

1	Persistent organic pollutants in tissues of the white-blooded Antarctic fish
2	Champsocephalus gunnari and Chaenocephalus aceratus
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- 25 Abstract
- 26

27 The global occurrence of persistent organic pollutants (POPs) continuously contributes to 28 their accumulation also in remote areas such as the Antarctic Ocean. Antarctic fish, which 29 hold high trophic positions but appear to possess low endogenous elimination rates for 30 chemicals, are expected to bioaccumulate POPs with rising anthropogenic pollution. 31 Using a chemical-analytical method, we measured concentrations of PCBs, PBDEs, HCBs, 32 HCH and DDTs and determined toxic equivalents (TEQs) and bioanalytical equivalents 33 (BEQs) in muscle and ovaries of Antarctic icefish caught in the Southern Ocean around 34 Elephant Island. We used two species with different feeding habits and trophic web positions: the planktivorous Champsocephalus gunnari and the piscivorous Chaenocephalus aceratus. 35 Our results revealed higher contaminant levels in ovary than in muscle tissues of both 36 37 species. Most analytes concentrations and the TEQs (0.2-0.5) and BEQs (0.2) were lower as in temperate species. Comparison with literature data points to higher PCB (20-22 ng g⁻¹ lipid 38 weight (lw)) and DDT (7-19.5 ng g^{-1} lw) concentrations than those measured in icefish in the 39 90's. For the other contaminants, we could not identify temporal trends. We found a higher 40 bioaccumulation of contaminants, particularly HCB and DDTs, in C. aceratus (6.2 & 19.5 ng 41 g⁻¹ lw, respectively) than in C. gunnari (3.8 & 7.0 ng g⁻¹ lw, respectively). However, there 42 43 was no general species-specific accumulation pattern of the different toxicant classes between 44 the two icefish. Thus, the expected link between contaminant burdens of C. aceratus and C. 45 gunnari and their ecological traits was only weakly supported for these species. 46 47 Highlights 48 49 PCB and DDT concentrations in icefish are higher than those measured in the late 90s _ 50 _ Mature, female icefish possess higher contaminant levels in ovaries than in muscle 51 POP levels are similar in fish from different sampling sites around the Antarctic _ 52 Peninsula

- 53 Piscivorous icefish display a higher toxicant accumulation than planktivorous icefish
- Bioanalytical results (BEQs) were similar to chemical analytical calculations (TEQs)
 in icefish
- 56
- 57 Keywords

58 Icefish, bioaccumulation, persistent organic pollutants, polychlorinated biphenyls (PCBs),

59 toxic equivalents (TEQs), DR CALUX bioanalytical equivalents (BEQs)

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- 61 62

1. Introduction

63

64 Antarctica has been less affected by human influences than other continents for a long time, 65 however, contamination with anthropogenic contaminants, in particularly persistent organic 66 pollutants (POPs) has increased progressively (Nash, 2011; UNEP/AMAP, 2011). Nowadays, 67 Antarctica serves as a major sink for highly persistent contaminants. Long-range atmospheric 68 transport, together with global distillation processes and cold condensation, are considered to 69 be the main mechanisms for the progressive contamination of the Antarctic ecosystem, 70 together with local sources such as fishing, tourism and research activities (Simonich and 71 Hites; Feely et al., 2008; Nash, 2011). Particularly POPs such as polychlorinated biphenyls 72 (PCBs) and polybrominated diphenyl ethers (PBDEs), and amongst them also the formerly 73 used insecticides such as γ -hexachlorcyclohexane (γ -HCH) and p, p'-DDT are ubiquitous 74 pollutants with a wide application spectrum. For example, polymer additives such as flame 75 retardants are found globally in building material, furniture, paint, textiles or plastics, which 76 are also used in Antarctic bases and vessels cruising in these regions (Hale et al., 2008; 77 Kohler et al., 2008). This worldwide abundance and the high persistence of POPs 78 continuously contribute to an accumulation in the ice masses and biota of polar regions 79 (Wania and Mackay, 1993; Chiuchiolo et al., 2004; Goerke et al., 2004; Corsolini et al., 80 2007; Bargagli, 2008; Borghesi et al., 2008; Borghesi et al., 2009; Xie et al., 2011; Wolschke 81 et al., 2015).

82

83 Furthermore, those lipophilic organic chemicals have a high potential to bioaccumulate in 84 aquatic biota, and particularly in Antarctic species, which generally possess low endogenous 85 elimination rates for those chemicals (Strobel et al., 2015). Additionally, it is expected that 86 climate warming will lead to the release of those pollutants trapped in glaciers and sea-ice, 87 which additionally contributes to increasing POP levels in the tissues of Antarctic animals (Weber and Goerke, 2003; Bogdal et al., 2010; Schmid et al., 2010; van den Brink et al., 88 89 2011; Cabrerizo et al., 2013; Goutte et al., 2013). As global chemical usage continues to 90 grow, including the usage of persistent and bioaccumulative compounds, it is to be expected 91 that contaminant intake into Antarctica will further increase in the future.

93 The endemic, Antarctic notothenioid fish are evolutionary well-adapted to the cold and stable

94 environment of the Southern Ocean. Adaptations involve e.g. high amounts of tissue lipids,

95 slow growth rates, long life spans and slow metabolism and elimination rates for xenobiotics

96 (Mintenbeck et al., 2012; Strobel et al., 2015). All these factors may favor the

97 bioaccumulation of lipophilic contaminants in the tissues of those fish.

98

99 Lipophilic, non-charged contaminants like PCBs can be taken up via a physico-chemically 100 driven, passive partitioning of the chemicals from the water phase into the lipid phase of the 101 organism and thereby accumulate differentially in tissues or fish species, depending on their 102 lipophilicity of the chemical and the lipid content or composition of the tissues (Nichols et 103 al., 2013). In fact, tissue-specific differences of POP levels are reported for notothenioid 104 species (Lana et al., 2014). Furthermore, bioaccumulation and biomagnification can be 105 related to an organism's habitat and its trophic position within the food web (Corsolini et al., 106 2002a; Weber and Goerke, 2003; Chiuchiolo et al., 2004). Particularly for highly lipophilic 107 compounds, oral uptake via the prey contributes by a major extent to the bioaccumulation of 108 toxicants in the tissues of a species. For example, Weber and Goerke (2003) found higher 109 contaminant levels in the piscivorous icefish Chaenocephalus aceratus than in the 110 planktivorous icefish *Champsocephalus gunnari*, which was apparently related to the 111 different food spectra of the two notothenioids (Goerke et al., 2004). Such studies highlight 112 that POPs are transferred within the Antarctic food web, leading to increasing POP 113 concentrations along the food chain up to high concentrations in top level predators 114 (Wolschke et al., 2015).

115

Considering the trends of rising POP concentrations in Antarctica (UNEP/AMAP, 2011 and their high potential to exert toxic effects in marine biota (Nash, 2011), it is very important to keep monitoring body burdens of the fish living in the Southern Ocean. Yet, we are far from having a solid understanding of the contamination status, time trends, the diversity of contaminants or their toxicity potential in Antarctic biota (UNEP, 2002; UNEP/AMAP, 2011). Particularly the bioaccumulation pattern of contaminants such as the polybrominated flame retardants (PBDEs) or insecticides (e.g. DDT) have hardly been measured in icefish.

124 The aim of this study was to determine levels of selected POPs in muscle and ovary tissue of 125 two white-blooded Antarctic notothenioid species in the study area around Elephant Island,

- 126 the South Shetland Islands and the Antarctic Peninsula, and to estimate temporal trends of 127 fish POP levels in this area. The analytically measured contaminant concentrations were 128 converted into toxic equivalents (TEQs) using standardized values. In order to examine 129 whether food web position or tissue lipid content has an influence on chemical 130 concentrations, we analyzed lipid content and POP levels in fish species of the same family 131 but with different feeding habits and trophic web positions, namely the planktivorous C. 132 gunnari and the piscivorous C. aceratus. 133 Extending the classical contaminants, this study includes polybrominated flame retardants 134 (PBDEs) and insecticides (i.e. DDT, HCB). In addition to chemical analytics, we also applied 135 bioanalytics using the DR-CALUX assay (Murk et al., 1996; Kuiper et al., 2006) to assess the 136 total accumulated dioxin-like activity in the fish tissues. 137 138 139 2. Methods 140 141 2.1 Study species 142 Two species of the Channichthyidae (white-blooded icefish) were fished with bottom trawls 143 down to 500 m, during a cruise with the research vessel 'Polarstern' (ANTXXVIII/4, March 144 13 to April 9, 2012; http://expedition.awi.de/expedition/ANT-145 XXVIII/4?alias=PS79#mapChart) at different, closely located sampling sites around Elephant Island and the South Shetland Islands (61.1°S, 55.1°W). Only fish netted alive and without 146 147 macroscopically visible damage were used for tissue sampling. The planktivorous mackerel 148 icefish, C. gunnari, shows a mainly bentho-pelagic feeding mode, while the piscivorous 149 Scotia Sea icefish C. aceratus is predominantly a benthos feeder (Weber and Goerke, 2003). Persistent organic pollutants were analyzed in muscle and ovary tissue of mature (stage III-150 151 IV), female fish only. Sex and maturity stage of the fish were verified histologically. Animal 152 weight and length are given in Table 1. All tissue samples were wrapped in aluminum foil 153 and stored immediately at -20 °C until used for analysis. 154 155 2.2 Sample preparation 156
- 157 Muscle and gonad samples were defrosted, cut into pieces and lyophilized at 33 Pa for 72
- 158 hours until constant weight. Dried tissue of muscle (5-10 g) or gonads (0.5-2 g) was ground

with anhydrous sodium sulfate and quartz sand in a ceramic mortar and pistil to obtain a finepowder.

This homogenate was then Soxhlet-extracted with a speed-extractor (E-914, Büchi, 161 Switzerland) in 120 mL extraction cells at a constant temperature and pressure of 100 °C and 162 163 100 bar, respectively (4 cycles, hold time 10 min., discharge 4 min), with ~150 ml n-164 hexane/dichloromethane 1:1 (v:v) (Hartmann, 2013). The extract was concentrated in a 165 Syncore evaporator (Büchi, Switzerland) und let dry completely by applying a gentle nitrogen stream. The residue accounted for the fat content of the sample. ${}^{13}C_{12}$ labeled 166 167 internal standards (Schmid et al., 2007) were added to the samples and after the addition of 2 168 -3 mL of *n*-hexane the solution was treated with 3 ml oleum (7% SO₃ in conc. sulfuric acid). 169 After centrifugation for 3 min at 5'000 rpm, the solvent layer with the lipophilic target 170 analytes was removed and the remaining suspension was re-extracted two times more with n-171 hexane. The pooled extracts were concentrated to 0.5 mL in a rotary evaporator at 45 °C and 172 300 mbar. Subsequently, the extract was purified on a multilayer mini silica gel column (from 173 top to bottom: 0.25 g anhydrous sodium sulfate, 0.25 g silica gel 60 with 44% sulfuric acid 174 and 0.25 g silica gel 60 activated at 130 °C). The sample was applied on the column and 175 eluted with 5 ml *n*-hexane, followed by 5 mL *n*-hexane/dichloromethane 1:1 (v:v). The eluate 176 was concentrated using a rotary evaporator to 0.5 mL. After transfer to a mini GC-Vial the 177 volume was further reduced to 30 μ L by the application of a gentle stream of nitrogen at room temperature. Finally the recovery standard ${}^{13}C_{12}$ labeled PCB 70 was added. Samples 178 179 were stored in toluene at -20 °C until analysis. Method blank levels for the whole analytical 180 procedure were determined in duplicates (Table 1 and A.2).

