



ECOSYSTEMS

Vertical distribution of the zooplankton in the Antarctic Peninsula during the austral summer of 2017

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Abstract: Zooplankton plays a crucial role as the primary consumers in the Southern Ocean and its ecological processes, particularly in the Antarctic Peninsula, influenced by regional glaciological and oceanographic changes. To assess the overall composition of these communities, vertical samples were collected at various depths using a Multinet at oceanographic stations in the Antarctic Peninsula during the XXXV OPERANTAR expedition in summer 2017. Abiotic data (temperature, salinity and chlorophyll-*a*) were collected using a CTD and a fluorometer. Organisms were identified to a higher level, with Chaetognatha and Euphausiacea identified to species whenever possible. Copepoda were the most abundant (73.4%), with Calanoida present in all samples and more abundant at 300m. Salpidae ranked as the second most abundant taxon (16.6%) up to 100m. Three species of Chaetognatha were identified, with *Eukrohnna hamata* being the most abundant, particularly at 300m. Two species of Euphausiacea were found, *Euphausia superba* and *Thysanoessa macrura*, with low abundances. Abiotic parameters showed significant relationships with the taxa. The region exhibits complex oceanography associated with zooplankton communities. The recorded data align with the zooplankton characterization of this region, uncovering a prevalence of Copepoda and surface abundant Salpidae, along with Chaetognatha (particularly *E. hamata*) and Euphausiacea.

Key words: abiotic parameters, community structure, key species, Southern Ocean.

INTRODUCTION

Zooplankton act as the primary link in the energy flow of the food web in the water column, playing a crucial role in primary productivity and other essential ecological processes, such as the biological carbon pump (Raymont 1983, Schnack-Schiel & Isla 2005, Steinberg & Landry 2017) particularly in polar regions (Infante 1988), these organisms are subject to environmental fluctuations, which are reflected in their abundances (Richardson 2008).

The specific conditions of the Southern Ocean (SO) directly influence zooplankton communities (Clarke & Peck 1991, Atkinson et al. 2004). Encompassing a vast area with

an extensive range of low temperatures and isolated at the southernmost extreme of the globe, the SO influences the global climate and is connected with all ocean basins (Lynne et al. 2011). The continuous eastward flow of the Antarctic Circumpolar Current (ACC) encircles the Antarctic Continent up to ~60°S. Recently it has been suggested that its temperature is rising and that the ice coverage is declining (Cook et al. 2005, Meredith & King 2005), influencing the stratification and vertical distribution of biological communities in the water column (Convey et al. 2014).

In the Antarctic Peninsula (AP) the vulnerability of ice fields and climatic and

oceanographic variability is evident (Meredith & King 2005), and the zooplankton populations have been the subject of many studies in order to evaluate the impact of these environmental changes (Steinberg et al. 2015, Tarling et al. 2018). The AP is a region of high biological productivity (Zhou et al. 1994), crucial for the reproduction and development of various zooplanktonic organisms, especially the Antarctic Krill (Brinton & Townsend 1991).

The order Euphausiacea plays a fundamental role in the Antarctic food web. Antarctic Krill, *Euphausia superba* is recognized as a key organism in biological structuring (Siegel 1988, 1989). They are found over approximately 32 million km² around the Southern Ocean (Siegel & Watkins 2016). It constitutes the primary food item for various Antarctic predators, such as seabirds, fish and large mammals (Newell & Newell 1963). In the AP region, other euphausiid species like *Euphausia crystallorophias* and *Euphausia frigida* can also be found (Montú et al. 1990, Pakhomov & Perissinotto 1994).

Chaetognatha is another important group in the SO, has a widespread distribution and high abundances (Hopkins 1985). They constitute the second most abundant group of zooplankton after Copepoda (El-Sayed 1985). *Eukrohnia hamata* is highlighted as the most abundant and frequent species in the AP (Oresland 1990). They are essential for the trophic energy transfer in polar regions (Oresland 1995), playing a significant role in the biological carbon pump during the austral summer (Raymont 1983, Giesecke et al. 2010) and serving as excellent bioindicators of water masses (Johnson & Terazaki 2004).

Since the 1970's, changes in some zooplankton populations in the SO have been reported (Flores et al. 2012), linked to alterations in oceanographic characteristics and the extent of sea ice (Atkinson et al. 2004). Zooplankton is considered an excellent

environmental bioindicator with its metabolic and reproductive processes associated with the water column temperature (Richardson 2008). Also, zooplankton's biogeographical distribution is linked to variations in oceanographic features such as the SO's fronts (Esquivel-Garrote et al. 2023, Esquivel-Garrote & Muxagata 2023) and is vulnerable to prevailing global and regional climate patterns. Changes in zooplankton communities and specific composition of groups such as Copepoda (Tarling et al. 2018), Chaetognatha (Takahashi et al. 2010), Euphausiacea (Atkinson et al. 2004), among others, have already been reported in the recent literature.

Considering the significance of studies on the zooplankton communities and their tremendous importance for the SO ecosystem and the global climate (Ratnarajah et al. 2023), this present study presents the composition and distribution of zooplankton, investigating the correlations of these assemblages with abiotic characteristics in different levels of the water column during the austral summer of 2017 in the AP region.

MATERIALS AND METHODS

The study area covers the AP region in the Atlantic sector of the SO, including the northeast sector of the Bellingshausen Sea (E#24) and the waters of the Gerlache Strait (E#07 e E#12) and Bransfield Strait (E#15, E#30, E#33 e E#49) (Fig. 1).

