First evidence of legacy chlorinated POPs bioaccumulation in Antarctic sponges from the Ross sea and the South Shetland Islands

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OUTRO

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## 28 Abstract

29 Antarctica is no longer pristine due to the confirmed presence of anthropogenic contaminants like 30 Persistent Organic Pollutants (POPs). Benthic organisms are poorly represented in contamination 31 studies in Antarctica although they are known to bioaccumulate contaminants. Sponges (Phylum 32 Porifera) are dominant members in Antarctic benthos, both in terms of abundance and biomass, and 33 are an important feeding source for other organisms, playing key functional roles in benthic 34 communities. To the best of our knowledge, legacy chlorinated POPs such as polychlorinated biphenyls (PCBs), hexachlorobenzene (HCB), and dichlorodiphenyltrichloroethane (DDT) and their 35 36 metabolites have never been investigated in this Phylum in Antarctica. The aim of this work was to 37 evaluate the bioaccumulation of PCBs, HCB, o,p'- and p,p'-DDT and their DDE and DDD isomers in 35 sponge samples, belonging to 17 different species, collected along the coast of Terra Nova Bay 38 39 (Adèlie Cove and Tethys Bay, Ross Sea), and at Whalers Bay (Deception Island, South Shetland 40 Islands) in Antarctica. Lipid content showed a significant correlation with the three pollutant classes. The overall observed pattern in the three study sites was  $\Sigma PCBs > \Sigma DDTs > HCB$  and it was found in 41 42 almost every species. The  $\Sigma$ PCBs,  $\Sigma$ DDTs, and HCB ranged from 54.2 to133.7 ng/g lipid weight (lw), 43 from 17.5 to 38.6 ng/g lw and from 4.8 to 8.5 ng/g lw, respectively. Sponges showed contamination 44 levels comparable to other Antarctic benthic organisms from previous studies. The comparison 45 among sponges of the same species from different sites showed diverse patterns for PCBs only in one 46 out of four cases. The concentration of POPs did not vary significantly among the three sites. The 47 predominance of lower chlorinated organochlorines in the samples suggested that long-range 48 atmospheric transportation (LRAT) could be the major driver of contamination as molecules with a 49 high long range transport potential (e.g. low chlorinated PCBs, HCB) prevails on heavier ones.

50

- 51 Keywords
- 52 PCBs
- 53 Chlorinated pesticides
- 54 Porifera
- 55 Southern Ocean
- 56 Bioaccumulation
- 57 Benthic organisms
- 58

## 59

Journal Prespos

## 60 Introduction

61

62 Due to its geographical isolation and the absence of human activities, except for research, industrial 63 fishing, and tourism, Antarctica and the Southern Ocean are usually regarded as one of the most 64 pristine regions on Earth (Kim et al. 2015; Vecchiato et al. 2015; Vergara et al. 2019). Nevertheless, 65 anthropogenic contaminants can reach Antarctica through long-range transport mechanisms. In fact, 66 semi-volatile compounds are subjected to the global distillation process consisting of repeated 67 evaporation and condensation events that can transport Persistent Organic Pollutants (POPs) far from 68 their emission sources (Wania & Mackay, 1993). Once in the Polar Regions, amplification 69 mechanisms such as cold condensation (Wania and Mackay, 1993) or snow scavenging (Casal et al. 2019) result in a preferential accumulation of POPs in both the Arctic and Antarctica. POPs fall out 70 71 through dry or wet depositions but also enter marine ecosystems transported by global ocean currents 72 (Casas et al. 2020; Casas et al. 2022) and from pack ice melting (Casal et al. 2019; Potapowicz et al. 73 2019).

74 Among legacy POPs, polychlorinated biphenyls (PCBs), hexachlorobenzene (HCB), and 75 dichlorodiphenyltrichloroethane (DDT) are the most studied worldwide (Stockholm Convention, 76 2004). These persistent and toxic compounds, even if banned or restricted decades ago, are still found 77 in every region worldwide, including Antarctica (Bargagli, 2008; Corsolini, 2009; Mello et al. 2016; 78 Morales et al. 2022). Legacy POPs that may have been stored in the deeper layers of glaciers, 79 perennial pack ice, and ice shelves may further be released during accelerated glacier melt due to 80 climate change, becoming available again for bioaccumulation in the food webs (Ma et al. 2011; 81 Potapowicz et al. 2019). Legacy contamination is increasingly being studied, as polar regions are experiencing some of the most rapid impacts of warming, acidification, and sea ice loss, and 82 83 impacting benthic communities (Meredith et al. 2019; Brasier et al. 2021; Di Giglio et al. 2020; 84 Figuerola et al. 2021, 2022). Bearing that in mind, it is valuable to keep studying the environmental

fate of legacy POPs, their transfer through the marine food webs and their potential effects in Arctic
and Antarctic ecosystems.

87 Antarctic food webs have peculiar characteristics: they are based on very few key species, such as 88 the Antarctic krill Euphausia superba Dana 1950, and the Antarctic silverfish Pleuragramma 89 antarctica Boulenger 1902, and thus they are likely fragile and vulnerable, with a very low resilience 90 (Corsolini, 2009; Corsolini et al. 2017). Antarctic biota contamination studies have often focused on 91 the most central species (E. superba, P. antarctica and penguins) of the pelagic food webs (Corsolini 92 et al. 2002; Corsolini et al. 2017). Nevertheless, Southern Ocean benthic organisms are highly 93 abundant, diverse, and able to bioaccumulate contaminants they are yet poorly represented in 94 contamination studies (Di Giglio et al. 2020; Brasier et al. 2021; De Castro et al. 2021). 95 Consequently, knowledge about pollutants accumulation among them is still scarce. This could be due to the complex logistic sampling in remote areas (e.g. scuba diving, trawling) and the 96 97 concentrations of POPs being usually lower than those in other regions (Krasnobaev et al. 2020). 98 Bates et al. (2017) found that HCB could be remobilised from benthic biota with increasing 99 temperatures, therefore, especially under global climate change, benthic communities deserve more 100 attention as they can represent potential secondary sources of legacy pollutants. In this context, there 101 is an urgent need to identify suitable benthic bioindicator species for environmental pollution 102 monitoring in polar regions.