181

182 2.3 Chemical analysis

183

184 PCBs (indicator PCB 28, 52, 101, 138, 153, and 180; dioxin-like PCBs 77, 81, 105, 114, 118,

185 123, 126 156, 157, 167, 169, and 189), DDT (*o*,*p*'-DDT, *p*,*p*'-DDT, and *p*,*p*'-DDE),

186 hexachlorobenzene (HCB), γ-hexachlorocyclohexane (γ-HCH), and PBDEs (BDE 28, 47, 99,

187 100, 153, 154, 183, and 209) were included in this study. Quantitative determination of the

188 target analytes in the extracts was achieved by gas chromatography/high resolution mass

- 189 spectrometry (GC/HRMS). Analyses were carried out on a Finnigan MAT95 high-resolution
- 190 mass spectrometer (Thermo Finnigan MAT, Bremen Germany) coupled to a Finnigan Trace

191 GC Ultra equipped with a Triplus auto sampler (Thermo Electron Corporation, Waltham, MA, USA). Samples were injected in splitless mode (splitless time 30 s) at an injector 192 193 temperature of 260 °C. For the gas chromatographic separation a RTX5 Sil-MS column (30 194 $m \times 0.25$ mm, film thickness 0.10 µm) was used with helium as carrier gas at a pressure of 195 100 kPa. The following temperature programs were used for the different compound classes. 196 For the PCBs, the initial column temperature was 100 °C. After 0.5 min, the temperature was 197 ramped at 20 °C/ min to 180 °C, followed by 3 °C/ min to 250 °C, and 20 °C/ min to 300 °C. 198 For pesticides, the initial column temperature was 100 °C. After 0.5 min, the temperature was 199 ramped at 10 °C/ min to 160 °C, followed by 4 °C/ min to 240 °C, and 20 °C/ min to 300 °C. 200 For the PBDEs, the initial column temperature was 100 °C. After 0.5 min, the temperature 201 was ramped at 20 °C/ min to 220 °C, followed by 6 °C/ min to 300 °C, and 10 °C/ min to 320 °C. The ion source was operated at 220 °C, the electron energy was 70eV, and the mass 202 203 spectrometer was tuned to a mass resolution of 8000-10000. The two most abundant signals of the molecular ion cluster of the analytes and the ${}^{13}C_{12}$ labeled internal standards were 204

- 205 recorded in the single ion monitoring mode.
- 206

207 The analytes were identified by comparing the retention times with those of the labeled

208 internal standards. Quantification was based on peak areas of the analytes and the labeled

209 reference compounds with known concentrations. More details about the method are

available in the literature (Zennegg et al., 2003; Schmid et al., 2007).

211 For the calculation of TEQs, the dioxin-like PCBs were determined in the same way, but as

212 part of the sample extract was used later on for the analysis by the bioassay (DR-CALUX),

213 no ${}^{13}C_{12}$ -labeled internal dl-PCBs standards could be used for the quantification, as the

214 isotope labeled analogues exhibit similar activity in the DR-CALUX. Therefore, quantitative

215 determination of dl-PCBs was based on the ${}^{13}C_{12}$ labeled indicator PCBs used as internal

216 standards and previously determined response factors to native dl-PCBs.

The blank values (in ng g^{-1} lipid weight) were all below the compound concentrations and are given in Table 1 and A.1.

219 The limit of detection (LOD) and the limit of quantification (LOQ) were set by definition at

signal to noise ratios of greater than three $(s/n \ge 3)$ and ten $(s/n \ge 10)$ respectively. All glass

221 ware used were cleaned with strongly alkaline detergents and backed out overnight in a

222 ceramic oven at 450 °C. Directly before use, the glass ware was rinsed with solvents (n-

223 hexane, dichloromethane).

- 225 2.4 Determination of dioxin-like toxic equivalents (TEQs)
- 226
- In order to assess total dioxin-like activity in the fish tissues, we calculated the 2,3,7,8-
- 228 tetrachlorodibezo-p-dioxin (TCDD) equivalent (TEQ) concentrations of dl-PCBs in the
- 229 muscle and gonad extracts of both icefish species using Toxic Equivalency Factors (TEFs)
- proposed by the World Health Organization (WHO) for fish (Van den Berg et al., 1998).
- TEQs were calculated as the sum of the TEFs of all dl-PCBs listed in Table A.1. Although
- the TEF values are not derived from studies with Antarctic fish, since TEFs are only available
- for salmonids (Van den Berg et al., 1998) they provide a useful tool for a reasonable estimateof toxicity effects of PCBs on Antarctic fish.
- 235
- 236 2.5 Determination of bioequivalent values (BEQs)
- 237
- The same extracts that were used for chemical analysis were also used in a standard bioassay,
- the DR-CALUX (Dioxin Responsive Chemically Activated Luciferase Gene Expression)
- assay. This cell and receptor based reporter gene bioassay measures the binding of dioxin-like
- 241 HAHs, e.g. dioxin-like PCBs (dl-PCBs), to the Aryl hydrocarbon receptor (AhR) via
- activation of a reporter gene. The assay was performed by BioDetection Systems b.v.
- 243 (Amsterdam, The Netherlands).
- 244

Bioassay-derived TEQ values, or bioequivalent values (BEQs), were compared with TEQ
values calculated from the chemical analytical data. This comparison revealed if there was a
significant higher activity than predicted from the chemical analysis. Such a combined
approach has been repeatedly used in studies on chemical contamination of the Arctic region
(Letcher et al., 2010), but it has not been used in studies with Antarctic fish so far.

- 250
- 251 2.5 Calculations and statistics
- 252
- 253 All analyses include the lipid content of the respective tissue. POP concentrations were
- normalized against both lipid weight (lw, ng g^{-1}) and fresh weight (fw, ng g^{-1}) of the tissue
- and provided as mean values \pm standard error of the mean (sem). Toxicant concentrations
- 256 were compared among icefish species and tissues using ANOVA with Tukey post-hoc test.

257	Influence of sampling site, fish weight and length, maturity stage and tissue lipid content was
258	tested with ANOVA. Data were considered to be statistically significant at $p < 0.05$. Normal
259	distribution of data was tested with Kolmogorov-Smirnov and equality of variances with
260	Bartlett's test. All statistical tests were performed with STATISTICA 12, StatSoft, Inc., and
261	GraphPad Prism 5, GraphPad Software, Inc.
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264	3. Results
265	
266	3.1 Interspecies comparison of contaminant patterns and levels
267	
268	In this study, we measured lipid content, PCB, DDT, HCB, γ -HCH, and PBDEconcentrations
269	in both muscle and ovary tissue of female, mature <i>C. gunnari</i> and <i>C. aceratus</i> from spatially
270	closely located sampling sites around Elephant Island and the South Shetland Islands. We
271	express the compound concentrations on lipid weight and the fresh weight basis.
272	Sampling site, maturity stage and tissue lipid content of the fish showed no correlation to the
273	accumulation of any of the compounds analysed in this study.
274	All samples contained detectable levels of the target compounds, which are presented in
275	Table 1. Analytes concentrations were about 15 - 110 times higher when calculated per lipid
276	weight than based on tissue fresh weight.
277	Overall, the PCBs were the predominant group among all compounds analysed in our study
278	(mean values \pm sem (ng g ⁻¹ lw): <i>C. gunnari</i> muscle 20.0 \pm 4.3, <i>C. gunnari</i> ovaries 47.8 \pm
279	10.2, <i>C. aceratus</i> muscle 21.9 ± 4.9 , <i>C. aceratus</i> ovaries 31.9 ± 7.3), followed by the DDTs
280	(mean values \pm sem (ng g ⁻¹ lw): <i>C. gunnari</i> muscle 7.0 \pm 1.0, <i>C. gunnari</i> ovaries 6.9 \pm 1.1, <i>C.</i>
281	<i>aceratus</i> muscle 19.4 \pm 8.1, <i>C. aceratus</i> ovaries 17.8 \pm 3.9), HCB (mean values \pm sem (ng g ⁻¹
282	lw): C. gunnari muscle 3.8 ± 0.5 , C. gunnari ovaries 2.8 ± 0.4 , C. aceratus muscle 6.2 ± 0.9 ,
283	<i>C. aceratus</i> ovaries 7.3 ± 2.0), the PBDEs (mean values \pm sem (ng g ⁻¹ lw): <i>C. gunnari</i> muscle
284	4.1 ± 0.5 , C. gunnari ovaries 0.7 ± 0.1 , C. aceratus muscle 5.1 ± 0.8 , C. aceratus ovaries 10.0
285	\pm 1.9) and γ -HCH (mean values \pm sem (ng g ⁻¹ lw): <i>C. gunnari</i> muscle 0.6 \pm 0.2, <i>C. gunnari</i>
286	ovaries 1.3 ± 0.3 , <i>C. aceratus</i> muscle 0.6 ± 0.2 , <i>C. aceratus</i> ovaries 1.2 ± 0.5) (Figure 1).
287	Within the PCBs, only PCB 28 and 101 were significantly different between the ovaries of <i>C</i> .
288	gunnari and C. aceratus (Table 1). When we calculated the percentage contribution of the
289	individual PCB congeners to \sum PCBs (lw) of both muscle and ovary tissue, there was no

290 difference in PCB congener composition between the two icefish species (two-way

- 291 ANOVA).
- Among the PCBs, PCB 153 contributed by 28% in muscle and by 26% in ovaries to all PCBs
- in both species. The second-most abundant PCBs were PCB 138 (~26%) and 101 (~21%)
- 294 (Figure 2).
- 295
- In the ovaries, levels of *p*,*p*'-DDE, *o*,*p*'-DDT and *p*,*p*'-DDT (per lw) were significantly
- 297 higher in *C. aceratus* compared to those of *C. gunnari*. Also *p,p*'-DDT concentrations in the
- 298 muscle of *C. aceratus* were higher than those in muscle of *C. gunnari* (per lw). Σ DDT (per
- lw) was significantly higher in muscle and ovaries of *C. aceratus* than in *C. gunnari* (Figure
- 300 1). Among the DDTs, *p*,*p*'-DDE showed the highest concentration of up to 57% in muscle of
- 301 *C. aceratus. p,p* '-DDE tended to be higher in muscle tissue, while *p,p* '-DDT was slightly
- 302 higher in ovary tissue in both species. *o*,*p*'-DDT was slightly higher in *C*. *gunnari* with
- 303 ~22%, while it was only ~16% in *C. aceratus*. Detailed information on the contribution of all
- individual compounds measured in this study is given in Table A.2 in the appendix.
- 305γ -HCH was only different between the muscle tissues of the two icefish (on fw basis).
- 306 HCB concentrations per lipid weight in both muscle and ovaries were higher in *C. aceratus*
- 307 than in *C. gunnari* (Figure 1). Except HCB, none of the analysed compounds exhibited a
- 308 correlation with fish weight or length. HCB showed a significant linear dependence of fish 309 weight and length in *C. aceratus*, except in the ovaries ($R^2=0.723$).
- 310
- In both species, the PBDEs were dominated by BDE 47 (~59% in muscle, ~54% in ovaries),
- followed by ~20% BDE 99. On both the lipid and the fresh weight basis, BDE 183 and 197
- 313 were significantly different between the two tissues types and fish species. Also BDE 99
- 314 showed different concentrations between the ovaries of *C. aceratus* and *C. gunnari* on the
- 315 lipid basis (Table 1, Table A.2, Figure 2).
- 316

317 3.2 Comparison of contaminant patterns and levels in different tissue

- 318
- 319 Lipid content was significantly higher in the ovaries of *C. gunnari* than in the ovaries of *C*.
- 320 *aceratus*. *C. gunnari* exhibited significantly higher lipid content in ovaries than in their
- 321 muscle tissue, while this was not the case for *C. aceratus* (Table 1).
- 322 A comparison between the compound concentrations in muscle and ovary tissue for each
- 323 icefish species revealed higher HCB concentrations in ovary than in muscle tissue of *C*.

