Earth System
Science
Data

1 **TITLE**

2 *The Italian contribution to the Synoptic Arctic Survey programme: the 2021 CASSANDRA cruise*
3 *(LB21) through the Greenland Sea Gyre along the 75°N transect*

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18 **ABSTRACT**

19 In September 2021, as part of the Italian Arctic research programme, a multidisciplinary cruise
20 along the 75th parallel north through the Greenland Sea Gyre was conducted aboard the Italian
21 icebreaker Laura Bassi as part of the CASSANDRA project, which also contributed to the Synoptic
22 Arctic Survey (SAS) 2020/22. The cruise took place during the period of the lowest summer sea ice
23 extent ever measured. The data show strong horizontal gradients with temperatures between 1.5 °C
24 and 9.0 °C and salinity between 30 and 35. Warm and salty Atlantic Water (AW, $\theta$ > 3.0 °C, S
25 around 35) dominates on the eastern side of the transect in the upper 500 m with surface
26 temperatures of 4.5–9.0 °C, while Polar Water (PW, $\theta$ < 0 °C, S < 33) occupies the surface layer
27 (50–80 m) in the west. The intermediate layer (100–500 m) consists of mixed water, and below 500
28 m the deep water of the Greenland Sea and the Norwegian Sea predominates. The oxygen
29 enrichment is higher in the intermediate layers, while the values in deep layers and western regions
30 are lower (< 300 µmol kg$^{-1}$). A stratified upper layer (30–50 m deep) with low surface nutrients,
31 especially nitrate, is observed, while an accumulation of silicate occurs in deep water masses. The
32 surface water in the eastern part of the transect has high pH$_T$ and total alkalinity values due to
33 photosynthesis and the presence of salty AW, while the fresh PW in the west has a lower alkalinity.
34 Respiratory activity and organic matter concentrations (particulate/dissolved organic carbon) vary
35 horizontally at the surface, decrease with depth, and increase slightly near the seafloor. A west-east
36 gradient is also observed for $\delta^{18}$O and $\delta$D, with the ratios indicating the influence of freshwater at
37 the surface near the Greenland coast. The abundance of prokaryotes decreases from the photic zone

38   (< 100 m depth) to the sea floor. Carbohydrates and carboxylic acids are identified as well-utilised
39   polymers at every station and in every layer. Overall, the microbial enzyme patterns show a decrease
40   from the surface to deeper layers, with some hotspots of metabolic activity at 20-40 m and in the
41   aphotic layer. The enzyme patterns vary spatially, with activity peaks at the ends and in the middle
42   of the transect. Phytoplankton biomass, measured as chlorophyll-*a*, varies across the transect, with
43   higher values at its extremities. Micro-phytoplankton fraction dominates in PW, replacing the nano-
44   phytoplankton fraction, which is prevalent in AW, even at the interface between the two water
45   masses. Data of phytoplankton communities show low abundances and a dominance of nano-sized
46   organisms, with diatoms being more abundant in the western part. Microzooplankton represents an
47   important fraction of the planktonic community in this area, with tintinnids being the most important
48   groups along the transect. Micrometazoans and aloricate ciliates are more abundant in the AW,
49   resulting in higher biomass values at the eastern stations. Copepods are the most abundant
50   mesozooplanktonic taxon both at the surface and in the upper 100 m water layer (97% and 94% of
51   total mesozooplankton abundance, respectively), mainly represented by the genus *Calanus*.

52   The data are publicly available at the Italian Arctic Data Centre (IADC), see section Data
53   availability.

## 1   INTRODUCTION

55   The Greenland Sea, in the north Atlantic, is a region of deep ocean convection that contributes to
56   the Atlantic Meridional Overturning (AMOC) and the exchange of water masses between the
57   Atlantic and Arctic Oceans. Its sensitivity to climate change remains uncertain, as the ecosystems
58   of the subarctic Atlantic are particularly sensitive to global warming (Whitt, 2023). In fact, the
59   Greenland Sea serves also as a hub for heat, salts, nutrients, carbon, and organisms between the
60   Arctic, subarctic, and lower latitudes. Arctic and subarctic regions have experienced warming twice
61   as fast as global warming over the past 50 years (Rantanen et al., 2022). This has led to significant
62   environmental changes, including increasing wetness, reduced sea ice thickness and coverage,
63   changes in snow cover, thawing of permafrost, and melting of the Greenland ice sheet (Carmack et
64   al., 2015; Polyakov et al., 2017, 2023; Babb et al., 2023). These changes have created a positive
65   feedback loop known as "Arctic amplification" which is likely to intensify in the future. Despite its
66   crucial role in the global climate system, the Arctic Ocean remains poorly understood due to its
67   remote location, harsh weather, and seasonal ice cover.  The Arctic Ocean receives heat through the
68   inflow of Atlantic Water (AW, with Temperature (T) > 0 °C). AW flows northward transported by
69   the Norwegian Atlantic Current (NAC) and the West Spitsbergen Current (WSC) along the eastern
70   slope of the North Atlantic and crosses the Greenland Sea and Fram Strait, where it is partly
71   deflected by the local circulation (Fig. 1). On the other hand, the Arctic outflow of Polar Water
72   (PW, T < 0 °C), together with the sea ice export from the Arctic, is driven by the East Greenland
73   Current (EGC, Chatterjee et al., 2018). Greenland freshwater flux shows a large seasonal variation,
74   with peaks in July (4-6 times higher than in winter), but also consistent increase since the 2000s
75   (Dukhovskoy et al., 2019). Changes in annual and multi-year sea ice trends are also an important
76   factor to consider when analysing the physical and biogeochemical conditions in the Greenland Sea.
77   Sea ice extent and thickness in the Arctic regions have continuously decreased over the last four
78   decades, and significant changes have also been observed in the Fram Strait since 2015 (de Steur et
79   al., 2023). Since 2020, however, sea ice extent in the Fram Strait and marginal seas has shown a
80   slight recovery in seasonal winter maxima (Onarheim et al., 2024).  Open-ocean convection also

81  occurs in the Greenland Sea during winter seasons. It is thought to represent a significant proportion
82  of dense water production for the regions, even though a large variability in this process has been
83  observed in recent decades, including changes in the depth of convection (Simpkins, 2019; Brakstad
84  et al., 2019). Overall, the predominant atmospheric conditions over the Arctic are characterised by
85  the presence of high pressure (i.e., Polar High) over the western Arctic and low pressure over the
86  Siberian region, which trigger the main anticyclonic wind regime. However, after 2007 a secondary
87  dipole, characterised by higher sea level pressure over the Beaufort Gyre and the Canadian
88  Archipelago along with lower sea level pressure over the Siberian Arctic, became dominantly
89  positive, favouring reduced flows into the Arctic through the Fram Strait along with enhanced
90  inflows through the Barents Sea Opening (Polyakov et al., 2023). Consequently, from 2007 to 2021,
91  the predominant cyclonic atmospheric regime over the Arctic Ocean pushed large amounts of
92  freshwater from the Siberian shelves into the Beaufort Gyre (Polyakov et al., 2023).

93  The cyclonic Greenland Sea Gyre (GSG) in the central Greenland Sea, which is mainly driven by
94  large-scale cyclonic winds, contributes to the regulation of the inflow and outflow of AW and PW
95  between the Atlantic and Arctic Oceans (Chatterjee et al., 2018). The GSG, and the North Atlantic
96  subpolar gyre (SPG), south of Iceland, have large implications for the large-scale changes in the
97  subpolar and polar marine environment. A strong GSG triggers a northwestward shift of the
98  subpolar front, which intensifies the poleward transport of Atlantic water towards the Fram Strait,
99  Barents Sea and Arctic Ocean (Chatterjee et al., 2018). In contrast, a weak phase (i.e., a negative
100 index, see Fan et al., 2023) in the SPG enables the northward expansion of subtropical warm and
101 saline waters, while a strong SPG feeds cold and fresh subpolar waters into the Atlantic inflow.

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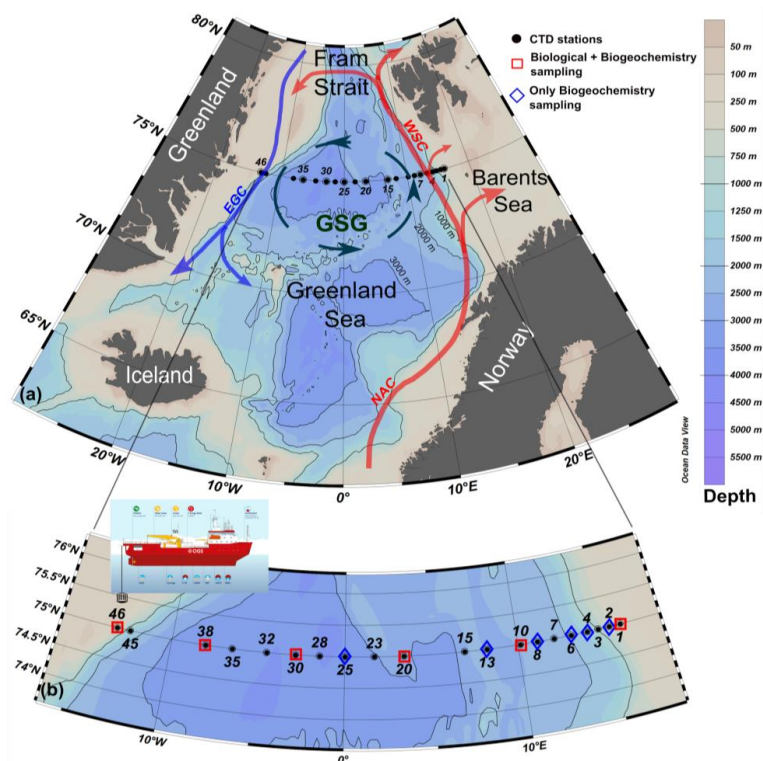


**Figure 1** - **(a)** Schematic of the general circulation in the Greenland Sea (**GSG**-Greenland Sea Gyre; **WSC**-West Spitsbergen Current; **EGC**-East Greenland Current; **NAC-** Norwegian Atlantic Current). **(b)** Distribution of hydrological (CTD) stations conducted during the LB21 CASSANDRA cruise (29 August -14 September 2021) with the positions of the physical (black dots), biogeochemical and biological stations (blue and red symbols).

