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Microbial functional structure and stable isotopic variation of leptocephali across three current zones in the western South Pacific

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ABSTRACT

The ecology of leptocephali remains poorly known but they appear to feed on marine snow that can vary spatially and temporally according to the food web dynamics. This study provided new information about the position of leptocephali within the functional structure of microbial plankton and other food web components of the western South Pacific (WSP) region at a large geographic scale including the New Caledonia, Fiji, and Samoa islands. The hydrographic structure varied with latitude, and nutrient levels were generally low but somewhat variable. Stable isotopic signatures were examined in relation to the 3 current zones of the eastward flowing South Equatorial Countercurrent (SECC; north), the westward South Equatorial Current (SEC; mid-latitudes), and the eastward South Tropical Countercurrent (STCC; south), and all zones were found to be primarily based on a heterotrophic planktonic functioning that were co-limited by inorganic nitrogen and phosphorus, with biomasses of planktonic groups varying with depth. Isotopic signatures of leptocephali were compared to the signatures of other mesozooplankton, micronekton, and *Trichodesmium* components of the food web, and in relation to the signatures of particulate organic matter (POM) that varied among the 3 collection depths. The isotopic signatures of six taxa of leptocephali, other taxonomic groups and POM showed interesting variability according to latitude and among some stations. The presence of *Trichodesmium* at the surface in the STCC zone influenced the isotopic signatures of POM and thus the signatures of leptocephali. The signatures of leptocephali were therefore linked with the overall food web and were consistent with the larvae feeding on marine snow components of POM. The two apparent groups of leptocephali with different isotopic signatures that have also been observed in other oceanic areas may be explained by feeding behavior at different depths or on different types of marine snow.

Key words: Western South Pacific, Leptocephali, meso-macro-zooplankton, micronekton, POM, isotopic signature, microbial loop

Highlights

- Western South Pacific microbial biomass/function and stable isotopes were examined
- Primary producers, whatever the depth, were N-P co-limited
- All zones were based on a heterotrophic planktonic functioning
- Stable isotope signatures of leptocephali, POM and other components of the food web were analyzed
- POM signatures varied with depth and latitude, influencing all taxonomic groups
- The presence of *Trichodesmium* may contribute to low southern $\delta^{15}\text{N}$ values

1. Introduction

Leptocephali are the highly transparent leaf-like larvae of anguilliform fishes and their relatives of the super-order Elopomorpha (Smith, 1989; Inoue et al., 2004), which contains more than 800 species (Nelson, 2006). The order Anguilliformes is composed of 15 families and 3 suborders Anguilloidei, Congroidei and Muraenoidei. This diversity is reflected by a widespread distribution, with anguilliforms being found in almost all oceans and habitats: from southern and northern hemisphere temperate zones to tropical latitudes (Castle, 1984; Böhlke, 1989a, b). This widespread distribution is possibly due to a diversity of ecological traits, including diadromous anguillid eel and migratory and sedentary marine eel life histories. Adults may be benthic in deeper waters (Nettastomatidae, Synphobranchinae) or mesopelagic offshore (Serrivomeridae, Nemichthyidae) and many species live in coastal or continental shelf areas (Muraenidae, Congridae, Ophichthidae) (Miller and Tsukamoto, 2004). Among them, the Anguillidae forms a unique catadromous family whose leptocephali use offshore areas for development and are transported to their recruitment areas by ocean currents (Miller and Tsukamoto, 2017). Despite this wide range of juvenile and adult habitats, all anguilliform larvae live together in the upper few hundred meters of the ocean (Castonguay and McCleave, 1987; Miller, 2009; Onda et al., 2017).

Leptocephali are likely present throughout oligotrophic tropical and sub-tropical oceanic areas (Miller et al., 2019a) where primary production and the amount of trophic resources available are much lower in comparison to temperate areas (McGowan, 1974), but they also occur along continental shelves where productivity is higher (Miller, 2009). At least some species make diel vertical migrations in the open ocean from within the upper 100 m at night to as deep as about 250 m during the day (Castonguay and McCleave, 1987; Otake et al., 1998). Diel vertical migration is a ubiquitous behavior of organisms in pelagic ecosystems (Hays, 2003). It allows species to feed in the upper food-rich layers of the water column during the night and to reduce the risk of visual predation occurring in the illuminated upper layer in daylight (Tarling, 2003). The degree to which leptocephali use these behaviors in each area is unknown, and some species can be caught at the surface in neuston nets (Miller et al., 2006a; Ross et al., 2007; Miller, 2009; Quattrini et al., 2019).

The yolk reserves of the newly hatched larvae, referred to as preleptocephali, of anguillid eels are consumed within one to two weeks after hatching, then larvae actively feed in their environment (Otake et al., 1995; Tanaka et al., 1995). However, which materials in the photic zone such as phytoplankton and zooplankton or decaying organic matter leptocephali use as a food resource has remained a mystery for many years. Like many fish larvae (Last, 1978; Post

and Kitchell, 1997), mesozooplankton (> 0.2 mm) could be a food source for leptocephali, but none of these organisms were observed in leptocephalus gut contents..

However, studies that directly analyzed the stomach contents of leptocephali and found a variety of materials including zooplankton fecal pellets and discarded appendicularian (larvacean) houses (Otake et al., 1993; Mochioka and Iwamizu, 1996). Ciliates were also reported to be present (Otake, 1990; Govoni, 2010). Similar materials were also seen in photographic studies of leptocephalus gut contents, along with the typical amorphous material that is present in their guts (Miller et al., 2011, 2019b). These, which coincide with the observations of Tomoda et al. (2015, 2018) and Chow et al. (2010), are all consistent with leptocephali ingesting POM-like materials including marine snow. The latter host a very rich and abundant flora and fauna (Alldredge and Silver, 1988; Shanks and Walters, 1997; Kjørbe, 2000; Bohdansky et al., 2017).

Studies using stable carbon and nitrogen isotopic signatures of their body tissues indicated that leptocephali feed at a low level within the food web (Otake et al., 1993; Kimura et Tsukamoto, 2006; Miyazaki et al., 2011; Feunteun et al., 2015; Liénart et al., 2016; Quattrini et al., 2019). Their low trophic position was also confirmed using amino acid isotope analyses (Miller et al., 2013b). Other studies used DNA sequencing techniques with the gut contents of small leptocephali and found that a broad range of DNA sequences, including from bacteria, fungi, radiolarians, crustaceans, chaetognaths and gelatinous zooplankton were contained in their intestinal contents (Riemann et al., 2010; Ayala et al., 2018). However, tissue materials of all of those types of living or dead planktonic organisms can aggregate into POM and marine snow (Alldredge and Silver, 1988; Shanks and Walters, 1997; Kjørboe, 2000).

Therefore, although small leptocephali may be able to directly ingest some types of small organisms, the primary hypothesis about the food source of leptocephali is that they feed on marine snow. Marine snow is omnipresent and very abundant in the ocean (Alldredge, 1972, 1976; Alldredge and Silver, 1988; Turner, 2002; Pilskaln et al., 2005) and would be a readily available food source for leptocephali. Due to the oligotrophic status of tropical ocean waters, marine snow represents a major component of the pelagic ecosystem and is also the main trophic link between the photic and aphotic zones as it sinks to the ocean floor (Fowler and Knauer, 1986). These particles are colonized and enriched by a succession of bacterial communities during their fall through the water column (Pomeroy and Deibel, 1980), representing a trophic source for many grazers and are an alternative to the herbivorous food web generally occurring in more eutrophic waters (Kjørboe, 2001, 2011; Schnetzer and Steinberg, 2002; Simon et al., 2002). Marine snow particles have also been recently found to

be colonized by the less-well known heterotrophic stramenopile protists of the Labyrinthulomycetes (Li et al., 2013; Bochdansky et al., 2017), which appear to have been present in the gut contents of leptocephali in the Sargasso Sea (Miller et al., 2019b).

There is still little direct evidence of what type of marine snow might be ingested or what components of it are being assimilated by leptocephali (Miller, 2009; Feunteun et al., 2015; Liénart et al., 2016). It is also unclear whether leptocephali are feeding specialists or if any type of available particulate matter, such as marine snow, is sufficient as a food source (Miller, 2009). Feunteun et al. (2015) suggested the interspecific variety of observed head and jaw shapes (Böhlke, 1989a, b; Miller and Tsukamoto, 1994; Miller, 2009) might be related to feeding on different types of POM including marine snow. That study also found that there were two groups of isotopic signatures for species that reach large body sizes and those that do not (Feunteun et al., 2015). Those differences might be caused by the depths where they feed in the water column, or by the larvae consuming different types of POM.

Recent studies that investigated the possible contribution of marine snow to the diet of leptocephali species have analyzed POM filtered from water that was collected at various depths with Niskin bottles to evaluate the role of POM using $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ isotopic ratios (Feunteun et al., 2015) or combined with fatty acids (Liénart et al., 2016). Both studies showed gaps between the POM baseline signatures and those of leptocephali that did not show clear linkages with the POM signatures. Various possible reasons for this were discussed that include leptocephali selecting specific types of POM, or soluble fractions of POM not being retained during the filtration process. The study of Liénart et al. (2016) found latitudinal differences in the isotopic signatures across the latitudes of the western south Pacific (WSP). The fatty acid compositions of leptocephali were also studied from across the continental slope of the northwest Australia region (Deibel et al. 2012) as part of the effort to understand leptocephalus ecology.

Studies on plankton in the WSP focused on the nutrient cycle and nitrogen fixation by diazotrophic microorganisms (Campbell et al., 2005; Barboza Tenório et al., 2018; Knapp et al., 2018; Leblanc et al., 2018; Moutin et al., 2018; Van Wambeke et al., 2018), picophytoplankton distribution (Blanchot and Rodier, 1996; DiTullio et al., 2003; Grob et al., 2007; Rii et al., 2016; Zhang et al., 2008) and mesozooplankton and micronekton distribution (Hannides et al., 2013; Menkes et al. 2015; Carlotti et al., 2018), but none have attempted to examine planktonic microbial functional structure and stable isotopic variation of leptocephali.

The present study builds on the Liénart et al. (2016) stable isotope and fatty acid composition study in the WSP by expanding the range of taxa of leptocephali used and by analyzing the other organism components in relation to POM collected from different depths. The aims of this study were: 1) to reexamine the position of leptocephali within the planktonic food web of the WSP region at a large geographic scale. To this end, we performed an isotopic study of a wide diversity of organisms including leptocephali, to examine variations among leptocephali families and to evaluate the possible role of POM as a food source according to the characteristics of the environment (depth, latitude); and 2) analyze the living parts of the POM (prokaryotes and eukaryotes) that are a part of marine snow, and to consider the functional structure of the microbial loop, possibly limited by nutrients and controlled by grazing. To this end, we performed onboard grazing and nutrient enrichment experiments.

2. Material and methods

2.1. Study area and hydrographic parameters

The study was conducted in the intertropical region of the WSP that is somewhat unique in having numerous islands and atolls surrounded by deep pelagic areas (Fig. 1). The ocean currents and general hydrographic structure of the WSP have been overviewed previously (Miller et al., 2006b; 2009; Ganachaud et al., 2014), but one of the main features is the westward flowing South Equatorial Current (SEC, from 5°N and 20°S of latitude). In between the two latitudinal zones of the SEC, there is a narrow seasonally occurring eastward South Equatorial Counter Current (SECC from about 5 to 10°S) (Kessler and Taft, 1987; Chen and Qiu, 2004) (Fig. 1). To the south of the SEC there is eastward flow that has been referred to as the South Tropical Counter Current (STCC).

Sampling was made during Leg 2 of the KH-13-2 research cruise of the R/V Hakuho Maru from 4–24 February 2013. A total of 34 stations were sampled mostly at every 2° of latitude intervals in the region from New Caledonia to east of Fiji and Samoa (5°S–0°, 165°E–170°W, Fig. 1). The average depth of the ocean floor at the sampling stations ranged from 1370 to 5537 m. Salinity and temperature profiles were performed using a conductivity–temperature–depth (CTD) and oxygen probes (Sea-Bird SBE 9) from the surface to 500 m depths. The fluorescence profiles of the water column observed during the CTD down-casts were evaluated to determine the depths of the maximum *in situ* fluorescence for water sampling there during the up-casts. Water samples from subsurface (5 m), the maximum fluorescence depth (at 80–145 m, depending on station) and at 200–260 m depths (depending on station) were collected using 12 L Niskin bottles. Water samples, from the maximum fluorescence

depth were filtered immediately for analyzing dissolved inorganic nutrients [nitrate (NO_3), nitrite (NO_2), ammonium (NH_4), silicate ($\text{Si}(\text{OH})_4$) and phosphate (PO_4)], using GF/F filters (0.7 μm , Whatman, 47 mm diameter). Filtrates were stored at -20°C for NO_3 , NO_2 , NH_4 , and PO_4 , and 4°C for $\text{Si}(\text{OH})_4$ until analysis with a Skalar autoanalyzer (Strickland and Parsons, 1972).

2.2. Determining composition of particulate organic matter (POM)

2.2.1. Sampling design

POM, defined as organic particles $> 0.2 \mu\text{m}$, is composed of a mixture of living microorganisms and decaying organic matter including marine snow, so our objective was to evaluate many aspects of the overall community that can contribute to the marine snow that leptocephali might consume. We analyzed the abundance and biomass of heterotrophic prokaryotic and eukaryotic nanoplankton (defined as $0.2\text{--}2 \mu\text{m}$ and $3\text{--}20 \mu\text{m}$ respectively) (see section 2.2.3), phytoplankton (chlorophyll a biomass as a proxy of phytoplankton biomass) by size classes (picophytoplankton: $< 3 \mu\text{m}$, nanophytoplankton: $3\text{--}20 \mu\text{m}$ and microphytoplankton: $> 20 \mu\text{m}$) directly from water samples or using different filter pore sizes (see section 2.2.4). The isotopic signature of POM was also analyzed from filtered water samples. Samples were obtained with Niskin bottles during the CTD casts at 13 stations (Fig. 1) and at the same 3 depths described above. Stations were chosen to cover each general area of the study region and of the latitudinal ranges that correspond to the main current systems.

2.2.2. Samples of total POM for isotopic analysis

At each station, 6 to 14 L of water was filtered depending on visual estimation of the concentration of POM. The water was filtered under $< 10 \text{ mm Hg}$ vacuum pressure using glass fiber filters (GF/F, 47 mm diameter, $0.7 \mu\text{m}$ mesh) that were pre-combusted (490°C , 2 h) to eliminate organic carbon content. Filters containing POM samples were dried at 60°C for 12 h and analyzed for their stable isotopic content with a mass spectrometer (see section 2.5).

2.2.3. Abundance/biomass of pico- and nano-plankton

Five ml water samples were fixed with 2% formaldehyde and then frozen in liquid N_2 for analysis of heterotrophic picoplankton (e.g. bacteria). Afterwards, samples were stored at -80°C until enumeration. Each sample was analyzed for 30 s at low flow speed with a FacsCanto II cytometer (3-laser, 8-color (4-2-2), BD Biosciences) equipped with a 20 mW

488 nm coherent sapphire solid state blue laser. Data were acquired using DIVA software provided by BD-Biosciences.