324	gunnari. Concentrations of γ -HCH and DDTs in muscle and ovary tissue (per fresh weight)
325	were significantly different in both species (Table 1).
326	In C. aceratus, all PCB concentrations and most of the PBDEs were different from each other
327	in both muscle and ovary tissue. In case of dl-PCB congeners, the ∑dl-PCBs were higher in
328	muscle of C. gunnari than in muscle C. aceratus on both lipid and fresh weight basis (Table
329	A.1).
330	In C. gunnari, all PBDEs apart from congener 183 did differ between muscle and ovary
331	tissue. In contrast, only PCB 28 and 180 were different between the tissues of C. gunnari, all
332	other PCB congeners showed no significant tissue differences (Table 1, Figure 1).
333	
334	3.3 Toxic equivalents (TEQs)
335	
336	The toxic equivalent levels of dioxin-like PCBs (WHO _{PCB} -TEQ g^{-1} fw) were higher in the
337	muscle of C. gunnari (0.5 ± 0.1) than in C. aceratus (0.2 ± 0.1) , while C. aceratus exhibited
338	higher TEQs (0.5 ± 0.9) in the ovaries than <i>C. gunnari</i> (0.1 ± 0.0) (Table 1).
339	
340	3.4 Bioanalytical equivalents (BEQs)
341	
342	The bioanalytical equivalents (BEQs) determined by DR-CALUX in the muscle of both
343	species were in a similar range (<i>C. aceratus</i> : 0.15 ± 0.07 , <i>C. gunnari</i> : 0.2 ± 0.14 BEQ g ⁻¹ fw)
344	to the WHO _{PCB} -TEQs g^{-1} lw, which were calculated using the concentrations of dioxin-like
345	PCBs. In C. gunnari, however, the BEQs measured in the ovaries were about 70-fold higher
346	(BEQ: 5.14 ± 3.6 , TEQ: 0.07 ± 0.04) than the calculated WHO _{PCB} -TEQ g ⁻¹ fw (Table 1).
347	
348	
349	4. Discussion
350	
351	4.1 Species-specific contaminant patterns and levels
352	4.1.1. Species-specific patterns
353	
354	Literature data on present POP concentrations in tissues of Antarctic fish are scarce, and
355	particularly those on contaminant levels in white-blooded icefish.
356	Earlier studies of the 80's and 90's measured PCB concentrations in the icefish C. gunnari
357	and C. aceratus around the Antarctic Peninsula (Corsolini et al. 2002a, 2002b, 2009), which

- 358 were two to four times lower (0.7 ng g^{-1} fw in muscle) than the ones of the present study.
- 359 HCB levels of our icefish were within a similar range to those measured previously. It is
- 360 known for sub-Antarctic fish that PCB levels can follow a seasonal trend, which is related to
- the release of pollutants from melting snow and ice during summer and the atmospheric
- 362 transport of loads of pollutants which precipitate in Antarctic regions and are released during
- 363 warming (Jaffal et al., 2011). Since sampling of our icefish took place in Austral autumn,
- after the seasonal ice melting in Antarctica, a seasonal release of POPs trapped in sea ice
- 365 could thereby be one contributor to the comparably high levels of PCBs in our icefish
- 366 samples. In fact, the PCB concentrations (per fw) of our icefish were within a similar range to
- those of salmon from the Baltic Sea (Isosaari et al., 2006).
- In the early 00's, Borghesi et al. (2008; 2009) sampled *C. hamatus* and *C. gunnari* in the
- Ross Sea at 74° South. They found \sum PCB concentrations in *C. hamatus* muscle of 0.35 ng g⁻¹
- 370 fw, \sum non-ortho PCBs of 5 ng g⁻¹ fw and \sum PBDEs 0.16 ng g⁻¹ fw. In muscle of *C. gunnari*,
- 371 they report \sum PBDEs of 0.44 ng g⁻¹ fw. Those values are about five times higher than the
- size class, this difference is likely related to the measurement of different PBDE congeners inthe studies.
- 375 Another recent study measured contaminant concentrations (in ng g^{-1} lw) in muscle, liver and
- 376 gonads of three red-blooded Antarctic species, Notothenia coriiceps, N. rossii and
- 377 *Trematomus newnesi*, from Potter Cove, Antarctic Peninsula (Lana et al., 2014). They
- 378 reported similar \sum DDT values in muscle, but also species differences in the ovaries of their
- 379 fish: while *N. coriiceps* and *T. newnesi* had similar values to *C. aceratus* measured in our
- 380 study, *N. rossii* had much higher values than our fish. They also measured γ -HCH values
- about six-times higher than in our fish. N. rossii displays a rather benthic lifestyle, but also
- feeds on pelagic species, and thus has a prey spectrum similar to C. gunnari. Nevertheless,
- 383 the DDTs and γ -HCH were much higher in *N. rossii* than in *C. gunnari*. In contrast, \sum PCB
- and \sum BDEs were highly variable among the species and tissues measured by Lana et al.
- 385 (2014), but were generally within the same order of magnitude compared to our data. Thus,
- 386 such strong differences in DDT and γ -HCH accumulation patterns between Antarctic fish
- 387 species could also be related to a selective metabolism for individual contaminant classes
- between the species (Storelli et al., 2009), and not only to their ecological traits.
- 389
- Amongst the PBDEs, BDE 47 showed the highest concentration (60% of all congeners) in
 our icefish amongst all PBDE congeners, followed by BDE 100 and BDE 99. This is in line

- 392 with the general picture of those congeners being the dominating PBDEs in fish around the
- globe, i.e. BDE 47 being recognized as the most important PBDE congener in marine biota
 (Zennegg et al., 2003; Isosaari et al., 2006; Kuiper et al., 2006).
- 395 In comparison to fish from non-Antarctic regions, PBDE concentrations on the fresh weight
- 396 basis were particularly low in icefish. For example, PBDE concentrations range from about
- 1.0 to 8 ng g⁻¹ fresh weight, or up to 64 ng g⁻¹ lipid weight in various fish species from the
- Baltic Sea (Isosaari et al., 2006), which are at least ten times higher than in the icefish.
- 399 Although the production of the former widely used brominated flame retardants penta- and
- 400 octabromodiphenyl ether (PentaBDE and OctaBDE) were banned by the European Union in
- 401 2004 and several states of the USA, toxic and persistent lower brominated PBDEs are still
- 402 produced in other areas of the world and redistributed globally, also to the Antarctic (Cox and
- 403 Efthymiou, 2003; Renner, 2004; Vives et al., 2004; Kuiper et al., 2006).
- 404 HCB concentrations (per fw) were also up to 30 times lower, and DDT concentrations (per
- 405 fw) several hundreds of times lower in our icefish than in fish from the Northern hemisphere
- 406 (Sharma et al., 2009). Despite a worldwide stop of the production of DDTs during the 70's, it
- 407 has been reintroduced in the 2000s as malaria control by the WHO, and about 6000 tons of
- 408 DDTs are still produced per year (UNEP, 2008). Due to its high persistence, bioaccumulation
- 409 potential and cold condensation processes, DDTs and its metabolites are nowadays found in
- 410 biota all over the world, and particularly in polar regions (Mirmigkou and de Boer, 2015).
- 411
- 412 The TEQs we calculated for *C. aceratus* and *C. gunnari* were in a similar range than TEQs
- 413 reported for muscle of the icefish C. hamatus (TEQ 0.01-0.1 pg g^{-1} wet weight) or red-
- 414 blooded Antarctic fish (TEQ ~ 0.1 pg g^{-1} wet weight) (Focardi et al.; Corsolini et al., 2002a;
- 415 Borghesi et al., 2008). Generally, the values for Antarctic fish were lower than those for other
- 416 organisms living in less remote parts of the world, e.g. WHO_{PCB} -TEQ (pg g⁻¹ fw) of 3 to 15
- 417 in muscle of salmon or Baltic herring. Yet, burbot from Bothnian bay (Baltic Sea) exhibit
- 418 muscle TEQs, which are in a similar range as in Antarctic fish (Isosaari et al., 2006).
- 419 In *C. aceratus*, we found that the BEQs were comparable to the TEQs calculated on the basis
- 420 on the WHO_{PCB}-TEFs for salmonid fish species (Van den Berg et al., 1998). In comparison to
- 421 fish from the northern hemisphere (Husain et al., 2014), Antarctic fish had about twenty
- 422 times lower BEQ values in their muscle. In contrast, the BEQs in the ovaries of *C. gunnari*
- 423 were much higher than the WHO_{PCB}-TEQs calculated for their ovaries. The toxicity effects
- 424 might thus be actually much higher than expected by the single usage of the calculated
- 425 WHO_{PCB}-TEQ. In fact, the BEQs in the ovaries of *C. gunnari* were within the same range as

- 426 BEQs of fish from temperate latitudes. Also Corsolini et al. (Corsolini et al., 2002a) stated
- 427 that the TEQ values they measured in Antarctic fish were already half as high as those values
- 428 which are considered to elicit toxicological effects, such as reproductive and immunological
- 429 disorders, in marine mammals or birds (Kannan et al., 2000).
- 430 Rising PCB concentrations as observed by us in tissues of Antarctic fish will thus
- 431 increasingly have the potential to exert their toxic effects on those fish.
- 432
- 433 4.1.2. Temporal trends in contaminant levels
- 434

435 Long-term observations on contaminant levels in Antarctic biota are scarce. Yet, Weber and

436 Goerke (2003; Goerke et al., 2004) measured contaminant levels in the liver of *C. aceratus*

- 437 and *C. gunnari* in the same sampling area as in the present study. From 1987 to 1996, the
- 438 authors measured an increase of PCB 153 and PCB 180 levels in *C. aceratus*, but not in *C.*
- 439 gunnari. In the present study, we found about three times higher concentrations of those PCB
- 440 congeners in the tissues of our two icefish species than in the previous study from 1996
- 441 (Weber and Goerke, 2003). In contrast, the HCB concentrations show a declining trend from
- the 1987 study to our current survey. A similar trend of stable or declining HCB levels by up
- to 2.5% per year has also been observed in Arctic biota, such as birds, fish or marine
- 444 mammals, since the late 80's (Barber et al., 2005; Rigét et al., 2010).
- 445

In liver of *C. aceratus*, concentrations of p,p'-DDE had already increased from 1987 to 1996, and the values we measured in muscle and ovaries of *C. aceratus* were almost twice as high as in 1996 (Weber and Goerke, 2003). In contrast, p,p'-DDE concentrations in *C. gunnari* remained at similar levels from 1987 over 1996 (study by Weber and Goerke (2003)) to the present study.

Nevertheless, overall DDT concentrations were increasing from 1987 to 1996 in both icefish species, and our values are also slightly higher than those in 1996, suggesting an increasing trend of DDTs in icefish around the Antarctic Peninsula. Despite a general global reduction of DTT as an insecticide, DDT is still produced at high volumes (see above, (UNEP, 2008)). Furthermore, climate warming on the one hand leads to an increased volatility and worldwide distribution of DDTs, and on the other hand local sources such as melting glaciers may

- additionally contribute to increasing DDT concentrations in tissues of Antarctic fish around
- 458 the Antarctic Peninsula (van den Brink et al., 2009; van den Brink et al., 2011).
- 459

- 460 4.2 Correlation of contaminant concentrations with ecological traits
- 461

462 4.2.1 Tissue-specific patterns

463

464 Since all POPs analyzed in this study are highly lipophilic substances, tissue differences in 465 the contaminant levels in C. aceratus and C. gunnari on fresh weight basis at first instance 466 should correlate to tissue lipid concentrations (Corsolini et al., 2002a; Weber and Goerke, 467 2003; Chiuchiolo et al., 2004). Indeed, C. gunnari, which possesses a clearly higher lipid 468 content in the ovaries than C. aceratus, accordingly showed significantly higher PCB 469 concentrations in its ovaries based on fresh weight compared to C. gunnari. Also HCB was 470 two times more concentrated in the fat-rich ovaries of C. aceratus than in C. gunnari, but 471 only on the lipid basis. Accordingly, the BEQs were higher in the ovaries of C. aceratus and 472 C. gunnari than in their muscle tissue. Also Lana et al. (2014), investigating POP 473 accumulation patterns in notothenioid species, reported that the highest levels were found in 474 the gonads of the fish.

