

## Distribution of transparent exopolymer particles (TEP) in distinct regions of the Southern Ocean

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2	distinct regions of the Southern Ocean
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Longitude

#### 25 Abstract

Transparent exopolymer particles (TEP) are an abundant class of suspended organic 26 particles, mainly formed by polysaccharides, which play important roles in 27 biogeochemical and ecological processes in the ocean. In this study we investigated 28 horizontal and vertical TEP distributions (within the euphotic layer, including the upper 29 30 surface) and their short-term variability along with a suite of environmental and 31 biological variables in four distinct regions of the Southern Ocean. TEP concentrations in the surface (4 m) averaged 102.3  $\pm$  40.4  $\mu$ g XG eq L<sup>-1</sup> and typically decreased with 32 depth. Chlorophyll a (Chl a) concentration was a better predictor of TEP variability 33 across the horizontal ( $R^2 = 0.66$ , p< 0.001) and vertical ( $R^2 = 0.74$ , p< 0.001) scales than 34 prokaryotic heterotrophic abundance and production. Incubation experiments further 35 confirmed the main role of phytoplankton as TEP producers. The highest surface TEP 36 concentrations were found north of the South Orkney Islands (144.4  $\pm$  21.7 µg XG eq L<sup>-</sup> 37 <sup>1</sup>), where the phytoplankton was dominated by cryptophytes and haptophytes; however, 38 the highest TEP:Chl a ratios were found south of these islands (153.4  $\pm$  29.8  $\mu$ g XG eq 39 (µg Chl a)<sup>-1</sup>, compared to a mean of 79.3 ± 54.9 µg XG eq (µg Chl a)<sup>-1</sup> in the whole 40 cruise, in association with haptophyte dominance, proximity of sea ice and high 41 exposure to solar radiation. TEP were generally enriched in the upper surface (10 cm) 42 respect to 4 m, despite a lack of biomass enrichment, suggesting either upward transport 43 by positive buoyancy or bubble scavenging, or higher production at the upper surface 44 by light stress or aggregation. TEP concentrations did not present any significant cyclic 45 diel pattern. Altogether, our results suggest that photobiological stress, sea ice melt and 46 turbulence add to phytoplankton productivity in driving TEP distribution across the 47 Antarctic Peninsula area and Atlantic sector of the Southern Ocean. 48

49 Keywords: Transparent exopolymer particles, phytoplankton, prokaryotes, solar
50 radiation dose, Southern Ocean

51

## 52 **1 Introduction**

53 Transparent exopolymer particles (TEP) are gel-like organic particles stainable with Alcian Blue, a specific dye for acidic polysaccharides (Alldredge et al., 1993), that have 54 55 deserved attention due to their influence in biogeochemical and ecological processes 56 (Passow, 2002b). TEP are partly formed by the abiotic self-assembly from dissolved precursors (Chin et al., 1998; Orellana and Verdugo, 2003), thus connecting the 57 dissolved to particulate organic matter continuum (Engel et al., 2004). In addition, TEP 58 affect the biological carbon pump, not only because TEP by themselves may comprise a 59 mean of 5-10 % of primary production synthates (Mari et al., 2017) that can sink into 60 61 the deep ocean, but also because they promote particle aggregation (Engel et al., 2004), 62 thus favouring the sinking of marine snow to the deep ocean (Burd and Jackson, 2009). TEP also influence air-sea gas exchanges like that of carbon dioxide (CO<sub>2</sub>) (Wurl et al., 63 64 2016; Jenkinson et al., 2018), since they tend to accumulate at the surface microlayer (SML, the uppermost water layer, Cunliffe et al. (2013)), either after ascending through 65 the water column (Azetsu-Scott and Passow, 2004) or upon direct production in this 66 layer (Wurl et al., 2011b). 67

Phytoplankton are believed to be the main source of TEP and their precursors
(Alldredge et al., 1993; Passow, 2002b; Van Oostende et al., 2012; Engel et al., 2015),
being diatoms (Passow et al., 1994; Mari and Burd, 1998; Passow et al., 2001; Passow,
2002a) and haptophytes, like *Phaeocystis* (Hong et al., 1997), the groups that seem to
release larger amounts. Prokaryotic heterotrophs (Stoderegger and Herndl, 1999;

Ortega-Retuerta et al., 2010) and other organisms, such as suspension feeders
(Heinonen et al., 2007), zooplankton (Prieto et al., 2001) and seagrass (Iuculano et al.,
2017a), can also produce TEP.

76 Sinks of TEP comprise photolysis by UV radiation (Ortega-Retuerta et al., 2009a), sinking to the deep ocean, entrance to the atmosphere (Kuznetsova et al., 2005), and 77 degradation and consumption by microorganisms (Ling and Alldredge, 2003). Many 78 79 variables influence either sources, sinks or both, hindering the prediction of TEP distribution in the ocean. For example, high solar radiation has a dual effect; it 80 stimulates TEP release by microbes (Iuculano et al., 2017b) but also causes TEP 81 photolysis (Ortega-Retuerta et al., 2009a) or affect positively (Shammi et al., 2017) or 82 83 negatively (Orellana and Verdugo, 2003) the abiotic self-assembly of dissolved exopolymers into TEP. Other variables such as temperature (Nicolaus et al., 1999; 84 Claquin et al., 2008), CO<sub>2</sub> concentration (Engel, 2002), nutrient availability (Mari et al., 85 86 2005) and viral infections (Nissimov et al., 2018) can also influence TEP cycling. Moreover, the TEP production per phytoplankton biomass increases along the bloom 87 stages (Pedrotti et al., 2010), thus adding complexity to the TEP dynamics. 88

The Southern Ocean (SO) is characterized by the presence of the Antarctic Circumpolar Current, which connects the waters of the Pacific, Indian and Atlantic Oceans, and by having strong upwellings that enrich surface waters with macronutrients (Sarmiento et al., 2004). Productivity is generally limited by the lack of iron in combination with deep surface mixing and low light and temperature (Moore et al., 2013; Hoppe et al., 2017), although some regions are locally supplied with iron, particularly in the vicinity of islands (Morris and Sanders, 2011).

The study of TEP distributions in the SO is of particular importance in light of these 96 ocean's peculiarities. The SO is considered the largest region for anthropogenic CO<sub>2</sub> 97 sequestration in the world (Frölicher et al., 2015), especially around the island systems 98 (Pollard 2009), through higher CO<sub>2</sub> solubility in cold waters and a relatively high 99 particulate organic carbon (POC) surface export flux in comparison to lower latitudes 100 101 (Boyd and Trull, 2007; Marinov et al., 2008). Indeed, C export fluxes have been reported to attain 30-50 % of net primary production (Buesseler, 2001). Since TEP may 102 103 account for an important fraction of POC mass (Engel, 2004; Zamanillo et al., 2019) and export flux (Passow, 2002b; Wurl et al., 2011a), their study in the SO is important 104 105 to better predict the magnitude of the biological carbon pump and the future dynamics 106 of atmospheric CO<sub>2</sub>.

In this study we describe the horizontal and vertical (within the euphotic layer) 107 108 distributions, and short-term (diel) variability of TEP, along with relevant physical, 109 chemical and biological variables, in four distinct regions of the SO. The four regions were characterized by distinct phytoplankton communities, bloom stages and 110 environmental and ecological properties. Our objectives were: (a) to identify the main 111 drivers of TEP distribution across contrasting environmental conditions, both 112 horizontally and vertically, and (b) to examine the short-term temporal variability of 113 114 TEP and biological variables, over diel cycles. Data co-variation analyses and experimental incubations were conducted to ascertain the role of microorganisms 115 116 (phytoplankton vs. heterotrophic prokaryotes) in TEP production.

117

#### 118 2. Material and methods

## 119 **2.1 Study site and sampling**

Sampling was performed during the PEGASO cruise, on board the Spanish R/V 120 Hespérides, conducted from 7<sup>th</sup> January to 3<sup>rd</sup> February, in the austral summer of 2015. 121 A total of 70 stations were sampled within the Southern Ocean (Fig. S1). Four regions 122 123 were occupied for several days, following a Lagrangian approach; the north of the South 124 Orkney Islands (NSO), the southeast of the South Orkney Islands (SSO), the northwest of South Georgia (NSG) and the west of Anvers Island (WA). NSO, NSG and WA were 125 selected due to their relatively high chlorophyll a (Chl a) concentrations, different 126 127 nutrient conditions and relatively slow currents with the absence of stable direction (Nunes et al., 2019). SSO was selected for its vicinity to the sea ice edge. In order to 128 129 track the studied water bodies, WOCE (World Ocean Circulation Experiment) standard drifters provided with Iridium communication system were used in the NSO, NSG and 130 WA regions; in SSO, two icebergs were used as natural Lagrangian drifters. 131

132 During the transit between the regions, seawater was collected from 4 m depth using the 133 ship's underway pump (BKMKC-10.11. Tecnium, Manresa, Spain), approximately at 134 9:00 and 15:00 local time (UTC-3). During the Lagrangian occupation of the main 135 stations, CTD casts were carried out using a SBE 911 Plus probe attached to a rosette of 24 12-L PVC Niskin bottles, at least once a day, generally at 9:00 and 15:00 local time 136 (UTC-3). The underway sampling was used when conditions were too rough for CTD 137 launch (a total of 3 times). A 36-h cycle was also sampled in each region, with CTD 138 casts every 4 h, generally from 9:30 to 17:00 (solar times) the day after. NASA Solar 139 Calculator (https://www.esrl.noaa.gov/gmd/grad/solcalc/, accessed on 15 December 140 141 2017) was used to perform solar time calculations (Nunes et al., 2019). The position of 142 the main hydrographic fronts was determined as described in Nunes et al. (2019).

