

## Chemical contaminants (trace metals, persistent organic pollutants) in albacore tuna from western Indian and south-eastern Atlantic Oceans: Trophic influence and potential as tracers of populations \*

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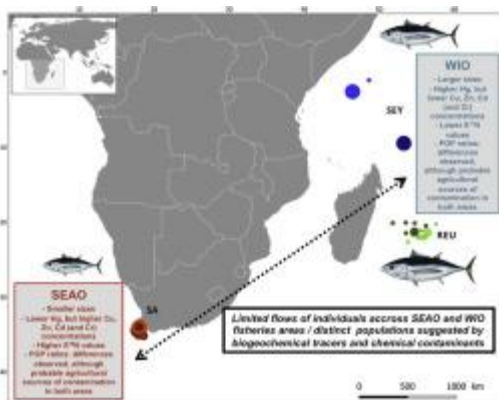
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### Abstract :

Albacore tuna (*Thunnus alalunga*) is a highly commercial fish species harvested in the world's Oceans. Identifying the potential links between populations is one of the key tools that can improve the current management across fisheries areas. In addition to characterising populations' contamination state, chemical compounds can help refine foraging areas, individual flows and populations' structure, especially when combined with other intrinsic biogeochemical (trophic) markers such as carbon and nitrogen stable isotopes. This study investigated the bioaccumulation of seven selected trace metals – chromium, nickel, copper (Cu), zinc (Zn), cadmium (Cd), mercury (Hg) and lead – in the muscle of 443 albacore tunas, collected over two seasons and/or years in the western Indian Ocean (WIO: Reunion Island and Seychelles) and in the south-eastern Atlantic Ocean (SEAO: South Africa). The main factor

that explained metal concentration variability was the geographic origin of fish, rather than the size and the sex of individuals, or the season/year of sampling. The elements Cu, Zn, Cd and Hg indicated a segregation of the geographic groups most clearly. For similar sized-individuals, tunas from SEAO had significantly higher concentrations in Cu, Zn and Cd, but lower Hg concentrations than those from WIO. Information inferred from the analysis of trophic markers ( $\delta^{13}\text{C}$ ,  $\delta^{15}\text{N}$ ) and selected persistent organic pollutants, as well as information on stomach contents, corroborated the geographical differences obtained by trace metals. It also highlighted the influence of trophic ecology on metal bioaccumulation. Finally, this study evidenced the potential of metals and chemical contaminants in general as tracers, by segregating groups of individuals using different food webs or habitats, to better understand spatial connectivity at the population scale. Limited flows of individuals between the SEAO and the WIO are suggested. Albacore as predatory fish also provided some information on environmental and food web chemical contamination in the different study areas.

### Graphical abstract



### Highlights

► 443 albacore tunas were analysed for their muscle concentrations in trace metals. ► Cu, Zn, Cd and Hg were the elements that most segregated the different groups. ► Trophic markers and organic contaminants confirmed the segregation observed. ► Differences in trace metal bioaccumulation were linked to fish trophic ecology. ► Inorganic elements can trace populations exploiting different food webs.

**Keywords:** Top predator, Bioaccumulation, Inorganic elements, Organic contaminants, Stable isotopes, Intrinsic markers

## 80 **1. Introduction**

81 Trace metals are inorganic elements that are naturally present on Earth. They have been used over a  
82 long time due to their properties such as mechanical resistance, electric or thermic conductivity, or  
83 biocidal properties. Since the industrial age, their increasing use in human activities has led to  
84 continuous release and contamination of all environmental compartments. Thus, trace metals are  
85 currently released into the environment from both natural (e.g., volcanism) and anthropogenic sources  
86 (e.g., industrial, urban, or agricultural). They reach the ocean through river inputs and atmospheric  
87 depositions, the atmospheric pathway at times transporting trace metals very far from the emission  
88 source (Mason, 2013). Some trace metals are recognized to be essential for organisms and form the  
89 basis of biochemicals, such as enzymes. However, they perform optimally in a relatively low range of  
90 concentrations and become either deficient or toxic at very low or high concentrations (e.g., copper  
91 (Cu), zinc (Zn)). Alternatively, some elements have no known biological role and are recognized for  
92 their toxicity towards most organisms (e.g., cadmium (Cd), mercury (Hg), lead (Pb)), even at low  
93 concentrations (Mason, 2013). Taxa- and species-specific regulation mechanisms of metals have been  
94 described for both essential and non-essential elements, influencing their storage or elimination by  
95 organisms (Wang and Rainbow, 2010). Their transfer between biogeochemical compartments, their  
96 bioaccumulation in organisms and/or biomagnification in food webs finally depend on the speciation  
97 of elements, which determines their bioavailability in both abiotic (habitat) and biotic (food sources)  
98 environments of organisms (Neff, 2002; Rainbow, 2002). Contrary to trace metals, persistent organic  
99 pollutants (POPs) such as polychlorinated biphenyls (PCBs) and dichlorodiphenyl-trichloroethane  
100 (DDT) are almost exclusively from anthropogenic origin. They are manufactured for, and are used  
101 widely by commercial sectors such as industry and agriculture, which released them into the  
102 environment. POPs volatilize easily and can be transported through the atmosphere over wide  
103 distances, and deposited in areas far from their point of emission (Jones and de Voogt, 1999; Bogdal et  
104 al., 2013). Moreover, these chemicals or their metabolites have a strong persistence in ecosystems, and  
105 they are well documented to biomagnify in food webs and to be toxic for organisms (e.g., Verreault et  
106 al., 2008).

107 In aquatic organisms and more specifically in marine top predators such as marine mammals, seabirds  
108 or large pelagic fish, the trophic pathway represents the main pathway for the intake of both trace  
109 metals and POPs (e.g., Fisk et al., 2001; Wang, 2002; Mathews and Fisher, 2009). Individual trophic  
110 ecology can thus largely affect the contaminant concentrations measured in a given organism. This  
111 includes feeding area, trophic level, or the type of prey consumed (Lahaye et al., 2005; Choy et al.,  
112 2009; Ramos et al., 2013; Teffer et al., 2014), some prey accumulating more contaminants than others,  
113 for instance (e.g., Bustamante et al., 1998; Pulster et al., 2005). Understanding the mechanisms leading  
114 to bioaccumulation of contaminants and/or interpreting contaminant concentrations measured in biota  
115 thus requires a good knowledge of the consumers' feeding habits and ecology.

116 Over the last decades, stable isotope analysis (SIA) of carbon ( $\delta^{13}\text{C}$ ) and nitrogen ( $\delta^{15}\text{N}$ ) in biological  
117 tissues has largely developed to study the trophic ecology of marine organisms such as top predators  
118 (Kelly et al., 2000; Newsome et al., 2010). It effectively represents an alternative or complementary  
119 tool to the traditional methods of dietary studies such as the analysis of stomach contents.  $\delta^{13}\text{C}$  values  
120 are generally considered a conservative tracer of the primary producer at the base of the food web  
121 supporting consumers, and consequently a tracer of their foraging habitat (France, 1995; Hobson,  
122 1999). Alternatively,  $\delta^{15}\text{N}$  values are generally used as a proxy of their trophic position (Post, 2002),  
123 although the interpretation of  $\delta^{15}\text{N}$  values should be food web-specific due to high variability in  $\delta^{15}\text{N}$   
124 baseline values between ecosystems (Sherwood and Rose, 2005; Ménard et al., 2007; Chouvelon et al.,  
125 2012a).

126 More recently, the combined use of parameters (i.e. SIA, fatty acid profiles, chemical contaminants  
127 including metals and/or POPs, etc.) measured in consumers' tissues as ecological tracers of marine  
128 predators' trophic position, dietary preferences or foraging areas, has drastically increased (e.g., Fisk et  
129 al., 2002; Iverson et al., 2004; Lahaye et al., 2005; Krahn et al., 2007; Mendez-Fernandez et al., 2013,  
130 2017; Cresson et al., 2015). Based on the assumption "I am what I eat", such ecological tracers are  
131 used to encompass the inherent difficulty of direct at-sea observations for these species. When  
132 combined to other tools such as geolocation devices and tags, or biological data such as genetic and  
133 morphological data, they may also be helpful in unravelling top predators' foraging strategies over  
134 time, spatial dynamics (migrations), or the use of different resources or habitats by the different  
135 populations of a given species (Ramos and González-Solís, 2012; Chouvelon et al., 2014; Cresson et  
136 al., 2015). To the best of our knowledge, such a combined approach has not yet been used in tunas,  
137 while these top predator fish represent an important commercially species harvested in the world's  
138 Oceans.

139 The commercial catch of albacore tuna (*Thunnus alalunga*) is the highest globally among the  
140 temperate tuna species and has contributed around 6% by weight of global tuna catches over the last  
141 decade (FAO, 2016). The state and the assessment of albacore stocks vary geographically. The  
142 estimated stock assessment has long been considered over-exploited in the South Atlantic Ocean  
143 (SAO), and not over-exploited in the Indian Ocean (IO) (ICCAT, 2016; IOTC, 2016), although large  
144 uncertainties remain (Guan et al., 2016) due to limited and low-quality data. Moreover, this species  
145 has been poorly studied in the IO in comparison with other Oceans (Nikolic et al., 2016). In the Pacific  
146 and Atlantic Oceans, the migration of albacore tunas between hemispheres is considered negligible  
147 (Nakamura, 1969; Lewis, 1990; Arrizabalaga et al., 2004), influenced by global intra-ocean circulation  
148 (e.g., oceanic gyres) that drives the oceanography of the northern and southern hemispheres. Due to  
149 the absence of such structures in the IO, albacore has been managed as one unique stock in this region  
150 (Chen et al., 2005), distributed between 5°N and 45°S. Additionally, the tip of South Africa has long  
151 been considered an impenetrable barrier for marine animals (Briggs, 1974). However, several recent  
152 studies have demonstrated that gene flow occurs between the SAO and western Indian Ocean (WIO)

153 for a number of species, including large vertebrates such as the green turtle (Bourjea et al., 2007), the  
154 scalloped hammerhead sharks (Duncan et al., 2006), and the tropical bigeye tuna (Durand et al., 2005).  
155 The question of connectivity between these two oceans for the albacore tuna is still pending. Indeed, a  
156 significant amount of juveniles of this species are caught each year in South African waters below  
157 30°S, and in the WIO between 30°S and 40°S. Assessing the origin and the fate of these southern  
158 juveniles, and the potential links (gene flows, individual migrations, etc.) that may exist between the  
159 SAO and WIO stocks is a crucial issue for sustainable management of albacore tunas. In addition,  
160 trace element data on the albacore tuna remain scarce (e.g., Das et al., 2000 in the north Atlantic; Chen  
161 et al., 2014 and Hisamichi et al., 2010 in the north Pacific; Storelli et al., 2002 in the Mediterranean  
162 Sea), and more specifically in the study areas (e.g., Bodin et al., 2017).

163 In this general context, the main objectives of the present study were: 1) to characterise the  
164 contamination in several trace metals of albacore tunas caught in the WIO (Reunion Island and  
165 Seychelles) and in the south-eastern AO (SEAO; South Africa); 2) to determine the main factors  
166 influencing the bioaccumulation of the selected inorganic elements in tunas (e.g., effects of size, sex,  
167 season/year of sampling, geographic origin and food webs exploited); 3) to assess the potential of  
168 these trace metals as biogeochemical tracers of albacore tuna populations; 4) to evaluate the input of  
169 SIA (i.e.  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  values, as trophic markers) and organic contaminants (i.e. selected POP ratios,  
170 as additional chemical tracers) for discriminating fish populations (i.e. do they corroborate,  
171 complement, or contradict the information provided by trace metals?). In addition, this study provides  
172 general information on the contamination state of the food webs supporting tunas in the different study  
173 areas.

