- Oxidative stress, metabolic activity and mercury concentrations in Antarctic krill *Euphausia superba* and myctophid fish of the Southern Ocean
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- 30 ABSTRACT

31 Indicators of oxidative stress and metabolic capacity are key factors in understanding the 32 fitness of wild populations. In this study, these factors were evaluated in the pelagic Southern 33 Ocean taxa Antarctic krill (Euphausia superba) and myctophid fish (Electrona antarctica, 34 Gymnoscopelus braueri and G. nicholsi) to establish a baseline record for future studies. 35 Mercury (Hg) concentrations in tissues were also analysed to evaluate its potential impacts 36 on species biochemical performance. E. superba had higher metabolic activity than the 37 myctophid species, which may explain the comparatively lower energy reserves found in the 38 former. The activity of antioxidant enzymes showed, generally, a lower level in E. superba 39 than in the myctophid species. The lack of any relationship between Hg levels and organism's 40 antioxidant and biotransformation defense mechanisms indicate that levels of Hg accumulated 41 in the studied species were not high enough to affect their biochemical processes adversely. 42 43 Keywords: Toxicity, Biochemical performance, Antioxidant capacity, Base line. 44

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47 The Southern Ocean ecosystem is distinctive for its low temperatures, large levels of 48 seasonal sea ice (Alberello et al., 2018), high nutrients concentrations (Brierley and Thomas, 49 2002), and the productive upwelling regions (Morrison et al., 2015). Globally, it is also 50 experiencing some of the highest levels of warming and ocean acidification (Freer et al., 2017; 51 IPCC, 2019; Rintoul et al., 2018; Turner et al., 2013), which can affect individuals at a 52 subcellular level and, in turn, alter patterns of distribution and food web structure (Atkinson et 53 al., 2004; Xavier and Peck, 2015). Warming increases the release of freshwater into the 54 Southern Ocean, particularly through accelerating the flow of glaciers, which liberates the 55 contaminants that they store. These contaminants have the potential to cause stress to 56 Southern Ocean fauna and it is important to describe and understand the impacts that they 57 may have, particularly to key biomass dominant taxa.

58 Antarctic krill Euphausia superba is a key species in the Southern Ocean trophic web, 59 being a major link between primary production and vertebrate predators (Everson, 2000). E. 60 superba is also a very important commercial species, with 260000 tonnes harvested in 2016 61 (Nicol et al., 2000; Tou et al., 2007); CCAMLR, 2017). This species is predominantly 62 herbivorous, feeding mainly on phytoplankton and rarely on some copepod crustaceans 63 (Everson, 2000). Whales, seals, penguins and flying seabirds are amongst those species that 64 consume high quantities of *E. superba* (Armstrong and Siegfried, 1991; Croxall et al., 1999; 65 Xavier et al., 2003). However, given evidence of a 50-80 % decline in E. superba over long 66 term (Atkinson et al., 2004), predators may have to switch to alternative prey groups. 67 Myctophid fish, the most abundant group in the mesopelagic fish community worldwide 68 (Gjøsaeter and Kawaguchi, 1980), can be considered a major alternative energy source to 69 predators in low *E. superba* abundance (Murphy et al., 2007; Saunders et al., 2018). In the 70 Southern Ocean, this group has an estimated biomass ranging between 70 and 200 million 71 tones (Collins et al., 2008; Suzuki et al., 2005). Myctophid fish are therefore an important 72 independent trophic link between primary consumers and a wide range of higher predators, 73 including king penguins (Olsson and North, 1997), albatrosses (Xavier et al., 2003), Antarctic 74 fur seals (Davis et al., 2006), squid (Kear, 1992) and Patagonian toothfish (Collins et al., 2007).