143 TEP, major nutrients, POC, prokaryotic heterotrophic abundance (PHA) and production144 (PHP), primary production (PP), phytoplankton recount and identification by

microscopy and flow citometry, extracellular enzyme activities and virus concentrations 145 146 were analyzed in surface samples. At 7 stations (three in NSO, two in SSO, one in NSG and one in WA), Chl a, phytoplankton pigment analyses and PHA were carried out for 6 147 148 different depths, generally from 4 to 120-150 m, while TEP were analysed for 4 depths, from 4 m to the second deepest depth. In some of these stations, phytoplankton 149 identification and recount by microscopy and flow cytometry were done from 2 to 4 150 depths, including the depth of maximum fluorescence. Mixed layer depth (MLD) 151 152 estimation was done from CTD profiles following Monterey and Levitus (1997).

A total of 14 samples (one in the transit between regions, seven in NSO, four in SSO, 153 154 one in NSG and one in WA) were collected from the upper surface (10 cm) using a 155 zodiac boat, away and upstream from the ship to avoid contamination. Sampling was carried out with a device of our own design, consisting of a 50 mL glass syringe 156 connected to a PTFE tube 1.5 m long, whose other end was fixed onto a floating item so 157 158 that the inlet of the tube remained at a depth of 10 cm regardless of the motion of the 159 sea surface. All material was prewashed with HCl (10 %) and Milli-Q water. Typical 160 collected volume was 400 mL; usually, sampling took place around 10:00 local time (UTC-3) during standard stations and at 11:00 and 17:00 local time during intensive 161 day-night cycles. TEP, Chl a, PHA, PHP and extracellular enzyme activities were 162 163 analysed at the upper surface. In order to compare these measurements with those at the nominal surface (4 m depth of the nearest CTD casts), we calculated an enrichment 164 factor (EF<sub>us</sub>), defined as: 165

 $EFus = \frac{[X]upper surface}{[X]surface}$ 

where [X] is the concentration of a given variable, in the upper surface and surface respectively.  $EF_{us} > 1$  indicates enrichment of a component in the upper surface and 168  $EF_{us} < 1$  indicates depletion. It is worth mentioning that the time difference between the 169 upper surface sampling and the nearest surface sampling was 1-3 h. When the upper 170 surface sampling occurred exactly between two CTD casts, the average of the two was 171 used.

#### 172 **2.2 Physical measurements**

173 Surface temperature and salinity were obtained continuously with the SBE21 Sea Cat Thermosalinograph, while vertical profiles were recorded down to 400 m with a CTD 174 175 SBE911 plus. Average diffuse attenuation coefficient in the euphotic zone for PAR broadband (400-700 nm; K<sub>d</sub> (PAR)), from both CTD and PRR-800 vertical profiles of 176 downwelling spectral irradiance (E<sub>d</sub>), were determined as the slope of the linear 177 regression between logarithmic irradiances and depth. Prior to analysis, the data were 178 179 carefully examined for irregularities (Mueller et al., 2003). Near-surface noise caused by smooth waves and ripples was eliminated from  $E_d$  (PAR) profiles. The obtained  $R^2$ 180 of regression analyses were always above 0.98 at all evaluated casts. The depth of the 181 euphotic layer was then calculated as 4.605/K<sub>d</sub> (PAR), being the depth at which K<sub>d</sub> 182 (PAR) is reduced to 1 % of its value just below the surface. Solar radiation dose was 183 184 calculated in every station using the following formula:

185 Solar radiation dose=
$$\frac{I}{Kd (PAR) \times MLD} \times (1 - e^{(-Kd (PAR) \times MLD)})$$

where, I is the average surface intensity radiation (W m<sup>-2</sup>) in the 24 h previous to sampling,  $K_d$  (PAR) is the average diffuse attenuation coefficient in the euphotic zone for PAR broadband (m<sup>-1</sup>), and MLD is the mixed layer depth (m).

## 189 2.3 Chemical and biological analyses

190 **2.3.1 Particulate organic matter (TEP and POC)** 

TEP were analysed by spectrophotometry following the method proposed by Passow 191 and Alldredge (1995). Samples were filtered under low constant filtration pressure 192 (~150 mm Hg) in duplicate (200-550 mL) using 25 mm diameter polycarbonate filters 193 194 (DHI) with a pore size of 0.4 µm. All sampling material was pre-washed with HCl (10 %) and Milli-Q water. Filters were immediately stained with 500  $\mu$ L of Alcian Blue 195 solution (0.02 %, pH 2.5) for 5 s, and then rinsed with Milli-Q water. Two empty filters 196 (blanks) were taken at every station and stained in the same way. All filters remained 197 198 frozen until further processing some months later at the home laboratory. There, filters were soaked in 80 % sulfuric acid (5 mL) for 3 h and shaked intermittently. Absorbance 199 of the acid solutions was then measured at 787 nm with a Varian Cary 100 Bio 200 201 spectrophotometer. The absorbance of filter blanks, taken in duplicate for every batch of 202 samples, was subtracted from the absorbance of samples. Calibration of the Alcian Blue 203 solution was conducted immediately after the cruise using xanthan gum (XG) solution 204 as a standard. The detection limit was set to 0.045 absorbance units, and the mean range 205 between duplicates was 25.5 %. TEP carbon content (TEP-C) was estimated using the 206 conversion factor of 0.51 µg TEP-C per µg XG eq (Engel and Passow, 2001).

For POC analysis, we filtered water samples (1000 mL) through pre-combusted (4 h, 450 °C) GF/F glass fiber filters (Whatman) that remained frozen (-20 °C) until further processing. Filters were dried, acidified to remove carbonates and analyzed with an elemental analyser (Perkin-Elmer 2400 CHN).

211 **2.3.2** Chl *a* 

For Chl *a* analyses, 250 mL of sea water were filtered through 25 mm diameter glass filters (Whatman GF/F). Filters were stored frozen (-20 °C) until further processing in the home laboratory. Ninety % acetone was used to extract pigments at 4 °C in the dark

for 24 h. The procedure of Yentsch and Menzel (1963) was followed to measurefluorescence of extracts, with a calibrated Turner Designs fluorometer.

#### 217 2.3.3 Inorganic nutrients

To analyse dissolved inorganic nutrients (nitrate, phosphate and silicate), unfiltered water samples were stored frozen (-20 °C) in 10 mL sterile polypropylene bottles and further processed in the home laboratory using a Skalar Autoanalyser and the standard segmented flow analyses with colorimetric detection elemental analyser (Hansen and Grasshoff, 1983).

## 223 2.3.4 HPLC pigment analysis and CHEMTAX

224 As described in Nunes et al. (2019), HPLC was used to determine pigment composition, 225 following Latasa (2014). Thirty-two pigments were identified at 474 and 664 nm. Version 1.95 of the CHEMTAX chemical taxonomy software was used to derive the 226 contribution of microalgal groups to the total Chl *a* biomass (ng Chl *a*  $L^{-1}$ ), from 227 pigment data (Mackey et al., 1996). Seven pigmentary classes were quantified: 228 229 chlorophytes, cryptophytes, diatoms, dinoflagellates, haptophytes, prasinophytes and 230 pelagophytes. Pigments were also used as indicators of phytoplankton 231 photoacclimation. Diadinoxanthin (Ddx) is the main light-protecting pigment in diatoms, dinoflagellates, haptophytes and pelagophytes. The Ddx:LHC ratio, between 232 233 Ddx and the sum of the main light-harvesting carotenoids (LHC: fucoxanthin, 19'butanoyloxyfucoxanthin, 19'-hexanoyloxyfucoxanthin and peridinin) was measured, 234 since it varies with the exposure of phytoplankton to underwater solar radiation 235 236 (Higgins et al., 2011; Nunes et al., 2019).

#### 237 2.3.5 Picophytoplankton abundance and biomass

Abundance of picoplankton was determined by flow cytometry in a FACS Calibur instrument (Becton and Dickinson), as described in Zamanillo et al. (2019). *Prochlorococcus* and *Synechococcus* were undetectable in all samples. Biomass of picoeukaryotic algae was measured using the average C:cell conversion factor from Simó et al. (2009):  $1319 \pm 813$  fg C cell<sup>-1</sup>.