A time lag of 3-5 years is expected for the thermohaline anomalies to propagate in the Nordic seas (Fan et al., 2023) from the sub polar regions. In other words: while the GSG regulates the AW inflow into the Arctic, the SPG modulates the proportion of subtropical and subpolar waters moving at high latitudes. Wind forcing and heat loss combine to drive the full variability of the flow and water mass transformations in the region (Smedsrud et al., 2022).

The anomalous inflow of AW from the Nordic seas and the subpolar regions is referred to a process called Atlantification, which has various direct and indirect effects on the Arctic environment. This process has intensified since the 2000s, leading to, among other effects, increasing anomalies in temperature and salinity in the upper layer, a reduction in sea ice cover, reduced stratification of the upper ocean, increased primary production and a shift of the summer bloom to earlier periods, changes in the phenology, distribution and community composition of zooplankton, and spreading of subarctic species (Polyakov et al., 2017; Ingvaldsen et al., 2021; Csapò et al., 2021). Microbial communities are also affected by the co-occurrence of Atlantification and Arctic amplification, as pointed out by recent observations (Ahme et al., 2023; Priest et al., 2023). Microorganisms are pivotal drivers in all the earth ecosystems, as primary degraders of organic matter and main players in nutrient cycling. They are thus particularly sensitive to the external environmental conditions and

125 as such, play a role of optimal sentinels of global changes and trends (Caruso et al., 2016). Distinct
126 bacterial communities have been reported between Atlantic- and Arctic-derived waters (Carter-
127 Gates et al., 2020), and a direct effect of organic matter dissolution from the sea ice on the microbial
128 diversity has been demonstrated, by documenting the occurrence of diverse bacterial assemblages
129 between sea-ice and seawater (Yergeau et al., 2017). The Greenland Sea, like other Nordic Seas, is
130 a sink for atmospheric $CO_2$ during the year (Skjelvan et al., 2005). The annual flux of $CO_2$ absorbed
131 by the sea has been estimated at 53 gC m$^{-2}$ yr$^{-1}$. Of this amount, about half was attributed to the flux
132 caused by heat loss and the other half to biological production (Anderson et al., 2000). Higher
133 estimates of annual fluxes, ranging from 40 to 85 gC m$^{-2}$ yr$^{-1}$, were presented by Skjelvan et al.
134 (2005 and reference therein). Total carbon in surface waters varies seasonally because of physical
135 and biological processes that influence the amount of carbon exported to deep waters (von
136 Bodungen et al., 1995; Noji et al., 1999). The mixing of the open ocean contributes to the transport
137 of carbon from the surface to the deep interior. As a result of the increasing $CO_2$ input from the
138 atmosphere, the pH in the Nordic Seas has decreased by ~0.0028 units per year in the period 1981-
139 2019 (Frasner et al., 2022).

140 Here, we present data and main results obtained in the framework of the CASSANDRA (*advanCing
141 knowledge on the present Arctic Ocean by chemical-phySical, biogeochemical and biological
142 obServAtioNs to preDict the futuRe ChAnges*) project funded by the Italian Arctic Research
143 Programme (*https://www.programmaricercaartico.it/*). CASSANDRA contributed to the Synoptic
144 Arctic Survey (SAS) by investigating the historical zonal transect at 75°N through the Greenland
145 Sea Gyre during the summer of 2021 (29 August - 14 September). The SAS initiative aims to
146 quantify the current state of the Arctic Ocean and its changes, focusing on water masses,
147 ecosystems, and the carbon cycle (see https://synopticarcticsurvey.w.uib.no/). SAS sees the
148 participation of 11 countries in 25 Arctic cruises.

## 2    DATA and METHODS

150 The LB21 CASSANDRA cruise (hereinafter, LB21 cruise) was carried out between 29 August 2021
151 (Longyearbyen, Svalbard) and 14 September 2021 (Bergen, Norway) on board the icebreaker Laura
152 Bassi (https://www.ogs.it/en/research-vessel-laura-bassi). All methods used for the CASSANDRA
153 activities were in line with the recommendations of the SAS and Go-Ship programme
154 (https://www.go-ship.org/), the latter including the 75°N transect. This approach was chosen to
155 obtain data that were as comparable as possible to other the SAS programme cruises. During the
156 LB21 cruise, a total of 28 vertical Conductivity-Temperature-Depth (CTD) profiles were conducted
157 at 20 hydrological stations (some stations include repeated casts). Of these, biogeochemical and
158 biological data were also collected at 12 and 6 stations, respectively (see Fig. 1). Unfortunately,
159 some technical problems and 2 days of adverse meteorological conditions in the middle of the cruise
160 prevented the completion of all planned stations. Some of the figures presented here were created
161 with Ocean Data View (Schlitzer, 2024).

### 2.1 Hydrographic data

163 All hydrographic profiles were recorded with a Seabird SBE911plus, equipped with some additional
164 sensors. CTD measurements provide vertical profiles of temperature (T) and conductivity (C)
165 approaching the seafloor to ~ 5–10 m, depending on sea conditions. Potential temperature ($\theta$),

166   salinity (S), and potential density anomaly ($\sigma_\theta$, referred to 0 dbar) were calculated from *in situ* data
167   using the MATLAB toolbox TEOS-10 (Gibbs SeaWater Oceanographic Toolbox) including the
168   thermodynamic equations for seawater (http://www.teos-10.org/software.htm). Dissolved Oxygen
169   concentration was measured using an SBE43 sensor. T and S data were quality checked and
170   averaged every 1 dbar, with overall accuracy within ± 0.002 °C for T, ± 0.005 for S and 2% of
171   saturation for oxygen. Fluorescence and turbidity in the water column were measured with optical
172   sensors WET Labs ECO-AFL/FL. Water sampling was carried out using a rosette system equipped
173   with 24 10-liter Niskin bottles.

174   **2.2 Biogeochemistry data**

175   The chemistry of seawater was investigated at 12 stations from discrete water samples (Fig. 1) by
176   measuring dissolved oxygen, nutrients (nitrite, nitrate, phosphate, silicate), dissolved and particulate
177   Carbon (DOC and POC), total dissolved nitrogen and phosphorus (TDN and TDP); $\delta^{18}O$ and $\delta D$ of
178   $H_2O$; inorganic carbonate system characterization by total alkalinity, $pH_T$ and derived parameters.
179   The dissolved oxygen concentration (DO) was determined by a potentiometric Winkler titration
180   (Oudot et al., 1988; Grasshoff et al., 1999). Samples for inorganic nutrients (nitrite - $NO_2$, nitrate -
181   $NO_3$, ammonium - $NH_4$, phosphate - $PO_4$ and silicate - $Si(OH)_4$) were collected and analysed as
182   described in Ingrosso et al. (2016a) using a four-channel Continuous Flow Analyzer QuAAtro (Seal
183   Analytical Inc., Mequon, WI, USA) autoanalyzer. Detection limits were 0.01 µmol $L^{-1}$, 0.02 µmol
184   $L^{-1}$, 0.03 µmol $L^{-1}$, 0.01 µmol $L^{-1}$ and 0.01 µmol $L^{-1}$ for $NO_2$, $NO_3$, $NH_4$, $PO_4$ and $Si(OH)_4$,
185   respectively. DON and DOP were calculated as the difference between dissolved total phosphorus
186   (TDP) and $PO_4$ and between dissolved total nitrogen (TDN) and dissolved inorganic nitrogen (DIN
187   $= NO_3 + NO_2 + NH_4$), respectively. TDP and TDN were determined as $PO_4$ and $NO_3$, respectively,
188   after quantitative conversion to inorganic P and N by persulfate oxidation (Hansen and Koroleff,
189   1999). The accuracy and precision of the analytical procedures are annually checked through the
190   quality assurance program (AQ1) QUASIMEME Laboratory Performance Studies (Wageningen,
191   The Netherlands) and internal quality control samples were used during each analysis. Samples for
192   $pH_T$ analysis were collected and analysed on board as described in Ingrosso et al. (2016b) and Urbini
193   et al. (2020) using a spectrophotometer (Varian Cary 50 UV-visible). The results were expressed
194   on the 'pH total hydrogen ion scale' ($pH_T$) at 25°C, with a reproducibility of 0.0048, determined by
195   replicates from the same Niskin bottles. To measure total alkalinity ($A_T$, µmol $kgSW^{-1}$), water
196   samples were collected and analysed as described in Ingrosso et al. (2016a) and Urbini et al. (2020)
197   using the seawater certified reference materials (CRMs) for $TCO_2$ and $A_T$ supplied by Prof. A.G.
198   Dickson, Scripps Institute of Oceanography, USA (Batch number #185) to calibrate HCl for
199   analyses. $A_T$ precision and the accuracy were less than ± 2.0 µmol $kg^{-1}$, assessed by analysing
200   CRMs. All other carbonate system parameters, including $pH_T$ at *in situ* temperature, seawater partial
201   pressure of $CO_2$ ($pCO_2$), $TCO_2$, and aragonite saturation state ($\Omega_{ar}$) were calculated using the
202   CO2Sys program as described in Urbini et al. (2020). The estimated uncertainties were: ±0.005 for
203   $pH_T$ at in situ temperature, ±7.8 µatm for $pCO_2$, ±5.6 µmol $kg^{-1}$ for $TCO_2$ and ±0.12 for aragonite
204   saturation state. Samples for stable isotopic composition of dissolved inorganic carbon ($\delta^{13}C$-DIC)
205   were collected in 12-mL Exetainer® (Labco Limited, Ceredigion, UK) evacuated glass tubes,
206   containing 2 µL of saturated $HgCl_2$. Samples were stored at 4 °C in the dark until analysis was
207   performed as described in Relitti et al. (2020). To determine the optimal extraction procedure for

208    water samples, two standard $Na_2CO_3$ solutions were prepared with a known $^{13}C$ value of $-10.8 \pm$
209    0.1 ‰ (k=1) and $-4.2 \pm 0.1$ ‰ (k=1), respectively. The stable isotopic composition of dissolved
210    inorganic carbon is given conventionally in δ-notation in per mil deviation (‰) from the Vienna
211    Peed Dee belemnite (VPDB) standard.