75 Myctophid fish prev mainly on zooplankton (Saunders et al., 2019) and undertake diurnal 76 vertical migration to feed and to avoid predators in surface waters during the day. Prev-77 selection differs between myctophid species: *Electrona antarctica* consumes the amphipod 78 Themisto gaudichaudii during summer time; Gymnoscopelus braueri preys on different 79 species, including Themisto gaudichaudii, Metridia spp. E. superba, Pleuromamma robusta 80 and ostracods; and G. nicholsi diet is dominated by Metridia spp. and E. superba during the 81 summer (Lourenço et al., 2017; Saunders et al., 2019; 2018). The vertical distribution of these 82 myctophid species varies from 0 to 700 m, with *E. antarctica* being the species with the widest 83 spread of depths through the water column (Collins et al., 2008; Saunders et al., 2019; 2014). 84 When exposed to stressful conditions, including the presence of pollutants, organisms 85 may resort to an overproduction of reactive oxygen species (ROS) in their cells, leading to a 86 state known as oxidative stress (Regoli and Giuliani, 2014). To prevent the establishment of 87 oxidative stress, cells possess an extensive antioxidant system, that includes enzymatic and 88 non-enzymatic forms such as the enzymes superoxide dismutase, catalase, glutathione 89 peroxidase, and reduced glutathione (Regoli et al., 2011). Depending on the stress level and 90 organism's antioxidant capacity, cellular damage (namely through lipid peroxidation) and loss 91 of redox balance (with increasing oxidation of reduced glutathione) may occur. To cope with 92 oxidative stress, organisms may need to increase their metabolic capacity to ramp up their 93 defence mechanisms, leading to increased electron transport system (ETS) activity and 94 expenditure of energy reserves (e.g. glycogen) (Cruz et al., 2016; Freitas et al., 2020). Several 95 pollutants, including mercury (Hg), have already been shown to cause oxidative stress, as well 96 as alterations on their metabolic capacity, in marine organisms (Coppola et al., 2018; Monteiro 97 et al., 2019).

Despite the ecological importance of these two groups (*E. superba* and myctophid fish) in the pelagic realm, there is still a crucial knowledge gap regarding their ecophysiology (Atkinson et al., 2002; Meyer, 2012; Quetin and Ross, 1991), with only a few studies considering this aspect in *E. superba*, and virtually none in Southern Ocean myctophid, only two looking into ETS as a proxy for respiration rates (Belcher et al., 2020; 2019). The main

103 goal of the present study is to describe the general oxidative stress and metabolic status of *E*.
104 superba and of other three species of Antarctic myctophid fish (*E. antarctica, G. braueri* and
105 *G. nicholsi*), to evaluate Hg concentration in these species and possible impacts on their
106 biochemical performance, and to establish a base record for future studies. For this,
107 antioxidant capacity, cellular damage, redox balance, metabolic capacity and energy reserves
108 content were evaluated in all the mentioned species.

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110 Samples were collected on board of the British research vessel RRS James Clark 111 Ross during the austral summers of 2015/2016 (December 2015 and January 2016) in the 112 Scotia Sea (cruise JR15004). Euphausia superba specimens were collected from the water 113 column using an 8 m² mouth-opening Rectangular Midwater Trawl (RMT8; mesh size reducing 114 from 4.5 mm to 2.5 mm in the cod end) (Roe and Shale, 1979). The net was rigged with two 115 nets that could be remotely opened and closed at different depths. Myctophid samples 116 (Electrona antarctica, Gymnoscopelus braueri and G. nicholsi) were collected using a similar net design, with 25 m² mouth-opening (RMT25; mesh size reducing from 8 mm to 4.5 mm in 117 118 the cod end) (Roe and Shale, 1979). Samples were preserved in individual sample bags at -119 80 °C.

120 Euphausia superba in the catches were identified and total length (TL) of each 121 individual was measured, from the anterior edge of the eye to the tip of the telson and rounded 122 down (Morris et al., 1992). Sex and maturity stage were determined with reference to the 123 presence of a petasma (males), thelycum (females) or absent (juveniles; individuals without 124 visible external sexual characteristics) (Ross and Quetin, 2000). Myctophids were identified 125 using published guides (Gon and Heemstra, 1990; Hulley, 1990) and measured for the nearest 126 mm using standard length (SL). Sex and maturity was determined whenever possible: in some 127 myctophid species (e.g. E. antarctica), there is sexual dimorphism associated with the location 128 of photophores, but when this was not possible, the gonads were examined following 129 dissection (Yamamoto, 1969).

From each species (*E. superba* (n= 20); *E. antarctica* (n= 5); *G. braueri* (n= 5); *G. nicholsi* (n= 3)), samples were homogenized and used for biochemical markers measurements and for total mercury (Hg) determination. For *E. superba*, whole individuals were used for biochemical and Hg analysis, while for the myctophids species only muscle was analysed to avoid the inclusion of any bone.