## 243 2.3.6 Microphytoplankton identification and biomass

We identified and quantified phytoplankton groups by microscopy, using the inverted microscope method (Utermöhl, 1958) as described in Nunes et al. (2019). We measured the biomass of phytoplankton groups using conversion equations of Menden-Deuer and Lessard (2000), as explained in Zamanillo et al. (2019). The total biomass of phytoplankton (Phyto B) was obtained as the sum of all the phytoplankton groups.

## 249 2.3.7 Primary production (PP)

Phytoplankton primary production rates at four depths within the water column were 250 determined by using 20 µCi of NaH<sup>14</sup>CO<sub>3</sub> (Steeman-Nielsen, 1952) in a series of 2-3 251 hours-long, on-deck incubations in flowing surface seawater and under simulated in situ 252 light levels. During the 36-h cycles, water was collected directly from the Niskin bottles 253 in 72 mL acid-washed polystyrene bottles at different sampling times (8:30, 16:30, 4:00 254 255 and 12:00 local time (UTC-3)). PP rates were estimated from radioisotope incorporation into particles retained on a 0.2 µm PC filter, after acid removal of inorganic <sup>14</sup>C and 256 substraction of the incorporation in dark controls. Daily rates were estimated by taking 257 258 into account the daylight period.

## 259 **2.3.8** Prokaryotic heterotrophic abundance (PHA) and production (PHP)

Prokaryotic heterotrophic abundance (PHA) was determined by flow cytometry following standard methods, after fixation with 1 % paraformaldehyde plus 0.05 % glutaraldehyde (Gasol and del Giorgio, 2000). PHP was estimated following the Kirchman et al. (1985) method after the centrifugation method described in Smith and Azam (1992). Final leucine concentration was 40 nM. The incubations were carried out in the dark at in situ temperature in a water bath during 3-4 h.

#### 266 **2.3.9 Viral abundance (VA)**

Viral abundance was quantified using flow cytometry (Brussaard, 2004). We
discriminated three viral subpopulations with different green fluorescence properties
(Brussaard et al., 2010) and classified them as "phage viruses" (PV), "large viruses"
(LV) and "total viruses" (TV).

## 271 2.3.10 Extracellular enzyme activities

We determined the activity of the extracellular enzymes  $\beta$ -glucosidase, esterase and fucosidase following the method detailed in Sala et al. (2016).

## 274 2.3.11 Maximum quantum efficiency of photosystem II photochemistry (F<sub>v</sub>:F<sub>m</sub>)

275 Maximum quantum efficiency of photosystem II photochemistry  $(F_v:F_m)$  was measured 276 using a Fast Repetition Rate fluorometer (FRRf; FASTracka, Chelsea Technologies, 277 Surrey, UK) on continuous seawater pumped from ca. 4 m deep, as described in Royer 278 et al. (2015).

## 279 2.4 TEP production by microbes during experimental incubations

280 To examine TEP production by microbes (phytoplankton and heterotrophic281 prokaryotes), we conducted two incubation experiments using surface water (4 m) from

282 the NSO (Experiment NSO) and the NSG (Experiment NSG). Seawater was prefiltered trough 200 µm to exclude the presence of mesozooplankton. Each experiment consisted 283 284 of two treatments: 1) unfiltered seawater (total (T) treatment), 2) seawater filtered through 0.8 µm pore-size Nucleopore filters to exclude most phytoplankton and 285 286 heterotrophic flagellates, leaving mainly prokaryotes (prokaryotic (P) treatment). Both treatments were set up in triplicates in 25 L teflonated containers under in situ irradiance 287 288 and temperature conditions. Treatments were incubated for 8 and 13 days for the experiments NSO and NSG, respectively. Experiment NSO was sampled at initial time 289 290  $(t_0)$  and after 3  $(t_1)$  and 8  $(t_2)$  days, and the experiment NSG was sampled at initial time 291  $(t_0)$  and after 2  $(t_1)$  and 13  $(t_2)$  days. The variables analysed were TEP, PHP and Chl a, 292 and we calculated TEP production rate per growth rate ( $\Delta$ TEP:  $\Delta$ Chl *a*).

#### 293 2.5 Statistical analyses

The packages *lmodel2* and *ggplot2* of R software were used to test for covariations and 294 295 to study the potential control of variables on TEP distribution in this study. Bivariate 296 analyses (ordinary least squares, OLS) between TEP concentrations and different 297 chemical, physical and biological variables were used to determine their main predictors. Although ranged major axis (RMA) regression would have been more 298 299 suitable, due to the presence of errors in our dependent and independent variables, we 300 performed OLS regressions to better compare the slopes between our study and others 301 present in the literature. We used pairwise Spearman correlation analyses to compare 302 different variables in the upper surface (10 cm) and in the surface (4 m). The data were log-transformed to fulfil the requirements of parametric tests. Non-parametric ANOVA 303 304 followed by post-hoc tests (Tukey) were carried out to compare variables among the 305 four regions. Principal component analysis (PCA) (devtools and ggbiblot packages in R) was applied to all samples after centering and scaling variables. One PCA was done 306

with a total number of 21 physical, chemical and biological variables, TEP included,
while the other one was done replacing TEP and Chl *a* by the TEP:Chl *a* ratio, which is
indicative of how prone the plankton community was to net TEP production. The map
in Fig. S1 was produced using the Ocean Data View software (version 4) (Schlitzer,
2017). The rest of plots were drawn using R programing software (version 3.5.1).

312

313 **3 Results** 

## 314 **3.1** General characterization of sea surface waters

The four study regions (NSO, SSO, NSG and WA, Fig. S1) presented distinct physical, 315 316 chemical and biological properties, summarized in Table 1. Surface temperatures in the whole transect ranged from -0.9 to 5.1 °C and salinity varied over a narrow range (33.1-317 34.2). Surface nitrate concentrations ranged from 14.6 to 32.5  $\mu$ mol L<sup>-1</sup> and covaried 318 with those of phosphate (1.0-2.5  $\mu$ mol L<sup>-1</sup>). Silicate concentrations were significantly 319 lower in the NSG region  $(2.0 \pm 0.4 \,\mu\text{mol L}^{-1})$  than in the other three regions  $(48.1 \pm 4.2)$ 320  $\mu$ mol L<sup>-1</sup>). Within the whole transect, surface Chl *a* concentrations ranged between 0.28 321 and 8.95  $\mu$ g L<sup>-1</sup> and averaged 2.36  $\pm$  1.92  $\mu$ g L<sup>-1</sup>, with a coefficient of variation of 81 %. 322 PHA ranged from 1.60 to 8.44  $\times 10^5$  cells mL<sup>-1</sup>, while PHP ranged between 0.09 and 323 2.65 mg C m<sup>-3</sup> d<sup>-1</sup>. POC ranged from 3.85 to 25.63  $\mu$ mol L<sup>-1</sup> and was higher in NSG and 324 WA (11.37-25.63  $\mu$ mol L<sup>-1</sup>). Primary production was highest in the NSG, averaging 325 111.6  $\pm$  85.4 mg C m<sup>-3</sup> d<sup>-1</sup>, while in the other three regions it averaged 16.2  $\pm$  15.8 mg C 326  $m^{-3} d^{-1}$ , being lower in the SSO (3.7 ± 0.8 mg C m<sup>-3</sup> d<sup>-1</sup>). Solar radiation dose and 327 Ddx:LHC ratio were higher in the NSO and SSO regions, whereas F<sub>v</sub>:F<sub>m</sub> was lower in 328 these regions. 329

The NSO was located within a meander of the Southern Boundary of the Antarctic 330 Circumpolar Current (Fig. S1) and two months before sampling, this region was 331 covered by > 25 % of sea ice (https://seaice.uni-bremen.de/start/). The average values of 332 the surface temperature were  $0.59 \pm 0.15$  °C (Fig. S2). The Chl *a* concentration 333 averaged 1.87  $\pm$  0.23 µg L<sup>-1</sup> (Fig. 1), and phytoplankton was dominated by 334 cryptophytes, haptophytes and diatoms (Fig. S2). The main planktonic cells identified 335 by microscopy at the surface of this region included the diatoms Corethron pennatum, 336 Fragilariopsis spp. and Thalassiosira spp., unidentified autotrophic cryptophytes and 337 and heterotrophic Gyrodinium autotrophic 338 nanoflagellates, (like spp.) and 339 dinoflagellates (Nunes et al., 2019).

340 The SSO was placed north of the Weddell Front (Fig. S1). During the sampling month, the ice cover was > 25 % in the region (https://seaice.uni-bremen.de/start/) (Nunes et al., 341 342 2019). The month before, the ice cover was > 50 %; therefore, we sampled waters of the 343 receding ice edge. The surface temperature was the lowest among the four regions (- $0.75 \pm 0.11$  °C). Surface Chl *a* averaged  $0.32 \pm 0.06 \ \mu g \ L^{-1}$ , i.e., significantly (p< 0.05) 344 lower than in the other regions (Fig. 1). Phytoplankton was dominated by haptophytes, 345 346 followed by cryptophytes, chlorophytes and diatoms (Fig. S2). The main planktonic cells identified by microscopy were the same as in the NSO but with a higher proportion 347 348 of Fragilariopsis spp., a characteristic diatom marker of sea ice influence. The SSO presented the highest Ddx:LHC ratio (0.58  $\pm$  0.09; Fig. 1) and the lowest  $F_v:\!F_m$  ratio 349 350  $(0.16 \pm 0.05)$ , consistently indicating exposure to higher solar radiation intensities in a 351 shallower mixed layer (Table 1).