103 Among benthic organisms, the sponges (Phylum Porifera) represent a predominant component in the 104 Antarctic benthos both in terms of abundance and biomass (Kersken et al. 2016), and are an important 105 feeding source for many species such as sea stars, sea urchins, and nudibranch molluscs, thus playing 106 a key role in the dynamics of the community (Dayton et al. 1974; Garcia et al. 1993; Iken et al. 2002; 107 McClintock 1987, 2005; Cardona et al. 2021). They are suspension-feeders, able to filter thousands 108 of litres of water per day (Vogel, 1977; Negri et al. 2006), with an excellent retention capacity 109 allowing them to capture particles in a range of  $0.2 - 50 \,\mu\text{m}$ , with a lower limit far less than most of 110 the other filter-feeders (Perez et al. 2004; Batista et al. 2013). Therefore, they may potentially

accumulate large amounts of organic pollutants both in dissolved and suspended phases (Perez et al.2004).

113 While sponges possess many traits of good bioindicators such as abundance, long lifespan (from years 114 to millennia, Dayton, 1989; Gatti, 2002), large size (up to meters, Moran & Woods, 2012; van Soest 115 et al. 2012), and efficient filtration capability (Rainbow, 1995), these organisms are less used than 116 other filter-feeders as sentinels in biomonitoring programs (Genta-Jouve et al. 2012). This is probably 117 due to their complex taxonomic identification compared to other common indicator species (Hooper 118 & van Soest, 2002). However, some authors have already pointed out their potential and usefulness 119 as indicators for trace elements and heavy metals (Perez et al. 2003; Negri et al. 2006; Batista et al. 120 2014; Gentric et al. 2016), POPs (Perez et al. 2004; Negri et al. 2006; Gentric et al. 2016), and polycyclic aromatic hydrocarbons (Negri et al. 2006; Batista et al. 2013; Gentric et al. 2016). 121 122 However, among the few studies currently available on POPs in Antarctic benthic organisms (e.g., 123 Corsolini et al. 2003; Borghesi et al. 2011; Goutte et al. 2013; Grotti et al. 2016; Krasnobaev et al. 124 2020), none includes sponges.

125 The main objective of this study was therefore to assess the bioaccumulation of nineteen congeners 126 of PCBs (including twelve dioxin-like congeners), the p,p'- and o,p'- isomers of DDT, and its main 127 metabolites, DDD and DDE, as well as HCB (one of the POPs with the greatest atmospheric long-128 range transport potential), in several species of Antarctic sponges collected in the Ross Sea between 129 2001 and 2005 (Adèlie Cove; Tethys Bay) and in the Bransfield Strait in 2017 (Whalers Bay, 130 Deception Island). Secondary objectives were: 1) to evaluate inter-specific differences in the 131 accumulation patterns and to compare individuals belonging to the same species collected in three distinct sites; 2) to compare pollutant levels in three differently impacted areas. We expected: i) low 132 133 levels of POPs in such organisms due to their low lipid content together with their trophic level, even 134 with their filtration capability; ii) to find differences in the species-specific pattern due to the 135 biological variability; iii) Whalers Bay to show higher concentrations than Ross Sea sites due to its 136 closer geographical position to South America and the number of local sources (increasing tourism

- and cruise ships in Deception Island, scientific stations, and its industrial past) that may affect POPsrelease.
- 139

## 140 Materials and methods

- 141
- 142 Study area and sponge species

143 Sponge samples were collected at Whalers Bay (Lat. 62°59'0" S, Long. 60°34'0" W, Port Foster, 144 Deception Island) in the South Shetland Islands archipelago (Bransfield Strait), and at Adèlie Cove 145 (Lat. 74°45′51″ S, Long. 164°0′34″ E, Terra Nova Bay) and Tethys Bay (Lat. 74°40′60″ S, Long. 146 164°4'0" E, Terra Nova Bay) in the Ross Sea. Sampling areas are showed in Figure 1. Whalers Bay is a sandy beach located on Deception Island, an active volcano with a safe natural harbour, that was 147 148 used by sealers as the first centre of their hunting activities during the 19th century (de Ferro et al. 149 2013). Nearly a century later, it was the most extensive docking station for whale processing factories 150 ships and housed the Hektor whaling station; the only land based commercial activity in Antarctic 151 history (Dibbern, 2010). Nowadays, Whalers Bay is one of the most frequently visited locations in 152 Antarctica by tourists (Dibbern, 2010; de Ferro et al. 2013) with >15,000 visitors per year (IAATO, 153 International Association of Antarctica Tour Operators, 2018). Whalers Bay also hosts a well-154 developed rocky area in the southernmost part, where a rich filter-feeder community is found 155 (Angulo-Preckler et al. 2018). Moreover, the South Shetland Islands archipelago presents one of the 156 highest concentrations of scientific stations in the world (Barnes et al. 2008) and Deception Island 157 hosts two summer scientific stations, one from Argentina and one from Spain (Roura, 2012; de Ferro et al. 2013). The Western Antarctic Peninsula, where Whalers Bay is located, also represents one of 158 159 the most impacted areas by industrial fishing (Aronson et al. 2011) that is also increasing in the 160 Southern Ocean (Chown et al. 2015).

Tethys Bay is a small inlet nearby the Italian Mario Zucchelli Station (MZS); here the sea bottom is
covered by littoral sediments that consists of coarse sands, pebbles, and gravel (Cerrano et al. 2009).

Adèlie Cove is a 70-m depth V-shaped bay along the coast of Terra Nova Bay (Povero et al. 2001), 163 164 with a bottom characterised by fine sediments rich in organic matter due to the presence of a breeding 165 colony of Adèlie penguins (Cattaneo-Vietti et al. 2000). The bay is separated from the open sea by a 12-15 m depth sill that represent a natural barrier to the in- and out-flows (Cattaneo-Vietti et al. 2000). 166 167 Outside of that sill the bottom becomes coarser and consists of large pebbles (Povero et al. 2001), 168 where benthic communities dominate and sponges show high diversity and biomass (Cattaneo-Vietti 169 et al. 2000). Adèlie Cove is located South of the Italian base and far from any other anthropogenic 170 contamination source.

