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1 **Element patterns in albatrosses and petrels: influence of trophic**
2 **position, foraging range, and prey type**

3
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16
17 *Trophic position, foraging range, and prey type were found to influence element*
18 *compositions and concentrations in Procellariiformes from South Georgia.*

19
20 **Abstract**

21

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22 We investigated the concentrations of 22 essential and non-essential elements
23 among a community of Procellariiformes (and their prey) to identify the extent to which
24 trophic position and foraging range governed element accumulation. Stable isotope analysis
25 (SIA) was used to characterise trophic ($\delta^{15}\text{N}$) and spatial patterns ($\delta^{13}\text{C}$) among species.
26 Few consistent patterns were observed in element distributions among species and diet
27 appeared to be highly influential in some instances. Arsenic levels in seabird red blood cells
28 correlated with $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$, demonstrating the importance of trophic position and
29 foraging range for arsenic distribution. Arsenic concentrations in prey varied significantly
30 across taxa, and in the strength of association with $\delta^{15}\text{N}$ values (trophic level). In most
31 instances, element patterns in Procellariiformes showed the clearest separation among
32 species, indicating that a combination of prey selection and other complex species-specific
33 characteristics (e.g. moult patterns) were generally more important determining factors than
34 trophic level *per se*.

35

36 *Key words:* Elements; Procellariiformes; Trophic Position; Diet; South Georgia

37

38 **1. Introduction**

39

40 Some metals and metalloids are essential elements (EE) involved in physiological
41 and biochemical processes (Abdulla and Chmielnicka, 1990), but can be toxic when
42 exposure and assimilation is excessive. Others, such as lead (Pb), cadmium (Cd), arsenic
43 (As) and mercury (Hg), are non-essential elements (NEE) that are also potentially toxic.

44 Marine organisms can be exposed to elements from anthropogenic and natural sources, and
45 exposure is governed by various factors, including foraging location and trophic position.

46 Although the Southern Ocean is a remote environment, it is subject to
47 anthropogenic metal inputs due to global transport of elements in the atmosphere and
48 through oceanic circulation (Gaiero et al., 2003). Seabirds provide an ideal model with
49 which to investigate contaminant dynamics in marine food webs (see Gilbertson et al.,
50 1987; Furness, 1993; Furness and Camphuysen, 1997) and the Procellariiformes exploit a
51 wide range of ecological niches in the Southern Ocean. Many species occupy high trophic
52 positions and so can act as sentinels of exposure, highlighting potential threats to other apex
53 predators. Previous studies have examined mercury (Hg) and other elements in lower order
54 species in the Southern Ocean (e.g. Bargagli et al., 1998; Negri et al., 2006) but, apart from
55 Hg, concentrations of other elements in biota at higher trophic levels have received much
56 less attention (although see Lock et al., 1992; Stewart et al., 1999; González-Solís et al.,
57 2002; Bocher et al., 2003).

58 Conventional approaches for quantifying dietary composition and assessing metal
59 intake, such as analysis of gut content or regurgitate, have a number of biases (see Votier et
60 al., 2003) and samples are often difficult to obtain over long periods. In this study, we
61 analysed stable isotopes to elucidate broad-scale, inter- and intra-specific dietary patterns in
62 Procellariiformes (Hobson et al., 1994), and so determine whether differences in foraging
63 strategy explained variation in elemental uptake. Use of stable isotopes for this purpose is
64 based on the predictable increase in the ratio of $^{15}\text{N}:^{14}\text{N}$ (by ca. 3.4‰) with each trophic
65 level, and so $\delta^{15}\text{N}$ can be used as a proxy for trophic position (e.g. DeNiro and Epstein,
66 1981; Hobson and Welch, 1992; Bearhop et al., 1999). $\delta^{13}\text{C}$ can vary depending on whether

67 animals feed inshore or offshore, or on benthic or pelagic prey (e.g. Chisholm et al., 1982;
68 Hobson et al., 1994); $\delta^{13}\text{C}$ values also vary with latitude (Cherel and Hobson, 2007) and, as
69 such, provide a coarse-scale proxy for foraging location in wide-ranging organisms.

70 The overarching aim of the present study was to investigate the distributions and
71 dynamics of toxic heavy metals, essential metals and metalloids in Southern Ocean
72 seabirds, and assess the extent to which they may be bioaccumulated or biomagnified
73 through foodchains (see Bargagli et al., 1996; McIntyre and Beauchamp, 2007). We
74 analysed red blood cells (hereafter blood) and feathers collected during the breeding season
75 to relate element assimilation to trophic position. The residence times of dietary isotopic
76 signatures are dependent on the regenerative time of the tissue analysed (Hobson and
77 Clarke, 1992a, 1992b; Hobson 1999). Blood cells are short-term integrators of dietary
78 isotope signatures and are derived from prey consumed within the past 2-3 weeks. Blood
79 metal concentrations also reflect recent exposure, concentrations equilibrating between the
80 blood and the body organs (Burger and Gochfeld, 1997). Hence, the isotopic signature and
81 element concentrations in blood are temporally matched and reflect foraging in the
82 breeding season. In contrast, most Procellariiformes grow feathers during the non-breeding
83 period and so feathers provide a means for characterising diet and element burdens
84 acquired at this time. Feathers, once grown, are metabolically inert. They therefore
85 provide a dietary isotope and element exposure ‘snap-shot’ for the period of feather growth
86 (Mizutani et al., 1990, 1992, Bearhop et al., 2002). By randomly sampling body feathers,
87 the measured isotope signature and element concentration is averaged over the moult period
88 (Bearhop et al., 2000). The temporal matching of the feather isotope signature and element
89 concentrations is unlikely to be as good as that for blood because feather data are based on

90 averaged values obtained over a relatively long time period. Overall, however, it is possible
91 to gather information on trophic level and foraging range for the breeding (blood stable
92 isotopes) and non-breeding season (feather stable isotopes), although it is also important to
93 determine isotope values for likely prey items (as determined from previous conventional
94 dietary studies) so that the extent to which certain prey types may affect metal
95 accumulation in birds can also be evaluated.

96 In this study, our specific objectives were to investigate: (1) current levels of
97 exposure of Procellariiformes to a range of EE and NEE by measuring concentrations in
98 blood and feathers; (2) the extent to which assimilation varies between species; and, (3) the
99 extent to which foraging area, diet, and trophic level explains intra- and inter-individual
100 variation in metal accumulation in Procellariiformes.