212    For POC concentrations filters were freeze-dried and subsampled by punching 18% of the 45 mm
213    filter area and fitted into silver capsules/boats. The subsamples were treated with 1M HCl to remove
214    inorganic carbon and then placed into an oven at 60 C until dry. Afterwards, the samples were
215    wrapped in tin capsules/boats to aid combustion during analysis. The samples were analyzed with
216    a Thermo Fisher elemental analyzer (FLASH 2000 CHNS = O) coupled with a Thermo Finnigan
217    DeltaC isotope ratio mass spectrometer (IRMS).

218    Water samples for DOC analyses were filtered aboard, immediately after collection, through
219    precombusted (4 h at 480 °C) Whatman GF/F glass fibre filters (0.7 um nominal pore size).
220    Filtration was performed by using a disposable polycarbonate syringe and a polypropylene 25 mm
221    filter holder (Nuclepore), to prevent atmospheric contamination. Filtered samples were stored in 25
222    mL high density polyethylene (HDPE) bottles (previously treated with $HNO_3$ 1.2 M at 50 °C for 1
223    h) which were quickly frozen in an aluminium block at -20 °C. In the laboratory, filtered samples
224    were thawed, acidified to pH = 2 with ultrapure HCl and purged with $N_2$ for about 10 min to remove
225    inorganic carbon, as outlined in Pettine et al. (2001). Dissolved organic carbon (DOC) was assayed
226    by high temperature catalytic oxidation (HTCO) using a Shimadzu TOC-L series analyser.

227    Samples for stable isotope ratio measurements in seawater were collected in 5 mL amber glass
228    bottles. The bottles were filled to avoid the presence of air, immediately sealed and stored at a
229    temperature of 4 °C until the analyses. Analyses were performed by means of a Thermo DeltaV-
230    Advantage mass spectrometer equipped with a gas-bench. For the analysis, a quantity of 200 $\mu L$ of
231    water sample was used in a glass vial firmly closed with a membrane cap. The samples were flushed
232    with a gas mixture of 2% H in helium with a purity of 99.998 and analysed to determine δD.
233    Immediately after the δD analyses, the same samples were flushed with a gas mixture of 0.4% $CO_2$
234    in helium with a purity of 99.998 and analysed to determine $\delta^{18}O$ after 20 hours of equilibration.
235    All samples were measured at least in triplicate and the isotopic data are the mean of consistent
236    results. The standard deviation of our results is always 0.50‰ and 0.06‰ or better for δD and $\delta^{18}O$,
237    respectively. The SMOW2 and SLAP2 isotopic standards were used as a reference together with a
238    'home-made' standard. The home-made standard is analysed every 3 measurements (3 replicates of
239    a single sample) to assess the stability of the measurements. The isotopic composition is expressed
240    as:

241                                  $\delta X = [(R_s-R_r)/R_r] \times 1000$

242    where δX represents δD or $\delta^{18}O$, R=D/H or $^{18}O/^{16}O$ in the sample (s) and in the reference (r),
243    respectively.

### 2.3 Phytoplankton data

#### 2.3.1 Total and size-fractionated chlorophyll a (chl-a)

Chlorophyll *a* (chl-*a*) and phaeopigment (phaeo) concentration were measured fluorometrically. Samples were filtered on Whatman GF/F glass-fiber and polycarbonate membranes to separate three size fractions: micro-phytoplankton (> 10 µm), nano-phytoplankton (10–2.0 µm) and pico-phytoplankton (2.0–0.45 µm) as reported in Decembrini et al. (2021).

#### 2.3.2 Utermöhl phytoplankton

For the determination of Utermöhl phytoplankton (i.e., all species/taxa detectable by light microscopy, thus excluding prokaryotic phytoplankton and the majority of picoeukaryotes < 1 µm), 500-mL water samples were collected in opaque polyethylene bottles and immediately fixed with pre-filtered and neutralised formaldehyde (1.6% final concentration) (Throndsen, 1978). Inverted microscopes equipped with phase contrast (Zeiss Axiovert 200M and Leica DMi8) were used for the taxonomic identification analysing a variable volume of sample (10-50 mL), according to the Utermöhl method (Zingone et al., 2010). Counting was performed along transects across the microscope chambers at a magnification of 400x for small (5-20 µm) or very abundant species and observing half of the sedimentation chamber at a magnification of 200x for less abundant microphytoplankton (> 20 µm). The abundance was expressed as the number of cells per liter (cells L$^{-1}$). The minimum value of the counted cells was 200 cells per sample for a confidence limit of 14% (Andersen & Throndsen, 2004).

### 2.4 Zooplankton data

#### 2.4.1 Microzooplankton

Microzooplankton (MZP) samples were collected in six stations at different depths depending on water column vertical profiles (Fig. 1). For MZP analyses, 10 L of seawater were reverse filtered through a 10 µm mesh to reduce the volume to 250 mL and immediately fixed with buffered formaldehyde (1.6% final concentration). Subsamples (50 mL) were then examined in a settling chamber using an inverted microscope (magnification 200x) (Leitz Labovert, Leica DMI 300B), following the Utermöhl method (1958). The entire surface of the chamber was examined. Among the MZP community, five main groups were considered: heterotrophic dinoflagellates, aloricate ciliates, tintinnids, micrometazoans and other rare protozoans. Tintinnids empty loricae were not differentiated from filled loricae because the tintinnid protoplasts are attached to the lorica by fragile strands that can easily detach during the collection and fixing of the samples. For each taxon, the biomass was estimated by measuring the linear dimensions of each organism with an eyepiece scale and relating the shapes to standard geometric figures. Cell volumes were converted into carbon values using the appropriate conversion factors, as follows: aloricate ciliates, pg C cell$^{-1}$ as µm$^3$ x 0.14 (Putt and Stoecker, 1989); tintinnids, pg C cell$^{-1}$ as µm$^3$ x 0.053 + 444.5 (Verity and Langdon, 1984); athecate heterotrophic dinoflagellates, pg C cell$^{-1}$ as µm$^3$ x 0.11 (Edler, 1979); thecate heterotrophic dinoflagellates, pg C cell$^{-1}$ as µm$^3$ x 0.13 (Edler, 1979); other protozoans, pg C cell$^{-1}$ as µm$^3$ x 0.08 (Beers and Stewart, 1970).

### 2.4.2 Mesozooplankton

Three hauls were conducted at six stations (Fig. 1): one horizontally at the surface with the Manta net (333 µm, 0.098 m² net opening) and 2 vertically (from 100 m depth to the surface) with the WP2 net. A Hydrobios flowmeter mounted in the net opening was used to measure the volume of seawater filtered through each net. Immediately after the catch, samples were treated to estimate biomass, taxa composition and abundance of the zooplanktonic community. Samples collected with the Manta net were split by using the Huntsman beaker technique (van Guelpen et al., 1982) and treated as follows: half of the sample was fractionated by passing it through a series of steel sieves with decreasing mesh size (> 2 mm; 2-1 mm; 1-0.5 mm; 0.5-0.2 mm) and immediately frozen at -20°C for biomass analysis, ¼ was fixed and preserved in a seawater-buffered formaldehyde solution (4% final concentration) for later determination of taxa composition and abundance and ¼ was fixed in 96% ethanol for molecular analysis (data not shown in this manuscript). Samples collected with the WP2 net were treated as follows: one sample was entirely fractionated and frozen at -20°C for biomass analysis using the same procedure as the Manta net samples, and one sample was split in half and fixed in seawater-buffered formaldehyde solution (4% final concentration) and 96% ethanol, respectively. In the laboratory, to estimate biomass (dry mass), each size fraction was resuspended in a small volume of filtered seawater and dewatered by vacuum filtration on pre-dried and weighed GF/C filters (47 mm diameter) after being briefly rinsed with distilled water to remove the salts of the seawater (Postel et al., 2000). Each filter was then placed in a small plastic Petri dish and dried in an oven at 60 °C for 24 hours or longer until completely dry and weighed on an electronic microbalance. The fixed samples were concentrated to remove the formaldehyde, and the organisms were suspended in filtered seawater and carefully passed through the same set of sieves used for the biomass. Depending on the abundance, the organisms present in the subsamples or in the entire fractionated sample were counted and identified (Copepoda, Chetognata, Mollusca, and others) under stereo-microscopes (Leica 165C :120x; Leica 205C: 160x).

### 2.5 Microbiological data

#### 2.5.1 Prokaryotic biomass, viable and dead cells, respiring cells

The microbial components were investigated by using different approaches. The detailed methodological procedures for assessing prokaryotic cell abundances, biomass and morphometric features are reported by La Ferla et al. (2012). The viability of prokaryotic cells quantified by the Live/Dead Bac Light Bacterial Viability Kit™ and the number of respiring cells quantified by the Bac Light Redox Sensor CTC Vitality Kit™ were estimated as reported by Azzaro et al. (2022).

### 2.5.2 Physiological profiles of microbial community

Physiological profiles were determined by the Biolog EcoPlate™ microplate assay. The metabolic potentials of bacterial assemblages were quantified as the optical density (OD) values of the formazan produced by oxidation of the 31 carbon sources included in the Biolog Ecoplates. The absorbance was recorded at 590 nm excitation wavelengths using a compact plate reader Byonoy Absorbance 9 and spectrophotometrically measured according to Azzaro et al. (2022) and references therein.

### 2.5.3 Microbial activities involved in organic matter decomposition and mineralization (enzymatic and respiratory activity rates)

The potential decomposition rates of organic polymers (proteins, polysaccharides and organic phosphates), mediated by the microbial enzymes leucine aminopeptidase (LAP), beta glucosidase (GLU) and alkaline phosphatase (AP) respectively, were estimated by incubation with fluorogenic substrates derived from methylcoumarin (MCA) and methylumbelliferone (MUF), as reported by Hoppe (1993), adapted by Caruso et al. (2020). Fluorescence readings were converted into enzymatic activity rates and expressed as the maximum rate (Vmax) of hydrolysis of the substrates, in $nM\ h^{-1}$.