The zooplankton samples used in this study were collected during the XXXV OPERANTAR (Operação Antártica Brasileira, Brazilian Antarctic Operation) on board of the Polar Ship Armirante Maximiano during the austral summer of 2017. A total of seven oceanographic stations were conducted using a Multinet Midi (Hidro-bios 0.24m²) equipped with five 300 µm nets (Table I). The vertical sampling covered five strata of

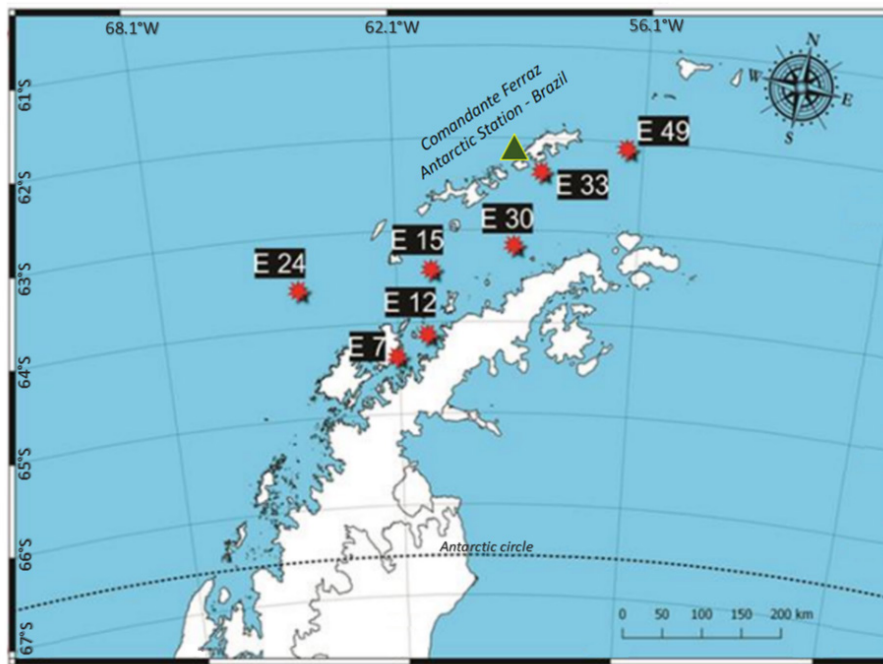


Figure 1. Map of the study area, indicating the sampling stations with the Multinet during the summer of 2017.

Table I. Oceanographic stations with Multinet during XXXV OPERANTAR. GS = Gerlache Strait; BS = Bransfield Strait; B = Bellingshausen Sea.

Station	Region	Date	Time	Latitude	Longitude	Local Depth	Max. Samp. Depth
#7	GS	16/02/2017	06:06	-64.570	-62.507	736m	500m
#12	GS	16/02/2017	17:30	-64.230	-61.587	457m	300m
#15*	BS	17/02/2017	04:38	-63.720	-61.391	757m	500m
#24	B	18/02/2017	08:20	-63.982	-65.007	440m	300m
#30	BS	19/02/2017	15:48	-63.277	-59.273	687m	500m
#33*	BS	21/02/2017	03:38	-62.604	-58.871	1501m	500m
#49*	BS	26/02/2017	01:36	-62.136	-56.818	778m	500m

(*) Samples collected during twilight/night period (sunrise: 05:38; sunset: 20:51).

100 meters (m) each (total of 500m sampled in the water column). Shallower locations were sampled with only three strata (300m depth sampled). The filtered volume of water was estimated using the Ocean Lab software (3.5.5.6) with an aid of a flowmeter attached to the net. All samples were preserved in a 4% formalin solution neutralized with borax according to Steedman (1976). Chlorophyll-a ($\mu\text{g L}^{-1}$), temperature ($^{\circ}\text{C}$) and salinity data were acquired

through a CTD (Conductivity, Temperature and Depth) and fluorimeter attached to the Multinet.

Once in the lab, all zooplankton organisms were counted and identified to major taxonomic groups using Bogorov chambers, a stereoscopic microscope and zooplankton identification guides (Brinton 1975, Kirkwood 1984, Mauchline & Fisher 1969, Mauchline 1971, 1980, O'Sullivan 1983, O'Sullivan & Hosie 1985). The subclass Copepoda were classified into different orders

according to Dussart & Defaye (1995) and sizes (large Copepoda equivalent to $>2000\mu\text{m}$; small Copepoda equivalent to $<2000\mu\text{m}$). Organisms from the Euphausiacea order and the Chaetognatha phylum were separated and identified at the lowest taxonomic level possible according to Baker et al. (1990), Bone et al. (1991), Lutschinger (1993) and Pierrot-Bults & Chidgey (1988). Samples containing large quantities of gelatinous organisms from the Salpidae order were meticulously examined and washed with 4% formalin, in order to remove organisms attached to their structures, and then quantified. Fragments of Siphonophorae from the Hydrozoa class found in the samples were summarized as one (1) organism. Pteropoda, Isopoda, Amphipoda, Mysida, Appendicularia, Foraminifera, Cnidaria, Siphonophorae, Echinodermata, Decapoda Cirripedia and non-identified (n.i.) larvae of Osteichthyes, were grouped in the category “others” due to their low abundances.