174

## 175 **2. Materials and methods**

### 176 **2.1. Sampling**

177 A total of 443 albacore tunas were collected from June 2013 to July 2014 by commercial fisheries in  
178 three distinct areas in the WIO and the SEAO: 128 were caught by semi-industrial drifting longliners  
179 around La Réunion Island (REU), 118 by tuna purse-seine fisheries operating in Seychelles waters  
180 (SEY), and finally 197 by tuna pole fisheries in waters of South Africa (SA) (Fig. 1). Depending on  
181 the area considered, sampling was performed over two seasons (S1: austral summer November-  
182 December, i.e. potential reproduction season of tunas; S2: austral winter April-July, i.e. potential  
183 feeding period or post-reproduction period; Nikolic et al. 2016), and/or years (Y1: 2013; Y2: 2014), in  
184 order to account for the variability of environmental conditions and for the different life history traits  
185 of albacore tuna. All fishes analysed in the present study were already dead by the time of sampling.  
186 For each individual, the fork length (from the tip of the snout to the fork of the tail, FL in cm) was  
187 recorded as well as the sex. Finally, around 15g of white muscle (without skin) were taken from the

188 dorsal musculature of each fish (sampled behind the head under the dorsal spine). All samples were  
189 frozen at  $-20^{\circ}\text{C}$ , freeze-dried and ground into a fine powder until further chemical analyses.

## 190 **2.2. Trace metal analysis**

191 Trace metal analysis was performed on all muscle samples of albacore tuna ( $N=443$ ).

192 Total Hg analyses were carried out individually with an Advanced Mercury Analyser (ALTEC AMA  
193 254), directly on subsamples (10–100 mg) of homogenised dry muscle tissue (untreated powder). For  
194 Hg determination, the metal is evaporated by progressive heating up to  $800^{\circ}\text{C}$ , then held under an  
195 oxygen atmosphere for 3 min, and finally amalgamated on a gold net. Afterwards, the net is heated to  
196 liberate the collected Hg, which is finally measured by atomic absorption spectrophotometry. Hg  
197 analyses were run according to a thorough quality control programme including the analysis of a  
198 certified reference material (CRM) IAEA-142 (mussel homogenate; International Atomic Energy  
199 Agency/IAEA).

200 For the analysis of the six other inorganic elements (Chromium (Cr), Nickel (Ni), Cu, Zn, Cd and Pb),  
201 subsamples ( $\sim 200$  mg) of homogenised dry muscle tissue were first digested with a mixture of 65%  
202  $\text{HNO}_3$ ,  $\text{H}_2\text{O}_2$  and milli-Q quality water, kept for a few hours at room temperature, and heated at  $85^{\circ}\text{C}$   
203 on a heating block for 6h. After the mineralization process, each sample was completed with a known  
204 quantity of internal standards and diluted to 50 ml with milli-Q quality water. Samples were re-diluted  
205 when necessary before analyses. Elemental analyses were finally carried out by Inductively Coupled  
206 Plasma-Mass Spectrometry (ICP-MS; Thermo X-Series I) and included the analysis of the two CRMs:  
207 IAEA-407 (whole fish homogenate; IAEA) and DOLT-3 (dogfish liver; National Research Council  
208 Canada).

209 Analytical performance details for the seven trace metals are summarized in Table 1. Metal  
210 concentrations are expressed in  $\text{mg}\cdot\text{kg}^{-1}$  dry mass (dm).

## 211 **2.3. Analysis of additional tracers**

212 From the total number of muscle samples of albacore collected for trace metal analysis, 335 were  
213 selected for SIA (96 from REU, 63 from SEY and 176 from SA), and 86 for POP analysis (43 from  
214 REU and 43 from SA).

215 SIA was performed according to the methods of Bodin et al. (2009) and Sardenne et al. (2016).  
216 Briefly, around 2 g of dried and ground samples were treated by an Accelerated Solvent Extraction  
217 system with dichloromethane.  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  were analysed together on dried lipid-free samples using  
218 a Delta V Advantage isotope ratio mass spectrometer interfaced to a Flash EA 1112 elemental analyser  
219 (Thermo Scientific). Analytical precision for both  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  was  $<0.15\%$  based on replicate  
220 measurements of internal laboratory standards (acetanilide and peptone, Thermo Scientific).

221 Details for POP analysis are found in Munschy et al. (2016). The concentrations and profiles of 18  
222 PCB congeners (namely, the 6 indicators congeners CB-28; -52; -101; -138; -153; -180, and the 12  
223 dioxin-like congeners -77; -81; -105; -114; -118; -123; -126; -156; -157; -167; -169; -189) were  
224 determined together with DDT compounds (namely, *p,p'*-DDT and *o,p'*-DDD, and its main  
225 degradation products *o,p'*-DDD, *p,p'*-DDD, and *p,p'*-DDE).

## 226 **2.4. Data treatment**

227 Statistics were only applied on trace metals for which almost 80% of individuals presented raw  
228 concentrations above the limits of quantification (LQ). These elements are further called  
229 “elements >LQ”, and Pb and Ni were thus excluded from data treatment and statistical analyses (Table  
230 1). All figures and statistics were performed using the software R version 2.15.2 (R Development Core  
231 Team, 2012), except the map of sampling locations (Fig. 1) that was done using ArcGIS software  
232 (www.arcgis.com).

233 In the case of POPs, whose detailed results were recently published (Munschy et al., 2016), only three  
234 ratios were considered here for comparison with trace metals, due to their ability to trace potential  
235 contamination sources:

236 - Sum DDTs/Sum PCBs, where “Sum PCBs” corresponds to the sum of the concentrations of the 18  
237 PCB congeners identified in albacore muscle samples, and “Sum DDTs” to the sum of the two parent  
238 DDT compounds and their three main metabolites. This ratio can generally distinguish the influence of  
239 sources from agricultural origin (when largely >1) vs. sources from industrial origin (Yogui et al.,  
240 2003; Lailson-Brito et al., 2011).

241 - *p,p'*-DDT/*p,p'*-DDE. This ratio can be used as a tracer of the residence time and degree of  
242 degradation of DDT in the environment, and consequently to distinguish new vs. old DDT sources  
243 (Suárez et al., 2013).

244 - *o,p'*-DDT/*p,p'*-DDT. This ratio can trace the DDT origin (close to the technical mixture (0.2–0.3), or  
245 not; Kalantzi et al., 2001).

### 246 *2.4.1 Classical statistics*

247 First, mean values in body sizes (FL), trace metal concentrations,  $\delta^{15}\text{N}$  and  $\delta^{13}\text{C}$  values and POP ratios  
248 were submitted to parametric Student t-tests or non-parametric Mann-Whitney-Wilcoxon tests to  
249 assess i) differences between males and females (within a given season/year and in a given area), and  
250 ii) differences between seasons/years (in a given area). Differences between geographic areas were  
251 checked using Kruskal-Wallis tests (KW) followed by multiple comparison tests with Holm’s  
252 adjustment method, because data never satisfied parametric conditions. All of the mean comparison  
253 tests were applied on raw data as well as size-normalised data (i.e. normalised to a 95cm FL  
254 individual) to account for the high variability in fish sizes between areas (Table 2). The level of  
255 significance for statistical analyses was always set at  $\alpha = 0.05$ .

#### 256 2.4.2 Generalized modelling

257 To further estimate the effect of potentially confounding explanatory variables on the variability of  
258 trace metal concentrations (i.e. body size, trophic position (through  $\delta^{15}\text{N}$ ), sex, season/year of  
259 sampling, and area), generalized modelling (considering each trace element separately) was applied  
260 using the mgcv package in R. Data effectively showed a marked departure from normality in most  
261 cases, preventing the application of multiple linear regressions.

262 Assuming potential migrations of fishes between areas and seasons and/or years, the two factors  
263 “area” and “season/year of sampling” were not considered separately in the models, as it could have  
264 been done in the case of true “nested data” (e.g., three areas considered, in which two seasons and/or  
265 years have been sampled). Indeed, with regard to potential flows of individuals between areas and  
266 seasons, S1 and S2 in the same area may be totally disconnected. The combination “Area/Season  
267 and/or Year of sampling” (Area/S-Y) was thus considered a single factor in the models, including 6  
268 modalities: REU/S1, REU/S2, SEY/S2-Y1, SEY/S2-Y2, SA/S1 and SA/S2. FL (when considering  
269 size-related trends; N=443 individuals) or  $\delta^{15}\text{N}$  values (when considering trophic position-related  
270 trends; N=335) were treated as the continuous explanatory variable in the models, while Sex and  
271 Area/S-Y were treated as categorical explanatory variables and/or added as an interaction term (for  
272 Area/S-Y).

273 Generalized Additive Models (GAMs), representing flexible non-parametric generalizations of  
274 (multiple) linear regressions, were first tested. They can capture and model both linear and complex  
275 non-linear relationships (Zuur, 2012), and as such they are said to be “data-driven” methods. GAMs  
276 were fitted on raw concentrations, with a Gamma distribution and a log link function. Applying GAMs  
277 on log-transformed concentrations directly (with a Gaussian distribution and an identity link function)  
278 substantially improved the models, as it is generally the case when dealing with contaminant data such  
279 as trace metals (e.g., Pierce et al., 2008; Mendez-Fernandez et al., 2013; Chauvelon et al., 2014).  
280 However, as the modelled trends with FL or  $\delta^{15}\text{N}$  appeared to be linear, Generalized Linear Models  
281 (GLMs) were finally applied. As in the case of GAMs, GLMs were first fitted on raw concentrations  
282 with a Gamma distribution and a log link function. However, Gaussian GLMs on log-transformed  
283 concentrations with an identity link function also gave slightly better results and were thus kept.

284 Differences between sexes were tested in the first models, but since there was no effect for most of the  
285 trace metals and conditions (see also boxplots and classical statistics; Fig. 2), we removed the factor  
286 Sex from the final GLMs. The general forms of the models performed for each trace metal were thus:

287 -  $\text{Log}[\text{Trace metal}] \sim \text{FL} + \text{Area/S-Y} [+ \text{interaction term FL:Area/S-Y}]$

288 -  $\text{Log}[\text{Trace metal}] \sim \delta^{15}\text{N} + \text{Area/S-Y} [+ \text{interaction term } \delta^{15}\text{N:Area/S-Y}]$

289 Results (fitted values) of the models were plotted on observed (log-transformed) data. Final models  
290 parameters (estimates, p-values, etc.) are given in supplementary material (Appendix A and  
291 Table A.1). For each model, we retained the variables that improved the relative goodness of fit in the  
292 GLM (most parsimonious model) based on the Akaike Information Criterion (AIC). When the AIC



293 was not significantly different between the last two nested models, the simplest model was preferred  
294 (Zuur et al., 2007). Finally, a model validation was systematically applied by checking normality and  
295 homogeneity in model residuals, with no violation of independence (Zuur et al., 2007; Zuur, 2012).  
296 Models not complying with all of these assumptions were not presented (i.e. for Cd). The percentage  
297 of total deviance explained (DE) by each final model was calculated as follows: Explained deviance =  
298 ((Null model deviance - final model residual deviance)/Null model deviance)\*100, with the null model  
299 that only contained the intercept terms (Mendez-Fernandez et al., 2013).

### 300 2.4.3 *Multivariate analyses*

301 Different principal component analyses (PCAs) were performed to: i) identify relationships among the  
302 different types of chemical components analysed and included in the different PCAs (trace metals,  
303 and/or trophic markers ( $\delta^{13}\text{C}$ ,  $\delta^{15}\text{N}$ ), and POP ratios); ii) apprehend global profiles of contamination of  
304 the different groups of individuals analysed (i.e. contaminants considered together, and not separately  
305 as in previous data treatment). Moreover, these PCAs assessed of the potential of individual variables  
306 and/or of the combination of variables to trace the different groups of individuals. As such, the PCA 1  
307 included all individuals for which trace metals were analysed (N=443); the PCA 2 included  
308 individuals for which both trace metals and trophic markers ( $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  values) were analysed  
309 (N=335); and the PCA 3 included individuals analysed for trace metals and POP ratios (N=86). All  
310 PCAs were based on correlation matrices and normalised data for each variable included (i.e. data  
311 centred and divided by the standard deviation).

312 Finally, to further evaluate the potential of trace metals and/or their combination with trophic markers  
313 (i.e. larger datasets, and datasets including the three areas) in discriminating individuals from different  
314 geographic origin, discriminant analyses (DAs) were performed. DAs were also performed on datasets  
315 that included only the two close areas, REU and SEY in the WIO, to evaluate how they discriminate.  
316 As for PCAs, DAs used normalised data. Correlation coefficients between the discriminant functions  
317 and each of the original variables included in DAs – also called canonical correlation coefficients (ccc)  
318 – were used to assess the relative importance of each variable for the discrimination of individuals  
319 along the axes (Zuur et al., 2007).