For the enumeration of nanoplankton (nanoflagellates=NF; comprising pigmented (PNF) and heterotrophic (HNF) nanoflagellates), 30 mL of seawater was preserved with 3 mL of 0.2 μm filtered paraformaldehyde (2% final concentration) and placed at 4°C until analysis. Samples were concentrated to 10 mL with a filtration tower mounted with 0.8 μm pore size black polycarbonate filters (Nucleopore) and stained with DAPI (2.5×10^{-4} g L⁻¹ final concentration). Enumeration of stained nanoplanktonic cells was performed using an epifluorescence microscope (Zeiss Axioskop, x 1000 magnification). This method permitted distinguishing PNF from HNF by repeatedly interchanging the filter sets of the microscope (Caron, 1983): phototrophic cells (crimson under UV 365 nm excitation and red colored under green 450–490 nm excitation) and heterotrophic cells (blue under UV excitation and invisible under green excitation) were enumerated separately. The abundance of each group was converted to carbon biomass ($\mu\text{g C mL}^{-1}$) according to the conversion factor used in Feunteun et al. (2015).

2.2.4. Phytoplankton biomass by size classes

The phytoplanktonic community was also studied using the proxy of biomass of chlorophyll *a* (chl_a). Water samples (100 to 6000 mL) were filtered in order to sort three functional classes: picophytoplankton (< 3 μm), nanophytoplankton (3–20 μm) and microphytoplankton (>20 μm), using three successive filtrations on three membranes with different porosity (Whatman GF/F (0.7, 3, 20 μm of porosity – 25 mm diameter). Each filter was stored at -20°C until analysis. Laboratory extraction used a Turner TD 700 Fluorimeter and followed the protocol proposed by Lorenzen (1967) and Aminot and K erouel (1982). Chlorophyll *a* biomass was expressed as $\mu\text{g L}^{-1}$ and converted into carbon biomass (50 $\mu\text{g C}$ corresponding to 1 $\mu\text{g chl}_a$).

2.3. On board experiments on microbial loop functioning

Water samples were taken from the depth of maximum fluorescence (between 80–145 m) at stations 39, 41, 47, 49, 51 and 55 (Fig. 1), with a minimum of one station per current region (SEC, SECC and STCC). Originally 11 grazing experiments were attempted (3 in STCC, 5 in SEC and 3 in SECC). Unfortunately, 5 of them did not work, due to a decrease of all the microorganisms in the experiments, so the final number of grazing experiments was 6 (1 in STCC, 4 in SEC and 1 in SECC, Fig. 1). These experiments were conducted according to the

methodology developed by Bouvy et al. (2011). Two bioassay series were conducted to compare the functioning and the responses of the microbial loop to inorganic enrichment (adding nutrients) with 100% of predators (nanoplankton, potential predators, using <10 µm filtered water) for the grazing rate measurement and with 1% of predators (using 99% of 0.22 µm-filtered water and 1% of <10 µm-filtered water) for the growth rate measurement. Each experimental group was duplicated in 2x100 mL Whirlpack polyethylene sterile bags, which allow the transmission of 70% UV radiation, and incubated for 24 h on the deck of the ship with continuous renewal of seawater pumping to maintain a constant temperature in the bath. At time zero (t₀), inorganic nitrogen (N: mixture of 5 µM as NH₄Cl and 5 µM as NO₃) and inorganic phosphorus (P: 1 µM as NaH₂PO₄) were added alone or combined: +N, +P, +NP; nothing was added to the controls. In all experiments, subsamples were removed at time zero (t₀) and after 24 h (t₂₄) incubation to measure the picoplankton (picophytoplankton and heterotrophic picoplankton) abundance. In addition to picoplankton, nanoplankton abundance was measured in the 100% predator experiments. The picoplankton growth rates (µ) in each duplicate were calculated as:

$\mu = (\ln N_{24} - \ln N_0)/t$, where t was the incubation time and N₂₄ and N₀ were the picoplankton concentration at the end and at the beginning of the incubation time, respectively. The grazing rates (g expressed in day⁻¹) was calculated using the same equation in the presence of predators (see Bouvy et al., 2011).

2.4. Sampling leptocephali, meso-macro-zooplankton, micronekton and primary producers

Leptocephali, meso-macro-zooplankton (0.2–20 mm) and micronekton (20–200 mm) were collected at each station using an Isaacs-Kidd Midwater Trawl (IKMT) with an 8.7 m² mouth opening and 0.5 mm mesh. In order to limit sampling bias due to daytime leptocephali net avoidance (Miller and McCleave, 1994; Miller et al., 2006b, 2013a) and their diel vertical distribution (Castonguay and McCleave, 1987), two strategies of trawl deployment were used: during the day, a one hour oblique tow was conducted from the surface to 200 m at each station, whereas during the night a step tow sampling every 50 m from the surface to 150 m was made.

Identification of leptocephali to the family, and when possible the genus levels, was conducted on board according to the criteria summarized in Tabeta and Mochioka (1988), Böhlke (1989a, b) and Miller and Tsukamoto (2004). Among the total number of leptocephali captured during the survey (N = 538), we selected 91 leptocephali (17% of the total number) that were preserved in 99% ethanol for isotopic analysis from 16 stations (Fig. 2) belonging to

the 4 most abundant families: Serrivomeridae (N = 19; Length Range (LR) = [6.0-154.4 mm], Length Mean (LM) \pm SD = 49.4 \pm 31.2 mm), Muraenidae (N = 28; LR = [23.2-83.0 mm], LM = 51.8 \pm 13.9 mm), Congridae (N = 15; LR = [53.4-81.4 mm], LM = 81.4 \pm 53.4 mm) and Nemichthyidae including *Nemichthys* (N = 22; LR = [37.2-259.0 mm], LM = 123.7 \pm 73.9 mm) and *Avocettina* (N = 31; LR = [20.3-223.0 mm], LM = 113.7 \pm 60.1 mm). Among the Congridae, *Ariosoma sp.* (N = 3; LR = [77.2-220.8 mm], LM = 143.3 \pm 72.5 mm) were separated from other Congridae (genus *Conger*, *Gnathophis*, *Gorgasia* and *Bathycongrus*) because they are a different subfamily, and because *Ariosoma* leptocephali can have different isotopic signatures than other species of the Congridae and other families that also have smaller maximum lengths (Miyazaki et al., 2011; Feunteun et al., 2015). We also used the two morphologically distinct genera of Nemichthyidae: *Avocettina sp.* (N = 19) and *Nemichthys sp.* (N = 10) for comparison of isotopic signatures. Not all taxa of leptocephali were evenly distributed among latitudes, which contributed to the imbalance of specimens among latitudes and sample sizes of some taxonomic groups (see Miller et al., 2006b, Liénart et al., 2016).

Meso- and macro-zooplankton were identified to class, subclass, order, family or species (copepods to genus or species when it was possible) for euphausiids (N = 26), chaetognaths (N = 7), cephalopods (N = 8), polychaetes (N = 5) and copepods (N = 49). Euphausiids were separated into two groups according to their size (< 10 mm and > 10 mm). Copepods were composed of Calanoidae that included the species *Metridia venusta* (N = 9), *Pleuromamma abdominalis* (N = 10), *Pleuromamma xiphias* (N = 9) and the genera *Candacia* (N = 8), *Temora* (N = 2) and *Anomalocera* (N = 1). Among micronekton, 3 groups of fish were considered: family Myctophidae (N = 12), tuna larvae (Scombridae) (N = 3) and unidentified fish larvae (N = 16). One genus of primary producer, a cyanobacteria *Trichodesmium sp.* (N = 2) that formed small aggregates, was collected from two IKMT station samples when it was present, and it was isolated under a binocular microscope for collection. All samples were stored in 70% ethanol for further isotopic analysis.

2.5. Stable isotope analyses

Filters containing POM were dried at 60°C for 12 h, and plankton (e.g. copepods, cephalopods...), micronekton and small fish were rinsed 3 times with distilled water and lyophilized before analysis. These organisms were analyzed without any dissection, except for big euphausiids for which shell and head were removed. Plankton samples were not decarbonated or delipidated because similar studies did not find any significant changes in $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ ratios before and after acidification (Bode et al., 2004; Chouvelon et al., 2014)

or the delipidation processes (Bodin et al., 2007; Chouvelon et al., 2014). Small pelagic fishes and larvae were used entirely, while only muscles were sampled in larger individuals. Leptocephali bodies without the head were analyzed, and were not decarbonated or delipidated because this appears not necessary for muscle (Pinnegar and Polunin, 1999). For whole individuals, preliminary tests did not show significant differences between treated and untreated samples (Bourillon, comm.pers).

Samples were analyzed with a Thermo Scientific Delta V Advantage mass spectrometer coupled to a Thermo Scientific Flash EA1112 elemental analyser to determine their carbon and nitrogen stable isotope contents. The carbon and nitrogen isotope ratios were reported according to the standard delta notation $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$, where: $\delta X = [(R_{\text{Sample}}/R_{\text{Reference}}) - 1] \times 1000$, where $X = \delta^{13}\text{C}$ or $\delta^{15}\text{N}$ and R is the ratio $^{13}\text{C}:^{12}\text{C}$ or $^{15}\text{N}:^{14}\text{N}$ in the sample and in the reference material. References are Vienna Pee Dee Belemnite (VPDB) (Craig, 1957) for C and to atmospheric nitrogen for N (Peterson and Fry, 1987; Ehleringer and Rundel, 1989) and results are expressed in units of ‰ \pm standard deviation (SD).

2.6. Statistical analysis

Before comparison analysis, data were evaluated for homoscedasticity (Levene-test) and normality of distribution (Shapiro-Wilk-test). If the data did not meet the assumptions, they were log or exponential transformed. When the assumptions were met, comparisons of two averages were performed using t-tests and comparison of more than two variances with ANOVA followed by post-hoc Tukey tests when ANOVA were significant ($p < 0.05$). In the case where assumptions were not met, non-parametric tests were used: two samples were compared with non-parametric Mann-Whitney-tests (MW), and more than two samples were compared with Kruskal-Wallis-test (KW) followed by a multiple comparison test (bilateral) when KW were significant ($p < 0.05$). All averages were expressed with their standard deviation (\pm SD). Statistical analyses were performed with Statistica software (Statsoft v.6.1, Tulsa, OK, USA). Cartography was made using “R” software and the package “map” and “marmap” (Fig 1.a).

3. Results

3.1. Hydrographic structure

Hydrographic sections constructed from the CTD casts along or near 180° (middle line of stations) and 170°W (eastern line of stations) (Fig. 1) showed the typical structure of temperature and salinity that occurs in the WSP (Miller et al., 2006b; Schabetsberger et al.,

2016). The warm water of the upper mixed layer (28–30°C) was deeper in the north (~ 100 m) and then steadily became shallower to the south, with surface temperatures of 26°C at the southernmost stations at 24°S (Figs. 3a, e). Temperatures at depths of around 250 m, which is likely to be the approximate maximum depth reached by leptocephali during the day if they vertically migrate deeper from the upper 100 m where they reside at night (Castonguay and McCleave, 1987), were 18–19°C. The deeper mixed layer in the north included the lower salinity water of the “Fresh Pool” that is typically present there, with salinities of about 34.2–35.5 (Figs. 3b, f). Deeper, at about 75–200 m the subsurface layer of Subtropical Underwater (STUW) was present (see Schabetsberger et al., 2016) that had maximum core-area salinities of 36.2 at 180° and 36.3 at 170°W. This water forms at the surface through evaporation farther to the east and is subducted into the thermocline as it moves to the west.

3.2. Chemical and biological water mass parameters

Average nutrient concentrations were generally low, but typical of tropical marine waters (Table 1). All nutrients, except NH_4 , which had less variability between the current zones, were higher in the SECC than SEC and STCC. However, concentrations of NO_3 ranged from 3.5 to 155.3 $\mu\text{g NO}_3 \text{ L}^{-1}$ and were variable in the SEC stations (range 3.5–45.3 $\mu\text{g NO}_3 \text{ L}^{-1}$) and the SECC (range 4.7–155.3 $\mu\text{g NO}_3 \text{ L}^{-1}$). Concentrations of NO_2 ranged from 0 to 19.1 $\mu\text{g NO}_2 \text{ L}^{-1}$ and were variable in all the current zones. Concentrations of PO_4 ranged from 27.1 to 77.8 $\mu\text{g PO}_4 \text{ L}^{-1}$ and were variable within the SECC (range 27.1–77.8 $\mu\text{g PO}_4 \text{ L}^{-1}$). Silicate concentrations ranged from 61.2 to 512.6 $\mu\text{g Si(OH)}_4 \text{ L}^{-1}$ and were variable in all the currents, with maximum variability occurring in the SEC (range 63.9 to 512.6 $\mu\text{g Si(OH)}_4 \text{ L}^{-1}$). Station 63 at the eastern side of the SECC zone had lower concentrations of all parameters compared to the 2 stations to the east, and particularly for NO_3 and NO_2 . Station 49 in the north-central SEC zone also had some lower nutrient values. Station 39 to the east of Vanuatu had an unusually high Si(OH)_4 value (512.6 $\mu\text{g L}^{-1}$; 137.5 $\mu\text{g L}^{-1}$ maximum for all other stations).

Maximum fluorescence of the chlorophyll occurred at depths of about 100 m across the 180° transect but the maximum values became lower and slightly deeper in the southern half of the 170°W transect in the east (Figs. 3c, g). The exact structure and variability of the chlorophyll maximum layer across these latitudes could not be observed because the CTD stations were too far apart, however vertical profiles for each CTD stations were available (vertical lines, Fig. 3). Fluorescence values were consistently lower in the southern half of the eastern most 170°W transect.

Oxygen concentration patterns differed from other parameters. A layer of highest concentrations occurred in the south at depths just above the chlorophyll maximum. However, this structure almost disappeared in the far north (Figs. 3d, h). The lowest oxygen levels occurred deeper than 250–300 m in the northern half of the section below the STUW.

Planktonic organisms from POM were separated by size groups (Fig. 4). POM was generally dominated by heterotrophic picoplankton (e.g. bacteria; significantly higher than other 4 groups; K-W test, $H_{(4, N=225)} = 119.4$, $p < 0.001$) that accounted for 50 to 90% of the biomass, followed by heterotrophic nanoflagellates (5–27%) and the three other size groups representing less than 15% (picophytoplankton (0–15%), nanophytoplankton (0.4–12%), and microphytoplankton (0–4%). Total biomass at each station was not significantly different according to currents (K-W test, $H_{(2, N=225)} = 0.67$, $p = 0.71$), but significantly differed according to depths (K-W test, $H_{(4, N=225)} = 40.56$, $p < 0.001$). At the deepest layer (200–260 m), planktonic biomass was lower than at the shallower depths (all comparison tests: $p < 0.001$). However, the composition of planktonic organisms changed according to the depths. At surface (5 m), only the heterotrophic picoplankton dominated with all other groups having low concentrations (Multiple comparison test, $p < 0.01$). At maximum fluorescence depth (80–145 m), heterotrophic picoplankton followed by heterotrophic nanoflagellates were dominant and microphytoplankton showed the lowest concentration (Multiple comparison test, $p < 0.001$). At the deepest layer, heterotrophic picoplankton followed by heterotrophic nanoflagellates were also dominant but the three other groups were almost absent (Multiple comparison test, $p < 0.05$).