The results were recorded and tabulated as number of organisms in 1000m^{-3} . The abundances were converted to $\log_{10}(x+1)$ to stabilize the variances. Pearson correlation coefficients were used to measure the strength of the association of taxa with the average abiotic parameters for each depth stratum. All analyses presented here were performed using Quantum Gis (3.32.2) and Ocean Data View (v5).

RESULTS

Biological data

A total of 15,676 organisms were counted and grouped into 21 taxa (Table II). Copepoda was the most abundant group in all samples (73,46%), followed by Salpidae (16,60%), Chaetognatha (3,36%), Euphausiacea (2,62%) and Ostracoda (2,33%).

Overall, the dominant portion of Copepoda was reported below 100m, with higher abundances at 300m in stations E#07 (39,391 org. 1000m^{-3}), E#12 (57,043 org. 1000m^{-3}) and E#15 (50,565 org. 1000m^{-3}) (Fig. 2c). Calanoida represented 87% of the group. The order Cyclopoida were recorded in higher abundances at GS stations, a total of 28,984 org. 1000m^{-3} while Harpacticoida amounted to only 77 org. 1000m^{-3} at station E#15 at 200m. 56% of small Copepoda ($<2000\mu\text{m}$) belonged to the Cyclopoida order. It is noticeable that large Copepoda ($>2000\mu\text{m}$) showed higher abundances in the deeper layers for all stations, numerically more abundant at 300m at E#12 and E#15.

Salpidae appeared in 94% of the samples, being dominant in the first 100m (Fig. 2a). They were co-dominators with Copepoda at some stations at greater depths. Their highest abundances (exceeding 10,000 org. 1000m^{-3}) were observed at E#33. This group dominates the surface layers of the BS. Only 5% of the total Salpidae appeared in the samples E#07, E#12 and E#24.

Ostracoda and Polychaeta were quite frequent groups recorded in higher abundances at depths deeper than 300m. The maximum values of Ostracoda (2,307 org. 1000m^{-3}) and Polychaeta (1,730 org. 1000m^{-3}) occurred in BS at the deepest stratum (Fig. 2e). Only 107,1 Ostracoda 1000m^{-3} and 74,1 Polychaeta 1000m^{-3} were recorded at E#24 in the Bellinghausen Sea, the lowest occurrences throughout all samples. The “Others” were dominated by Appendicularia, Foraminifera and Pteropoda, that accounted for 52%, 26% and 9% of the total organisms in this category.

A total of 511 Chaetognatha were found in the samples, with 438 identified at species level. From those *Eukrohnia hamata* represented 94,4% of all the identified Chaetognatha, followed by *Pseudosagitta maxima* (1,14%) and

Table II. List of Taxa and their Frequency of Occurrence (FO%), Relative Abundance (RA%) and Maximum density of Organisms (org. 1000m⁻³). (n.i. = not-identified).

Taxa	FO (%)	RA (%)	Max. density (org. 1000m ⁻³)
Phylum Foraminifera	45	0.18	2,912
Phylum Cnidaria	40	0.02	158
Order Siphonophorae	68	0.00	21
Class Polychaeta	81	0.48	7,577
Class Pteropoda	32	0.17	1,090
Infra Class Cirripedia (Cipris n.i.)	10	0.09	596
Class Ostracoda	77	2.33	14,740
Sub Class Copepoda	100	73.46	463,376
Order Calanoida	100	64.11	404,358
Order Cyclopoida	93	9.32	58,941
Order Harpacticoida	3	0.01	77
Order Isopoda	13	0.03	250
Order Mysida	3	0.00	42
Order Amphipoda	64	0.31	1,959
Order Euphausiacea	48	2.62	16,576
<i>Euphausia superba</i> Dana, 1850	23	0.05	326
<i>Thysanoesa macrura</i> G.O. Sars, 1883	26	0.23	1,365
Order Decapoda (larvae n.i.)	3	0.01	77
Phylum Echinodermata (larvae n.i.)	6	0.01	82
Phylum Chaetognatha	74	3.36	21,195
<i>Eukrohnia</i> sp.	29	0.14	924
<i>Eukrohnia hamata</i> (Möbius, 1875)	68	2.74	17,957
<i>Pseudosagitta</i> spp.	10	0.01	79
<i>Pseudosagitta maxima</i> (Conant, 1896)	13	0.03	207
<i>Pseudosagitta gazellae</i> (Ritter-Záhony, 1909)	6	0.01	82
Chaetognatha n.i.	52	0.31	1,945
Class Appendicularia	97	0.37	5,856
Order Salpidae	93	16.60	104,742
Order Osteichthyes (larvae n.i.)	13	0.03	240

Pseudosagitta gazellae (0,45%). Overall, the highest abundances of Chaetognatha were collected in the 200m stratum (Fig. 3b) at stations E#07 (3,478 org. 1000m⁻³) and E#12 (2,782 org. 1000m⁻³) in the GS. *Pseudosagitta maxima* was found at three stations (E#07, E#12 and E#15)

below 200m. *Pseudosagitta gazellae* was found only at E#15 in the BS, with densities below 47 org.1000m⁻³.

