sample, in the 200 – 500 µm size class, from each region, as a baseline (TL = 2), since this size fraction was consistently dominated by copepods, and the fact that this  $\delta^{12}$  class had the lowest  $\delta^{15}N$  values. Oceanic areas (average baseline  $\delta^{15}N = 5.8\%$ ) in a la higher baseline  $\delta^{15}N$  than the shelf (average = 3.9%) and the slope (average = 5.1%). In spite of differing baselines, the  $\delta^{15}N$  data produced a very consistent pattern of increase in trophic level with increasing size, in all regions (Figure 6). Assuming TEF = 3.2% TL<sup>-1</sup>, the mean trophic level of the zooplankton was very similar with TL = 2.2, 2.3 and 2.1 for shelf, slope and oceanic islands, respectively. For TEF = 2.3% TL<sup>-1</sup>, the mean were TL = 2.3, 2.5 and 2.2, in these sampling are,  $\delta^{12}N = 2.3\%$  TL<sup>-1</sup>, the mean were TL = 2.3, 2.5 and 2.2, in

Linear regression analysis revealed a log-linear relationship between  $\log_{10}$  (body size and TL (Figure 6). Linear models were always highly significant (p < 0.001), for shelf, slope and oceanic islands areas. Slopes of the log (body size) vs TL relationships (Table 3) were not significantly different among areas (ANCOVA, p = 0.07). Overall average slope was  $0.59 \pm 0.08$  TL  $\mu m^{-1}$  with TEF = 2.3 and  $0.42 \pm 0.59$  TL  $\mu m^{-1}$  with TEF = 3.2.

The choice of input TEF values only slightly changed the log<sub>10</sub> (body size) *vs* TL slopes, but this choice had a considerable effect on the estimates of predator/prey size ratio (PPSR) and predator/prey mass ratio (PPMR). For example, the estimate of PPMR at Oceanic Islands was 470 times lower when using TEF = 2.3 than when using TEF = 3.2. For TEF = 2.3, PPSR was 49, and PPMR was 121,547. For TEF = 3.2, these estimated were much higher, especially for PPMR. With TEF = 3.2, PPSR was 240, and an extremely high PPMR estimate of 13,894,955.

#### 4. Discussion

The present study revealed important variations in stable isotope ratios of zooplankton in the western tropical Ada tic that were conspicuously and significantly related to body size and geographical areas. Size-structured stable isotope analyses proved to be a useful approach to describe the structure and functioning of the systems (Fry and Quiñones, 1994; Montova et al., 2002; Lee et al., 2013; Hunt et al., 2015). It is the first study analyzing size classes and stable isotope composition of a zooplankton community carried out in tropical shelf, slope and oceanic waters, providing several important new insights into these pelagic ecosystems. Also, our study showed that TEF choice has a profound effect on the resulting prey-predator mass and size ratios. Lower TEF values than those widely used, clearly lead to more realistic results for this zooplankton community.

#### 4.1 Variation of POM and zooplankton stable isotope composition

In the present study, zooplankton presented higher  $\delta^{13}$ C and  $\delta^{15}$ N values off oceanic islands than in shelf and slope areas. Conversely,  $\delta^{13}$ C POM showed a clear coastal-offshore gradient, with  $^{13}$ C-enriched values at shelf areas, followed by slope and oceanic islands. Stable isotope composition of primary producers typically varies with spatio-temporal and physical features as a result of differences in biogeochemical processes that occur in each environment (Graham et al., 2010). Thus,  $\delta^{13}$ C values of POM followed an expected pattern, being more  $^{13}$ C-enriched in diatom-rich shelf environments than offshore, as observed in previous crucies in the study area (Schwamborn et al., 1997; Schwamborn et al., 1999) and cisewhere (Fry and Wainright, 1991).

An unexpected result was observed in relation to zooplankton  $\delta^{13}$ C values, which were conspicuously and significantly different from simultaneously obtained POM  $\delta^{13}$ C values. This discrepancy between PON, and zooplankton carbon isotope composition could be explained by two different phenomena. First, vertically migrating zooplankton may be feeding at depths that are not represented by the sampling of POM. Second, zooplankton generally display feeding selectivity (Sailley et al., 2015; Benedetti et al., 2016), which makes then isotope composition difficult to compare directly with POM values. POM encompa ses a highly variable mixture of a diversity of living organisms and non-living particles, among which only few may be selectively ingested by zooplankton (Lee et al., 2004). Our results indicate a selective use of  $\delta^{13}$ C-rich food sources (e.g., diatoms) by zooplankton.