A total of 35 sponge specimens were collected in the Ross Sea (n = 25) and at Deception Island (n = 25)171 172 10) (Table S1). The Ross Sea samples were collected during the austral summers 2001/2002 at Tethys Bay and 2004/2005 at Adèlie Cove, in the framework of the XVII and the XX Italian Expedition of 173 174 National Research Program in Antarctica (PNRA), respectively. The sampling was conducted along 175 longitudinal transects (at Adèlie Cove it was conducted outside of the described sill) at a depth of 60-120 m by bottom trawls; samples were then stored in polyethylene bags. The Deception Island 176 177 samples were collected by scuba diving at 15-20 m depth during the DISTANTCOM-2 Antarctic cruise in February 2017, wrapped individually in aluminium foils and stored in polypropylene bags. 178 179 All the samples were stored at -20°C until laboratory analyses.

180 All samples were identified at species-level (Table S1). The sponge species belong to two main 181 classes: Hexactinellida and Demospongiae, most of them belonging to the second group (Table S1). 182 Four species were found in both the Ross Sea sites (Table S1). Sponge samples were processed by 183 standard methods (Rützler, 1978). Skeletal architecture was examined by light microscope. Hand-cut 184 sections of the ectosome and choanosome were made following Hooper (2000). Taxonomic 185 identifications were made using the Systema Porifera (Hooper & van Soest, 2002), the revision of 186 Porifera classification of Morrow & Cárdenas (2015), and the World Porifera Database (WPD) (de 187 Voogd et al. 2022).

188

## 189 Chemicals and residue analysis

190 Samples were analysed for 19 PCB congeners including the IUPAC numbers 28, 52, 101, 138, 153,

191 180, 194, and the dioxin-like IUPAC numbers 77, 81, 105, 114, 118, 123, 126, 156, 157, 167, 169,

192 189; HCB; the *o*,*p* <sup>'</sup> and *p*,*p* <sup>'</sup> isomers of DDT, DDE, and DDD.

193 Acetone, hexane (Scharlau, Sentmenat Spain), and dichloromethane (Honeywell Riedel-de-Haën<sup>TM</sup>) 194 purity grade >99.9% were used for glassware washing. During the sample preparation Acetone 195 Pestinorm<sup>®</sup> supplied by VWR Chemicals (Leuven, Belgium) and *n*-hexane ultra resi-analyzed<sup>®</sup> 196 supplied by J.T.Baker® (Gliwice, Poland) were used. Labelled compounds solutions were prepared 197 with *n*-nonane Picograde® LGC Standards (Wesel, Germany). Sodium sulfate anhydrous (mesh 12-198 60) Ultra resi-analyzed® and silica gel (mesh 70-230) for column chromatography were supplied 199 respectively by J.T.Baker® (Center Valley, PA, U.S.A) and Merck (Darmstadt, Germany). Labelled 200 standard solutions were purchased by Cambridge Isotope Laboratories Inc. (Andover, USA).

201 Firstly, samples were lyophilised at – 80 °C and 0.2 mbar for 48 h with a Cryodos, Telstar Industrial, 202 S.L. (Terrassa, Spain) and weighed to calculate the water content. Then, they were manually grounded 203 with a ceramic mortar and pools of organisms were prepared when the amount was too low (number 204 of pooled individuals is shown in Table S1); therefore, the total number of samples analysed was 23. 205 Sample weight was about 5 g (3.50 - 5.03 g) (Table S1). Before extraction, procedural blanks and samples were spiked with a known amount of a solution containing the following <sup>13</sup>C-labelled 206 207 compounds: PCB-28, -52, -101, -138, -153, -180, -209, *p*,*p*'-DDE, *o*,*p*'-DDT, *p*,*p*'-DDT, and HCB. 208 The extraction of the analytes was carried out by matrix solid-phase dispersion and the clean-up using 209 multi-layer silica gel columns as previously described in Roscales et al. (2016b). Samples were 210 transferred into vials and concentrated under a gentle nitrogen stream, then reconstituted with 20 µL of injection standard containing <sup>13</sup>C<sub>12</sub>-PCB-111, -170, -178. The lipid content was determined 211 212 gravimetrically using 0.5 g of each sample and following the same procedure used for the analytes 213 extraction. The extract was rotary evaporated to nearly dryness and then dried at 80 °C until steady 214 weight.

Target compounds were identified and quantified by gas chromatography coupled with low resolution 215 216 mass spectrometry (GC-LRMS) following Roscales et al. (2016a). The analyses were performed 217 using an Agilent 7890A gas chromatograph coupled with an Agilent 5975C mass spectrometer 218 (Agilent, Palo Alto, CA, USA) in selected ion monitoring (SIM) mode with electron ionization (EI) 219 at an electron voltage of 70 eV. The injector temperature was 250 °C and the injected volume was 1 220 µL in splitless mode, the carrier gas was He (0.8 mL/min constant flux at a pressure of 17.9 psi). The 221 GC was equipped with a BPX5 low bleed (SGE Analytical Science) capillary column ( $60 \text{ m} \times 0.5$ 222 mm i.d.  $\times$  0.25 µm film thickness). Oven temperature program started at 120°C, held for 2 min, 223 increased to 250°C at 35°C/min and held for 30 min, and finally ramped to 310 °C at 15°C/min and 224 held for 30 min. The transfer line was set at a temperature of 280°C, the source at 230°C, and the quadrupole at 150°C. The identification was based on the detection at the corresponding retention 225 226 time of at least two m/z ions. The relative abundance of the monitored ions was respected. Native 227 compounds quantification was based on the construction of a linear seven-point calibration curve (1  $-200 \text{ pg/}\mu\text{L}$ ) using the isotopic dilution technique. 228