3.3. Microbial loop functioning

Growth and grazing rates were assessed for picophytoplankton and heterotrophic picoplankton (Figs. 5, 6) in water samples collected in the 3 current zones after nutrients were added or not. In the control, picophytoplankton net growth rate was negative or low, ranging from $-2.1 \pm 2.6 \text{ day}^{-1}$ in the SEC to $0.43 \pm 0.60 \text{ day}^{-1}$ in the STCC (Fig. 5). Grazing rates on picophytoplankton ranged from no grazing in the SECC to $1.4 \pm 1.3 \text{ day}^{-1}$ in the SEC (Fig. 6). Net growth rates of heterotrophic picoplankton were higher and less variable than for picophytoplankton, and ranged from $3.9 \pm 0.01 \text{ day}^{-1}$ in the STCC to $5.2 \pm 0.008 \text{ day}^{-1}$ in the SECC (Fig. 5). The grazing rate on heterotrophic picoplankton ranged from 3.4 ± 0.03 in the SECC to $4.3 \pm 0.55 \text{ day}^{-1}$ in SEC (Fig. 6).

The addition of N, P or NP (both inorganic nitrogen and phosphorus) did not change the net growth rate of picophytoplankton (KW: $H_{(3, N=48)} = 0.79$, $p > 0.05$) while net growth rate

of the heterotrophic picoplankton increased when adding a NP combination (KW: $H_{(3, N=48)} = 20.61$, $p < 0.001$ and multiple comparison test: $p < 0.05$; Fig. 5). This pattern suggested a co-limitation of nitrogen and phosphorus for the development of heterotrophic picoplankton in the 3 current zones.

The addition of N, P or NP did not change grazing rate on picophytoplankton but increased grazing rate on heterotrophic picoplankton in the presence of the NP combination in the STCC, and inversely decreased grazing rate of heterotrophic picoplankton in the presence of the NP combination in the SECC (Fig. 6).

To summarize, heterotrophic picoplankton was dominant in terms of biomass (Fig. 4), had a better growth (Fig. 5) and was grazed in the 3 current zones (Fig. 6). In contrast, picophytoplankton biomass was low (Fig. 5), grew only in the STCC (Fig. 5), was weakly grazed in the SEC and STCC and was not detectable in the SECC experimental treatments (Fig. 6). Therefore, fluxes inside the food web were mainly based on the heterotrophic pathway. However, the heterotrophic picoplankton exhibited a co-limitation of N and P for its development in the 3 current zones.

3.4. Isotope analyses of POM, meso-macro-zooplankton, micronekton and leptocephali

3.4.1. POM isotopic signatures

Isotopic signatures of POM were highly variable according to currents and depths (Fig. 7). Mean values varied from -26.2 to -23.16‰ in $\delta^{13}\text{C}$ ($-25.1 \pm 1.24\text{‰}$ overall) and from 1.04 to 9.93‰ in $\delta^{15}\text{N}$ ($5.57 \pm 2.74\text{‰}$) (Fig. 7). A significant effect of depth was observed for $\delta^{13}\text{C}$ (two-way ANOVA, $F_{2,36} = 59.79$, $p < 0.001$ for depth, $F_{2,36} = 1.88$, $p > 0.05$ for current zones, $F_{4,36} = 2.21$, $p > 0.05$ for interaction), and more depleted $\delta^{13}\text{C}$ values were found for the deep and maximum of fluorescence POM compared to at the surface (Tukey: $p < 0.001$ for both) (Fig. 7). $\delta^{13}\text{C}$ values were not different between deep and maximum fluorescence (Tukey: $p > 0.05$). For $\delta^{15}\text{N}$, significant effects were found both according to depth and current zones (two-way ANOVA, $F_{2,36} = 36.95$, $p < 0.001$ for depth, $F_{2,36} = 13.09$, $p < 0.001$ for currents, $F_{4,36} = 0.47$, $p > 0.1$ for interaction). More $\delta^{15}\text{N}$ enriched values were found for POM from the SEC and SECC compared to the STCC (Tukey: $p < 0.01$ and $p < 0.001$ for SEC and SECC, respectively), but $\delta^{15}\text{N}$ values were not different between the SEC and SECC (Tukey: $p > 0.05$) (Figs. 7, 8). Finally, POM from deep samples was significantly more enriched in $\delta^{15}\text{N}$ than POM from the maximum fluorescence and surface layers (Tukey: $p < 0.001$ for both), with those layers being not significantly different (Tukey: $p > 0.05$) (Fig. 7).

The isotopic landscapes of the region showed a tendency for a north south gradient of $\delta^{15}\text{N}$ at all depths, with a more patchy distribution at the maximum fluorescence and deeper depths than at surface depths (Fig. 8 left panels). Highest values were observed in the north for the 3 layers and lowest values in the south also for the 3 layers. The deep layer had higher $\delta^{15}\text{N}$ values (11‰) while they only reached 6.8‰ at surface and max fluorescence layers as also reflected in Figure 7. Lowest values of $\delta^{15}\text{N}$ were observed at the surface southern stations with values of 1.2‰ areas with the lowest values increasing from the surface to the deep layer (1.2; 2.4 and 6‰ at the surface, maximum fluorescence and deep, respectively).

$\delta^{13}\text{C}$ isotopic landscapes ranged over approximately 2‰ but they were more patchy regardless of the depth compared to $\delta^{15}\text{N}$ (Fig. 8, right panels). At the surface, lowest values of POM (-24.8‰ to -24‰) were observed along a diagonal roughly from south of New Caledonia to south of the Samoan islands between areas of higher $\delta^{13}\text{C}$ (about -23‰) to the north and southeast. At maximum fluorescence depths, $\delta^{13}\text{C}$ of POM was patchier and partially inverted compared to northern latitudes, with lowest values of -26‰ to -25.2‰ observed along a diagonal going from the southwest to the north east of the study area. At the deep layer, higher $\delta^{13}\text{C}$ POM values occurred along southern latitudes and patches of higher $\delta^{13}\text{C}$ values were present within the relatively lower values (-26.5‰) in the north.

3.4.2. Meso-macro-zooplankton isotopic signatures

A variety of meso-macro-zooplankton taxa with different feeding ecologies were analyzed for isotopic signatures to compare to leptocephali ones (Fig. 9). For euphausiids, no difference was found in both $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ isotopic ratios between small euphausiids (< 10 mm) and bigger euphausiids (> 10 mm) (t-test: $t_{24} = 0.62$, $p > 0.05$ and $t_{24} = 0.62$, $p > 0.05$, respectively for $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$). Thus, they were then pooled into one single group, the euphausiids (Fig. 9).

Within each of the 3 current zones most taxa of meso- and macro- zooplankton $\delta^{15}\text{N}$ isotopic ratio values were similar among the taxa due to high variability (ANOVAs: $p > 0.05$) despite the fact that large predatory species (Myctophidae, Sagittoidea, Polychaeta and Cephalopoda) tended to have higher averages (notably SECC and STCC) relative to other species. The crustacean taxa (copepods and euphausiidae) had relatively higher values in the SECC zone (Fig. 9). Most individual taxa showed decreasing $\delta^{15}\text{N}$ values from the north (SECC) to the south (STCC) similar to the pattern observed for POM. However, a few taxa such as chaetognaths and cephalopods had similar values in the SECC and SEC zones and did

not follow that decreasing trend. This was similar to the little difference observed in $\delta^{15}\text{N}$ values between POM in the maximum fluorescence layer for the SECC and SEC zones. There were also few consistent differences in the $\delta^{13}\text{C}$ values of each taxa between the SECC and SEC zones, but most taxa had lower values in the STCC zone. The cyanobacteria, *Trichodesmium* sp. was only collected at stations 55 and 56 in the STCC zone (likely due to a bloom making them visible) and had high $\delta^{13}\text{C}$ isotopic values ($-14.59 \pm 3.43\text{‰}$) and very low $\delta^{15}\text{N}$ ratios ($0.54 \pm 0.39\text{‰}$) (Fig. 9).

Only the taxa present in sufficient numbers (*i.e.* copepods and euphausiids) in the three zones were used to compare the isotopic ratios according to the currents. For euphausiids, differences were found in both $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ isotopic values among currents (ANOVA: $F_{2,23} = 10.29$, $p < 0.001$ and ANOVA: $F_{2,23} = 8.30$, $p < 0.005$, respectively) (Fig. 9). Individuals captured in the STCC had lower $\delta^{13}\text{C}$ (Tukey: $p < 0.01$) compared to the two other zones and euphausiids captured in the SECC had higher $\delta^{15}\text{N}$ isotopic values (Tukey: $p < 0.05$) relative to the SEC and STCC zones. For copepods, differences in their $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ ratios among zones were found (KW: $H_{(2, N=48)} = 7.80$, $p < 0.05$ and ANOVA: $F_{2,45} = 36.84$, $p < 0.001$, respectively). Individuals in the STCC had lower $\delta^{13}\text{C}$ than those of the SEC (multiple comparison test: $p < 0.05$). Copepods $\delta^{13}\text{C}$ values in the SECC zone were at an intermediate level between the two other zones. $\delta^{15}\text{N}$ ratios of copepods in the three zones were significantly different (Tukey: $p < 0.005$) with the STCC showing the lowest $\delta^{15}\text{N}$ values, the SEC intermediate, and the SECC the highest ones.

3.4.3. Isotope data analysis of mesopelagic fishes from micronekton

A range of sizes of small fish were analyzed, with undetermined larvae being the smallest (around 10 mm), tuna larvae being medium-sized (around 20 mm) and Myctophidae being largest (around 40 mm). No difference was found among the $\delta^{13}\text{C}$ values (KW: $H_{(2, N=31)} = 1.03$, $p > 0.05$) of these fishes because they all had relatively similar values in each of the current zones, except that fish larvae and Myctophidae had relatively high values in the STCC zone in the south (Fig. 11). Significant differences of $\delta^{15}\text{N}$ signatures were noticed among the three groups of fish (ANOVA: $F_{2,28} = 5.48$, $p < 0.01$) with the Myctophidae being characterized by significantly higher values of $\delta^{15}\text{N}$ compared to fish larvae (Tukey, $p < 0.05$). The $\delta^{15}\text{N}$ isotopic ratios of fish larvae and Myctophidae both decreased from north to south, and the tuna larvae in the STCC were also low.

For the Myctophidae, $\delta^{13}\text{C}$ ratios were different among the three zones (KW: $H_{(2, N=12)} = 6.58$, $p < 0.05$), but the multiple comparison test could not identify which area(s) were

different from other(s) ($p > 0.05$). Their $\delta^{15}\text{N}$ signatures were not different among the three zones (ANOVA: $F_{2,9} = 4.11$, $p > 0.05$) even though there was a decreasing trend to the south. For the fish larvae, no difference was found for their $\delta^{13}\text{C}$ values (ANOVA: $F_{2,13} = 1.73$, $p > 0.05$), but significant differences were found for their $\delta^{15}\text{N}$ signatures (ANOVA: $F_{2,13} = 7.52$, $p < 0.01$). Fish larvae in the STCC had the lowest $\delta^{15}\text{N}$ ratios ($4.92 \pm 1.79\text{‰}$), which were significantly different from those of the SECC (Tukey: $p < 0.05$) that had the highest values ($10.64 \pm 2.98\text{‰}$). Individuals captured in the SEC were at medium levels ($7.50 \pm 2.02\text{‰}$) between the two others. Tuna larvae were only collected in the STCC (Supplementary Table), so no comparisons between currents was possible.

3.4.4. Isotopic composition of leptocephali

Similar to the plankton and other nekton taxa, the leptocephali showed significant variability of $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ signatures (KW: $H_{(2, N=91)} = 14.55$, $p < 0.001$; KW: $H_{(2, N=91)} = 39.92$, $p < 0.001$; respectively) among the 3 currents and the taxa themselves (KW: $H_{(5, N=91)} = 36.49$, $p < 0.001$; KW: $H_{(5, N=91)} = 34.80$, $p < 0.001$; respectively; Fig. 10). As was observed in the western Indian Ocean (Feunteun et al., 2015), the leptocephali could be separated into two groups of taxa based on their isotopic signatures. On average, Group 1 leptocephali (Congridae, Muraenidae, Serrivomeridae) (Fig. 10, blue colors) had lower $\delta^{13}\text{C}$ and higher $\delta^{15}\text{N}$ values (MW: $U_{(1, N=91)} = 459$, $Z = 4.03$, $p < 0.001$; MW: $U_{(1, N=91)} = 433$, $Z = 4.24$, $p < 0.001$; respectively) compared to the Group 2 leptocephali (*Ariosoma*, *Avocettina*, *Nemichthys*) (Fig. 10, red colors), which are species that grow to larger sizes than those of Group 1. Although there was some overlap in the signatures of individual larvae of each group and there were different sample sizes among taxa, these results were verified for both $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ isotopic ratios within each current zone (MW-tests, $p < 0.01$ for all tests). Serrivomerid leptocephali had the highest $\delta^{15}\text{N}$ signatures in both the SECC and SEC where they were present. In the SEC with the most larvae, the $\delta^{15}\text{N}$ values of Groups 1 ($8.60 \pm 2.49\text{‰}$) and 2 ($4.97 \pm 1.05\text{‰}$) were significantly different and $\delta^{13}\text{C}$ values between Group 1 and 2 (MW-tests, $p < 0.01$ for all tests). The individual taxa within each group also showed both decreased $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ mean values from north to south in the 3 current zones (Supplementary Table).

Interestingly, the signatures of most of the leptocephali in each of the 3 zones had a negative relationship between $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ (shaded ovals in Fig. 10) when outliers from a few stations were excluded. In the SECC an *Avocettina* had a very high $\delta^{15}\text{N}$ (from the northeastern-most station 64) and other individuals had much lower values that were from a

northeastern station. In the SEC, the leptocephali had a similar range of $\delta^{13}\text{C}$ values, but most had lower $\delta^{15}\text{N}$ signatures than those of the SECC. The exception were the larvae from station 65 that was located just southwest of the island of Samoa, which had a higher range of $\delta^{15}\text{N}$. Other outliers were Muraenidae larvae from the southeast area of the SEC. In the STCC, the Congridae larvae (station 34 and 37 in the southwest) showed the highest $\delta^{15}\text{N}$ values, but the Muraenidae and nemichthyid leptocephali overlapped in their signatures with the Group 2 larvae. A 114 mm *Bathycongrus* leptocephalus had lower $\delta^{15}\text{N}$ and C values than other congrids.