135 Biochemical parameters were analysed in each species. For this, 20 individuals of E. 136 superba (10 females, 10 males), and muscle tissue from 5 G. braueri (2 males, 3 unknow), 3 137 G. nicholsi (2 females, 1 male) and 5 E. antarctica (3 females, 2 males) (Table 1) were homogenized using a mortar and pestle with liquid nitrogen and sonicated for 15 s at 4 °C, 138 139 after buffer addition (1:2) (Carregosa et al., 2014). To determine the activity of superoxide 140 dismutase (SOD), catalase (CAT), glutathione peroxidase (GPx), glutathione S-transferases 141 (GSTs) enzymes and the content of glycogen (GLY) and protein (PROT), supernatants were 142 extracted with a potassium phosphate buffer (50 mmol/L KH₂PO₄, 1 mmol/L ethylenediamine 143 tetraacetic acid disodium salt hydrate (EDTA), 1 % (v/v) Triton X-100, 1 mmol/L dithiothreitol 144 (DTT), pH 7.0). For lipid peroxidation (LPO) assessment, supernatants were obtained using 145 20 % (w/v) trichloroacetic (TCA). For electron transport system (ETS) activity evaluation the 146 homogenizing buffer (0.1 mol/L Tris-HCl, 15 % (w/v) polyvinylpyrrolidone (PVP), 153 mmol/L 147 Mg SO₄, and 0.2 % (v/v) Triton-X 100, pH 8.5) was used. Samples were centrifuged for 20 148 min at 10 000 g (3 000 g for ETS) and 4° C (Carregosa et al., 2014), and supernatants were 149 preserved at -80 °C or analysed immediately.

150 The GLY content was determined according to the sulfuric acid method (DuBois et al., 151 1956), using glucose standards (between 0-10 mg/mL) to obtain a calibration curve. 152 Absorbance was measured at 492 nm after 30 min incubation at room temperature and results 153 were expressed in mg per g of fresh weight (FW). The PROT content was determined 154 according to Robinson and Hogden (1940), following the Biuret method that uses bovine 155 serum albumin (BSA) as standard (0 to 40 mg/mL) to obtain a calibration curve. After 10 min 156 incubation at 30 °C, the absorbance was read at 540 nm. The results were expressed in mg 157 per q of FW.

Metabolic capacity was assessed by measuring the ETS activity, following the method
of King and Packard (1975) and modifications by De Coen and Janssen (1997). Absorbance
was measured during 10 min at 490 nm in 25 s intervals and the extinction coefficient (ε)
15,900/M/cm was used to calculate the amount of formazan formed per unit time. Results
were expressed in nmol min per g of FW.

Cellular damage was measured by the quantification of LPO levels following the
method described in Ohkawa et al. (1979) with modifications referred by Carregosa et al.
(2014). Absorbance was measured at 535 nm and LPO was determined using the extinction
coefficient (ε) 156/mM/cm and results expressed in nmol of MDA equivalents formed per g of
FW.

168 The activity of SOD was quantified based on the method of Beaucham and Fridovic 169 (1971). SOD standards (0.25-60 U/mL) were used to generate a calibration curve. After 20 170 min incubation at room temperature, absorbance was measured at 560 nm. Results were 171 expressed in U per g FW and by U per mg of PROT. The activity of CAT was quantified 172 following Johansson and Borg (1988). Formaldehyde standards (0-150 µM) were used to 173 produce a calibration curve. Absorbance was measured at 540 nm and results expressed in 174 U per g FW and per mg of PROT. The activity of GPx was determined following the method 175 of Paglia and Valentine (1967). Absorbance was measured at 340 nm during 5 min in 10 s 176 intervals. Enzyme activity was calculated using the extinction coefficient (ε) 6.22/mM/cm and 177 the results were expressed in U per g FW and per mg of PROT. The activity of GSTs was 178 determined at room temperature using 1- chloro-2,4-dinitrobenzene (CDNB) as substrate 179 according to the method described by Habig et al. (1974) with modifications described in 180 Carregosa et al. (2014). The activity was determined spectrophotometrically at 340 nm using 181 the extinction coefficient (ε) 9.6/mM/cm and absorbance was measured in intervals of 10 s 182 during 5 min. The GSTs activity expressed in U per q FW and by U per mg of PROT.

183 Mercury concentrations were determined by thermal decomposition atomic absorption 184 spectrometry with gold amalgamation, using a LECO AMA-254 (Advanced mercury analyser) 185 following Coelho et al. (2008) methodology. Analytical quality control was performed using

186 certified reference material (CRM; in this case TORT-2 and TORT-3 [lobster hepatopancreas, 187 National Research Council, Canada] for E. superba; DORM-4 [Fish protein, National 188 Research Council, Canada] for myctophids). The obtained values (mean ± standard deviation 189 for the whole of the CRM analyses (n=13) provided recoveries ranging from 84 to 96 % (TORT-190 2: 88 ± 3 %; TORT-3: 89 ± 7 %; DORM-4: 91 ± 12%). The mass of CRM used for quality 191 control analyses was adjusted to be within the range of total Hg present in the samples. Blanks 192 were analysed at the beginning of each set of samples and the analyses were always 193 performed at least in duplicate, until coefficient of variation were below 10%.