The NSG was located right south of the Polar Front, north of the subAntarctic South Georgia Islands (Fig. S1). Sea surface temperature was the highest among regions (4.77  $\pm 0.45$  °C), and the MLD was deepest. The relatively low silicate concentrations found

in this region coincided with the dominance of well-silicified diatoms in the 355 phytoplankton community (like Eucampia Antarctica, Thalassiosira and Porosira spp. 356 and Odontella weissflogii; Fig. S2) and a smaller proportion of haptophytes and 357 pelagophytes. Surface Chl *a* concentration averaged 4.59  $\pm$  1.97 µg L<sup>-1</sup> (Fig. 1). The 358 phytoplankton community was complemented by haptophytes and pelagophytes. Total 359 viral abundances (TVA), PHP, PHA and  $F_{v}$ :  $F_{m}$  were significantly higher (p< 0.05) in 360 this region than in the others, whereas the solar radiation dose and Ddx:LHC were the 361 362 lowest.

The WA region was placed at the southernmost limit of the Antarctic Circumpolar Current (Southern Boundary) (Fig. S1). The average surface temperature was 1.46  $\pm$ 0.09 °C. Chl *a* concentration was slightly lower than in the NSG, with an average of 4.05  $\pm$  0.48 µg L<sup>-1</sup> (Fig. 1). The phytoplankton community was dominated by cryptophytes, haptophytes and diatoms (Fig. S2).

# 368 **3.2** TEP concentrations, TEP:Chl *a*, TEP:Phyto B and TEP:PP ratios and 369 contribution of TEP to POC in the sea surface of the study regions

Surface TEP concentrations in the entire study regions and transects ranged from 39.2 to 177.6  $\mu$ g XG eq L<sup>-1</sup> and averaged 102.3 ± 40.4  $\mu$ g XG eq L<sup>-1</sup>, with a coefficient of variation of 39 %. Mean TEP concentrations were significantly different among regions (p< 0.05; Table 1). NSO presented the maximum TEP concentrations (144.4 ± 21.7  $\mu$ g XG eq L<sup>-1</sup>) and SSO presented the lowest ones (48.1 ± 6.5  $\mu$ g XG eq L<sup>-1</sup>), while NSG and WA presented similar TEP concentrations (125.5 ± 21.1 and 111.6 ± 13.0  $\mu$ g XG eq L<sup>-1</sup>, respectively; Fig. 1).

The TEP:Chl *a* ratios were also significantly different among regions (p< 0.05; Fig. 1). SSO presented the highest TEP:Chl *a* values (153.4  $\pm$  29.8), while NSG and WA presented the lowest ratios  $(32.3 \pm 15.0 \text{ and } 28.2 \pm 4.8, \text{ respectively; Fig. 1})$ . The ratio in NSO was 76.7 ± 10.6. In the case of TEP:Phyto B and TEP:PP ratios, the SSO presented the highest values and the NSG the lowest ones (Table 1).

- 382 TEP and POC were significantly and positively correlated across the entire transect
- $(R^2 = 0.70, p < 0.001, n = 68; Table 2)$ . The contribution of TEP-C to the POC pool (TEP-
- 384 C%POC) ranged between 17.9 and 97.3 % (average 38.8 ± 12.3 %). TEP-C%POC in
- the NSO and the SSO were similar (44.7  $\pm$  6.7 and 38.5  $\pm$  8.7 %, respectively) and

higher than in NSG ( $30.6 \pm 6.2 \%$ ) and WA ( $25.2 \pm 5.3 \%$ ; Fig. 1).

## 387 **3.3 TEP in relation with other variables at the sea surface**

TEP were significantly and positively related to Chl *a* across the entire study area ( $R^2$ = 0.66, p< 0.001, n= 65; Table 2) and phytoplankton biomass ( $R^2$ = 0.66, p< 0.001, n= 67; Table 2). The regression equation for log converted TEP vs Chl *a* was log TEP = 1.90 (± 0.02) + 0.35 (± 0.03) × log Chl *a*.

TEP also presented significant positive relationships with temperature, salinity (p< 0.001) and all the phytoplankton groups (p< 0.05; Table 2), with explained variances higher than 60 % for prasinophytes, dinoflagellates and diatoms. By contrast, TEP were negatively related to the solar radiation dose, PHA:Chl *a*, Ddx:LHC, nutrients and wind speed at the sampling time (1-h average; Table 2). TEP did not present any significant relationship with PHA, although it was weakly but significantly related to PHP (Table 2).

399 PCA(a) and PCA(b) were performed to visualize the differences between regions and 400 the subset of variables that better explained TEP and TEP:Chl *a* patterns (Fig. 2). At 401 PCA(a) principal components 1 and 2 explained 57.1 % and 16.5 % of the variability, 402 respectively. SSO and NSG occupied the extremes of the PC1 gradient, characterized by

high solar radiation dose and Ddx:LHC ratio on one side (SSO) and higher TEP 403 concentrations and phytoplankton abundance, and lower macronutrient availability on 404 the other (NSG). WA and NSO showed intermediate values of PC1, but their position in 405 406 relationship with PC2, indicating an association with high cryptophytes and low wind speed values, was opposite to that of SSO and NSG. At PCA(b), where TEP and Chl a 407 were replaced by TEP:Chl a, the regions were separated along PC1 and 2 similarly to 408 PCA(a). The major loadings to component 1 positively related to Ddx:LHC, TEP:Chl a 409 410 and solar radiation dose, and negatively related to diatoms, temperature and POC. The major loadings to component 2 included cryptophytes, chlorophytes and, negatively, 411 wind speed,  $F_v$ :  $F_m$  and TVA. 412

## 413 **3.4. TEP vertical distribution and relationship with other variables**

414 TEP concentrations were generally higher within the upper mixed layer (average 119.2  $\pm$  39.0 µg XG eq L<sup>-1</sup>) than below it (average 53.1  $\pm$  45.2 µg XG eq L<sup>-1</sup>), and decreased 415 with depth in 4 out of 7 stations. In the other three stations, TEP maxima were found at 416 417 the deep Chl a maximum (DCM) or close to it (Fig. 3). The vertical distribution of Chl a differed among regions, but Chl a concentration was generally higher in the upper 50 418 m of the water column. TEP vertical distributions were significantly related to those of 419 Chl *a* considering profiles of all regions ( $R^2 = 0.74$ , p< 0.001, n= 26). The equation 420 obtained for the TEP-Chl *a* vertical relationship is  $\log_{10}$  (TEP) = 1.78 (± 0.04) + 0.67 (± 421  $(0.08) \times \log_{10}$  (Chl a). By contrast, the relationship between TEP and PHA was weak 422  $(R^2 = 0.18, n = 24, p < 0.05)$ . The Ddx:LHC ratio decreased strongly below 20-40 m 423 424 depth in all regions. TEP:Chl a ratios increased towards the surface in the NSO and 425 SSO, while in the other regions were constant or with the opposite pattern (Fig. S3). 426 TEP:Phyto B ratios, when available, generally followed a similar pattern than TEP:Chl *a* ratios in the profiles (data not shown). 427

In NSO and WA, TEP vertical distribution was highly coupled with haptophytes and cryptophytes (Fig. 3). In the NSO diatoms were higher below TEP maximum. In the SSO, TEP maxima were found above or coincident with the Chl *a* peaks and always above PHA maxima, and the most important phytoplankton groups (haptophytes and diatoms) increased their concentration deeper in the water column (Fig. 3). In NSG, the vertical TEP profile was similar to that of diatoms, which dominated the phytoplankton assemblage at all depths (Fig. 3).

## 435 **3.5.** Organic matter accumulation in the upper surface (10 cm)

Among all the parameters measured in the upper surface, TEP concentration (r= 0.71, n= 14), Chl *a* concentration (r= 0.70, n= 13), PHA (r= 0.71, n= 13), fucosidase (r=0.56, n= 14) and esterase activity (r= 0.82, n= 12) were significantly correlated (Spearman)

with the measurements of the same variables at 4 m depth (p < 0.05).

At most stations, TEP and the activity of the enzymes β-glucosidase, esterase and fucosidase were enriched in the upper surface (10 cm; Fig. 4) relative to 4 m. TEP-EF<sub>us</sub> averaged  $1.19 \pm 0.19$ , and the EF<sub>us</sub> average for the enzymatic activities were  $2.75 \pm 3.24$ (β-glucosidase),  $4.75 \pm 1.19$  (esterase) and  $31.40 \pm 100.78$  (fucosidase). TEP-EF<sub>us</sub> was positively related to the 1 h-average wind speed at the time of the upper surface sampling (R<sup>2</sup>= 0.58, p< 0.005, n= 14) (log<sub>10</sub> transformed), which varied among 0.5 and 8.6 m s<sup>-1</sup>, with an average of  $5.4 \pm 2.5$  m s<sup>-1</sup>.