### 2.5.4 Respiratory activity

The respiration rates were measured by the Electron Transport System activity (ETS) assay. This method is based on the conversion of tetrazolium salt into formazan. The detailed methodological procedures were reported by Azzaro et al. (2006, 2021) and references therein.

## 3 RESULTS

The LB21 cruise (see Fig. 1) was conducted in early September, when the seasonal minimum of sea ice extent is recorded in the Arctic and sub-Arctic regions (Fig. 2). The ice extent in the Fram Strait fluctuates from year to year. In the long term, the lowest extent in September was recorded exactly in 2021, the highest in 1987. After 2021, a recovery of the summer seasonal sea ice extent was observed. For winter, the lowest extent was observed in 2006, the highest in 1986. Our cruise therefore took place during the period of the lowest summer sea ice extent measured to date.

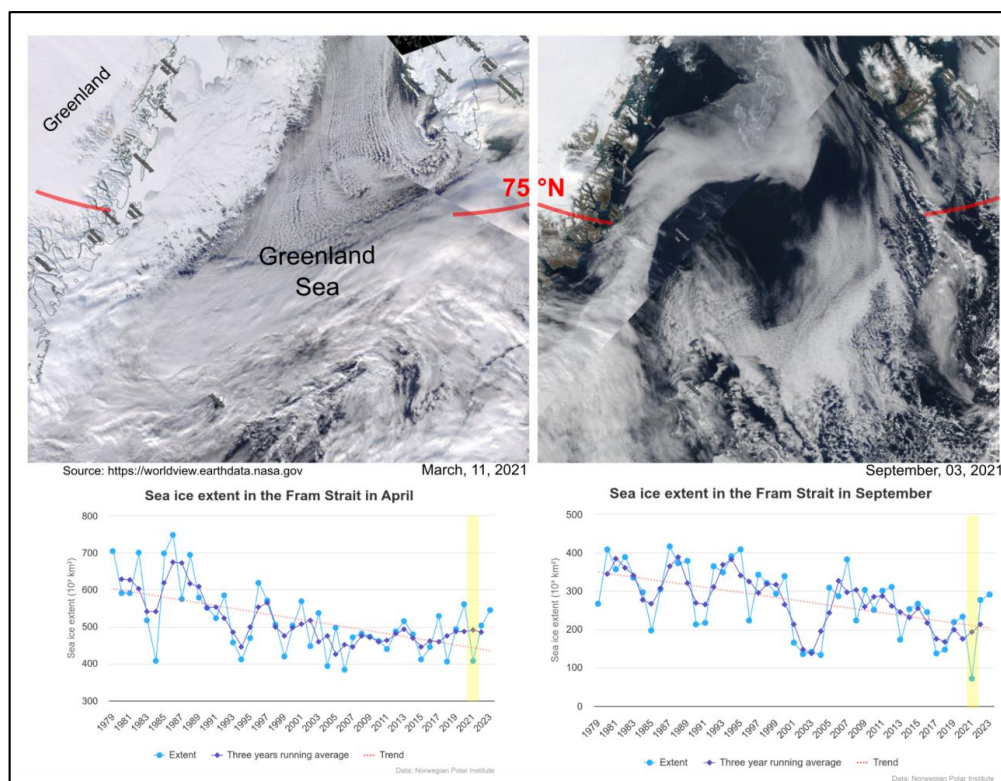
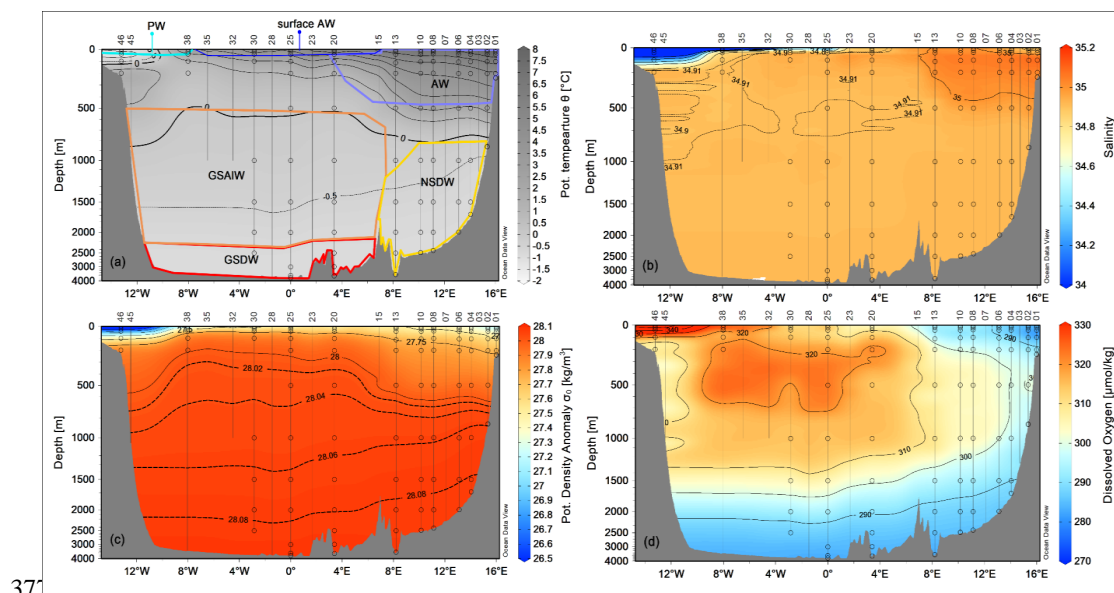



**Figure 2** - **Upper panels**: satellite images of the Greenland Sea and the Fram Strait in March and September 2021 (MODIS Corrected Reflectance Imagery, source NASA). **Bottom panels**: April and September mark the annual maximum and minimum sea ice extension in the Fram Strait (source: Norwegian Polar Institute, 2024, Environmental monitoring of Svalbard and Jan Mayen (MOSJ). URL: https://mosj.no/en/indikator/climate/ocean/sea-ice-extent-in-the-barents-sea-and-fram-strait/). The yellow vertical bar indicates the year 2021.

### 3.1 Physical Oceanography: thermohaline properties distribution

We grouped our data according to the definition of main water masses from Rudels et al. (1999) and Wang et al. (2021) and adjusted them to include most of our data. Due to some data gaps, it was not possible to define all relevant water masses in this area more precisely. Along the zonal transect at 75 °N, the ocean temperature shows a very pronounced horizontal gradient, with typical values of AW ($\theta > 3.0$ °C, S > 35) in the uppermost 500 m of the water column on the eastern side (Fig. 3). At the surface, the AW has temperatures between 4.5 °C and 9.0 °C (Fig. 4). The AW extends to the west and gradually becomes shallower, so that from station 20 to station 35 it only occupies the uppermost 50-80 metres. In contrast, the surface layer on the western side, from station 38 to 46, has a thin (about 40-80 m) layer of Polar Water (PW) with temperature and salinity values that are typical of sea ice meltwater ($\theta < 0$ °C, S $\leqq$ 33). The intermediate zone between 100 m and 500 m is largely occupied by mixed water and we refer to it as the transition layer. The deep layer below 500 m depth is occupied by Greenland Sea Arctic Intermediate Water (GSAIW, $-0.9 < \theta < 0$ °C, S ~ 34.9), Greenland Sea Deep Water (GSDW, $\theta < -1$ °C, S < 34.9) in the central and westernmost part, respectively, and Norwegian Sea Deep Water (NSDW, $\theta$ ~ -1.0 °C, S ~ 34.9) in

362   the easternmost part of the section. We find that the deep layer below 500 m has homogeneous
363   thermohaline properties, with a slightly pronounced difference at the easternmost edge where the
364   NSDW flows northwards. Isotherm at 0 °C and the overall distribution of isopycnals (Fig. 3 a, c)
365   show a classic dome shape with an upwelling in the central part caused by the effect of the GSG,
366   which tends to lift the intermediate water towards the surface due to its cyclonic (i.e., anticlockwise)
367   sense of rotation. In addition, we are certain that the strong horizontal shear and local meteorological
368   conditions can induce the formation of several mesoscale structures (i.e., eddies, hardly shown by
369   our spatial resolution ranging from about 20 km (along the sides) to 40-60 km (in the centre of the
370   transect). They can change the internal distribution and even trap nutrients and other chemical-
371   biological properties. The dissolved oxygen values show a higher oxygen enrichment (> 350 µmol
372   kg$^{-1}$) in the intermediate layer between stations 15 and 38 (Fig. 3d), while the maximum values are
373   found in the upper layer near the Greenland coasts, where the PW flows southwards. In contrast,
374   lower oxygen values (< 300 µmol kg$^{-1}$) are found below 1500 m depth and in the westernmost part
375   of the section where AW and NSDW occur (Fig. 3d). Overall, θ spans from 1.5 °C to almost 9 °C,
376   while S spans from 30 (melting waters) to 35 (AW, Fig. 4).