229

## 230 Quality assurance/quality control (QA/QC)

231 Results were presented on lipid weight (lw) basis because a significant positive correlation 232 (Spearman's correlation test, p < 0.05) was found between the lipid content and the dry weight-based 233 analyte concentrations (PCB: p=0.0129, r=0.5100; HCB: p=0.0003, r=0.6995; DDT: p=0.0311, r=234 0.4604). Dry, lipid and wet-weight based concentrations are reported in SI (Tables S2-S7). Analytes 235 were identified according to: i) retention times of the selected m/z ions within  $\pm 0.1$  min of those 236 found in standard compounds; ii) variations in the relative abundances of the targeted ions  $\leq 10$  % of 237 the mean values obtained for the calibration standards. Recoveries of labelled compounds were 238 satisfactory in all cases (mean  $\pm$  standard deviation): 93  $\pm$  3 % for HCB, 89  $\pm$  13 % for the PCB 239 congeners nos. 28, 52, 101, 153, 138, 180, 209 ( $\Sigma_7$ PCBs) and 101 ± 13 % for the *o*,*p*'-DDT, *p*,*p*'-240 DDT and  $p_{,p}$ '-DDE ( $\Sigma_3$ DDTs) (Table S8); correspondence between labelled and native compounds

for identification and quantification is included in Table S9. One procedural blank was analysed with 241 242 each batch, which consisted of 4 or 5 samples to check for laboratory interferences. Limit of detection 243 (LOD) and limit of quantification (LOQ) were calculated with the signal to noise (s/n) ratio approach 244 and defined as 3 and 10 times the s/n value, respectively. The average LOD values were in the range 245 0.8 - 6.0 ng/g lw for PCBs, between 3.3 and 4.6 ng/g lw for DDTs, and 0.8 ng/g lw for HCB. The 246 LOQs averaged values ranged between 2.8 - 19.9 ng/g lw for PCBs, 11.1 - 15.2 ng/g lw for DDTs, 247 and 2.5 ng/g lw for HCB. See the SI for detailed LOD and LOQ values and detection frequencies 248 (Table S10, S11).

249

## 250 Data analysis

Statistical analyses were performed with Excel 2016 (Microsoft®), GraphPad Prism 5.01 (GraphPad 251 252 Software), and XLStat 2016 (Addinsoft©). Values below LOD were substituted with ½LOD. 253 Compounds below LOD in all samples were excluded from statistical comparisons and total 254 concentration calculations. Concentrations of the analytes were corrected subtracting the 255 corresponding procedural blank mean value. No corrections were applied according to recovery 256 measures since the isotopic dilution technique was used for quantification. Data distribution was 257 evaluated with Shapiro-Wilk test and was not normal even after a log<sub>n</sub> transformation. Thus, 258 concentration variations among samples collected in different sites were evaluated through the non-259 parametric Kruskal-Wallis test (significance level: p < 0.05). The comparison of homologue patterns 260 among sites and the evaluation of the contribution of PCB congeners or pollutants classes among 261 species, were based on descriptive statistics.

262

263

264 **Results and discussion** 

265

The lipid contents in the studied sample composites ranged from 0.7 to 8.5% (Table S1), in agreement 266 267 with the values reported by McClintock (1987) and Batista et al. (2013) for different sponge species 268 collected at McMurdo Sound (Antarctica) and along the Brazilian coast, respectively. The significant 269 positive correlation described above, between the lipid contents and the contaminants, suggests that 270 they are an important factor for the bioaccumulation. The water content of the samples ranged from 271 36 to 88% (Table S1). For both lipid and water content, the range observed may reflect a species-272 specific variability. All pollutant families showed detectable concentrations in all the samples with a 273 common concentration pattern:  $\Sigma PCB > \Sigma DDT > HCB$  in the three study sites (Table 1). Moreover, 274 the pattern is confirmed in every individual (including the only Hexactinellid specimen) except for 275 Neopetrosia similis (HCB >  $\Sigma$ PCB >  $\Sigma$ DDT) and one sample of Dendrilla antarctica ( $\Sigma$ DDT >  $\sum$ PCB > HCB) (Figure 2). In fact, it is interesting to note that the species *N. similis* collected at Adèlie 276 277 Cove showed the highest HCB percentage, exceeding the 60%; this pattern might be due to individual 278 variability. For this reason, this outlier value was not included in the statistical calculations. 279 Noteworthy, **SDDT** percentage in the genus *Dendrilla* ranged from 30 to 50% while in the other 280 genus it was between 4 and 30% (Figure 2), regardless of the sampling site, perhaps suggesting a 281 peculiar inability to degrade the pesticide. Thus, concerning the accumulation pattern, species-282 specific variability was relevant in less cases than expected. No other relevant differences can be 283 observed about the pattern in individuals of the same species collected in different site.

284

285 PCBs

Among PCBs, 11 out of 19 were detected at least in one sample (Table 1) including the seven indicator PCBs -28, -52, -101, -118, -138, -153, -180 and, among the coplanar dioxin-like congeners (other than -118) three mono-ortho -105, -123, -167 and the non-ortho -126, one of the most toxic congeners together with the -169 and -77 that resulted <LOD in all the samples. Detection frequencies of indicator PCBs were always above 90% and in 5 out of 7 cases (only excluding PCB-28 and PCB- 180) reached the 100% confirming the ubiquity of these POPs in the environment (Montone et al.2003) (Table 1).