Data obtained from toxicology, biochemical analyses and biological factor (size, weight, sex and species) were submitted to permutational multivariate analysis of variance with the PERMANOVA+add-on in PRIMER v6 (Anderson et al., 2009). The pseudo-F values in the PERMANOVA main tests were evaluated in terms of significance. When the main test revealed statistical significant differences (p < 0.05), pairwise comparisons were performed. The null hypothesis tested for each parameter was: for each biochemical parameters and Hg concentration, no significant differences existed among species.

Biochemical responses and Hg concentrations for the different species were used to calculate the Euclidean distance similarity matrix. This matrix was simplified through the calculation of the distance within the centroid matrix based on species, which was then submitted to ordination analysis, performed by Principal Coordinates (PCO). Pearson correlation vectors (r > 0.85) of physiological and biochemical descriptors and Hg concentrations were provided as supplementary variables being superimposed on the top of the PCO graph.

For all the studied species, both sexes were initially analysed separately and since no significant differences were observed in terms of biochemical performance the total number of organisms per species was pooled.

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212 In terms of biochemical performance and Hg concentration in all the analysed 213 organisms, the PCO analysis clearly highlights differences between species, with PCO1

explaining 64.8% of the total variation and distinguishing between *E. superba* (KRI) on the
negative side from the myctophids species in the positive side. PCO2 explains 26.0% of the
total variation, separating *E. antarctica* (ELN) and *G. braueri* (GYB) in the negative side from *G. nicholsi* (GYN) and *E. superba* (KRI) in the positive side (Figure 1).

218 In term of Hg levels, most probably due to lower trophic level and lifespan, E. superba 219 had significantly lower Hg concentrations than all the myctophid fish species analysed (Table 220 2), with Hg concentrations in *E. superba* within the range of values observed in previous 221 studies with the same species (0.008 to 0.077 µg/g, Seco et al., 2019). The results obtained 222 also showed that Hg concentrations did not significantly differ among myctophid species. To 223 our knowledge, there is only two previous studies that reports Hg concentration in Southern 224 Ocean myctophids, with reported concentrations within the same range as the ones obtained 225 in the current study (Cipro et al., 2018; Seco et al., 2020).

226 In terms of energy reserves, individuals of *E. superba* had a relatively low PROT 227 content (Table 2) when compared with other crustaceans (Aristeus antennatus, Parapenaeus 228 longirostris and Nephrops norvegicus) but the content of GLY was higher than in other 229 crustacean species (Rosa and Nunes, 2003). These findings can be related to the fact that 230 the current study was performed during the austral summer, when Southern Ocean 231 inhabitants tend to build up energy reserves for winter time. For the 3 myctophid species, 232 significant differences of the quantity of PROT (*E. antarctica* \leq *G. braueri* \leq *G. nicholsi*) were 233 observed between E. antarctica and G. nicholsi, which may be related to the difference in size between both species' individuals (G. nicholsi individuals were 2 times the length of E. 234 235 Antarctica individuals; Table 1). This result is corroborated by the PCO analysis, where a close 236 relationship between PROT content and size and weight were found (Figure 1). No significant 237 differences on the PROT content were observed between G. braueri and the other 2 238 myctophid species. In terms of GLY content, no significant differences were observed among 239 the 3 myctophid species, while E. superba had lower GLY values than G. nicholsi but there 240 were no significant differences between *E. superba* and the other two myctophid species.

241 PROT and GLY contents were lower in *E. superba* compared to the three myctophid fish, 242 which may indicate higher energetic requirements by the crustacean compared with the fish 243 species. However, in the present study, E. superba was not the species with the highest 244 metabolic activity and, therefore, lower GLY and PROT concentrations did not result from their 245 expenditure in response to increased metabolism. In fact, in terms of metabolic capacity, no 246 significant differences were observed between E. superba and both Gymnoscopelus species 247 (Table 2). Also, the highest ETS activity observed in G. nicholsi was not accompanied by 248 higher energy expenditure as this species presented the highest GLY and PROT contents. 249 Low energy reserves content in *E. superba* can be related with lower production and/or 250 accumulation capacity in the crustacean compared with the fish species, however there is still 251 a lack of knowledge regarding the energy cycle and reverses in this species. We can also 252 hypothesize that differences in the energy reserves between *E. superba* and myctophid fish 253 may be related with the dietary differences between the groups as E. superba feeds on 254 phytoplankton (Everson, 2000) whereas myctophid fish feed mainly in zooplankton (e.g.: E. 255 superba, Metridia spp. and on Rhincalanus gigas (Saunders et al., 2018)).

