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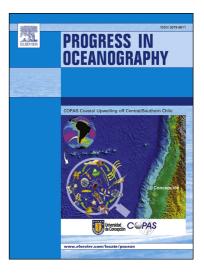
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Trophic niches of sympatric tropical tuna in the Western Indian Ocean inferred by

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2 3 stable isotopes and neutral fatty acids

Fany Sardenne^{1*}, Nathalie Bodin¹, Emmanuel Chassot¹, Aurélien Amiel^{1,2}, Edwin Fouché^{1,2}, 4 Maxime Degroote¹, Stéphanie Hollanda³, Heidi Pethybridge⁴, Benoit Lebreton⁵, Gaël 5 Guillou⁵ & Frédéric Ménard⁶ 6 7 ¹ IRD, UMR 248 MARine Biodiversity Exploitation and Conservation (IRD, Ifremer, UM, 8 CNRS), BP 570, Victoria, SEYCHELLES. 9 ² INRA, UMR 1331 TOXALIM (INRA), Research Centre in Food Toxicology, Toulouse, FRANCE. ³ Seychelles Fishing Authority (SFA), BP 449, Victoria, SEYCHELLES. 10 ⁴ CSIRO Oceans and Atmosphere Flagship, Hobart, AUSTRALIA. 11 ⁵ CNRS-Université de la Rochelle, UMR 7266 Littoral Environnement et Sociétés (LIENSs), 12 La Rochelle, FRANCE. 13 ⁶ IRD, Mediterranean Institute of Oceanography (Aix-Marseille Université, CNRS/INSU, 14 Université de Toulon, IRD), Marseille, FRANCE. 15 16 17 Highlights 18 • Sympatric tropical tuna (bigeye, yellowfin and skipjack tuna) were sampled in the Indian 19 Ocean; 20 • Trophic niche partitioning was quantitatively assessed using time-integrated tracers; 21 • Intra- and inter-species resource overlap in three tropical tuna species was detected; 22 • Diet shifts at ~ 100 cm fork length were detected in the larger tuna species (bigeye and 23 yellowfin); 24 • New insights into the mechanisms involved in tropical tuna coexistence are provided. 25 26 27 28 ^{*}Corresponding author: <u>Fany.Sardenne@hotmail.fr</u> 29 30 31 32

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33 ABSTRACT

34 This study examined the trophic ecology of three sympatric tropical tuna species (bigeye 35 BET, skipjack SKJ, and yellowfin YFT) sampled in the Western Indian Ocean throughout 2013. Specifically we explored inter-specific resource partitioning and ontogenetic variability 36 using neutral fatty acids and stable isotope analysis of liver and muscle from small (≤ 100 cm 37 fork length, FL) and large (>100 cm FL) tuna collected in mixed schools at the surface by 38 purse-seine. Both biochemical tracers were used to calculate trophic niche indices that 39 collectively revealed high potential for resource overlap, especially among small tuna. 40 Resource overlap appeared strongest between BET and YFT, with SKJ tissues having high 41 carbon isotope (δ^{13} C) values (-17±0.3%), lower nitrogen isotope (δ^{15} N) values (11.4±0.6%), 42 and higher relative proportion of poly-unsaturated fatty acids (PUFA) than the two other 43 species, indicating a different diet. Size was found to be a strong predictor for most 44 biochemical tracers in the three species with δ^{13} C, δ^{15} N and total lipid content in the liver. In 45 the larger species (YFT and BET), proportions of mono-unsaturated fatty acids typically 46 increased with size, while quantities of PUFA decreased. In addition to ontogenetic 47 variability, trophic markers were shown to vary between sampling area and season: higher 48 lipid reserves and $\delta^{15}N$ values, and lower $\delta^{13}C$ values occurred during monsoon periods 49 around Seychelles than in the Mozambique Channel (parted from about 1500 km). Our multi-50 51 tracer approach reveals the magnitude of potential competitive interactions in mixed tropical tuna schools at both small and large sizes and demonstrates that ontogenetic niche 52 differentiation acts as a major factor of coexistence in tropical tuna. 53

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55 Keywords: niche area; resource overlap; mixed-school; ontogenetic changes; trophic tracers.

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66 **1. INTRODUCTION**

Three principal market tropical tuna species, bigeye (Thunnus obesus; BET), skipjack 67 (Katsuwonus pelamis; SKJ) and yellowfin (T. albacares; YFT), occur in the pelagic 68 environment in schools, often mixed, that are largely targeted by industrial tuna fisheries 69 including purse seine. In all oceans, tropical tuna schools display specific compositions with 70 SKJ, whose size does not exceed one meter, generally recovered with juvenile BET and YFT, 71 72 while adults of these two species (>100 cm) can be found together (Fonteneau et al., 2000; 73 Hare et al., 2015). Indeed, diving abilities increase with size in BET and YFT and result in 74 deeper vertical distribution with age, whereas SKJ is more vertically confined to the surface 75 layers due to the lack of swim-bladder and higher oxygen requirement (Graham and Dickson, 76 2004). In the purse seine fishery of the Western Indian Ocean, small tuna (fork length; FL≤100 cm) mainly aggregate under floating objects in mixed schools and large tuna 77 78 (FL>100 cm) mainly form free-swimming schools, although some large fishes are also 79 associated with floating objects (Chassot et al., 2015). Sympatry of tropical tuna in mixed schools suggests possible competition for trophic resources, particularly where resources are 80 81 limited.

The trophic ecology of tropical tuna has mainly been investigated through stomach content 82 83 analysis which suggests that all species collectively display opportunistic predatory 84 behaviour, feeding during the day on a large diversity of prey composed of small fish, crustaceans and cephalopods (Potier et al., 2004; Olson et al., 2014). Globally, diet 85 composition has been shown to be influenced by local prey availability, fishing gear that 86 corresponds to specific environmental characteristics (e.g. depth, time of the catch), and 87 environmental conditions (e.g. season, geographical area) (Young et al., 2010b; Olson et al., 88 89 2014). In the Indian Ocean, crustaceans, mainly the swimming crab *Charybdis smithii* and the 90 stomatopod Natosquilla investigatoris, were found to be the main prey of the three species, 91 especially during the day time in September-October (Romanov et al. 2009), followed by 92 small fish like Cubiceps pauciradiatus and small scombridae (Potier et al., 2001, 2004; 93 Grande, 2013; Zudaire et al., 2015). Cephalopods, especially the ommastrephids Sthenoteuthis 94 oualaniensis and Ornithoteuthis volatilis, occurred mainly in BET and YFT diet (Potier et al., 95 2001; Ménard et al., 2013). In addition, feeding habits of tuna have been shown to vary between large individuals that are shallow and deep-dwelling (Potier et al., 2004). While large 96 BET and YFT continue to feed on small prey, at equal size, BET hunt on larger prey than 97 YFT (Ménard et al., 2006), and the average prey size increases with predator size in the 98 tropical tuna (Jaquemet et al., 2011). Limited information is available on diet shift in tropical 99

tuna, globally, with the exception of a study by Graham et al. (2006) who detected anontogenetic shift in coastal YFT from Hawaii at around 50 cm FL.

102 Since stomach content analysis only provides a snapshot of the animal's diet (Cortés, 1997), alternative techniques capable of examining trophic relationships over longer time periods 103 have been progressively introduced. Stable isotopes (SI) and fatty acids (FA) are two of the 104 main trophic tracers currently used in trophic ecology as they have the advantage of reflecting 105 nutrients assimilated over period of weeks to multi-years, depending on the rate at which they 106 are incorporated from diet into consumer tissues and the turn-over of the tissues (Dalsgaard et 107 al., 2003; Buchheister and Latour, 2010). SI of carbon (expressed as δ^{13} C, %) details the 108 origins of primary producers' with different values allowing primary feeding habitats to be 109 differentiated (i.e., continental vs. near shore marine vs. offshore marine; Clementz and Koch, 110 2001). Within an ecosystem, SI of nitrogen (expressed as $\delta^{15}N$, ‰) provides a proxy of an 111 112 animal's trophic position due to a predictable stepwise enrichment from the heavy to the light 113 isotope between trophic levels. Profiles of FA can be used to examine the dynamics of predator-prey relationships, food web structure and energy fluxes (Iverson et al., 2004). 114 Increasingly the importance for dietary studies to focus on FA stemming from storage lipids 115 (i.e. neutral fatty acids, NFA) has been emphasized (Lee et al., 2006; Pethybridge et al., 2014) 116 117 as they provide fuel for energetic demands and are transferred with limited modifications from prey to predator in comparison to structural lipids or other nutrients (Robin et al., 2003). 118

Trophic niche metrics (including nestedness, total niche area, niche widths, and niche 119 overlap) derived from functional ecology have been recently adapted to stable isotope ecology 120 (Layman et al., 2007; Cucherousset and Villéger, 2015) and have also been explored using FA 121 (Pethybridge et al., 2014). Several species-specific studies have used SI or FA to investigate 122 123 either tuna's trophic ecology (SI: Graham et al., 2006; Ménard et al., 2007; Zudaire et al., 2015; FA: Saito et al., 1996; Parrish et al., 2013) or the effect of environmental changes 124 125 (Lorrain et al., 2015; Pethybridge et al., 2015b, 2015a). To our knowledge, no study has undertaken inter-specific comparisons of the three tropical tuna' diet at the same time (their 126 127 niche partitioning) despite the ecological and economic significance of them co-occurring in 128 mixed schools. This study aimed to examine the extent of resource partitioning among 129 sympatric tropical tuna and with ontogeny for a given species in the Western Indian Ocean. By using a multi-tracer (SI and NFA) and multi-tissue (muscle and liver) approach, our 130 comparative study provides valuable insights into how these co-occurring marine top-131 predators compete or partition space and prey resources. 132

133

134 2. MATERIAL AND METHODS

135 **2.1. Tuna and tissue collection**

136 Tuna sampling was conducted throughout 2013 in the Western Indian Ocean during the unloading of the purse seiners at Victoria port, Seychelles. Fish selection was based on two 137 criteria: (i) the capture area from January to December to cover possible geographic and 138 seasonal trophic changes (sampling realized according to the information provided by the 139 purse-seine logbook; the mean coordinates were used when several activities, always 140 comprised within a maximum of 5° square, were grouped in purse-seine wells); and (ii) tuna 141 142 size, to examine ontogenetic dietary shifts. A total of 81 BET, 94 SKJ and 93 YFT were collected and processed at the laboratory where FL (in cm, taken from the tip of the snout to 143 144 the fork of the tail), and sex (Indeterminate, I; Female, F; Male, M) were recorded for each individual. Maturity stage was based on macroscopic examination of the gonads and was used 145 146 to group fish into two phases: developing (Dev, i.e., immature to developing fish), and 147 spawning (Spa, i.e., spawning capable to resting fish; Brown-Peterson et al., 2011). Table S1 gives details of the number of samples analyzed for each fish descriptor, i.e. size class, sex, 148 season, capture area and maturity stage. Two sub-samples of frozen fish of around 2 g (wet 149 weight, ww) were taken from the dorsal white muscle (sampled under the dorsal spine on the 150 151 left side) and the liver of each fish, and frozen within 20 min. All samples were stored at -152 80°C until further analyses.