320

## 321 **3. Results**

322 Characteristics of the albacore tunas considered in this study and the mean trace metal concentrations  
323 ( $\pm$  standard deviation) measured in the muscle are given in Table 2. 100% of individuals from REU  
324 and 88% from SEY had FL  $\geq$ 90 cm (min–max: 90–113 cm and 82–108 cm for REU and SEY,  
325 respectively), and females were significantly smaller than males ( $p < 0.001$  for both REU/S1 and  
326 REU/S2;  $p = 0.022$  and  $p = 0.001$  for SEY/S1-Y1 and SEY/S2-Y2, respectively). Individuals from SA  
327 (min–max: 74–118 cm FL) were smaller on average than those from REU and SEY (Table 2), with

328 most of the individuals having FL <90 cm (72% of individuals) and the majority of the fish with  
329 FL ≥90 cm that were captured in S2 (Table 2). No significant size difference between sexes was found  
330 for SA individuals ( $p = 0.608$  and  $p = 0.798$  for SA/S1 and SA/S2, respectively).

331 Overall, Cr individual concentrations varied between  $0.32 \text{ mg.kg}^{-1} \text{ dm}$  and  $0.74 \text{ mg.kg}^{-1} \text{ dm}$  (min–  
332 max), while the range of values was  $<0.03\text{--}0.28 \text{ mg.kg}^{-1} \text{ dm}$  for Ni (i.e.  $<\text{LQ}\text{--}0.28 \text{ mg.kg}^{-1} \text{ dm}$ ),  $0.38\text{--}$   
333  $3.67 \text{ mg.kg}^{-1} \text{ dm}$  for Cu, and  $6.7\text{--}77.5 \text{ mg.kg}^{-1} \text{ dm}$  for Zn. For non-essential elements, concentrations  
334 varied between  $<0.05 \text{ mg.kg}^{-1} \text{ dm}$  and  $1.55 \text{ mg.kg}^{-1} \text{ dm}$  for Cd,  $0.386\text{--}4.665 \text{ mg.kg}^{-1} \text{ dm}$  for Hg, and  
335  $<0.05\text{--}0.29 \text{ mg.kg}^{-1} \text{ dm}$  for Pb (Table 2).

### 336 **3.1. Differences in trace elements between sexes, seasons/years, and areas**

337 Among the five elements analysed  $>\text{LQ}$ , with the raw data considered, significant differences between  
338 sexes were only observed in one case for Zn (with females on average presenting slightly higher  
339 values than males in REU/S2) and in some cases for Hg (slightly higher concentrations measured in  
340 males than in females in REU/S2, SEY/S2-Y2, SA/S1). Some differences were also observed for Cr  
341 (with females presenting slightly higher values than males) when FL-normalised data were used  
342 (Table 2; Fig. 2). Differences between seasons/years of sampling were more often significant than  
343 between sexes and were very similar whether raw or FL-normalised data were used, although no  
344 general pattern appeared. Indeed in REU, there was no significant difference between seasons for Hg,  
345 but significantly lower concentrations in Cr, Cu, Zn and Cd were found in individuals from S1 than in  
346 individuals from S2. In SEY, significantly lower concentrations in Cr and Hg, associated with slightly  
347 higher concentrations in Cu, Zn and Cd were found in individuals captured the first year (S2-Y1)  
348 relative to those captured at the second year (S2-Y2). Finally in SA, significantly higher  
349 concentrations in Cu, Zn and Cd, along with significantly lower concentrations in Cr and Hg were  
350 measured in individuals from S1 compared with individuals from S2 (Table 2; Fig. 2). Concerning  
351 geographical differences, for all elements except Cr, the significant differences observed also remained  
352 similar whether raw or FL-normalised data were considered. Trace metal concentrations measured in  
353 individuals from the three areas were always significantly different in Cu, Zn and Hg, in line with the  
354 order of concentrations:  $\text{SA} > \text{REU} > \text{SEY}$  for Cu and Zn; and  $\text{REU} > \text{SEY} > \text{SA}$  for Hg. For Cd,  
355 mean concentrations differed significantly only between individuals captured in the WIO and those  
356 caught in the SEAO (i.e. similar and lower concentrations on average in REU and SEY vs. SA). Cr  
357 showed no significant difference between areas when considering raw data, while the order of  
358 concentrations was  $\text{SA} > \text{REU} > \text{SEY}$  when considering size-normalised data (Table 2; Fig. 2).  
359 Finally, although statistical tests could not be applied due to a large number of individuals below LQ  
360 for Ni and Pb, slightly higher Ni concentrations were observed in individuals from SA (Table 2). In  
361 fact, despite the low concentrations observed, most individuals from SA presented Ni concentrations  
362  $>\text{LQ}$ , while this element was poorly detected in the muscle of individuals from REU and SEY.

363 Considering the five trace elements in a single analysis, the PCA 1 corroborated the low influence of  
364 sex in explaining metal concentrations variability, while the sampling season and/or sampling year  
365 (especially for SA individuals), as well as the geographic origin, were much more significant factors.  
366 As such, the first two axes of the PCA 1 explained 70.1% of the variability observed in the dataset.  
367 The elements Cu, Zn, Cd and Hg (in the order), and Cr to some extent (i.e. for SA individuals),  
368 contributed the most to the cluster of individuals belonging to different groups, and consequently, to  
369 the dispersion of individuals captured in the different areas and/or sampled at different seasons/years  
370 (Fig. 3). With the five trace metals >LQ included, the DAs 1 and 2 further evidenced the significance  
371 of Cu, Zn, Cd and Hg (in that order) in clearly classifying individuals into the three areas. Cu also  
372 appeared to be a significant discriminant element between individuals from REU vs. SEY (Table 3).

### 373 **3.2. Size (FL)- and $\delta^{15}\text{N}$ -related trends**

374 In the final GLMs performed, the total deviance explained (DE) in log-transformed metal  
375 concentration variability was quite high, ranging between 45.9% (for  $\delta^{15}\text{N}$ -related trends in Hg  
376 variability) and 84.3% (for FL-related trends in Cu variability). The AIC scores were systematically  
377 lower in models performed with FL as single continuous explanatory variable than in those with  $\delta^{15}\text{N}$   
378 (details in Appendix A and Table A.1).

379 Considering final models with FL, the effect of the factor Area/S-Y was always highly significant. The  
380 effect of size (FL) was significant for Cr and Cu only ( $p=0.010$  and  $p<0.001$  respectively), despite a  
381 low p-value also observed for Hg ( $p=0.075$ ). The interaction term (i.e. model assuming that the form  
382 of the relationship between (log-transformed) concentrations and FL can be different between  
383 modalities of the factor Area/S-Y) was significant for Cr and Hg only (p-values  $<0.05$ ). Overall,  
384 GLMs results indicated i) variable trends for Cr (i.e. decreasing or increasing concentrations with  
385 increasing size of individuals), depending on the modality of the factor Area/S-Y; ii) a trend to  
386 decreasing Cu concentrations with increasing size of individuals; iii) no clear size-related trends for Zn  
387 and Cd, although no model complying with model validation could be kept for Cd; and iv) a slight  
388 trend to enhanced Hg bioaccumulation with increasing FL, especially for the modalities of the factor  
389 Area/S-Y with a large size range (Fig. 4). GLMs also revealed that for similar-sized individuals, SA  
390 individuals had significantly higher (log-transformed) concentrations in Cu and Zn than those from  
391 REU and SEY. For similar-sized individuals, those sampled in S2 in SA also had significantly higher  
392 Cr concentrations than all other individuals (Fig. 4; Appendix A and Table A.1). For similar-sized  
393 individuals sampled in REU, those captured in S2 had higher Cu and Zn concentrations than those  
394 sampled in S1, and had also higher Cu and Zn concentrations than individuals from SEY. Finally,  
395 there was no significant difference between individuals from S2-Y1 and S2-Y2 individuals in SEY  
396 (Fig. 4).

397 In the final models with  $\delta^{15}\text{N}$ , the effect of the factor Area/S-Y in explaining metal variability was still  
398 highly significant for the four elements complying with model validation. As for models with FL, the  
399 effect of  $\delta^{15}\text{N}$  was slightly significant for Cr and Cu only ( $p = 0.010$  and  $p = 0.027$  respectively), despite  
400 a quite low p-value also observed for Hg ( $p = 0.090$ ). The interaction term was only significant for Cr  
401 (all p-values  $< 0.05$ ). A slight trend to enhanced concentrations with increasing  $\delta^{15}\text{N}$  values was  
402 predicted for Cu, and to a lesser extent Zn and Hg, and variable trends for Cr. Overall, for individuals  
403 with similar- $\delta^{15}\text{N}$  values, geographical and seasonal/inter-annual differences in metal concentrations  
404 were analogous to those for similar-sized individuals (Fig. 4, Appendix A and Table A.1).

### 405 3.3. Stable isotope values and POP ratios

406  $\delta^{13}\text{C}$  values varied between  $-18.50\%$  and  $-16.30\%$ , while the range in  $\delta^{15}\text{N}$  values was  $10.71\text{--}15.11\%$   
407 (Table 4). Concerning  $\delta^{13}\text{C}$  values, a significant difference between sexes (all p-values  $< 0.05$ ) was  
408 observed in REU only but for both sampling seasons, with males presenting on average higher values  
409 than females. Significant differences between seasons/years of sampling were found within the three  
410 areas considered (all p-values  $< 0.01$ ), with individuals sampled at S2 (or S2-Y2 for SEY) having  
411 higher  $\delta^{13}\text{C}$  values. No difference in  $\delta^{13}\text{C}$  was found between individuals from REU and SA, while  
412 SEY individuals presented slightly lower  $\delta^{13}\text{C}$  values. For  $\delta^{15}\text{N}$  values, there was generally no  
413 significant difference between sexes and seasons/years of sampling. As in the case of metallic  
414 contamination, the spatial influence was more important.  $\delta^{15}\text{N}$  values significantly differed between  
415 individuals captured in the WIO and those caught in the SEAO (i.e., similar and lower concentrations  
416 on average in REU and SEY, vs. SA). Finally, within each group of individuals considered, no  
417 correlation was found between isotope values and size of individuals (results not shown).

418 POP ratios were investigated in selected individuals from REU and SA sampled in S1 (Table 4). No  
419 differences were found between sexes in SA individuals. In REU, the “Sum DDTs/Sum PCBs” and the  
420 “*p,p'*-DDT/*p,p'*-DDE” ratios were significantly lower in males than in females, while no sexual  
421 difference was observed for the “*o,p'*-DDT/*p,p'*-DDT” ratio. Similarly to metal contamination,  
422 significant differences between geographic areas were evidenced. Individuals from REU presented  
423 significantly lower values of the “*p,p'*-DDT/*p,p'*-DDE” ratio and higher values of the “*o,p'*-DDT/*p,p'*-  
424 DDT” ratio, although no spatial difference was found for the “Sum DDTs/Sum PCBs” ratio.

425 Finally, the PCAs 2 and 3 (including the five trace metals  $> \text{LQ}$ , and the trophic markers or POPs)  
426 confirmed the results from PCA 1 in terms of clustering of individuals belonging to different groups.  
427 As such, in all PCAs performed, individuals from SA clearly segregated from individuals from REU  
428 and SEY, and seasonal differences among individuals from SA were further evidenced in PCA 2  
429 (including trophic markers). The variability explained by the first two axes of the PCAs 2 and 3 ( $63.0$   
430 and  $65.7\%$ , respectively) was in the same order of PCA 1. Lastly, the DAs 3 and 4 (including trace  
431 metals + trophic markers) emphasized the highly significant contribution of  $\delta^{15}\text{N}$  values in

432 discriminating individuals from the three different areas (i.e. increasing the percentage of well-  
433 classified individuals for each area), along with the significant contribution of Cu, Zn, Cd and Hg  
434 (Table 3).