- 447 By contrast, Chl *a*, PHA and PHP were not usually enriched in the upper surface. Their
- enrichment factors averaged 0.97  $\pm$  0.37 (Chl *a*), 1.00  $\pm$  0.27 (PHA) and 0.77  $\pm$  0.47
- (PHP) (Fig. 4, 7). TEP-EF<sub>us</sub> and Chl *a*-EF<sub>us</sub> were not significantly related (p > 0.05), but
- 450 TEP:Chl *a* ratios were generally higher at the upper surface than at the surface (average
- 451 TEP:Chl *a*-EF<sub>us</sub>  $1.37 \pm 0.44$ ).

#### 452 **3.6. TEP variations in the 36-h cycles**

We did not detect recurrent diel patterns of Chl *a*, PHP or TEP in the diel cycle studies (Fig. S5). TEP showed a relative amplitude (max-min/mean) of 0.31, 0.20, 0.29 and 0.31 in NSO, SSO, NSG and WA, respectively. Chl *a* relative amplitude (0.33, 0.07, 0.95 and 0.37, respectively) was highest in NSG. Looking at covariations (Spearman) of TEP with other biological variables in diel cycles, TEP were only significantly (p<

458 0.05) coupled with Chl *a* (r=0.81, n=8) in NSG.

#### 459 **3.7. TEP production by microbes during experimental incubations**

In the two incubation experiments (conducted in NSO and NSG), and in both treatments 460 (T and P), all measured variables (TEP, Chl a and PHP) increased over time, except 461 PHP in experiment NSG (Fig. 5). In the T treatments, the increases were steeper (higher 462 daily increase rates), and TEP evolved in parallel with Chl a concentration. The TEP 463 464 production rate per growth rate was 78 in the experiment NSO and 18 in the experiment 465 NSG. These values were very similar to the TEP:Chl a ratio average of the regions where the water was collected from (76.7  $\pm$  10.6 in the NSO and 32.5  $\pm$  14.5 in the 466 467 NSG).

468

#### 469 **4 Discussion**

#### 470 **4.1 TEP concentrations in the Southern Ocean**

In the present study, we advance the existing knowledge on TEP variability across environmental conditions in the Southern Ocean, with a combination of surface observations, vertical profiles, short-term variability and changes associated with microbial growth during laboratory incubations. The few previous studies describing TEP distributions in the Southern Ocean were located around the Antarctic Peninsula 476 (Passow et al., 1995; Corzo et al., 2005; Ortega-Retuerta et al., 2009b), the Drake
477 Passage (Corzo et al., 2005) and the Ross Sea (Hong et al., 1997). Our study was carried
478 out in four distinct regions of the peninsular area, and the PCA analyses (Fig. 2)
479 confirmed a clear separation of these regions.

Our surface TEP concentrations are in the upper range of published values from around 480 481 the Antarctic Peninsula (Corzo et al., 2005; Ortega-Retuerta et al., 2009b; Table S1). In 482 contrast, the TEP:Chl a ratios in our study are not substantially higher than those reported in previous works. It is worth mentioning that Ortega-Retuerta et al. (2009b) 483 484 found remarkably lower TEP and TEP:Chl a ratios in the western Weddell Sea area 485 compared with our SSO data (northern Weddell Sea), although the two sampling sites 486 are geographically close. Difference in the sampling periods can account for this variability, since the previous study was conducted later in the season, when TEP may 487 488 have been consumed or degraded. The values reported by Hong et al. (1997) in the Ross 489 Sea were higher than our observations in NSO and SSO, probably because they found a 490 bloom of colonial *Phaeocystis antarctica* with higher Chl a concentrations and the characteristic presence of mucilage flocs. Regarding vertical TEP distribution, the 491 492 general pattern of higher TEP concentrations in the upper mixed layer has already been reported in previous studies in the SO (Hong et al., 1997; Corzo et al., 2005; Ortega-493 494 Retuerta et al., 2009b) and potential causes for this pattern will be discussed below.

## 495 **4.2 TEP distribution and its main drivers in the Southern Ocean**

Phytoplankton appears to be the main driver of TEP distribution in this study at both the horizontal and vertical scales, as suggested by the positive relationships between TEP and Chl a and phytoplankton biomass. However, it is worth mentioning that the coefficient of variation of surface Chl a concentration along the entire cruise track (81

%) was much higher than that of TEP (39 %). This suggested that TEP concentrations 500 were relatively more stable than those of their main sources: total phytoplankton 501 502 biomass expressed as Chl a concentration and biomass of the different phytoplankton 503 groups. Considering surface (4 m) samples only, the slope of the log-converted TEP-Chl *a* relationship for the entire study ( $\beta$ = 0.35 (± 0.03); Table 2, Fig. 6) was very 504 similar to that obtained in the Antarctic Peninsula area (Corzo et al., 2005; Ortega-505 Retuerta et al., 2009b) but lower than those observed in the Ross Sea (Hong et al., 506 507 1997), driven by the aforementioned bloom of the TEP-producing *Phaeocystis*, and in the Gerlache Strait (Corzo et al., 2005). Concerning surface waters, TEP presented 508 509 significant positive relationships with all the phytoplankton groups. At the vertical 510 scale, the positive relationship between TEP and Chl *a* corroborates findings of previous studies in the area (Corzo et al., 2005; Ortega-Retuerta et al., 2009b) and suggests that 511 512 phytoplankton abundance, generally higher in the upper mixed layer, is the main cause 513 for the general higher TEP concentrations found in this layer too. Our work presents for 514 the first time the vertical TEP distribution along with the phytoplankton group 515 distributions in the SO, showing that haptophytes and cryptophytes were closely associated with TEP vertical distribution in the NSO and WA regions, while diatoms 516 did in the NSG region. 517

The fact that TEP:Chl *a* and TEP:Phyto B ratios were different across regions and depths, further supports the role of drivers other than phytoplankton biomass on TEP distribution. On this line, phytoplankton light stress appeared to control TEP distributions in our study area. The SSO presented the highest solar radiation doses and Ddx:LHC ratios, and the lowest  $F_v:F_m$  values, which indicates stronger sunlight exposure and the need for active photoacclimation (Bergmann et al., 2002; Kaiblinger et al., 2007). The NSO also presented relatively high Ddx:LHC ratios and solar radiation

doses (Table 1). In these regions, TEP:Chl a ratios were the highest, which could be 525 related to a decrease of Chl a per cell due to photoadaptation (Kiefer et al., 1976) but 526 also to a stimulation of TEP production, as previously suggested by Iuculano et al. 527 528 (2017b) and Agustí and Llabrés (2007). The fact that TEP:Phyto B and TEP:PP ratios were also higher in these regions than in the NSG and WA is in line with this second 529 explanation. In the PCA(b) analysis (Fig. 2), we could clearly determine that high 530 TEP:Chl a ratios corresponded to regions with higher macronutrient availability, solar 531 532 radiation dose and Ddx:LHC ratio, highlighting the role of light stress in driving TEP:Chl a ratios. 533

534 The occurrence of sea ice cover and sea ice melting also seemed to influence TEP 535 concentration in the SSO region. It is known that microorganisms living in sea ice release organic substances (Assmy et al., 2013; Vancoppenolle et al., 2013), including 536 537 TEP, as a survival strategy (Ewert and Deming, 2013). In fact, TEP concentrations measured in sea ice samples during our cruise were > 600  $\mu$ g XG eq L<sup>-1</sup>, i.e., way above 538 539 concentrations in sea water, although it must be noticed that the selection of ice chunks was based on their colour, picking the brownish as indicative of colonisation by algae 540 (Dall'Osto et al., 2017). The lower salinity at the surface and the larger proportion of 541 Fragilariopsis spp. were indicative of sea ice melt influence in SSO (Cefarelli et al., 542 543 2011). Consequently, the highest TEP:Phyto B ratios in surface waters of SSO may have partially resulted from an enhancement of TEP release due to sea ice melt. This 544 545 result agrees with previous studies suggesting sea ice melt being a source of TEP into 546 seawater (Assmy et al., 2013; Galgani et al., 2016), although this role should be studied more in detail if we are to predict TEP concentrations in polar regions. 547

548 Nutrient depletion and the phase stage of the phytoplankton blooms could have also 549 influenced TEP distribution too, but we cannot assess this hypothesis with the data at 550 hand. NSG was the only case that presented macronutrient-deplete conditions (silicate restriction; Nunes et al., 20189). The other regions probably were iron-limited, but this 551 micronutrient was not monitored. Regarding the phase stage of the phytoplankton 552 553 blooms, since in NSO the sea ice retreat occurred earlier in the season, we hypothesize that NSO hosted an advanced phase of the SSO bloom, favoring TEP production (Corzo 554 et al., 2000). Therefore, the bloom phase as sea ice recedes, the influence of sea ice 555 melt, and the aforementioned high insolation in shallow mixed layer, all possibly 556 557 contributed to TEP distribution.