377

378   **Figure 3 -** Vertical distribution of (a) potential temperature (°C), (b) salinity, (c) potential density anomaly (kg m$^{-3}$), and
379   (d) dissolved oxygen (µmol kg$^{-1}$) along the zonal transect at 75° N in September 2021. The empty dots indicate the
380   sampling points of the Niskin bottles. Panel a show also the distribution of principal water masses according to their core
381   values [**AW** - Atlantic Water; **GSAIW** - Greenland Sea Arctic Intermediate water; **NSDW** - Norwegian Sea Deep Water;
382   **GSDW** - Greenland Sea Deep Water; **PW** - Polar Water or Melting Water].
383

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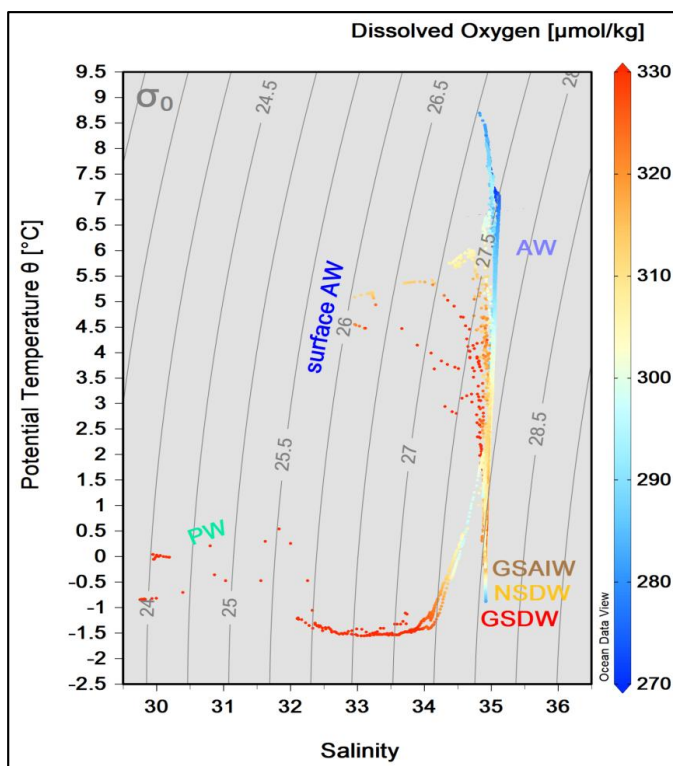

384
385  **Figure 4 -** θ/S diagram from the CTD data collected along the 75 °N section, during the LB21 cruise in September
386  2021. The colorbar on the right-hand side refers to the values for dissolved oxygen concentration (μmol kg⁻¹). [For
387  water masses acronyms see caption of figure 3].

388  **3.2 Biogeochemistry**

389  Along the entire transect, except for the two edges, the seasonal warming in summer had created a
390  well-stratified upper layer about 30-50 metres deep. At the surface, the central Greenland Sea
391  appears to be almost nutrient poor (Fig. 5). The western side of the transect is characterised by
392  higher concentrations of phosphate and silicate, good indicators of the upper halocline of Arctic
393  surface water along the Greenland slope, whereas nitrate concentrations are very low. At depth,
394  NSDW and GSDW (see Fig. 3, 5) were enriched with silicate.

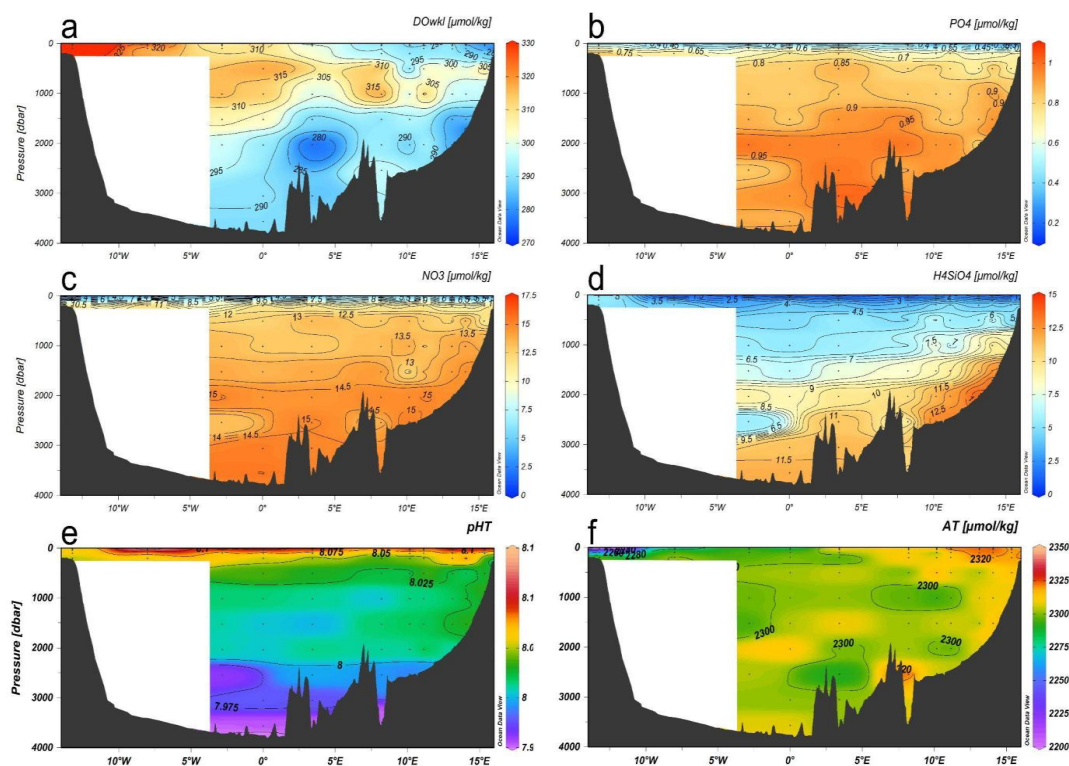
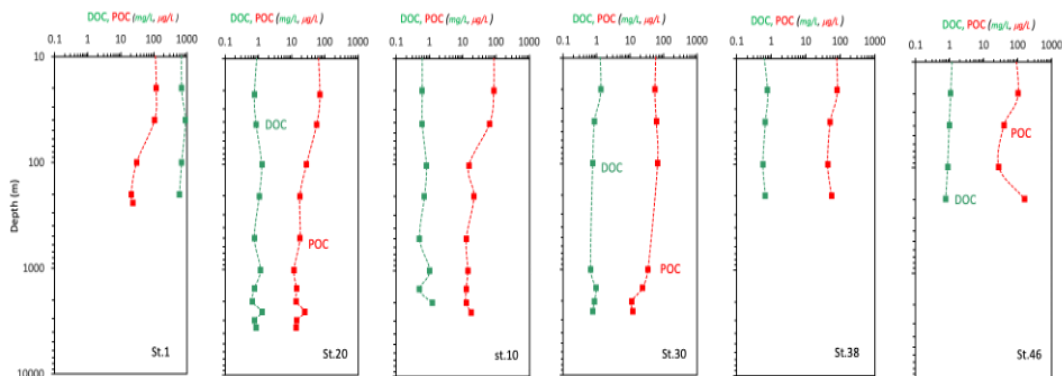



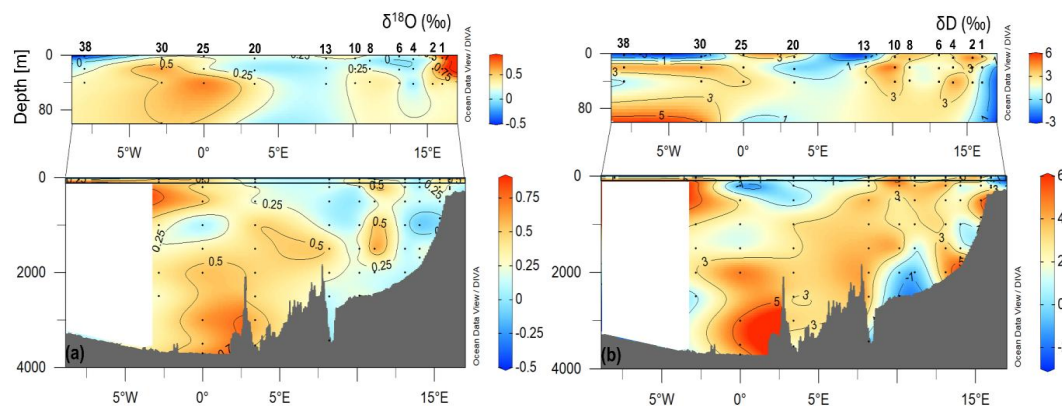
**Figure 5** - Vertical distribution of (a) dissolved oxygen concentration (µmol/kg), (b, c, d) nutrients concentration (µmol/kg), (e) pH$_T$, and (f) Total Alkalinity (µmol/kg) along the zonal transect at 75° N during the LB21 cruise in September 2021. Data are obtained from the laboratory analyses on water samples.

A marked vertical gradient is found along the entire transect with higher pH$_T$ values (up to 8.227, Fig. 5e) in the photic layer (< 100 m depth) decreasing with increasing depth and reaching the lowest values (7.946-7.997) in the deep layer of the Greenland Sea, presumably due to the degradation of settling organic matter. High total alkalinity values (Fig. 5f) are found on the easternmost side where the AW and NSDW flow northwards. The highest values (A$_T$ up to 2320-2348 µmol kg$^{-1}$) are found in the higher salinity AW and particularly in the photic layer also due to the contribution of photosynthetic activity. The lowest values are instead associated with fresh Polar Water found at the surface on the westernmost part of the section.

At the surface layer, a high variability in the rates of respiratory activity and organic matter pool (POC and DOC) was observed (Fig. 6); a decreasing trend up to 1000 m and an interesting increase towards greater depths characterised the vertical profiles of both respiratory activity and DOC. POC concentrations ranged from 11.2 µg C L$^{-1}$ (St 30, at 2002 m depth; 0.93 µM C) to 160 µg C L$^{-1}$ (St 46, at 198 m depth; 13.3 µM C). A greater variability with depth was observed (Coefficient of Variation higher than 65). The highest values were generally found at the surface and within the 20–40 m depth interval. The highest mean concentrations were measured at station 46 (mean 80.5 ± 53.4 µg C L$^{-1}$), mean values at stations 1, 38 and 30 (67.2 ± 46.8; 60.0 ± 15.3 and 41.0 ± 23.1 µg C L$^{-1}$, respectively) and the lowest at stations 10 and 20 (34.1 ± 31.0 and 28.2 ± 20.5 µg C L$^{-1}$, respectively). DOC showed classic vertical profiles, with the highest concentrations (1.22 mg C L$^-$

419    $^1$; 101.7 µM C) at station 30 at 20 m depth and a decrease to a minimum of 0.5 mg C L$^{-1}$ (41.7 µM
420    C) at station 10 at 499 m depth. Between 200 and 1000 m depth, the values remained low (mean
421    $0.83 \pm 0.22$ mg C L$^{-1}$; $69.2 \pm 18.3$ µM C), while a slight increase was observed in the bottom water,
422    particularly at stations 20 and 10 (Fig. 6).