101

102 **2. Materials and methods**

103

104 *2.1 Sample collection and preparation*

105

106 Fieldwork was undertaken on Bird Island, South Georgia (54° 00'S, 38° 03'W) from
107 December 2001 for a period of four months. Ten species were sampled in total: Antarctic
108 prion (*Pachyptila desolata*), black-browed albatross (*Thalassarche melanophrys*), blue
109 petrel (*Halobaena caerulea*), common diving petrel (*Pelecanoides urinatrix*), South
110 Georgian diving petrel (*Pelecanoides georgicus*), grey-headed albatross (*T. chrysostoma*),
111 northern giant petrel (*Macronectes halli*), southern giant petrel (*M. giganteus*), white-

112 chinned petrel (*Procellaria aequinoctialis*) and wandering albatross (*Diomedea exulans*).
113 Blood and feathers from adults of each species were taken from surface-nesting birds at the
114 end of the brood-guard period and from burrowing birds whilst mist-netting adjacent to
115 breeding colonies. Whole blood (0.2-1.0ml) (from the tarsal vein) was spun to separate
116 cells and plasma, immediately frozen, subsequently freeze-dried to a constant mass and
117 then again frozen until analysis. Only red blood cells were analysed for isotopes and trace
118 elements. Six to 8 feathers sampled at random were taken from the mantle region of each
119 bird. Feathers were stored in plastic bags at room temperature until analysed. Fresh prey
120 samples (muscle tissue) were obtained from seabird regurgitates, or from fisheries vessels
121 operating in waters surrounding South Georgia, and frozen at -20°C after identification.
122 Cephalopods were identified from beaks according to Clarke (1986) and Rodhouse et al.
123 (1992), fish from external morphology according to Gon and Heemstra (1990), or from
124 otoliths according to Reid (1996), and crustaceans from Kirkwood (1984). Identifications
125 were checked against BAS reference collections. Prey were of unknown size and age.

126

127 2.2 Stable isotope analyses

128

129 All blood and feather samples were homogenised in a freezer mill prior to analysis
130 and oven-dried for >24hrs at 50°C to a constant mass. In the case of prey samples, lipids
131 were extracted prior to SIA over a 4 hour period using Soxhlet apparatus containing 1:1
132 methanol to chloroform solvent mixture. Stable carbon and nitrogen isotope analyses were
133 performed by continuous flow isotope ratio mass spectrometry (CF-IRMS) on 0.7mg sub-
134 samples of material loaded into tin cups and combusted in a Costech ECS 4010 elemental

135 analyser coupled to a Thermo Finnigan Delta Plus XP mass spectrometer. Every 8-10
136 samples were followed by two laboratory standards allowing correction for drift. Isotope
137 ratios are expressed in standard δ notation against international reference standards, vPDB
138 (virtual Pee Dee Belemnite) for $\delta^{13}\text{C}$ and atmospheric nitrogen for $\delta^{15}\text{N}$ according to the
139 equation:

140

$$141 \quad \delta X = [(R_{\text{sample}}/R_{\text{standard}}) - 1] \times 1000$$

142

143 where X is ^{15}N or ^{13}C and R is the corresponding ratio $^{15}\text{N}:^{14}\text{N}$ or $^{13}\text{C}:^{12}\text{C}$. Precision for
144 both $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ was routinely estimated to be $\leq 0.2\%$.

145

146 *2.3 Element analyses*

147

148 Samples were oven-dried for 24hrs at 50°C to a constant mass. Between 0.05 and
149 0.1g dry weight of sample (mean weights for each tissue type; 0.6g for prey, 0.09g for
150 blood, and 0.05g for feathers) was weighed accurately and added to 2ml nitric acid and left
151 to cold digest for 24hrs. Samples were then hot digested for 50 minutes at 120°C, after
152 which 0.5ml of hydrogen peroxide was added and left for a final 15 minutes. The samples
153 were made up to 5ml with deionised water. Total absolute concentrations for 22 essential
154 and non-essential elements were measured using an ELAN 6100 dynamic reaction cell
155 inductively coupled plasma mass spectrometer (DRC ICP-MS) (PerkinElmer, Connecticut,
156 US) with a reference material and internal standard run with every 10 unknowns. Spiked
157 samples and blanks were also run. Limits of detection (LoD) for particular elements are in

158 Table 1. Precision and accuracy (Table 2) were measured using replicate samples and
159 certified reference material (TORT-2 lobster hepatopancreas, NRCC, Canada). Element
160 concentrations are expressed throughout as mean \pm SD in ng.g⁻¹ on a dry weight basis,
161 unless otherwise specified.

162

163 *2.4 Data analyses*

164

165 Elements with mean concentrations below LoD (beryllium (Be) for prey; aluminium
166 (Al), scandium (Sc), cobalt (Co), zinc (Zn), antimony (Sb), tungsten (W), and Be for blood;
167 and Sc, caesium (Cs), lithium (Li), Pb, and Be for feathers) are reported in the summary
168 statistics but are excluded from further consideration or statistical analyses. Among the
169 remaining elements, concentrations in some samples in some species were below the LoD
170 (maximum 13% of samples for rubidium (Rb) in feathers, 1-3% of samples for vanadium
171 (V), iron (Fe), molybdenum (Mo), nickel (Ni), Cs, Hg, manganese (Mn), and selenium (Se)
172 in various tissue types) and were assigned a value equal to half the LoD for the particular
173 tissue. Where element concentrations were below the LoD in >30% of samples overall for a
174 particular tissue type (V, barium (Ba), uranium (U), Li, and Pb for blood; V, Sb, and W for
175 feathers), those elements were also included in summary statistics but excluded from
176 subsequent statistical analyses, as in other studies (Borgå et al., 2006). Fe was excluded
177 from all statistical analyses of blood data because of its physiological function in this tissue.

178 For elements that were included in the statistical analysis, we first used principal
179 component analyses (PCA) in Canoco 4.5 for Windows[®] to identify co-variance among
180 elements, and between elements and isotope ratios (the latter entered as passive variables,

181 following Borgå et al., 2006) in seabird tissues. PCA produces ordination axes that reduce
182 the total residual sum of squares among all the variables included (i.e. all elements, $\delta^{15}\text{N}$,
183 and $\delta^{13}\text{C}$). The primary axis (PC1) represents the greatest proportion of component
184 variation among the samples. The secondary axis (PC2) explains the greatest proportion of
185 any remaining variation that is uncorrelated with PC1. Variables are shown as lines, with
186 the length of each line demonstrating relative contribution of a particular variable to the
187 separation of samples in ordination space. Angles between lines represent a greater or lesser
188 degree of co-variance between elements; with acute angles representing the greatest degree
189 of co-variance, a 180° angle demonstrating a negative correlation, and a 90° angle represent
190 elements that are completely un-correlated. All PCA were performed on standardised
191 values, so that those elements with vastly different ranges could be analysed together. We
192 then used univariate general linear models (GLMs) to test the significance of relationships
193 between species, $\delta^{15}\text{N}$, $\delta^{13}\text{C}$, and elements of high factor loadings that were identified
194 through PCA, (elements with ≥ 0.50 factor loadings for PC1 and/or PC2). These analyses
195 were applied to both blood and feather data separately and the most parsimonious
196 univariate models were determined using Akaike Information Criterion (AIC) (Akaike
197 1973). Absolute element concentrations were logarithmically transformed (\log_{10}) in some
198 instances to reduce skewness and heterogeneity. Model residuals were also examined by
199 way of Q-Q plots to assess goodness of fit. Statistical tests were performed in statistical
200 packages R ver. 2.6.2 and SPSS[®] ver. 14.