475

476 4.2.2 Ecological-related patterns

477

In addition to body lipid contents, our results point to an influence of habitat and trophic levelon POP levels in white-blooded icefish. The benthic-living *C. aceratus* had two-times higher

480 concentrations of almost all DDT congeners and HCB than the bentho-pelagic *C. gunnari*.

481 Also the previous study by Weber and Goerke (2003) report higher (lipid-based) contaminant

482 burdens in *C. aceratus* than in *C. gunnari* and highlight the higher tendency of *C. aceratus* to

483 accumulate DDTs in its tissues than *C. gunnari* over the time.

484 Since POPs accumulate in sediments, benthic fish species are generally thought have a higher

485 exposure and uptake of lipophilic contaminants (Goerke et al., 2004; Borghesi et al., 2008).

486 In addition to uptake from water or sediment, also the feeding habit thus plays a role, with

487 species at higher tropic levels tending to show higher contaminant accumulation due to

488 biomagnification. A recent study by Wolschke et al. (2015) also highlights the

489 biomagnification of POPs from lower to higher trophic levels in the Antarctic food chain,

490 which can be attributed to the diets of the animals.

491 From these observations, the benthic, piscivorous *C. aceratus* was expected to have higher

492 contaminant burdens than the bentho-pelagic, planktivorous C. gunnari. However, only HCB

- 493 and DDTs, but none of the other congener classes, were higher in the predominantly benthic
- 494 *C. aceratus* than in the bentho-pelagic *C. gunnari* on the lipid weight basis.
- 495
- 496

497 Conclusion

- 498
- 499

500 Overall, PCB and DDT concentrations tend to rather increase than decrease in tissues of the 501 two white-blooded icefish species C. aceratus and C. gunnari around Elephant Island and the 502 South Shetland Islands, when compared to earlier studies. Our results thereby support the 503 global transportation of POPs to the Southern Ocean and their bioaccumulation in the local 504 marine fish, and point to a trend of increasing concentrations of POPs in Antarctic icefish. 505 Our data also suggest that worldwide climate change effects may contribute to an increased 506 volatilization and release of POPs trapped in glaciers, sea- or pack-ice, thereby leading to an 507 ongoing contamination of the Southern Ocean and its biota.

- 508 Furthermore, we found differences in POP accumulation patterns between the two icefish
- 509 species, which were weakly correlated to their trophic position. The piscivorous *C. aceratus*
- 510 showed a higher potential to accumulate contaminants in its tissue than the planktivorous C.
- 511 *gunnari*. This species difference highlights the influence of intake of POPs via the specific
- 512 prey of individual fish species. However, the expected link between the contaminant burdens
- 513 of *C. aceratus* and *C. gunnari* and their ecological traits could not be fully supported.
- 514 Additional factors, such species differences in toxicant metabolism rates and selective
- 515 metabolism for single contaminant classes, may also play an important role in defining
- 516 chemical bioaccumulation patterns in Antarctic fish species in the long term. In the end,
- 517 Antarctic fish are a central link between the benthic community and top level predators,
- 518 concomitantly POPs bioaccumulated in their tissues are likely to contribute to a progressive
- 519 biomagnification of POPs along Antarctic food webs.
- 520 521

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- 527 Tables & Figures
- 528
- 529 Tables
- 530
- Table 1: Lipid content (%) and levels of organic contaminants (ng g⁻¹ lipid weight; ng g⁻¹ fresh weight) (mean \pm sem) in tissues of two Antarctic icefish species.
- 533

	C. aceratus (n=	=10)					C. gun	nari (n=	11)						
Length (cm)	47-66						33-50								
Weight (g)	636-3620		252-888												
	lipid weight (ng	g g ⁻¹ lw)	fresh v	fresh weight (ng g^{-1} fw)				eight (n	g g ⁻¹ lw)		fresh weight (ng g ⁻¹ fw)				Blank
Tissue	muscle	ovaries	muscle	2	ovaries	5	muscle	;	ovarie	S	muscle	e	ovaries	5	
	mean sem	mean sem	mean	sem	mean	sem	mean	sem	mean	sem	mean	sem	mean	sem	
TEQs			0.16	0.1	0.53	0.94					0.47	0.14*	0.07	0.04*	
BEQs			0.15	0.07	0.57	0.75					0.20	0.14*	5.14	3.60*	

НСВ	6.22	0.85#	7.37	1.98#	0.09	0.02	0.16	0.04	3.82	0.55#	2.84	0.37#	0.07	0.01*	0.18	0.04*	0.06
ү-НСН	0.58	0.21*	1.18	0.49*	0.01	< 0.01 ^{#,*}	0.03	0.01*	0.59	0.20	1.25	0.31	0.01	< 0.01 ^{#,*}	0.07	0.02^{*}	0.07
<i>p,p'</i> -DDE	13.11	5.99	8.76	2.46#	0.14	0.04 ^{#,*}	0.19	0.03*	3.19	0.58	2.36	0.38#	0.06	0.01 ^{#,*}	0.16	0.04*	0.52
<i>o,p'-</i> DDT	3.20	0.98	2.83	0.65#	0.04	0.01*	0.07	0.01*	1.65	0.36	1.64	0.38#	0.03	< 0.01*	0.09	0.02^{*}	0.07
<i>p,p'</i> -DDT	5.31	1.61#	6.21	1.10#	0.06	0.02*	0.15	0.02*	2.19	0.42#	2.98	0.52#	0.04	0.01*	0.19	0.04*	0.32
Σ DDTs	19.45	8.14 [#]	17.80	3.91 [#]	0.22	0.07	0.40	0.05	7.04	1.00 [#]	6.98	1.14 [#]	0.12	0.02*	0.44	0.10*	
28	0.95	0.06*	1.50	0.26*	0.01	< 0.01*	0.04	0.01 ^{#,*}	1.23	0.11*	2.84	0.53*	0.04	0.01*	0.18	0.05#,*	0.12
52	2.15	0.79*	3.29	0.98*	0.02	0.01*	0.09	0.03*	1.56	0.38	7.72	2.31	0.36	0.34	0.37	0.09	0.25
101	5.12	1.48*	7.10	1.69*	0.06	0.01*	0.19	0.05 ^{#,*}	3.99	0.97	10.74	2.72	0.68	0.63	0.60	0.12#	0.26
138	5.74	1.21*	8.20	1.83*	0.07	0.01*	0.22	0.05*	5.47	1.23	11.06	2.69	0.60	0.52	0.68	0.18	0.40
153	6.00	1.26*	8.72	2.10*	0.07	0.01*	0.23	0.06*	5.94	1.40	11.96	2.52	0.41	0.33	0.76	0.20	0.49
180	1.94	0.31*	3.06	0.67*	0.02	< 0.01*	0.08	0.02*	1.86	0.35*	3.51	0.50*	0.07	0.04	0.25	0.08	0.17
Σ PCBs	21.91	4.98*	31.87	7.26*	0.26	0.03*	0.85	0.21#,*	20.04	4.27	47.82	10.24	2.17	1.86	2.85	0.63#	
28	0.21	0.09	0.83	0.37	< 0.01	< 0.01*	0.02	0.01*	0.22	0.08^{*}	0.87	0.33*	< 0.01	< 0.01*	0.05	0.02^{*}	< 0.01

%lipid					1.45	0.18	2.64	0.23 [#]					2.13	0.35*	6.93	1.32 ^{#,*}	
Σ PBDEs	5.01	0.82*	10.03	1.86*	0.07	0.01*	0.29	0.06*	4.08	0.54*	0.72	0.14*	0.08	0.01*	0.05	0.02*	
197	0.13	0.03#,*	0.27	0.08 ^{#,*}	< 0.01	< 0.01*	0.01	<0.01 ^{#,*}	0.07	0.01#,*	0.33	0.10 ^{#,*}	< 0.01	< 0.01*	0.02	0.01#,*	0.01
183	0.08	0.01#	0.16	0.07	< 0.01	< 0.01 ^{#,*}	< 0.01	< 0.01 ^{#,*}	0.04	0.01#	0.53	0.09	< 0.01	< 0.01 ^{#,*}	0.04	0.02 ^{#,*}	0.01
153	0.11	0.02	0.21	0.07	< 0.01	< 0.01*	0.01	< 0.01*	0.07	0.01*	0.53	0.09*	< 0.01	< 0.01*	0.04	0.02*	0.01.
99	0.99	0.47*	1.97	0.49 ^{#,*}	0.01	0.02*	0.06	0.02*	0.86	0.12*	5.28	1.01#,*	0.02	< 0.01*	0.39	0.19*	0.05
100	0.59	0.12*	1.27	0.27^{*}	0.01	< 0.01*	0.04	0.01*	0.45	0.10*	2.17	0.42*	0.01	< 0.01*	0.15	0.05*	0.11
47	2.90	0.50*	5.32	1.10*	0.04	< 0.01*	0.15	0.03*	2.46	0.39*	10.14	1.41*	0.04	0.01*	0.70	0.26*	0.33

534 In **bold**: sum (Σ) of all DDT, PCB & PBDE congeners. Lw: lipid weight, fw: fresh weight. TEQ: toxic equivalents, pg WHO-TEQ ⁻¹g fw,

calculated by using the toxic equivalency factors recommended by Van den Berg (1998). BEQ: bioequivalent values, BEQ g^{-1} fw. The # denotes a

significant difference between *C. gunnari* and *C. aceratus* in the given tissue at $p \le 0.05$. The * denotes a significant difference between tissues for

537 each species at $p \le 0.05$.

538 Figure Captions

- 540 Figure 1 Mean (\pm sem) concentration of Σ PCBs, Σ BDEs, HCBs, γ -HCHs and Σ DDTs (ng
- g^{-1} lipid weight), in muscle and ovaries of the two icefish species, *C. gunnari* (*n*=11) and *C*.
- *aceratus* (*n*=10). The # denotes a significant difference between C. gunnari and C. aceratus in
- 543 the given tissue at $p \le 0.05$. The * denotes a significant difference between tissues for each
- 544 species at $p \le 0.05$.
- **Figure 2:** Congener composition (percentage) of PCBs, DDTs and BDEs in muscle (m) and
- 547 ovaries (ov) of *C. gunnari* (*Cg*) and *C. aceratus* (*Ca*).

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Highlights

- PCB and DDT concentrations in icefish are higher than those measured in the late 90s
- Mature, female icefish possess higher contaminant levels in ovaries than in muscle
- POP levels are similar in fish from different sampling sites around the Antarctic Peninsula
- Piscivorous icefish display a higher toxicant accumulation than planktivorous icefish
- Bioanalytical results (BEQs) were similar to chemical analytical calculations (TEQs) in icefish

1	Persistent organic pollutants in tissues of the white-blooded Antarctic fish
2	Champsocephalus gunnari and Chaenocephalus aceratus
3	
4	
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- 25 Abstract
- 26

27 The global occurrence of persistent organic pollutants (POPs) continuously contributes to 28 their accumulation also in remote areas such as the Antarctic Ocean. Antarctic fish, which 29 hold high trophic positions but appear to possess low endogenous elimination rates for 30 chemicals, are expected to bioaccumulate POPs with rising anthropogenic pollution. 31 Using a chemical-analytical method, we measured concentrations of PCBs, PBDEs, HCBs, 32 HCH and DDTs and determined toxic equivalents (TEQs) and bioanalytical equivalents 33 (BEQs) in muscle and ovaries of Antarctic icefish caught in the Southern Ocean around 34 Elephant Island. We used two species with different feeding habits and trophic web positions: the planktivorous Champsocephalus gunnari and the piscivorous Chaenocephalus aceratus. 35 Our results revealed higher contaminant levels in ovary than in muscle tissues of both 36 37 species. Most analytes concentrations and the TEQs (0.2-0.5) and BEQs (0.2) were lower as in temperate species. Comparison with literature data points to higher PCB (20-22 ng g⁻¹ lipid 38 weight (lw)) and DDT (7-19.5 ng g^{-1} lw) concentrations than those measured in icefish in the 39 90's. For the other contaminants, we could not identify temporal trends. We found a higher 40 bioaccumulation of contaminants, particularly HCB and DDTs, in C. aceratus (6.2 & 19.5 ng 41 g⁻¹ lw, respectively) than in C. gunnari (3.8 & 7.0 ng g⁻¹ lw, respectively). However, there 42 43 was no general species-specific accumulation pattern of the different toxicant classes between 44 the two icefish. Thus, the expected link between contaminant burdens of C. aceratus and C. 45 gunnari and their ecological traits was only weakly supported for these species. 46 47 Highlights 48 49 PCB and DDT concentrations in icefish are higher than those measured in the late 90s _ 50 _ Mature, female icefish possess higher contaminant levels in ovaries than in muscle 51 POP levels are similar in fish from different sampling sites around the Antarctic _ 52 Peninsula