423

424    **Figure 6 -** Vertical profiles of Dissolved Organic Carbon (DOC, mg L$^{-1}$) and Particulate Organic Carbon (POC, µg L-
425    1) concentrations; Data are represented in natural log (*ln*) scale.

426    The isotope values in the total of 96 analysed samples range from -0.43 to 0.8‰ for δ$^{18}$O and from
427    -2.51 to 5.36‰ for δD. Along the transect, a gradient from west to east can be seen in the surface
428    waters, with lower values for δ$^{18}$O and δD on the westernmost side (Fig. 7). The lowest values at
429    the surface in the westernmost part of the section could be related to fresh PW, while the higher
430    values in the easternmost part could be related to northward flowing AW. At depths between 500
431    and 1000 m, isotope values drop to a relative minimum, while maxima occur near the bottom at
432    stations 20 and 25, where GSDW is identified. In addition, a minimum is observed at depth at
433    stations 8 and 10, in an area occupied by NSDW.

434

435    **Figure 7 -** Vertical distribution of δ$^{18}$O (a) and δD (b) along the zonal transect at 75° N during the LB21 cruise in
436    September 2021.

### 3.3 Phytoplankton

#### 3.3.1 Total and size-fractionated chlorophyll *a* (chl-*a*)

Integrated total chl-*a* in the euphotic layer of the water column (0-100 m) averages 55.4 mg m$^{-2}$ while the highest concentrations (73.1 mg m$^{-2}$) were found in the stations located at the ends of the transect (65.3 mg m$^{-2}$) close to the continental shelf (Fig. 8) with the maximum at the easternmost station (st. 46). Overall, the concentration of chl-*a* ranges between 0.20 mg m$^{-3}$ (St.1, 100 m) and 2.90 mg m$^{-3}$ (St.1, 20 m) with a mean value of $0.63 \pm 0.64$ mg m$^{-3}$. Degraded pigments (phaeo) are around 40 % with respect to chl-*a*. The size spectrum of the phytoplankton community biomass along the water column, shows different percentage contributions to the total with 65% for the nano-phytoplankton, 21% for the micro-fraction and 14% for the pico-phytoplankton (Fig. 8). Exception to this composition is observed in the westernmost station (st. 46) where the micro-fraction dominates, replacing the nano-phytoplankton.

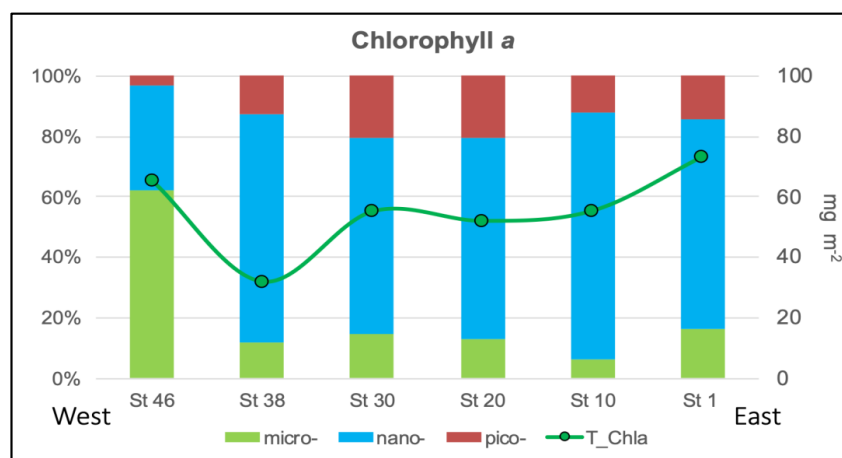



**Figure 8 -** Integrated total chl-*a* concentration (0-100 m depth) at the stations along the 75°N zonal transect (green line) with the percentage of contribution of micro- nano- and pico-phytoplankton size-fractions (histogram).

#### 3.3.2 Phytoplankton composition and abundances

The phytoplankton analyses do not reveal a clear pattern along the transect, although some differences in abundance and composition were observed. The integrated abundances of phytoplankton, ranging from 7.00 (st. 30) to 23.48 x 10$^4$ cells L$^{-1}$ (st. 20), are higher at the easternmost stations than at the westernmost ones (on average 19.33 x 10$^4$ cells L$^{-1}$ at stations 1, 10, 20 and 9.44 x 10$^4$ cells L$^{-1}$ at stations 30, 38, 46, see Fig. 9). This is mainly due to a more even vertical distribution of abundance at the eastern stations. In contrast, the westernmost stations have even higher phytoplankton abundances, but limited to subsurface maxima, like 50.40 x 10$^4$ cells L$^{-1}$ at 38 m at station 20, while abundances in the rest of the water column are very low. The phytoplankton community along the transect is characterised by the dominance of the flagellate group (70% of the total phytoplankton), mainly represented by small (< 10 µm) forms with uncertain taxonomic identification (on average, 61%). Diatoms (on average, 9% of the total phytoplankton) are present in very low abundances in the easternmost stations (on average, 0.43 x 10$^4$ cells L$^{-1}$ at stations 1, 10, 20, 30), while higher values were recorded in the two westernmost stations 38 and 46 (on average, 3.24 x 10$^4$ cells L$^{-1}$). Finally, dinoflagellates accounted for an average of 21% of

468   the total phytoplankton, with higher abundances occurring in the easternmost stations than in the
469   westernmost stations ($3.32 \times 10^4$ cells L$^{-1}$ and $1.42 \times 10^4$ cells L$^{-1}$, respectively).

470

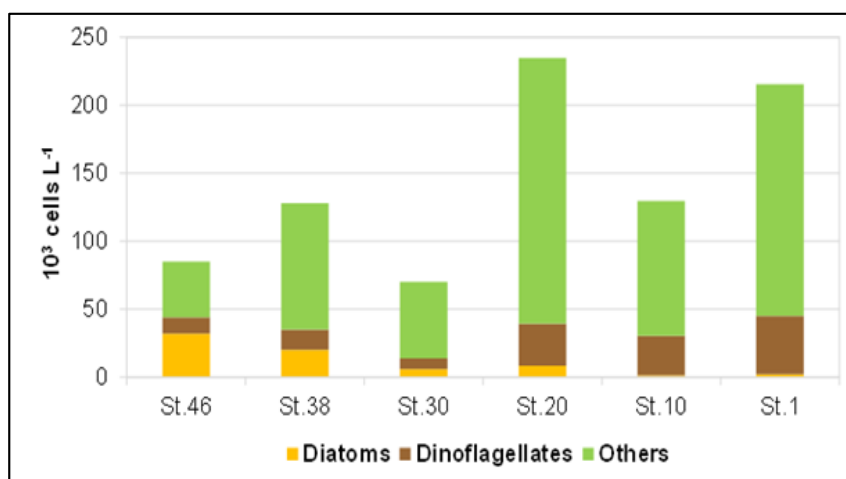



**Figure 9 -** Integrated contribution of the main phytoplankton groups (diatoms, dinoflagellates, and others) to phytoplankton abundance at sampled stations along the 75°N transect.

### 3.4 Zooplankton

### 3.4.1 Microzooplankton abundance and biomass

475   The MZP abundance in the study area varies between 721.5 ind. L$^{-1}$ (St. 38, at 0 m) and 6.25 ind. L$^{-1}$ (St. 20, at 3500 m). The carbon content shows higher values at the surface with a maximum value of 1.7 µgC L$^{-1}$ (St. 38, 0 m) and a minimum of 0.01 µgC L$^{-1}$ at 2520 m depth (St. 10). The vertical distribution of organisms, based on their abundance along the water column, shows higher abundances in the first 200 m, compared to the zone > 200 m. In particular, the integrated abundance of MZP in the upper layer reaches its maximum at St. 38, mainly due to the higher presence of tintinnids, while the lowest values are found at St. 20, in the centre of the transect (Fig. 10). The highest carbon content is recorded at St. 10, which is due to the high abundance of other protists and micrometazoans (Fig. 10). Tintinnids are the most abundant taxa in the study area, followed by heterotrophic dinoflagellates and aloricate ciliates (Fig. 10).

485

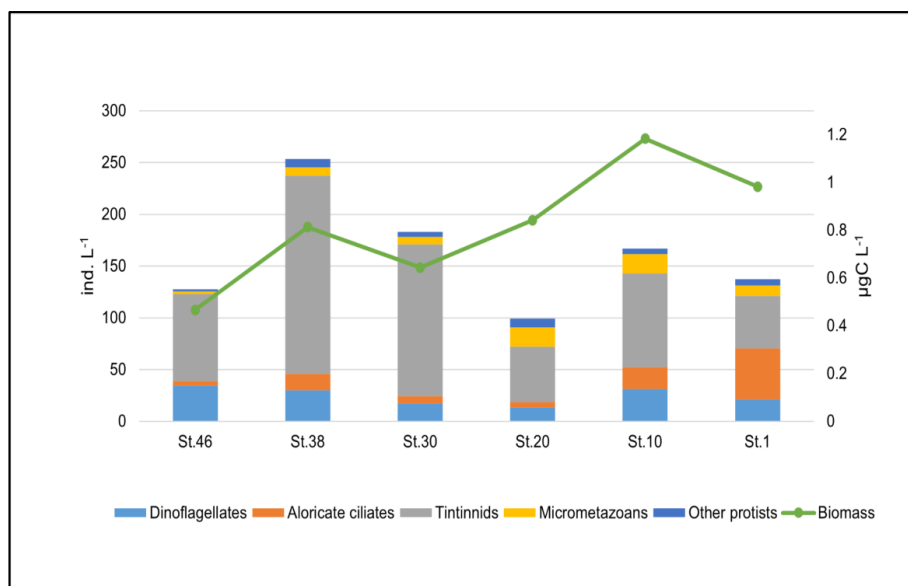



**Figure 10** - Integrated values of abundance (ind. L$^{-1}$) and biomass (µgC L$^{-1}$) of microzooplankton in the upper layer (< 200 m depth).