Euphausiacea was found predominantly in the first 100m of the water column, with only two identified species. *Thysanoesa macrura* was

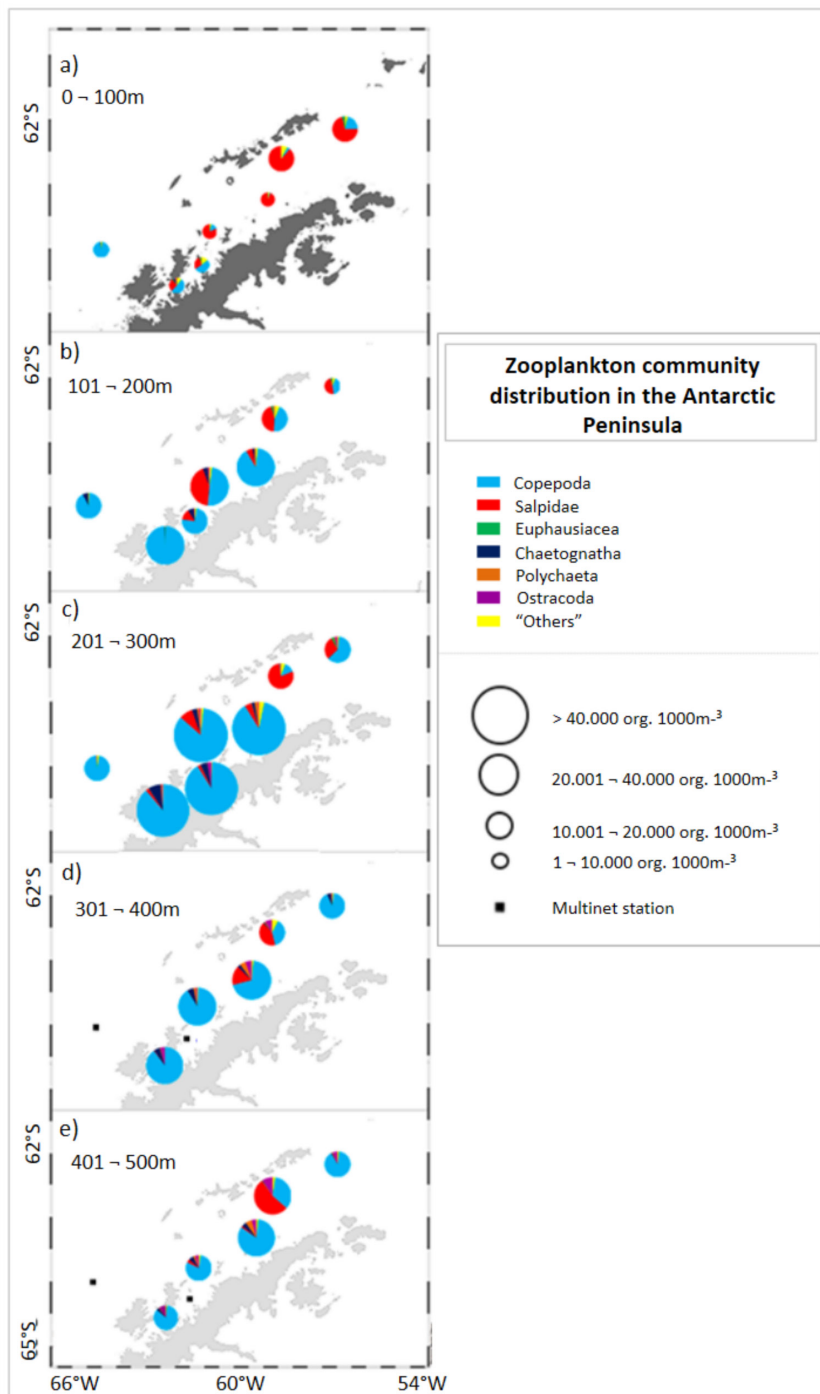


Figure 2. Spatial distribution of the zooplankton community across the Antarctic Peninsula. a) Stratum 0 ~ 100m depth; b) Stratum 101 ~ 200m depth; c) Stratum 201 ~ 300m depth; d) Stratum 301 ~ 400m depth; e) Stratum 401 ~ 500m depth.

the most abundant species, accounting for 82% Euphausiacea. Adult individuals of *T. macrura* were numerically greater in BS. Their maximum densities reached 360 org. 1000m⁻³ at E#49. Adult individuals of *E. superba* occurred only at E#07 (400m) and E#15 (100m) (Fig. 4). Larval stages

Calyptops I e II of *T. macrura* were recorded in the 200m strata at E#07 and E#33. Below 300m, only two specimens of *T. macrura* (E#33) and one of *E. superba* (E#07) were recorded, with numbers below 77 org. 1000m⁻³.