153

154 2.2. Trophic tracer analyses

155 The detailed procedures are presented in the Supplementary material.

A total of 518 samples (250 livers and 268 muscles) were analyzed for total lipid content 156 (TLC), δ^{13} C and δ^{15} N according to the methods of Bodin et al. (2009) and Sardenne et al. 157 (2015b). Briefly, TLC of each sample, expressed in % of dry weight (dw), was estimated by 158 Accelerated Solvent Extraction with dichloromethane. $\delta^{13}C$ and $\delta^{15}N$ were analyzed together 159 on dried lipid-free samples using a Delta V Advantage isotope ratio mass spectrometer 160 interfaced to a Flash EA 1112 elemental analyzer (Thermo Scientific). Analytical precision 161 for both δ^{13} C and δ^{15} N was <0.15% based on replicate measurements of internal laboratory 162 163 standard (acetanilide and peptone, Thermo Scientific).

A total of 367 samples (180 livers and 187 muscles) were selected for the analysis of NFA, to cover a range of sizes. Compounds were separated on a TRACE 1310 gas chromatograph equipped with a FAMEWAXTM column (30 m, 0.32 mm internal diameter, Restek) and a

167 flame-ionization detector (Thermo Scientific). NFA results were expressed in % as a relative168 abundance of total identified compounds.

169

170 **2.3. Data analyses**

Variations in TLC, δ^{13} C and δ^{15} N values, and NFA among tuna species were investigated to 171 test the effects of fish size, habitat (i.e., fishing area) and season. Two distinct size classes 172 based on the aggregative behavior were considered (FL ≤ 100 cm and >100 cm; Chassot et al., 173 2015), with the second class only including BET and YFT. As previous studies highlighted 174 the peculiarity of the Mozambique Channel (MOZ) in term of productivity (Potier et al., 175 2014), this sampling area was analyzed separately from the Western-Central Indian Ocean 176 (WCIO; Fig. 1). Finally, the monsoon circulation results in a strong seasonal variability in 177 oceanographic features and biological productivity in the Indian Ocean (Schott and 178 179 McCreary, 2001), with subsequent effects on tuna prey such as mesopelagic fishes (Vipin et 180 al., 2012). Hence, four distinct seasons were considered in subsequent analyses: North-Eastern Monsoon (NEM) from mid-November to mid-March, South-Western Monsoon 181 (SWM) from mid-May to mid-September, Spring Inter-Monsoon (SIM) from mid-March to 182 mid-May, and Autumn Inter-Monsoon (AIM) from mid-September to mid-November. 183 184 However, season was excluded from area/season interaction tests because the sampling was 185 unbalanced between these two variables (Table S1) in relation to tropical tuna seasonal migration (Kaplan et al., 2014). 186

A total of 27 NFA accounting each for >0.8% in all samples cumulated (i.e., the most 187 representative) were kept for statistical analyses and arc-sinus root squared transformed to 188 improve normality and homoscedasticity (Underwood, 1997). Permutational multivariate 189 190 analyses of variance (PERMANOVA) based on Euclidean distance matrix were first 191 performed on NFA profiles of muscle and liver separately to test the effects of species, size 192 class, area and season. Non-metric multidimensional scaling (NMDS), also based on the 193 Euclidean distance matrix, was then used to compare the samples through the relative 194 dissimilarity in NFA profiles among tested groups. NMDS is a rank-based approach that aims 195 to summarize the information provided by a large set of dimensions. Here, a set of three 196 dimensions was considered and an optimal configuration of the individuals was based on a 197 maximum limit value of 0.1 for the stress (i.e., goodness-of-fit index). The samples coordinates in the three dimensions of the NMDS were subsequently used to investigate the 198 ontogenetic changes in NFA composition in each tuna species. Differences in individual NFA 199 200 compounds were assessed using Wilcoxon's tests, pairwise tests being used when more than

two means were compared. Analysis of variance (ANOVA) was finally performed on δ^{13} C, δ^{15} N and TLC, and followed by Tukey's honestly significant difference (HSD) post hoc tests. Ontogenetic changes in trophic tracers in each tissue were investigated for each species using Generalized Additive Models (GAMs; Hastie and Tibshirani, 1990). FL was used as a proxy of age because of the difficulties associated with age estimation in tropical tuna (Sardenne et al., 2015a). Sex of the fish (*S*) and maturity stage (*D*) were included in the model in addition to sampling area (*A*) and season (*M*) following:

208 $T_i = a + s(FL_i) + A_i + M_i + S_i + D_i + \varepsilon_i$,

with T is the trophic tracer (δ^{13} C, δ^{15} N, TLC or NMDS scores of NFA profiles), a the 209 intercept, s the spline function smoother (with no preset value concerning the amount of 210 smoothing), *i* indicates individual fish of a given species, and ε is the random noise term 211 212 assumed to be normally distributed with mean zero and constant variance. The best model was selected using Akaïke Information Criterion (AIC), and ANOVAs were used to detect 213 214 significant effects of covariates. When a significant size effect was detected on NFA profiles 215 in a given tissue, the initial model was also applied on the three main NFA families (monounsaturated FA, MUFA; poly-unsaturated FA, PUFA; and saturated FA, SFA) and essential 216 NFA (i.e. 22:603 (DHA); 20:503 (EPA); 20:406 (AA); 18:206; 18:303; Arts et al., 2001) to 217 investigate how they vary with size. 218

Species-specific indices of nestedness based on SI or NFA (i.e. convex hull areas or volumes) 219 220 were calculated to compare tuna' trophic niches as described in Cucherousset and Villéger (2015). Convex hull allows integration of organisms positioned at the edge of SI or NFA 221 222 niches. The indice of nestedness is the ratio between the niche volume shared and the minimal volume filled by a group. It ranges between 0 and 1 with 0 indicating no overlap between the 223 trophic niches and 1 indicating a perfect overlap or a full inclusion of the lowest volume in the 224 225 larger one. It was computed on an isotopic two-dimensional standardized space, where each axis was scaled to have the same range (e.g. 0-1) for each stable isotope (INes; Cucherousset 226 227 and Villéger, 2015), and on a NFA three-dimensional space (FANes), based on NMDS 228 coordinates. These two indices were calculated to compare species trophic niches, according 229 to size class and sampling area, using bootstrapping to overcome differences in the number of 230 individuals per species.

All statistical analyses were performed using R 3.0.2 software. The *vegan* library (Oksanen et al., 2015) was used for PERMANOVA and NMDS and the *mgcv* library (Wood and Wood,

233 2015) was used for fitting GAMs. Indices of nestedness were calculated using the published R
234 code of Cucherousset and Villéger (2015).

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236 **3. RESULTS**

3.1. 3.1.

3.1.1. Total lipid content comparisons

Inter-specific differences

TLC was significantly higher in liver $(15\pm8\%)$ than muscle $(3\pm2\%)$, Tukey HSD, P<0.0001; 239 Table 1). Muscle TLC did not vary with season, area, size class or species (Table 2). Liver 240 241 TLC significantly varied with size class and season, but similarly to muscle, no inter-specific 242 differences were highlighted in liver samples either (Table 1). According to post-hoc Tukey HSD comparisons, large tuna presented higher lipid-rich livers than small tuna (TLC =243 $22\pm11\%$ and $14\pm6\%$, respectively, P<0.0001). Tuna liver was also characterized by 244 245 significant lower TLC during SIM than during the monsoon periods with tuna sampled during 246 this period being $4\pm7\%$ leaner than during NEM (P<0.05) and $5\pm2\%$ leaner than during SWM (P<0.001). While ANOVA showed that there was a significant influence of area on TLC in 247 both tissues in each size class (Table 2), post-hoc tests detected no TLC differences between 248 area in each size class in liver (small and large size classes, P=0.80 and P=0.29 respectively) 249 250 and muscle (P=0.12 and P=0.05 respectively).

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252 3.1.2. Neutral fatty acids profiles

NFA profiles for all tuna were consistently dominated by saturated compounds (means 253 ranging from 42 to 49%, Table 1), that remained constant across size classes. However, 254 varying importance of PUFA (17–37%) and MUFA (13–35%, Table 1) was observed. Among 255 256 the species, the most common NFA in both tissues, in decreasing order of importance, were typically: 16:0 ($32\pm7\%$), 22:6 ω 3 (DHA; 19 $\pm7\%$), 18:1 ω 9 (13 $\pm7\%$) and 18:0 (11 $\pm3\%$). There 257 258 were significant differences in NFA profiles between tissues, with muscle having higher proportions of PUFA and especially of DHA than liver (PUFA: 33±9% vs. 29±9%, W=13379, 259 260 P<0.001; DHA: 21±7% vs. 17±6, W=10364, P<0.0001).

For both tissues, differences in NFA profiles were observed among size classes, species, areas and seasons, in decreasing order of importance. Interactions among size classes and species and among species and area were also detected in muscle (both P<0.05), when a significant interaction between size classes and area was detected in liver (P<0.01; Table 2).