4.2 Spatial patterns of zooplankton abundance, biovolume, composition, size and stable isotope ratios

We observed higher zooplankton abundance and biovolume over the slope than the shelf and in offshore areas (Figure 2). These results contradict the typical coastaloceanic gradient, previously found in the study area (Bueno et al., 2017; Campelo et al., 2018; Santana et al., 2018) and elsewhere (Dai et al., 2016; Giering et al., 2018). Indeed it is typically assumed that zooplankton abundance follows a continuous gradient with higher abundance nearshore (Neumann-Leitão et al., 2008; Marcolin et al., 2013; Leitão et al., 2019). This common pattern was mostly attributed to continental runoff, specifically the input of nutrients and organisms from large escaries, and resuspension from shelf sediments (Schwamborn et al., 1999). Numert inputs boost the primary production and may consequently generate a coastal ocean gradient of zooplankton density with higher abundance of organisms in coasal environments than oligotrophic oceanic waters. The lower biomass values overved at the shelf than at the slope may occur because of a series of factors. First, during the sampling period (September to October), there was a low continental influence (low river runoff during the dry season), low wind intensities (little resuspersion from shelf sediments). Also, sampling was conducted off the main rep. ductive season for most coastal invertebrates (low larval inputs from coastal ecosystaris), which is generally from January to March, in the study area. In addition, processes like eddies may allow a higher retention of plankton at the continental slope, leading to higher zooplankton biomass and densities (Franco et al., 2006; Katsuragawa et al., 2014). Our stations with highest abundance were located in a region close to continental slope, with mixed layer depth and upper thermocline shallower than expected for the season (Assunção et al., 2020). This leads to a shallowing of the nutricline and likely an increase in primary productivity sustaining the higher density of zooplankton.

In this oligotrophic ecosystem we demonstrated the existence of zooplankton accumulation at the continental slope. The observed pattern also agrees with the finding that any direct estuarine influence in the study area is generally limited to approximately 10 km from the coast (Schwamborn et al., 1999, 2002). The shelf break is known to play an important role in transporting and retaining zooplankton (Genin, 2004; Zhu et al., 2009). Indeed, interactions between topography and currents aggregate zooplankton seaward of the shelf-break zone (e.g. Genin, 2004; Cotté and Simard, 2005; Swartzman et al., 2005). Also, the whole slope area, up to several up off the shelf break, is characterized by strong turbulence and current shear between the base of the mixed layer and the upper thermocline that can transport nutrients from deep water masses upwards into the euphotic layer. This shear and turbuance is caused by the strong North Brazil Undercurrent (NBUC) that flows north words along the shelf break (Stramma et al., 1995; Schott, et al., 2005). Firth rmore, significantly higher abundance and biovolume at the slope, as observed in this study, may be due to small-scale mechanisms that were hitherto ignored, e.g., upwelling at submarine canyons (Kämpf, 2007; Howattand and Allen, 2013), and zooplankton accumulations at fronts and eddies (Schwamborn et al., 2001, Maps et al., 2015).

Zooplankton composition differed drastically between size-classes. Early life stages, such as invertebrate (probably mostly copepod) eggs, copepod nauplii, and gastropods dominated the smallest size class (< 200 μm), as found in many other regions, such as in the Mediterranean Sea (Bănaru et al., 2014). Similarly, in coastal areas of the Brazilian Northeast, Neumann Leitão et al. (2019) observed that nauplii and veliger larvae were the most frequent taxonomic groups in the microzooplankton (64 – 120 μm mesh net samples). Mesh size effects have already been documented in numerous zooplankton community studies (Tseng et al., 2011; Tosetto et al., 2019),

which demonstrate that mesh size drastically affects the representation of the abundance, composition, and diversity of zooplankton communities.

Zooplankton composition in the size fraction from 500 to 2000 µm is generally dominated by copepods (Mauchline, 1998), as observed by Neumann–Leitão et al. (2019) and Campelo et al. (2018) in coastal and oceanic waters of the western tropical Atlantic as well as in numerous other regions, such as in the south Atlantic (Boltovskoy, 1999), the Mediterranean (Bănaru et al., 2014; Espinasse et al., 2014), the subtropical north Pacific (Dai et al., 2016), the western tropical north Facin c (Yang et al., 2016) and the eastern Atlantic (Marcolin et al., 2013).