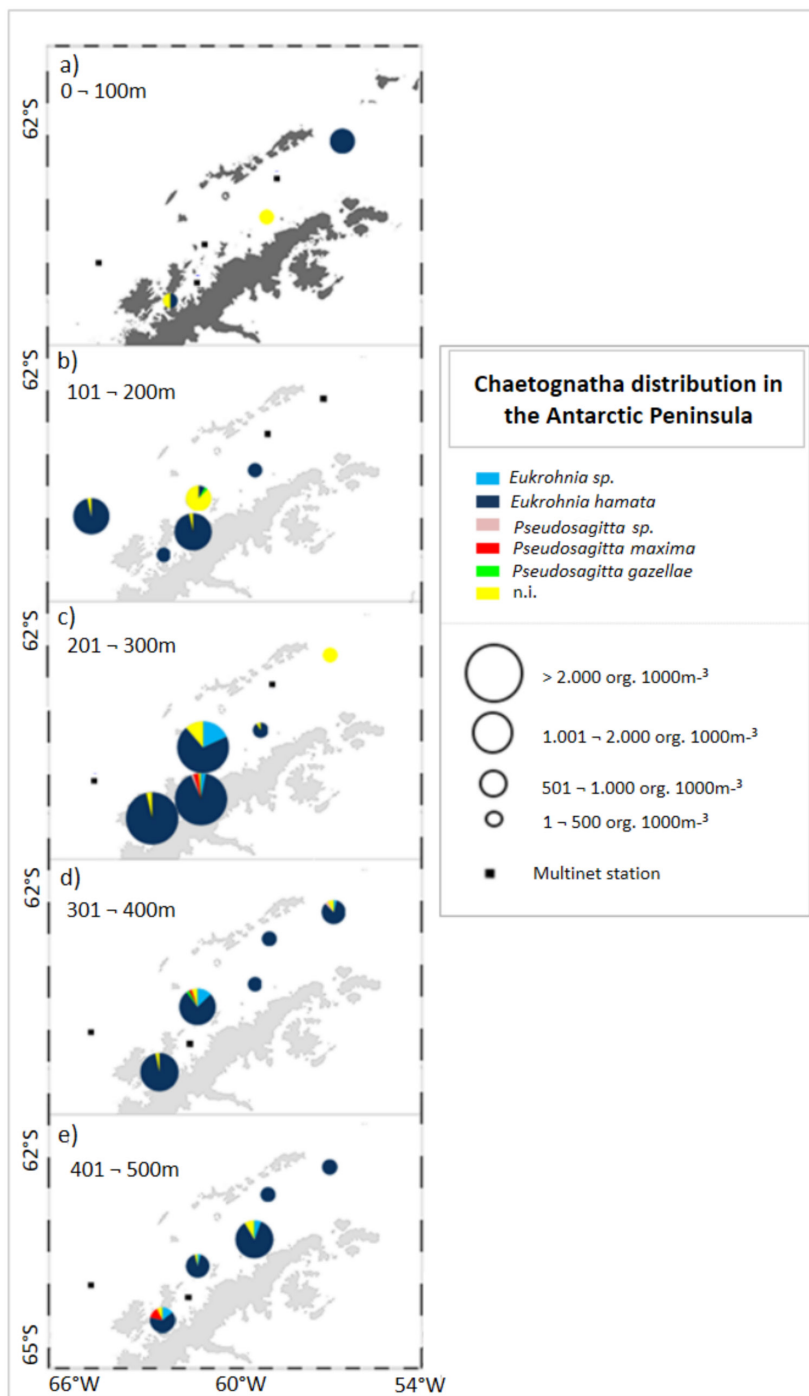


Figure 3. Spatial distribution of *Chaetognatha* species across the Antarctic Peninsula.

a) Stratum 0 ~ 100m depth;
 b) Stratum 101 ~ 200m depth;
 c) Stratum 201 ~ 300m depth;
 d) Stratum 301 ~ 400m depth;
 e) Stratum 401 ~ 500m depth.
 (n.i. = not-identified).

Environmental parameters and analysis

The highest temperature value was recorded in the surface layer at E#24 (3.19°C). Only BS stations reached near-minimum values of -1°C below 100m (E#25, E#33 and E#49). Low temperatures were recorded at E#07 in GS at

deep layers. Salinity values ranged between 33.3 and 34, with the highest values at E#24. Surface waters showed lower values, especially at E#12 in GS. The highest surface chlorophyll-*a* value was also recorded at E#12 (5.55µg L⁻¹), followed by