In both tissues, small tuna ($\leq 100 \text{ cm FL}$) generally had higher proportions of PUFA ($32\pm8\%$ in liver and $35\pm9\%$ in muscle, the three species combined) than larger tuna (PUFA: $22\pm10\%$

in liver and 29±8% in muscle, YFT and BET combined; both p<0.0001), and especially of 267 essential PUFA (e.g. in liver: DHA: 18±5% vs. 12±6%; EPA: 5±2% vs. 3±2% and AA: 4±1% 268 269 vs. $3\pm 2\%$). In contrast, MUFA were higher in large tuna, within both liver ($32\pm 10\%$ vs. $21\pm8\%$, W=5315, p<0.0001) and muscle ($25\pm6\%$ vs. $18\pm6\%$, W=5442, P<0.0001). Levels of 270 SFA remained similar between size class: $46\pm7\%$ in liver (W=2954, P=0.28) and $47\pm9\%$ in 271 muscle (W=2886, P=0.16), with limited differences in individual SFA (mean difference <2% 272 for 15:0 and 17:0 in liver and 14:0, 18:0 and 20:0 in muscle). 273 Among the species, higher PUFA proportions were found in SKJ liver and muscle (32±8% 274

and $37\pm9\%$, respectively) than in the two other species (on average $27\pm9\%$ and $31\pm8\%$ for 275 liver and muscle respectively for YFT and BET, small and large specimens combined, all 276 277 P<0.0001). DHA mainly contributed to these inter-specific discriminations, with higher proportions in SKJ (19±6% and 24±6% in liver and muscle, respectively), while BET and 278 279 YFT had similar proportions (on average $15\pm6\%$ and $20\pm6\%$ in liver and muscle, 280 respectively). In contrast, regardless of size class, in both tissues MUFA proportions were highest in BET followed by YFT then SKJ (e.g. in liver, 30±10% in BET, 25±11% in YFT 281 and 18±5% SKJ, small and large specimens combined for BET and YFT, all P<0.05). Those 282 differences were mainly attributed to 18:1ω9, 18:1ω7 and 20:1ω9 (Table 1). Overall, SFA 283 284 were in comparable proportions among species, despite some specific differences in liver (e.g. 285 16:0 was higher, 36±7% in SKJ compared to 34±5% in YFT and 31±5% in BET small and large specimens combined for BET and YFT). 286

The influence of sampling area on tuna's NFA was highlighted by post-hoc Wilcoxon's tests, 287 but different results were obtained for each tissue. Slightly higher proportions of SFA were 288 found in muscle of tuna from WCIO $(47\pm9\%)$ than those from MOZ $(44\pm9\%)$; W=4043, 289 290 P<0.05), while proportions of PUFA and MUFA were similar (P=0.15 and P=0.10). In tuna' 291 liver, proportions of PUFA were higher in MOZ $(34\pm7\%)$ than in WCIO $(28\pm10\%)$; W=1927, 292 P<0.005, mainly driven by changes in DHA (W=3714, P<0.01) but not EPA (P=0.09), the 293 second most relevant PUFA. In contrast, MUFA were higher in WCIO (25±11%) than in 294 MOZ ($20\pm8\%$; W=3765, P<0.005) while proportions of SFA were the same (P=0.718). 295 Seasonal effects were observed in tuna' liver due to variations in $\omega 6$ proportions between SIM 296 (7±2%) and SWM (6±2%; P<0.01) and in EPA proportions between AIM (6±2%) and NEM $(5\pm 2\%)$; P<0.05). Seasonal effects were also observed in tuna' muscle largely due to 297 differences in MUFA proportions between SIM (22±6%) and NEM (18±7%; P<0.01). 298 Several interactions among factors influenced tissue NFA profiles. In liver, only an 299

300 interaction between size class and area was detected (Table 2). Thus, large YFT in the WCIO

had significantly different liver NFA profiles than similar sized YFT in the MOZ (PERMANOVA, Pseudo-F=16.9, P<0.001), with on average 14% more MUFA than YFT from MOZ (W=20, P<0.005), mainly attributed to $18:1\omega9$ (12% more; W=18, P<0.005), but on average 15% less PUFA (W=137, P<0.001). In contrast, in the liver of small tuna almost no differences between areas were detected: only ω 6-PUFA family was found higher of 1% in MOZ (MOZ: 8±1%, WCIO: 7±1%; W=2096, P<0.001).

In muscle, inter-specific size-relevant differences from both areas were detected. In small tuna 307 from WCIO, higher proportions of MUFA were reported in BET compared to YFT (mean 308 309 difference of 5%, P<0.05) and SKJ (mean difference of 9%, P<0.001). Such differences were mainly attributed to 18:109 (P<0.001). No significant inter-specific differences were detected 310 among small tuna for PUFA, ω 3- and ω 6-PUFA families, however there were significantly 311 higher contributions of EPA, DHA, AA, and 22:506 observed in SKJ compared to both YFT 312 313 and BET (P<0.05) which showed similar profiles. Only one SFA, 18:0, was found to differ 314 among muscle of small tuna, with lowest proportions found in BET (12% of total NFA) than in SKJ (14%, P<0.05) and YFT (14%, P<0.05). For large tuna in the WCIO, no differences in 315 NFA profiles were noted among species (PERMANOVA, Pseudo-F=1.0, P=0.36): only 316 18:3w3 was found to differ inter-specifically in the muscle. Similar inter-specific size-317 318 comparable differences were also detected among tuna sampled in the MOZ, with lower 319 proportions of PUFA recorded in BET muscle compared to SKJ (mean difference of 14%, P<0.001) and YFT (mean difference of 8%, P<0.005), the highest proportions being measured 320 in SKJ muscle (e.g. DHA on average 8% and 5% lower in BET than SKJ and YFT 321 respectively). The highest MUFA proportions were observed in BET and the lowest in SKJ, 322 mainly attributed to the 18:109, on average 5% higher in BET than in SKJ for muscle 323 324 (P<0.05).

325

326 3.1.3. Stable isotope values

327 δ^{13} C values varied with area and size class in liver and by size class, species and season in 328 muscle. δ^{15} N values significantly varied with area, season, species and according to a species-329 area interaction in both tissues, but varied with size class in muscle only and with a size class-330 species interaction in liver (Table 2).

331 Muscle δ^{13} C values were slightly lower during NEM (-17.1±0.2‰) than AIM (-16.8±0.2‰)

and SWM (-16.9±0.4%) (P<0.05) but no difference was observed in liver. δ^{15} N values in tuna

muscle were significantly higher in WCIO ($12.2\pm0.9\%$) than in MOZ ($10.2\pm1.3\%$, P<0.001),

with a similar trend displayed in the liver (WCIO: $11.4\pm0.7\%$; MOZ: $9.3\pm1.2\%$, P<0.001).

 δ^{15} N values in both tissues were lower during SIM than during the other seasons (i.e. in liver: mean value in SIM was 1.9% lower than AIM, 1.8% lower than NEM and 1.6% lower than SWM, P<0.001; and in muscle: mean value in SIM was 1.3% lower than AIM, 1.0% lower than NEM and 1.5% lower than SWM, P<0.001).

Size-class differences were reported for both δ^{13} C and δ^{15} N values in muscle with higher values reported in large tuna in comparison to small tuna (mean difference of 0.2‰ and 0.8‰ respectively, P<0.001). In liver, δ^{13} C values were significantly higher in large tuna than in small tuna (mean difference of 0.3‰, P<0.001), but no difference in δ^{15} N values was observed (P=0.784).

Inter-specific size-comparable differences from the WCIO were found using post-hoc Tukey 344 HSD comparisons. In small tuna, $\delta^{13}C$ values were slightly lower in BET compared to YFT 345 and SKJ in muscle (mean difference of 0.2%, P<0.01), while no inter-specific differences 346 were detected in liver (both P>0.1). In contrast, δ^{15} N values were significantly higher in BET, 347 348 closely followed by YFT, and finally SKJ in both tissues (Table 1), with mean differences among species ranging from 0.3 to 1.1% (P<0.05, except between BET and YFT in muscle: 349 P=0.12). In large tuna, significantly lower δ^{13} C values were found in BET muscle than in 350 YFT muscle (on average 0.5%, P<0.001), while the low differences (of 0.2%) in liver were 351 not significant (P=0.09). BET presented significantly higher δ^{15} N values than YFT for both 352 tissues (on average 1% and 0.8% higher, respectively, P<0.001). In MOZ, δ^{13} C values were 353 slightly lower in small BET compared to small YFT and SKJ, as in WCIO (mean difference 354 of 0.2‰, P<0.05). On the contrary, small BET had higher muscle δ^{15} N values (1.7‰ higher) 355 than YFT (P<0.005) and SKJ (P<0.001), while small YFT and SKJ exhibited similar muscle 356 δ^{15} N values (P=0.986). In liver, the highest δ^{15} N values were recorded for small BET 357 followed by small YFT and finally SKJ, with mean differences ranging from 0.7 to 1.9% 358 (P<0.05). Post-hoc Tukey HSD comparisons revealed significantly lower δ^{15} N values in large 359 360 YFT from MOZ than in large YFT from WCIO in muscle and liver (mean difference of 2.3‰, P<0.001 and 1.9‰; P<0.001, respectively). δ^{13} C values in muscle of large YFT were 361 also lower in MOZ than in WCIO (mean difference of 0.3%, P<0.001), while no difference 362 363 was detected in liver (P=0.386).

364

365 3.1.4. Biochemical trophic niche metrics

Trophic niche overlap based on SI (INes) and FA (FANes) was explored among the three tuna species differentiating patterns between tissues, size classes and sampling areas. In WCIO, index values showed less variation in liver (ranging from 0.30 to 0.81) than in muscle (0.09 to

0.81, Table 3), while they were similar in MOZ (from 0.08 to 0.73 in liver and from 0.06 to
0.63 in muscle). Although some differences were evident between INes and FANes, the
degree of overlap among species agreed, with the lowest trophic space shared by BET and
SKJ.

According to these indices in both tissues of small tuna, and in particular muscle, SKJ and 373 BET shared limited trophic space, whereas the highest niche overlap was between BET and 374 YFT. Indeed, in muscle of small tuna sampled in the WCIO, the NFA niche volume defined 375 for YFT overlapped with that of BET, while the niche area of SKJ differed mainly due to 376 PUFA (Fig. 2a). A similar pattern was observed for INes in liver, with large overlap between 377 YFT and BET (0.81) due to higher δ^{15} N values compared to SKJ (Fig. 3a). However, in 378 muscle, the highest INes was observed between small YFT and SKJ (0.81). In large tuna, 379 380 FANes in both tissues showed quite large trophic overlaps between BET and YFT (Table 3, 381 Fig. 2b). On the contrary, differences in INes were apparent with no overlap of trophic space 382 between large BET and YFT in muscle (INes=0.0) and a limited overlap in liver (INes=0.34) 383 (Fig. 3b).

In MOZ, SKJ and small YFT shared most of their trophic space, while small BET and small YFT shared almost a third of their trophic space (with FANes and INes in both tissues ranging from 0.40 to 0.73 and 0.29 to 0.44, respectively, Table 3). In comparison, small BET and SKJ exhibited reduced overlaps according to both FANes and INes in both tissues.

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3.2. Intra-specific variability focusing on ontogenetic and sexual shifts

390 *3.2.1. Bigeye tuna*

GAM predicted TLC in BET remained constant in muscle (3±2 % dw) while it generally 391 392 increased in liver with size from around 10 to 20% dw between 29.5 and 160.8 cm FL, with 393 no influence of other tested variables, as sex (Table S2). The dominance of the predicted 394 proportions of MUFA and PUFA (predominantly represented by 18:1009 and DHA 395 respectively) were reversed in liver at 91 cm FL with MUFA higher in larger individuals (Fig. 396 4a). Proportions of MUFA increased linearly with size, while PUFA proportions decreased 397 (23.6% of deviance explained with size alone), mainly due to a decrease of DHA proportion 398 in comparison to other essential NFA (Fig. 4b). No change was observed for SFA in liver, which remained the main NFA class throughout the BET size range. Finally, deviance 399 explained for the main NFA families in BET muscle was poor (ranging from 5.3% for PUFA 400 to 23.4% for MUFA), but there was also a slight increase of MUFA proportion with size. No 401 402 influence of maturity or sex was observed in NFA in either tissues (Table S3a and S3b).