293

294 Sponges had similar PCB concentrations to those reported (for most sampling years) by Grotti et al. 295 (2016) in the mollusc Adamussium colbecki (E. A. Smith, 1902) (Table S12), collected near Mario 296 Zucchelli station, likely due to its filter-feeding habits. In contrast, sponges showed lower levels of 297 PCBs (one order of magnitude) than the sea star Odontaster validus (Koehler, 1906) and the sea 298 urchin Sterechinus neumayeri (Meissner, 1900), previously collected near Mario Zucchelli station 299 (Borghesi et al. 2011) and near Durmont D'Urville French station (Goutte et al. 2013) (Table S12), as expected by their different dietary habits (Corsolini et al. 2003a), being relevant predators in 300 301 Antarctic ecosystems (Dayton et al. 1974). Ko et al. (2018) reported concentrations two to three 302 orders of magnitude higher than this study in the brittle star Ophionotus victoriae (Bell, 1902) and S. 303 neumayeri from Chinese (Chun-Shan) and Australian (Davis) stations (Table S12), this is probably 304 due to their proximity to that permanent research stations. Our findings also confirm that Antarctica 305 is one of the least contaminated regions on Earth as the PCB levels found here were three to four 306 orders of magnitude lower compared to Mediterranean sponge specimens (Perez et al. 2003) (Table 307 S12). In spite of the major role of anthropogenic activities in Whalers Bay and its closeness to the 308 American continent compared to the rest of sites, spatial differences were not statistically significant 309 (p=0.2829). The Ross Sea sites showed the highest levels of PCBs (Table 1). In contrast, the only 310 local input of PCBs at Adélie Cove could be the presence of a large Adèlie penguin rookery in the 311 cove. Wildlife may have a role in the POP redistribution and local amplification, as already reported in Polar Regions (Evenset et al. 2007; Roosens et al. 2007). In fact, penguins, being intermediate 312 313 predators, could accumulate lipophilic pollutants through biomagnification and release them in the 314 surrounding environment by excreta, abandoned or unhatched eggs, and carcasses (Roosens et al. 315 2007; Cipro et al. 2019; Corsolini et al. 2019; Morales et al. 2022). Concerning Tethys Bay samples, 316 the observed values could be related to the presence of local inputs of PCBs from the near research

station (Cabrerizo et al. 2012; Chen et al. 2015; Vecchiato et al. 2015). The absence of differences among the sites, nevertheless their different characteristics, could be due to a regional scale redistribution of the pollutants, due to both oceanic and atmospheric transport, making them more available than expected for the bioaccumulation in Ross Sea sponges. However, this result has to be evaluated carefully taking into account that it could be affected by other factors, such as the different number of samples, species-specific differences and temporal differences in the sampling time.

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325 The abundance of the PCB homologues was similar in the studied samples. The PCB homologue 326 pattern was penta- > hexa- > tetra- > tri- > hepta-CBs in samples from Adèlie Cove and Whalers Bay, 327 and penta- > hexa- > tetra- = hepta- > tri-CBs for those from Tethys Bay (Figure 3). In the samples 328 from the Tethys Bay, the presence of high-chlorinated and less volatile congeners like the hexa-CBs 329 (-138, -153, -167) and hepta-CB (-180), accounting for more than 40% of the total residue (Figure 330 S1), might confirm a local contamination source from near scientific stations (Chen et al. 2015; 331 Vecchiato et al. 2015). However, the lower chlorinated congeners nos. 28, 52 and 101 made up about 332 40% of the residue (Figure S1), also confirming a contribution by the LRAT. Corsolini et al. (2002, 333 2003b) reported a similar pattern to that observed for *E. superba* and *P. antarcticum* collected in the 334 same area and highlighted its similarity to the Kanechlor technical mixtures (KC-500 and -1000) 335 profile, used in Asian countries, perhaps suggesting a long-range transportation from those areas. 336 The Adèlie Cove and Whalers Bay samples showed a high presence of low-chlorinated PCBs: -28, -

52, -101, -105, -118, -123, -126, accounting for more than 70% and 80%, respectively (Figure S1). It
is interesting to note that PCB-101 is the most abundant congener in the Whalers Bay samples, far
exceeding the 60% of the total residue. Consistent with our results, the PCB-101 shows a higher
bioaccumulation potential (Log Kow 6.19; Ballschmiter et al. 2005) respect to other prevailing
congeners like PCB-28 (Log Kow 5.58; Ballschmiter et al. 2005) and PCB-52 (Log Kow 5.91;
Ballschmiter et al. 2005) and it has been already reported as one of the dominant congeners in

Antarctic air (Montone et al. 2003) as well as in penguins (Corsolini et al. 2007). Moreover, the 343 344 overall abundance of penta- and hexa-chlorinated congeners was already reported in some lower 345 trophic level organisms, such as molluscs A. colbecki (Grotti et al. 2016), sea cucumbers 346 (Heterocucumis steineni Ludwig, 1898), ascidians (Cnemidocarpa verrucosa Lesson, 1830), sea stars 347 (O. validus), limpets (Nacella concinna Strebel, 1908) and sea urchins (S. neumayeri) (Krasnobaev 348 et al. 2020). Our results also agree well with a previous study by Goutte et al. (2013), reporting the 349 predominance of penta- over hexa-CBs in Antarctic benthic species such as the starfish Saliasterias 350 brachiata Koehler, 1920 and the sea urchin S. neumayeri.

351 Comparing the individuals belonging to the four species collected at both the Ross Sea sites (Figure 352 4), the general observed pattern was mostly confirmed; samples from Tethys Bay showed higher 353 percentage of the heaviest congeners compared to Adèlie Cove specimens in three out of four cases. 354 However it is interesting to note that Artemisina tubulosa showed a slightly inverted pattern with the 355 percentages for Tethys Bay moved towards lighter congeners than Adèlie Cove; this could be due to a species-specific ability to transform and excrete selected congeners by the sponge itself or its 356 357 associated microorganisms, like hypothesized for some PCBs in Spongia officinalis (Perez et al. 358 2003).

359

Five DDT isomers were >LOD in 30% of samples; the o,p '-DDD isomer was <LOD in all samples (Table 1). DDTs were mostly undetectable in the samples from the Ross Sea, showing 73% and 77% of the values <LOD in Tethys Bay and Adèlie Cove samples (excluding the o,p '-DDD isomer), respectively (Table 1). The DDT isomer concentrations were not reported in one sample from Tethys Bay due to a co-eluting unknown compound that made the quantification uncertain. Instead, samples from Deception Island showed values <LOD in 20% of the cases (Table 1).