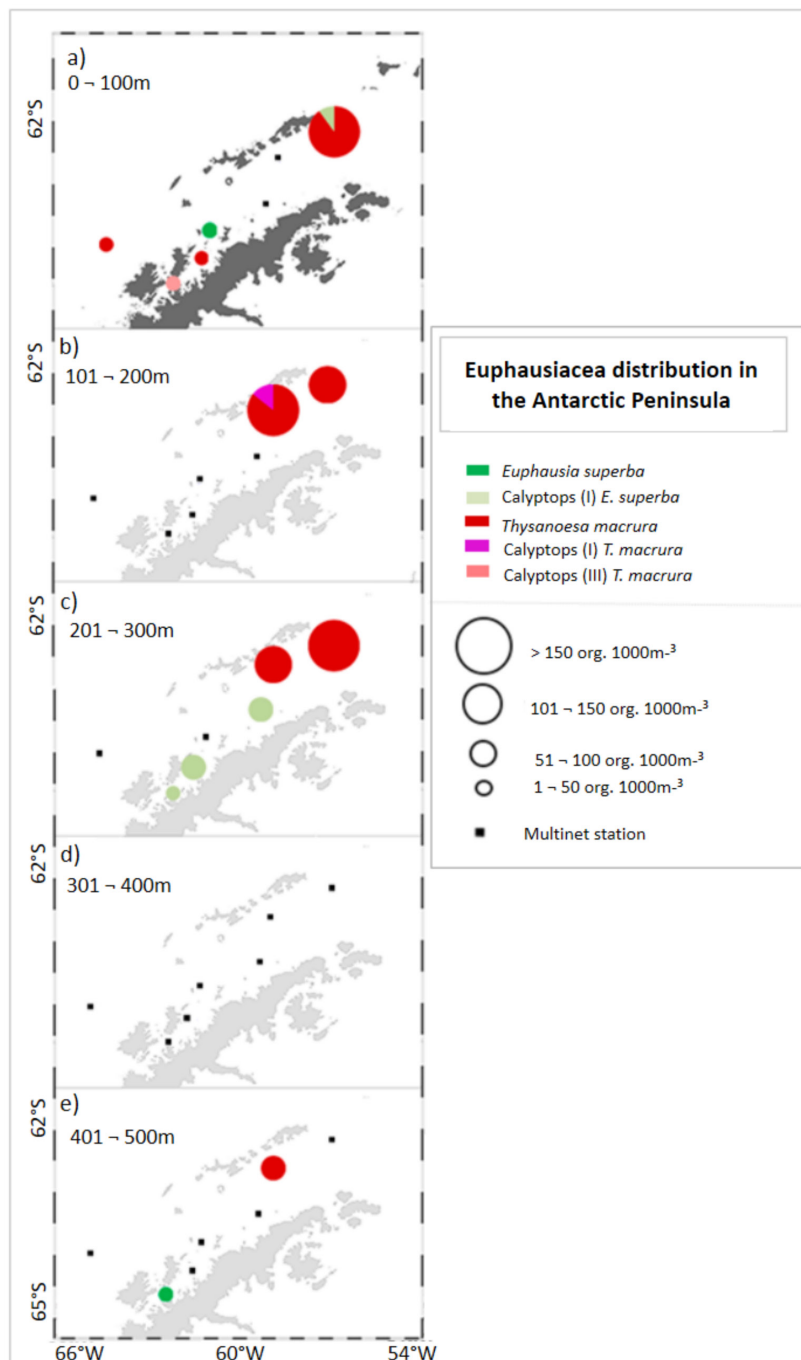


Figure 4. Spatial distribution of Euphausiacea species across the Antarctic Peninsula.

a) Stratum 0 ~ 100m depth;
b) Stratum 101 ~ 200m depth;
c) Stratum 201 ~ 300m depth;
d) Stratum 301 ~ 400m depth;
e) Stratum 401 ~ 500m depth.

E#33 ($1.53\mu\text{g}\text{L}^{-1}$) in the BS. Layers below 100m did not exceed $0.69\mu\text{g}\text{L}^{-1}$ of Chl-*a* (Fig. 5).

The Pearson Correlation Analysis (Table III) showed significant and negative correlations of environmental parameters with all major taxonomic groups identified. High and negative correlations were observed between

temperature, Chl-*a* and depth with Copepoda, Chaetognatha, Ostracoda, Polychaeta and "others". A weak and negative correlation appeared between salinity and *T. macrura*. Concentrations of Salpidae showed a significant and negative correlation with Copepoda and *P. maxima*. In contrast, high concentrations of

Table III. Pearson Correlation Analysis indicating the relationships between taxa and biological and abiotic parameters collected in the AP during the summer of 2017.

Taxa	T (°C)	S	Chl-a	Depth	Salpidae	Copepoda
Copepoda	-0.45**	NS	-0.73**	-0.72**	-0.33*	-
L Copepoda (>2000µm)	-0.40**	NS	-0.65**	-0.65	NS**	-
S Copepoda (<2000µm)	-0.39*	NS	-0.65**	-0.65**	-0.44**	-
Salpidae	NS	NS	NS	NS	-	-0.33*
Chaetognatha	-0.49**	NS	-0.53**	-0.53**	NS	NS
<i>Eukrohnia sp.</i>	-0.40*	NS	-0.34*	-0.34**	NS	NS
<i>Eukrohnia hamata</i>	-0.45**	NS	-0.55**	-0.55**	NS	NS
<i>Pseudosagitta spp.</i>	NS	NS	NS	NS	NS	NS
<i>Pseudosagitta máxima</i>	NS	NS	NS	NS	-0.35*	NS
<i>Pseudosagitta gazellae</i>	NS	NS	NS	NS	NS	NS
Chaetognatha n.i.	NS	NS	NS	NS	NS	NS
Euphausiacea	NS	-0.35*	NS	NS	NS	NS
<i>Euphausia superba</i>	NS	NS	NS	NS	NS	NS
<i>Thysanoesa macrura</i>	NS	-0.44*	NS	NS	NS	-0.40*
Ostracoda	-0.73**	NS	-0.66**	-0.66*	NS	NS
Polychaeta	-0.67**	NS	-0.50**	-0.50**	NS	NS
"Others"	-0.43**	NS	NS	NS	NS	NS

(*) p-values < 0.05; (**) p-values < 0.01; (NS) Not significant; (-) Not applicable; (n.i.) not-identified. T = Temperature; S = Salinity.

Copepoda exhibited significant and negative correlations with Salpidae and *T. macrura*.

DISCUSSION

The AP is characterized by a complex oceanography and frontal systems within a crucial transitional environment, where distinct water masses with varying abiotic parameters interact (for further information, see Orsi et al. 1995, Sangrá et al. 2011, 2017). This abiotic scenario influences the structure and composition of the zooplankton community (Atkinson et al. 2012, Su et al. 2022). Moreover, the GS and BS are described as the most productive regions in the AP (Zhou et al. 2002). Changes in the sea ice and

sea shelves patterns lead to increased primary production in the AP, especially in regions near the coast (Canals et al. 2016, Aracena et al. 2018).