Interestingly, in most cases,  $\delta^{13}$ C values  $\alpha_1^{-1}$  tot vary significantly with size, indicating that, in contrast to other studies (e.g., Cchwamborn and Giarrizzo, 2015), the primary carbon sources of zooplankten 11 these size classes were similar, and carbon isotopic fractionation was negligib!•.

Although most zooplankton size fractions were dominated by copepods, the largest size fraction (> 2000  $\mu$ m) showed the greatest richness in taxonomic groups, with many large-sized organisms cuci. as fish larvae, euphausiids, mysids and many gelatinous organisms, such as solps and chaetognaths. The higher abundance and biovolume of large-sized predators (e.g., fish larvae and chaetognaths) in the largest size class explains well why  $\delta^{15}N$  and TL increased with size, showing a good consistency within and between our data sets.

Taxonomic compositions of both data sets (measured size classes and taxonomy *vs* sieving and isotope analysis) were most likely very similar, since they were obtained concomitantly at each station. Also, the measured size has an obvious relation to the retention in sieves. Furthermore, the observed distributions of taxonomic groups within

size groups were as expected (e.g., copepods being dominant in the smaller size fractions). In addition, with the approach we used, the taxonomic composition does not need to be absolutely identical in both datasets (i.e., only the size distribution has to be similar). Taxonomic information is presented here for illustration and to aid in interpreting the stable isotope results. Most importantly, the increase in  $\delta^{15}N$  with size in this study was expected, since it agrees well with previous studies on zooplankton food webs in the southwestern subtropical Pacific (Hunt et al., 2015) and in the western tropical north Pacific (Yang et al., 2016).

# 4.3 Body size as determinant of trophic level

In previous trophic ecology studies, the size classes chosen to determine the baseline differed widely between authors, e.g., 1°.5 – 250 μm (Fry and Quinones, 1994), 200 – 500 μm (Hauss et al., 2013) or 250 – 500 μm (Montoya et al., 2002). Assuming a size-based food web structure, in our study, we choose the 200 – 500 μm size fraction, once this size class was composed mainly by copepods, assumed to be mostly filter feeders and presented the smallest S1′N values. Thus, their isotopic composition should be close to primary consumer. (1°L2). In addition, copepods have better integrative properties than POM, since their turnover rate is much lower than most POM.

A positive relationship between  $\delta^{15}N$  and size class within the zooplankton community was also observed in the Mediterranean Sea by Koppelman et al (2009) and Bănaru et al. (2014), except for the largest size class (> 2000  $\mu$ m), which had lower  $\delta^{15}N$  values, probably due to the contribution of low-TL salps, in contrast to the dominance of high-TL chaetognaths in the present study. Their  $\delta^{15}N$  data for size-fractioned zooplankton were generally very similar to the present study except for the

largest zooplankton size class. A series of factors affect the trophic position and size of plankton. For example, variations in nutrient inputs regulate the size of primary producers (Kiørboe, 2008), thus strongly impacting species composition and size structure of herbivorous zooplankton and pelagic food webs.

#### 4.4 TEF choice and predator – prey size ratios

The TEF is known to differ widely between groups of caralisms (McCutchan et al., 2003; Vanderklift and Ponsard, 2003). Schwamborn an Garrizzo (2015) found a lower ecosystem-wide TEF value and much lower TEFs for higher TLs, which may support criticism of the use of fixed fractionation factors Hear, we compared results for a TEF of 3.2 (Post, 2002; Ménard et al., 2014) and 2.2 (Schwamborn and Giarrizzo, 2015). A lower TEF in ecosystem-based studies such as in Schwamborn and Giarrizzo (2015), than in laboratory feeding studies (Post, 2002), is to be expected since growing and migrating animals in real ecosystems will be constantly shifting their diets and TLs. Thus, most organisms in narral ecosystems will not attain full equilibrium with their current food source. In his st laboratory feeding experiments, only the final TEF (difference in isoto, e composition between diet and consumer), under perfect equilibrium, is reported. TEFs that occur in natural ecosystems will not to represent this kind of situation. Thus, lower, ecosystem-based TEF values (e.g., 2.3% TL<sup>-1</sup>) may be more realistic for the use in zooplankton food web analyses (Schwamborn and Giarrizzo, 2015) than laboratory-derived TEF estimates. Our study showed that TEF choice has a huge influence on PPMR and that the use of TEF above 3 will lead to a gross overestimation of PPMR and subsequently underestimation of ecosystem trophic efficiency (TE). Hunt et al. (2015) obtained similar results for the meso-,

macrozooplankton and micronekton communities in the subtropical Pacific. They also concluded that a TEF of 3.4‰ TL<sup>-1</sup>, that they used, likely overestimated PPMR and underestimated TE.