Regarding the vertical distribution, the increase of Ddx:LHC above 20-40 m in all cases, indicative of protection against light stress, suggests that light stress could have also triggered TEP production near surface of some regions (Iuculano et al., 2017b). NSO and SSO, the regions with the strongest solar radiation doses, presented a general increase of TEP:Chl *a* ratio towards the surface (Fig. S1), and also of TEP:Phyto B ratio in the single profile from SSO. However, caution must be taken due to the limited data available of TEP:Phyto B ratios across the profiles.

Heterotrophic prokaryotes did not contribute significantly to explain TEP variations 565 566 across the horizontal or vertical scales, given the weak relationship between TEP and 567 PHA and PHP (Table 2). Other studies have also found a lack or a negative relationship between TEP and PHA or PHP (Passow and Alldredge, 1994; Bhaskar and Bhosle, 568 2006; Zamanillo et al., 2019). In contrast, some works, including two in the Southern 569 570 Ocean, have found a positive relationship between these variables (Passow et al., 2001; Hung et al., 2003; Santschi et al., 2003; Corzo et al., 2005; Ortega-Retuerta et al., 571 572 2009b; Ortega-Retuerta et al., 2010; Zamanillo et al., 2019).

Viral abundances did not significantly contribute to explain TEP variations neither in
the horizontal nor in the vertical scale (Table 2), although some studies have found a
link between TEP production and the viral infection of different phytoplankton taxa,
such as *Emiliania huxleyi* (Nissimov et al., 2018), *Phaeocystis globosa*-(Grossart et al.,
1998; Stoderegger and Herndl, 1999; Passow, 2002b; Brussaard et al., 2005; Mari et al.,
2005) and *Micromonas pusilla*-(Lønborg et al., 2013). This result suggests the
predominance of other factors driving TEP distribution in this study.

Few studies have measured and compared the distribution of TEP and other organic 580 matter compounds within the first few meters of the ocean surface (Wurl et al., 2011a; 581 582 Bélanger et al., 2013; Taylor et al., 2014; Thornton et al., 2016). The significant positive 583 correlations we found for most of the measured parameters (TEP, Chl a, PHA, esterase and fucosidase activities) at 10 cm vs 4 m, indicates that the horizontal variability of 584 these parameters at the surface (4 m) is mirrored at the upper surface (10 cm). TEP were 585 586 generally enriched in the upper surface. The TEP-EF<sub>us</sub> average  $(1.19 \pm 0.19)$  was lower 587 than that found in the SML with respect to underlying water in most previous studies (Wurl and Holmes, 2008; Cunliffe et al., 2009; Wurl et al., 2009; Wurl et al., 2011b; 588 Engel and Galgani, 2016; Galgani et al., 2016). This is somehow expected owing to the 589 distinct properties of the SML, which forms by the accumulation of surface-active 590 compounds (Wurl et al., 2017). However, despite the modest TEP enrichment at 10 cm, 591 TEP could be responsible for the enhancement of the activities of the ectoenzymes  $\beta$ -592 glucosidase, fucosidase and esterase (Fig. 4), which are secreted by heterotrophic 593 prokaryotes to hydrolyze polysaccharides such as TEP into smaller, bio-available 594 macromolecules (Bar-Zeev and Rahav, 2015). 595

596 The enrichment of TEP in the upper surface was not associated with higher abundances 597 of microorganisms or higher prokaryotic heterotrophic activity (Fig. S4). Chl *a* and

PHA were not generally enriched in the upper surface. Therefore, reasons for the TEP 598 enrichment could be the following; (1) the increase of per-cell TEP production in the 599 upper surface due to high solar radiation (Ortega-Retuerta et al., 2009a; Iuculano et al., 600 2017b); (2) an allochthonous source of TEP and/or their precursors at 10 cm, namely 601 sea ice melt waters of lower density; (3) ascending TEP from underlying waters, 602 603 embedded in low density particles or captured and transported by air bubbles (Zhou et al., 1998); and (4) an increase of TEP self-assembly of dissolved precursors in the upper 604 605 surface due to higher turbulence (Passow, 2000; Beauvais et al., 2006). The last two mechanisms are the most plausible since we found a positive relationship between TEP-606  $EF_{us}$  and wind speed ( $R^2 = 0.58$ ) (Fig. S4). Winds stronger than about 5 m s<sup>-1</sup> cause the 607 entrainment of bubbles down from breaking waves (Andreas and Monahan, 2000; 608 609 Deane and Stokes, 2002), and when these bubbles rise they can bring TEP or their 610 precursors towards the surface, among other items such as prokaryotic cells (Mayol et al., 2017), where aggregation is favoured by enhanced turbulence. A recent study 611 612 (Robinson et al., 2019) has shown that deliberate bubbling of seawater (mimicking the 613 effect of wind in the field) increases TEP concentrations in the SML, and although they did not observe an increase of TEP concentration at 1 m, it is plausible that an increase 614 at 10 cm was also occurring, which is in line with our observations. 615

## 616 **4.5 TEP variations in the 36-h cycle**

We did not find a cyclic diel pattern of any of the measured variables (TEP, Chl *a* and PHP) similarly to the outcome of the only published study on short-term TEP variation (Ortega-Retuerta et al., 2017). This result illustrates the influence of many processes in driving TEP distribution.

#### 621 **4.6 TEP production by microbes in incubation experiments**

622 The results of the incubation experiments supported the idea that phytoplankton are the main TEP producer in the study area (Fig. 5). In treatments with only prokaryotes, TEP 623 624 were formed at low rates by abiotic self-assembly and/or the action of heterotrophic 625 prokaryotes and archaea. In unfiltered treatments, TEP increased much faster and nearly 626 parallel to the Chl a increase. Net TEP production rates per net phytoplankton growth rate varied between experiments and were very similar to the average of TEP:Chl a 627 ratios of the respective regions, confirming our observations that phytoplankton were 628 629 the main TEP producers in the study area. Interestingly, TEP production rate per phytoplankton growth rate was higher in the NSO experiment than in the NSG 630 experiment, despite the higher biomass and dominance of diatoms in the latter. This 631 contradicts the common view that diatoms are highly specific TEP producers (Passow, 632 2002a), and supports the aforementioned discussed idea that environmental conditions, 633 634 associated physiological stress and the phytoplankton bloom stage influence TEP 635 production by phytoplankton beyond biomass, primary productivity and taxonomical 636 composition.

637

#### 638 **5** Conclusions

Our study expands the existing knowledge on TEP distribution in the Southern Ocean and for the first time provides information on TEP variability in the first meters of the ocean surface and with high temporal resolution. Phytoplankton abundance was the main predictor of TEP distribution in both the horizontal and vertical scales, and the outcome of experimental incubations further supported this observation in situ. Photoacclimation and sea ice melt played complementary but important roles driving differences in TEP distribution across regions. Phytoplankton composition was also an

important driver, especially along the vertical scale, but diatoms were not the main 646 producers. TEP were generally enriched at centimetres under the surface, likely due to 647 scavenging by bubbles and turbulence-derived aggregation, contrasting with the null 648 649 enrichment of microorganism abundances. We highlight the need to carry out more field 650 studies in the area, extending to temporal dynamics along the entire ice-receding and ice-free season, to better predict TEP concentration in the Southern Ocean and 651 quantitatively evaluate their effects in biogeochemical processes such as the biological 652 653 carbon pump and organic aerosol formation.

654

#### 655 Author Contributions

MZ conducted the study, analysed samples, processed and analysed the data. EOR and RS designed the study and analysed data. SN, MEs, MS, SJR, MEm, DV, CM and DLS analysed samples and provided data. MZ, EOR and RS wrote the manuscript with the help of all the co–authors.

#### 660 Competing interests

661 The authors declare that they have no conflict of interest

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## 1032 Figure captions

1033 **Figure 1**: Boxplots of transparent exopolymer particles (TEP), chlorophyll *a* (Chl *a*), TEP:Chl a, contribution of TEP-C to the POC (particulate organic carbon) pool (TEP-1034 C%POC), solar radiation dose, Ddx:LHC ratio (Ddx: Diadinoxanthin, LHC: sum of the 1035 main light-harvesting carotenoids), prokaryotic heterotrophic abundance (PHA) and 1036 prokaryotic heterotrophic production (PHP) in the surface of the four visited regions 1037 1038 :North of the South Orkney Islands (NSO), South of the South Orkney Islands (SSO), Northwest of South Georgia Island (NSG), West of Anvers Island (WA). 1039 The horizontal lines of the boxes represent 25%, 50% (median) and 75% percentiles (from 1040 1041 bottom to top). Whiskers represent minimum and maximum values.