256 As demonstrated previously, organisms tend to reduce their metabolism as a strategy 257 to avoid accumulation of toxic substances, reducing for example their filtration rate and, 258 consequently, ingestion of contaminants, as reported in estuarine bivalves (Almeida et al., 259 2014; 2015; Pinto et al., 2019). Nevertheless, in the present study, the lowest ETS activity (E. 260 antarctica) was not associated with the lowest Hg concentrations and, in the same way, the 261 highest ETS activity observed in *G. nicholsi* did not correspond to higher Hg concentrations. 262 The ETS activity in myctophids showed to be positively correlated with size, as demonstrated 263 in a previous study looking into ETS as a respiration rate, larger myctophids had higher ETS 264 (Belcher et al., 2020). These preliminary data suggest that both groups of organisms may not 265 decrease their metabolism as a strategy to avoid accumulation of pollutants or, most probably, 266 Hg concentration in the environment was not the factor that conditioned species metabolism, 267 since accumulated levels were very low.

268 Many Southern Ocean cold waters inhabitants have generally slow activity and low 269 metabolic rates (Abele and Puntarulo, 2004). In theory, this should result in lower rates of 270 reactive oxygen species (ROS) formation in ectotherm species. It is well known that 271 mitochondria respiration system is responsible for the generation of ROS which are 272 responsible for cellular damage (including LPO). In the present study although lower metabolic 273 rate (identified by lower ETS values) observed in E. antarctica was not accompanied by lower 274 LPO values, higher LPO levels observed in *G. nicholsi* may result from higher ETS activity 275 recorded in this species. In particular, LPO levels varied inter-specifically (G. nicholsi $\geq E$. 276 antarctica > E. superba > G. braueri). Euphausia superba presented significantly higher LPO 277 levels than G. braueri, but lower than G. nicholsi. In the fish group, E. antarctica and G. nicholsi 278 had higher levels of LPO than *G. braueri* (Table 2). It is well described that LPO may occur as 279 a consequence of pollutants exposure due to overproduction of ROS and inefficiency of 280 antioxidant mechanisms (among others, Regoli and Giuliani, 2014). In the present study, the 281 highest LPO levels identified in G. nicholsi did not correspond to higher Hg tissue 282 concentrations, which, once again, may corroborate the hypothesis that Hg concentrations 283 observed in organisms were not high enough to induce cellular alterations. Also, a study on 284 oxidative stress profiles on Dicentrarchus labrax, demonstrated that in some cases higher 285 contamination levels do not result in LPO increase (Hg 0.04 μ g/g and 0.08 μ g/g) (Mieiro et al., 286 2011).

287 Regarding the activity of antioxidant enzymes (Table 2, Figure 1), significantly lower values 288 were observed in E. superba than in some of the myctophid group (E. antarctica and G. 289 nicholsi for SOD and G. braueri for CAT), while no significant differences were found among 290 the three myctophid species. GPx activity presented no changes among all the analysed 291 species whereby it may not be influenced by environmental conditions, with similar response 292 in all studied species. Detoxification enzymes, like GSTs, are related to elimination routes of 293 contaminants (e.g., Hg) (Elia et al., 2003). In the present study GSTs levels showed no 294 significant differences among species, with no relationship with the Hg concentrations in

295 organism's tissues. The results obtained for GSTs activity were lower than the ones observed 296 by other authors for *E. superba* (Tremblay and Abele, 2015). Once again, these results may 297 indicate that due to the low Hg concentration in seawater, and consequently low accumulation 298 levels in the studied species, defence mechanisms were not responding to Hg tissues 299 concentrations. From the literature published on this topic it is possible to conclude that 300 antioxidant responses in E. superba vary between studies: SOD levels were lower in the 301 present study, but CAT activity was higher in samples collected in 2011, also around South 302 Georgia (Tremblay and Abele, 2015). GPx was also lower than in individuals captured in the 303 eastern Antarctic sector in 2006, when compared with the control individuals of the study 304 performed by Dawson 2017. Thus, the obtained results demonstrated that the activity of 305 antioxidant and biotransformation enzymes was similar among species, regardless the LPO 306 levels and Hg concentration. Such response may indicate low stress levels in the organisms 307 caused by low Hg concentrations accumulated by the organisms, with increased LPO levels 308 resulting from increased metabolic capacity rather than contamination levels.