435

## 436 **4. Discussion**

### 437 **4.1. Fish size differences between areas**

438 Albacore has an estimated length at 50% maturity (L50) of around 90 cm FL in all Oceans, apart from  
439 the Mediterranean Sea where estimates are smaller (Juan-Jordá et al., 2016). More recently in the  
440 WIO, the L50 of female albacores was estimated at  $85.3 \pm 0.7$  cm FL (mean  $\pm$  standard error;  
441 Dhurmeea et al., 2016). In REU and SEY, almost all individuals had  $FL \geq 90$ cm and were thus mature  
442 individuals. In both areas, the significant difference in fish sizes between adult males and females  
443 (with females being significantly smaller than males) was consistent with the documented sexual  
444 dimorphism in this tuna species (e.g., Karazulak et al., 2011; Dhurmeea et al., 2016). Alternatively,  
445 although some fish sampled in SA were  $FL \geq 90$  cm (especially during S2), most individuals from this  
446 area were immature, probably explaining the absence of significant size difference between sexes in  
447 SA. More generally, further consideration of size (FL) in statistical analyses and interpretation of trace  
448 metal concentrations was therefore crucial, to avoid potential masking of other important factors in  
449 contamination differences (e.g., the geographic factor).

### 450 **4.2. General trends in trace metal concentrations and confrontation to health guidelines**

451 Overall, Ni and Pb concentrations were very low in all individuals analysed (mostly  $<LQ$ ), while  
452 concentrations in Cr, Cu, Zn, Cd and Hg were more variable between areas (see below). For the three  
453 non-essential elements Cd, Hg and Pb, the European Commission (EC) fixed the following safety  
454 concentrations in tuna flesh (in wet mass, wm):  $0.10 \text{ mg.kg}^{-1} \text{ wm}$  for Cd,  $1.00 \text{ mg.kg}^{-1} \text{ wm}$  for Hg, and  
455  $0.30 \text{ mg.kg}^{-1} \text{ wm}$  for Pb (EC, 2006). Considering 70% moisture content in tuna flesh (Kojadinovic et  
456 al., 2006; Munsch et al., 2016), this corresponds to maximum safety concentrations of  $0.33 \text{ mg.kg}^{-1}$  in  
457 dry mass (dm) for Cd,  $3.33 \text{ mg.kg}^{-1} \text{ dm}$  for Hg, and  $1.00 \text{ mg.kg}^{-1} \text{ dm}$  for Pb. These limits are indicated  
458 in Fig. 2 for Cd and Hg. Pb concentrations measured in all tunas from the present study were far below  
459 EC health guidelines. For Cd, 67 (34%) on the 197 individuals sampled in SA (both males and females  
460 and mostly from S1) displayed concentrations above the EC guidelines. Seven individuals (nearly 6%)  
461 of the 118 sampled in SEY also had Cd concentrations above the guidelines, while none of the  
462 individuals from REU were of a similar concern. Finally for Hg, only six individuals of the 443  
463 analysed, that were of a larger size class collected in REU and SA, presented concentrations slightly  
464 above the safety limits.

465 Hence globally, trace metal concentrations measured in the present study were in the same order of  
466 magnitude, or even in the low range of contamination concentrations by metallic elements compared  
467 to those reported in the literature for large pelagic fish such as tuna species worldwide, including  
468 albacore (e.g., Storelli et al., 2002, 2005; Kojadinovic et al., 2006, 2007; Choy et al., 2009; Chen et al.,  
469 2014; Ruelas-Inzunza et al., 2014; Bodin et al. 2017; and associated references). Similar observations  
470 were made in the case of POPs (Munschy et al., 2016), emphasizing a global low impact of  
471 anthropogenic contaminants in albacore tuna in the study areas.

#### 472 **4.3. Significant factors affecting trace metal bioaccumulation**

473 Categorical factors explaining the variability in trace metal concentrations showed the following order  
474 of influence: sex < season/year of sampling < geographic area. The influence of the continuous factors  
475 FL (fish size) and  $\delta^{15}\text{N}$  was intermediate, depending on the element considered.

476 The low effect of sex for most of the elements likely suggests poor elimination of trace metals during  
477 reproduction, contrary to organic contaminants for which this phenomena has been widely  
478 documented in adult females in particular (e.g., Jones and de Voogt, 1999; Bodiguel et al., 2009;  
479 Munschy et al., 2016). This may be nuanced in the case of Hg, for which differences between sexes  
480 were more often significant, although differences in Hg concentrations between males and females  
481 may be also due to sexual differences in the trophic ecology (e.g., type of prey, trophic level).  
482 However, potential sexual differences in trophic ecology were not supported by stable isotope values  
483 (i.e. no sexual differences revealed in  $\delta^{15}\text{N}$  values; Table 4), though different prey compositions can  
484 lead to similar isotopic signatures in a predator's tissues (Bearhop et al., 2004).

485 Within each ecosystem and especially those where fish with a large size range could be collected, Hg  
486 was the only element for which a trend to higher Hg concentrations with increasing size was found,  
487 while the trend was generally inverse or null for the other elements. The influence of the age of  
488 organisms (for which individual size may be generally considered a proxy) on Hg bioaccumulation in  
489 muscle is effectively a well-documented phenomenon. This is due to the very low elimination of this  
490 metal over time once it is linked to sulfhydryl groups of muscular proteins (Wang and Wong, 2003).  
491 The influence of individual trophic position (here regarded through  $\delta^{15}\text{N}$  values within a specific  
492 ecosystem) – which generally increases with age, size and body mass in fish – was also expected,  
493 because of the known biomagnification of Hg in food webs (Gray, 2002; Chen et al., 2008).  
494 Alternatively, for the other trace metals, there is still no consensus on a potential increase in the  
495 concentrations observed with size or trophic level of organisms, nor biomagnification in food webs;  
496 although some authors reported such phenomenon in tuna species (Ruelas-Inzunza et al., 2014) and in  
497 particular cases in both marine and freshwater food webs (Wang, 2002, Croteau et al., 2005; Cheung  
498 et Wang, 2008). On the contrary, the present results (such as those for Cu) rather suggest a dilution of  
499 muscle metal burden with individual growth, and/or that the concentrations measured and the

500 variability observed are more probably due to differences in the trophic ecology of individuals or  
501 populations. In all analyses performed, the significance of geographic origin for explaining variability  
502 in metal concentrations was effectively the highest, and notably once the effect of size had been taken  
503 into account. Due to the large predominance of the trophic pathway for the intake of metallic  
504 contaminants in marine organisms (Wang, 2002) and in top predators in particular, geographic  
505 differences in tuna trace metal concentrations are therefore most likely due to differences in the  
506 different types of prey and/or food webs exploited.

507 As a direct comparison, concentrations of several trace metals in different tissues were reported for the  
508 yellowfin tuna *T. albacares* in the WIO area, with higher muscle Hg concentrations found in  
509 individuals from Reunion Island relative to those captured in the Mozambique Channel for similar-  
510 sized individuals (Kojadinovic et al., 2006, 2007). Yellowfin tunas from Reunion Island also showed  
511 slightly higher Cu and Zn muscle concentrations relative to those captured southern in the  
512 Mozambique Channel, while concentrations in Cd were similar. Kojadinovic et al. (2007) then linked  
513 the spatial differences observed to the main prey consumed by tunas in the different zones. Indeed, in  
514 the case of Hg, mesopelagic prey species were found to present enhanced Hg bioaccumulation in  
515 various ecosystems, in comparison to epipelagic species (Monteiro et al., 1996; Choy et al., 2009;  
516 Chouvelon et al., 2012b). Moreover, most fish and piscivorous species mainly accumulate Hg under  
517 its organic form methyl-Hg (Bloom, 1992; Cossa et al., 2012), which is the most stable form, the most  
518 bioavailable and thus the most bioaccumulable form of Hg by organisms (Fitzgerald et al., 2007). The  
519 proportion of methyl-Hg relative to total-Hg in other types of organisms and prey (e.g., crustaceans,  
520 cephalopod molluscs, bivalve molluscs, or lower trophic level organisms in general) was shown to be  
521 less important and/or more variable (e.g., Claisse et al., 2001; Bustamante et al., 2006; Cossa et al.,  
522 2012). As a consequence, piscivorous predators may be more exposed to Hg than predators feeding on  
523 other prey types. A recent study reported a positive correlation between the frequency of occurrence of  
524 fish prey in the diet of dolphinfish, tunas and sharks, and the mean Hg concentrations measured in the  
525 muscle of these predators (Teffer et al., 2014). The diet and feeding habits of tunas from the present  
526 study indicated a higher consumption of fish prey (reconstituted prey weight) for individuals captured  
527 in SEY (i.e. relative to those from REU and SA; Nikolic et al., 2015; M. Potier and N. Bodin, personal  
528 communication). The mean size of prey (notably fish prey) was also found to be larger (i.e. prey of  
529 potentially higher trophic level) for individuals from REU and SEY than those from SA. Lastly,  
530 juvenile tunas were reported to be gregarious and found more in surface than adult tunas, the latter  
531 presenting more individual behaviours and being generally found deeper (e.g., Domokos et al., 2007,  
532 Cosgrove et al., 2014; reviewed in Nikolic et al., 2016), potentially feeding more on deeper prey  
533 species presenting higher Hg burdens. All this information may therefore explain the order of  
534 concentrations found for Hg, i.e. REU (fish diet important but not exclusive, prey size important and  
535 deeper behaviour reported by Nikolic et al., 2015) > SEY (almost exclusively fish diet and prey size  
536 important, but surface behaviour reported) >> SA (fish diet important but not exclusive, lower prey

537 size and surface behaviour reported). Furthermore, tunas from SA were shown to ingest greater  
538 proportions of both crustaceans and cephalopods relative to tunas from other areas, although  
539 cephalopod prey were also significant in terms of occurrence and reconstituted prey weight in the diet  
540 of REU tunas (Nikolic et al., 2015; M. Potier and N. Bodin, personal communication). This may  
541 account for the significantly higher Cu and Cd concentrations found in individuals from SA, regardless  
542 of the size of fish. Indeed, Cu is one constituent of crustaceans' respiratory pigment (hemocyanin), as  
543 it is the case for some molluscs such as cephalopods (Eisler, 2010). Crustacean and cephalopod prey  
544 therefore accumulate Cu in relatively high proportions in their tissues (e.g. Amiard et al., 1980;  
545 Miramand and Bentley, 1992; Eisler, 2010) and may contribute to the intake of Cu by their predators.  
546 Similarly, invertebrate prey species such as crustaceans or cephalopods exhibit higher Cd  
547 concentrations relative to fish (Amiard et al., 1980; Cossa and Lassus, 1989; Lahaye et al., 2005), and  
548 cephalopods (notably oceanic species) were proved to be a main vector of Cd for predators such as  
549 marine mammals (Caurant and Amiard-Triquet 1995; Bustamante et al., 1998; Lahaye et al., 2005).  
550 Finally, in addition to seasonal and/or geographical differences in the trophic ecology of individuals to  
551 explain the tuna concentrations observed, potential differences i) in trace metal environmental (abiotic)  
552 concentrations, or ii) in the bioavailability of elements at the base of the different food webs exploited,  
553 may constitute supplemental factors leading to the observations made. For instance, REU and SEY  
554 beyond to the same biogeographic province (following Oliver and Irwin, 2008), are more oligotrophic  
555 areas compared with SA (Longhurst, 1998; Sherman and Hempel, 2008), and oligotrophic conditions  
556 in the Mediterranean Sea were supposed to enhance the bioavailability of Hg (through methylation by  
557 bacteria) (Heimbürger et al., 2010). This may also contribute to the higher Hg concentrations observed  
558 in REU and SEY tunas relative to SA. Determination of trace metal concentrations in water and low  
559 trophic level organisms (i.e. plankton) would enable testing of this hypothesis, especially as abiotic  
560 and low trophic level chemical composition would directly affect the concentrations that can be found  
561 in the different prey species consumed in the different areas, and in predators. Indeed, in the case of Cr  
562 for instance, higher inputs and environmental concentrations in SA marine waters may be expected,  
563 because SA has been recognized for a long time and at the worldwide level a major reservoir for  
564 chromite mining (i.e. main ore from which industrially produced Cr comes from; Chiffoleau, 1994).