The results presented in this study highlight Copepoda as the most abundant group in the AP, consistent with findings from Atkinson & Ward (2012) and Canals et al. (2016) which emphasize that Copepoda is the most significant and diverse group in this region. The Copepoda can increase their biomass and biodiversity by up to 80% during the austral summer in surface layers (Bradford-Grieve et al. 1999), particularly in coastal areas with high phytoplankton availability (Smetacek et al. 1990). Due to their varying sizes and feedings habits (Hopkins 1985)

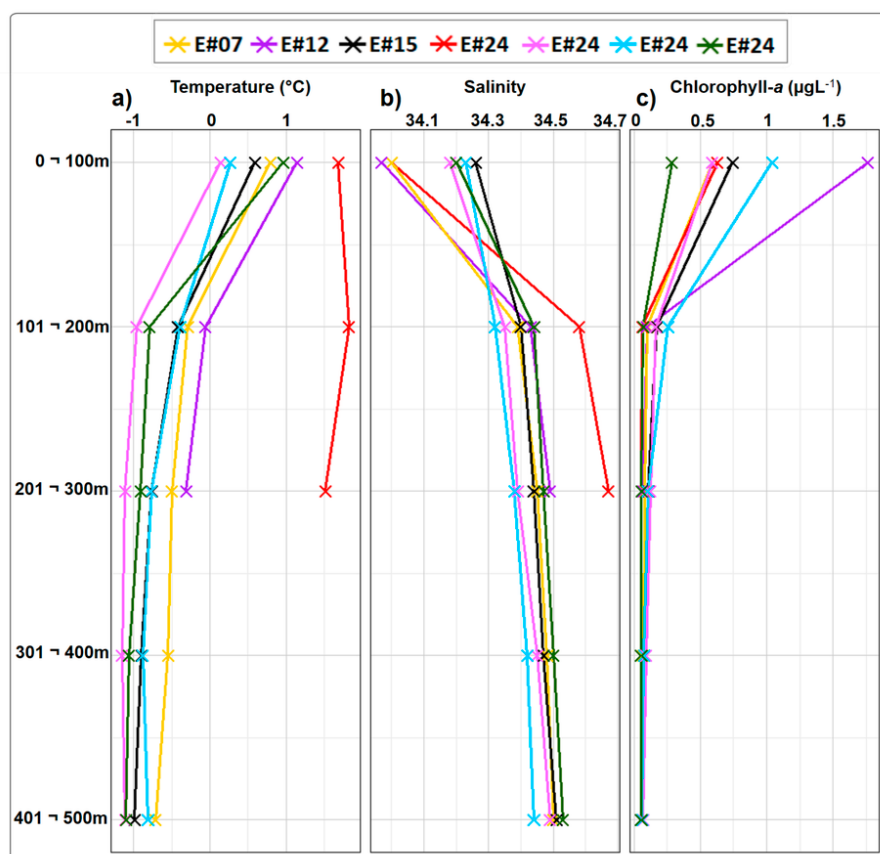


Figure 5. Abiotic parameters for each strata for the Multinet stations. a) Average temperature (°C); b) Average salinity; c) Average chlorophyll-a (µg L⁻¹).

they can contribute in different ways sustaining the AP's food web (Zhou et al. 2002).

Some studies indicate shifts in Copepoda size structure where smaller species dominate early in the summer while larger species prevail toward the end of stratification and into early fall (García et al. 2016, 2020). In our study, there was an evident dominance of the order Calanoida, consistent with other authors' findings (Tarling et al. 2004, Atkinson et al. 2012). Larger Calanoida Copepoda such as *Metridia gerlachei*, were the most abundant during the austral summer (Tarling et al. 2018). On the other hand, Marrari et al. (2011) emphasizes that smaller copepods from the order Cyclopoida are also abundant in this region. These smaller copepods persist for brief periods during their season, prevailing in waters with higher concentrations of chlorophyll-a (Fransz 1988) and playing a crucial role in the

energy flow of the AP's food web (Atkinson et al. 2004, Thompson et al. 2013).

Euphausiacea holds the title of keystone species in the Antarctic marine ecosystem (Siegel 1988, 1989). Atkinson et al. (2017) records that the highest densities of Euphausiacea in the AP occur up to 200m, decreasing when the depth increases. According to the literature, the most abundant species for the AP are *E. superba*, *E. frigida*, *E. crystallorophias* and *T. macrura* (Makarov et al. 1990, Ross et al. 1996, Atkinson et al. 2012). Here in our work we found two of these important species, *Euphausia superba* and *T. macrura*, which were also identified as the most abundant species by Steinberg et al. (2015) reporting great patches near the surface (Raymont 1983). Seasonal ecological successions are marked by these organisms throughout the AP (Atkinson et al. 2012, Conroy et al. 2020).

As our results pointed out that *T. macrura* prevailed throughout the AP, typically found in northern oceanic regions of AP and decreasing in their numbers further south (Atkinson et al. 2017). *Tysanoessa macrura* also plays a fundamental role in the diet of various fish, penguins, seals and whales in the SO (Marrari et al. 2011). Siegel et al. (2013) reported a significant concentration of juvenile Euphausiacea southern in the AP, coinciding with the higher phytoplankton concentration areas (Nordhausen 1992, 1994, Wiebe et al. 2011). Larval stages of *E. superba* are often observed in the GS and near the South Shetland Islands (Makarov et al. 1990, Atkinson et al. 2012, Rombolá et al. 2019), crucial areas for the growth of diverse populations (Zhou et al. 1994).