201

202 **3. Results**

203

204 In total, we analysed samples for 22 essential and non-essential elements. Mercury
205 was detected in all three tissue types but, because of the particular biomagnification
206 properties and risks associated with this element, the results have been examined in greater
207 detail and reported elsewhere; essentially, mercury levels were found to increase with
208 trophic level, indicating strong biomagnification among Procellariiformes (Anderson et al.
209 2009). Fourteen elements were quantifiable in blood (Table 3) and seventeen in feathers
210 (Table 4). Species varied in the extent to which they accumulated different elements and no
211 single species accumulated the highest concentrations of all elements (Tables 3-5). Of all
212 the elements, Fe and Se had the highest concentrations in blood (Table 3), and Fe and Zn
213 the highest concentrations in feathers (Table 4).

214 In prey samples, 21 elements were detected, only Be was below detection limits.
215 Iron and Zn had the highest concentrations of all elements tested for (Table 5). Aluminium
216 concentrations were generally an order of magnitude higher in crustaceans than in fish and
217 cephalopods, and Cd concentrations were likewise some 10-fold higher in several squid
218 species and in *Themisto gaudichaudii*, compared with other species (Table 5). Benthic-
219 feeding fish, such as Patagonian toothfish (*Dissostichus eleginoides*) (García de la Rosa et
220 al., 1997), often had the highest As concentrations. Isotopic analyses indicated a high
221 degree of variation in trophic position ($\delta^{15}\text{N}$) and foraging range ($\delta^{13}\text{C}$) between seabird
222 species (Fig. 1), and considerable trophic overlap between fish and cephalopod prey. The
223 greatest degree of spatial variation (largest range in $\delta^{13}\text{C}$ ratios) in prey was between fish
224 species (Fig. 1).

225 Initial PCA analyses identified strong associations between As concentrations and
226 both $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ in blood (Fig. 2a). There was a similar but weaker association for
227 caesium (Cs), as indicated by the shorter line on the PCA plot (Fig. 2a), but no similar
228 associations for other elements in blood. The distribution of As in ordination space was
229 predominantly governed by white-chinned petrel, suggesting that blood As concentrations
230 were elevated in this species during the breeding season; this association did not occur in
231 feathers (Fig. 2b). There were other similar associations between specific elements and
232 seabird species. The positions of Rb and Cd in ordination space were largely governed by
233 grey-headed albatross, indicating that blood concentrations of these elements were highest
234 in this species during the breeding season (Fig. 2a); again there were no such associations
235 for feathers (Fig. 2b). The ordinal positions of white-chinned petrel and wandering
236 albatross were influenced by $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ in both blood and feathers. Overall, As and Cd
237 (NEE) contributed most to PC1 in blood, while Se (EE), Rb (NEE), and Mo (EE)
238 contributed most to PC2 (Table 2, supplementary material). In contrast, Mn (EE), Fe (EE),
239 Co (EE), As (NEE), and Mo (EE) contributed most to PC1 in feathers, while Se and U
240 (NEE) contributed most to PC2.

241 With the exception of blood As, all elements contributing ≥ 0.50 to factor loadings
242 in the PCA were found to differ significantly between seabird species (Table 6). Univariate
243 GLM analyses identified significant differences between species for a number of elements.
244 Blood concentrations of Cd, Se, Rb, and Mo all differed significantly between species
245 (Table 6), and in feathers, significant differences between species were found in the levels
246 of Mn, Fe, Co, As, Mo, Se, and U.

247 Univariate GLMs demonstrated that there was a significant and positive association
248 between As and $\delta^{15}\text{N}$ in blood, although the pattern of this relationship varied between
249 species, as demonstrated by the significant interaction term in the model (Table 6). Blood
250 As concentrations were negatively associated with $\delta^{13}\text{C}$ across all species (i.e., at
251 community level, Table 6) but the exact nature of this relationship may also have varied
252 between species, the interaction term in the model approaching significance (Table 6).
253 There was a significant relationship between Se and $\delta^{13}\text{C}$ in feathers (Table 6). There were
254 no other significant relationships between element concentrations and trophic position
255 and/or foraging location in either tissue type (Table 6).

256 The positive association between blood As concentrations and trophic position
257 ($\delta^{15}\text{N}$) was consistent with the concept that As is bioconcentrated along food chains. To
258 investigate this further, we examined how As concentrations in prey species varied with
259 $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values (Table 7). Arsenic concentrations differed significantly between
260 taxonomic groups but also varied with $\delta^{15}\text{N}$ (Table 7). However, associations between $\delta^{15}\text{N}$
261 and As concentrations were not consistent between prey groups (Table 7), indicating that
262 associations between the trophic position of species and As are complex (Fig. 3).

263

264 **4. Discussion**

265

266 The large number of elements analysed in this study required us to focus the
267 discussion only on those elements for which a significant relationship was identified among
268 the parameters in the univariate analyses. We have also focused particularly on those

269 elements for which a significant relationship occurred in blood as blood is the better tissue
270 for characterising trophic and spatial patterns in relation to element composition within
271 these birds. This is because feathers have the potential to acquire element burdens both
272 directly through diet and from remobilisation of elements from internal tissues (Lewis and
273 Furness, 1991; Monteiro, 1996). This may be why the relationships identified between
274 stable isotope ratios and elements in blood in the present study were not always mirrored
275 with feathers.

276

277 *4.1 Arsenic*

278

279 Of the elements analysed in seabird blood, only As showed significant interactions
280 with $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ at the community level (Table 6), suggesting that trophic position
281 and/or foraging range affect As assimilation by Procellariiformes during the breeding
282 season. The significant and positive association between blood As and $\delta^{15}\text{N}$ may be due to
283 bioconcentration or reflect changes in isotopic baseline that coincide with variation in
284 environmental As concentrations. We found no evidence of bioconcentration of As through
285 the foodchain. Although As concentrations varied significantly between prey groups
286 (crustacea, cephalopods, and fish), there was no consistent relationship between As
287 concentrations and trophic position in prey (Table 7, Fig. 3).

288 Given the lack of clear evidence of bioconcentration, it is possible that foraging
289 location is the prime factor affecting As assimilation in Procellariiformes during the
290 breeding season, and any associations with $\delta^{15}\text{N}$ may be coincidental. In this respect, As
291 strongly influenced the position of white-chinned petrel in the PCA ordination space (Fig.