3.4.5. Comparison of leptocephali isotopic ratios to other groups

The plots of all the mean values of the different taxa of leptocephali, crustaceans, micronekton, POM and *Trichodesmium* in each of the 3 current zones showed some interesting consistencies but also some variability among the zones (Fig. 11). The cyanobacteria, *Trichodesmium* sp., were only collected in the STCC zone (although likely present), had the highest $\delta^{13}\text{C}$ ratio and the lowest $\delta^{15}\text{N}$ ratio. The $\delta^{13}\text{C}$ gap between POM from the 3 depths and the taxa of marine organisms was about the same within the northern current zones but was reduced in the STCC. The overlap between the $\delta^{15}\text{N}$ values of POM and organisms increased from north to south. In the SECC many taxa had higher $\delta^{15}\text{N}$ than the deep POM, but in the STCC almost no taxa had higher values.

The relationship between the taxa of leptocephali, crustaceans and micronekton also showed some variability among the current zones (Fig. 11). The micronekton usually had higher $\delta^{13}\text{C}$ values than leptocephali, and their $\delta^{15}\text{N}$ values were mostly higher than leptocephali in the SECC and SEC and showed more overlap in the STCC. The crustacean taxa showed various types of overlapping values with lower $\delta^{13}\text{C}$ of some taxa compared to leptocephali in the SECC or mostly overlapping values in the SEC. In the SEC, *Avocettina* and *Nemichthys* had the lowest $\delta^{15}\text{N}$ values, which were close to the values of POM at the surface and maximum fluorescence (Fig. 11). In the STCC, the $\delta^{15}\text{N}$ values of *Ariosoma* and *Avocettina* were almost the same low value, which was slightly higher than the POM at the surface and slightly lower than the POM at the maximum fluorescence (Fig. 11).

4. Discussion

4.1. Pelagic environment in the WSP region

The hydrographic structure observed in the WSP in February 2013 included the low salinity Fresh Pool in the STCC zone and part of the northern SEC, and the warmer water

mixed layer of surface water became gradually shallower and mostly disappeared in the southern STCC zone, which are typical features of the region (Miller et al., 2006b; Schabetsberger et al., 2016). The highest salinities of the STUW centered on the 100–200 m layer were only present in the STCC and SEC zones. The maximum fluorescence layer was centered at about 100 m and deeper in the east. Therefore, the 3 latitudinal current zones we used for analyses had different hydrographic characteristics. Our zonal separations were only approximations and there is likely some overlap in water masses that results from eddy exchange and countercurrent jets as can be inferred from the complexity of the currents in the region (Ganachaud et al., 2014), which might account for some of the variability that was observed. We also compared some microbial biomasses, the functioning of the microbial loop, and POM signatures in the warm surface layer (5 m) above the STUW, the maximum fluorescence layer near the top of the thermocline within the STUW and the deeper layer in the middle of the thermocline and below the STUW.

The concentration of nutrients in our study showed the typical characteristics of oligotrophic ecosystems, as was observed in another recent study in the WSP (Bonnet et al., 2018). But nitrogen and phosphate concentrations were slightly higher in New Caledonian waters during an El Niño episode (Tenório et al., 2018) or in the Southwestern Pacific Ocean (Campbell et al., 2005) compared to our study. However, there was some variability in nutrient levels among our stations in 2013 such as the unusually high silicate value (St. 39) in an area where there could be eastward flowing water of the Coral Sea Countercurrent (Ganachaud et al., 2014) passing through the Vanuatu Islands. Upwelling of high silicates (Leblanc et al., 2018) and nitrates (Blanchot and Rodier, 1996) has been observed in that general area. DiTullio et al. (2003) also found some variability in nutrient levels along 170°W in the South Pacific Ocean.

4.2. Characteristics of the microbial loop structure

The three depths sampled in the upper 260 m of the water column were dominated by heterotrophic organisms that were primarily heterotrophic picoplankton (e.g. bacteria). In the North Pacific Ocean (137–155°E), Yamaguchi et al. (2004) showed the importance of bacterial biomass, which was similar to our study. Secondly, heterotrophic nanoflagellates also were relatively abundant especially in the maximum fluorescence depths (80–145 m depending of stations).

Within the primary producers, small phytoplanktonic cells (pico-nano-phytoplankton) were more abundant compared to the very low biomass of microphytoplankton. Due to the

low nutrient concentrations in this area, microphytoplankton organisms were generally scarce along 170°W (our easternmost transect) in the offshore parts of the gyre compared to higher latitudes during the summer season (DiTullio et al., 2003). In contrast, picophytoplankton dominated across all 3 of our current zones as was observed in the studies of Blanchot and Rodier (1996), DiTullio et al. (2003), Campbell et al. (2005) and Grob et al. (2007). Their abundance in previous studies was highest between the surface (5 m) and around 80–100 m (corresponding to a maximum fluorescence depths) in our study.

The presence of *Trichodesmium* (i.e. a diazotroph cyanobacteria, which fixes atmospheric N₂) was detected in the STCC zone in February 2013 in our study, as was observed in previous studies (e.g. Bonnet et al., 2018). However, Campbell et al. (2005) also found *Trichodesmium* in the SEC in March-April. The occurrence of *Trichodesmium* fluctuates near New Caledonia (Tenório et al., 2018), maybe due to a limitation by nitrate availability (Moutin et al., 2005). In oligotrophic ecosystems that are “nitrogen-depleted”, diazotroph organisms can overcome the N-limitation and are thus important contributors to the primary productivity (e.g. Bonnet et al., 2018).

The microbial loop at the maximum fluorescence layers was examined by onboard experiments. The net growth rate and grazing rate of picophytoplankton and heterotrophic picoplankton were determined using water from the maximum fluorescence depth. Consequently, these results cannot be extrapolated to the entire photic zone. The main results showed the relatively high abundance, growth rate of heterotrophic microorganisms and grazing rate on heterotrophic microorganisms. As far as we know, this is the first time that these kinds of measurements have been performed on components of the microbe food web in this region. Carlotti et al. (2018) studied mesozooplankton structure and functioning in the south western Pacific, but not the associated microbe food web, so no comparison is possible. Despite the low nutrient concentration observed *in situ*, no growth of phytoplankton community was induced when adding nitrogen and phosphorus. Heterotrophic picoplankton grew well and was well grazed, as demonstrated in an oligotrophic aquatic system (Samuelsson and Andersson, 2003). Therefore, fluxes inside the microbe food web were mainly based on the heterotrophic pathway. However, in the 3 current zones, the development of heterotrophic picoplankton was co-limited by nitrogen and phosphorus. The picophytoplankton community did not respond to the addition of nitrogen and phosphorus possibly because heterotrophic picoplankton are much more competitive than picophytoplankton (Thingstad and Rassoulzadegan, 1995).

Even if all the grazing experiments did not work, the tendency was clear that the water column was dominated by heterotrophic organisms, with relatively high growth rates of heterotrophic microorganisms and also high grazing rates on heterotrophic microorganisms. In addition, primary producers were N-P co-limited. All of our observations appeared to be consistent with typical characteristics of oligotrophic areas.

4.3. Variability of POM signatures with latitude and depth

One of the most interesting findings of our study was that several patterns were observed in the isotopic signatures of the POM (including marine snow aggregates), which was analyzed from the water samples filtered with 0.7 μm mesh at different depths and latitudes. A patchy isotopic landscape of $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ that was observed that could be a result of the eddies that occur within the southwest Pacific Ocean (e.g. DiTullio et al., 2003). The POM of the SECC and SEC zones was enriched in $\delta^{15}\text{N}$ compared to the STCC zone in the south. There was some east-west variability in the basic latitudinal trend among the zones, but the $\delta^{15}\text{N}$ values were generally higher in the north and lower in the south. Similar latitudinal changes in isotopic signatures have been observed in other regions and in highly mobile marine animals such as tunas (Estrada et al., 2005; Ménard et al., 2007; Lorrain et al., 2015). Distinct geographic differences in the $\delta^{15}\text{N}$ signatures of POM were found by Houssard et al. (2017), which are consistent with the decreasing $\delta^{15}\text{N}$ values of POM and other organisms from north to south in our study. In the STCC, the *Trichodesmium* present in this area would be a greatly depleted source of $\delta^{15}\text{N}$, especially for the isotopic signatures of surface POM (our results; Waite et al., 2007 ; Somes et al., 2010; Bonnet et al., 2018).)Nevertheless, Campbell et al. (2005) found *Trichodesmium* in the SEC in March-April and this may affect the isotopic POM signatures of all the food webs in this area during this season.

In addition, depth patterns were observed with differences in both $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ signatures occurring according to the sampling depth of POM. POM composition in the deep layer (200–260 m) that was likely dominated by heterotrophic organisms (pico-nanoplankton) was enriched in nitrogen compared to the surface and maximum fluorescence (80–145 m) depths, and higher $\delta^{13}\text{C}$ values were observed in the surface POM. Miyazaki et al. (2011) also found the highest $\delta^{15}\text{N}$ signatures of POM in the deeper layers, and their signatures at the surface (5 m) overlapped with intermediate depths. Hannides et al. (2013) found that POM also had the lowest $\delta^{15}\text{N}$ near the surface ($< 2\text{‰}$ above 50 m) and the values remained much higher ($> 8\text{‰}$) below about 120 m in the North Pacific subtropical gyre. The same basic trend was also seen in the southeastern North Pacific for POM (0.7–53 μm) where there were high

N values ($> 8\text{‰}$) below about 100 m (Williams et al., 2014). In contrast, POM at the surface (5 m, 5‰) also had slightly higher $\delta^{15}\text{N}$ values than at 100 m (4‰) in the western Indian Ocean in January-February 2010 (Feunteun et al., 2015). These different $\delta^{15}\text{N}$ isotopic signatures were likely reflecting the different relative biomasses of the various sizes of heterotrophic and autotrophic microorganisms detected at the different depths.

4.4. Representation of the food web in the western South Pacific Ocean

Our 3 approximated current zones were spread across a region with many island groups and various latitudinal gradients in many parameters such as temperature, salinity, and productivity (e.g. DiTullio et al., 2003), so they only roughly characterized the diversity of biological oceanographic conditions in those regions of the WSP. As a result, signatures likely differed among some stations within each designated zone because of factors such as eddies and current jets. Despite that likelihood, some overall patterns in the isotopic signatures of organisms were observed with depth and latitude that are reflecting changes in some aspects of the trophic structure of the food webs there.

Firstly, Minagawa and Wada (1984) found that the trophic enrichment factor ($\text{TEF}_{\text{prey-predator}}$) for nitrogen isotopic values between two trophic level is generally about $3.4 \pm 1.1\text{‰}$ (see also Post, 2002), and many authors consider the carbon enrichment to be $\text{TEF}_{\text{prey-predator}} = 1\text{‰}$ (DeNiro and Epstein, 1978; Rau and Anderson, 1981; Fry and Arnold 1982; Peterson and Fry, 1987). If these TEF factors are applied to our results, it suggests there were two kinds of sources of the food web, possibly in part because of *Trichodesmium* was only observed in STCC zone. Our isotopic analyses showed some expectable patterns such as the mean values of the micronekton signatures being partially separated from the crustaceans and leptocephali, and they were most distinctly separated with both higher $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ signatures in the SEC zone. Cephalopods (Hanlon, 1987), chaetognaths (Terazaki and Marumo, 1982), annelids and myctophids are carnivores that feed on macro-mesozooplankton and small fish larvae. Small fishes are often known to feed on meso- microzooplankton (Last, 1978; Mehner et al., 1996; Post and Kitchell, 1997). Both leptocephali and crustaceans however, showed various patterns of signatures compared to each other in the 3 current zones. Some copepods can be considered as carnivorous like *Pleuromamma abdominalis*, *Pleuromamma xiphias* (Gueredrat, 1971), *Metridia venusta* and some species of Calanoidae since they showed relatively high values of $\delta^{15}\text{N}$. Adult stages of the genus *Euphausia* are generally carnivorous, but the species found in this study were small and they can consume POM (Schnetzler and

Steinberg, 2002). Fish larvae can also consume POM (Larson and Shanks, 1996), which might explain their intermediate $\delta^{15}\text{N}$ values in our study.

A large gap in $\delta^{13}\text{C}$ between POM and grazers was observed in our data with each current zone. A similar gap between POM and some leptocephali and potential grazers was observed in similar isotopic analyses made to the west of the Mascarene Plateau in the Indian Ocean (Feunteun et al., 2015; Lienart et al. 2016). Consequently, if “grazers” do feed on POM, they probably select or assimilate some particular kinds of POM (Feunteun et al., 2015; Lienart et al. 2016), and this is discussed more for leptocephali in the next section. A complementary hypothesis is that the POM grazers would likely be feeding on “marine snow”, and not on the free-living component of POM, which was dominated by heterotrophic picoplankton in our study. This may be one explanation for the large gap in $\delta^{13}\text{C}$ between POM and grazers in our data within each current zone. In the south (STCC), most taxa showed lower $\delta^{15}\text{N}$ values similar to POM as influenced by the presence of *Trichodesmium*. Therefore, our results indicate there are food webs based on a mixture between POM and *Trichodesmium*, that form two kinds of sources of food webs in the southern WSP. Carlotti et al. (2018) estimated that mesozooplankton remove daily 1.5% to 22% of N_2 stocks fixed by phytoplankton in the western tropical South Pacific along 20°S, suggesting considerable variability is present, as we observed in our study.

4.5. Linkage of leptocephali and POM isotopic signatures within the WSP food webs

The analyses of the stable isotopic signatures of the 6 major taxa of leptocephali in relation POM and other food web components the 3 current zones showed some interesting trends not observed in previous studies on fewer taxa or in more limited geographic areas. Within each current zone of the WSP, the signatures showed similar patterns for the two groups of taxa of leptocephali (as designated by Feunteun et al., 2015), with Group 1 (high $\delta^{15}\text{N}$, low $\delta^{13}\text{C}$; Congridae, Muraenidae, Serrivomeridae) and Group 2 (low $\delta^{15}\text{N}$, $\delta^{13}\text{C}$ high; species that grow to large larval sizes: *Avocettina*, *Nemichthys* and *Ariosoma*) having similar relationships within each current zone. This relationship between the 2 groups is essentially the same as what was observed in leptocephali collected in a tight grid of stations located on the downstream side of the Mascarene Plateau where the flow of the Indian Ocean SEC passed through the shallow banks where some eels were spawning and other eels were spawning offshore (Feunteun et al., 2015; Miller et al., 2015, 2019b). In addition, an *Ariosoma* species (Group 2) also showed clearly lower $\delta^{15}\text{N}$ values compared to *Anguilla* (Group 1) in the North Equatorial Current of the western Pacific (Miyazaki et al., 2011). The same clear relationship

between many Group 1 taxa of leptocephali and *Ariosoma* and *Nemichthys* (Group 2) was also seen in a larger study of the isotopic signatures of leptocephali from the NEC in November 2013 (Onda, 2017; Onda et al., 2017), and it was observed in the Gulf of Mexico, along with a few outlier values (Quattrini et al., 2019). These differences do not appear to be related to the actual size of the larvae being isotopically analyzed in our study (not shown) or in a previous study that examined isotopic signatures in relation to larval size (Feunteun et al., 2015). This suggests these groups may constitute a typical pattern of isotopic signatures among leptocephali from oligotrophic ocean gyres, despite the fact that the long larval duration of leptocephali allows for the mixing of larvae from different geographic origins.