#### 4.5 PPMR as a key parameter for food webs and size spectra analyses

Our PPMR calculations assume isometry (i.e., size-invariant shape) and size-invariant density, which are hardly fulfilled in nature. In the sanse, Lins et al. (2019) showed that on average, density is not significantly different from 1 in tropical zooplankton samples, and that these relationships do not change with biomass. We are therefore confident that our basic assumptions are sufficiently well fulfilled for an assessment of PPMR. However we acknowledge that the investigation of these important aspects deserves further effects.

The large review by Hansen et al. (1994) reported typical PPSR values of 18 to 50 for mesozooplankton. In our gudy, extremely high PPSR and PPMR estimates obtained with TEF = 3.2 (Table 4), way be considered unrealistic, since they were much higher than previously publi hea estimates (except for the slope area, where the lowest PPSR was observed). Conver ely, the application of TEF = 2.3 produced realistic estimates for shelf and slope data, and for the overall mean. The zooplankton communities around oceanic islands had very high PPSR, being above the Hansen et al. (1994) range, with both TEF values (TEF = 2.3 and 3.2). This high PPSR is consistent with our observation of very abundant large organisms, such as fish larvae and gelatinous predators, in the waters around oceanic islands.

When using TEF = 3.2, we would obtain an overall mean PPSR value of 240, which is far above any known estimates for mesozooplankton. Yet, when using TEF =

2.3, our study indicates an overall mean PPSR value of 49, which is within the PPSR range given by Hansen et al. (1994). Similarly to the study of Hansen et al. (1994), we also used ESD to obtain PPSR estimates, so that both PPSR assessments seem well comparable, thus further strengthening the point for ESD as a standard measure of size in plankton research.

Community-wide PPSR above 1,000 should be considered extremely unlikely, such as those obtained with TEF = 3.2. This suggests that a realistic TEF for the zooplankton community sampled in this study would rather be close to 2.3, as observed by Schwamborn and Giarrizzo (2015). Since there are still ve y few studies available that attempt to estimate PPMR and PPSR based or suble isotopes and size-structured zooplankton sampling, our results may be used as a baseline for future studies.

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#### Supplementary material

**Table MS1**. Zooplankton size classes stable isotope composition between the three areas  $\delta^{13}C$  and  $\delta^{15}N$  minimum, maximum mean and standard deviation values.

Cina Classes	$\delta^{13}$ C	$\delta^{13}$ C	$\delta^{13}$ C	$\delta^{13}$ C	S <sup>15</sup> N	$\delta^{15}N$	$\delta^{15}N$	$\delta^{15}N$
Size Classes	Min	Max	Mean	sd	Niii	Max	Mean	sd
	Coast							
< 200 μm	-23.23	-20.92	-22.08	1.63	3.17	4.20	3.69	0.73
$200-500\;\mu m$	-21.49	-20.20	-21.08	0.38	2.87	5.09	3.91	0.69
$500-1000~\mu m$	-21.39	-20.32	-20.72	0.32	3.88	6.12	4.59	0.79
$1000-2000~\mu m$	-21.87	-20.43	-21.07	0.57	4.43	6.42	5.05	0.63
>2000 µm	-22.17	-20.79	-21 +6	0.43	4.51	6.17	5.08	0.52
	Slope							
< 200 μm	-21.79	-21.71	-21 74	0.05	4.40	5.39	4.92	0.50
$200-500~\mu m$	-22.39	-20.48	-21.46	0.63	1.07	4.03	3.06	0.94
$500-1000~\mu m$	-21.76	-20.05	-21.14	0.48	2.24	5.25	4.03	0.92
$1000-2000~\mu m$	-23.21	-20.44	-21.58	0.86	2.54	5.26	4.13	0.79
>2000 µm	-22.76	-20 22	-21.75	0.66	3.28	6.92	5.04	1.09
	Oceanic Islands							
< 20 0µm	-21.95	-20 23	-20.94	0.55	5.10	7.52	5.99	0.60
$200-500\;\mu m$	-21.02	-20.29	-20.64	0.26	5.07	6.84	5.79	0.52
$500-1000~\mu m$	-21.75	19.96	-20.72	0.43	5.54	6.91	6.00	0.38
$1000-2000~\mu m$	-21.22	-20.24	-20.62	0.26	5.88	8.25	6.52	0.62
>2000 µm	-20.65	-18.69	-20.02	0.50	5.88	8.89	6.67	0.68