367

<sup>360</sup> *DDTs* 

Samples collected at Whalers Bay and Tethys Bay showed concentration values lower than those of 368 369 O. validus and higher than those of S. neumayeri reported previously by Borghesi et al. (2011) (Table 370 S12). However, samples from Adèlie Cove showed values on the same order of magnitude than those 371 detected in the sea urchin from the same study (Table S12). Focusing on the p,p'-DDE isomer, 372 sponges from Whalers and Tethys Bays showed similar values to O. validus and higher than A. 373 colbecki and S. neumayeri as reported by Corsolini et al. (2003a) (Table S12). Adèlie Cove samples 374 showed concentrations lower than the sea stars and similar to molluscs and sea urchins from the same 375 study (Table S12).

As discussed for PCBs, dietary differences may explain these results when comparing them to those from the literature (e.g. lower concentrations in sponges than in predators like *O. validus*) (Corsolini et al. 2003a). However, more studies are needed to interpret differences among species.

379

380 Differences in DDT concentrations among sites were not statistically significant (p=0.1575), although 381 they were higher in sponges from Whalers Bay (Table 1). On one hand, concentrations found in 382 Whalers Bay could be influenced by its proximity to South America, where this pesticide has been 383 used along history (Montone et al. 2003; Dickhut et al. 2005; Corsolini et al. 2007) and from which 384 it could be transported via LRAT to Antarctica (Dickhut et al. 2003; Montone et al. 2005). On the 385 other hand, local inputs, such as the penguin rookery near Adèlie Cove, could contribute to increasing 386 concentrations in this site. Inputs from these sources at each site may have flattened the expected 387 differences among the two areas. Moreover, the frequency of values <LOD in the two areas seems to 388 be in line with the expected results being higher in the Ross Sea than at Whalers Bay (70% and 20% 389 respectively). An explanation for these apparently contrasting results could be found in the species-390 specific characteristics; noteworthy, in fact, the only Ross Sea samples in which DDTs were found 391 were of the same genus of the Whalers Bay samples (Dendrilla). However, again, other factors such 392 as the different number of samples analysed, and the year of sampling have to be taken into account.

The p, p'-DDE showed the highest values in all samples from each site, followed by its precursor p, p'-393 394 DDT; thus the ratio p,p'-DDT/p,p'-DDE was <1 (Figure S2), indicating an old contamination event 395 (Ricking & Schwarzbauer, 2012). Nonetheless, the detection of p,p'-DDT in all samples from 396 Deception Island and in four samples from the Ross Sea could be related to the current use of this 397 pesticide against the mosquitoes Anopheles (Stockholm Convention, 2004; Pozo et al. 2017; Zanardi-398 Lamardo et al. 2019), vector of the malaria disease, as well as other current applications like 399 antifouling paints (Pozo et al. 2017; Zanardi-Lamardo et al. 2019) and the following LRAT from 400 those countries where it is applied notwithstanding the Stockholm Convention. Geisz et al. (2008) 401 also suggested the melting glaciers as a possible secondary mechanism for DDTs to enter the marine 402 Antarctic ecosystem. Since the Antarctic Peninsula is suffering the highest warming events due to 403 climate change (Turner et al. 2005), this mechanism could also support Whalers Bay sponges 404 presenting higher frequencies of detection of DDTs than the Ross Sea samples. An uncompleted 405 degradation of DDTs by sponges or by their symbiotic bacteria associations may be another reason 406 of its detection. For example, Krasnobaev et al. (2020) reported concentrations <LOD for p,p'-DDT 407 in some benthic invertebrates (sea cucumbers, ascidians, sea stars, limpets, and sea urchins) collected 408 in 2017 (the same year we collected our Whalers Bay samples), near Rothera Point (Western Antarctic 409 Peninsula), suggesting a complete transformation of DDTs into p,p'-DDE instead of a lack of the still 410 debated recent input (Van den Brink et al. 2009). Further studies are needed to clarify if our results 411 were determined mostly by the scarce degradation capability of sponges following an old 412 contamination event or by a new LRT event due to its continued use in countries where DDT is still 413 crucial to control malaria.

414

415 HCB

416 HCB values were <LOD only in one sample collected at Adèlie Cove (Table 1), confirming its global</li>
417 distribution, persistence, and wide past usage (Bailey, 2001; Wang et al. 2010).

The HCB concentrations were lower than those previously reported in the seastar O. validus and 418 419 higher than in the sea urchin S. neumayeri from Antarctica (Borghesi et al. 2011) (Table S12). 420 However, our values were lower than in the sea star and sea urchin and of the same order of magnitude 421 of those reported for the bivalve A. colbecki in a previous study (Corsolini et al. 2003a) (Table S12). 422 Again, these contrasting results suggest that not only different dietary habits, but also metabolism, 423 season of sampling, and environmental concentrations could play a key role in determining these 424 interspecific variabilities. In addition, these comparison results, being not consistent in terms of prey-425 predator patterns, did not allow further considerations on biomagnification processes as expected for a benthic trophic web (Evenset et al. 2016; Romero-Romero et al. 2017). 426

427 HCB concentrations were of the same order of magnitude in all samples with no significant 428 differences among sites (p=0.2719) except for some samples from Adèlie Cove and Whalers Bay, 429 which showed concentrations one order of magnitude higher (Table 1). The lack of significant spatial 430 variations could be related to the physical-chemical properties of the pesticide: its vapour pressure combined with water solubility and persistence, in fact, make it widespread globally (Bailey, 2001). 431 432 Furthermore, other factors could contribute to the result; for example, in Whalers Bay, changes in the 433 frequency of snowfalls may locally amplify the HCB concentration, as suggested by Krasnobaev et 434 al. (2020), and the same may happen by biological transportation in Adèlie Cove.