Our leptocephali data had some outliers from the main range of signatures in each zone, which were very large larvae (*Avocettina*, *Nemichthys*, *Bathycongrus*) that would have been drifting for many months. Considering the geographic variation in $\delta^{15}\text{N}$ values shown in our study and by Houssard et al. (2017) in the WSP, these large larvae may have come from areas with different signatures than the zones they were collected in. Another interesting finding was that all the leptocephali and other taxa at St. 65 within the SEC zone had higher $\delta^{15}\text{N}$ values more like the signatures in the SECC. It is unclear if the water in that area had been transported there from the north, but the POM in the upper sampling layers had higher signatures than at the stations to the east. The high $\delta^{15}\text{N}$ signatures at that station added some variability into our comparisons of the various taxa among the 3 zones, but the general position of leptocephali in the plots of average $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ signatures of all taxa and POM were similar within each current zone.

Our distribution of stations was not perfect for examining the latitudinal variation in isotopic signatures due to several stations being very close to the zone boundaries, the presence of east-west variations in the POM signatures, and some taxa such as Serrivomeridae were more abundant in the north than the south (see Liénart et al., 2016). However, there was an overall trend of $\delta^{15}\text{N}$ signatures decreasing from north to south in all the taxa examined, and the leptocephali generally followed this trend. The signatures of leptocephali were therefore linked with the overall food web.

Interestingly, understanding how leptocephali are positioned within the food web needs to be approached differently than for most other marine animals whose feeding niches are more well defined, because of their unique feeding ecology. Direct observations of the gut contents of leptocephali (Otake et al., 1993; Mochioka and Iwamizu, 1996; Miller et al., 2011, 2019; Tomoda et al., 2018) and the diversity of taxa detected using DNA sequencing of anguillid leptocephali gut contents (Riemann et al., 2010; Ayala et al., 2018) have shown that

leptocephali consume marine snow materials. The guts of leptocephali are usually partially or mostly empty, probably due to their soft bodies and the capture process, but when food is present it appears as amorphous materials or includes detrital-like marine snow or roundish objects (Mochioka and Iwamizu, 1996; Miller et al., 2011; Tomoda et al., 2018). A study of leptocephali gut contents in the Sargasso Sea suggests that some of these roundish objects are thraustochytrid protists (Miller et al., 2019a). These protists and other groups of the Labyrinthulomycetes cannot be detected genetically unless appropriate PCR primers are used (e.g. Li et al., 2013), so they were not detected in the gut contents of leptocephali in the Sargasso Sea (Riemann et al., 2010; Ayala et al., 2018). They have been found within marine snow aggregates (Li et al., 2013; Bochdansky et al., 2017), so they may be a previously unrecognized component of some types of marine snow that is consumed by leptocephali.

Feeding on marine snow, therefore makes the diet of leptocephali difficult to define clearly, because marine snow is a broad category of detrital materials that originates from a wide range of sources and is colonized by many types of small organisms (Alldredge and Silver, 1988; Shanks and Walters, 1997; Kiørboe, 2000; Bochdansky et al., 2017) including bacteria (Pomeroy and Deibel, 1980) and can contain a wide diversity of DNA sequences (Ayala et al., 2018). In addition, some abundant marine organisms, such as protists will require new techniques to be detected. Consequently, the content of marine snow particles will directly depend on the activity within the food web at each location and on what material is being released at each depth. Further complicating the question of what each type of materials leptocephali feed on is that they can ingest distinguishable objects that contribute to marine snow such as appendicularian houses (Mochioka and Iwamizu, 1996; Miller et al., 2011), which are discarded frequently after their filters clog up with materials present in the water that is being filtered to obtain food. Marine snow also contains carbohydrates that originate from transparent exoparticles (TEP) that are released from phytoplankton or bacteria (Alldredge et al., 1993; Passow, 2002; Engel, 2004). In addition, the widely abundant small calycophoran siphonophores are seemingly contributing some kind of material to what is consumed by leptocephali, at least in the Sargasso Sea of the North Atlantic (Ayala et al., 2018). This means that different sizes or types of marine snow can include materials ranging from being directly from primary producer phytoplankton (TEP), or being from secondary consumers (siphonophores), or from a variety of sources aggregated into individual marine snow particles, which can be diverse in their content in oligotrophic areas (Ayala et al., 2018).

Considering these factors, perhaps it is not surprising that different isotopic signatures were observed among leptocephali. Despite the long larval life of leptocephali and the

likelihood of some larvae from different geographic areas mixing together, the two isotopic groups of leptocephali appears to be a consistent pattern observed in each study that has included those taxa (Miyazaki et al., 2011; Feunteun et al., 2015; Onda, 2017; Quattrini et al., 2019). Because Group 2 consists of the species that grow to much larger sizes (*Avocettina*, *Nemichthys* and *Ariosoma*) than all other anguilliform leptocephali, it cannot be assumed there are not some kind of physiological differences in isotopic fractionation between the two groups. Isotopic fractionation is a poorly understood process that might change according to species, tissues, or trophic levels. Organisms that have different assimilation rates or also different tissue turnover rates depending on the type or organisms and physiological status/growth can differ in their trophic level fractionation (Ambrose and Norr, 1993), and different degrees of fractionation of nitrogen isotopes have been detected along trophic linkages (Gannes et al., 1998; McCutchan et al., 2003). Furthermore, individuals belonging to the same family with the same diet, have shown differences reaching more than 2‰ for $\delta^{13}\text{C}$ isotopic signatures (DeNiro and Epstein, 1978; Voigt et al., 2003). It therefore seems possible that some of the differences between the two groups is resulting from physiological differences related to growing to larger sizes.

If this is not the case, then an obvious alternative explanation clearly suggested by the present study is that the two groups may be feeding at different depths where the marine snow has different isotopic signatures. Serrivomeridae followed by Congridae generally had the highest $\delta^{15}\text{N}$ levels, followed by Muraenidae that had more variable levels, so these species may tend to feed at deeper depths than *Avocettina*, *Nemichthys* and *Ariosoma*, which could feed more at shallower depths. This is consistent with the low POM $\delta^{15}\text{N}$ signatures at the surface or within the fluorescence maximum, but the high $\delta^{13}\text{C}$ signatures of the Group 2 larvae suggests they may feed closer to the surface where our POM had clearly higher $\delta^{13}\text{C}$ values. Leptocephali are known to sometimes vertically migrate (Castonguay and McCleave, 1987) like other planktonic organisms (Hays, 2003; Tarling, 2003), and they are usually found in the upper 100 m at night (Castonguay and McCleave, 1987; Otake et al., 1998; Miller, 2009, 2015; Onda et al., 2017). Different taxa of leptocephali have also been found at different depth zones within the upper 100 m at night including at individual stations (Miller, 2009, 2015; Onda et al., 2017). Information on the depths used by leptocephali during the day, when they are likely feeding is more limited however due to possible net avoidance (see Miller et al., 2013a). In the Sargasso Sea, anguillid, muraenid, chlopsid and serrivomerid leptocephali (Group 1), seemed to move from < 100 m depths during nighttime to depths mostly < 250 m during the day (Castonguay and McCleave, 1987) where there can be high

$\delta^{15}\text{N}$ POM signatures (our study; Miyazaki et al., 2011; Hannides et al., 2013; Williams et al., 2014; Feunteun et al., 2015). In contrast that study found no evidence of Group 2 species vertically migrating to deeper depths during the day, which is consistent with the possibility that they feed at shallower depths compared to Group 1 species.

While the general relationships of higher or lower values of leptocephali and POM from different depths compared to each other show possible correspondences, the gaps between the two do not seem to correspond to traditional fractionation values (DeNiro and Epstein, 1978; Rau and Anderson, 1981; Fry and Arnold, 1982; Peterson and Fry, 1987; Post, 2002). This is typical of the isotopic studies of leptocephali, zooplankton, and POM (Miyazaki et al., 2011; Feunteun et al., 2015; Dupuy et al., 2016; Lienart et al. 2016). Those studies, the present study, and typical studies including POM (e.g. Dupuy et al., 2016) use a $0.7\ \mu\text{m}$ filter pore size, but the marine snow particles that leptocephali can feed on are much larger than that (more on the scale of millimeters). This means that the POM that is analyzed includes free-living plankton taxa, which are not included in the size of marine snow that leptocephali feed on. Waite et al. (2007) showed that $\text{POM} > 20\ \mu\text{m}$ was both about 2‰ $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ higher than $\text{POM} < 5\ \mu\text{m}$, so size of the POM particles consumed by leptocephali is likely an important factor. Therefore, there is no clear expectation that the signatures of leptocephali (feeding on marine snow) and POM (marine snow and many other materials $\geq 0.7\ \mu\text{m}$) should match precisely according to normal fractionation values, because as pointed out previously, leptocephali are likely selecting some components of the total POM samples (Feunteun et al., 2015; Liénart et al., 2016). In the future, an Underwater Vision Profiler coupled with a “Appinet” or marine snow catcher could be used to gather in situ diversity and abundance information about marine snow particles as was done by Duret et al. (2019) and Lundgreen et al. (2019).

4.6. Conclusions

The region of the WSP that we studied during February 2013 varied hydrographically and biologically with latitude, and different isotopic compositions and microbial compositions were found at the three depths we examined. Heterotrophic picoplankton was dominant at all three depths and phytoplankton (mainly small size) and heterotrophic nanoflagellates were mostly present within the maximum fluorescence layer. The pico-nano-microplankton was both N and P limited within the maximum fluorescence layer and the 3 currents zones were based on a heterotrophic planktonic functioning. All of our observations appeared to be

consistent with typical characteristics of oligotrophic subtropical gyres; however, this particular region is interesting because of the alternating east and west current zones as well as other smaller current jets, which likely result in mixing of water masses. Generally, consistent patterns of the isotopic signatures of POM, leptocephali, zooplankton and micronekton were observed, with the baseline $\delta^{15}\text{N}$ signatures of all components generally decreasing from north to south, with some east-west variation also seen in POM values. Furthermore, the presence of *Trichodesmium* in the STCC surface zone would influence the isotopic signatures of POM and consequently leptocephali. The signatures of leptocephali were therefore linked to the overall food web because all evidence indicates that they feed on the larger size range of the marine snow. The apparent two groups of leptocephali with different isotopic signatures may be explained by feeding behavior at different depths or on different types of marine snow. Further studies need to examine marine snow particles at different depths directly to determine what types of particles each type and size of leptocephali may feed on to begin to fully understand the feeding ecology of leptocephali in the WSP and each region of the world. In addition, more research is needed using stable isotope analysis of leptocephali and other food web components, such as small mesozooplankton (< 0.5 mm) that were not sampled in this study, across a wider horizontal scale to gain a better understanding of the feeding ecology of this interesting type of fish larvae.

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Author contributions

C.D., A.C., E.F., A.C., and CLi. collected the specimens and data during the sampling survey that was designed and managed by T.O., K.T., J.A., and S.W. The leptocephali were identified on board by M.J.M. and M.K., the specimen data was analyzed by L.G., C.D., E.F., CLi., and Cle., and L.G. and M.J.M. made the figures. C.D., L.G., M.J.M., and E.F. drafted the manuscript and all coauthors approved the manuscript

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Journal Pre-proofs

Table 1. Measured values of nutrients ($\mu\text{g L}^{-1}$) within the maximum fluorescence depth layer at individual stations within the South Equatorial Countercurrent (SECC), South Equatorial Current (SEC), and South Tropical Countercurrent (STCC) current zones shown in Figure 1. The geographic location of some stations is indicated as northeast (NE), southwest (SW) etc.

Currents/ Stations	NO ₃	NO ₂	PO ₄	Si(OH) ₄	NH ₄
SECC					
St 44	127.8	19.1	65.1	137.5	19.3
St 47	155.3	11.0	77.8	109.0	14.3
St 63 (NE)	4.7	0.0	27.1	91.9	10.4
Mean±S.D.	95.9±80.2	10.1±9.6	56.7±26.4	112.8±23.1	14.7±4.5
SEC					
St 39	28.3	4.7	44.8	512.6	12.9
St 41	45.3	2.4	44.9	76.0	20.6
St 49	3.5	0.3	29.3	67.9	9.5
St 51	11.7	2.8	42.5	88.4	19.4
St 60 (E)	9.7	1.9	34.9	63.9	25.7
Mean±S.D.	19.7±17.0	2.4±1.6	39.3±6.9	161.8±196.3	17.6±6.4
STCC					
St 36 (SW)	15.3	2.4	31.1	99.9	15.3
St 55	21.2	2.9	33.2	61.2	17.3
St 56 (SE)	29.0	7.5	31.5	70.2	11.7
Mean±S.D.	21.8±6.9	4.3±2.8	31.9±1.1	77.1±20.2	14.8±2.9

Figures

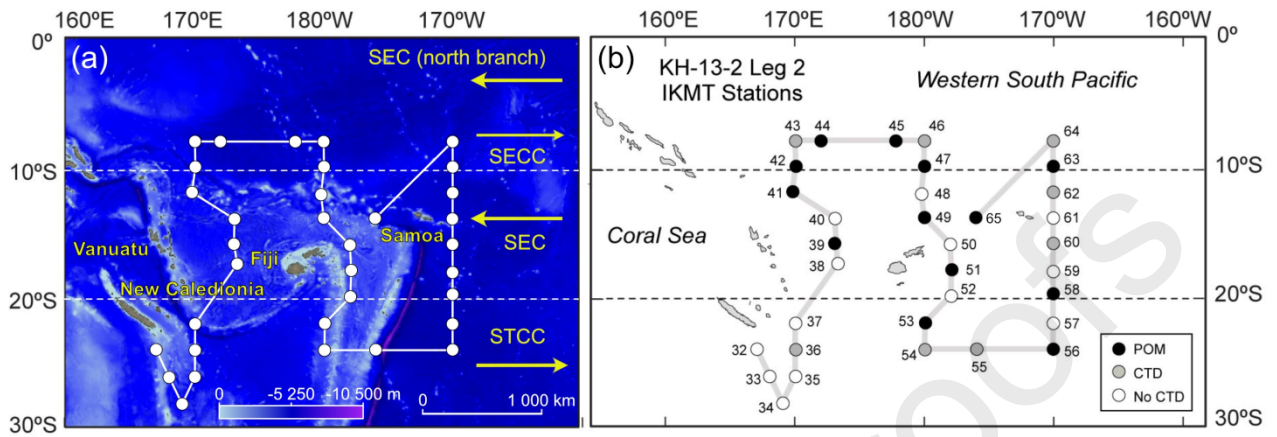


Figure 1. Maps of the KH-13-2 sampling survey of the R/V Hakuho Maru in February 2013 in relation to bathymetry and major current systems (a) and showing the IKMT stations for collecting leptocephali and other organisms, the station numbers (St. 32–65), CTD stations where particulate organic matter (POM) was collected (dark circles), and where other CTD casts made (grey circles) (b). Water samples for microbial loop functioning experiments were collected at stations 39, 41, 47, 49, 51 and 55. Darker colors indicate deeper areas (mid-range blues about 2000-3000 m, dark blues about 4000-5000 m) and lighter colors shallow areas (white about 500 m or less) in (a).