#### 565 **4.4. Relationships with trophic markers and POP patterns**

566 The results obtained for the other potential tracers (stable isotope and POP tracers) were consistent  
567 with trace metals. As such,  $\delta^{15}\text{N}$  values clearly segregated individuals captured in the WIO (REU and  
568 SEY) from those collected in SA. For  $\delta^{13}\text{C}$  values, the greater significance of seasonal/year differences  
569 than the geographic factor may be linked to shorter-term variations in C sources or baseline  $\delta^{13}\text{C}$   
570 values (reflected in higher trophic levels) than for contaminants. Several factors, linked to the season,  
571 may effectively induce variations in  $\delta^{13}\text{C}$  values of particulate organic matter (POM) or of primary



572 producers directly, including: i) temperature (influencing carbon isotopic fractionation in  
573 phytoplankton, e.g., Fontugne and Duplessy, 1981; Goericke and Fry, 1994; Rau et al., 1997);  
574 ii) partial pressure and molecular CO<sub>2</sub> concentration in ambient waters (Rau et al., 1997);  
575 iii) phytoplankton cell size, cell surface area/volume or cell geometry (Popp et al., 1998); iv) POM  
576 and/or phytoplankton composition (Darnaude et al., 2004; Harmelin-Vivien et al., 2008), with diatoms  
577 presenting higher  $\delta^{13}\text{C}$  values than other phytoplankton groups such as dinoflagellates (Fry and  
578 Wainwright, 1991). Furthermore, the temporal integration of C and N stable isotopes in biological  
579 tissues may differ slightly from those of trace metals. Animals can exhibit variable integration time of  
580 C and N signals from their food during tissue biosynthesis (Martínez del Rio et al., 2009), depending  
581 on species or on individual physiology. For trace metals, several factors can also affect the muscle  
582 bioaccumulation of contaminants observed *in situ* by organisms (Luoma and Rainbow, 2005),  
583 depending for instance on organisms' regulation mechanisms for the different contaminants. Trace  
584 metal concentrations in the muscle are generally thought to be less variable and/or reflect metal  
585 exposure on the relatively long term, i.e. order of several weeks or months, in comparison with tissues  
586 such as liver or kidneys that probably reflect contamination more immediately following contaminant  
587 incorporation, and/or metal storage at more variable temporal scales (i.e. detoxification organs,  
588 involving metallothionein proteins; Reinfelder et al., 1998; Wang and Rainbow, 2010). The muscle  
589 burden in some metals might thus reflect longer-term exposure than the seasonal scale, for instance.  
590 However, estimating metal kinetics (both accumulation and elimination) in the tissues of *in situ*  
591 organisms remains very complex. Indeed, experimental studies have shown that in fish for instance, it  
592 strongly depends on species, on metals, and on their potential interactions, on the exposure dose and  
593 on the exposure pathway (dissolved vs. trophic), etc. Moreover, experimental studies are generally  
594 performed on small species and/or juvenile individuals, on a single species at time or during relatively  
595 short periods of time, and sometimes results are contradictory (e.g., Kargin and Çoğun, 1999; Kim et  
596 al., 2004; Kraemer et al., 2005; Łuszczek-Trojnar et al., 2013). This makes very difficult to extrapolate  
597 to top predators or long-lived species such as the tunas analysed here.

598 Despite these potential artefacts, trophic markers and more specifically  $\delta^{15}\text{N}$  values provided  
599 supplemental evidence that great spatial variations occur in  $\delta^{15}\text{N}$  values at the base of different food  
600 webs (along with  $\delta^{13}\text{C}$  values), and that  $\delta^{15}\text{N}$  cannot be used independently as a proxy of the trophic  
601 positions of organisms unless the data are compared within a given environment (Jennings and Warr  
602 2003; Ménard et al., 2007; Chouvelon et al., 2012a; Pethybridge et al., 2015). Moreover,  $\delta^{15}\text{N}$  values  
603 measured in consumers may indicate different trophic functioning at the base of food webs  
604 (e.g., oligotrophy vs. mesotrophy; low vs. high primary production; etc.). As for  $\delta^{13}\text{C}$  values,  
605 differences in baseline  $\delta^{15}\text{N}$  values occur naturally (Montoya, 2007). They are intimately linked to  
606 processes occurring at the dissolved inorganic nitrogen (DIN) level, or to POM and/or phytoplankton  
607 composition, which reflect in higher trophic levels. Some of these processes lead to higher  $\delta^{15}\text{N}$   
608 values, including: i) when DIN demand is higher than the supply of nutrients in a particular area (i.e.

609 high primary production/bloom events), primary producers may be faced with a  $^{15}\text{N}$ -enriched N source  
610 (e.g., ‘‘recycled’’ or ammonium-enriched, especially if it comes from higher trophic levels), which is  
611 then reflected in the local food chain (Montoya, 2007); ii) when most of the organic matter is dissolved  
612 in particular environments, microbial food webs are predominantly supported (Biddanda et al., 2001)  
613 and microbial processing or remineralization of organic matter at the base of food webs usually results  
614 in higher  $\delta^{15}\text{N}$  values (Jennings and Warr, 2003; MacKenzie et al., 2014). Alternatively, other  
615 processes can lead to lower  $\delta^{15}\text{N}$  values measured, including the fixation of atmospheric  $\text{N}_2$  by  
616 diazotrophic organisms (e.g., cyanobacteria) in offshore or oligotrophic waters, which lower the  $\delta^{15}\text{N}$   
617 values of the residual  $\text{NO}_3^-$  pool available for primary producers in these areas (Montoya, 2007) and  
618 result in low  $\delta^{15}\text{N}$  values in consumers. As such, SIA results of the present study highlighted the  
619 higher primary productivity in SA waters (i.e. higher  $\delta^{15}\text{N}$  values measured in tunas from this  
620 ecosystem subject to the Benguela and Agulhas currents), while REU and SEY, located in the Indian  
621 South Subtropical Gyre province, face more oligotrophic conditions (i.e. lower  $\delta^{15}\text{N}$  values measured  
622 in tunas from these areas) (Longhurst, 1998; Oliver and Irwin, 2008; Sherman and Hempel, 2008).  
623 Therefore, in addition to complement the information given by trace metals (i.e. segregation between  
624 the different groups of tunas), SIA in tuna muscle appeared powerful in providing information on local  
625 biogeochemistry (C and N sources), including trophic functioning at the base of food webs. Recently  
626 in the southwest Pacific Ocean, Pethybridge et al. (2015) similarly reported a latitudinal gradient of  
627  $\delta^{15}\text{N}$  values in albacore tuna, and linked this gradient to nutrient cycling dynamics and oceanographic  
628 features in both oceans (i.e. lower  $\delta^{15}\text{N}$  values in environments with more oligotrophic conditions).  
629 The POP patterns further supported the spatial segregation of tunas between REU and SA (i.e.  
630 between the WIO and the SEAO). Higher PCB and DDT concentrations in tunas from SA in  
631 comparison with REU were previously reported by Munsch et al. (2016), and were attributed to  
632 higher dietary inputs and higher total lipid content in SA. In the present study, the POP tracers also  
633 provided complementary information on potential differences in the sources of contamination across  
634 the different areas. For instance, the ‘‘*o,p'*-DDT/*p,p'*-DDT’’ ratio (commonly used to determine the  
635 DDT origins) showed significant different mean values (0.30 in REU vs. 0.17 in SA, Table 4),  
636 although both were close to those characterising the technical mixture (0.2–0.3, Kalantzi et al., 2001).  
637 Similarly, a significant difference in the ‘‘*p,p'*-DDT/*p,p'*-DDE’’ ratio was found between individuals  
638 from these two areas. This ratio is generally used to trace the resident time and degree of degradation  
639 of *p,p'*-DDT (Suarez et al., 2013). Despite the differences observed, the values of this ratio (i.e. <0.5)  
640 indicate old DDT inputs in both areas (Suarez et al., 2013). Alternatively, the ‘‘Sum DDTs/Sum PCBs’’  
641 ratio, used to distinguish the influence of sources from agricultural origin (when >1) vs. sources from  
642 industrial origin (Lailson-Brito et al., 2011), exhibited high values in both areas. Therefore, it  
643 highlighted the important agricultural use of DDT (in comparison with industrial sources) in the  
644 southern hemisphere, including both REU and SA, while DDT has been banned for several years or  
645 even decades in northern countries.

#### 646 **4.5. Conclusions and further work perspectives**

647 Our results demonstrated that inorganic elements, when considered in isolation, are quite efficient in  
648 discriminating individuals from different geographic origins. However, the coupling with trophic  
649 markers (SIA) and with information derived from stomach contents analysis (e.g., Nikolic et al., 2015)  
650 appeared essential for explaining the metal contamination profiles observed, and for distinguishing  
651 more clearly the different populations exploiting different food webs. Information derived from POP  
652 analyses were also very complementary, including the ratios used in the present study and the detailed  
653 contamination results reported previously by Munschy et al. (2016). It strengthened the metal results  
654 by supporting the spatial segregation of tunas observed between the WIO and the SEAO as well, by  
655 evidencing the influence of the trophic ecology on the contamination observed (i.e. different food  
656 webs exploited by the tuna populations), and it also provided substantial information on the potential  
657 sources of contamination (i.e. agricultural vs. industrial sources) in the study areas. When using and  
658 interpreting biogeochemical tracers, since different prey compositions in the diet can lead to similar  
659 isotopic signatures or contaminant concentrations in a predator's tissues (e.g. Bearhop et al., 2004),  
660 one must effectively keep in mind that only differences are really informative.

661 The investigation of metal isotope ratios would be useful to understand the origin and the sources of  
662 metal contamination (e.g., Cu or Hg isotopes; El Azzi et al., 2013; Perrot et al., 2010). Other  
663 interesting perspectives would include further work on environmental matrices (e.g., speciation of  
664 metals in the water) and on lower trophic level organisms in the different areas, to thoroughly  
665 understand the processes of metal transfer leading to the differences observed in top predators.  
666 Additionally, apprehending the mechanisms of metal regulation by the different fish populations,  
667 through the analysis of metallothioneins in detoxification organs and/or through the analysis of genetic  
668 markers of metallothioneins, would also be of interest. Finally, the analysis of emerging contaminants  
669 could help to define more comprehensive or exhaustive chemical contamination profiles, although this  
670 study demonstrated that classically and historically monitored contaminants are already very  
671 informative.

672 Our results are also important in a fisheries management perspective, through the assessment of the  
673 potential links (gene flows, individuals migrations, etc.) between Atlantic and Indian Oceans stocks,  
674 managed by different Regional Fisheries Management Organisations. Indeed, one of the key issues to  
675 improve the management of albacore tuna is a better understanding of the spatial dynamics and  
676 population connectivity. At the integration time scale represented by this study of metal  
677 bioaccumulation and additional trophic and chemical tracers in tuna muscle, the differences observed  
678 suggested limited flows of individuals between the SEAO and the WIO. In the near future, further  
679 coupling of this information derived from chemical/ecological tracers with genetic data should help  
680 refine management units and stocks of albacore tunas in the southern hemisphere, and more  
681 specifically in the Indian and Atlantic Oceans.

682

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999

1000 **Table 1:** Results obtained for the certified reference materials (CRMs) used for trace metal analyses. Values are means  $\pm$  standard deviation, in mg.kg<sup>-1</sup> dry  
 1001 mass (dm). The number of replicates for each CRM was 37 for Hg, and 18 for the other trace metals. Certified values are in italics. The recovery rate (in %),  
 1002 the limit of quantification (LQ, in mg.kg<sup>-1</sup> dm) and the percentage of individuals below the LQ (on the 443 analysed) are also indicated.