### 3.4.2 Mesozooplankton biomass and abundance

Biomass and abundance of mesozooplankton have the same distribution along the transect both at the surface (ca. 20 cm depth, Fig. 11a) and in the upper layer (0-100 m depth, Fig. 11b), with the highest values found in the central part of the transect (St. 20). Nevertheless, the surface samples are richer in organisms than those collected in the water column (mean abundance: Manta net: 1257 ± 1110 ind. m$^{-3}$; WP2 net: 492 ± 387 ind. m$^{-3}$), which also corresponds to a higher biomass (mean biomass: Manta net: 41 ± 32 mgDM m$^{-3}$; WP2 net: 10 ± 7 mgDM m$^{-3}$). Overall, organisms 1-2 mm in size account for 61 % of the mesozooplanktonic biomass in the surface waters and are the most abundant at almost all stations (Fig. 11a), while in the samples collected with the WP2 net, the biomass fractions consisting of organisms 1-2 mm and > 2 mm in size were the most abundant, accounting for 35 and 38 % of the total biomass, respectively (Fig. 11b). Copepods are the most abundant taxon both at the surface and in the upper layer (97 % and 94 % of the total mesozooplankton abundance, respectively), mainly represented by the genus *Calanus*. Chaetognaths, although much less abundant than copepods, are found along the entire transect both at the surface and in the upper layer, being most numerous at stations 10, 20, and 30. Molluscs are almost absent in the surface water and are mainly found in the water column at the eastern stations of the transect (St. 1 and St.10).

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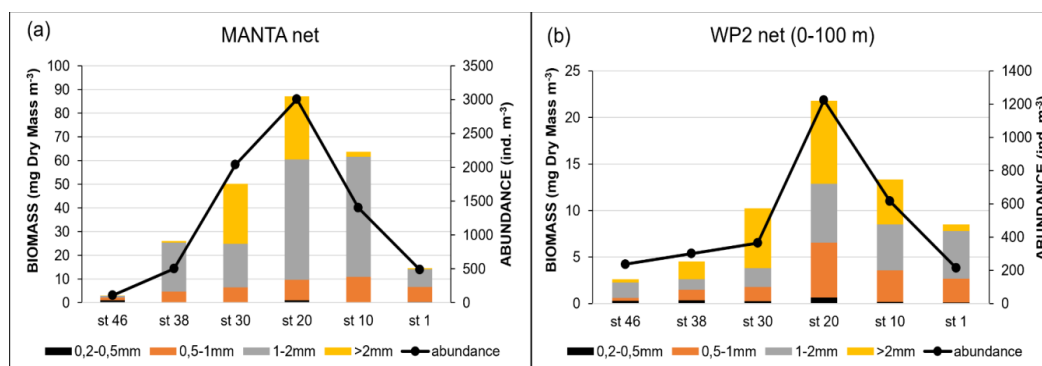
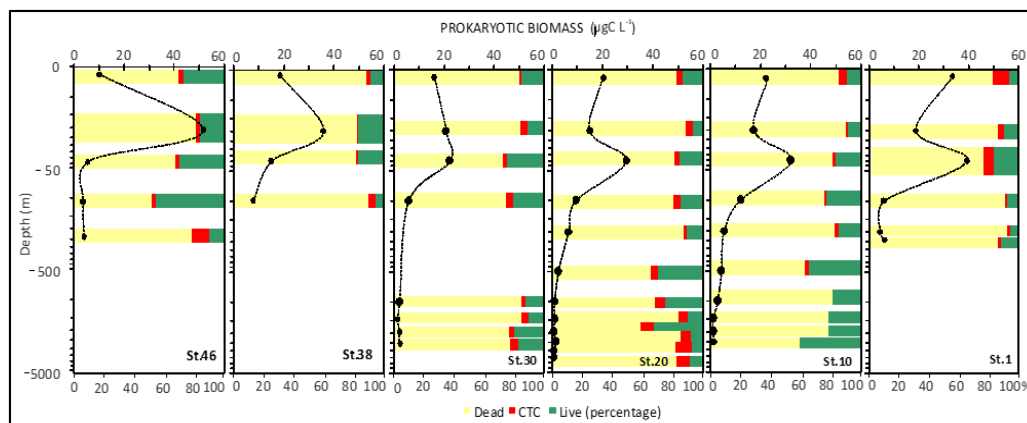
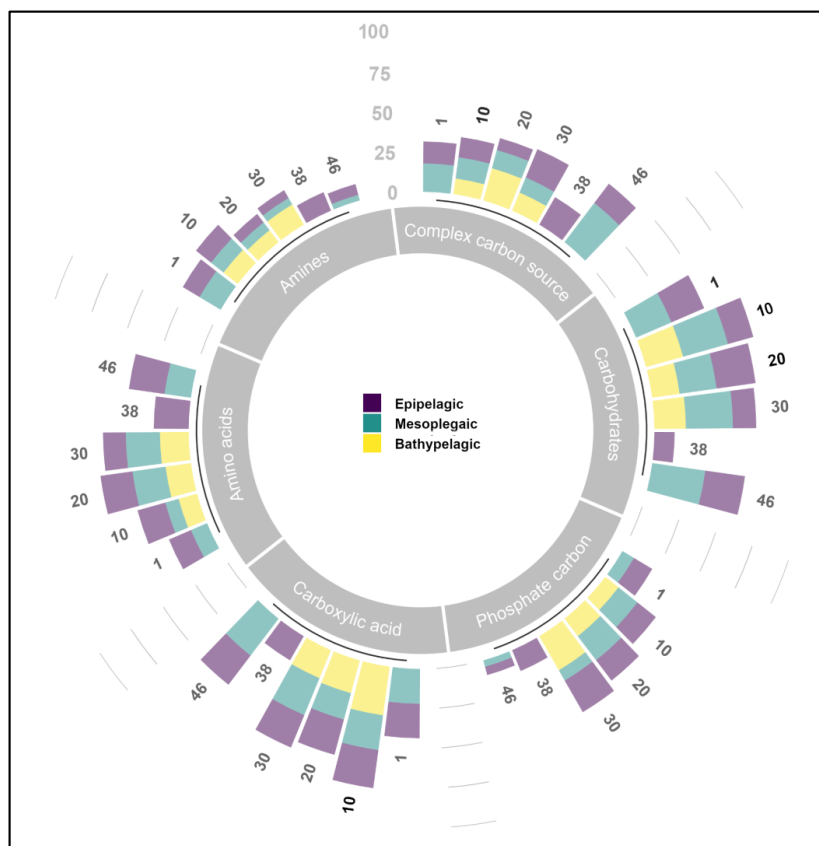

506 **Figure 11** - Biomass and abundance of mesozooplankton: (a) Manta net sampling at surface and (b) WP2 net sampling
507 in the 0-100 m layer.

## 3.5 Microbial compartment: abundance, biomass, and activities

509 The abundance of prokaryotes is high in the photic (0-100 m depth) layer (range: 5.57 to 11.3 x
510 $10^5$ cells m$^{-3}$) and decreases with depth. The highest values are measured at stations 20 and 1. From
511 100 to 1000 m depth, the abundance ranges from 1.75 to 3.21 x $10^5$ cells m$^{-3}$ and from 0.53 to 0.64
512 x $10^5$ cells m$^{-3}$ at greater depths. The cell volume ranges from 0.049 to 0.098 µm$^3$, with a mean
513 value of 0.072 ± 0.018 µm$^3$. In the photic and aphotic layers, the cell volumes vary in a similar
514 range (0.08 ± 0.02 µm$^3$). The highest volumes at great depths characterise stations 30 and 10, where
515 large, curved rods are observed. Apart from stations 30 and 10, the data show a similar vertical
516 profile in the size distribution. The highest percentage of live cells (about 33%) is observed at
517 station 46 in the photic layer. However, peaks are also found in the deeper layers. The number of
518 respiring cells (CTC+) is in the order of 104 cells ml$^{-1}$. Variability between the layers is observed
519 at all stations. In general, the high proportion of respiring cells below 100 m depth is observed at
520 stations 46, 30 and 20. Conversely, the higher percentages are found in the photic layer at stations
521 St. 10 and 1 (Fig. 12).

523 **Figure 12 -** Vertical profiles of prokaryotic biomass, viable and dead cells (Live/Dead) and respiring cells (CTC); Data
524 on y axis are represented in natural log (*ln*) scale.

525   Overall, the average percentages of carbon source utilisation determined by Biolog Ecoplates show
526   that carbohydrates and carboxylic acids were well-utilised polymers at each station and in each
527   layer. Especially in the aphotic layer, the percentage utilisation is highest at most stations. Complex
528   carbon sources and phosphate carbon are utilised at similar percentages throughout the water
529   column; conversely, amino acids are preferentially utilised in the aphotic layer. Amines are only
530   little used (Fig. 13).

531

532   **Figure 13** - Carbon substrate utilisation (as percentage of the total) determined in the epi-, meso-, and bathy-pelagic
533   layers.

534   Enzymatic activity measurements yield values of LAP, GLU and AP ranging from 0.072 to 8.08
535   nM h$^{-1}$, from 0.007 to 0.35 nM h$^{-1}$ and from 0.001 to 0.36 nM h$^{-1}$ respectively (Fig. 14). Total
536   enzymatic patterns depict vertical trends generally decreasing from surface towards deep layers,
537   with some hot spots of metabolic activity at 20-40 m and in the aphotic layer. LAP activity peaks
538   at the lateral ends of the transect as well as at St. 20. AP and GLU decrease from the Western
539   (Greenland) side moving towards St 20, then increase again towards the Eastern side (St. 10) of
540   the transect.