- 53 Piscivorous icefish display a higher toxicant accumulation than planktivorous icefish
- Bioanalytical results (BEQs) were similar to chemical analytical calculations (TEQs)
 in icefish
- 56
- 57 Keywords

58 Icefish, bioaccumulation, persistent organic pollutants, polychlorinated biphenyls (PCBs),

59 toxic equivalents (TEQs), DR CALUX bioanalytical equivalents (BEQs)

- 60
- 61 62

1. Introduction

63

64 Antarctica has been less affected by human influences than other continents for a long time, 65 however, contamination with anthropogenic contaminants, in particularly persistent organic 66 pollutants (POPs) has increased progressively (Nash, 2011; UNEP/AMAP, 2011). Nowadays, 67 Antarctica serves as a major sink for highly persistent contaminants. Long-range atmospheric 68 transport, together with global distillation processes and cold condensation, are considered to 69 be the main mechanisms for the progressive contamination of the Antarctic ecosystem, 70 together with local sources such as fishing, tourism and research activities (Simonich and 71 Hites; Feely et al., 2008; Nash, 2011). Particularly POPs such as polychlorinated biphenyls 72 (PCBs) and polybrominated diphenyl ethers (PBDEs), and amongst them also the formerly 73 used insecticides such as γ -hexachlorcyclohexane (γ -HCH) and p, p'-DDT are ubiquitous 74 pollutants with a wide application spectrum. For example, polymer additives such as flame 75 retardants are found globally in building material, furniture, paint, textiles or plastics, which 76 are also used in Antarctic bases and vessels cruising in these regions (Hale et al., 2008; 77 Kohler et al., 2008). This worldwide abundance and the high persistence of POPs 78 continuously contribute to an accumulation in the ice masses and biota of polar regions 79 (Wania and Mackay, 1993; Chiuchiolo et al., 2004; Goerke et al., 2004; Corsolini et al., 80 2007; Bargagli, 2008; Borghesi et al., 2008; Borghesi et al., 2009; Xie et al., 2011; Wolschke 81 et al., 2015).

82

83 Furthermore, those lipophilic organic chemicals have a high potential to bioaccumulate in 84 aquatic biota, and particularly in Antarctic species, which generally possess low endogenous 85 elimination rates for those chemicals (Strobel et al., 2015). Additionally, it is expected that 86 climate warming will lead to the release of those pollutants trapped in glaciers and sea-ice, 87 which additionally contributes to increasing POP levels in the tissues of Antarctic animals (Weber and Goerke, 2003; Bogdal et al., 2010; Schmid et al., 2010; van den Brink et al., 88 89 2011; Cabrerizo et al., 2013; Goutte et al., 2013). As global chemical usage continues to 90 grow, including the usage of persistent and bioaccumulative compounds, it is to be expected 91 that contaminant intake into Antarctica will further increase in the future.

93 The endemic, Antarctic notothenioid fish are evolutionary well-adapted to the cold and stable

94 environment of the Southern Ocean. Adaptations involve e.g. high amounts of tissue lipids,

95 slow growth rates, long life spans and slow metabolism and elimination rates for xenobiotics

96 (Mintenbeck et al., 2012; Strobel et al., 2015). All these factors may favor the

97 bioaccumulation of lipophilic contaminants in the tissues of those fish.

98

99 Lipophilic, non-charged contaminants like PCBs can be taken up via a physico-chemically 100 driven, passive partitioning of the chemicals from the water phase into the lipid phase of the 101 organism and thereby accumulate differentially in tissues or fish species, depending on their 102 lipophilicity of the chemical and the lipid content or composition of the tissues (Nichols et 103 al., 2013). In fact, tissue-specific differences of POP levels are reported for notothenioid 104 species (Lana et al., 2014). Furthermore, bioaccumulation and biomagnification can be 105 related to an organism's habitat and its trophic position within the food web (Corsolini et al., 106 2002a; Weber and Goerke, 2003; Chiuchiolo et al., 2004). Particularly for highly lipophilic 107 compounds, oral uptake via the prey contributes by a major extent to the bioaccumulation of 108 toxicants in the tissues of a species. For example, Weber and Goerke (2003) found higher 109 contaminant levels in the piscivorous icefish Chaenocephalus aceratus than in the 110 planktivorous icefish *Champsocephalus gunnari*, which was apparently related to the 111 different food spectra of the two notothenioids (Goerke et al., 2004). Such studies highlight 112 that POPs are transferred within the Antarctic food web, leading to increasing POP 113 concentrations along the food chain up to high concentrations in top level predators 114 (Wolschke et al., 2015).

115

Considering the trends of rising POP concentrations in Antarctica (UNEP/AMAP, 2011 and their high potential to exert toxic effects in marine biota (Nash, 2011), it is very important to keep monitoring body burdens of the fish living in the Southern Ocean. Yet, we are far from having a solid understanding of the contamination status, time trends, the diversity of contaminants or their toxicity potential in Antarctic biota (UNEP, 2002; UNEP/AMAP, 2011). Particularly the bioaccumulation pattern of contaminants such as the polybrominated flame retardants (PBDEs) or insecticides (e.g. DDT) have hardly been measured in icefish.

124 The aim of this study was to determine levels of selected POPs in muscle and ovary tissue of 125 two white-blooded Antarctic notothenioid species in the study area around Elephant Island,

- 126 the South Shetland Islands and the Antarctic Peninsula, and to estimate temporal trends of 127 fish POP levels in this area. The analytically measured contaminant concentrations were 128 converted into toxic equivalents (TEQs) using standardized values. In order to examine 129 whether food web position or tissue lipid content has an influence on chemical 130 concentrations, we analyzed lipid content and POP levels in fish species of the same family 131 but with different feeding habits and trophic web positions, namely the planktivorous C. 132 gunnari and the piscivorous C. aceratus. 133 Extending the classical contaminants, this study includes polybrominated flame retardants 134 (PBDEs) and insecticides (i.e. DDT, HCB). In addition to chemical analytics, we also applied 135 bioanalytics using the DR-CALUX assay (Murk et al., 1996; Kuiper et al., 2006) to assess the 136 total accumulated dioxin-like activity in the fish tissues. 137 138 139 2. Methods 140 141 2.1 Study species 142 Two species of the Channichthyidae (white-blooded icefish) were fished with bottom trawls 143 down to 500 m, during a cruise with the research vessel 'Polarstern' (ANTXXVIII/4, March 144 13 to April 9, 2012; http://expedition.awi.de/expedition/ANT-145 XXVIII/4?alias=PS79#mapChart) at different, closely located sampling sites around Elephant Island and the South Shetland Islands (61.1°S, 55.1°W). Only fish netted alive and without 146 147 macroscopically visible damage were used for tissue sampling. The planktivorous mackerel 148 icefish, C. gunnari, shows a mainly bentho-pelagic feeding mode, while the piscivorous 149 Scotia Sea icefish C. aceratus is predominantly a benthos feeder (Weber and Goerke, 2003). Persistent organic pollutants were analyzed in muscle and ovary tissue of mature (stage III-150 151 IV), female fish only. Sex and maturity stage of the fish were verified histologically. Animal 152 weight and length are given in Table 1. All tissue samples were wrapped in aluminum foil 153 and stored immediately at -20 °C until used for analysis. 154 155 2.2 Sample preparation 156
- 157 Muscle and gonad samples were defrosted, cut into pieces and lyophilized at 33 Pa for 72
- 158 hours until constant weight. Dried tissue of muscle (5-10 g) or gonads (0.5-2 g) was ground

with anhydrous sodium sulfate and quartz sand in a ceramic mortar and pistil to obtain a finepowder.

This homogenate was then Soxhlet-extracted with a speed-extractor (E-914, Büchi, 161 Switzerland) in 120 mL extraction cells at a constant temperature and pressure of 100 °C and 162 163 100 bar, respectively (4 cycles, hold time 10 min., discharge 4 min), with ~150 ml n-164 hexane/dichloromethane 1:1 (v:v) (Hartmann, 2013). The extract was concentrated in a 165 Syncore evaporator (Büchi, Switzerland) und let dry completely by applying a gentle nitrogen stream. The residue accounted for the fat content of the sample. ${}^{13}C_{12}$ labeled 166 167 internal standards (Schmid et al., 2007) were added to the samples and after the addition of 2 168 -3 mL of *n*-hexane the solution was treated with 3 ml oleum (7% SO₃ in conc. sulfuric acid). 169 After centrifugation for 3 min at 5'000 rpm, the solvent layer with the lipophilic target 170 analytes was removed and the remaining suspension was re-extracted two times more with n-171 hexane. The pooled extracts were concentrated to 0.5 mL in a rotary evaporator at 45 °C and 172 300 mbar. Subsequently, the extract was purified on a multilayer mini silica gel column (from 173 top to bottom: 0.25 g anhydrous sodium sulfate, 0.25 g silica gel 60 with 44% sulfuric acid 174 and 0.25 g silica gel 60 activated at 130 °C). The sample was applied on the column and 175 eluted with 5 ml *n*-hexane, followed by 5 mL *n*-hexane/dichloromethane 1:1 (v:v). The eluate 176 was concentrated using a rotary evaporator to 0.5 mL. After transfer to a mini GC-Vial the 177 volume was further reduced to 30 μ L by the application of a gentle stream of nitrogen at room temperature. Finally the recovery standard ${}^{13}C_{12}$ labeled PCB 70 was added. Samples 178 179 were stored in toluene at -20 °C until analysis. Method blank levels for the whole analytical 180 procedure were determined in duplicates (Table 1 and A.2).

181

182 2.3 Chemical analysis

183

184 PCBs (indicator PCB 28, 52, 101, 138, 153, and 180; dioxin-like PCBs 77, 81, 105, 114, 118,

185 123, 126 156, 157, 167, 169, and 189), DDT (*o*,*p*'-DDT, *p*,*p*'-DDT, and *p*,*p*'-DDE),

186 hexachlorobenzene (HCB), γ-hexachlorocyclohexane (γ-HCH), and PBDEs (BDE 28, 47, 99,

187 100, 153, 154, 183, and 209) were included in this study. Quantitative determination of the

188 target analytes in the extracts was achieved by gas chromatography/high resolution mass

- 189 spectrometry (GC/HRMS). Analyses were carried out on a Finnigan MAT95 high-resolution
- 190 mass spectrometer (Thermo Finnigan MAT, Bremen Germany) coupled to a Finnigan Trace

191 GC Ultra equipped with a Triplus auto sampler (Thermo Electron Corporation, Waltham, MA, USA). Samples were injected in splitless mode (splitless time 30 s) at an injector 192 193 temperature of 260 °C. For the gas chromatographic separation a RTX5 Sil-MS column (30 194 $m \times 0.25$ mm, film thickness 0.10 µm) was used with helium as carrier gas at a pressure of 195 100 kPa. The following temperature programs were used for the different compound classes. 196 For the PCBs, the initial column temperature was 100 °C. After 0.5 min, the temperature was 197 ramped at 20 °C/ min to 180 °C, followed by 3 °C/ min to 250 °C, and 20 °C/ min to 300 °C. 198 For pesticides, the initial column temperature was 100 °C. After 0.5 min, the temperature was 199 ramped at 10 °C/ min to 160 °C, followed by 4 °C/ min to 240 °C, and 20 °C/ min to 300 °C. 200 For the PBDEs, the initial column temperature was 100 °C. After 0.5 min, the temperature 201 was ramped at 20 °C/ min to 220 °C, followed by 6 °C/ min to 300 °C, and 10 °C/ min to 320 °C. The ion source was operated at 220 °C, the electron energy was 70eV, and the mass 202 203 spectrometer was tuned to a mass resolution of 8000-10000. The two most abundant signals of the molecular ion cluster of the analytes and the ${}^{13}C_{12}$ labeled internal standards were 204

- 205 recorded in the single ion monitoring mode.
- 206

207 The analytes were identified by comparing the retention times with those of the labeled

208 internal standards. Quantification was based on peak areas of the analytes and the labeled

209 reference compounds with known concentrations. More details about the method are

available in the literature (Zennegg et al., 2003; Schmid et al., 2007).