Trace metal	<i>Certified value</i>	Measured value	Recovery rate	<i>Certified value</i>	Measured value	Recovery rate	LQ	% of individuals <LQ
	<b>IAEA-407</b>			<b>DOLT-3</b>				
<b>Cr</b>	<i>0.73 ± 0.06</i>	0.76 ± 0.05	104	3.5*	2.5 ± 0.3	71	0.03	0
<b>Ni</b>	<i>0.60 ± 0.05</i>	0.57 ± 0.27	95	<i>2.72 ± 0.35</i>	2.34 ± 0.32	86	0.03	81
<b>Cu</b>	<i>3.28 ± 0.08</i>	3.00 ± 0.07	92	<i>31.2 ± 1.0</i>	30.8 ± 0.6	99	0.25	0
<b>Zn</b>	<i>67.1 ± 0.8</i>	66.5 ± 1.2	99	<i>86.6 ± 2.4</i>	90.3 ± 1.4	104	2.5	0
<b>Cd</b>	<i>0.189 ± 0.004</i>	0.179 ± 0.009	95	<i>19.4 ± 0.6</i>	19.2 ± 0.3	99	0.05	21
<b>Pb</b>	<i>0.12 ± 0.02</i>	0.11 ± 0.03	92	<i>0.319 ± 0.045</i>	0.377 ± 0.191	118	0.05	98
	<b>IAEA-142</b>							
<b>Hg</b>	<i>0.126 ± 0.007</i>	0.124 ± 0.002	98	—	—	—	0.015	0

1003 \* Value only given for information on the certificate.

1004

1005 **Table 2:** Fish body size (fork length, FL, in cm) and trace metal concentrations (in mg.kg<sup>-1</sup> dry mass) measured in the muscle of albacore tunas (N =443  
 1006 individuals) collected in 2013-2014 in the western Indian Ocean and in the south-eastern Atlantic Ocean. Values are means ± standard deviation. Italic values  
 1007 are those per season and/or year of sampling, and bold values are those per geographic area.

Area; Season and/or Year of sampling; Sex	N	FL	Cr	Ni *	Cu	Zn	Cd	Hg	Pb *
<b>Reunion Island (REU)</b>	<b>128</b>	<b>102 ± 5</b>	<b>0.53 ± 0.06</b>	<b>0.01 ± 0.01</b>	<b>0.84 ± 0.20</b>	<b>14.5 ± 7.1</b>	<b>0.07 ± 0.03</b>	<b>1.708 ± 0.629</b>	<b>0.01 ± 0.03</b>
<i>Season 1 (Nov-Dec 2013)</i>	<i>64</i>	<i>104 ± 4</i>	<i>0.51 ± 0.06</i>	<i>0.02 ± 0.02</i>	<i>0.71 ± 0.15</i>	<i>11.2 ± 2.2</i>	<i>0.06 ± 0.02</i>	<i>1.773 ± 0.677</i>	<i>0.01 ± 0.01</i>
Females	31	101 ± 3	0.52 ± 0.05	0.02 ± 0.02	0.74 ± 0.18	11.0 ± 2.1	0.06 ± 0.02	1.721 ± 0.630	0.01 ± 0.01
Males	33	107 ± 3	0.50 ± 0.07	0.02 ± 0.02	0.69 ± 0.13	11.5 ± 2.2	0.06 ± 0.02	1.821 ± 0.725	0.01 ± 0.01
<i>Season 2 (May-July 2014)</i>	<i>64</i>	<i>101 ± 5</i>	<i>0.54 ± 0.05</i>	<i>0.01 ± 0.00</i>	<i>0.98 ± 0.14</i>	<i>17.7 ± 8.7</i>	<i>0.08 ± 0.03</i>	<i>1.644 ± 0.574</i>	<i>0.01 ± 0.04</i>
Females	32	99 ± 4	0.54 ± 0.05	0.00 ± 0.00	0.96 ± 0.15	20.0 ± 11.6	0.07 ± 0.02	1.405 ± 0.392	0.00 ± 0.00
Males	32	104 ± 4	0.55 ± 0.05	0.01 ± 0.00	1.00 ± 0.12	15.4 ± 2.7	0.08 ± 0.03	1.882 ± 0.631	0.01 ± 0.05
<b>Seychelles (SEY)</b>	<b>118</b>	<b>96 ± 5</b>	<b>0.52 ± 0.04</b>	<b>0.01 ± 0.01</b>	<b>0.64 ± 0.11</b>	<b>11.3 ± 2.2</b>	<b>0.09 ± 0.16</b>	<b>1.403 ± 0.406</b>	<b>0.00 ± 0.00</b>
<i>Season 2-Year 1 (June-July 2013)</i>	<i>50</i>	<i>96 ± 5</i>	<i>0.50 ± 0.04</i>	<i>0.01 ± 0.01</i>	<i>0.67 ± 0.14</i>	<i>12.3 ± 2.9</i>	<i>0.16 ± 0.22</i>	<i>1.310 ± 0.306</i>	<i>0.01 ± 0.00</i>
Females	20	95 ± 4	0.51 ± 0.05	0.01 ± 0.01	0.64 ± 0.12	12.5 ± 2.0	0.15 ± 0.13	1.302 ± 0.220	0.01 ± 0.00
Males	30	98 ± 5	0.50 ± 0.03	0.01 ± 0.01	0.68 ± 0.15	12.2 ± 3.4	0.17 ± 0.27	1.316 ± 0.356	0.01 ± 0.01
<i>Season 2-Year 2 (April 2014)</i>	<i>68</i>	<i>96 ± 5</i>	<i>0.53 ± 0.03</i>	<i>0.01 ± 0.01</i>	<i>0.62 ± 0.09</i>	<i>10.6 ± 1.0</i>	<i>0.04 ± 0.01</i>	<i>1.471 ± 0.457</i>	<i>0.00 ± 0.00</i>
Females	36	94 ± 5	0.53 ± 0.03	0.01 ± 0.01	0.61 ± 0.10	10.7 ± 1.1	0.04 ± 0.01	1.364 ± 0.427	0.00 ± 0.00
Males	32	99 ± 4	0.53 ± 0.03	0.01 ± 0.01	0.63 ± 0.08	10.4 ± 0.9	0.04 ± 0.01	1.592 ± 0.465	0.00 ± 0.00
<b>South Africa (SA)</b>	<b>197</b>	<b>87 ± 8</b>	<b>0.53 ± 0.08</b>	<b>0.04 ± 0.04</b>	<b>1.55 ± 0.38</b>	<b>23.3 ± 8.3</b>	<b>0.34 ± 0.26</b>	<b>0.958 ± 0.443</b>	<b>0.01 ± 0.02</b>
<i>Season 1 (Nov-Dec 2013)</i>	<i>98</i>	<i>83 ± 4</i>	<i>0.46 ± 0.04</i>	<i>0.04 ± 0.04</i>	<i>1.69 ± 0.43</i>	<i>26.4 ± 9.5</i>	<i>0.37 ± 0.25</i>	<i>0.795 ± 0.151</i>	<i>0.01 ± 0.02</i>
Females	50	83 ± 4	0.46 ± 0.04	0.04 ± 0.04	1.70 ± 0.42	25.0 ± 7.7	0.37 ± 0.26	0.759 ± 0.143	0.01 ± 0.02
Males	48	83 ± 4	0.45 ± 0.04	0.03 ± 0.03	1.69 ± 0.45	27.9 ± 10.9	0.38 ± 0.23	0.833 ± 0.151	0.01 ± 0.02
<i>Season 2 (April-May 2014)</i>	<i>99</i>	<i>90 ± 9</i>	<i>0.59 ± 0.05</i>	<i>0.04 ± 0.04</i>	<i>1.42 ± 0.25</i>	<i>20.2 ± 5.4</i>	<i>0.30 ± 0.26</i>	<i>1.120 ± 0.562</i>	<i>0.02 ± 0.02</i>
Females	51	90 ± 9	0.60 ± 0.05	0.04 ± 0.05	1.44 ± 0.23	20.8 ± 6.8	0.30 ± 0.22	1.093 ± 0.444	0.01 ± 0.01
Males	48	91 ± 10	0.59 ± 0.05	0.03 ± 0.02	1.40 ± 0.27	19.5 ± 3.4	0.30 ± 0.30	1.150 ± 0.669	0.02 ± 0.02

1008 \* Values only given for information, due the high percentage of individuals below the limit of quantification for Ni and Pb (see section 2.4 and Table 1).  
 1009

1010 **Table 3:** Results for the discriminant analyses (DAs) applied to evaluate the potential of trace metals and/or their combination with trophic markers ( $\delta^{13}\text{C}$  and  
 1011  $\delta^{15}\text{N}$  values) in discriminating albacore tunas from different geographic areas. The correlation coefficients between the discriminant functions and each of the  
 1012 original variables (trace metal or isotope value) included in DAs – also called « canonical correlation coefficients » (ccc) – are given, and help assess the  
 1013 relative importance of each variable for the discrimination of individuals along the axes. The coefficients of linear discriminants for axis 1 and/or 2 – also  
 1014 called « standardised discrimination coefficients » (sdc) – are given as well, and can be used for reconstructing equations of the DAs (Zuur et al., 2007).  
 1015 Variables with  $\text{ccc} > 0.50$  (absolute value) are in bold.

DA: variables used; number of areas considered (number of individuals)	% of well-classified individuals for each area	Order of importance of variables (ccc; sdc) for discrimination along the first axis of DA	Order of importance of variables (ccc; sdc) for discrimination along the second axis of DA
<b>DA 1: 5 trace metals; 3 areas (N=443)</b>	REU = 69% SEY = 83% SA = 93%	<b>Cu</b> (0.98; 1.49) > <b>Zn</b> (0.72; 0.16) > <b>Cd</b> (0.65; 0.10) > <b>Hg</b> (-0.57; -0.32) > Cr (0.05; 0.17)	<b>Hg</b> (0.71; 1.00) > Cd (-0.30; -0.65) > Zn (0.17; 0.20) > Cu (0.14; 0.87) > Cr (0.10; 0.02)
<b>DA 2: 5 trace metals; 2 areas (N=246)</b>	REU = 73% SEY = 89%	<b>Cu</b> (-0.86; -1.14) > <b>Zn</b> (-0.46; -0.04) > <b>Hg</b> (-0.45; -0.45) > Cd (0.18; 0.44) > Cr (-0.12; 0.09)	—
<b>DA 3: 5 trace metals + <math>\delta^{13}\text{C}</math> + <math>\delta^{15}\text{N}</math>; 3 areas (N=335)</b>	REU = 84% SEY = 60% SA = 100%	<b><math>\delta^{15}\text{N}</math></b> (0.92; 1.57) > <b>Cu</b> (0.86; 0.74) > <b>Zn</b> (0.68; 0.16) > <b>Hg</b> (-0.56; -0.33) > <b>Cd</b> (0.54; -0.01) > <b><math>\delta^{13}\text{C}</math></b> (-0.15; -0.38) > Cr (0.06; 0.14)	Hg (0.44; 0.78) > <b><math>\delta^{13}\text{C}</math></b> (-0.37; -0.39) > Cu (0.31; 1.21) > Zn (0.24; 0.15) > <b><math>\delta^{15}\text{N}</math></b> (-0.17; -0.56) > Cd (0.13; -0.49) > Cr (0.07; 0.21)
<b>DA 4: 5 trace metals+ <math>\delta^{13}\text{C}</math> + <math>\delta^{15}\text{N}</math>; 2 areas (N=159)</b>	REU = 83% SEY = 78%	<b>Cu</b> (-0.63; -0.80) > <b>Hg</b> (-0.44; -0.56) > <b><math>\delta^{13}\text{C}</math></b> (0.38; 0.59) > <b><math>\delta^{15}\text{N}</math></b> (0.35; 0.39) > <b>Zn</b> (-0.35; -0.25) > <b>Cd</b> (0.26; 0.45) > Cr (-0.08; 0.07)	—

1016

1017

1018 **Table 4:** Values for additional tracers considered and measured on a subsample of individuals analysed for trace elements: trophic markers ( $\delta^{13}\text{C}$ ,  $\delta^{15}\text{N}$ , in ‰)  
 1019 and ratios of persistent organic pollutants (POPs). Values are mean  $\pm$  standard deviation in the muscle tissue. N =number of individuals. Italic values are those  
 1020 per season and/or year of sampling, and bold values are those per geographic area.