Predicted values of both isotopes were mainly affected by size and area in both BET tissues 403 (Table S2). Season also influenced δ^{13} C values in both tissues, but no influence of sex or 404 maturity on isotopic data was observed. No clear pattern with size was obtained for liver $\delta^{13}C$ 405 values, with a mean increase from around -18.2±0.2% to -17.4±0.2% between 30 and 76 cm 406 FL before oscillating around -17.4±0.5% until 160 cm FL (Fig. 5a). δ^{13} C values in muscle 407 displayed an increase from 30 to 77 cm FL (from around -17.5±0.1% to -17.0±0.1%) and 408 then remained constant at -17±0.1‰ until 160 cm FL (Fig. 5b). Liver δ^{15} N values slowly 409 increased with BET size between 30 and 69 cm FL (i.e. mean increase of 0.7% in 39 cm, 410 411 from around 11.3±0.2% to 12±0.1%), then remained stable until 160 cm FL (Fig. 5a). In muscle, the main variation of δ^{15} N was observed from 30 to 54 cm FL (i.e. mean increase of 412 1.7% in 24 cm, from around 11.0 ± 0.2 to $12.7\pm0.2\%$), followed by relatively constant values 413 until 140 cm and a slight increase between 140 and 160 cm (i.e. from around 13.4±0.2% to 414 415 13.7±0.3%; Fig. 5b).

416

417 3.2.2. Yellowfin tuna

TLC in YFT muscle and liver was affected by size (Table S2) but with unpredictable trends, 418 and the fitted GAM performed poorly for muscle with only 11.6% of deviance explained. In 419 liver, TLC oscillated with size, and males had a slightly higher TLC than females and 420 421 indeterminate individuals combined (23±11% vs. 15±8%, P<0.05). NFA profiles were mainly affected by size (Table S3a). In liver, the mean proportions of MUFA and PUFA were 422 reversed at 109 cm FL (Fig. 4a): MUFA increased by about 20% from 109 to 160 cm FL 423 (from around 50% to 70%), when PUFA subsequently decreased (from around 55% to 35%). 424 Size explained 38% of MUFA and 51.5% of PUFA variability in liver, but showed minor 425 426 influence on SFA (Table S3b). Among PUFA, three essential NFA contributed strongly to the 427 observed PUFA decrease with size in liver: DHA, EPA and AA (Fig. 4b). Although minor 428 contributors, $18:3\omega3$ and $18:2\omega6$ also significantly decreased with size (P<0.0001). In muscle, PUFA and SFA generally remained constant with size $(38\pm3\%)$ and $44\pm5\%$, respectively). In 429 430 contrast, size explained 58.4% of MUFA variability in muscle with proportions showing a progressive increase from $14\pm 2\%$ at 32 cm FL to $29\pm 2\%$ at 157 cm FL. Despite a maturity 431 432 influence in NFA profiles of liver and muscle, no influence was detected when considering 433 the main NFA families individually (Table S3b). Isotopic compositions in both tissues were influenced by all tested variables, with the exception of sex. In liver, GAM predicted δ^{13} C 434 values increased ca. -17.7±0.1% to -16.6±0.2%, and δ^{15} N values from ca. 12.1±0.2% to 435 $12.5\pm0.3\%$ % between 29 and 157 cm FL (Fig. 5a). In muscle, minor variations of δ^{13} C 436

437 values were observed over the range size (<0.6‰), while δ^{15} N values strongly increased 438 between 29 and 53 cm (from 10.7±0.3‰ to 12.9±0.3‰), then progressively increased until 439 157 cm to reach 13.9±0.4‰, with a stronger increase between 140 and 157 cm (Fig. 5b). 440 Muscle δ^{13} C values were also highest in YFT spawning (-16.5±0.2‰) compared to those 441 developing (-17±0.3‰, P<0.001).

442

443 **3.2.3.** Skipjack tuna

GAM predicted TLC was not affected by any of the tested variables in muscle of SKJ (Table 444 445 S2) but varied with size in liver (min-max=2-30% dw), despite no obvious trend observed. In 446 SKJ tissues, differences in NFA profiles were mainly attributed to sex and area, whereas size 447 had no significant influence in muscle and a limited influence in liver (Table S3a). There was also no influence of maturity in both tissues (Table S3a). Wilcoxon tests revealed that the 448 449 liver of females contained more SFA (53±7% vs. 46±7%; W=218, P<0.05), mainly 16:0 450 (W=238, P<0.005), and more MUFA than males $(21\pm5\% vs. 15\pm5\%; W=240, P<0.005)$ mainly 16:1 ω 7 (W=250, P<0.001) and 18:1 ω 9 (W=234, P<0.01) (Table S4). In contrast, liver 451 of males had higher relative amounts of essential NFA including EPA, DHA and AA (all 452 P<0.01; Table S4) than females: mean differences between males and females ranged from 453 1% for 22:506 to 8% for DHA (Table S4). PUFA proportion in liver was higher in males and 454 indeterminate individuals (W=201, P<0.05) than in females (W=173, P<0.001, Table S4). 455

In contrast, no differences in isotopic compositions were observed between sexes in either tissue (Table S4). No clear trend was observed for δ^{13} C values which remains overall constant (around -17.5% in liver and -17% in muscle), whereas δ^{15} N values increased by 2.0% in muscle from 11%, and by 1.1% in liver from 11.1% between 30 and 78 cm FL (Fig. 5). For both isotopic values in both tissues, no influence of maturity was noted (Table S2).

461

462 **4. DISCUSSION**

Using complementary trophic tracers, this study provides greater understanding of trophic 463 464 relationships and food partitioning among tropical tuna from the Western Indian Ocean. Due 465 to their biological proximity and habitat sharing, overlaps in their trophic niches were 466 detected, especially in NFA compositions of large individuals (Fig. 2b). However, significant inter- and intra-specific differences were also noticed: higher PUFA proportions in SKJ than 467 in small BET and YFT (Fig. 2a); separated isotopic niches occupied by large YFT and BET 468 (Fig. 3b); increased MUFA proportions with size in the liver of BET and YFT (Fig. 4), while 469 a difference between sexes was observed only in SKJ (higher PUFA proportions in males). 470

471

472 4.1.FA and SI compositions of tuna prey

473 The origins of these inter-specific differences are likely explained through the FA and SI compositions of their main prey and this section gives an overview of these compositions, 474 used in the following sections. Only a few studies report NFA profiles for tuna prey in the 475 Western Indian Ocean, but some information exists for other oceans. High MUFA proportions 476 are characteristic of myctophid fishes (Saito and Murata, 1998), including in tropical waters 477 (Sebastine et al., 2011; Baby et al., 2014), due to their high consumption of copepods rich in 478 479 20:1 MUFA (Saito and Kotani, 2000). Myctophids are small pelagic fishes, inhabiting deep layers (over 400 m) during the day and that migrate to surface layers (around 5-100 m) each 480 night to feed (Vipin et al., 2012). They are important prey of squids. Indeed, middle-sized 481 squid Sthenoteuthis oualaniensis, a common tuna prey in the region under study (Ménard et 482 483 al., 2013), ascends between the surface and 100 meters depth only at night to feed on 484 myctophids (Shulman et al., 2002). Despite S. oualaniensis feed on myctophids rich in MUFA, its mantle contains high proportions of DHA (Shulman et al., 2002) and PUFA (52% 485 of total FA; Young et al., 2010a). Cigarfish C. pauciradiatus is another small fish prey rich in 486 MUFA (Young et al., 2010a), on which YFT mainly forage during the NEM period in WCIO 487 488 (Zudaire et al., 2015). Globally, less information is available concerning the FA profile of 489 crustaceans, including the swimming crab Charybdis smithii and the stomatopod Natosquilla investigatoris, that are also known prey of tropical tuna (Potier et al., 2004). In the Pacific 490 491 Ocean a related crab species, *Charybdis hawaiiensis*, is reported to be mainly composed of 492 PUFA (42% of total FA) (Piché et al., 2010). In addition, C. smithii come up in surface layers 493 mainly at night in the Indian Ocean (Romanov et al., 2009), while N. investigatoris was 494 occasionally found in surface layers at daytime (Losse and Merrett, 1971).

495 In contrast to NFA profiles, isotopic data of the main tuna prey groups were available for the 496 Western Indian Ocean, but more so for the Mozambique Channel. Squids and small fishes have been shown to occupy similar trophic levels with mean δ^{15} N values around 9.4±0.8% 497 and 9.5±1.6%, respectively, while crustaceans occupy a slightly lower trophic level with 498 mean δ^{15} N values around 8.7±1.3‰ (Ménard et al., 2014). However, a great variability was 499 observed among the small fishes: cigarfish had lower mean δ^{15} N values (9.0±2.2‰) than most 500 myctophids such as *Diaphus metopoclampus* with higher δ^{15} N values (12.2±0.4‰) but lower 501 δ^{13} C values (-19.3±0.2‰; Ménard et al., 2014). δ^{15} N values of main tuna prey have also been 502 shown to increase with size (in myctophids, the cigarfish and squid S. oualaniensis; Parry, 503 2007; Ménard et al., 2014). However, limited variability in δ^{13} C values are displayed among 504

tuna prey with mean values ranging from $-18.6\pm0.4\%$ in small fish to $-18.2\pm0.6\%$ in crustaceans. This suggests that any differences in δ^{13} C values in tuna from this area should be interpreted with caution.