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#### Ingures and tables legends

**Figure 1.**Sampling tation in the tropical Atlantic off northeastern Brazil, in September and October 2015. Color codes indicate sampling stations on the continental shelf, at the slope, and off two oceanic islands (Fig. Fernando de Noronha Archipelago and RA: Rocas Atoll).

**Figure 2.** Mean (± standard deviations) zooplankton abundance and biovolume between continental shelf, slope and oceanic islands in the tropical Atlantic off northeastern Brazil. All values shown represent the sum of all size classes

**Figure 3.**Relative abundance and biovolume of zooplankton classified within four size classes (0-IV). I:  $200-500~\mu m$ ; II:  $500-1000~\mu m$ ; III:  $1000-2000~\mu m$  and IV:  $>2000~\mu m$ . Samples were collected on the continental shelf, at the slope and off oceanic islands in the tropical Atlantic off northeastern Brazil.

- **Figure 4.** Mean  $\delta^{13}$ C and  $\delta^{15}$ N ( $\pm$  standard deviation) of particulate organic matter (POM) and size-fractionated zooplankton sampled on the continental shelf, at the slope and off oceanic islands in the tropical Atlantic off northeastern Brazil.
- **Figure** 5:  $\delta^{13}$ C and  $\delta^{15}$ N (in ‰) of particulate organic matter (POM) and zooplankton size fractions (0, < 200 µm; I: 200 500 µm; II: 500 1000 µm; III: 1000 2000 µm; IV > 2000 µm) on the shelf, at the slope and off oceanic islands in the western tropical Atlantic.
- **Figure 6**: Trophic level (TL) and  $\log_{10}$  (mean ESD size,  $\mu$ m) for each zooplankton size class in the western tropical Atlantic. TL was calculated from  $\delta^{15}N$ , assuming a trophic enrichment factor of 3.2 and 2.3% TL<sup>-1</sup>. Grey area: 95% confidence enrelege for the linear regression slope.
- **Table 1.** Results of Kruskal-Wallis ANOVA (p-values) and Dunn's post-hoc test on Particulate Organic Matter (POM), zooplankton  $\delta^{13}$ C,  $\delta^{15}$ N, abundance and biovolume according to local (S: Shelf; SB: Slope and OI: Oceanic Islands) and the classes (0: < 200  $\mu$ m; II: 200 500  $\mu$ m; II: 500 1000  $\mu$ m; III: 1000 2000  $\mu$ m, IV > 200  $^{\circ}$  unit n.s. not significant (i.e., p > 0.05).
- **Table 2.** Results of Kruskal -Wallis ANOVA and Dunns's post-hoc test with zooplankton  $\delta^{13}$ C and  $\delta^{15}$ N, classified into five size classes ( $2 < 200 \mu m$ ; I:  $200 500 \mu m$ ; II:  $500 1000 \mu m$ ; III:  $1000 2000 \mu m$ ; IV  $> 2000 \mu m$ ) for each area: Continental Shelf, Slope and Oceanic islands.
- **Table 3**. Slope values of cophic level (TL) vs body size (ESD,  $\mu$ m), testing two different trophic enrichment factors (TEF = 3.2 and 2.3), for each area ( $\pm$  standard errors).
- **Table 4.** Predator/prey mass ratio (PPMR) and predator/prey size ratio (PPSR) testing two different trophic enrichment factors (TEF = 3.2 and 2.3), for each area.

#### Supplementary material

**Table MS1**. Zooplankton size classes stable isotope composition between the three areas  $\delta^{13}$ C and  $\delta^{15}$ N minimum, maximum mean and standard deviation values.

#### **Tables**

Table 1.