309 Overall, the present findings highlighted that G. nicholsi presented higher levels in 310 almost all the analysed biochemical parameters. This performance may be due to a difference 311 in size, compared with the other analysed species. To the best of our knowledge, this is the 312 first study reporting biochemical parameters in Southern Ocean myctophid fish. The unique 313 environmental features of the Southern Ocean, and its highly specialized inhabitants, are likely 314 to make it very sensitive to environmental change. With ocean warming, increased levels of 315 glacial melt and higher amounts of freshwater input, levels of contaminants (like Hg) may 316 increase further into the future. Oxidative stress and metabolic capacity will be among the first 317 biological responses of resident species to contaminants. So, it is important to describe levels 318 of natural variation in these parameters for future comparisons.

The present study provides values for a number of metabolic parameters (PRO, GLY, ETS, LPO, CAT, ETS, GPx, SOD, GSTs) for *E. superba* and three biomass dominant myctophid species, all of which play key roles in Southern Ocean ecosystem function. At present, none of the studied biochemical parameters shows any positive or negative

relationship with levels of Hg found within tissues. Nevertheless, these values provide an important baseline to establish whether any future increases in contamination levels are having a notifiable effect on species' metabolic capacity and biochemical performance.

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Table 1. Total number, mean length and weight values for each collected species and location (latitude and longitude) of sampling areas on the Scotia sea, surveyed during the austral summer of 2015/2016.

Species	Ν	<i>Total length</i> (mm)	<i>Weight</i> (g)	Latitude	Longitude	
Euphausia superba	20	5.1 (± 0.3)	1.2 (±0.3)	-60.3131	-46.8488	
Electrona antarctica	5	82 (± 14)	6.3(±5.4)	-59.9861	-47.22192	
Gymnoscopelus braueri	5	104 (± 11)	9.1 (±2.9)	-59.9861	-47.22192	
Gymnoscopelus nicholsi	3	148 (± 8)	14.7 (±5.9)	-60.33097	-46.67431	

Table 2. Biomarkers and mercury (Hg) concentration measured in each species. Biomarkers: PROT: protein content; LPO: lipid peroxidation levels; ETS: Electron transport system; GLY: Glycogen content; GPx: Glutathione peroxidase; GSTs: Glutathione S-transferases activity; SOD: superoxide dismutase activity; CAT: catalase activity. Units are presented per gram of fresh weight (FW) and per mg of protein (PROT) for comparison with result from the literature.