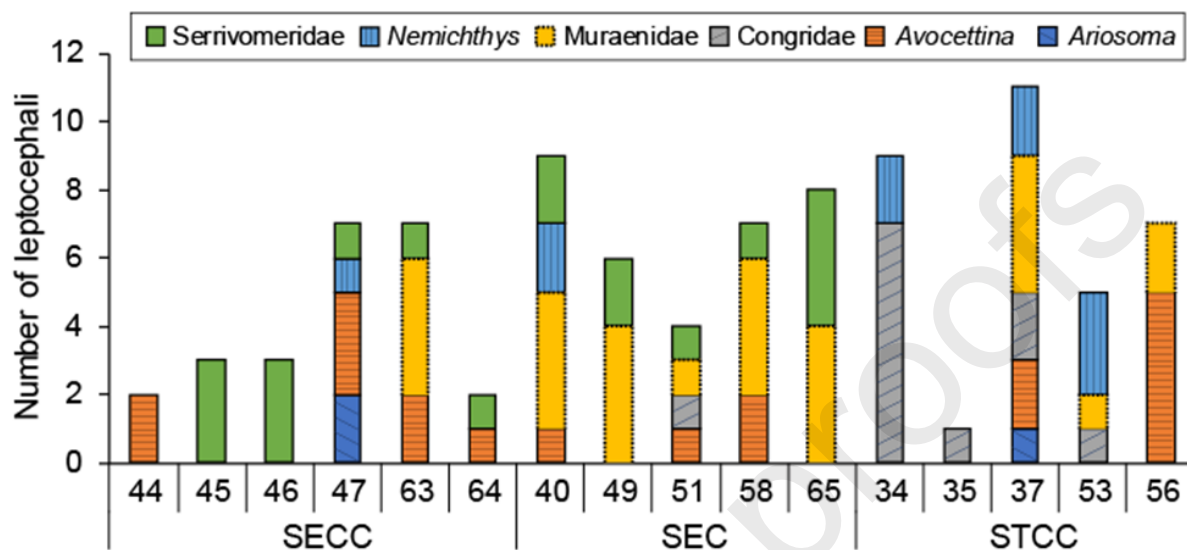


Figure 2. Number of 6 taxa of leptocephali that were analyzed for their isotopic signatures during the R/V Hakuho Maru cruise at each station within the 3 current zones of South Equatorial Countercurrent (SECC), South Equatorial Current (SEC), and South Tropical Countercurrent (STCC).

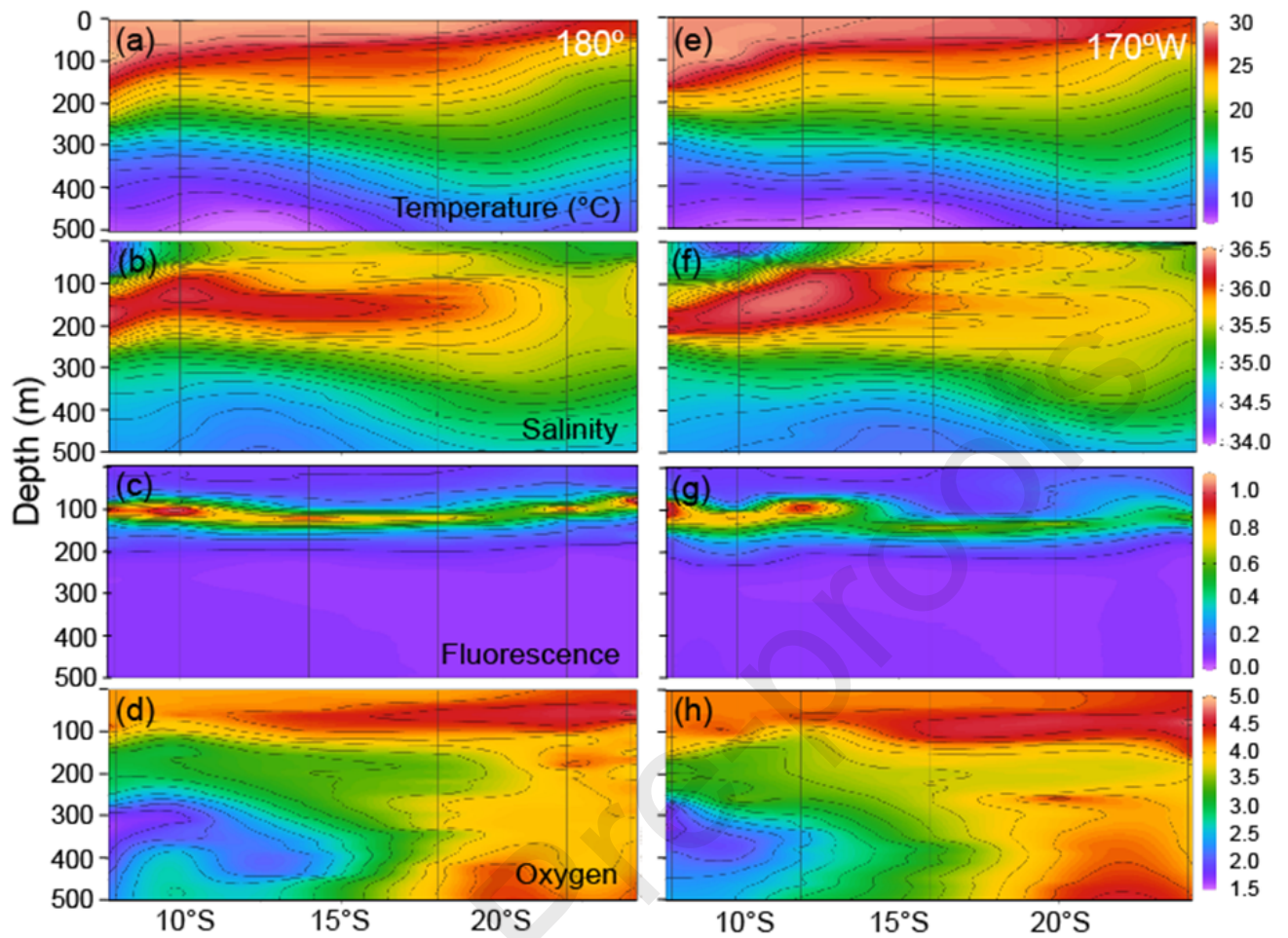


Figure 3. Hydrographic sections plotted from the CTD data from stations along or near 180° (middle transect) and 170°W (eastern transect) for different parameters measured in the upper 500 m.

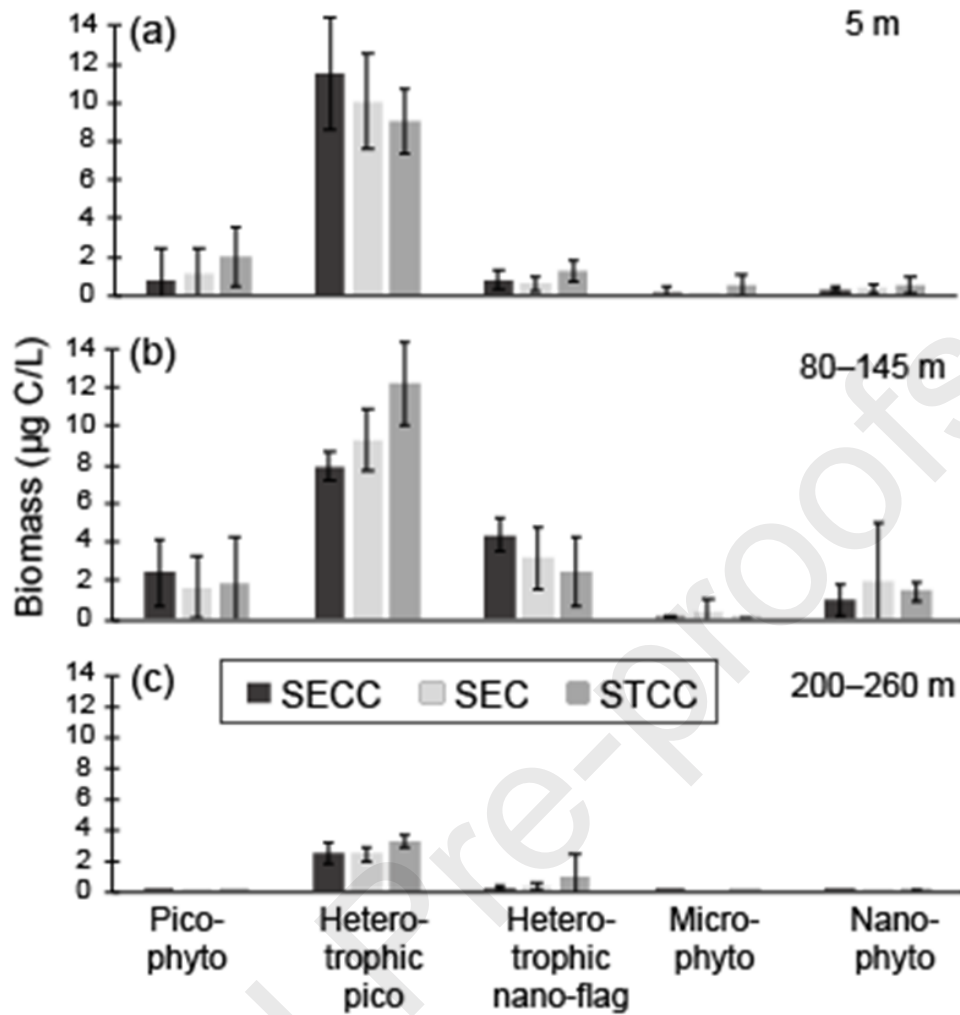


Figure 4. Biomass ($\mu\text{g C. L}^{-1} \pm \text{SD}$) of the major groups and size categories of planktonic organisms contributing to POM at 3 depths in the 3 current zones of the South Equatorial Countercurrent (SECC), South Equatorial Current (SEC), and South Tropical Countercurrent (STCC).

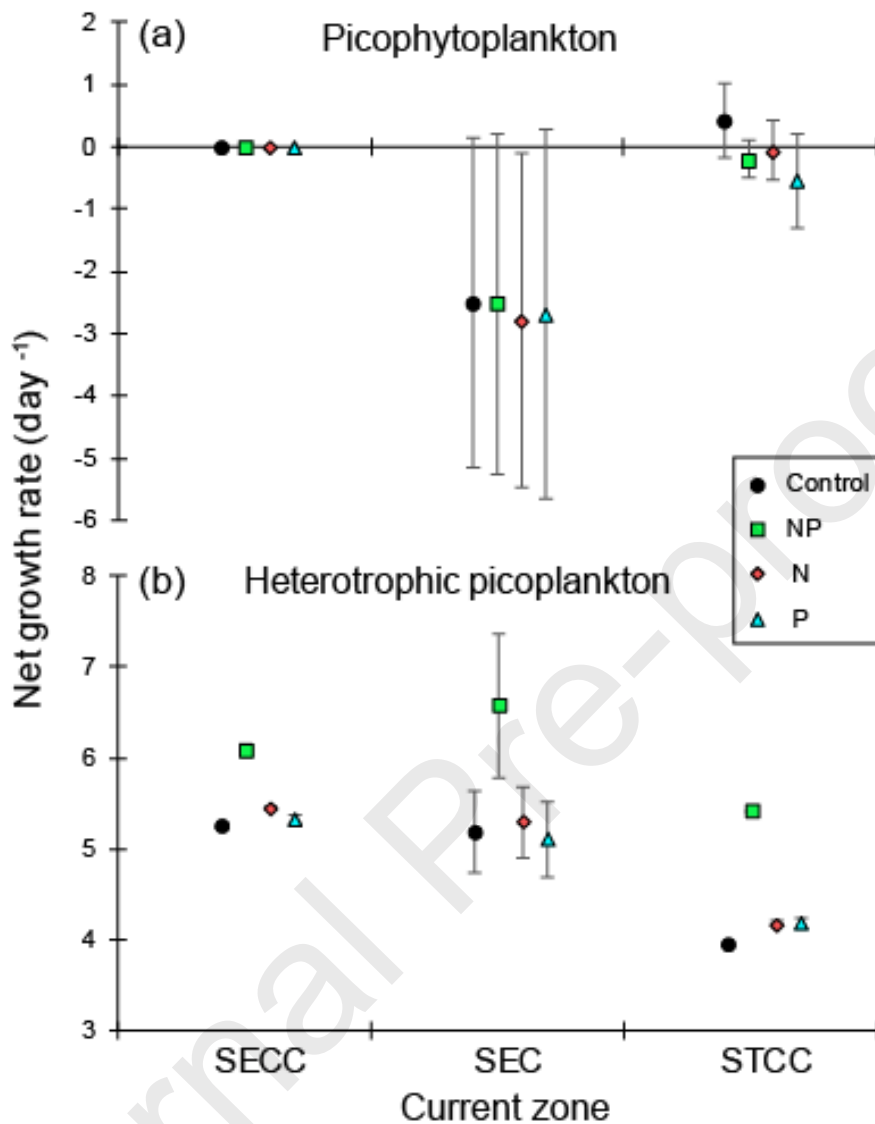


Figure 5. Net growth rate (day⁻¹ ± SD) of (a) picophytoplankton and (b) heterotrophic picoplankton with the addition of inorganic nitrogen (N), inorganic phosphorus (P) and a combination of nitrogen and phosphorus (NP) and a control (no additions) using water from the 3 current zones of the South Equatorial Countercurrent (SECC), South Equatorial Current (SEC), and South Tropical Countercurrent (STCC) at the maximum fluorescence depth.