Response variable	Factor	P	Post hoc
δ <sup>13</sup> C Zooplankton	Area	< 0.001	OI > S > SB
o e zoopiankion	Size class	n.s.	n.s
$\delta^{15}$ N Zooplankton	Area	< 0.001	OI > S = SB
o in Zoopiankion	Size class	< 0.001	I > 0 = III = IV; I > IV
213 a p 2 2 5		0.004	
δ <sup>13</sup> C POM	Area	< 0.001	3 > 9B > 01
$\delta^{15}$ N POM	Area	n.s.	n.s
Abundance	Area	< 0.01	SB = S > OI
Biovolume	Area	<05	S = SB; S = OI; SB > OI
Abundance	Area Area	< 0.001  n.s. < 0.01	S > SB > OI  n.s $SB = S > OI$

**Table 2.** Results of Kruskal -Wallis  $\Delta NO^{-7}A$  and Dunns's post-hoc test with zooplankton  $\delta^{13}C$  and  $\delta^{15}N$ , classified into five size classes (0, < 200  $\mu$ m; I: 200 – 500  $\mu$ m; II: 500 – 1000  $\mu$ m; III:  $1000 - 2000 \mu$ m; IV > 2000  $\mu$ m; for each area: Continental Shelf, Slope and Oceanic islands.

Re sponse variable	<i>P</i> -value	Post hoc				
Shelf						
δ <sup>13</sup> C	< 0.01	$II \neq IV; II \neq 0$				
$\delta^{15}N$	< 0.01	$I \neq III = IV$				
Slope						
$\delta^{13}$ C	>0.05	n.s				
$\delta^{15}N$	< 0.001	$I \neq 0$ ; $IV = II = III > I$				
Oceanic islands						
$\delta^{13}$ C	< 0.001	IV>0=I=II=III				
$\delta^{15}$ N	>0.05	n.s				

**Table 3**. Slope values of trophic level (TL) vs body size (ESD,  $\mu$ m), testing two different trophic enrichment factors (TEF = 3.2 and 2.3), for each area ( $\pm$  standard errors).

	Slope (TL log <sub>10</sub> (μm) <sup>-1</sup> )			
	TEF 2.3	TEF 3.2		
Shelf	$0.58 \pm 0.14$	$0.42 \pm 0.10$		
Slope	$0.85 \pm 0.18$	0.61 ± 0.1°		
Oceanic Islands	$0.44 \pm 0.08$	0.31 ±0.00		
All	$0.59 \pm 0.08$	0.42 ±0.59		

**Table 4.** Predator/prey mass ratio (PPMR) and p. eda or/prey size ratio (PPSR) testing two different trophic enrichment factors (TEF = 3.2 and 2.3), for each area.

	TE. 2 3		TEF 3.2		
	PPMR	PSR	PPMR	PPSR	
Shelf	142.773	52	13,625,858	293	
Slope	3 27 6	14	75,932	42	
Oceanic Islands	(,41 <sup>7</sup> .294	185	3,021,138,554	1,445	
All	121,547	49	13,894,955	240	

# **Highlights**

- Maximum zooplankton abundance and biovolume was found at the continental slope.
- POM showed lower  $\delta^{13}$ C than zooplankton, indicating selective feeding.
- Zooplankton  $\delta^{15}$ N and trophic level increased w. 1 body size.
- Using a  $\delta^{15}N$  trophic enrichment factor above 2.3 leads to unrealistic estimates.