211 For the calculation of TEQs, the dioxin-like PCBs were determined in the same way, but as

212 part of the sample extract was used later on for the analysis by the bioassay (DR-CALUX),

213 no ${}^{13}C_{12}$ -labeled internal dl-PCBs standards could be used for the quantification, as the

214 isotope labeled analogues exhibit similar activity in the DR-CALUX. Therefore, quantitative

215 determination of dl-PCBs was based on the ${}^{13}C_{12}$ labeled indicator PCBs used as internal

216 standards and previously determined response factors to native dl-PCBs.

The blank values (in ng g^{-1} lipid weight) were all below the compound concentrations and are given in Table 1 and A.1.

219 The limit of detection (LOD) and the limit of quantification (LOQ) were set by definition at

signal to noise ratios of greater than three $(s/n \ge 3)$ and ten $(s/n \ge 10)$ respectively. All glass

221 ware used were cleaned with strongly alkaline detergents and backed out overnight in a

222 ceramic oven at 450 °C. Directly before use, the glass ware was rinsed with solvents (n-

223 hexane, dichloromethane).

- 225 2.4 Determination of dioxin-like toxic equivalents (TEQs)
- 226
- In order to assess total dioxin-like activity in the fish tissues, we calculated the 2,3,7,8-
- 228 tetrachlorodibezo-p-dioxin (TCDD) equivalent (TEQ) concentrations of dl-PCBs in the
- 229 muscle and gonad extracts of both icefish species using Toxic Equivalency Factors (TEFs)
- proposed by the World Health Organization (WHO) for fish (Van den Berg et al., 1998).
- TEQs were calculated as the sum of the TEFs of all dl-PCBs listed in Table A.1. Although
- the TEF values are not derived from studies with Antarctic fish, since TEFs are only available
- for salmonids (Van den Berg et al., 1998) they provide a useful tool for a reasonable estimateof toxicity effects of PCBs on Antarctic fish.
- 235
- 236 2.5 Determination of bioequivalent values (BEQs)
- 237
- The same extracts that were used for chemical analysis were also used in a standard bioassay,
- the DR-CALUX (Dioxin Responsive Chemically Activated Luciferase Gene Expression)
- assay. This cell and receptor based reporter gene bioassay measures the binding of dioxin-like
- 241 HAHs, e.g. dioxin-like PCBs (dl-PCBs), to the Aryl hydrocarbon receptor (AhR) via
- activation of a reporter gene. The assay was performed by BioDetection Systems b.v.
- 243 (Amsterdam, The Netherlands).
- 244

Bioassay-derived TEQ values, or bioequivalent values (BEQs), were compared with TEQ
values calculated from the chemical analytical data. This comparison revealed if there was a
significant higher activity than predicted from the chemical analysis. Such a combined
approach has been repeatedly used in studies on chemical contamination of the Arctic region
(Letcher et al., 2010), but it has not been used in studies with Antarctic fish so far.

- 250
- 251 2.5 Calculations and statistics
- 252
- 253 All analyses include the lipid content of the respective tissue. POP concentrations were
- normalized against both lipid weight (lw, ng g^{-1}) and fresh weight (fw, ng g^{-1}) of the tissue
- and provided as mean values \pm standard error of the mean (sem). Toxicant concentrations
- 256 were compared among icefish species and tissues using ANOVA with Tukey post-hoc test.

257	Influence of sampling site, fish weight and length, maturity stage and tissue lipid content was
258	tested with ANOVA. Data were considered to be statistically significant at $p < 0.05$. Normal
259	distribution of data was tested with Kolmogorov-Smirnov and equality of variances with
260	Bartlett's test. All statistical tests were performed with STATISTICA 12, StatSoft, Inc., and
261	GraphPad Prism 5, GraphPad Software, Inc.
262	
262	
263	
264	3. Results
265	
266	3.1 Interspecies comparison of contaminant patterns and levels
267	
268	In this study, we measured lipid content, PCB, DDT, HCB, γ -HCH, and PBDEconcentrations
269	in both muscle and ovary tissue of female, mature <i>C. gunnari</i> and <i>C. aceratus</i> from spatially
270	closely located sampling sites around Elephant Island and the South Shetland Islands. We
271	express the compound concentrations on lipid weight and the fresh weight basis.
272	Sampling site, maturity stage and tissue lipid content of the fish showed no correlation to the
273	accumulation of any of the compounds analysed in this study.
274	All samples contained detectable levels of the target compounds, which are presented in
275	Table 1. Analytes concentrations were about 15 - 110 times higher when calculated per lipid
276	weight than based on tissue fresh weight.
277	Overall, the PCBs were the predominant group among all compounds analysed in our study
278	(mean values \pm sem (ng g ⁻¹ lw): <i>C. gunnari</i> muscle 20.0 \pm 4.3, <i>C. gunnari</i> ovaries 47.8 \pm
279	10.2, <i>C. aceratus</i> muscle 21.9 ± 4.9 , <i>C. aceratus</i> ovaries 31.9 ± 7.3), followed by the DDTs
280	(mean values \pm sem (ng g ⁻¹ lw): <i>C. gunnari</i> muscle 7.0 \pm 1.0, <i>C. gunnari</i> ovaries 6.9 \pm 1.1, <i>C.</i>
281	<i>aceratus</i> muscle 19.4 \pm 8.1, <i>C. aceratus</i> ovaries 17.8 \pm 3.9), HCB (mean values \pm sem (ng g ⁻¹
282	lw): C. gunnari muscle 3.8 ± 0.5 , C. gunnari ovaries 2.8 ± 0.4 , C. aceratus muscle 6.2 ± 0.9 ,
283	<i>C. aceratus</i> ovaries 7.3 ± 2.0), the PBDEs (mean values \pm sem (ng g ⁻¹ lw): <i>C. gunnari</i> muscle
284	4.1 ± 0.5 , C. gunnari ovaries 0.7 ± 0.1 , C. aceratus muscle 5.1 ± 0.8 , C. aceratus ovaries 10.0
285	\pm 1.9) and γ -HCH (mean values \pm sem (ng g ⁻¹ lw): <i>C. gunnari</i> muscle 0.6 \pm 0.2, <i>C. gunnari</i>
286	ovaries 1.3 ± 0.3 , <i>C. aceratus</i> muscle 0.6 ± 0.2 , <i>C. aceratus</i> ovaries 1.2 ± 0.5) (Figure 1).
287	Within the PCBs, only PCB 28 and 101 were significantly different between the ovaries of <i>C</i> .
288	gunnari and C. aceratus (Table 1). When we calculated the percentage contribution of the
289	individual PCB congeners to \sum PCBs (lw) of both muscle and ovary tissue, there was no

290 difference in PCB congener composition between the two icefish species (two-way

- 291 ANOVA).
- Among the PCBs, PCB 153 contributed by 28% in muscle and by 26% in ovaries to all PCBs
- in both species. The second-most abundant PCBs were PCB 138 (~26%) and 101 (~21%)
- 294 (Figure 2).
- 295
- In the ovaries, levels of *p*,*p*'-DDE, *o*,*p*'-DDT and *p*,*p*'-DDT (per lw) were significantly
- 297 higher in *C. aceratus* compared to those of *C. gunnari*. Also *p,p*'-DDT concentrations in the
- 298 muscle of *C. aceratus* were higher than those in muscle of *C. gunnari* (per lw). Σ DDT (per
- lw) was significantly higher in muscle and ovaries of *C. aceratus* than in *C. gunnari* (Figure
- 300 1). Among the DDTs, *p*,*p*'-DDE showed the highest concentration of up to 57% in muscle of
- 301 *C. aceratus. p,p* '-DDE tended to be higher in muscle tissue, while *p,p* '-DDT was slightly
- 302 higher in ovary tissue in both species. *o*,*p*'-DDT was slightly higher in *C*. *gunnari* with
- 303 ~22%, while it was only ~16% in *C. aceratus*. Detailed information on the contribution of all
- individual compounds measured in this study is given in Table A.2 in the appendix.
- 305γ -HCH was only different between the muscle tissues of the two icefish (on fw basis).
- 306 HCB concentrations per lipid weight in both muscle and ovaries were higher in *C. aceratus*
- 307 than in *C. gunnari* (Figure 1). Except HCB, none of the analysed compounds exhibited a
- 308 correlation with fish weight or length. HCB showed a significant linear dependence of fish 309 weight and length in *C. aceratus*, except in the ovaries ($R^2=0.723$).
- 310
- In both species, the PBDEs were dominated by BDE 47 (~59% in muscle, ~54% in ovaries),
- followed by ~20% BDE 99. On both the lipid and the fresh weight basis, BDE 183 and 197
- 313 were significantly different between the two tissues types and fish species. Also BDE 99
- 314 showed different concentrations between the ovaries of *C. aceratus* and *C. gunnari* on the
- 315 lipid basis (Table 1, Table A.2, Figure 2).
- 316

317 3.2 Comparison of contaminant patterns and levels in different tissue

- 318
- 319 Lipid content was significantly higher in the ovaries of *C. gunnari* than in the ovaries of *C*.
- 320 *aceratus*. *C. gunnari* exhibited significantly higher lipid content in ovaries than in their
- 321 muscle tissue, while this was not the case for *C. aceratus* (Table 1).
- 322 A comparison between the compound concentrations in muscle and ovary tissue for each
- 323 icefish species revealed higher HCB concentrations in ovary than in muscle tissue of *C*.