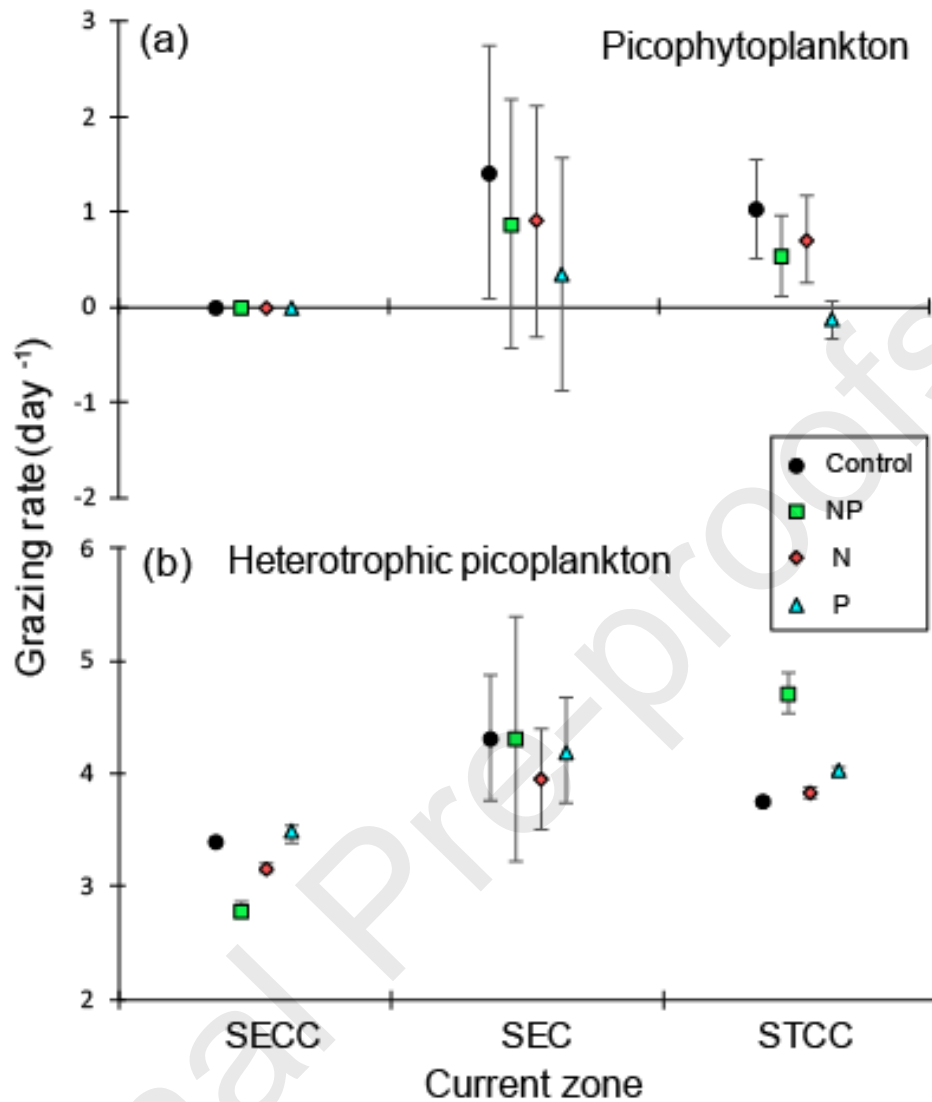


Figure 6. Grazing rate ($\text{day}^{-1} \pm \text{SD}$) of (a) picophytoplankton and (b) heterotrophic picoplankton with the addition of inorganic nitrogen (N), inorganic phosphorus (P) and a combination of nitrogen and phosphorus (NP) and a control (no additions) using water from the 3 current zones of the South Equatorial Countercurrent (SECC), South Equatorial Current (SEC), and South Tropical Countercurrent (STCC) at the maximum fluorescence depth.

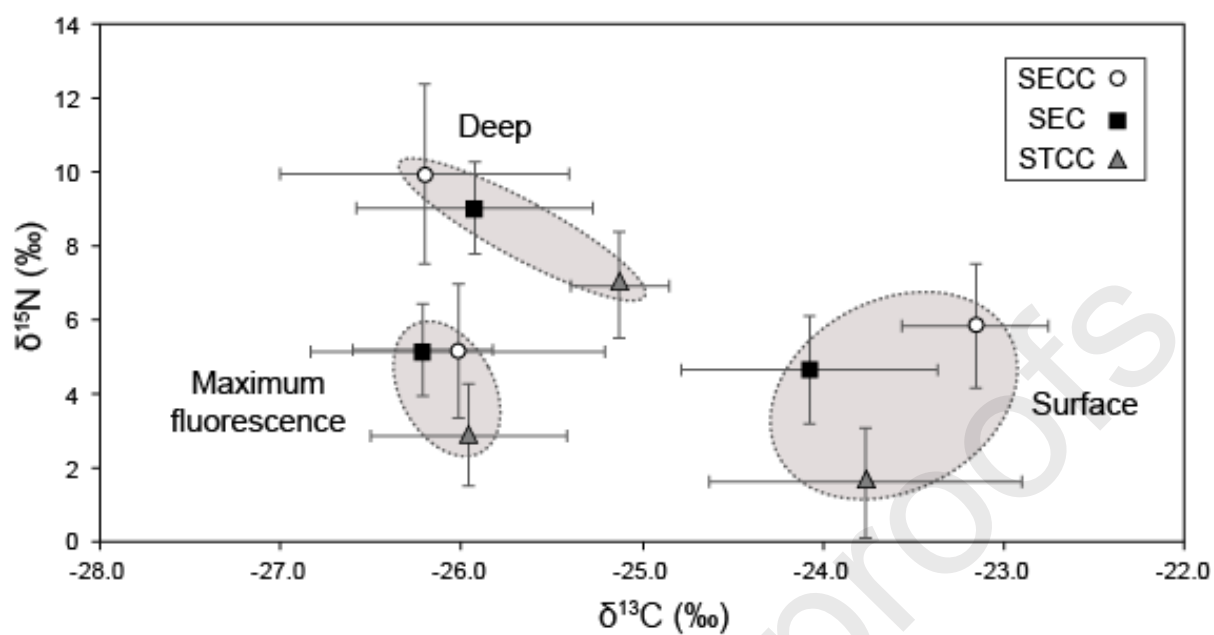


Figure 7. Mean carbon and nitrogen stable isotope values ($\text{‰} \pm \text{SD}$) of POM at different depths (depending on stations) from the 3 current zones of the South Equatorial Countercurrent (SECC), South Equatorial Current (SEC), and South Tropical Countercurrent (STCC). Grey shaded ovals highlight the distribution of mean values at the 3 POM sampling depths.

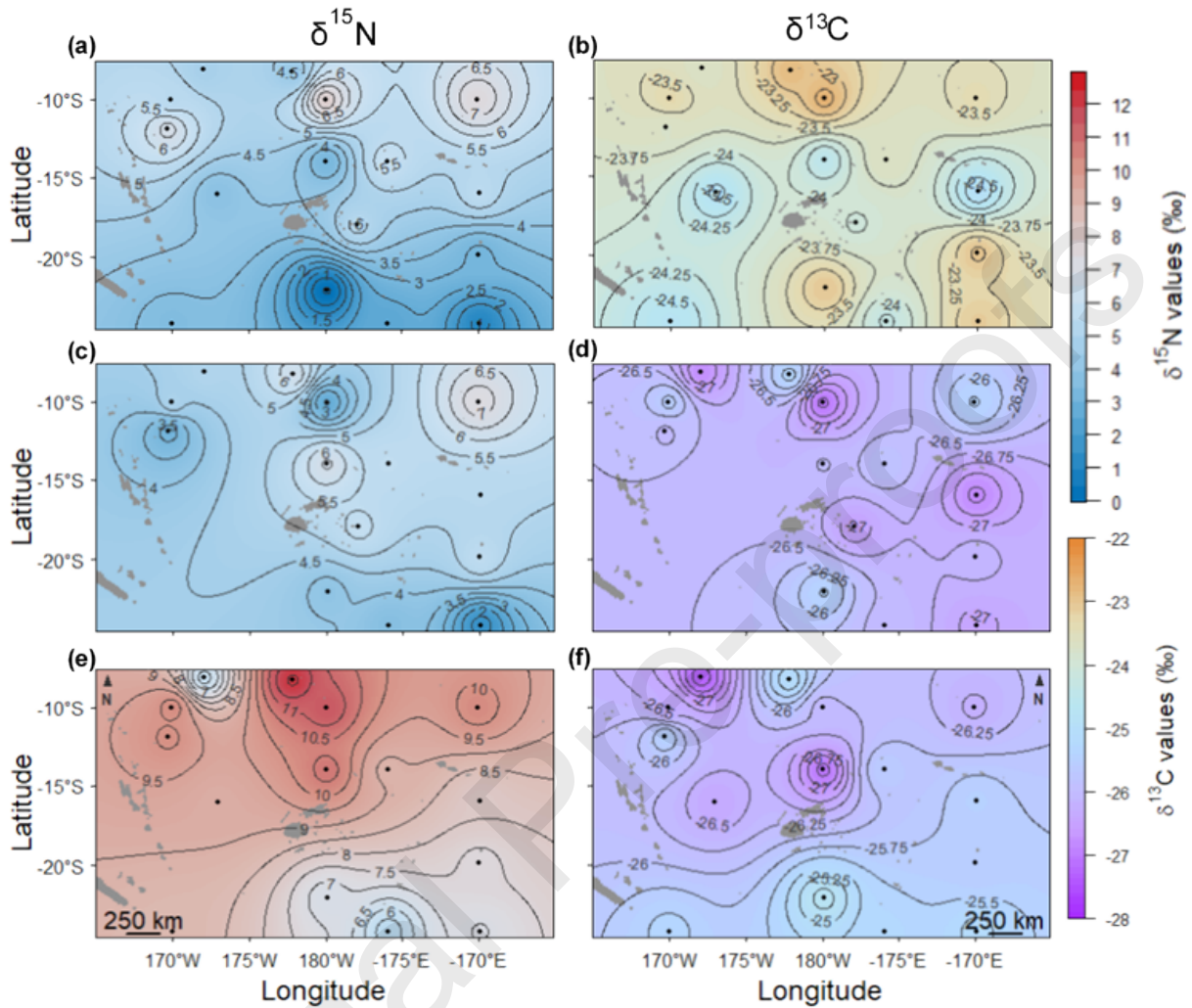


Figure 8. Contour plots of $\delta^{15}\text{N}$ (left panels) and $\delta^{13}\text{C}$ (right panels) isotope values (Inverse Distance Weighted Interpolation, distance coefficient = 2) of POM at the surface (5 m, **a** and **b**), maximum fluorescence (80 – 145 m, **c** and **d**) and at the bottom (200 – 260, **e** and **f**) in the western South Pacific Ocean during the KH-13-2 cruise (February 2013, southern summer). The dots represent POM sampling stations, blue or purple colors show lower isotopic signatures than red or orange colors for $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$, respectively. The observed $\delta^{15}\text{N}$ values range from 0 to 12.1‰ and from -27.3 to -22.7‰ for $\delta^{13}\text{C}$ values.

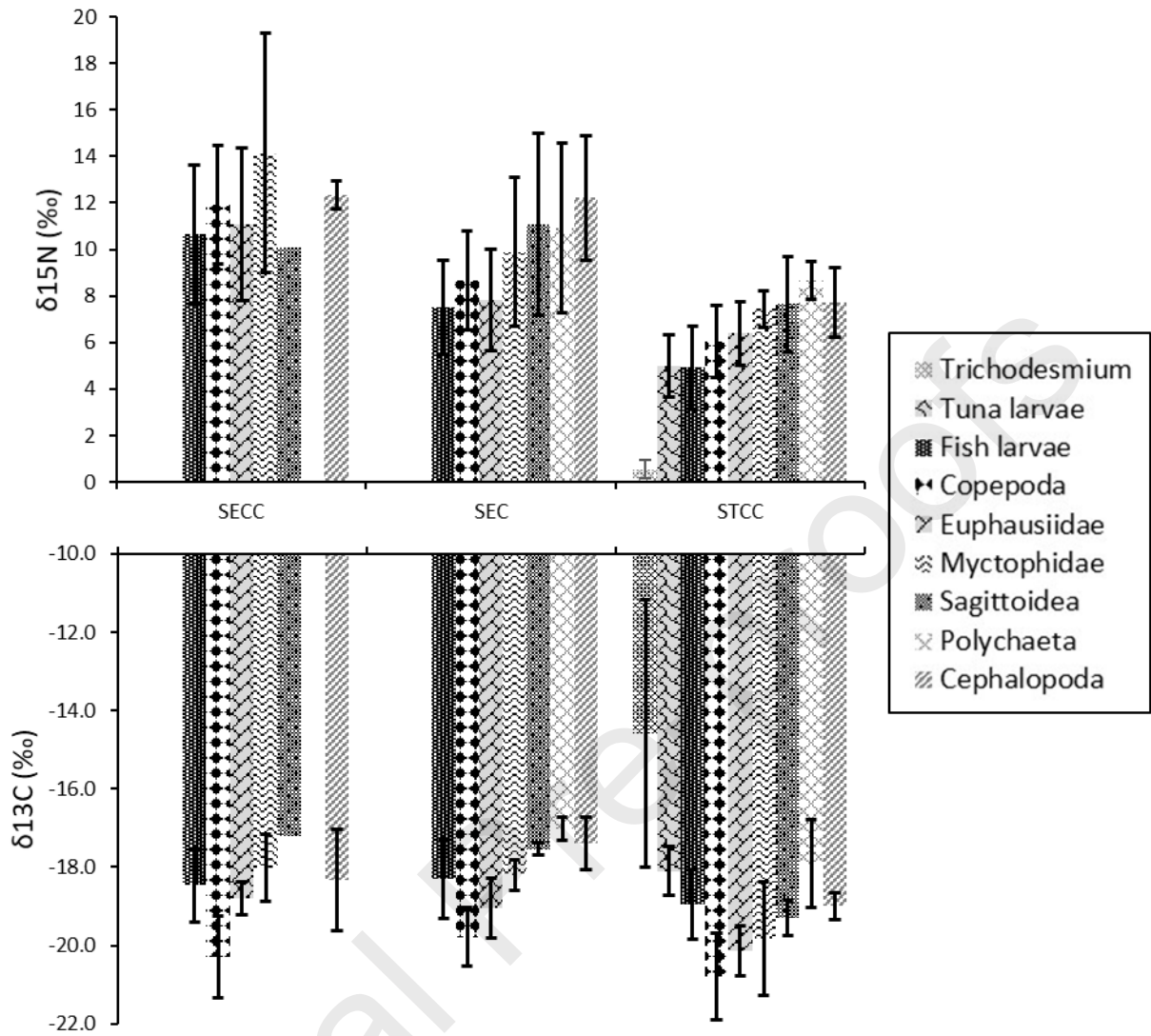


Figure 9. Mean carbon and nitrogen stable isotope values ($\text{‰} \pm \text{SD}$) of different taxa of meso- and macro-zooplankton, micronekton and the cyanobacterium *Trichodesmium*.

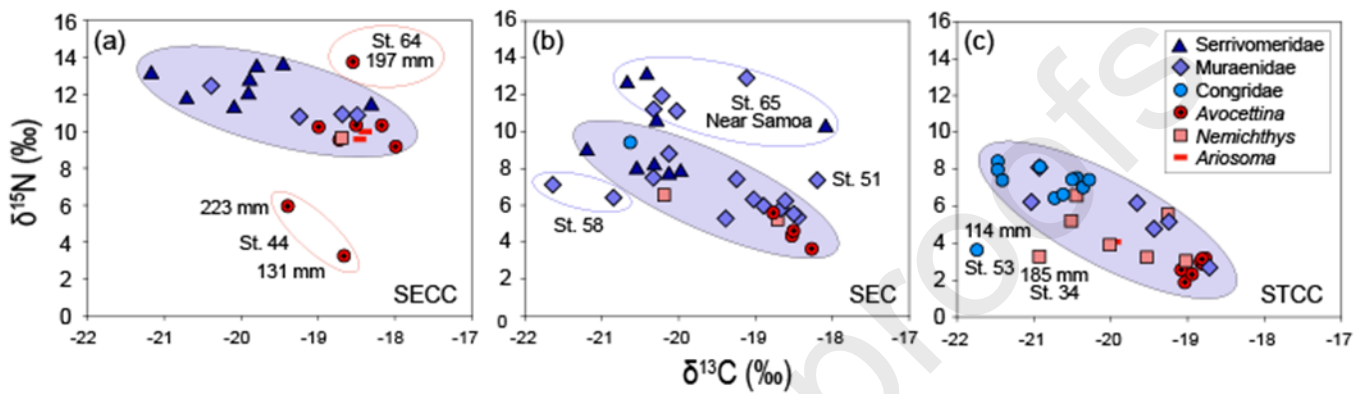


Figure 10. Mean carbon and nitrogen stable isotope values ($\text{‰} \pm \text{SD}$) of different taxa of leptocephali from the 3 current zones of the South Equatorial Countercurrent (SECC), South Equatorial Current (SEC), and South Tropical Countercurrent (STCC). Shaded ovals highlight the main group of signatures for comparison among current zones and some larvae from individual stations are indicated by open ovals. The points of Group 1 leptocephali are shown in blue shades and Group 2 with red shades.