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509 **4.2.Trophic partitioning in tuna aggregations**

Total lipid content data suggests that none of the three tuna species appears to store lipids in 510 muscle, as TLC values remained low and did not vary with the considered factors. The liver 511 showed more variability in TLC, which was mainly explained by season and size class 512 differences, suggesting this organ could be used to store lipids. In the three species, fattest 513 514 livers were sampled during the two monsoon periods; however, modest seasonal differences 515 were detected in SKJ. This could be related to a specific diet during monsoons, as YFT have been recorded to forage predominantly on fatty small fishes (7-39% dw of lipids; Young et 516 517 al., 2010a), such as C. pauciradiatus during NEM in the studied region (Zudaire et al., 2015). 518 Partial niche overlaps were observed in the liver of small tuna (Fig. 2a and 3a). As tuna still form multispecific schools, partial trophic sharing does not appear to reduce the advantages of 519 schooling behavior (e.g. reduced predation, faster food detection; Pavlov and Kasumyan, 520 2000). When trophic niches are large (i.e., diet behaviors of individuals are diverse in each 521 522 species), diet overlap among species does not really increase the competition intensity among 523 them, because only a few individuals in each species are impacted; Bolnick et al. (2011) called this "niche complementarity". This phenomenon might reduce predator-prey 524 dependency and dampen consumers-resources oscillations (McCann et al., 1998), which may 525 offer a long-term advantage in nutrient poor habitats such as tropical offshore waters. As these 526 small tuna were caught under drifting fish aggregating devices, the use of which has recently 527 528 intensified (Chassot et al., 2015) and is thought to act as an ecological trap (e.g. Hallier and 529 Gaertner, 2008), the possibility that devices promote competition among small tuna cannot be 530 excluded (Jaquemet et al., 2011)

Calculated FANes showed greater trophic overlap between YFT and BET than between YFT 531 532 and SKJ, mainly based on differences in MUFA and PUFA proportions. The high proportions 533 of PUFA in SKJ suggest a lower consumption of myctophids, squid consuming myctophids, 534 and small fish in general, but a higher consumption of crustaceans (see section 4.1 for details, and Grande (2013) who investigated stomach contents of SKJ in the Western Indian Ocean). 535 By contrast, PUFA proportions in small BET and in YFT suggest a low consumption of 536 crustaceans that seems to decrease with size with higher proportions of MUFA in large 537 individuals (Fig. 4a). In addition, mean δ^{15} N values in both tissues were higher in small BET 538

than in small YFT and SKJ. This suggests that small BET likely feeding on prey with higher 539 540 trophic positions and/or of a larger size, as already observed by comparing the size of prey 541 from stomach contents of YFT and BET (Ménard et al., 2006). No studies have specifically focused on tuna prey size distribution in the Indian Ocean, however, Potier et al. (2008) 542 noticed that surface YFT and BET fed more on smaller cigarfish than sub-surface YFT and 543 BET of the same size, and suggested ontogenetic changes in the depth distribution of 544 cigarfish. On the other hand, diving capacities differences among small tuna (further 545 discussed in section 4.3) also suggest the possibility of vertical sharing in mixed schools of 546 547 small tuna, with small BET occupying the deeper layers of schools.

In large tuna, only minor differences in the proportions of individual NFA were observed 548 (Table 1). This suggests that the two species BET and YFT fed on similar proportions of the 549 550 same prey species, which is in agreement with stomach content analyses of large tuna caught 551 by purse-seine (Potier et al., 2004). However, the stable isotope compositions were different 552 between these two species and suggested a higher trophic level in BET than in YFT (Fig. 3b). As mentioned for small tuna, differences in prev size could explain differences in δ^{15} N values 553 between large tuna. These results align with stomach content data from the same area, which 554 showed that squid and cigarfish individuals were larger in the stomachs of BET than similar 555 sized YFT (Potier et al., 2008; Ménard et al., 2013). The present study also reported large 556 557 MUFA proportions in the tissues of the larger tropical tuna (BET and YFT) which is likely associated to direct consumption of myctophids or indirect myctophid consumption passing 558 through the digestive gland of various squids (Phillips et al., 2002; Pethybridge et al., 2013). 559 Indeed, large BET in deeper water layers have been found to feed more on squids than similar 560 sized YFT (Potier et al., 2004). As indicated by differences in δ^{13} C values, we suspect that 561 BET fed, directly or indirectly (i.e., through squids), on specific myctophids, such as 562 D. metopoclampus, with lower δ^{13} C values. The particular isotopic position of this myctophid 563 species might increase the isotopic position of large BET, which has higher δ^{15} N values and 564 lower δ^{13} C values than large YFT, in muscle and liver (Fig. 3b). Thus, despite large BET and 565 YFT co-occurring and focusing on similar prey groups in surface waters when caught by 566 567 purse-seine, in long-term (as integration times of trophic tracers used in the present study 568 approach several months; Budge et al., 2011; Madigan et al., 2012), a specific orientation based on prey size and/or specific myctophids predators could limit their competition. In 569 addition, the same inter-specific differences in isotopic compositions were observed in both 570 tissues with different integration time (Fig. 3a). This suggests that the aggregations of large 571

individuals in mixed schools are not durable at the scale of isotopic turnover rates (around 6months in muscle and 3 in liver; Madigan et al., 2012).

574 As noticed by Young et al. (2010b) in the Pacific Ocean for large individuals, diet differences reflect diel variations in foraging among tuna, with large BET thought to feed more at night 575 than YFT, when myctophids and squids ascend from deeper layers (see section 4.1). In 576 contrast, SKJ feed mainly during the day, but night foraging has been reported (Romanov et 577 al., 2009; Grande, 2013). Such diel foraging behaviors may be reflected in the biochemical 578 579 signatures and suggest limited tuna competition. Indeed, considering the main tuna prey, myctophids, squids and the crab C. smithii come up in surface layers mainly at night in the 580 581 Indian Ocean), whereas the stomatopod, N. investigatoris, could be found in surface layers at daytime (see section 4.1) and in stomach content of surface tuna in Indian Ocean (Potier et al., 582 2008; Grande, 2013). Thus, a myctophid dominant diet, as suggested in BET with the high 583 584 MUFA proportions, is probably more related to nocturnal feeding and/or twilight feeding.

Environmental variables (i.e., fishing area and season) influenced trophic tracers in both tissues, attesting to their non-negligible impacts on tuna diet. For example, the cigarfish constitutes most of the tuna diet in the Indian Ocean especially during NEM (Fonteneau et al., 2008; Potier et al., 2008) and appears to be an important energy source in the reproduction of YFT in WCIO (Zudaire et al., 2015). Although this could explain the higher MUFA proportion in the liver of tuna from WCIO than from MOZ, the large variability among individuals does not allow to properly characterizing these environmental influences.

Lower δ^{15} N values were found in tuna from MOZ than from WCIO, as reported in previous 592 studies (Ménard et al., 2007; Zudaire et al., 2015). This is generally attributed to a lower base 593 line in Mozambique Channel (Ménard et al., 2007), due to diazotrophic organisms that lower 594 the $\delta^{15}N$ value of particulate organic matter (Dupuy et al., 2016). An influence on NFA 595 profiles was also possible, as diatoms, with high EPA biosynthesis capacities, have been 596 597 shown to predominate in mesoscale eddies (e.g. Brown et al. (2008) in subtropical Pacific), which are abundant in MOZ (Potier et al., 2014). In contrast, higher proportions of DHA, 598 599 characteristic of dinoflagellates, were found in liver of tuna from MOZ, suggesting a more 600 complex response to mixing events of nutrient, and overall a mixing signal through multiple 601 trophic levels from phytoplankton to higher trophic levels (Kainz et al., 2004).

602

603 **4.3. Diet changes during tuna ontogeny**

Patterns of NFA and SI observed in each tuna species were slightly different with size,
 probably related to diet differences but also to physiological differences. For the three species,

the liver was found to be more informative than the white muscle to detect intra-specific 606 607 changes, as it is a reactive tissue to rapid diet changes in relation with fast turnover rates 608 (around 6 months in muscle and 3 months in liver for stable isotopes; Madigan et al., 2012). In addition, TLC was relatively lower in white muscle than in liver, attesting to the known 609 role of liver in energy storage. Therefore, NFA profiles in muscle appear to be less 610 informative for ontogenetic studies in leaner fish such as these tropical tuna. By contrast, 611 NFA profiles in liver were found to be of valuable interest when examining dietary changes 612 613 with size.

In both BET and YFT, PUFA proportions decreased in liver with size while MUFA 614 concurrently increased (at around 91 and 109 cm in BET and YFT; Fig. 4a). In contrast, no 615 616 changes were observed with size in SKJ, but differences between sexes were noticed, especially in liver, with higher PUFA proportion in males (Table S4). Two main hypotheses 617 618 could support differences among the tuna species: a biological process response (e.g. 619 reproduction) and/or a change of habitat (e.g. from surface to deeper layers). First, concerning reproduction, the size at which 50% of the population matures was estimated at 100 cm FL 620 for BET in the Pacific Ocean (Farley et al., 2006), 75 cm for YFT (Zudaire et al., 2013) and 621 39 cm for SKJ (Grande et al., 2014), both studies from the Western Indian Ocean. Thereby, 622 623 PUFA of mature fish would be allocated to germinal cell production, especially in females. 624 Indeed, vitellogenin synthesis in ovary has been shown to require higher quantities of PUFA, especially in EPA and DHA (Silversand and Haux, 1995). However, no significant influence 625 of maturity on the proportions of the main NFA classes was detected in this study. The 626 influence of reproduction on NFA profiles is however more likely in SKJ, as it has been 627 shown to be an income-breeder species, i.e. which reinvest the majority of its acquired energy 628 629 to reproduction, and breeding all year round (Grande et al., 2014). This suggests that in females, PUFA from liver are rapidly allocated to the ovary after acquisition; while males 630 631 have fewer PUFA requirements and can store them into the liver explaining the higher proportion of PUFA in males than in females SKJ observed in the present study, and, 632 633 potentially explaining higher TLC in liver of males. In the same way, in liver, PUFA 634 proportion was found to be higher in indeterminate individuals (i.e. the smaller fishes) than in 635 females (Table S4). Indeed, according to this assumption, PUFA would be allocated to the gonads. These differences between sexes could also make females SKJ more sensitive to 636 environmental changes, especially if dietary essential PUFA become less available, as 637 predicted for the south west Pacific Ocean (Pethybridge et al., 2015a). 638

Concerning habitat usage, the shift in main NFA families in BET and YFT could correspond 639 640 to a habitat change for deeper environments providing rich-MUFA prey as myctophids or the 641 myctophids-predators squids, which inhabit deeper layers (see section 4.1). Indeed, this shift 642 could correspond to the sufficient development of the swim bladder and regional endothermy (Graham and Dickson, 2004), allowing BET and YFT to reach deeper layers with size, while 643 SKJ is restricted to the surface due to the lack of swim bladder. Magnuson (1973) observed an 644 allometric growth of swim bladder for YFT until 8-10 kg, corresponding approximately to a 645 70-75 cm FL fish. Bertrand and Josse (2000), who observed larger swim bladder in BET than 646 647 in YFT at similar size, observed also an increase in the swim bladder volume over 120 cm FL in BET. This would suggest that BET could reach deeper layers at smaller size than YFT, as 648 emphasized with the earlier shift in NFA profile observed in BET than in YFT (i.e., 91 vs. 109 649 cm FL). 650

 δ^{15} N values should be interpreted in context of specific differences in amino acids 651 composition that are nitrogen compounds (i.e. slow turnover of essential amino acids; Popp et 652 al., 2007), the tissue growth and the turnover rate (Martinez del Rio et al., 2009). Indeed, the 653 fast increase with size of δ^{15} N values in muscle of the three species highlighted the influence 654 655 of metabolic process and the importance of selected tissue for the trophic level estimation in food webs studies. In juvenile bluefin tuna, T. orientalis, Madigan et al. (2012) showed that 656 the half-lives of ¹⁵N in white muscle and liver were 167 and 86 days respectively, and 255 and 657 162 days for ¹³C, respectively, highlighting influences of turn-over on isotopic composition. 658 In addition, in different marine fish, a growth rate increase promotes a decrease of δ^{15} N values 659 in muscle, and high metabolic levels promote a faster equilibration with diet (Trueman et al., 660 661 2005).