324	gunnari. Concentrations of γ -HCH and DDTs in muscle and ovary tissue (per fresh weight)
325	were significantly different in both species (Table 1).
326	In C. aceratus, all PCB concentrations and most of the PBDEs were different from each other
327	in both muscle and ovary tissue. In case of dl-PCB congeners, the ∑dl-PCBs were higher in
328	muscle of C. gunnari than in muscle C. aceratus on both lipid and fresh weight basis (Table
329	A.1).
330	In C. gunnari, all PBDEs apart from congener 183 did differ between muscle and ovary
331	tissue. In contrast, only PCB 28 and 180 were different between the tissues of C. gunnari, all
332	other PCB congeners showed no significant tissue differences (Table 1, Figure 1).
333	
334	3.3 Toxic equivalents (TEQs)
335	
336	The toxic equivalent levels of dioxin-like PCBs (WHO _{PCB} -TEQ g^{-1} fw) were higher in the
337	muscle of C. gunnari (0.5 ± 0.1) than in C. aceratus (0.2 ± 0.1) , while C. aceratus exhibited
338	higher TEQs (0.5 ± 0.9) in the ovaries than <i>C. gunnari</i> (0.1 ± 0.0) (Table 1).
339	
340	3.4 Bioanalytical equivalents (BEQs)
341	
342	The bioanalytical equivalents (BEQs) determined by DR-CALUX in the muscle of both
343	species were in a similar range (<i>C. aceratus</i> : 0.15 ± 0.07 , <i>C. gunnari</i> : 0.2 ± 0.14 BEQ g ⁻¹ fw)
344	to the WHO _{PCB} -TEQs g^{-1} lw, which were calculated using the concentrations of dioxin-like
345	PCBs. In C. gunnari, however, the BEQs measured in the ovaries were about 70-fold higher
346	(BEQ: 5.14 ± 3.6 , TEQ: 0.07 ± 0.04) than the calculated WHO _{PCB} -TEQ g ⁻¹ fw (Table 1).
347	
348	
349	4. Discussion
350	
351	4.1 Species-specific contaminant patterns and levels
352	4.1.1. Species-specific patterns
353	
354	Literature data on present POP concentrations in tissues of Antarctic fish are scarce, and
355	particularly those on contaminant levels in white-blooded icefish.
356	Earlier studies of the 80's and 90's measured PCB concentrations in the icefish C. gunnari
357	and C. aceratus around the Antarctic Peninsula (Corsolini et al. 2002a, 2002b, 2009), which

- 358 were two to four times lower (0.7 ng g^{-1} fw in muscle) than the ones of the present study.
- 359 HCB levels of our icefish were within a similar range to those measured previously. It is
- 360 known for sub-Antarctic fish that PCB levels can follow a seasonal trend, which is related to
- the release of pollutants from melting snow and ice during summer and the atmospheric
- 362 transport of loads of pollutants which precipitate in Antarctic regions and are released during
- 363 warming (Jaffal et al., 2011). Since sampling of our icefish took place in Austral autumn,
- after the seasonal ice melting in Antarctica, a seasonal release of POPs trapped in sea ice
- 365 could thereby be one contributor to the comparably high levels of PCBs in our icefish
- 366 samples. In fact, the PCB concentrations (per fw) of our icefish were within a similar range to
- those of salmon from the Baltic Sea (Isosaari et al., 2006).
- In the early 00's, Borghesi et al. (2008; 2009) sampled *C. hamatus* and *C. gunnari* in the
- Ross Sea at 74° South. They found \sum PCB concentrations in *C. hamatus* muscle of 0.35 ng g⁻¹
- 370 fw, \sum non-ortho PCBs of 5 ng g⁻¹ fw and \sum PBDEs 0.16 ng g⁻¹ fw. In muscle of *C. gunnari*,
- 371 they report \sum PBDEs of 0.44 ng g⁻¹ fw. Those values are about five times higher than the
- size class, this difference is likely related to the measurement of different PBDE congeners inthe studies.
- 375 Another recent study measured contaminant concentrations (in ng g^{-1} lw) in muscle, liver and
- 376 gonads of three red-blooded Antarctic species, Notothenia coriiceps, N. rossii and
- 377 *Trematomus newnesi*, from Potter Cove, Antarctic Peninsula (Lana et al., 2014). They
- 378 reported similar \sum DDT values in muscle, but also species differences in the ovaries of their
- 379 fish: while *N. coriiceps* and *T. newnesi* had similar values to *C. aceratus* measured in our
- 380 study, *N. rossii* had much higher values than our fish. They also measured γ -HCH values
- about six-times higher than in our fish. N. rossii displays a rather benthic lifestyle, but also
- feeds on pelagic species, and thus has a prey spectrum similar to C. gunnari. Nevertheless,
- 383 the DDTs and γ -HCH were much higher in *N. rossii* than in *C. gunnari*. In contrast, \sum PCB
- and \sum BDEs were highly variable among the species and tissues measured by Lana et al.
- 385 (2014), but were generally within the same order of magnitude compared to our data. Thus,
- 386 such strong differences in DDT and γ -HCH accumulation patterns between Antarctic fish
- 387 species could also be related to a selective metabolism for individual contaminant classes
- between the species (Storelli et al., 2009), and not only to their ecological traits.
- 389
- Amongst the PBDEs, BDE 47 showed the highest concentration (60% of all congeners) in
 our icefish amongst all PBDE congeners, followed by BDE 100 and BDE 99. This is in line

- 392 with the general picture of those congeners being the dominating PBDEs in fish around the
- globe, i.e. BDE 47 being recognized as the most important PBDE congener in marine biota
 (Zennegg et al., 2003; Isosaari et al., 2006; Kuiper et al., 2006).
- 395 In comparison to fish from non-Antarctic regions, PBDE concentrations on the fresh weight
- 396 basis were particularly low in icefish. For example, PBDE concentrations range from about
- 1.0 to 8 ng g⁻¹ fresh weight, or up to 64 ng g⁻¹ lipid weight in various fish species from the
- Baltic Sea (Isosaari et al., 2006), which are at least ten times higher than in the icefish.
- 399 Although the production of the former widely used brominated flame retardants penta- and
- 400 octabromodiphenyl ether (PentaBDE and OctaBDE) were banned by the European Union in
- 401 2004 and several states of the USA, toxic and persistent lower brominated PBDEs are still
- 402 produced in other areas of the world and redistributed globally, also to the Antarctic (Cox and
- 403 Efthymiou, 2003; Renner, 2004; Vives et al., 2004; Kuiper et al., 2006).
- 404 HCB concentrations (per fw) were also up to 30 times lower, and DDT concentrations (per
- 405 fw) several hundreds of times lower in our icefish than in fish from the Northern hemisphere
- 406 (Sharma et al., 2009). Despite a worldwide stop of the production of DDTs during the 70's, it
- 407 has been reintroduced in the 2000s as malaria control by the WHO, and about 6000 tons of
- 408 DDTs are still produced per year (UNEP, 2008). Due to its high persistence, bioaccumulation
- 409 potential and cold condensation processes, DDTs and its metabolites are nowadays found in
- 410 biota all over the world, and particularly in polar regions (Mirmigkou and de Boer, 2015).
- 411
- 412 The TEQs we calculated for *C. aceratus* and *C. gunnari* were in a similar range than TEQs
- 413 reported for muscle of the icefish C. hamatus (TEQ 0.01-0.1 pg g^{-1} wet weight) or red-
- 414 blooded Antarctic fish (TEQ ~ 0.1 pg g^{-1} wet weight) (Focardi et al.; Corsolini et al., 2002a;
- 415 Borghesi et al., 2008). Generally, the values for Antarctic fish were lower than those for other
- 416 organisms living in less remote parts of the world, e.g. WHO_{PCB} -TEQ (pg g⁻¹ fw) of 3 to 15
- 417 in muscle of salmon or Baltic herring. Yet, burbot from Bothnian bay (Baltic Sea) exhibit
- 418 muscle TEQs, which are in a similar range as in Antarctic fish (Isosaari et al., 2006).
- 419 In C. aceratus, we found that the BEQs were comparable to the TEQs calculated on the basis
- 420 on the WHO_{PCB}-TEFs for salmonid fish species (Van den Berg et al., 1998). In comparison to
- 421 fish from the northern hemisphere (Husain et al., 2014), Antarctic fish had about twenty
- 422 times lower BEQ values in their muscle. In contrast, the BEQs in the ovaries of *C. gunnari*
- 423 were much higher than the WHO_{PCB}-TEQs calculated for their ovaries. The toxicity effects
- 424 might thus be actually much higher than expected by the single usage of the calculated
- 425 WHO_{PCB}-TEQ. In fact, the BEQs in the ovaries of *C. gunnari* were within the same range as

- 426 BEQs of fish from temperate latitudes. Also Corsolini et al. (Corsolini et al., 2002a) stated
- 427 that the TEQ values they measured in Antarctic fish were already half as high as those values
- 428 which are considered to elicit toxicological effects, such as reproductive and immunological
- 429 disorders, in marine mammals or birds (Kannan et al., 2000).
- 430 Rising PCB concentrations as observed by us in tissues of Antarctic fish will thus
- 431 increasingly have the potential to exert their toxic effects on those fish.
- 432
- 433 4.1.2. Temporal trends in contaminant levels
- 434

435 Long-term observations on contaminant levels in Antarctic biota are scarce. Yet, Weber and

436 Goerke (2003; Goerke et al., 2004) measured contaminant levels in the liver of *C. aceratus*

- 437 and *C. gunnari* in the same sampling area as in the present study. From 1987 to 1996, the
- 438 authors measured an increase of PCB 153 and PCB 180 levels in *C. aceratus*, but not in *C.*
- 439 gunnari. In the present study, we found about three times higher concentrations of those PCB
- 440 congeners in the tissues of our two icefish species than in the previous study from 1996
- 441 (Weber and Goerke, 2003). In contrast, the HCB concentrations show a declining trend from
- the 1987 study to our current survey. A similar trend of stable or declining HCB levels by up
- to 2.5% per year has also been observed in Arctic biota, such as birds, fish or marine
- 444 mammals, since the late 80's (Barber et al., 2005; Rigét et al., 2010).
- 445

In liver of *C. aceratus*, concentrations of p,p'-DDE had already increased from 1987 to 1996, and the values we measured in muscle and ovaries of *C. aceratus* were almost twice as high as in 1996 (Weber and Goerke, 2003). In contrast, p,p'-DDE concentrations in *C. gunnari* remained at similar levels from 1987 over 1996 (study by Weber and Goerke (2003)) to the present study.

Nevertheless, overall DDT concentrations were increasing from 1987 to 1996 in both icefish species, and our values are also slightly higher than those in 1996, suggesting an increasing trend of DDTs in icefish around the Antarctic Peninsula. Despite a general global reduction of DTT as an insecticide, DDT is still produced at high volumes (see above, (UNEP, 2008)). Furthermore, climate warming on the one hand leads to an increased volatility and worldwide distribution of DDTs, and on the other hand local sources such as melting glaciers may

- additionally contribute to increasing DDT concentrations in tissues of Antarctic fish around
- 458 the Antarctic Peninsula (van den Brink et al., 2009; van den Brink et al., 2011).
- 459

- 460 4.2 Correlation of contaminant concentrations with ecological traits
- 461

462 4.2.1 Tissue-specific patterns

463

464 Since all POPs analyzed in this study are highly lipophilic substances, tissue differences in 465 the contaminant levels in C. aceratus and C. gunnari on fresh weight basis at first instance 466 should correlate to tissue lipid concentrations (Corsolini et al., 2002a; Weber and Goerke, 467 2003; Chiuchiolo et al., 2004). Indeed, C. gunnari, which possesses a clearly higher lipid 468 content in the ovaries than C. aceratus, accordingly showed significantly higher PCB 469 concentrations in its ovaries based on fresh weight compared to C. gunnari. Also HCB was 470 two times more concentrated in the fat-rich ovaries of C. aceratus than in C. gunnari, but 471 only on the lipid basis. Accordingly, the BEQs were higher in the ovaries of C. aceratus and 472 C. gunnari than in their muscle tissue. Also Lana et al. (2014), investigating POP 473 accumulation patterns in notothenioid species, reported that the highest levels were found in 474 the gonads of the fish.

475

476 4.2.2 Ecological-related patterns

477

In addition to body lipid contents, our results point to an influence of habitat and trophic levelon POP levels in white-blooded icefish. The benthic-living *C. aceratus* had two-times higher

480 concentrations of almost all DDT congeners and HCB than the bentho-pelagic *C. gunnari*.

481 Also the previous study by Weber and Goerke (2003) report higher (lipid-based) contaminant

482 burdens in *C. aceratus* than in *C. gunnari* and highlight the higher tendency of *C. aceratus* to

483 accumulate DDTs in its tissues than *C. gunnari* over the time.

484 Since POPs accumulate in sediments, benthic fish species are generally thought have a higher

485 exposure and uptake of lipophilic contaminants (Goerke et al., 2004; Borghesi et al., 2008).

486 In addition to uptake from water or sediment, also the feeding habit thus plays a role, with

487 species at higher tropic levels tending to show higher contaminant accumulation due to

488 biomagnification. A recent study by Wolschke et al. (2015) also highlights the

489 biomagnification of POPs from lower to higher trophic levels in the Antarctic food chain,

490 which can be attributed to the diets of the animals.