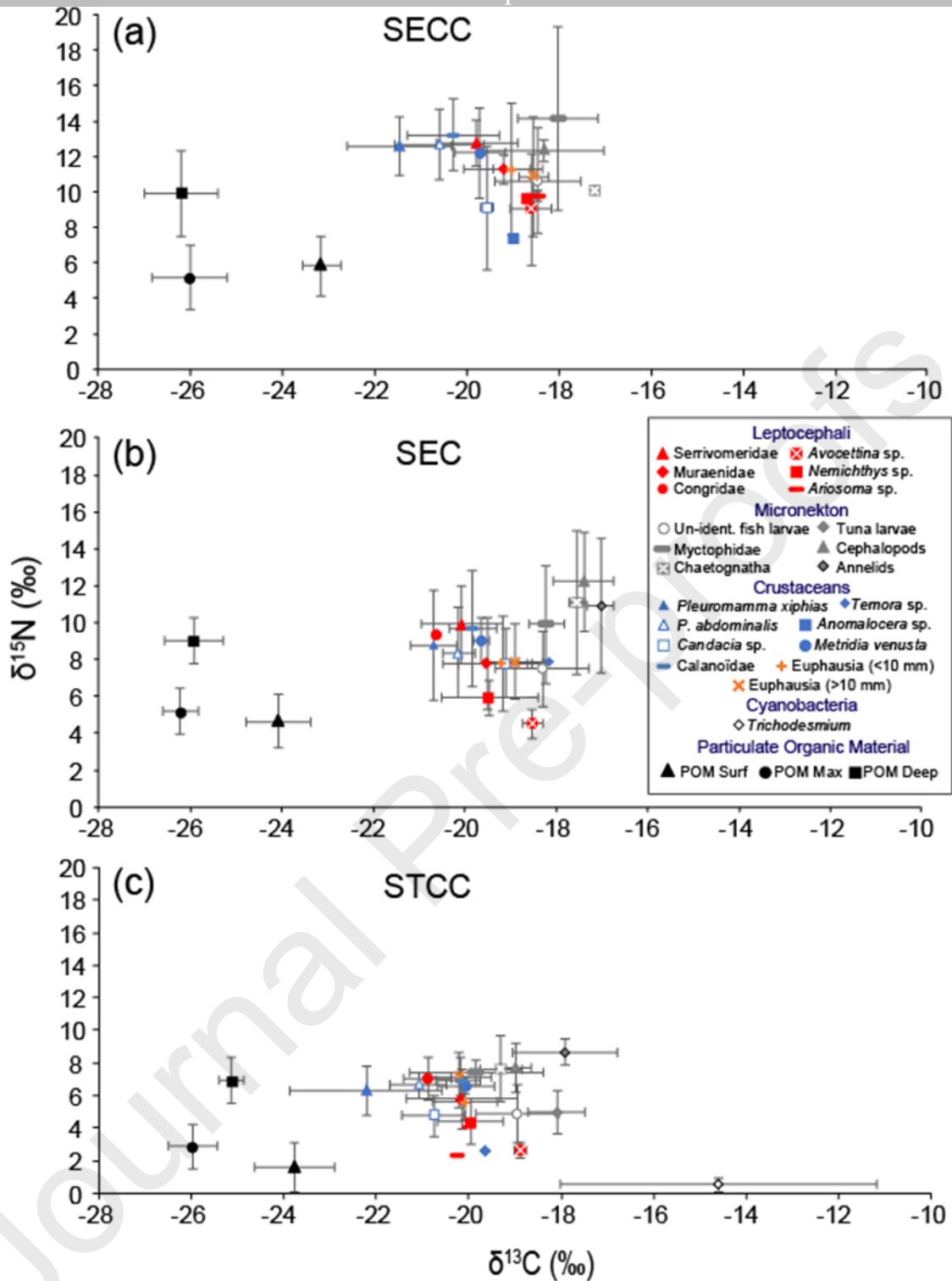


Figure 11. Mean carbon and nitrogen stable isotope values ($\text{‰} \pm \text{SD}$) of different taxa of leptocephali (red), meso- and macro-zooplankton, micronekton and POM at different depths from the 3 current zones of the South Equatorial Countercurrent (SECC), South Equatorial Current (SEC), and South Tropical Countercurrent (STCC).

Supplementary table. Values of carbon and nitrogen stable isotope values ($\text{‰} \pm \text{SD}$) of different taxa of leptocephali, meso- and macro-zooplankton, micronekton and POM at different depths within the 3 current zones of the South Equatorial Countercurrent (SECC), South Equatorial Current (SEC), and South Tropical Countercurrent (STCC).

Taxonomic groups	SECC					SEC					STCC					
	N	$\delta^{15}\text{N}$ mean \pm S.D	$\delta^{15}\text{N}$ range	$\delta^{13}\text{C}$ mean \pm S.D	$\delta^{13}\text{C}$ range	N	$\delta^{15}\text{N}$ mean \pm S.D	$\delta^{15}\text{N}$ range	$\delta^{13}\text{C}$ mean \pm S.D	$\delta^{13}\text{C}$ range	N	$\delta^{15}\text{N}$ mean \pm S.D	$\delta^{15}\text{N}$ range	$\delta^{13}\text{C}$ mean \pm S.D	$\delta^{13}\text{C}$ range	
Leptocephali																
Nemichthyidae	<i>Nemichthys</i>	1	9.6	-	-18.7	-	2	5.9 \pm 0.9	5.2–6.6	-19.5 \pm 1.0	-20.2–-18.7	7	4.4 \pm 1.4	3.0–6.6	-20.0 \pm 0.7	-20.9–-19.0
	<i>Avocettina</i>	8	9.0 \pm 3.2	3.2–13.7	-18.6 \pm 0.4	-19.4–-18.0	4	4.5 \pm 0.8	3.6–5.6	-18.5 \pm 0.2	-18.8–-18.3	7	2.7 \pm 0.5	1.8–3.1	-18.9 \pm 0.1	-19.1–-18.8
Serrivomeridae		9	12.7 \pm 1.3	11.2–15.3	-19.8 \pm 0.9	-21.2–-18.3	10	9.9 \pm 2.1	7.7–13.1	-20.1 \pm 0.9	-21.2–-18.1	0	-	-	-	-
Muraenidae		4	11.3 \pm 0.8	10.8–12.5	-19.2 \pm 0.9	-20.4–-18.5	17	7.8 \pm 2.5	5.3–12.9	-19.5 \pm 1.0	-21.6–-18.2	7	5.8 \pm 1.8	2.7–8.1	-20.2 \pm 1.2	-22.0–-18.7
Congridae	<i>Conger</i>	0	-	-	-	-	1	9.4	-	-20.6	-	6	7.4 \pm 0.6	6.6–8.1	-20.7 \pm 0.4	-21.5–-20.3
	<i>Gnathophis</i>	0	-	-	-	-	0	-	-	-	-	2	7.9 \pm 0.7	7.4–8.4	-21.4 \pm 0.0	-21.5–-21.4
	<i>Gorgasia</i>	0	-	-	-	-	0	-	-	-	-	2	6.9 \pm 0.7	6.4–7.4	-20.6 \pm 0.2	-20.7–-20.5
	<i>Ariosoma sp.</i>	2	9.8 \pm 0.3	9.6–10.0	-18.5 \pm 0.1	-18.5–-18.4	0	-	-	-	-	1	4.1	-	-20.0	-
	<i>Bathycongrus</i>	0	-	-	-	-	0	-	-	-	-	1	3.6	-	-21.7	-
Micronekton																
	Polychaeta	0	-	-	-	-	2	10.9 \pm 3.6	8.4–13.5	-17.0 \pm 0.3	-17.2–-16.8	3	8.7 \pm 0.8	8.1–9.6	-17.9 \pm 1.1	-19.2–-17.0
	Cephalopoda	2	12.4 \pm 0.6	11.9–12.8	-18.3 \pm 1.3	-19.2–-17.4	2	12.2 \pm 2.7	10.3–14.1	-17.4 \pm 0.7	-17.9–-16.9	4	7.7 \pm 1.5	6.3–9.7	-19.0 \pm 0.3	-19.2–-18.5
	Sagittoidea	1	10.1 \pm -	-	-17.2 \pm -	-	3	11.1 \pm 4.0	8.3–15.6	-17.5 \pm 0.2	-17.6–-17.4	3	7.6 \pm 2.0	5.9–9.9	-19.3 \pm 0.5	-19.8–-19.0
	Myctophidae	4	14.1 \pm 5.1	8.5–20.8	-18.0 \pm 0.9	-18.7–-16.8	4	9.9 \pm 3.2	7.0–12.8	-18.2 \pm 0.4	-18.6–-17.7	4	7.4 \pm 0.8	6.5–8.2	-19.8 \pm 1.5	-22.0–-18.7
	Tuna larvae	0	-	-	-	-	0	-	-	-	-	3	5.0 \pm 1.3	3.7–6.3	-18.1 \pm 0.6	-18.8–-17.6
	Fish larvae	3	10.6 \pm 3.0	7.2–12.6	-18.5 \pm 0.9	-19.5–-17.6	7	7.5 \pm 2.0	5.2–10.8	-18.3 \pm 1.0	-20.5–-17.4	6	4.9 \pm 1.8	2.7–7.2	-19.0 \pm 0.9	-20.7–-18.3
Crustaceans																
	<i>Metridia venusta</i>	3	12.2 \pm 2.5	9.5–14.5	-19.7 \pm 0.5	-20.3–-19.3	3	9.0 \pm 1.2	8.0–10.4	-19.6 \pm 0.2	-19.8–-19.5	3	6.6 \pm 0.5	6.1–7.1	-20.1 \pm 0.6	-20.6–-19.4
	<i>Pleuromamma abdominalis</i>	3	12.7 \pm 2.0	10.4–14.3	-20.6 \pm 1.0	-21.7–-19.8	4	8.4 \pm 2.5	5.7–11.1	-20.1 \pm 0.3	-20.4–-19.8	3	6.6 \pm 0.8	5.9–7.5	-21.1 \pm 0.6	-21.8–-20.7
	<i>Pleuromamma xiphias</i>	3	12.6 \pm 1.7	10.7–13.9	-21.5 \pm 1.1	-22.4–-20.2	3	8.8 \pm 3.0	6.2–12.1	-20.7 \pm 0.5	-21.3–-20.4	3	6.3 \pm 1.5	5.0–7.9	-22.2 \pm 1.6	-23.3–-20.3
Calanoida	<i>Candacia sp.</i>	2	9.1 \pm 3.5	6.7–11.6	-19.6 \pm 0.1	-19.7–-19.5	3	7.8 \pm 1.9	6.1–9.8	-19.1 \pm 0.5	-19.5–-18.6	3	4.8 \pm 1.2	3.8–6.1	-20.7 \pm 0.7	-21.5–-20.3
	<i>Temora sp.</i>	0	-	-	-	-	1	7.9	-	-18.1	-	1	2.7	-	-19.7	-
	<i>Anomalocera sp.</i>	1	7.4	-	-19.0	-	0	-	-	-	-	0	-	-	-	-
	Unidentified Calanoida	4	13.2 \pm 2.0	11.3–15.2	-20.3 \pm 1.0	-21.2–-18.9	3	9.7 \pm 3.2	6.0–12.0	-19.8 \pm 0.5	-20.2–-19.2	3	7.0 \pm 1.7	5.4–8.7	-20.2 \pm 0.5	-20.6–-19.7
	Euphausiidae (>10 mm)	4	10.9 \pm 3.4	8.6–15.9	-18.5 \pm 0.3	-18.8–-18.1	5	7.9 \pm 2.0	5.8–11.3	-18.9 \pm 0.6	-19.8–-18.5	4	7.2 \pm 1.2	6.2–8.8	-20.2 \pm 0.7	-21.1–-19.6
Euphausiacea	Euphausiidae (<10 mm)	4	11.3 \pm 3.7	7.0–15.8	-19.0 \pm 0.4	-19.5–-18.6	5	7.8 \pm 2.6	4.8–10.8	-19.2 \pm 0.9	-20.3–-18.3	4	5.6 \pm 1.1	4.6–6.7	-20.1 \pm 0.7	-21.0–-19.4
Cyanobacteria																
	<i>Trichodesmium sp.</i>	0	-	-	-	-	0	-	-	-	-	2	0.5 \pm 0.4	0.3–0.8	-14.6 \pm 3.4	-17.0–-12.2
POM																
	Surface	5	5.8 \pm 1.7	4.2–7.8	-23.2 \pm 0.4	-23.7–-22.7	7	4.6 \pm 1.4	2.9–6.7	-24.1 \pm 0.7	-24.9–-23.0	4	1.6 \pm 1.5	-0.04–3.3	-23.8 \pm 0.9	-24.7–-23.0
	Maximum fluorescence	5	5.1 \pm 1.8	2.5–7.2	-26.0 \pm 0.8	-27.0–-25.4	6	5.2 \pm 1.3	2.8–6.6	-26.2 \pm 0.4	-26.8–-25.8	3	2.9 \pm 1.4	1.3–3.8	-26.0 \pm 0.5	-26.5–-25.4
	Deep	5	9.9 \pm 2.4	5.8–12.1	-26.2 \pm 0.8	-27.3–-25.1	6	9.0 \pm 1.2	7.4–10.7	-25.9 \pm 0.7	-27.1–-25.5	4	6.9 \pm 1.4	5.3–8.7	-25.1 \pm 0.3	-25.4–-24.7

**Manuscript: Microbial functional structure and stable isotopic variation of leptocephali across three current zones in the western South Pacific
(Revision requested for PROOCE_2019_94)**

Declaration of interests

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

The authors declare the following financial interests/personal relationships which may be considered as potential competing interests:

Christine Dupuy
03/10/2019



Highlights

- Western South Pacific microbial biomass/function and stable isotopes were examined
- Primary producers, whatever the depth, were N-P co-limited
- All zones were based on a heterotrophic planktonic functioning
- Stable isotope signatures of leptocephali, POM and other components of the food web were analyzed
- POM signatures varied with depth and latitude, influencing all taxonomic groups
- The presence of *Trichodesmium* may contribute to low southern $\delta^{15}\text{N}$ values

Journal Pre-proofs