In the three species, limited variations of δ^{13} C values in muscle indicated no change of carbon 662 sources through life cycle. However, δ^{13} C values in liver were more difficult to interpret due 663 664 to irregular variations, which could be related to metabolic activities as previously mentioned. In both tissues, an increase in δ^{15} N values with size was observed in the three species with a 665 666 strongest signal in muscle (Fig. 5). This increase could be linked to slowdown in growth rate 667 and/or to an increase of prey size. Indeed, tropical tuna are opportunistic feeders and the mean 668 size of their prey increased with tuna size (Ménard et al., 2006; Jaquemet et al., 2011), besides, $\delta^{15}N$ values of prey also increase with prey size (Ménard et al., 2014). Finally, in 669 BET and YFT, no breakpoints in isotopic values matched specific maturity or growth events, 670 although Zudaire et al. (2015) observed higher δ^{15} N values in liver of spawning YFT. 671

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673 Conclusion

This study provides a greater understanding of trophic interactions among and within three 674 675 prominent tropical tuna species. We found that there is a resource partitioning among sympatric tuna and particularly among tuna ≤100 cm FL. Based on the biochemical tracers, 676 we show evidence of ontogenetic dietary shifts in the larger tuna species (i.e. BET and YFT). 677 Generally, larger tuna individuals were shown to have higher $\delta^{15}N$ and $\delta^{13}C$ values than 678 smaller individuals, suggesting higher trophic positions for the larger individuals, and higher 679 680 proportions of MUFA commonly high in myctophid fish. Future work should focus on reporting of NFA profiles for potential prey in the Western Indian Ocean that to date was 681 particularly scarce of such data, and detailing specific turnover rates of tissues and 682 683 compounds (SI as NFA).

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937 FIGURE CAPTION

Figure 1. Location of skipjack tuna (SKJ), yellowfin tuna (YFT) and bigeye tuna (BET)
caught in the Western Indian Ocean throughout 2013. The limit of the two study areas,
Mozambique Channel (MOZ) and Western-Central Indian Ocean (WCIO), is indicated with a
dashed line.

Figure 2. Scatterplots and associated convex hull of three-dimensional nonmetric multidimensional scaling (NMDS) using Euclidean distance matrix for neutral fatty acid profiles in the muscle of skipjack tuna (SKJ), yellowfin tuna (YFT) and bigeye tuna (BET) caught in the Western-Central Indian Ocean (WCIO), according to size class: (a) fish with fork length less than 100 cm (FL \leq 100 cm) and (b) fish with fork length longer than 100 cm (FL>100 cm). Arrows are significant variables (P<0.05) correlated with the ordination. Dim: dimension.

Figure 3. Two-dimensional scaled isotopic space (δ^{15} N and δ^{13} C) in the muscle and liver of skipjack tuna (SKJ), yellowfin tuna (YFT) and bigeye tuna (BET) caught in the Western-Central Indian Ocean (WCIO), according to size class: (a) fish with fork length less than 100 cm (FL \leq 100 cm), (b) fish with fork length longer than 100 cm (FL>100 cm).

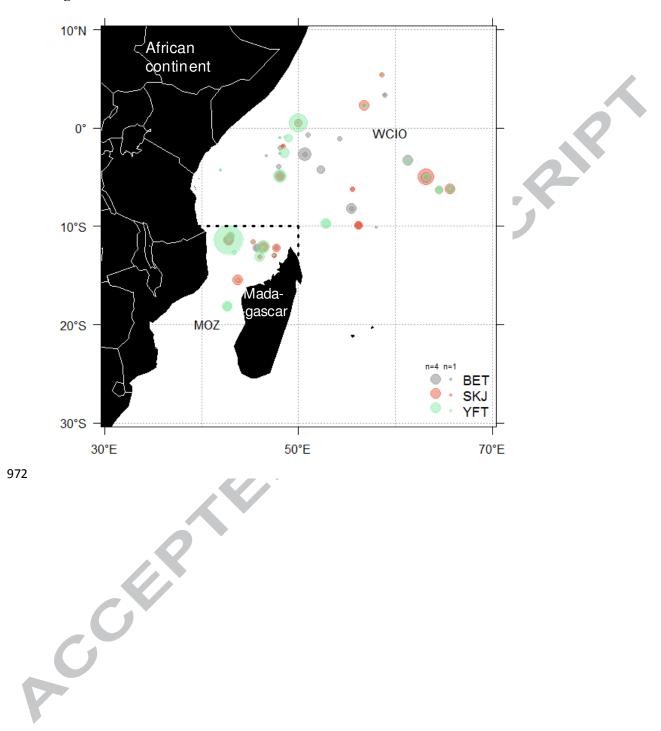
Figure 4. Smoother plots illustrating the relationship between observed fork length (FL) and 953 954 GAM predicted values for (a) the main fatty acid (FA) families (b) six individuals neutral 955 fatty acid (NFA) of which five are essential (DHA, EPA, AA, 18:3ω3 and 18:2ω6) in the liver of BET and YFT sampled in the Western-Central Indian Ocean throughout 2013. Vertical 956 dashed lines represent the size at which a proportion change in the main FA families occurs 957 (91 and 109 cm FL for BET and YFT respectively). Solid areas showed the confidence 958 interval (1.96*standard error) around the predicted value (line). Short vertical lines on the x-959 960 axis represent the sampled values; BET: Bigeye; SKJ: Skipjack; YFT: Yellowfin; EPA: 961 Eicospentaenoic acid ($20:5\omega 3$); DHA: Docosahexaenoic acid ($22:6\omega 3$); AA: Arachinoic acid 962 $(20:4\omega 6).$

Figure 5. Smoother plots illustrating the relationship between observed fork length (FL) and GAM predicted values for stables isotopes values (δ^{13} C in blue and δ^{15} N in black) in (a) liver and (b) muscle of bigeye (BET), yellowfin (YFT) and skipjack (SKJ) from Western-Central Indian Ocean. Solid areas showed the confidence interval (1.96*standard error) around the predicted value (line). Short vertical lines on the x-axis represent the sampled values.

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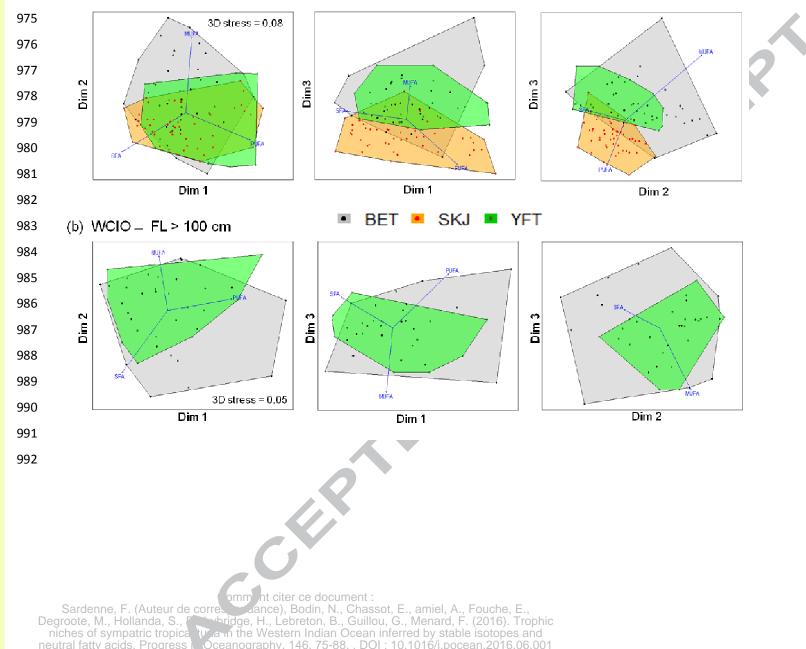
971 **Figure 1.**

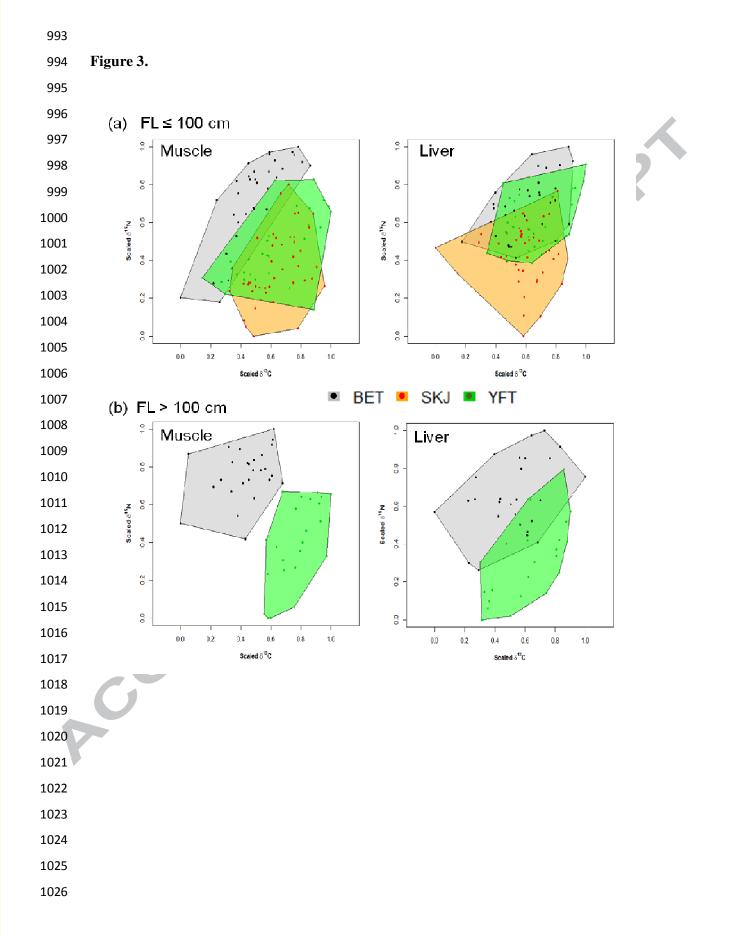


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973 Figure 2.