Hg	GLY	PROT	ETS LPO		SOD		CAT		GPx		GSTs		
μg g ⁻¹ DW	mg/g FW	mg/g FW	nmol/min/g FW	nmol MDA/g FW	nmol.mg .PROT	U/g FW	U.mg. PROT	U/g FW	U.mg. PROT	U/g FW	U.mg. PROT	U/g FW	U.mg. PROT
<u>0.04</u> (± 0.01) ^a	<u>18.65</u> (<u>±3.27)</u> ª	<u>41.36</u> (±2.48)ª	33.80 (±8.37)ª	24.82 (±3.90)ª	0.82 (±0.21)	<u>1.96</u> (±0.92)ª	0.02 (±0.005)	<u>10.79</u> (<u>±3.83)</u> ª	0.13 (±0.05)	5.15 (±3.22)ª	0.25 (±0.16)	0.05 (±0.02) ^a	0.78 (±0.57)
0.22 (± 0.08) ^b 0.17 (± 0.03) ^b 0.17 (± 0.04) ^b	21.19 (±4.52) ^{a, b} 24.03 (±4.38) ^{a, b} 25.54 (±1.41) ^b	64.81 (±9.17) ^b 85.34 (±22.57) ^{b, c} 118.99 (±38.51) ^c	<u>17.88</u> (±5.77) ^b 22.36 (±8.82) ^{a, b} 42.45 (±10.71) ^{a, b}	36.19 (±8.41) ^{a, c} 15.77 (± 9.10) ^b 76.86 (±7.99) ^c	0.28 (±0.11) 0.26 (±0.05) 0.34 (±0.17)	4.19 (±0.90) ^b 2.74 (±0.81) ^{a, b} 3.73 (±0.84) ^b	$\begin{array}{c} 0.03 \\ (\pm 0.01) \\ \hline 0.01 \\ (\pm 0.009) \\ \hline 0.01 \\ (\pm 0.005) \end{array}$	17.45 (±6.80) _{a, b} 22.60 (±7.96) ^b 20.11 (±10.20) ^{a, b}	$\begin{array}{c} 0.13 \\ (\pm 0.05) \\ 0.13 \\ (\pm 0.02) \\ 0.09 \\ (\pm 0.05) \end{array}$	2.03 (±1.97) ^a 3.48 (±2.27) ^a 5.07 (±1.92) ^a	$\begin{array}{c} 0.07 \\ (\pm 0.07) \\ 0.09 \\ (\pm 0.06) \\ 0.11 \\ (\pm 0.04) \end{array}$	0.05 (±0.01) ^a <u>0.04</u> (±0.02) ^a 0.13 (±0.04) ^a	$\begin{array}{c} 1.23 \\ (\pm 0.22) \\ 0.77 \\ (\pm 0.53) \\ 1.79 \\ (\pm 1.66) \end{array}$
	Hg μg g ⁻¹ DW 0.04 (± 0.01) ^a 0.22 (± 0.08) ^b 0.17 (± 0.03) ^b 0.17 (± 0.04) ^b	Hg GL Y μ g g ⁻¹ DW mg/g FW $0.04 \\ (\pm 0.01)^a$ $\frac{18.65}{(\pm 3.27)^a}$ 0.22 (\pm 0.08)^b $21.19 \\ (\pm 4.52)^{a,b}$ 0.17 (\pm 0.03)^b $24.03 \\ (\pm 4.38)^{a,b}$ 0.17 (\pm 0.04)^b $(\pm 1.41)^b$	HgGLYPROT $\mu g g^{-1} DW$ mg/g FWmg/g FW $0.04 (\pm 0.01)^a$ $\frac{18.65}{(\pm 3.27)^a}$ $\frac{41.36}{(\pm 2.48)^a}$ $0.22 (\pm 0.08)^b$ $(\pm 4.52)^{a,b}$ $(\pm 9.17)^b$ $(\pm 0.08)^b$ $(\pm 4.32)^{a,b}$ $(\pm 2.57)^{b,c}$ $0.17 (\pm 0.03)^b$ $(\pm 4.38)^{a,b}$ $(\pm 22.57)^{b,c}$ $0.17 (\pm 0.04)^b$ $(\pm 1.41)^b$ $(\pm 38.51)^c$	HgGLYPROTETS $\mu g g^{-1} DW$ mg/g FWmg/g FWnmol/min/g FW $0.04 (\pm 0.01)^a$ $\frac{18.65}{(\pm 3.27)^a}$ $\frac{41.36}{(\pm 2.48)^a}$ 33.80 $(\pm 8.37)^a$ $0.22 (\pm 0.08)^b$ $(\pm 4.52)^{a,b}$ $(\pm 9.17)^b$ $\frac{17.88}{(\pm 5.77)^b}$ $0.17 (\pm 0.03)^b$ $(\pm 4.38)^{a,b}$ $(\pm 22.57)^{b,c}$ $(\pm 8.2)^{a,b}$ $0.17 (\pm 0.03)^b$ $(\pm 1.41)^b$ $(\pm 38.51)^c$ $(\pm 10.71)^{a,b}$	HgGLYPROTETSI $\mu g g^{-1} DW$ mg/g FWmg/g FWnmol/min/g FWnmol MDA/g FWnmol MDA/g FW $0.04 (\pm 0.01)^a$ $\frac{18.65}{(\pm 3.27)^a}$ $\frac{41.36}{(\pm 2.48)^a}$ 33.80 $(\pm 8.37)^a$ 24.82 $(\pm 3.90)^a$ $0.22 (\pm 0.08)^b$ $(\pm 4.52)^{a,b}$ $(\pm 9.17)^b$ $\frac{17.88}{(\pm 5.77)^b}$ 36.19 $(\pm 8.41)^{a,c}$ $0.17 (\pm 0.03)^b$ $(\pm 4.38)^{a,b}$ $(\pm 22.57)^{b,c}$ $(\pm 8.2)^{a,b}$ 15.77 $(\pm 9.10)^b$ $0.17 (\pm 0.04)^b$ $(\pm 1.41)^b$ $(\pm 38.51)^c$ $(\pm 10.71)^{a,b}$ $(\pm 7.99)^c$	HgGLYPROTETSLPO