292 2a), reflecting their high blood As concentrations (2500 ng g⁻¹ dry weight) relative both to
293 other species in this study and to other species elsewhere; reference values for blood As in
294 birds from uncontaminated areas is 20 ng g⁻¹ wet weight (Burger and Gochfeld, 1997),
295 approximately equivalent to a dry weight red blood cell concentration of 400 ng g⁻¹.
296 White-chinned petrels have amongst the greatest foraging distances of any seabird and
297 breeding birds from South Georgia regularly forage on the northern Patagonian shelf,
298 >2000km from the colony (Phillips et al., 2006). Prey food webs on the Patagonian shelf
299 have greater trophic complexity than those in waters around South Georgia (Forero et al.,
300 2004, 2005), and so δ¹⁵N levels in birds feeding in that region may be elevated, reflecting
301 longer food chain lengths. The Patagonian shelf region is also more likely to be subject to
302 greater anthropogenic inputs from riverine influxes, mineral activities, and industry than the
303 more remote regions of the Southern Ocean (González-Solís et al., 2002), which could
304 explain the elevated arsenic levels in birds that feed there. It is also possible, however, that
305 enhanced environmental exposure to As may stem from natural processes. For example,
306 disruption in the phosphorus/nitrogen ratio could lead to greater As accumulation by
307 phytoplankton in a particular region and subsequently enhance As levels in the food chain.

308 In our study, there was no relationship between feather As concentrations and δ¹⁵N.
309 Feather As concentrations were highest in Antarctic prion (APR), but only about half the
310 concentrations measured in black-footed albatross (*Phoebastria nigripes*) feathers from the
311 North Pacific (Fujihara et al., 2004). Low feather As concentrations in Antarctic
312 Procellariiformes may reflect exposure during the non-breeding season to relatively low-
313 level natural levels of As. In our study, As concentrations in prey tended to be highest in
314 benthic/demersal species (Table 5). This suggests that a major source of As in the prey of

315 Antarctic Procellariiformes around South Georgia may stem from metallic mineralisation of
316 the seabed rather than from atmospheric transport of particles. Indeed, marine sediments are
317 thought to be the largest geochemical reservoir of arsenic with residence times of ca. 100
318 million years (Maher and Butler 1988). Furthermore, manganese (Mn) concentrations were
319 also elevated in benthic species, again suggesting a natural source of As; elevated As levels
320 *without* matching high levels of Mn are thought to be indicative of anthropogenic As input
321 (Peterson and Carpenter, 1986).

322 Although there was a significant relationship between $\delta^{13}\text{C}$ and blood As
323 concentrations in the Procellariiformes in our study, no such relationship occurred among
324 prey species. This may be because prey were collected from a small geographic area,
325 relative to the foraging areas of the birds. However, there was still considerable variation in
326 $\delta^{13}\text{C}$ ratios between prey species (Fig. 1) and this probably reflects the influence of depth
327 on prey $\delta^{13}\text{C}$ ratios. The lack of association between $\delta^{13}\text{C}$ and As in prey indicates that As
328 uptake by seabirds is unlikely to be influenced by the proportion of mesopelagic prey in the
329 diet, unlike patterns of Hg uptake which are heavily influenced by prey type (see Monteiro
330 et al., 1996; Thompson et al., 1998). The association observed between $\delta^{13}\text{C}$ and As in
331 Procellariiformes is most likely due to geographical variation in background As
332 concentrations within the marine environment.

333

334 4.2 Cadmium

335

336 Blood cadmium concentrations in seabirds varied significantly between species but
337 not with isotopic signatures, indicating that neither trophic position nor foraging location

338 dictated Cd burdens during the breeding season. Our results match those of Gonzalez-Solis
339 et al. (2002) in that Cd concentrations were greater in northern than southern giant petrels
340 (Table 3). Gonzalez-Solis et al. (2002) postulated that the former are exposed to higher
341 levels through foraging over the Patagonian shelf, as the region is thought to be more
342 exposed to heavy metal contamination than the waters immediately adjacent to South
343 Georgia. However, if latitudinal variation in foraging location was the major factor
344 affecting Cd assimilation in Antarctic Procellariiformes, we would anticipate a correlation
345 between Cd concentrations in birds and $\delta^{13}\text{C}$. It may be that foraging location does dictate
346 Cd uptake, but that any such influence reflects the existence of Cd ‘hot spots’ rather than a
347 latitudinal gradient *per se*. In fact, Nygård et al. (2001) linked deep ocean upwelling, which
348 is often naturally Cd-rich (Holm-Hansen, 1985), to high Cd concentrations in Antarctic krill
349 *Euphausia superba* and other prey of Procellariiform seabirds (Petri and Zauke, 1993).

350 Previous studies have attributed high Cd concentrations in Procellariiformes to
351 consumption of squid (Muirhead and Furness, 1988; Honda and Marcovecchio, 1990;
352 Elliott, 2005; Stewart et al., 1999). Our results support this, in that higher Cd levels were
353 found in the blood of grey-headed albatrosses and white-chinned petrels (Table 3). Both
354 white-chinned petrels and grey-headed albatrosses from South Georgia consume large
355 amounts of squid, the latter feeding in particular on *Martialia hyadesi*, and, also on lamprey
356 *Geotria australis* (Berrow and Croxall, 1999, Xavier et al., 2003). In our study, both these
357 prey types had relatively elevated Cd concentrations (Table 5). Hence, high Cd burdens in
358 Procellariiformes do not necessarily indicate anthropogenic contamination. Levels of
359 cadmium in feathers, for example, were largely similar to those in similar species, e.g.

360 black-footed and Laysan albatrosses (*Phoebastria immutabilis*) in the more contaminated
361 North Pacific (Burger and Gochfeld, 2000).

362 Squid are thought to accumulate high Cd concentrations naturally (Kurihara et al.,
363 1993; Gerpe et al., 2000). Elevated burdens in Procellariiformes can also result from
364 amphipod consumption (Cheng et al., 1984; Elliott and Scheuhammer, 1997), and in our
365 study, it was notable that we recorded the highest Cd levels of any prey item in *Themisto*
366 *gaudichaudii* (Table 5). Rainbow (1989) concluded that *Themisto* species are a significant
367 source of Cd in the diets of some seabirds, and our findings are consistent with this as the
368 highest blood Cd levels in birds in our study were in blue petrels, a known predator of *T.*
369 *gaudichaudii* (Prince, 1980). Moreover, hyperiid amphipods may also contribute to the diet
370 of some squid species (Croxall and Prince, 1980) which can then be eaten by seabirds.
371 Thus, dietary preferences, rather than foraging area, may comprise the dominant influence
372 on Cd burdens in Procellariiformes.

373

374 4.3 Selenium

375

376 Blood (hence breeding season) concentrations of Se, an essential element, varied
377 significantly between seabird species, but did not show a significant relationship with
378 trophic position ($\delta^{15}\text{N}$) and foraging location ($\delta^{13}\text{C}$) (Table 6). Therefore, it seems probable
379 that Se does not biomagnify among Procellariiformes. Additionally, our data suggest that
380 foraging location does not dictate Se burdens in Southern Ocean Procellariiformes,
381 although this may in fact reflect the constrained foraging ranges of many species during the
382 breeding season rather than a true absence of association between the two parameters.