974 (a) WCIO – FL ≤100 cm

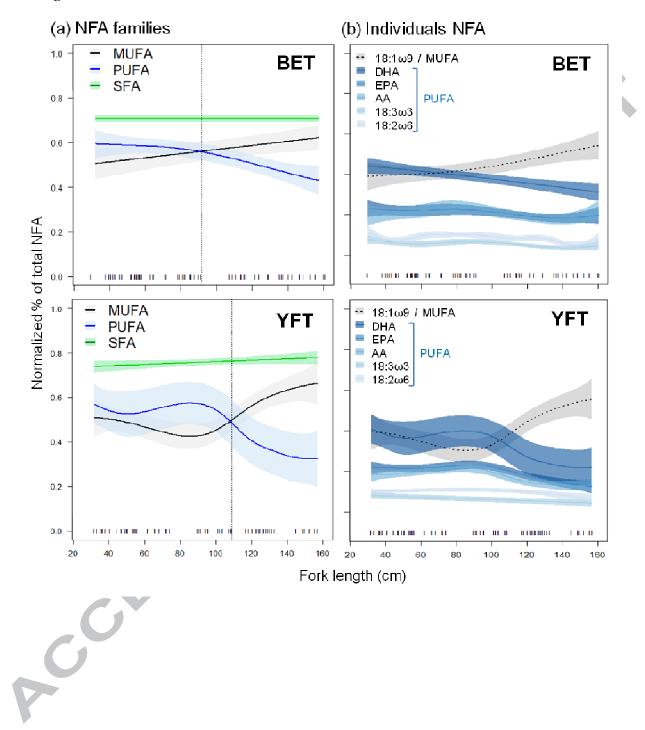




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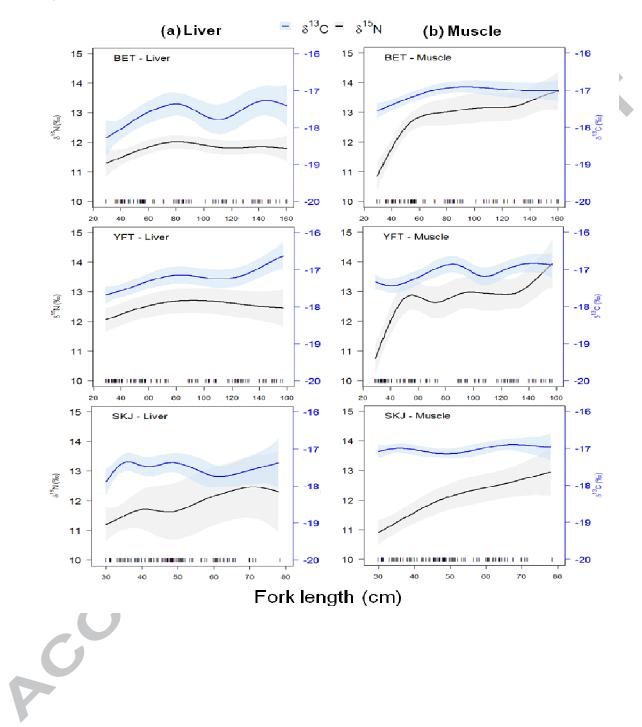
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1027 **Figure 4.**



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Table 1. Mean values (±standard deviation) of carbon and nitrogen isotopic data (δ^{13} C and δ^{15} N, ‰), total lipid content (TLC, % dry weight) and fatty acid profiles (neutral lipids as a % of total fatty acids) of the liver and muscle of skipjack tuna (SKJ), yellowfin tuna (YFT) and bigeye tuna (BET) in the Western-Central Indian Ocean, according to size class (FL≤100 cm and FL>100 cm). N is the number of samples for stable isotopes (SI) and neutral fatty acids (NFA) analyses. Superscripts represent significantly δ^{13} C, δ^{15} N, TLC and NFA differences among species (P<0.05 for Wilcoxon's test): ^B different from BET; ^S different from SKJ; ^Y different from YFT.

	,		Small tunas	· · ·			Large tunas (FL>100 cm)							
		Liver			Muscle		Live	er	Muscle					
	BET	SKJ	YFT	BET	SKJ	YFT	BET	YFT	BET	YFT				
N (SI II NFA)	33 26	43 46	29 24	33 28	43 47	29 23	27 24	23 19	26 21	23 19				
δ ¹³ C	-17.6 ± 0.4	-17.7 ± 0.4	-17.5 ± 0.4	-17.2 ± 0.3 ^{SY}	-17.0 ± 0.3 ^B	-17.0 ± 0.4 ^B	-17.4 ± 0.5	-17.1 ± 0.5	-17.0 ± 0.2 ^Y	-16.6 ± 0.3 ^B				
δ ¹⁵ N	11.8 ± 0.6 ^{SY}	10.8 ± 0.6 ^{BY}	11.4 ± 0.5 ^{BS}	12.5 ± 0.7 ^{SY}	11.4 ± 0.6 ^B	11.7 ± 0.6 ^B	11.8 ± 0.5 ^B	11.0 \pm 0.5 ^Y	13.4 ± 0.4 ^Y	12.4 ± 0.6 ^B				
TLC (% dw)	13.8 ± 5.0	14.4 ± 5.7	13.9 ± 8.3	2.7 ± 1.9	3.1 ± 3.5	2.0 ± 1.0	20.0 ± 8.8 ^B	27.3 ± 11.2 ^Y	2.2 ± 1.0	2.5 ± 1.3				
14:0	1.3 ± 0.9	1.7 ± 0.9 ^Y	1.2 ± 0.6 ^s	1.2 ± 1.2 ^s	0.6 ± 0.5 ^B	0.7 ± 0.5	1.1 ± 0.6	1.4 ± 0.8	0.8 ± 0.4	1.1 ± 0.9				
15:0	0.8 ± 0.4	1.0 ± 0.4	0.8 ± 0.4	0.5 ± 0.4	0.4 ± 0.4	0.4 ± 0.3	0.5 ± 0.4	0.5 ± 0.3	0.4 ± 0.3	0.6 ± 0.4				
16:0	30.6 ± 4.5 ^s	36.1 ± 7.0 ^B	33.4 ± 4.7	30.8 ± 8.0	32.8 ± 8.7	30.6 ± 9.4	$31.7 \pm 4.0^{\circ}$	35.7 ± 4.3 ^B	29.5 ± 8.8	29.6 ± 8.8				
17:0	1.4 ± 0.5	1.6 ± 0.7	1.6 ± 0.7	1.3 ± 0.3	1.2 ± 0.4	1.3 ± 0.3	1.0 ± 0.6	1.2 ± 0.5	1.2 ± 0.2	1.2 ± 0.4				
18:0	7.6 \pm 1.4 ^Y	8.3 ± 1.7 ^Y	10.2 ± 1.3 BS	12.0 ± 3.9 ^{SY}	13.8 ± 2.6 ^B	14.4 ± 3.1 ^B	8.3 ± 1.4	8.8 ± 1.9	12.4 ± 3.4	11.6 ± 2.5				
20:0	0.1 ± 0.1	0.1 ± 0.1	0.1 ± 0.1	0.4 ± 0.4	0.3 ± 0.4	0.3 ± 0.4	0.1 ± 0.1	0.2 ± 0.1	0.6 ± 0.4	0.6 ± 0.4				
22:0	0.0 ± 0.1	0.0 ± 0.0	0.0 ± 0.0	0.1 ± 0.3	0.2 ± 0.5	0.3 ± 0.4	0.0 ± 0.0	0.0 ± 0.0	0.4 ± 0.6	0.3 ± 0.5				
ΣSFA	41.9 ± 5.3 ^{sy}	48.9 ± 7.5 ^B	47.3 ± 5.2 ^B	46.4 ± 8.4	49.4 ± 9.1	48.0 ± 8.9	42.7 ± 5.0 ^Y	47.8 ± 4.3 ^B	45.2 ± 8.7	45.0 ± 8.3				
16:1ω7	2.9 ± 1.1	2.9 ± 1.2	2.4 ± 1.0	2.2 ± 1.7	1.6 ± 0.9	1.3 ± 1.0	3.9 ± 1.2	4.2 ± 1.0	2.2 ± 1.8	3.5 ± 2.1				
17:1ω7	0.7 ± 0.5 ^{SY}	0.5 ± 0.4 ^B	0.5 ± 0.4 ^B	0.7 ± 0.3 ^{SY}	0.5 ± 0.3^{B}	0.4 ± 0.3 ^B	0.8 ± 0.5	0.7 ± 0.4	0.6 ± 0.3	0.7 ± 0.4				
18:1ω9	16.4 ± 7.3 ^{SY}	10.4 ± 4.2 ^B	12.9 ± 8.8 ^B	13.1 ± 4.2 ^{SY}	7.8 ± 1.5 ^B	10.2 ± 3.2 ^B	23.8 ± 10.0	24.0 ± 6.0	15.0 ± 5.4	16.1 ± 2.6				
18:1ω7	3.2 ± 0.9 ^{SY}	2.2 ± 0.7 ^B	2.5 ± 1.0 ^B	2.8 ± 0.6 ^{SY}	1.9 ± 0.4 ^{BY}	2.5 ± 0.5^{BS}	3.6 ± 0.9	3.4 ± 0.4	3.2 ± 0.8	3.2 ± 0.6				
20:1ω9	1.0 ± 0.5 ^{SY}	0.3 ± 0.2 ^B	0.6 ± 0.5 ^B	1.0 ± 0.4 ^{SY}	0.3 ± 0.3^{BY}	0.6 ± 0.4^{BS}	1.0 ± 0.5	0.8 ± 0.3	0.9 ± 0.6	0.8 ± 0.4				
22:1ω9	0.1 ± 0.1	0.1 ± 0.1	0.1 ± 0.1	2.4 ± 2.8 ^S	0.9 ± 0.7^{BY}	2.2 ± 1.3 ^s	0.1 ± 0.1	0.1 ± 0.1	2.3 ± 3.0	2.0 ± 1.6				
24:1ω9	2.4 ± 2.2	2.0 ± 1.9	2.2 ± 2.3	0.3 ± 0.3	0.2 ± 0.4	0.4 ± 0.4	1.6 ± 1.5	0.9 ± 0.9	0.4 ± 0.5	0.4 ± 0.3				
ΣMUFA	26.8 ± 8.5 ^{sy}	18.2 ± 5.4 ^B	21.0 ± 10.2 ^B	22.6 ± 6.7 ^{SY}	13.3 ± 2.1 ^{BY}	17.5 ± 4.1 ^{BS}	34.8 ± 11.0	34.2 ± 6.9	24.6 ± 7.8	26.8 ± 3.9				
18:3ω3	0.5 ± 0.2 ^{SY}	0.2 ± 0.2 ^B	0.3 ± 0.1 ^B	0.3 ± 0.3	0.2 ± 0.2	0.3 ± 0.2	0.2 ± 0.1	0.3 ± 0.2	$0.3 \pm 0.4^{\circ}$	0.4 ± 0.3 ^B				
20:5ω3 (EPA)	4.9 ± 1.4	5.6 ± 1.9	5.1 ± 1.6	3.9 ± 0.7 ^s	5.0 ± 0.9^{BY}	3.8 ± 0.8 ^S	3.4 ± 1.5	2.5 ± 1.3	3.5 ± 0.9	3.5 ± 0.9				
22:5ω3	1.2 ± 0.5 ^s	0.9 ± 0.6 ^B	1.0 ± 0.8	0.7 ± 0.5	0.4 ± 0.5	0.9 ± 0.7	1.0 ± 0.6	1.1 ± 0.6	0.7 ± 0.6	0.8 ± 0.5				
22:6ω3 (DHA)	16.6 ± 5.3	19.1 ± 5.6	17.6 ± 4.6	20.1 ± 5.1 ^s	23.6 ± 6.2^{B}	22.7 ± 6.7	12.0 ± 5.6	9.4 ± 4.0	18.9 ± 6.3	16.7 ± 5.2				
18:2ω6	0.9 ± 0.4	0.8 ± 0.3 ^Y	1.1 ± 0.3 ^s	1.0 ± 0.4 ^{SY}	1.1 ± 0.3 ^{BY}	1.4 ± 0.4 ^{BS}	0.6 ± 0.4	0.7 ± 0.5	1.1 ± 0.4	1.3 ± 0.4				
18:3ω6	0.3 ± 0.4	0.3 ± 0.3	0.2 ± 0.2	0.1 ± 0.2	0.1 ± 0.2	0.2 ± 0.4	0.1 ± 0.1^{9}	0.2 ± 0.1 ^B	0.2 ± 0.2	0.3 ± 0.5				
20:2ω6	0.3 ± 0.2 ^{SY}	0.1 ± 0.1 ^B	0.2 ± 0.1 ^B	0.2 ± 0.2	0.2 ± 0.4	0.1 ± 0.3	0.1 ± 0.1	0.1 ± 0.1	0.2 ± 0.2	0.1 ± 0.2				
20:3ω6	0.1 ± 0.1 ^{SY}	0.1 ± 0.1 ^B	0.1 ± 0.1 ^B	0.0 ± 0.1	0.1 ± 0.1	0.1 ± 0.3	0.1 ± 0.1^{9}	0.1 ± 0.1 ^B	0.1 ± 0.1	0.1 ± 0.3				
20:4ω6 (AA)	4.5 ± 1.6	3.7 ± 1.1	4.2 ± 1.6	2.9 ± 0.6 ^S	4.0 ± 1.1 ^{BY}	2.7 ± 0.6 ^S	$3.4 \pm 1.8^{\circ}$	1.9 ± 0.8 ^B	3.1 ± 0.9	2.6 ± 1.0				
22:2ω6	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.2	0.1 ± 0.2	0.0 ± 0.1	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.1	0.1 ± 0.2				
22:4ω6	0.4 ± 0.3 ^S	0.2 ± 0.2 ^B	0.3 ± 0.3	0.1 ± 0.2	0.0 ± 0.1	0.0 ± 0.1	0.4 ± 0.3	0.3 ± 0.2	0.3 ± 0.5	0.2 ± 0.3				
22:5ω6	1.0 ± 0.3	1.2 ± 0.5	1.1 ± 0.3	1.4 ± 0.4 ^s	2.2 ± 0.7 ^{BY}	1.8 ± 0.5 ^s	0.8 ± 0.4	0.7 ± 0.2	1.5 ± 0.6	1.4 ± 0.6				
Σω3	23.2 ± 6.0	25.8 ± 7.0	24.0 ± 5.7	24.9 ± 5.3	29.3 ± 7.0	27.6 ± 7.4	16.5 ± 7.4	13.4 ± 5.7	23.4 ± 7.4	21.5 ± 5.9				
Σω6	7.5 ± 1.9	6.5 ± 1.6	7.1 ± 1.8	5.8 ± 1.0	7.7 ± 2.1	6.4 ± 1.4	5.5 ± 2.4	4.1 ± 1.3	6.4 ± 1.9	6.2 ± 1.7				
ΣΡυγΑ	30.7 ± 7.6	32.2 ± 8.3	31.1 ± 7.4	30.8 ± 5.8	37.0 ± 8.5	34.1 ± 8.1	22.0 ± 9.6	17.4 ± 6.9	29.8 ± 8.9	27.6 ± 7.4				