435

436 Several studies have found that among legacy POPs, HCB predominates in the Antarctic atmosphere, 437 mainly due to the wide use, high volatility, and persistence of this chemical (Cincinelli et al. 2009; 438 Kallenborn et al. 2013; Bengtson Nash et al. 2017). Some studies have shown that this pattern 439 sometimes is also reflected in wildlife, being HCB the most abundant compound in various marine 440 organisms, such as fish and krill (Corsolini, 2009; Corsolini & Sarà, 2017). Particularly, Corsolini et 441 al. (2003a) and Krasnobaev et al. (2020) found HCB concentrations above those of DDTs in some 442 marine invertebrate species collected in 1999/2000 in the Ross Sea and in 2017 in the Western 443 Antarctic Peninsula. In our study, HCB was the less abundant pollutant in the three study sites

444 (ΣPCBs>ΣDDTs>HCB). The peculiarity of sponges in terms of feeding habits, biodegradation
445 capability, and longevity may be responsible of these diverse POP bioaccumulation profiles and
446 deserves further efforts to better understand trophodynamic, transportation, and fate of these
447 pollutants.

448

## 449 **Conclusions**

450

To the best of our knowledge, no published data are available on the presence of HCB, DDTs and 451 452 PCBs in Antarctic Porifera. Sponges showed legacy POP levels comparable to other benthic 453 organisms from the same habitat and, as expected, much lower than sponge from northern temperate latitudes, confirming the Southern Ocean as one of the less contaminated ecosystems on Earth. The 454 455 samples from the Ross Sea showed, in general, lower concentrations respect to the South Shetland 456 Island samples, although differences were not statistically significant. In general, long-range 457 atmospheric transport was confirmed as the major driver for contamination in the Antarctic areas 458 where the study was performed. However, human presence and activities connected with research 459 stations, as well as wildlife amplification and ice melting could also affect the bioaccumulation 460 pattern found in these sponges. Future studies should also focus on increasing threats like tourism 461 activities and fishing to better understand how and to which extent they could act synergically with 462 other impacts in affecting the Antarctic ecosystems. While evaluation of species-specific patterns 463 showed a few interesting results (peculiar patterns observed in the genus Dendrilla and in the N. 464 similis individuals), further research is needed to clarify which mechanisms are involved in 465 determining the observed inter- and intraspecific differences. Our results indicate that sponges may 466 be suitable bioindicators for the benthic marine habitat. Moreover, they provide baseline data for 467 future monitoring and contamination trend studies that, in the light of climate change, may well 468 represent valid tools to understand and make predictions on the threats Antarctica has to cope with.

469

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471

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774 **Table 1**. Concentrations of HCB, PCB congeners and DDT isomers in the sponge samples from the 775 three study sites (n=number of samples; ng/g lipid weight; mean  $\pm$  standard deviation, minimum and 776 maximum values in brackets) and values < LOD (LOD %).

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	<b>Tethys Bay</b>	Adèlie Cove	Whalers Bay	<lod< th=""></lod<>
	( <b>n=7</b> )	(n=13)	(n=3)	%
PCB-28	$7.5\pm6.7$	$14.9 \pm 14.9$	$3.5\pm1.9$	9
	( <lod -="" 21.2)<="" td=""><td>(<lod -="" 52.4)<="" td=""><td>(1.9 - 5.6)</td><td></td></lod></td></lod>	( <lod -="" 52.4)<="" td=""><td>(1.9 - 5.6)</td><td></td></lod>	(1.9 - 5.6)	
<b>PCB-52</b>	$17.5\pm18.9$	$21.1 \pm 18.9$	$6.0 \pm 1.9$	4
	( <lod -="" 43.5)<="" td=""><td>(4.4 - 68.6)</td><td>(4.9 - 8.2)</td><td></td></lod>	(4.4 - 68.6)	(4.9 - 8.2)	
PCB-101	$12.5\pm12.4$	$30.3\pm51.1$	$21.5\pm10.0$	4
	( <lod -="" 36.1)<="" td=""><td>(0.3 – 196.2)</td><td>(10.5 – 30.1)</td><td></td></lod>	(0.3 – 196.2)	(10.5 – 30.1)	
PCB-105	$3.2 \pm 1.4$	$6.4 \pm 4.0$	$4.1 \pm 0.3$	74
	( <lod 5.1)<="" td="" –=""><td>(<lod -="" 16.5)<="" td=""><td>(<lod 4.4)<="" td="" –=""><td></td></lod></td></lod></td></lod>	( <lod -="" 16.5)<="" td=""><td>(<lod 4.4)<="" td="" –=""><td></td></lod></td></lod>	( <lod 4.4)<="" td="" –=""><td></td></lod>	
PCB-118	$4.3 \pm 3.2$	$10.6 \pm 14.0$	$5.1\pm0.9$	0
	(0.7 - 9.8)	(0.8 – 52.1)	(4.1 - 5.8)	
PCB-123	$5.0 \pm 2.4$	$9.0 \pm 4.8$	$2.7\pm0.0$	91
	( <lod -="" 8.5)<="" td=""><td>(<lod 17.0)<="" td="" –=""><td>(<lod 2.7)<="" td="" –=""><td></td></lod></td></lod></td></lod>	( <lod 17.0)<="" td="" –=""><td>(<lod 2.7)<="" td="" –=""><td></td></lod></td></lod>	( <lod 2.7)<="" td="" –=""><td></td></lod>	
PCB-126	$3.0 \pm 1.7$	$6.1 \pm 3.2$	$1.7\pm0.0$	96
	( <lod 5.4)<="" td="" –=""><td>(<lod -="" 10.9)<="" td=""><td>(<lod 1.7)<="" td="" –=""><td></td></lod></td></lod></td></lod>	( <lod -="" 10.9)<="" td=""><td>(<lod 1.7)<="" td="" –=""><td></td></lod></td></lod>	( <lod 1.7)<="" td="" –=""><td></td></lod>	
PCB-138	$10.0 \pm 6.8$	$12.5 \pm 13.0$	$4.4 \pm 0.9$	0
	(2.6 – 21.6)	(1.5 – 46.3)	(3.6 - 5.5)	
PCB-153	$10.4 \pm 12.8$	$12.5 \pm 14.2$	$2.9\pm0.3$	0
	(1.5 – 38.5)	(1.5 – 41.9)	(2.7 – 3.2)	
PCB-167	$2.0 \pm 1.4$	$2.2 \pm 1.2$	$0.7\pm0.0$	87
	( <lod -="" 4.9)<="" td=""><td>(<lod -="" 4.2)<="" td=""><td>(<lod 0.7)<="" td="" –=""><td></td></lod></td></lod></td></lod>	( <lod -="" 4.2)<="" td=""><td>(<lod 0.7)<="" td="" –=""><td></td></lod></td></lod>	( <lod 0.7)<="" td="" –=""><td></td></lod>	
PCB-180	$17.5 \pm 30.8$	$8.2\pm8.1$	$1.4 \pm 1.4$	4
	( <lod -="" 86.2)<="" td=""><td>(0.5 - 26.0)</td><td>(0.4 - 3.0)</td><td></td></lod>	(0.5 - 26.0)	(0.4 - 3.0)	
∑PCB	$92.8\pm47.6$	$133.7 \pm 119.3$	$54.2\pm10.8$	
	(27.7 - 164.7)	(39.8 - 482.8)	(41.9 - 62.0)	
<i>o,p</i> '-DDT	$2.5 \pm 2.8$	$2.9 \pm 1.4$	$3.4 \pm 2.8$	78
	( <lod -="" 8.2)<="" td=""><td>(<lod 5.1)<="" td="" –=""><td>(1.5 – 6.6)</td><td></td></lod></td></lod>	( <lod 5.1)<="" td="" –=""><td>(1.5 – 6.6)</td><td></td></lod>	(1.5 – 6.6)	
<i>p,p</i> '-DDT	$4.8 \pm 6.2$	$5.1 \pm 3.6$	$11.0 \pm 2.1$	70