383 Selenium concentrations were the highest of any element in Procellariiformes blood except
384 Fe. Moreover, blood Se levels in both giant petrel species were consistent with those
385 previously recorded by González-Solís et al. (2002). Blue petrels and Antarctic prions had
386 the highest blood Se concentrations of the birds we analysed and both species feed
387 predominantly on crustacea, including Antarctic krill (Prince, 1980; Croxall et al., 1997).
388 Of the prey species we analysed, crustacea, especially Antarctic krill and *T. gaudichaudii*,
389 had high Se concentrations relative to other prey species and groups. The physico-chemical
390 form in which Se is sequestered by different prey may also vary and affect subsequent
391 bioavailability. Hence, it would appear that inter-specific variation in blood Se
392 concentrations in seabirds most likely reflects species differences in prey selection.

393 Selenium concentrations in feathers varied considerably between species and were
394 also significantly and negatively related to foraging location (i.e. $\delta^{13}\text{C}$) (Table 6). This was
395 unexpected as concentrations of Se might be expected to increase with decreasing latitude,
396 as birds move into potentially more contaminated waters north of the sub-Antarctic polar
397 front. The negative relationship in this instance is difficult to explain, but could potentially
398 stem from natural sources of Se being more elevated in specific regions closer to the
399 Antarctic continent. In other words, Se burdens accumulated by Procellariiformes in the
400 non-breeding period may be dictated by naturally occurring Se in the marine environment,
401 potentially stemming from Antarctic intermediate and bottom waters which are sporadically
402 rich in organic Se (Cutter and Cutter, 2001). Evidence to support this hypothesis, comes
403 from the comparable levels of Se in the species covered by this study, to those of other
404 studies. Burger and Gochfeld (2000) examined Se in feathers from black-footed and Laysan

405 albatrosses in the North Pacific, and identified levels comparable to those found in this
406 study.

407

408 *4.4 Rubidium and Molybdenum*

409

410 Two other elements, Rb and Mo, also varied between seabird species, but not with
411 either $\delta^{15}\text{N}$ or $\delta^{13}\text{C}$, indicating that trophic position and foraging location were not key
412 determinants in the distribution of these two elements within Procellariiformes. Rb is
413 known to bioaccumulate through the foodchain (Nyholm and Tyler, 2000), and in fact,
414 Campbell et al. (2005) reported a significant correlation between Rb and $\delta^{15}\text{N}$ in marine
415 organisms, potentially suggesting biomagnification. While our results were not significant
416 in the presence of other parameters, there was a significant, albeit weak, positive
417 relationship between Rb and $\delta^{15}\text{N}$ in blood (linear regression, $F_{1,140} = 9.9$, $p = 0.002$, $r^2 =$
418 0.07). However, whether this is evidence of biomagnification of Rb within this particular
419 system is doubtful given there was no significant relationship between Rb concentrations
420 and $\delta^{15}\text{N}$ in prey (linear regression, $F_{1,82} = 1.5$, $p = 0.229$, $r^2 = 0.02$). Given that the
421 accumulated evidence concerning Rb bioconcentration through foodchains is unclear,
422 further investigations into Rb distributions and bioaccumulation in the marine environment
423 appear warranted.

424

425 **5. Conclusions**

426

427 This study presents new data on the distributions and dynamics of heavy metals,
428 essential metals and metalloids in Southern Ocean seabirds. Our results indicate that there
429 are considerable differences in residues accumulated by different species. For arsenic,
430 residues in Procellariiformes appear to be partially explained by changes in foraging range
431 and trophic positioning, although these patterns were not always consistent across species.
432 Arsenic was accumulated at higher concentrations in Procellariiformes feeding at higher
433 trophic levels, although this may be confounded to some extent by individuals foraging
434 across different food webs with varying $\delta^{15}\text{N}$ baselines. There was little or no evidence that
435 trophic level was important in the accumulation of other elements. Our data suggest that
436 differences between species in feeding preferences for prey and the location of the foraging
437 sites are the likely major factors determining element uptake by seabirds in the South
438 Atlantic.

439

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441

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633 **Fig. 1.** Range plots for $\delta^{15}\text{N}$ (a) and $\delta^{13}\text{C}$ (b) values for all Procellariiformes (blood) and
634 prey species. Species are ordered by increasing $\delta^{15}\text{N}$ values within each of the four
635 groupings. Data is shown for Procellariiform blood samples and prey muscle samples.

636

637 **Fig. 2.** Biplot of Procellariiform species mean scores extracted by principal component
638 analyses (PCA) and element loadings on the two principal axes (PC1 and PC2), with
639 standardised element and isotope values in (a) blood and (b) feathers from ten species.
640 APR=Antarctic prion, BBA=Black-browed albatross, BLP=Blue petrel, CDP=Common
641 diving petrel, GDP=South Georgian diving petrel, GHA=Grey-headed albatross,
642 NGP=Northern giant petrel, SGP=Southern giant petrel, WCP=White-chinned petrel,
643 WNA=Wandering albatross.

644

645 **Fig. 3.** Patterns in \log_{10} arsenic concentrations in muscle tissue samples from three
646 Procellariiform prey groups in relation to (a) $\delta^{15}\text{N}$ values and (b) $\delta^{13}\text{C}$ values from the same
647 tissue samples.

648 **Table 1.** Limits of Detection (LoD) for prey, blood, and feather samples (ng g⁻¹).

Element	Prey LoD	Blood LoD	Feather LoD
Al	199	1357	2365
As	3	18	32
Ba	20	136	237
Be	1	7	12
Cd	1	5	8
Co	2	14	24
Cs	1	5	8
Cu	7	45	79
Fe	332	2261	3942
Li	1	7	12
Mn	2	14	24
Mo	10	68	118
Ni	3	23	39
Pb	20	136	237
Rb	1	5	8
Sb	3	23	39
Sc	17	113	197
Se	10	68	118
U	1	5	8
V	7	45	79
W	3	23	39
Zn	332	2261	3942

649

650 **Table 2.** TORT-2 lobster hepatopancreas reference material for elements; expected and
 651 obtained CRM values ($\mu\text{g g}^{-1}$ dry wt.). All elements, where possible, were CRM checked
 652 against an expected value. However, not all elements tested had a reported certified
 653 concentration in the CRM. For these elements only the obtain CRM value is reported.