SFA: Saturated fatty acids; MUFA: Monounsaturated fatty acids; PUFA: Polyunsaturated fatty acids; EPA: Eicospentaenoic acid; DHA: Docosahexaenoic acid; AA: Arachinoic acid; w3: Omega-3 fatty acids; w6: Omega-6 fatty acids;

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Table 2. ANOVA and PERMANOVA results of tests for differences in total lipid content (TLC, % dw), fatty acid profiles (neutral lipids as a % of total fatty acids), and nitrogen and carbon isotope values (δ^{15} N and δ^{13} C, %₀) in the muscle and liver of tropical tuna. Tested factors are season (NEM, SWM, SIM, AIM), area (WCIO or MOZ), size class (FL≤100 cm or FL>100 cm), species (SKJ, YFT and BET) and their interactions when possible. NEM: North-Eastern Monsoon; SWM: South Western Monsoon; SIM: Spring Inter-Monsoon; AIM: Autumn Inter-Monsoon; WCIO: Western-Central Indian Ocean; MOZ: Mozambique channel; BET: Bigeye; SKJ: Skipjack; YFT: Yellowfin. df: freedom degree; sqs: squares, F and Pseudo F: statistics of the tests.

		TLC				FATTY ACIDS				δ ¹⁵ N				δ ¹³ C							
Tissue	Factors	df	Sums of sqs	Mean sqs	F	P-value	df	Sums of sqs	Mean sgs	Pseudo-F	P-value	df	Sums of sqs	Mean sqs	F	P-value	df	Sums of sqs	Mean sqs	F	P-value
	Season	3	16.2	5.4	1.1	0.333	3	0.30	0.10	2.43	0.006	3	106.5	35.5	56.8	0.000	3	1.0	0.3	4.9	0.003
	Area	1	1.5	1.5	0.3	0.568	1	0.13	0.13	3.20	0.021	1	150.3	150.3	240.3	0.000	1	0.0	0.0	0.0	0.877
	Size class	1	4.3	4.3	0.9	0.342	1	0.51	0.51	12.37	0.001	1	32.2	32.2	51.5	0.000	1	2.8	2.8	40.2	0.000
Muscle	Species	2	3.2	1.6	0.3	0.715	2	0.49	0.24	5.84	0.001	2	79.5	39.7	63.6	0.000	2	3.3	1.6	23.2	0.000
MUSCIE	Size class * Species	1	14.7	14.7	3.1	0.079	1	0.09	0.09	2.25	0.043	1	0.7	0.7	1.1	0.296	1	0.1	0.1	1.2	0.276
	Size class * Area	1	60.9	60.9	12.9	0.000	1	0.07	0.07	1.78	0.120	1	1.7	1.7	2.7	0.103	1	0.5	0.5	6.4	0.012
	Species * Area	2	15.2	7.6	1.6	0.201	2	0.22	0.11	2.66	0.011	2	6.5	3.2	5.2	0.006	2	0.0	0.0	0.3	0.735
	Residuals	254	1198.9	4.7			176	7.32	0.04	0.80		242	151.4	0.6			242	17.0	0.1		
	Season	3	1101.5	367.2	7.8	0.000	3	0.30	0.10	2.85	0.006	3	159.5	53.2	139.0	0.000	3	1.1	0.4	2.5	0.060
	Area	1	199.8	199.8	4.2	0.041	1	0.19	0.19	5.28	0.002	1	109.5	109.5	286.2	0.000	1	1.2	1.2	8.1	0.005
	Size class	1	2508.1	2508.1	53.0	0.000	1	1.27	1.27	35.70	0.001	1	0.0	0.0	0.1	0.760	1	5.3	5.3	34.8	0.000
Liver	Species	2	230.4	115.2	2.4	0.090	2	0.54	0.27	7.56	0.001	2	53.9	26.9	70.4	0.000	2	0.5	0.2	1.5	0.215
LIVCI	Size class * Species	1	153.6	153.6	3.2	0.073	1	0.03	0.03	0.97	0.334	1	3.0	3.0	7.7	0.006	1	0.0	0.0	0.2	0.646
	Size class * Area	1	689.0	689.0	14.6	0.000	1	0.22	0.22	6.31	0.002	1	0.0	0.0	0.0	0.849	1	0.4	0.4	2.4	0.121
	Species * Area	2	95.9	48.0	1.0	0.365	2	0.05	0.03	0.77	0.583	2	8.7	4.4	11.4	0.000	2	0.6	0.3	1.9	0.149
	Residuals	241	11407.1	47.3			168	5.97	0.04	0.70		227	86.8	0.4			227	34.4	0.2		

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Table 3. Biochemical trophic niches indices (INes for stable isotopes and FANes for neutral fatty acids profiles) calculated for small (≤ 100 cm) and large tuna (> 100 cm), in Western-Central Indian Ocean (WCIO) and Mozambique Channel (MOZ) separately. Values correspond to the degree of overlap among species with 0 = no overlap and 1 = complete overlap. BET: Bigeye; SKJ: Skipjack; YFT: Yellowfin.

Area	Size class FL≤100cm FL>100cm	Tissue Muscle Liver	Groups BET, SKJ BET, YFT SKJ, YFT BET, SKJ BET, YFT	FANes (3D) 0.12±0.07 0.62±0.08 0.07±0.10 0.30±0.05	INes (2D) 0.09±0.03 0.37±0.03 0.81±0.05 0.43±0.07
WCIO		Liver	BET, YFT SKJ, YFT BET, SKJ BET, YFT	0.62±0.08 0.07±0.10 0.30±0.05	0.37±0.03 0.81±0.05
wcio		Liver	SKJ, YFT BET, SKJ BET, YFT	0.07±0.10 0.30±0.05	0.81±0.05
wcio			BET, SKJ BET, YFT	0.30±0.05	
WCIO			BET, YFT		0.43±0.07
	FL>100cm		-	0 00 0 00	
	FL>100cm			0.60±0.08	0.81±0.04
	FL>100cm		SKJ, YFT	0.51±0.07	0.49±0.03
	1 L>1000m	Muscle	BET, YFT	0.71±0.06	0.00±0.00
		Liver	BET, YFT	0.53±0.14	0.34±0.06
			BET, SKJ	0.06±0.08	0.10±0.02
		Muscle	BET, YFT	0.29±0.13	0.36±0.06
MOZ	all sizes		SKJ, YFT	0.41±0.07	0.63±0.05
WOZ		Liver	BET, SKJ	0.08±0.04	0.15±0.04
			BET, YFT	0.34±0.22	0.44±0.13
			SKJ, YFT	0.40±0.06	0.73±0.05
0					

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Highlights

• Sympatric tropical tuna (bigeye, yellowfin and skipjack tuna) were sampled in the Indian Ocean;

- Trophic niche partitioning was quantitatively assessed using time-integrated tracers;
- Intra- and inter-species competition in three tropical tuna species were detected;
- Diet shifts at ~ 100 cm FL were detected in the larger tuna species (bigeye and yellowfin);
- • New insights into the mechanisms involved in tropical tuna coexistence were provided.

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