Figure 1

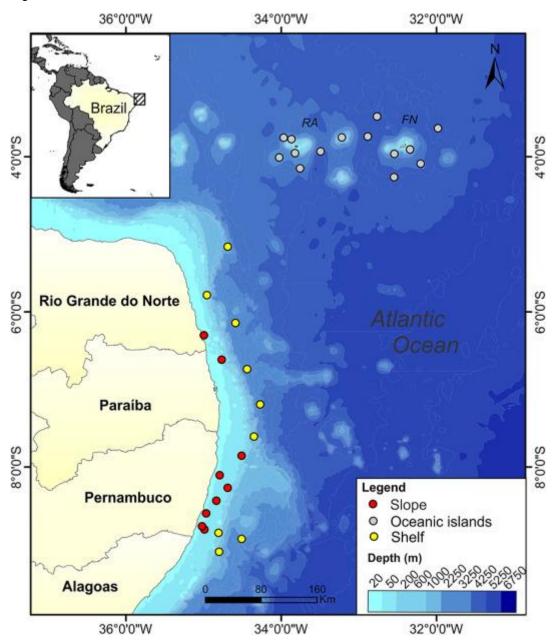


Figure 2

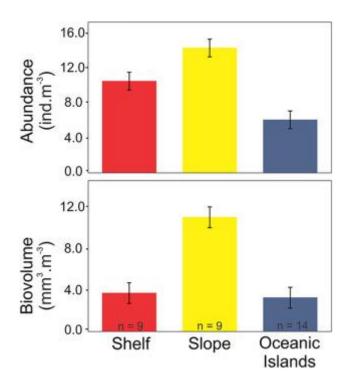


Figure 3

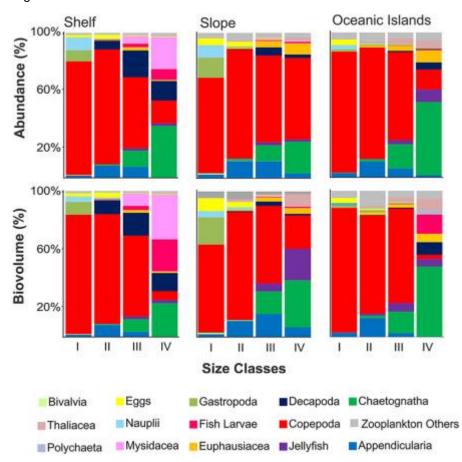


Figure 4

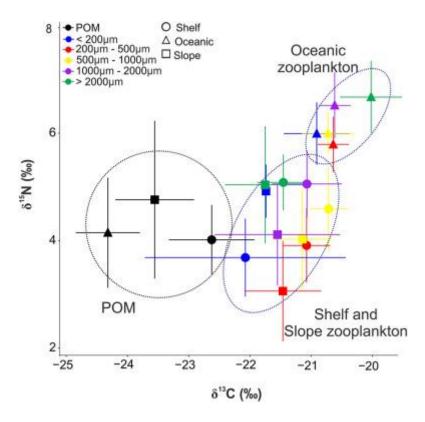


Figure 5

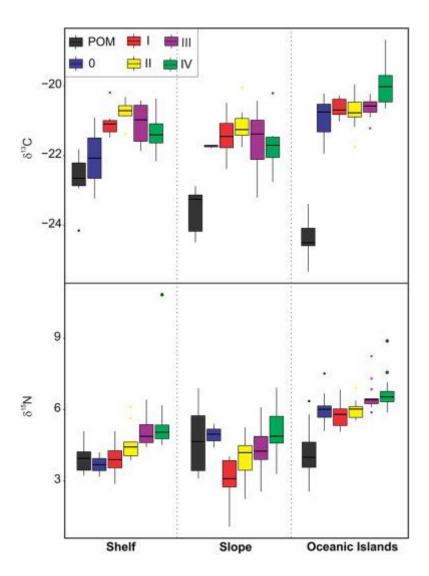
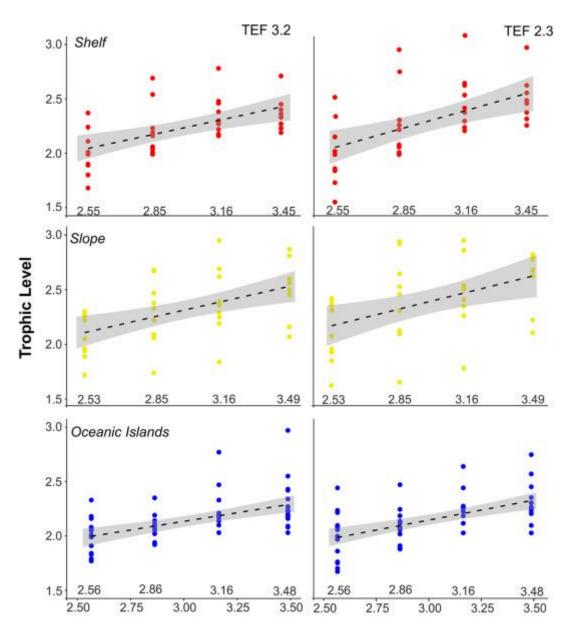


Figure 6



log₁₀(Mean size, ESD, μm)