Element	Expected CRM Value	$\pm 95\%$ CI	Obtained CRM Value	$\pm 95\%$ CI
Al			17.0	0.9
As	21.6	1.8	26.4	0.8
Ba			2.0	0.1
Be			0.01	0.0
Cd	26.7	0.6	28.8	0.8
Co	0.5	0.1	0.5	0.1
Cs			0.0.2	0.0
Cu	106.0	10.0	93.3	3.0
Fe	105.0	13.0	126.0	10.8
Li			0.3	0.0
Mn	13.6	1.2	12.8	0.4
Mo	0.9	0.1	1.1	0.1
Ni	2.5	0.2	2.4	0.1
Pb			0.2	0.0
Rb			2.7	0.1
Sb			0.1	0.0
Sc			0.1	0.0
Se	5.6	0.7	7.5	0.3
U			0.1	0.0
V	1.6	0.2	1.9	0.1
W			0.01	0.0
Zn	180.0	6.0	191.7	12.2

654 **Table 3.** Mean concentrations of elements (ng g⁻¹ dry wt.) ± SD in blood from 10 species of Procellariiformes.

Species	<i>n</i>	Al	As	Ba	Be	Cd	Co	Cs	Cu	Fe	Li	Mn	Mo	Ni	Pb	Rb	Sb	Sc	Se	U	V	W	Zn
Antarctic prion	16	<LoD	431 ± 186	345 ± 289	<LoD	48 ± 58	<LoD	5 ± 4	414 ± 366	2604580 ± 168540	1990 ± 7849	132 ± 66	226 ± 172	2140 ± 723	1052 ± 3203	4161 ± 523	<LoD	<LoD	198425 ± 59904	8 ± 5	85 ± 130	<LoD	<LoD
Black-browed albatross	15	<LoD	373 ± 267	192 ± 234	<LoD	86 ± 177	<LoD	15 ± 10	815 ± 579	4330039 ± 6571356	6 ± 10	150 ± 128	84 ± 113	1728 ± 1286	140 ± 172	4185 ± 1898	<LoD	<LoD	102335 ± 85049	11 ± 15	140 ± 126	<LoD	<LoD
Blue petrel	2	<LoD	388 ± 47	191 ± 174	<LoD	346 ± 27	<LoD	<LoD	71 ± 68	2440536 ± 75575	3 ± 0	91 ± 3	225 ± 46	1652 ± 276	68 ± 0	4368 ± 178	<LoD	<LoD	261757 ± 131638	12 ± 3	<LoD	<LoD	<LoD
Common diving petrel	15	<LoD	389 ± 129	187 ± 170	<LoD	19 ± 29	<LoD	<LoD	639 ± 1010	2265035 ± 355674	3 ± 0	128 ± 78	217 ± 131	1442 ± 589	322 ± 718	3612 ± 625	<LoD	<LoD	84842 ± 29419	9 ± 6	162 ± 285	<LoD	<LoD
S. Georgian diving petrel	15	<LoD	476 ± 197	310 ± 324	<LoD	27 ± 32	<LoD	14 ± 6	196 ± 186	2570513 ± 582271	29 ± 20	190 ± 76	88 ± 68	2007 ± 553	281 ± 446	3824 ± 967	<LoD	<LoD	118897 ± 60846	10 ± 5	60 ± 97	<LoD	<LoD
Grey-headed albatross	15	<LoD	896 ± 380	303 ± 198	<LoD	228 ± 81	<LoD	15 ± 10	670 ± 905	2483096 ± 239518	106 ± 46	252 ± 430	68 ± 80	1936 ± 530	235 ± 648	4526 ± 777	<LoD	<LoD	194381 ± 43946	6 ± 5	<LoD	<LoD	<LoD
Northern giant petrel	16	<LoD	417 ± 787	<LoD	<LoD	88 ± 213	<LoD	7 ± 4	638 ± 246	2333908 ± 659977	60 ± 23	97 ± 38	94 ± 26	1131 ± 966	<LoD	4991 ± 819	<LoD	<LoD	151170 ± 39472	<LoD	75 ± 188	<LoD	<LoD
Southern giant petrel	16	<LoD	184 ± 156	<LoD	<LoD	37 ± 105	<LoD	5 ± 3	690 ± 109	2447292 ± 698505	111 ± 70	202 ± 494	90 ± 13	1480 ± 990	<LoD	5052 ± 712	<LoD	<LoD	151432 ± 77353	<LoD	<LoD	<LoD	<LoD
White-chinned petrel	16	<LoD	2492 ± 1375	<LoD	<LoD	179 ± 60	<LoD	10 ± 5	241 ± 250	2482217 ± 219636	46 ± 31	192 ± 69	127 ± 71	1866 ± 561	<LoD	4532 ± 904	<LoD	<LoD	97951 ± 40289	<LoD	<LoD	<LoD	<LoD
Wandering albatross	15	<LoD	686 ± 377	<LoD	<LoD	50 ± 60	<LoD	17 ± 6	889 ± 455	2398513 ± 681723	98 ± 47	86 ± 17	92 ± 50	1312 ± 970	<LoD	3913 ± 653	<LoD	<LoD	80187 ± 21270	<LoD	<LoD	<LoD	<LoD

655

656

657 **Table 4.** Mean concentrations of elements (ng g⁻¹ dry wt.) ± SD in feathers from 10 species of Procellariiformes.