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Figure 2: Principal component analyses (PCA) of surface (4 m) samples. The % of the 1043 overall explained variance is given on each principal component axis. The plot (b) is the 1044 same as (a) plot after replacing transparent exopolymer particles (TEP) and chlorophyll 1045 a (Chl a) by TEP:Chl a. Note that the opposite direction of axes in PCA(a) and (b) is 1046 only due to calculations Chlorophytes (Chlor), cryptophytes (Crypt), diatoms (Diat), 1047 1048 dinoflagellates (Dinof), haptophytes (Hapt), prasinophytes (Pras), pelagophytes 1049 (Pelago), Ddx:LHC (see Fig. 1 for abbreviations), Maximum quantum efficiency of photosystem II photochemistry (FvFm), temperature (Temp), total viral abundance 1050 (TVA), prokaryotic heterotrophic production (PHP) and particulate organic carbon 1051 (POC). 1052

Figure 3: Vertical profiles of temperature (Temp), salinity (Sal), concentration of the
different phytoplankton groups derived from CHEMTAX (see Fig. 2 for abbreviations),
transparent exopolymer particles (TEP), chlorophyll *a* (Chl *a*) and prokaryotic
heterotrophic abundance (PHA) in every region (see Fig. 1 for abbreviations of regions).

**Figure 4**: Boxplots of  $EF_{us}$  of transparent exopolymer particles (TEP), chlorophyll *a* (Chl *a*), TEP:Chl *a*, prokaryotic heterotrophic abundance (PHA), prokaryotic heterotrophic production (PHP), and fucosidase, esterase and  $\beta$ -glucosidase activities across the study regions (n= 14). See Fig. S4 for the definition of  $EF_{us}$ .

**Figure 5**: Incubation experiments at NSO (top) and NSG (bottom). Transparent exopolymer particles (TEP) concentration (blue circle), prokaryotic heterotrophic production (PHP) (pink triangle) and chlorophyll *a* (Chl *a*) (green square) are plotted over time (h). Filled symbols and continuous lines: unfiltered water (treatment T). Empty symbols and dashed lines: water filtered through 0.8  $\mu$ m, hence, only with prokaryotes (treatment P).

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1068 Figure 6: Log-log relationship between transparent exopolymer particles (TEP) and chlorophyll a (Chl a) concentration from the PEGASO cruise, with the linear regression 1069 1070 line (regression equation in the text). Four regions are distinguished: NSO (empty circles), SSO (empty triangles), NSG (filled circles), WA (crosses) (see Fig. 1 for 1071 abbreviations of regions), T: transect (plus). Regression lines from the literature in the 1072 Southern Ocean are also shown for comparison.  $\alpha$  and  $\beta$  indicate the intercept and the 1073 slope, respectively; log TEP ( $\mu$ g XG eq L<sup>-1</sup>) =  $\alpha + \beta \times \log$  Chl a ( $\mu$ g L<sup>-1</sup>); [a]  $\alpha = 2.25$ 1074 and  $\beta = 0.65$ , (Hong et al., 1997); [b]  $\alpha = 1.63$  and  $\beta = 0.32$ , (Corzo et al., 2005); [c]  $\alpha =$ 1075

1076 1.52 and  $\beta = 0.67$ , (Corzo et al., 2005); [d]  $\alpha = 1.08$  and  $\beta = 0.38$ , (Ortega–Retuerta et al., 2009b).

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**Figure S1**: Study area. Ship trajectory (blue line) and the four case studies with Lagrangian occupation (circles, see Fig. 1 for abbreviations of regions). Main hydrographic fronts (black lines) along the cruise: Antarctic Polar Front (PF), Southern Antarctic Circumpolar Current Front (SACCF), Southern Boundary of the Antarctic Circumpolar Current (SB), and Weddell Scotia Confluence Zone (WSCF). Figure produced with the Ocean Data View software.

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**Figure S2**: Average concentration of each phytoplankton group in the sea surface of the four visited regions derived from CHEMTAX (contribution to the total Chl *a*, ng Chl *a*  $L^{-1}$ ) (see Fig. 1 for abbreviations of regions).

**Figure S3**: Vertical profiles of TEP:Chl *a* ratio (transparent exopolymer particles (TEP), chlorophyll *a* (Chl *a*)) in every study region (see Fig. 1 for abbreviations of regions).

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**Figure S4**: Enrichment factor between the upper surface (10 cm) and the surface (4 m) ( $EF_{us}= [X]$  upper surface/ [X] surface) for transparent exopolymer particles (TEP), chlorophyll *a* (Chl *a*), prokaryotic heterotrophic abundance (PHA) and prokaryotic heterotrophic production (PHP) (bars) in the different regions (see Fig. 1 for abbreviations of regions). Horizontal line denotes  $EF_{us}= 1$ . Wind speed is indicated by filled circles. T: transect.

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1100Figure S5: Transparent exopolymer particles (TEP) concentration (blue circle),1101prokaryotic heterotrophic production (PHP) (pink triangle) and chlorophyll a (Chl a)1102(green square) in the 36-h cycles (local time, UTC-3) in every visited region (see Fig. 11103for abbreviations of regions). Grey shaded areas indicate night by enclosing the period1104when global solar radiation was < 10 W m<sup>-2</sup>.

**Table 1.** Mean  $\pm$  standard deviation of several variables and dominant phytoplankton groups in the sea surface (4 m depth) of the four study regions. MLD: mixed layer depth. Phyto B: Phytoplankton biomass. PP: Primary production. POC: Particulate organic carbon. PHA and PHP: Prokaryote heterotrophic abundance and production, respectively. TVA and LVA: Total and large viral abundance, respectively.  $F_v:F_m$ : Maximum quantum efficiency of photosystem II photochemistry. Ddx:LHC: Ddx, diadinoxanthin; LHC, sum of the main light-harvesting carotenoids. TEP: Transparent exopolymer particles. See Fig. 1 for abbreviations of regions.

	NSO (n= 16)	SSO (n= 15)	NSG (n= 14)	WA (n= 9)
Date (d/m/y)	10-15/01/2015	16-20/01/2015	23-25/01/2015	02-03/02/2015
Temperature (°C)	$0.59\pm0.15~^a$	-0.75 $\pm$ 0.11 <sup>b</sup>	$4.77\pm0.45^{c}$	$1.46\pm0.09^{d}$
Salinity	$33.83\pm0.08^{a}$	$33.15\pm0.05^{\text{ b}}$	$33.74 \pm 0.02^{\circ}$	$33.41\pm0.03^{d}$
MLD (m)	$28.9\pm11.8^{\text{ a}}$	$15.8\pm5.4^{\text{ b}}$	$49.3\pm11.8^{c}$	$22.9\pm5.6^{\ ab}$
Ice cover (%)	0 <sup>e</sup>	> 25	0	0
Solar radiation dose (W m <sup>-2</sup> )	$80.3\pm40.4~^a$	$101.7\pm20.3^{\text{ a}}$	$22.4\pm5.4^{\text{ b}}$	$47.0{\pm}~10.6^{\text{ b}}$
Nitrate ( $\mu$ mol L <sup>-1</sup> )	$27.3\pm1.9~^{a}$	$27.5\pm3.2~^{a}$	$17.2\pm1.6^{\ b}$	$18.7\pm0.9^{b}$
Phosphate ( $\mu$ mol L <sup>-1</sup> )	$2.0\pm0.2^{aa}$	$2.1\pm0.3^{\ ab}$	$1.3\pm0.2^{c}$	$1.8\pm0.2^{\;aa}$
Silicate (µmol L <sup>-1</sup> )	$47.9\pm4.1~^a$	$47.3\pm4.7~^{a}$	$2.0\pm0.4^{b}$	$49.7{\pm}3.7^{a}$
Chl $a$ (µg L <sup>-1</sup> )	$1.87 \pm 0.23^{a}$	$0.32\pm0.06^{\text{ b}}$	$4.59\pm1.97^{c}$	$4.05\pm0.48^{c}$
Phyto B ( $\mu$ g C L <sup>-1</sup> )	$95.7\pm21.0^{\text{ a}}$	$22.3\pm4.4^{\text{ b}}$	$222.1 \pm 53.1$ <sup>c</sup>	$94.4 \pm 37.5^{a}$
PP (mg C $m^{-3} d^{-1}$ )	$13.0\pm4.0^{\:a}$	$3.7\pm0.8^{a}$	$111.6 \pm 85.4^{b}$	$34.6 \pm 15.7^{a}$
POC ( $\mu$ mol L <sup>-1</sup> )	$13.8\pm2.6^{\ a}$	$5.5\pm1.2^{b}$	$18.0\pm4.4^{\rm c}$	$19.3\pm3.5^{\circ}$
PHA (cells mL <sup>-1</sup> $\times$ 10 <sup>5</sup> )	2.34±0.46 <sup>a</sup>	2.89±0.70 <sup>ab</sup>	5.68±0.98 °	$3.43 \pm 0.95^{b}$
PHP ( $\mu g C L^{-1} d^{-1}$ )	$0.50\pm0.16^{\:a}$	$0.32\pm0.19^{\text{ a}}$	$1.23\pm0.66^{\:b}$	$0.40 \pm 0.14^{\ a}$
TVA (virus mL <sup>-1</sup> $\times$ 10 <sup>5</sup> )	44.8±21.8 <sup>a</sup>	32.3±17.1 <sup>a</sup>	211.2±99.8 <sup>b</sup>	137.1±42.5 °
LVA (virus mL <sup>-1</sup> $\times$ 10 <sup>5</sup> )	2.8±0.9 <sup>a</sup>	2.5±2.3 <sup>a</sup>	25.3±19.5 <sup>b</sup>	8.9±5.1 <sup>a</sup>
Dominant phytoplankton	Cryptophytes >	Haptophytes >	Diatoms	Cryptophytes
groups	diatoms	chlorophytes ~		> diatoms
F <sub>v</sub> :F <sub>m</sub>	$0.18\pm0.06^{\text{ a}}$	$0.16\pm0.05^{a}$	$0.29 \pm 0.05^{b}$	$0.21 \pm 0.06^{a}$
Ddx:LHC	$0.36\pm0.06^{a}$	$0.58 \pm 0.09$ <sup>b</sup>	$0.19\pm0.04^{\rm \ c}$	$0.25 \pm 0.04$ <sup>c</sup>
TEP ( $\mu g XG eq L^{-1}$ )	$144.4\pm21.7^{a}$	$48.1\pm6.5^{b}$	$125.5 \pm 21.1^{\circ}$	$111.6 \pm 13.0^{\circ}$
TEP:Chl $a$ (µg XG eq µg <sup>-1</sup> )	$76.7\pm10.6^{a}$	$153.4 \pm 29.8$ <sup>b</sup>	$32.3\pm15.0^{\text{ c}}$	$28.2\pm4.8^{c}$