Several international projects such as the Convention for the Conservation of Antarctic Marine Living Resources (CCAMLR), funded in 1982, and Circumpolar Data on Antarctic Krill (KRILLBASE) (Atkinson et al. 2017), funded in 1990, monitor Euphausiid populations revealing various positive and negative anomalies in the AP (Steinberg et al. 2015). Atkinson et al. (2004) highlight a correlation between krill abundances and the extent of winter sea ice, which has been changing in recent years (Stammerjohn et al. 2008). Additionally, due their high mobility, vertical samplings with nets like Multinet may not be efficient for capturing these organisms, which could explain the low records in this study. Despite this, it is crucial to recognize that, although these populations are changing and being recorded in smaller numbers, potentially affecting the entire Southern Ocean ecosystem (Flores et al. 2012), our findings suggest that Copepoda can also play a significant role in the water column in terms of ecosystem functioning. They may even have a greater impact than krill, which are often regarded as the primary

contributors, thereby maintaining a vital balance within this environment.

Chaetognatha are the main consumers of Copepoda biomass (Oresland 1990, Giesecke et al. 2010). The Chaetognatha species recorded here are consistent with most publications for the AP (Pakhomov et al. 2002, Kruse et al. 2010) and are found to be the second most abundant group after Copepoda (Raymont 1983). They can represent portions between 5 to 30% of zooplankton biomass on de SO (Pakhomov et al. 2002). These voracious zooplankton predators dominate the SO, preferentially in the deeper layers of the water column (Duró & Gili 2001, Kruse et al. 2010). Larger species such as *P. maxima* and *P. gazellae* are often found in deeper layers at shelf regions (Kruse et al. 2010). *Eukrohnia hamata* is the most abundant and frequent specie in the AP, followed by *Solidosagitta marri* and *P. gazellae* according to Duró & Gili (2001). *Pseudosagitta maxima* goes next, in lower numbers (Oresland 1990). Giesecke et al. (2010) shows that *P. gazellae* contributes to 30% of the biomass predation in the SO, and it's essential in the biological pump during the austral summer. Steinberg et al. (2015) point out a trend of decreasing of Chaetognatha abundances in the AP related to the reduction in the extent of sea ice.

Organisms of the Salpidae order play a significant role in the structure of zooplankton communities, standing out for their efficiency in filtering particles on the water column (Raymont 1983) and having the potential to consume large concentrations of phytoplankton in the surface layers (Atkinson et al. 2012). Reviewed studies indicate an increase in the capture of these organisms around the SO (Atkinson et al. 2004, Pakhomov et al. 2006, Bernard et al. 2012). McClintock et al. (2008) explains the importance of the ACC flow, carrying Salpidae to the western shelf of the AP. The most common species in

the SO is *Salpa thompsoni*, representing about 95% of the group and reported in abundance in the AP (Perissinotto & Pakhomov 1998, Słomska et al. 2021). Positive and negative anomalies in Salpidae abundance are reported in Steinberg et al. (2015)'s review, related to changes in SO oceanographic characteristics, such as variations in sea surface temperature, sea ice extent and climate patterns (Barlett et al. 2018, Słomska et al. 2021). The highest concentrations of Salpidae recorded in this data set corresponded to low concentrations of Chl-*a* in surface layers, Raymont (1983) suggest that, in polar regions, these efficient filters have effective biological mechanisms to maintain abundance in the water column, like their alternate life cycles.

The contrast between “salp years” and “krill years” in the AP region has been consistently documented in numerous publications (Perissinotto & Pakhomov 1998, Pakhomov et al. 2002, Ross et al. 2014). Raymont (1983) already reported a competitive relationship between these two efficient Antarctic filter feeders. The literature extensively discusses the alternation of *S. thompsoni* and *E. superba* abundances (Atkinson et al. 2004, 2017, Steinberg et al. 2015) and the coexisting groups in time and space (Pakhomov et al. 2002, Ross et al. 2008, Bernard et al. 2012) in the SO. Furthermore, efforts are made to explain the factors that drives this relationship (Pakhomov et al. 2002) such as environmental variations in the AP's complex oceanography, along with changes in the phytoplankton composition (for example Mendes et al. 2013). Henschke et al. (2016) rethink the act of Salpidae with the advance of climate change in the SO, emphasizing their crucial role in the vertical carbon flux and potential fecal pellet production, enhancing the overall biological pump around the SO (Henschke et al. 2016, Iversen et al. 2016).

The study recorded the vertical distribution of the main zooplankton groups, with emphasis on Chaetognatha and Euphausiacea, for an extremely complex region that has shown clear evidence of climate variations in the global scenario (Cook et al. 2005, Meredith & King 2005). The results indicated the dominance of Copepoda throughout the water column, alternating on the surface with high abundances of Salpidae. Overall, Chaetognatha were frequent and abundant, with three species identified, *E. hamata* particularly prevalent. Euphausiacea were recorded in low abundances and low species diversity, appearing punctually in their larval stages. Further studies in time and space are necessary for understanding and describing these important Antarctic communities and their responses to climate change, facilitating the assessments on global marine biodiversity (Ratnarajah et al. 2023) especially in the SO.

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Both authors contributed to the study conception and design. CML data collection and analysis, writing, preparation of figures and tables, reviewing and editing. EM data collection and analysis, supervision, reviewing and editing. Both authors read and approved the final manuscript.