Species	<i>n</i>	Al	As	Ba	Be	Cd	Co	Cs	Cu	Fe	Li	Mn	Mo	Ni	Pb	Rb	Sb	Sc	Se	U	V	W	Zn
Antarctic prion	16	28396 ± 25222	411 ± 227	1654 ± 1930	<LoD	59 ± 110	605 ± 1295	<LoD	185292 ± 644966	1010868 ± 3457595	<LoD	5254 ± 16921	3411 ± 1216 3	36011 ± 68146	<LoD	153 ± 79	120 ± 116	<LoD	7319 ± 2192	19 ± 9	610 ± 2060	285 ± 571	113658 ± 364227
Black-browed albatross	16	17924 ± 13028	121 ± 39	1481 ± 1802	<LoD	578 ± 246	270 ± 481	<LoD	8613 ± 11995	610216 ± 1657667	<LoD	2704 ± 6669	1686 ± 4057	3284 ± 6669	<LoD	194 ± 27	84 ± 43	<LoD	3381 ± 923	17 ± 3	321 ± 782	128 ± 160	39592 ± 25311
Blue petrel	5	18704 ± 22291	164 ± 88	1065 ± 291	<LoD	74 ± 90	498 ± 329	<LoD	8745 ± 1953	968705 ± 1285689	<LoD	4972 ± 5981	3462 ± 4877	7862 ± 9365	<LoD	123 ± 110	121 ± 82	<LoD	6597 ± 2029	21 ± 7	545 ± 799	47 ± 40	6953 ± 10470
Common diving petrel	16	37468 ± 70275	177 ± 126	1226 ± 1108	<LoD	376 ± 349	140 ± 115	<LoD	31254 ± 58194	130911 ± 119824	<LoD	954 ± 837	581 ± 437	6534 ± 15075	<LoD	140 ± 157	192 ± 283	<LoD	5050 ± 3878	19 ± 14	207 ± 525	21 ± 4	301098 ± 481053
S. Georgian diving petrel	16	65277 ± 34649	261 ± 163	1402 ± 981	<LoD	300 ± 353	419 ± 241	<LoD	20176 ± 15166	791151 ± 572700	<LoD	5077 ± 3490	2979 ± 2133	25442 ± 55886	<LoD	162 ± 115	80 ± 84	<LoD	9797 ± 4917	42 ± 15	865 ± 504	95 ± 94	22283 ± 37587
Grey-headed albatross	16	21372 ± 27043	125 ± 29	769 ± 605	<LoD	196 ± 163	161 ± 101	<LoD	5661 ± 1820	229219 ± 190078	<LoD	1291 ± 940	893 ± 730	12683 ± 14224	<LoD	220 ± 46	61 ± 59	<LoD	5395 ± 1467	17 ± 5	163 ± 113	56 ± 46	50115 ± 20426
Northern giant petrel	16	11944 ± 8969	145 ± 72	641 ± 454	<LoD	83 ± 51	90 ± 31	<LoD	6211 ± 1559	103719 ± 54442	<LoD	608 ± 244	552 ± 229	3811 ± 10126	<LoD	243 ± 55	59 ± 45	<LoD	8377 ± 2346	16 ± 4	124 ± 71	25 ± 14	67557 ± 28067
Southern giant petrel	17	13939 ± 14005	112 ± 36	815 ± 631	<LoD	289 ± 93	101 ± 60	<LoD	6877 ± 3788	95208 ± 42181	<LoD	587 ± 184	509 ± 142	930 ± 1118	<LoD	246 ± 35	<LoD	<LoD	9757 ± 3539	17 ± 6	111 ± 65	25 ± 10	90208 ± 23541
White-chinned petrel	16	13355 ± 15522	127 ± 69	502 ± 407	<LoD	138 ± 132	101 ± 56	<LoD	13110 ± 17785	262076 ± 195334	<LoD	1399 ± 992	1007 ± 646	19449 ± 25581	<LoD	116 ± 21	<LoD	<LoD	4819 ± 1202	8 ± 2	177 ± 120	50 ± 38	77646 ± 17983
Wandering albatross	16	7126 ± 7482	93 ± 30	547 ± 344	<LoD	317 ± 478	114 ± 123	<LoD	6032 ± 2175	166858 ± 276706	<LoD	991 ± 1514	772 ± 1275	1720 ± 2043	<LoD	151 ± 66	<LoD	<LoD	4574 ± 878	<LoD	149 ± 176	40 ± 30	58160 ± 52039

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662 **Table 5.** Mean concentrations of elements (ng g⁻¹ dry wt.) ± SD in muscle tissue of 17 prey species.

Species	n	Al	As	Ba	Be	Cd	Co	Cs	Cu	Fe	Li	Mn	Mo	Ni	Pb	Rb	Sb	Sc	Se	U	V	W	Zn
Crustaceans																							
<i>Euphausia</i>	10	240691	2344	46063	<LoD	176 ± 40	155 ± 83	33 ± 16	22123	372020	463	7797	142 ± 49	1192	167	1960	14 ± 14	299 ± 75	5486	61	1178	37 ± 101	49485
<i>superba</i>		± 194587	± 251	± 27443					± 5644	± 230259	± 284	± 4172	± 605	± 123	± 515	± 14	± 75	± 1141	± 20	± 761	± 2	± 51472	± 18771
<i>Themisto</i>	7	162821	4119	10519	<LoD	19229	4269	10 ± 2	18279	395128	389	3927	692 ± 919	1397	255	724	6 ± 10	516 ± 169	5142	145	971	2 ± 0	± 51472
<i>gaudichaudii</i>		± 56608	± 465	± 1519		± 2453	± 10614		± 4106	± 152043	± 154	± 992	± 1362	± 391	± 116			± 711	± 34	± 314		± 25824	
Fish																							
<i>Champsocephalus</i>	16	8046 ± 7507	3795	1257	<LoD	29 ± 16	38 ± 28	65 ± 40	2053	82261	284	1108	41 ± 45	281	33 ± 88	1830	7 ± 5	65 ± 19	3043	10	189	4 ± 5	43330
<i>gunnari</i>			± 2288	± 1266					± 1734	± 50544	± 221	± 796	± 113	± 923	± 8	± 137	± 14	± 1716	± 518	± 9	± 141	± 2	± 11550
<i>Dissostichus</i>	2	1036 ± 118	9226	168 ± 31	<LoD	1 ± 0	14 ± 1	117 ± 495	924 ± 113	47900	113	923 ± 180	8 ± 2	137	6 ± 6	1716	5 ± 2	63 ± 3	2224	2 ± 0	11 ± 2	2 ± 0	± 11550
<i>eleginoides</i>			± 2770					± 21	± 6075	± 17		± 110		± 224				± 258				± 2978	
<i>Geotria</i>	11	5787 ± 3719	1698	1060	<LoD	1948	30 ± 31	4 ± 3	4241	43360	39 ± 31	558 ± 242	49 ± 24	86 ± 78	19 ± 15	486	5 ± 2	40 ± 21	1815	4 ± 4	207	3 ± 3	± 26612
<i>australis</i>			± 822	± 856		± 1151			± 1370	± 21458				± 184				± 337		± 146		± 12891	
<i>Parachaenichthys</i>	1	4348 ^a	4161 ^a	2241 ^a	<LoD	426 ^a	77 ^a	103 ^a	3089 ^a	92029 ^a	241 ^a	2641 ^a	34 ^a	212 ^a	10 ^a	2081 ^a	7 ^a	48 ^a	2801 ^a	6 ^a	168 ^a	4 ^a	41845 ^a
<i>georgianus</i>																							
<i>Patagonotothen</i>	8	10750	5968	2965	<LoD	168 ± 148	91 ± 115	79 ± 88	2480	285291	930	7854	42 ± 43	476	38 ± 47	2791	14 ± 19	131 ± 158	4100	33	442	11 ± 10	± 63892
<i>guntheri</i>		± 11364	± 6388	± 2770					± 3288	± 369982	± 1170	± 9008	± 705	± 3064				± 4509	± 41	± 564		± 68792	
<i>Pseudochaenichthys</i>	5	4151 ± 1720	5152	426 ± 226	<LoD	14 ± 14	31 ± 18	155 ± 37	1011	104317	190	449 ± 181	9 ± 3	118	20 ± 12	2065	8 ± 2	69 ± 23	2622	4 ± 3	128	2 ± 1	± 33603
<i>georgianus</i>			± 882					± 161	± 75579	± 67		± 67		± 144				± 203		± 180		± 12931	
Cephalopods																							
<i>Galiteuthis</i>	3	19409	1434	14951	<LoD	2105	44 ± 3	8 ± 3	61979	66581	121	1161	793 ± 1109	704	682	578	55 ± 11	117 ± 100	3703	231	379	17 ± 2	± 85928
<i>glacialis</i>		± 8627	± 509	± 10737		± 1745			± 31048	± 11678	± 102	± 213	± 446	± 547	± 230		± 11	± 100	± 346	± 109	± 374		± 34899
<i>Gonatus</i>	2	6378 ± 165	2065	1378	<LoD	901 ± 676	15 ± 1	6 ± 0	16853	19097	18 ± 23	1058	1423	549	10 ± 0	353	39 ± 33	26 ± 25	3162	75	401	2 ± 0	± 94001
<i>antarcticus</i>			± 596	± 835		± 676			± 13774	± 10696	± 23	± 471	± 1894	± 329	± 0	± 39	± 33	± 25	± 271	± 48	± 409		± 111988
<i>Kondakovia</i>	2	7472 ± 3945	1706	2133	<LoD	9216	30 ± 4	23 ± 14	13216	46122	122	1021	181 ± 176	473	24 ± 20	1212	13 ± 10	<LoD	4734	118	246	2 ± 0	± 91585
<i>longimana</i>			± 113	± 1636		± 6413			± 1778	± 11729	± 28	± 117	± 176	± 262	± 20	± 52	± 10		± 381	± 129	± 184		± 35517
<i>Martialia</i>	2	4056 ± 1167	4828	82 ± 14	<LoD	14524	46 ± 5	18 ± 2	28745	19018	139	1083	68 ± 3	353	51 ± 31	4997	1 ± 0	55 ± 8	3498	8 ± 2	109	2 ± 0	± 69918
<i>hyadesi</i>			± 1290	± 844		± 1805			± 10952	± 400	± 25	± 2	± 14	± 464	± 31	± 0	± 8	± 347	± 2	± 34		± 10882	
<i>Moroteuthis</i>	4	4830 ± 3376	2944	976 ± 844	<LoD	10009	32 ± 17	12 ± 5	17517	46102	129	1652	345 ± 551	328	90 ± 148	1888	22 ± 37	46 ± 26	4709	47	222	17 ± 31	± 95067
<i>knipovitchi</i>			± 1689			± 10296			± 16023	± 40016	± 106	± 231	± 142	± 1203				± 1817	± 63	± 183		± 36352	
<i>Psychroteuthis</i>	2	9130 ± 5520	1397	4598	<LoD	2248	37 ± 13	2 ± 2	16654	27134	93 ± 130	774 ± 383	1176	297	61 ± 72	513	10 ± 12	51 ± 62	4024	153	577	2 ± 0	± 24226
<i>glacialis</i>			± 94	± 4137		± 460			± 5875	± 2526		± 1447	± 107	± 72	± 339			± 617	± 152	± 166		± 34026	