Area; Season and/or Year of sampling; Sex	N	Trophic markers ( $\delta^{13}\text{C}$ , $\delta^{15}\text{N}$ ) <sup>a</sup>		N	POPs (ratios) <sup>a</sup>		
		$\delta^{13}\text{C}$	$\delta^{15}\text{N}$		Sum DDTs / Sum PCBs <sup>b</sup>	<i>p,p'</i> DDT / <i>p,p'</i> DDE <sup>c</sup>	<i>o,p'</i> DDT / <i>p,p'</i> DDT <sup>d</sup>
<b>Reunion Island (REU)</b>	<b>96</b>	<b>-17.42 <math>\pm</math> 0.33</b>	<b>12.06 <math>\pm</math> 0.56</b>				
<i>Season 1 (Nov-Dec 2013)</i>	<i>49</i>	<i>-17.54 <math>\pm</math> 0.34</i>	<i>12.01 <math>\pm</math> 0.58</i>	<b>43</b>	<b>2.83 <math>\pm</math> 1.11</b>	<b>0.26 <math>\pm</math> 0.14</b>	<b>0.31 <math>\pm</math> 0.09</b>
Females	24	-17.74 $\pm$ 0.32	11.96 $\pm$ 0.63	21	3.44 $\pm$ 1.05	0.35 $\pm$ 0.13	0.28 $\pm$ 0.09
Males	25	-17.35 $\pm$ 0.23	12.07 $\pm$ 0.54	22	2.25 $\pm$ 0.83	0.17 $\pm$ 0.09	0.34 $\pm$ 0.09
<i>Season 2 (May-July 2014)</i>	<i>47</i>	<i>-17.30 <math>\pm</math> 0.27</i>	<i>12.11 <math>\pm</math> 0.55</i>	—	—	—	—
Females	23	-17.38 $\pm$ 0.26	12.02 $\pm$ 0.50	—	—	—	—
Males	24	-17.22 $\pm$ 0.27	12.20 $\pm$ 0.59	—	—	—	—
<b>Seychelles (SEY)</b>	<b>63</b>	<b>-17.26 <math>\pm</math> 0.27</b>	<b>12.31 <math>\pm</math> 0.43</b>	—	—	—	—
<i>Season 2-Year 1 (June-July 2013)</i>	<i>26</i>	<i>-17.40 <math>\pm</math> 0.33</i>	<i>12.47 <math>\pm</math> 0.44</i>	—	—	—	—
Females	11	-17.44 $\pm$ 0.38	12.37 $\pm$ 0.58	—	—	—	—
Males	15	-17.37 $\pm$ 0.29	12.54 $\pm$ 0.30	—	—	—	—
<i>Season 2-Year 2 (April 2014)</i>	<i>37</i>	<i>-17.16 <math>\pm</math> 0.17</i>	<i>12.20 <math>\pm</math> 0.39</i>	—	—	—	—
Females	24	-17.18 $\pm$ 0.18	12.18 $\pm$ 0.41	—	—	—	—
Males	13	-17.14 $\pm$ 0.16	12.24 $\pm$ 0.35	—	—	—	—
<b>South Africa (SA)</b>	<b>176</b>	<b>-17.47 <math>\pm</math> 0.43</b>	<b>13.80 <math>\pm</math> 0.53</b>				
<i>Season 1 (Nov-Dec 2013)</i>	<i>78</i>	<i>-17.62 <math>\pm</math> 0.41</i>	<i>13.65 <math>\pm</math> 0.50</i>	<b>43</b>	<b>3.02 <math>\pm</math> 0.66</b>	<b>0.40 <math>\pm</math> 0.07</b>	<b>0.17 <math>\pm</math> 0.02</b>
Females	43	-17.59 $\pm$ 0.44	13.67 $\pm$ 0.53	22	3.18 $\pm$ 0.80	0.40 $\pm$ 0.07	0.17 $\pm$ 0.02
Males	35	-17.65 $\pm$ 0.38	13.63 $\pm$ 0.46	21	2.85 $\pm$ 0.44	0.40 $\pm$ 0.07	0.16 $\pm$ 0.02
<i>Season 2 (April-May 2014)</i>	<i>98</i>	<i>-17.34 <math>\pm</math> 0.40</i>	<i>13.92 <math>\pm</math> 0.53</i>	—	—	—	—
Females	51	-17.38 $\pm$ 0.40	13.89 $\pm$ 0.54	—	—	—	—
Males	47	-17.30 $\pm$ 0.39	13.96 $\pm$ 0.52	—	—	—	—

1021 <sup>a</sup> N=335 and 86 in total, for trophic markers and POPs respectively.

1022 <sup>b</sup> “Sum PCBs/Sum DDTs” ratio: can generally distinguish the influence of sources from agricultural origin (when largely >1) vs. sources from industrial origin  
 1023 (Yogui et al., 2003; Lailson-Brito et al., 2011).



1024 <sup>c</sup> “*p,p'*-DDT/*p,p'*-DDE” ratio: can trace the residence time and degree of degradation of DDT in the environment, and consequently distinguish new vs. old  
1025 DDT sources (Suárez et al., 2013).  
1026 <sup>d</sup> “*o,p'*-DDT/*p,p'*-DDT” ratio: can trace the DDT origin (close to the technical mixture (0.2–0.3), or not; Kalantzi et al., 2001).  
1027

1028 **Caption to figures:**

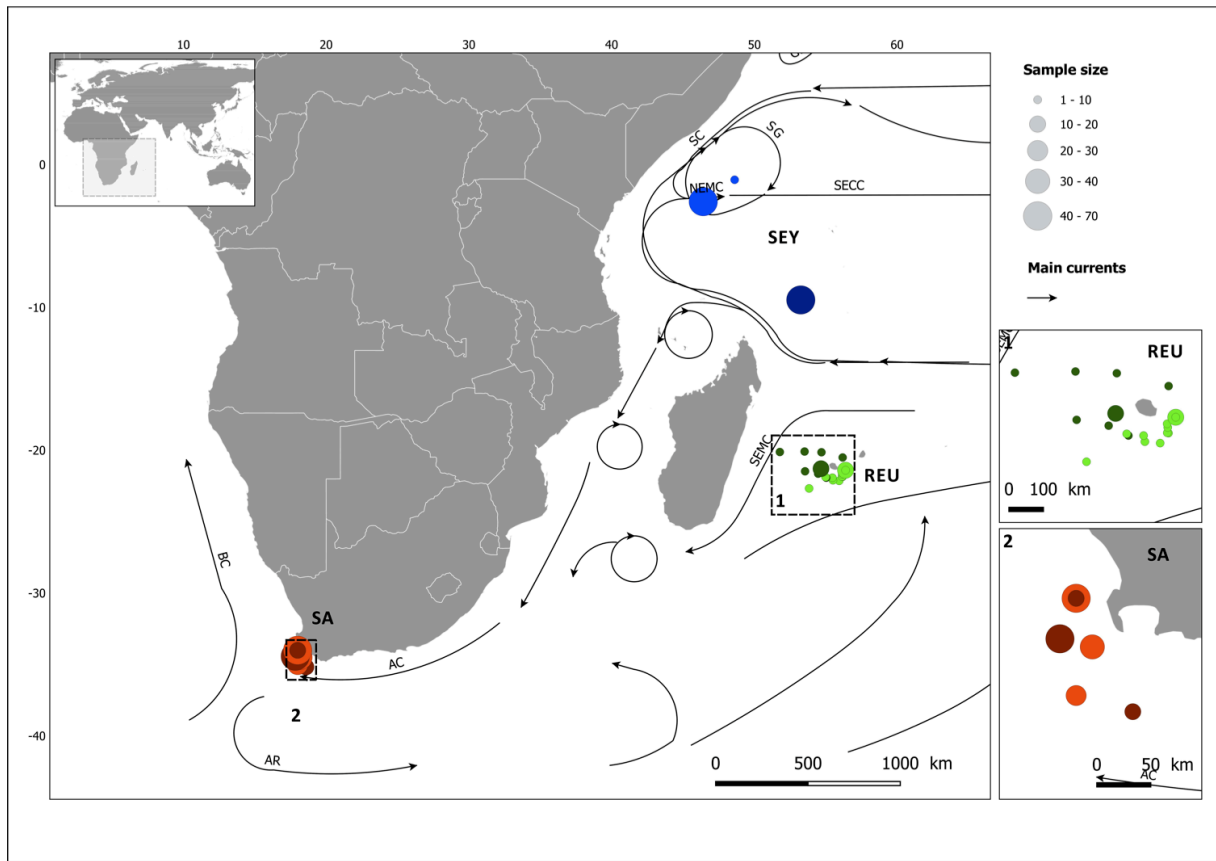
1029

1030 **Figure 1:** Map of tuna sampling in different areas in the western Indian Ocean and in the south-eastern  
1031 Atlantic Ocean, and over two seasons and/or years between June 2013 and July 2014.  
1032 REU =Reunion Island; SEY =Seychelles; SA =South Africa. Light colours correspond to a sampling  
1033 over Season 1 (for REU and SA) or Season 2-Year 1 (for SEY), and dark colours to a sampling over  
1034 Season 2 (for REU and SA) or Season 2-Year 2 (for SEY). Main regional currents (from Schott et al.,  
1035 2009) are also indicated: SECC =South Equatorial Counter Current; NEMC =Northeast Madagascar  
1036 Current; SEMC =Southeast Madagascar Current; SC: =Somali Current; SG =Southern Gyre;  
1037 AC =Agulhas Current; AR =Agulhas Current Retroflexion; BC =Benguela Current. "Sample size"  
1038 indicates the number of individuals collected in each area.  
1039

1040 **Figure 2:** Boxplots of raw and size-normalised (to a 95cm-individual) trace metal concentrations  
1041 measured in the study tunas (N =443), excluding the elements Ni and Pb for which most of values  
1042 were under limits of quantification (see section 2.4). The box width is function of the number of  
1043 individuals in each group considered. The box length represents the interquartile, the bar length  
1044 represents the range, and the horizontal lines in bold are median values. A same letter indicates that  
1045 areas are not significantly different (results of the post-hoc multiple comparison test with Holm  
1046 adjustment method, after Kruskal-Wallis test). Following Student t-tests or Mann-Whitney-Wilcoxon  
1047 tests performed, stars indicate significant differences between sexes (in a given season and/or year,  
1048 and in a given area), or significant differences between seasons/years of sampling in a given area (with  
1049 \* p <0.05; \*\* p <0.01; \*\*\* p <0.001). For the non-essential elements Cd and Hg, the maximum safety  
1050 concentration limits (EC, 2006) are also indicated in red dashed lines, considering average 70%  
1051 moisture content in tuna flesh. F =Females; M =Males; S1 =Season 1; S2 =Season 2; S2-  
1052 Y1 =Season 2-Year 1; S2-Y2 =Season 2-Year 2.  
1053