	(3.0 – 7.3)	( <lod -="" 19.6)<="" th=""><th>(5.7 – 13.1)</th><th></th></lod>	(5.7 – 13.1)	
НСВ	$\textbf{4.8} \pm \textbf{1.8}$	*8.1 ± 5.0	8.5 ± 4.0	4
	(5.6 – 53.7)	(9.3 – 59.2)	(31.3 – 46.1)	
∑DDT	$17.5 \pm 18.3$	$\textbf{27.2} \pm \textbf{17.1}$	$\textbf{38.6} \pm \textbf{7.4}$	
	( <lod 1.9)<="" th="" –=""><th>(<lod -="" 4.5)<="" th=""><th>(<lod -="" 0.7)<="" th=""><th></th></lod></th></lod></th></lod>	( <lod -="" 4.5)<="" th=""><th>(<lod -="" 0.7)<="" th=""><th></th></lod></th></lod>	( <lod -="" 0.7)<="" th=""><th></th></lod>	
<i>p,p</i> '-DDD	$1.3 \pm 0.6$	$2.4\pm1.2$	$0.7\pm0.0$	91
	( <lod -="" 24.5)<="" th=""><th>(<lod 38.3)<="" th="" –=""><th>(13.9 – 23.6)</th><th></th></lod></th></lod>	( <lod 38.3)<="" th="" –=""><th>(13.9 – 23.6)</th><th></th></lod>	(13.9 – 23.6)	
<i>p,p</i> '-DDE	$7.3\pm8.9$	$14.1\pm12.5$	$19.6\pm5.0$	30
	( <lod 2.2)<="" th="" –=""><th>(<lod -="" 5.2)<="" th=""><th>(1.8 - 5.3)</th><th></th></lod></th></lod>	( <lod -="" 5.2)<="" th=""><th>(1.8 - 5.3)</th><th></th></lod>	(1.8 - 5.3)	
<i>o,p</i> '-DDE	$1.5\pm0.6$	$2.7 \pm 1.4$	$3.9\pm1.8$	78
	( <lod 17.3)<="" th="" –=""><th>(<lod -="" 14.7)<="" th=""><th>(9.6 – 13.4)</th><th></th></lod></th></lod>	( <lod -="" 14.7)<="" th=""><th>(9.6 – 13.4)</th><th></th></lod>	(9.6 – 13.4)	

778	*mean calculated on 12 samples
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# 789 Figure legends

790	Figure 1: a) Antarctic continent with the indication of the two areas where the sampling site are
791	located; b) Deception Island area in the South Shetland Archipelago (n=10, year of sampling 2017);
792	c) coastal area of Victoria Land in the Ross Sea (Tethys Bay: n=7, year of sampling 2001-2002;
793	Adèlie Cove: n=18, year of sampling 2004-2005). Black stars show the sampling site. Red symbols
794	indicate summer-only stations or facilities (e.g., Enigma Lake and Browning Pass airstrips) and blue-
795	red symbols year-round stations. Blue dots indicate important bird areas.
796 797	Figure 2: Contributions (%) of PCBs, DDTs, and HCB in the 23 Antarctic sponge species from the three study sites (Whalers Bay, Tethys Bay, Adèlie Cove).
798	Figure 3: Homologue pattern (%) in sponges from the three study sites (Adèlie Cove; Tethys Bay;
799	Whalers Bay).
800	Figure 4: Percentage contribution of PCB congeners to the total residue (%) in eight Antarctic sponges
801	belonging to four different species and collected from Adèlie Cove (AC) and Tethys Bay (TB).
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Figure 4

## Highlights

- First data about legacy chlorinated POPs in Antarctic sponges are reported
- Antarctic sponges are suitable organisms for legacy POPs contamination studies
- Sponges showed levels of contamination comparable to other benthic organisms
- DDTs and HCB concentrations in sponges: South Shetland Island > Ross Sea

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## Author statement

Nicolas Pala: Conceptualization, Formal analysis, Investigation, Data curation, Methodology, Writing - original draft.

Begoña Jiménez: Conceptualization, Resources, Supervision, Funding Acquisition, Writing – review & editing.

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Blanca Figuerola: Conceptualization, Methodology, Investigation, Resources, Writing – review and editing.

Conxita Avila: Conceptualization, Methodology, Funding acquisition, Investigation, Resources, Writing – review and editing.

Simonetta Corsolini: Conceptualization, Resources, Supervision, Funding Acquisition, Writing – review & editing.

## **Declaration of interests**

☑ The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

□ The authors declare the following financial interests/personal relationships which may be considered as potential competing interests:

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