663 ^a No SD available due to low sample size (included in table for completeness but excluded in statistical analyses).

664 **Table 6.** Univariate GLM results for element concentrations in blood and feathers of 10
 665 Procellariiform species. Model components were selected by AIC from all possible model sub-sets
 666 with two factorial levels. Parameter estimates (β) shown as Na if non-significant or from multi-level
 667 parameters. Significance values represent values for the parameter within the overall model.

Tissue	Element	Parameter	df	F	<i>p</i>	β	s.e.		
Blood	LogAs ^a	$\delta^{15}\text{N}$	1	13.8	<0.001***	4.2	1.1		
		$\delta^{13}\text{C}$	1	13.9	<0.001***	-2.8	0.7		
		Species	8	1.9	0.061	Na	Na		
		Species* $\delta^{13}\text{C}$	8	1.9	0.067	Na	Na		
		Species* $\delta^{15}\text{N}$	8	2.9	0.006**	Na	Na		
		$\delta^{13}\text{C}$ * $\delta^{15}\text{N}$	1	13.4	<0.001***	0.2	0.1		
	LogCd ^b	Species	9	13.1	<0.001***	Na	Na		
		Se ^c	$\delta^{13}\text{C}$	1	2.8	0.103	-19944.2	23857.4	
	LogRb ^d	LogMo ^e	Species	9	2.4	0.014*	Na	Na	
			Species* $\delta^{13}\text{C}$	1	2.5	0.011*	Na	Na	
			Species	9	4.9	<0.001***	Na	Na	
	Feathers	LogMn ^f	$\delta^{13}\text{C}$	1	3.9	0.052	-0.1	0.03	
			Species	9	7.5	<0.001***	Na	Na	
		LogFe ^g	LogCo ^h	Species	9	4.6	<0.001***	Na	Na
				Species	9	5.0	<0.001***	Na	Na
LogAs ⁱ		LogMo ^j	Species	9	7.45	<0.001***	Na	Na	
			Species	9	5.8	<0.001***	Na	Na	
Se ^k		LogU ^l	$\delta^{15}\text{N}$	1	2.3	0.129	0.03	0.02	
			Species	9	2.2	0.029*	Na	Na	
			$\delta^{15}\text{N}$	1	2.6	0.112	0.2	0.1	
LogU ^l		LogU ^l	Species* $\delta^{15}\text{N}$	9	2.0	0.043*	Na	Na	
			$\delta^{13}\text{C}$	1	5.1	0.025*	-551.4	243.2	
LogU ^l		LogU ^l	Species	9	6.9	<0.001***	Na	Na	
			Species	9	14.2	<0.001***	Na	Na	

668 ^a R² = 0.69 ^g R² = 0.24
 669 ^b R² = 0.48 ^h R² = 0.32
 670 ^c R² = 0.51 ⁱ R² = 0.30
 671 ^d R² = 0.25 ^j R² = 0.34
 672 ^e R² = 0.37 ^k R² = 0.44
 673 ^f R² = 0.23 ^l R² = 0.48
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676 **Table 7.** Univariate GLM results for \log_{10} transformed arsenic concentrations of tissue samples from
 677 prey groups (see Table 5 for groups). Model components specified in table as selected by AIC from all
 678 possible model sub-sets with two factorial levels. Parameter estimates (β) shown as Na for multi-level
 679 parameters.

Parameter	df	F	<i>p</i>	β	s.e.
Group	2	8.9	<0.001***	Na	Na
$\delta^{15}\text{N}$	1	5.6	0.021*	-0.2	0.1
$\delta^{13}\text{C}$	1	3.8	0.054	0.2	0.1
Group* $\delta^{15}\text{N}$	2	9.4	<0.001***	Na	Na
Cephalopod* $\delta^{15}\text{N}$	1	Na	<0.001***	-0.3	0.1
Crustacea* $\delta^{15}\text{N}$	1	Na	0.010*	-0.3	0.1
Fish* $\delta^{15}\text{N}$	1	0 ^b	0 ^b	0 ^b	0 ^b
$\delta^{13}\text{C}$ * $\delta^{15}\text{N}$	1	5.6	0.021*	-0.02	0.01

680 ^a $R^2 = 0.43$

681 ^b This parameter is set to zero because it is redundant.

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