491 From these observations, the benthic, piscivorous *C. aceratus* was expected to have higher

492 contaminant burdens than the bentho-pelagic, planktivorous C. gunnari. However, only HCB

- 493 and DDTs, but none of the other congener classes, were higher in the predominantly benthic
- 494 *C. aceratus* than in the bentho-pelagic *C. gunnari* on the lipid weight basis.
- 495
- 496

497 Conclusion

- 498
- 499

500 Overall, PCB and DDT concentrations tend to rather increase than decrease in tissues of the 501 two white-blooded icefish species C. aceratus and C. gunnari around Elephant Island and the 502 South Shetland Islands, when compared to earlier studies. Our results thereby support the 503 global transportation of POPs to the Southern Ocean and their bioaccumulation in the local 504 marine fish, and point to a trend of increasing concentrations of POPs in Antarctic icefish. 505 Our data also suggest that worldwide climate change effects may contribute to an increased 506 volatilization and release of POPs trapped in glaciers, sea- or pack-ice, thereby leading to an 507 ongoing contamination of the Southern Ocean and its biota.

- 508 Furthermore, we found differences in POP accumulation patterns between the two icefish
- 509 species, which were weakly correlated to their trophic position. The piscivorous *C. aceratus*
- 510 showed a higher potential to accumulate contaminants in its tissue than the planktivorous C.
- 511 *gunnari*. This species difference highlights the influence of intake of POPs via the specific
- 512 prey of individual fish species. However, the expected link between the contaminant burdens
- 513 of *C. aceratus* and *C. gunnari* and their ecological traits could not be fully supported.
- 514 Additional factors, such species differences in toxicant metabolism rates and selective
- 515 metabolism for single contaminant classes, may also play an important role in defining
- 516 chemical bioaccumulation patterns in Antarctic fish species in the long term. In the end,
- 517 Antarctic fish are a central link between the benthic community and top level predators,
- 518 concomitantly POPs bioaccumulated in their tissues are likely to contribute to a progressive
- 519 biomagnification of POPs along Antarctic food webs.
- 520 521

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526

- 527 Tables & Figures
- 528
- 529 Tables
- 530
- Table 1: Lipid content (%) and levels of organic contaminants (ng g⁻¹ lipid weight; ng g⁻¹ fresh weight) (mean \pm sem) in tissues of two Antarctic icefish species.
- 533

	C. aceratus (n=	=10)					C. gunnari (n=11)											
Length (cm)	47-66							33-50										
Weight (g)	636-3620		252-888															
	lipid weight (ng	g g ⁻¹ lw)	fresh weight (ng g- ¹ fw)				lipid w	eight (n	g g ⁻¹ lw)		fresh weight (ng g ⁻¹ fw)				Blank			
Tissue	muscle ovaries		muscle		ovaries		muscle		ovaries		muscle		ovaries					
	mean sem	mean sem	mean	sem	mean	sem	mean	sem	mean	sem	mean	sem	mean	sem				
TEQs			0.16	0.1	0.53	0.94					0.47	0.14*	0.07	0.04*				
BEQs			0.15	0.07	0.57	0.75					0.20	0.14*	5.14	3.60*				

НСВ	6.22	0.85#	7.37	1.98#	0.09	0.02	0.16	0.04	3.82	0.55#	2.84	0.37#	0.07	0.01*	0.18	0.04*	0.06
ү-НСН	0.58	0.21*	1.18	0.49*	0.01	< 0.01 ^{#,*}	0.03	0.01*	0.59	0.20	1.25	0.31	0.01	< 0.01#,*	0.07	0.02^{*}	0.07
<i>p,p'</i> -DDE	13.11	5.99	8.76	2.46#	0.14	0.04 ^{#,*}	0.19	0.03*	3.19	0.58	2.36	0.38#	0.06	0.01 ^{#,*}	0.16	0.04*	0.52
<i>o,p'-</i> DDT	3.20	0.98	2.83	0.65#	0.04	0.01*	0.07	0.01*	1.65	0.36	1.64	0.38#	0.03	< 0.01*	0.09	0.02^{*}	0.07
<i>p,p'</i> -DDT	5.31	1.61#	6.21	1.10 [#]	0.06	0.02*	0.15	0.02*	2.19	0.42#	2.98	0.52#	0.04	0.01*	0.19	0.04*	0.32
Σ DDTs	19.45	8.14 [#]	17.80	3.91 [#]	0.22	0.07	0.40	0.05	7.04	1.00 [#]	6.98	1.14#	0.12	0.02*	0.44	0.10*	
28	0.95	0.06*	1.50	0.26*	0.01	< 0.01*	0.04	0.01 ^{#,*}	1.23	0.11*	2.84	0.53*	0.04	0.01*	0.18	0.05 ^{#,*}	0.12
52	2.15	0.79*	3.29	0.98*	0.02	0.01*	0.09	0.03*	1.56	0.38	7.72	2.31	0.36	0.34	0.37	0.09	0.25
101	5.12	1.48*	7.10	1.69*	0.06	0.01*	0.19	0.05 ^{#,*}	3.99	0.97	10.74	2.72	0.68	0.63	0.60	0.12#	0.26
138	5.74	1.21*	8.20	1.83*	0.07	0.01*	0.22	0.05*	5.47	1.23	11.06	2.69	0.60	0.52	0.68	0.18	0.40
153	6.00	1.26*	8.72	2.10*	0.07	0.01*	0.23	0.06*	5.94	1.40	11.96	2.52	0.41	0.33	0.76	0.20	0.49
180	1.94	0.31*	3.06	0.67^{*}	0.02	< 0.01*	0.08	0.02*	1.86	0.35*	3.51	0.50^{*}	0.07	0.04	0.25	0.08	0.17
Σ PCBs	21.91	4.98*	31.87	7.26*	0.26	0.03*	0.85	0.21 ^{#,*}	20.04	4.27	47.82	10.24	2.17	1.86	2.85	0.63#	
28	0.21	0.09	0.83	0.37	< 0.01	< 0.01*	0.02	0.01*	0.22	0.08^{*}	0.87	0.33*	< 0.01	< 0.01*	0.05	0.02^{*}	< 0.01

%lipid					1.45	0.18	2.64	0.23 [#]					2.13	0.35*	6.93	1.32 ^{#,*}	
Σ PBDEs	5.01	0.82*	10.03	1.86*	0.07	0.01*	0.29	0.06*	4.08	0.54*	0.72	0.14*	0.08	0.01*	0.05	0.02*	
197	0.13	0.03#,*	0.27	0.08 ^{#,*}	< 0.01	< 0.01*	0.01	<0.01 ^{#,*}	0.07	0.01#,*	0.33	0.10 ^{#,*}	< 0.01	< 0.01*	0.02	0.01#,*	0.01
183	0.08	0.01#	0.16	0.07	< 0.01	< 0.01 ^{#,*}	< 0.01	<0.01 ^{#,*}	0.04	0.01#	0.53	0.09	< 0.01	< 0.01 ^{#,*}	0.04	0.02 ^{#,*}	0.01
153	0.11	0.02	0.21	0.07	< 0.01	< 0.01*	0.01	< 0.01*	0.07	0.01*	0.53	0.09*	< 0.01	< 0.01*	0.04	0.02*	0.01.
99	0.99	0.47*	1.97	0.49 ^{#,*}	0.01	0.02*	0.06	0.02*	0.86	0.12*	5.28	1.01#,*	0.02	< 0.01*	0.39	0.19*	0.05
100	0.59	0.12*	1.27	0.27^{*}	0.01	< 0.01*	0.04	0.01*	0.45	0.10*	2.17	0.42*	0.01	< 0.01*	0.15	0.05*	0.11
47	2.90	0.50^{*}	5.32	1.10*	0.04	< 0.01*	0.15	0.03*	2.46	0.39*	10.14	1.41*	0.04	0.01*	0.70	0.26*	0.33

534 In **bold**: sum (Σ) of all DDT, PCB & PBDE congeners. Lw: lipid weight, fw: fresh weight. TEQ: toxic equivalents, pg WHO-TEQ ⁻¹g fw,

calculated by using the toxic equivalency factors recommended by Van den Berg (1998). BEQ: bioequivalent values, BEQ g^{-1} fw. The # denotes a

significant difference between *C. gunnari* and *C. aceratus* in the given tissue at $p \le 0.05$. The * denotes a significant difference between tissues for

537 each species at $p \le 0.05$.

538 Figure Captions

- 540 Figure 1 Mean (\pm sem) concentration of Σ PCBs, Σ BDEs, HCBs, γ -HCHs and Σ DDTs (ng
- g^{-1} lipid weight), in muscle and ovaries of the two icefish species, *C. gunnari* (*n*=11) and *C*.
- *aceratus* (*n*=10). The # denotes a significant difference between C. gunnari and C. aceratus in
- 543 the given tissue at $p \le 0.05$. The * denotes a significant difference between tissues for each
- 544 species at $p \le 0.05$.
- **Figure 2:** Congener composition (percentage) of PCBs, DDTs and BDEs in muscle (m) and
- 547 ovaries (ov) of *C. gunnari* (*Cg*) and *C. aceratus* (*Ca*).

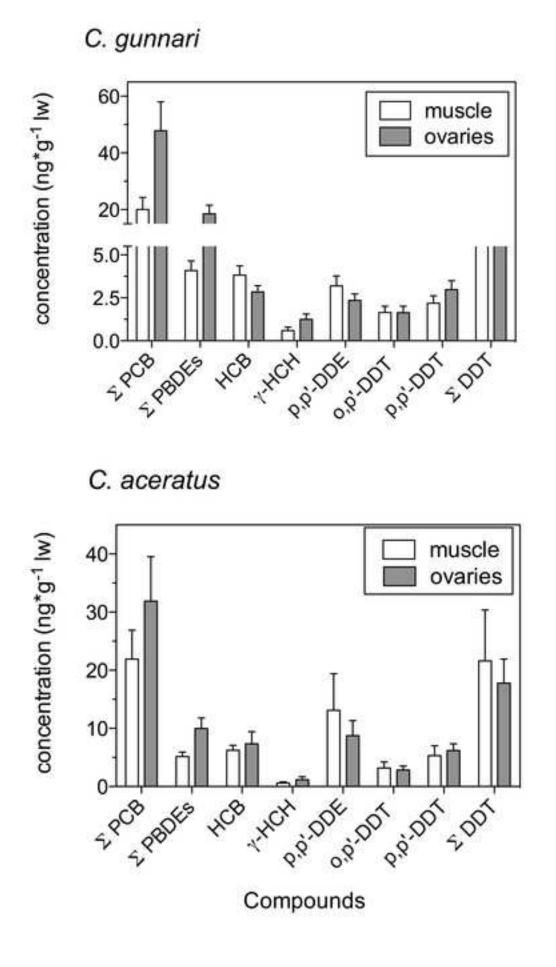
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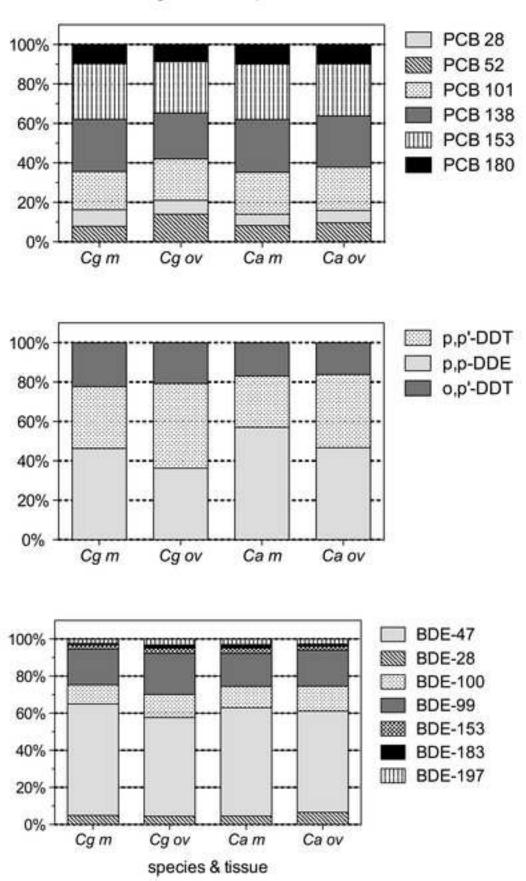
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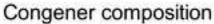
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