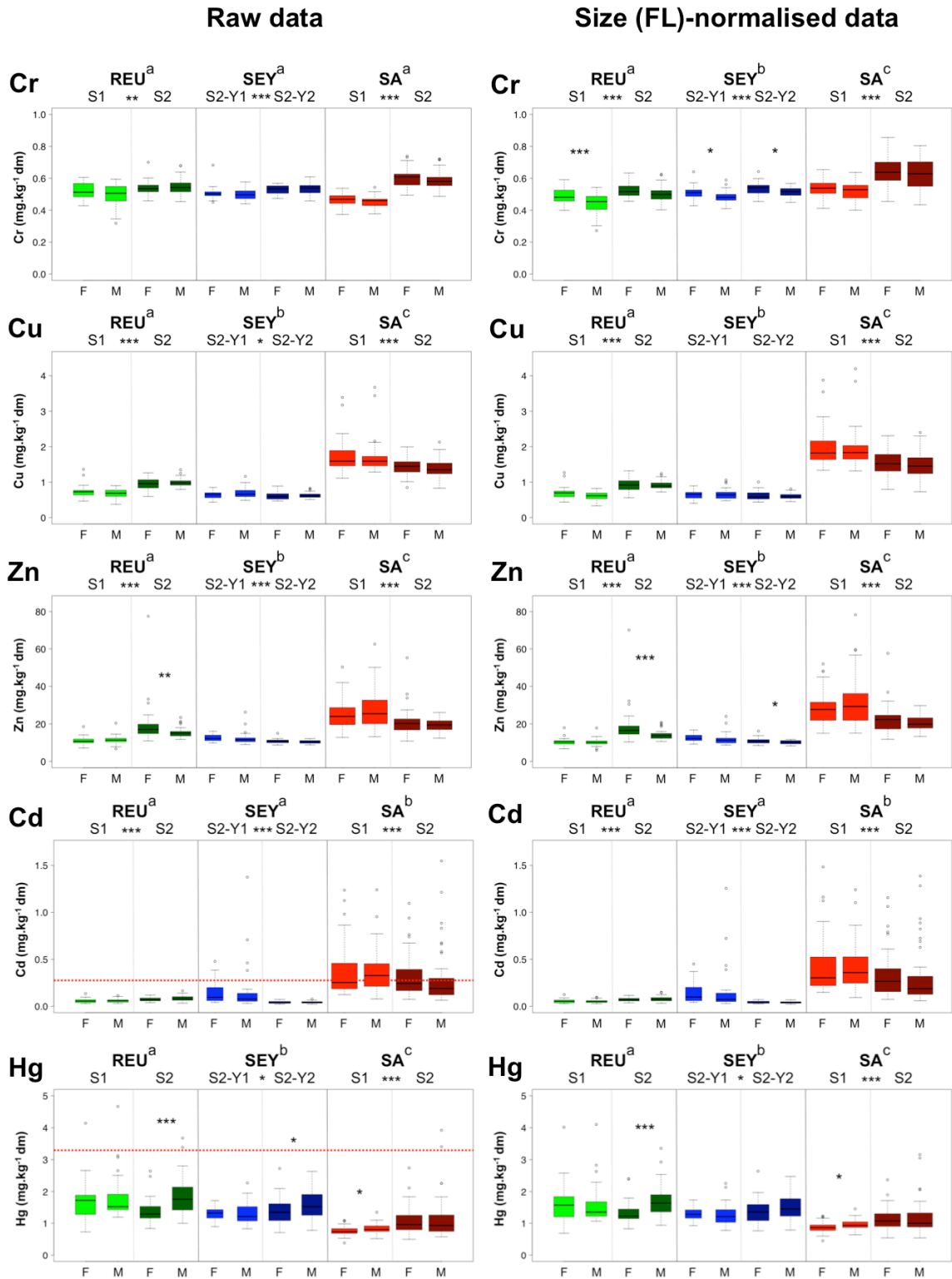
1054 **Figure 3:** Projection of variables and individuals on the first two components resulting from the  
1055 principal component analyses (PCAs) 1–3 performed (PCA 1: including only the 5 trace metals above  
1056 limits of quantification; PCA 2: 5 trace metals + trophic markers  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$ ; PCA 3: 5 trace metals  
1057 + POP ratios). For each PCA, the top figure corresponds to the correlation biplot showing the  
1058 distribution of variables. For a given variable, the length of the line shows how well it is represented  
1059 by the 2-dimensional approximation, and reflects its contribution to the first two principal components  
1060 (horizontal axis: principal component 1; vertical axis: principal component 2; and the variability  
1061 explained by each axis is given). Variables pointing in the same direction display a high positive  
1062 correlation; variables pointing in the opposite direction have a high negative correlation; and variables  
1063 with an angle of  $90^\circ$  have a small correlation close to 0. For each PCA, the middle figure corresponds  
1064 to the projection of individuals on the correlation biplot, with a grouping of individuals by sex. The  
1065 bottom figure finally corresponds to the projection of individuals on the correlation biplot as well, but  
1066 with a grouping of individuals by area and season and/or year of sampling. N=number of individuals  
1067 included in each PCA; F =Females; M =Males; S1 =Season 1; S2 =Season 2; S2-Y1 =Season 2-  
1068 Year 1; S2-Y2 =Season 2-Year 2; REU =Reunion Island; SEY =Seychelles; SA =South Africa.  
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1070 **Figure 4:** Relationships between muscle log-transformed metal concentrations and body size (FL; left  
1071 panel) or  $\delta^{15}\text{N}$  values (right panel) in the study tunas (N =443 when FL considered; N =335 when  
1072  $\delta^{15}\text{N}$ ). Results from the GLM models (lines) are plotted on observed data (models fitted to individual  
1073 log-transformed metal concentrations to identify size-related or diet  $\delta^{15}\text{N}$ -related trends, and the  
1074 confounding effect of sex, season/year and area of sampling for explaining trace metal concentrations  
1075 variability). As the factor Area/Season-Year of sampling (Area/S-Y) was always significant in the  
1076 final GLMs, one line per modality of the factor Area/S-Y is presented. S1 =Season 1; S2 =Season 2;  
1077 S2-Y1 =Season 2-Year 1; S2-Y2 =Season 2-Year 2. When the interaction term was not significant,  
1078 lines are parallel. Results for models that did not comply with necessary assumptions are not presented  
1079 (i.e. for Cd).  
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Figure 1



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Figure 2

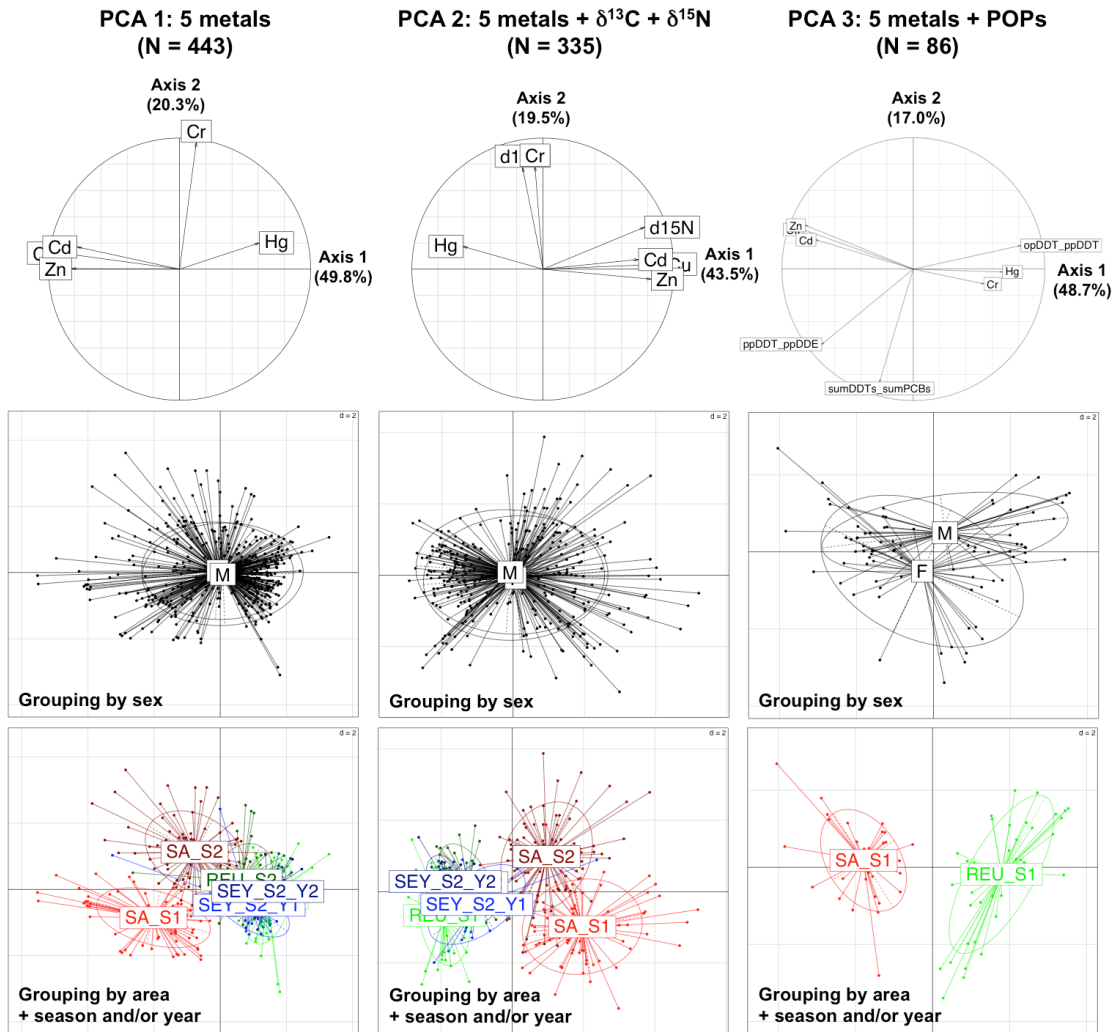
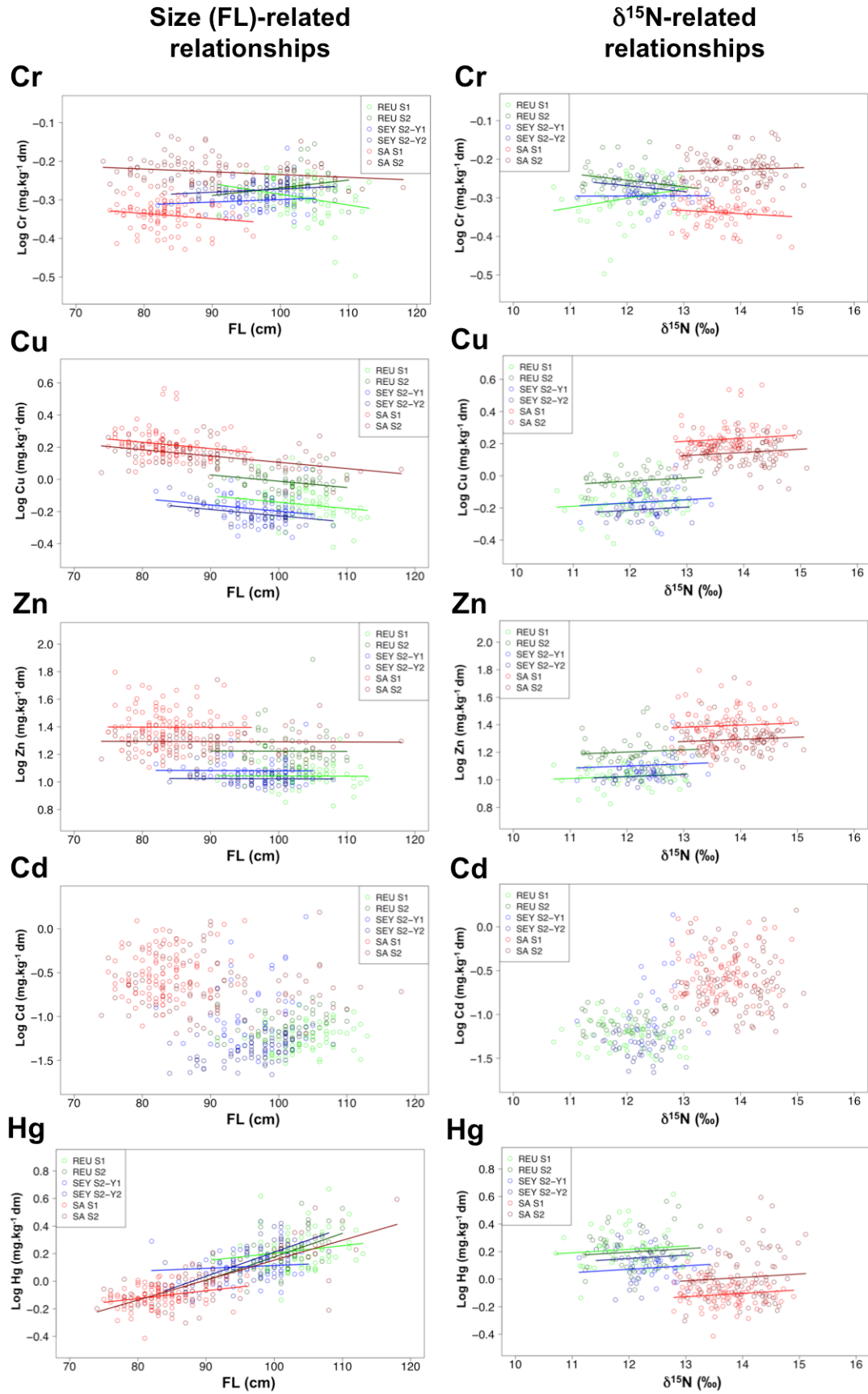


Figure 3

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Figure 4