TEP:Phyto B	1.56 ±0.35 <sup>a</sup>	$2.24 \pm 0.54$ <sup>b</sup>	$0.61 \pm 0.18$ <sup>c</sup>	$1.26\pm0.36^{a}$
TEP:PP	$12.5 \pm 3.1^{a,b}$	$14.0\pm5.7~^{a}$	$1.8 \pm 1.3^{\mathrm{c}}$	$4.6\pm4.1^{\text{ b,c}}$
TEP:POC ( $\mu$ g XG $\mu$ g C <sup>-1</sup> )	0.9±0.1 <sup>a</sup>	$0.8{\pm}0.2^{a}$	0.6±0.1 <sup>b</sup>	0.5±0.1 <sup>b</sup>

<sup>a, b, c, d</sup> significantly different (p< 0.05) groups using non-parametric ANOVA followed by post-hoc tests (Tukey).

<sup>e</sup> ice cover was > 25 % two months before our study

**Table 2.** Regression equations and statistics describing the relationship between transparent exopolymer particles (TEP) (dependent variable) and several independent variables across surface samples in the entire PEGASO cruise (note all variables were  $log_{10}$ -transformed). Regressions with p< 0.05 are highlighted. PVA: phage viral abundance. See Table 1 for abbreviations.

Dep. var.	Independent var.	$R^2$	р	Intercept	Slope	n
TEP	Temperature <sup>a</sup>	0.27	< 0.001	-78.1	32.8	66
	Salinity	0.25	< 0.001	-33.74	23.38	66
	Wind speed	0.08	< 0.05	2.13	-0.19	67
	Nitrate	0.10	< 0.05	2.84	-0.63	69
	Phosphate	0.14	< 0.05	2.18	-0.77	69
	Silicate	0.06	< 0.05	2.08	-0.08	69
	Solar radiation dose	0.24	< 0.001	2.65	-0.38	46
	Chl a	0.66	< 0.001	1.90	0.35	65
	Phyto B	0.66	< 0.001	1.17	0.43	67
	PP	0.43	< 0.05	1.67	0.25	19
	РНА	0.01	> 0.05			71
	РНР	0.18	< 0.001	2.07	0.25	68
	TVA	0.16	< 0.001	0.45	0.22	71
	LVA	0.08	< 0.05	1.27	0.12	71
	PVA	0.18	< 0.001	0.41	0.23	71
	POC <sup>b</sup>	0.70	< 0.001	-1.00	1.03	68
	PHA:Chl a	0.71	< 0.001	4.07	-0.39	64
	F <sub>v</sub> :F <sub>m</sub>	0.04	> 0.05			71
	Ddx:LHC	0.18	< 0.001	1.74	-0.42	68
	Prasinophytes <sup>c</sup>	0.68	< 0.001	1.86	0.14	37
	Chlorophytes <sup>c</sup>	0.17	< 0.05	1.72	0.23	38
	Dinoflagellates <sup>c</sup>	0.67	< 0.001	1.57	0.32	38
	Cryptophytes <sup>c</sup>	0.47	< 0.001	1.65	0.18	38
	Diatoms <sup>c</sup>	0.67	< 0.001	1.62	0.18	38
	Pelagophytes <sup>c</sup>	0.58	< 0.001	1.83	0.15	38

Haptophytes <sup>c</sup>	0.52	< 0.001	0.78	0.51	38	

 $R^2$ : explained variance; sample size; p: level of significance; n: sample size <sup>a</sup> Kelvin degrees, <sup>b</sup> POC as dependent variable, <sup>c</sup> ng Chl *a* L<sup>-1</sup> **Table S1.** Compilation of published transparent exopolymer particles (TEP) concentrations (mean  $\pm$  SE and ranges;  $\mu$ g XG eq L<sup>-1</sup>), chlorophyll *a* (Chl *a*) concentrations (mean  $\pm$  SE and ranges;  $\mu$ g L<sup>-1</sup>) and TEP:Chl *a* ratios (mean  $\pm$  SE and ranges;  $\mu$ g XG eq ( $\mu$ g Chl *a*)<sup>-1</sup>) in the Southern Ocean. Bdl: below detection limit.

Location	Comments	TEP	Chl a	TEP:Chl a	Reference
Anvers Island	Summer (November 1994-February	15 - >500	-	-	Passow et al. (1995)
	1995)				
	2- 6 m, coastal samples				
	Two blooms observed				
	(Cryptomonads and diatoms)				
Kita-no-seto Strait	Mid-January 1994	<b>26 <math>\pm</math>6-41<math>\pm</math>4 particles mL<sup>-1</sup></b>	-	-	Marchant et al. (1996)
	15 m				
Ross Sea	Summer (November, December	Surface 308 (0-2800)	3.61 (0.27-8.81)	89.1 (surface)	Hong et al. (1997)
	1994) 0-150 m,		(surface)		
D	Phaeocystis and diatoms bloom				
Bransfield Strait	Summer (13 December 1990- 3	$56.77 \pm 54.50$ (bdl-345.9)	$0.98 \pm 0.83 \ (0.05 - 4.81)$	51.0	Corzo et al. (2005)
Gerlache Strait	January 2000)	38 (0-283)	1.16	32.7	
Drake Passage	0-100 m	35 (0-157)	1.17	29.9	
Antarctic Peninsula	Summer (February 2005),	$15.4 \pm 10.0 \text{ (bdl-}48.9)$	0.01-5.36	$40.9 \pm 157.8$ (bdl-1492)	Ortega-Retuerta et al.
(all the study)	0-200 m		_		(2009b)
Bellingshausen Sea		14.3±9.5 (bdl-33.8)		84.2±257.5 (bdl-1492)	
Weddell Sea		16.3±12.5 (bdl-48.9)		9.8±7.4 (1.2–28.4)	
Bransfield and		15.8±8.9 (bdl-35.8)		15.0±20.4 (3.0–18.0)	
Gerlache Strait					
Southern Ocean (all	Summer (7 January-3 February	$102.3 \pm 40.4 (39.2 - 177.6)$	$2.36 \pm 1.92 \ (0.28 - 8.95)$	$79.3 \pm 54.9 \ (10.9-239.0)$	This study
transect)	2015)				
	4 m				
South Orkney Islands		$144.4 \pm 21.7 (97.8-177.6)$	$1.87 \pm 0.23  (1.58 - 2.21)$	$76.7 \pm 10.6 \ (60.4-97.5)$	
(NSO)					
southeast of the		48.1 ± 6.5 (39.2-63.8)	$0.32 \pm 0.06 \ (0.28 - 0.45)$	$153.4 \pm 29.8 \ (102.5 - 211.6)$	
South Orkney Islands					
(SSO)					1
northwest of South		$125.5 \pm 21.1 \ (93.6-157.4)$	$4.59 \pm 1.97 \ (1.92 - 8.95)$	32.3 ± 15.0 (17.6-70.0)	
Georgia (NSG)					
west of Anvers		$111.6 \pm 13.0 (90.5 \text{-} 125.0)$	$4.05 \pm 0.48$ (3.41-4.91)	$28.2 \pm 4.8 \ (18.4-34.9)$	
Island (WA)					





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Figure 1













![](_page_59_Figure_0.jpeg)

![](_page_60_Figure_0.jpeg)

![](_page_60_Figure_1.jpeg)

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![](_page_61_Figure_1.jpeg)

![](_page_62_Figure_0.jpeg)

Figure 6

![](_page_63_Figure_0.jpeg)

Figure S1

Figure S2

![](_page_64_Figure_1.jpeg)

Figure S3

![](_page_65_Figure_1.jpeg)

![](_page_66_Figure_0.jpeg)

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