μ g g-1 DWmg/g FWmg/g FWnmol/min/g FWnmol MDA/g FWnmol.mg NDA/g FWnmol.mg NDA/g FWnmol.mg NDA/g FW $0.04(\pm 0.01)^a$ $\frac{18.65}{(\pm 3.27)^a}$ $\frac{41.36}{(\pm 2.48)^a}$ 33.80 ($\pm 8.37)^a$ 24.82 ($\pm 3.90)^a$ 0.82 (± 0.21) $0.22(\pm 0.08)^b$ $21.19(\pm 4.52)^{a,b}$ 64.81 ($\pm 9.17)^b$ $\frac{17.88}{(\pm 5.77)^b}$ 36.19 ($\pm 8.41)^{a,c}$ 0.28 (± 0.11) 0.17 ($\pm 0.03)^b$ 24.03 ($\pm 4.38)^{a,b}$ 85.34 ($\pm 22.57)^{b,c}$ 22.36 ($\pm 8.82)^{a,b}$ 15.77 ($\pm 9.10)^b$ 0.26 (± 0.05) 0.17 ($\pm 0.04)^b$ 25.54 ($\pm 1.41)^b$ 118.99 ($\pm 38.51)^c$ 42.45 ($\pm 10.71)^{a,b}$ 76.86 ($\pm 7.99)^c$	HgGLYPROTETSLPOSC $\mu g g^{-1} DW$ mg/g FWmg/g FWnmol/min/gnmol/min/gnmol/mDA/gnmol.mgU/g FW $0.04 (\pm 0.01)^a$ $\frac{18.65}{(\pm 3.27)^a}$ $\frac{41.36}{(\pm 2.48)^a}$ 33.8024.820.82 $\frac{1.96}{(\pm 0.21)}$ $(\pm 0.08)^b$ $(\pm 3.27)^a$ $\frac{41.36}{(\pm 2.48)^a}$ $\frac{17.88}{(\pm 5.77)^b}$ $(\pm 8.41)^{a, c}$ (± 0.21) $\frac{1.96}{(\pm 0.92)^a}$ 0.22 (± 1.19) $(\pm 4.52)^{a, b}$ $(\pm 9.17)^b$ $\frac{17.88}{(\pm 5.77)^b}$ $(\pm 8.41)^{a, c}$ (± 0.11) $(\pm 0.90)^b$ 0.17 24.03 85.34 22.36 15.77 0.26 2.74 $(\pm 0.03)^b$ $(\pm 4.38)^{a, b}$ $(\pm 22.57)^{b, c}$ $(\pm 8.82)^{a, b}$ $(\pm 9.10)^b$ (± 0.05) $(\pm 0.81)^{a, b}$ 0.17 25.54 118.99 42.45 76.86 0.34 3.73 $(\pm 0.04)^b$ $(\pm 1.41)^b$ $(\pm 38.51)^c$ $(\pm 10.71)^{a, b}$ $(\pm 7.99)^c$ (± 0.17)	HgGLYPROTETSLPOSOD $\mu g g^{-1} DW$ mg/g FWmg/g FWnmol/min/gnmol/min/gnmolmmol.mgU/g FWU.mg. 0.04 ($\pm 0.01)^a$ 18.65 ($\pm 3.27)^a$ 41.36 ($\pm 2.48)^a$ 33.80 ($\pm 8.37)^a$ 24.82 ($\pm 3.90)^a$ 0.82 (± 0.21) 1.96 (± 0.21) 0.02 ($\pm 0.92)^a$ 0.22 ($\pm 0.08)^b$ 21.19 ($\pm 4.52)^{a,b}$ 64.81 ($\pm 9.17)^b$ 17.88 ($\pm 5.77)^b$ 36.19 ($\pm 8.41)^{a,c}$ 0.28 (± 0.11) 4.19 ($\pm 0.90)^b$ 0.03 (± 0.01) 0.17 ($\pm 0.03)^b$ 85.34 ($\pm 22.57)^{b,c}$ 22.36 ($\pm 8.82)^{a,b}$ 15.77 ($\pm 9.10)^b$ 0.26 (± 0.05) 2.74 ($\pm 0.81)^{a,b}$ 0.01 (± 0.009) 0.17 ($\pm 0.04)^b$ 118.99 ($\pm 1.41)^b$ 42.45 ($\pm 38.51)^c$ 76.86 ($\pm 7.99)^c$ 0.34 (± 0.17) 0.01 ($\pm 0.84)^b$	HgGLYPROTETSLPOSODCA $\mu g g^{-1} DW$ mg/g FWmg/g FWnmol/min/gnmol/min/gnmol/mol/mol/mol/mol/mol/mol/mol/mol/mol/	HgGLYPROTE1SLPOSODCAT $\mu g g^{-1} DW$ mg/g FWmg/g FWnmol/min/g FWnmol/min/g FWnmol/mDA/g FWnmol.mg .PROTU/g FWU.mg. 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Figure 1. Ordination diagram based on biomarkers, biological characteristic and mercury (Hg) concentration. ELN - *Electrona Antarctica*, GYB - *Gymnoscopelus braueri*; GYN - *Gymnoscopelus nicholsi*; KRI - *Euphausia superba*; PROT - protein content; LPO - lipid peroxidation levels; ETS - Electron transport system activity; GLY - Glycogen content; GPx - Glutathione peroxidase; GSTs - Glutathione S-transferases activity; SOD - superoxide dismutase activity; CAT - catalase activity; Hg - Mercury concentration.

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