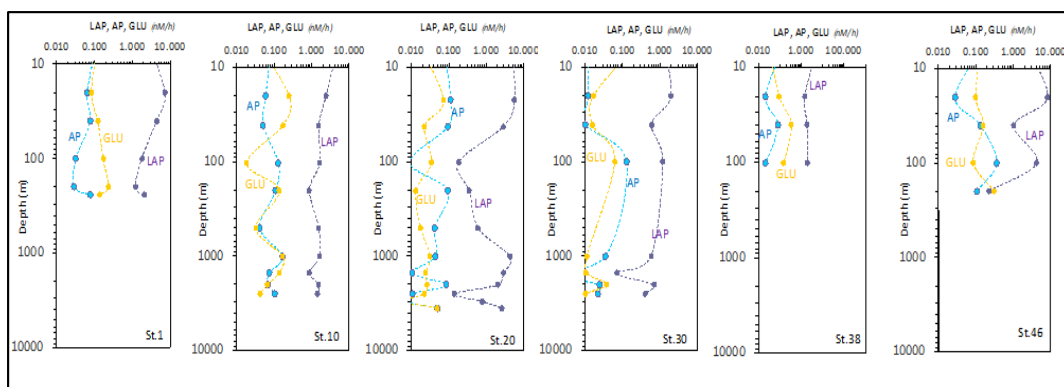



**Figure 14** - Vertical profiles of the enzymatic activity rates measured at the sampled stations (LAP, leucine aminopeptidase; AP, alkaline phosphatase; GLU, beta glucosidase). Data are represented in natural log (*ln*) scale.

The respiration rates (ETS) range from 0.0290 to 0.329 $\mu L\ O_2\ h^{-1}\ L^{-1}$ (Fig. 15). The respiratory activity values generally decrease from the surface to 1000 m depth and then increase in the deeper layer. A high value of respiratory activity was also determined in the deepest sample of St. 46.

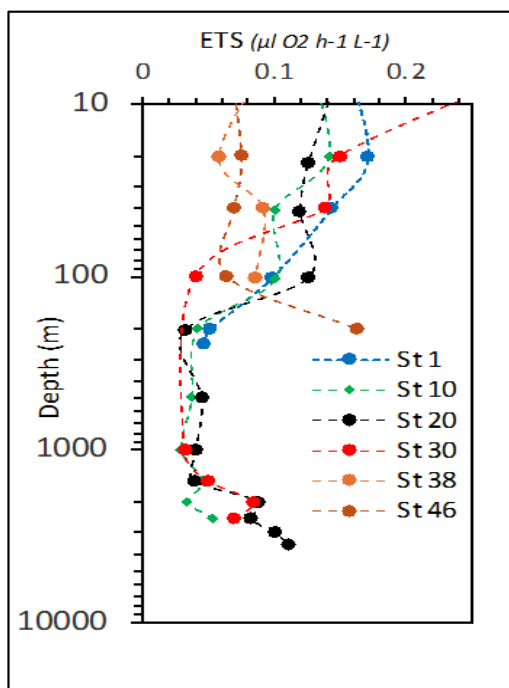



**Figure 15** - Vertical profiles of Electron Transport System (ETS) respiratory activity. Data on the y axis are represented in natural log (*ln*) scale.

## 4    Summary and conclusions

Here we present the main data and results of the multidisciplinary oceanographic campaign carried out between 29 August and 14 September 2021 on board the Italian icebreaker Laura Bassi in the Greenland Sea (along the 75° N latitude section) as part of the Italian project CASSANDRA, funded by the Italian Ministry of Research in the framework of the Italian Arctic Programme. The Greenland Sea, in the north Atlantic, is a region of deep ocean convection that contributes to the AMOC and the exchange of water masses between the Atlantic and Arctic Oceans. On its easternmost side, it is dominated by the presence of AW, while on the westernmost side, by the presence of Polar waters. Both large scale patterns, local meteorological conditions, and ice extent can influence physical and biogeochemical properties. Different phases of the North Atlantic Oscillation index (NAO, i.e., atmospheric pressure difference between the Azores high and the subpolar low at sea level) can modulate basin-wide changes in the intensity and position of the North Atlantic jet stream and storm track, as well as influence patterns of zonal and meridional heat and moisture transport, which in turn can affect temperature and precipitation patterns over the Arctic and sub-Arctic regions. Increasingly positive phases of the NAO are associated with increased AW inflow, as was the case in the late 1980s/early 1990s (Dickson et al., 2000). The winter of 2020/2021, which preceded the cruise, had a slightly negative NAO index (-0.72), after two years of predominantly positive values. A negative NAO means weaker westerly winds in the mid-latitude regions in terms of climatology and stronger winds in the North Atlantic west of Iceland. Sea ice in the Fram Strait fluctuates from year to year. The lowest extent in September was recorded exactly in 2021, the highest in 1987. In the long term, the regions of the Norwegian Sea and the Fram Strait experienced a drop in temperature from 2018 to 2020, which rose again in 2021. Instead, a strong freshening phase set in after 2013, which is continuing. Continuous warming was also observed in the deep-water layer of the Greenland Sea at a depth of 3000 metres. In particular, the GSDW temperature shows a relatively steady increase from -1.18 °C to -0.86 °C between 1993 and 2021 (ICES Report on Ocean Climate 2021, available at https://ices-library.figshare.com/articles/report/ICES_Report_on_Ocean_Climate_2021/24755574?file=43769571). Our measurements report temperature values of about -0.9 °C for the GSDW. Relatively warm and saline Atlantic Water (AW, $\theta > 3.0$°C, S about 35), with highest $A_T$ concentrations, dominates on the eastern side in the upper 500 m with surface temperatures of 4.5–9.0 °C. A dome-shaped isotherm distribution indicates upwelling from the Greenland Sea Gyre, while several mesoscale structures such as eddies seem to be responsible for the large spatial variability in the upper layer. At the surface, the central Greenland Sea is almost nutrient poor. On the western side, however, higher phosphate and silicate values are good indicators for the upper halocline of the Arctic surface water along the Greenland slope. Nitrate levels remained very low there. The deep waters NSDW and GSDW, presented the lowest $pH_T$ values and the highest enrichment of silicate. Phytoplankton biomass along the euphotic water column, expressed as chl-*a*, showed a quali-quantitative difference between the central-eastern and western sectors with greater abundances at the extremes of the transect. The dimensional structure of the phytoplankton community characterizes the PW with a predominance of the micro-phytoplankton fraction that almost entirely replaces the nano-phytoplankton fraction, more abundant in the AW, even in the front station between these two water masses. The plankton communities were analysed with more detail in the upper layer (0-100 m), where the phytoplankton and zooplankton diversity reflected the different water masses (AW and PW). Diatoms increased at the western stations affected by PW, while dinoflagellates and small

596 flagellates were more abundant at the eastern stations affected by AW. The higher MZP abundance
597 was recorded at St. 38, where the layer below 30-40 m depth was still occupied by AW, while the
598 surface layer was affected by PW. The MZP abundance and biomass decreased drastically in the
599 presence of cold PW on the Greenland slope. Micrometazoans and aloricate ciliates increased
600 towards the easternmost side of the section, where the stations were characterised by higher
601 temperatures. Copepods of the genus *Calanus* were the main taxa observed, but the structure of the
602 mesozooplankton communities changed during the transect and polar taxa increased westwards.
603 The $\delta^{18}O$ and $\delta D$ isotope ratios indicate the influence of freshwater at the surface level near the
604 Greenland shelf. We also highlight a marked difference in $\delta^{18}O$ and $\delta D$ isotope ratios between
605 GSDW and NSDW, which occupies the bottom region of the studied area, with NSDW showing
606 lighter values compared to GSDW. Prokaryotic abundance and microbial enzymes generally
607 depicted vertical decreasing trends with peaks of cells and activity recorded at station 20 as well as
608 at the ends of the transect. While living cells prevailed at station 46 in the photic layer, actively
609 respiring cells were quite variable in their distribution. Large, curved rods were found at stations 30
610 and 10. A high utilization of carbohydrates and carboxylic acids regardless of the examined station
611 or depth characterized the microbial community metabolism. Amino acids were actively
612 metabolised in the aphotic layer, while no differences were found in the utilization of complex
613 carbon sources and phosphate carbon compounds along the water column.

614 This study emphasises the significant spatial and vertical variability of water properties, nutrient
615 distribution and biological communities caused by local and seasonal oceanographic dynamics in a
616 region characterised by a strong exchange of water masses between the Arctic and Atlantic Oceans
617 and a major influence of atmospheric teleconnections between the polar and subpolar regions.

618 **5 Data availability**

619 Data presented here are available through the repository Italian Arctic Data Center (IADC), at the
620 following links: CTD casts, available at https://doi.datacite.org/dois/10.71761%2Fc082c3ca-40bf-
621 42b1-a61a-7b3697ab2c5a (Bensi et al., 2024). Physical, biological, and biogeochemical analyses
622 on water samples from Niskin bottles available at
623 https://doi.datacite.org/dois/10.71761%2Ff7474404-3331-43e5-883b-25755e94956d (Azzaro et
624 al., 2024). Satellite data used in the work are freely available at
625 https://worldview.earthdata.nasa.gov/. Data on sea ice extension are available at
626 https://mosj.no/en/indikator/climate/ocean/sea-ice-extent-in-the-barents-sea-and-fram-strait/.

627 **Declaration of Competing Interest**

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

630 **Acknowledgements**

631 The field activities carried out on board the icebreaker Laura Bassi were partly funded by the Italian
632 Ministry of University and Research through the Italian Arctic Research Programme
633 (https://www.programmaricercaartico.it/), the National Institute of Oceanography and Applied
634 Geophysics (OGS) and the National Research Council (CNR). The research was conducted as part
635 of the project "AdvanCing knowledge on the presentArctic Oceanby chemical-phySical,

636  biogeochemical and biological obServAtioNs to preDict the futuRe chAnges" (CASSANDRA),
637  which was funded by the Italian Arctic Research Programme (CASSANDRA project;
638  PRA21_0001). This study was supported by the Synoptic Arctic Survey (SAS) – an international
639  research programme that coordinates the data collection of essential ocean variables measured
640  during Arctic research cruises. SAS is partly funded by the European Union's Horizon 2020
641  research and innovation programme through the Arctic PASSION project under the grant agreement
642  10100347. We would also like to thank Lidia Urbini, Paolo Mansutti, Leonardo Langone, Patrizia
643  Giordano, Warren Cairns, Matteo Feltracco, the entire technical staff of the OGS, the captain and
644  crew of the Laura Bassi for their support during the data collection. We also thank Alenka Goruppi
645  and Giulia Peloso (OGS, Trieste, Italy) for their help in processing the zooplankton data and Nives
646  Ogrinc (Jožef Stefan Institute Ljubljana, Slovenia) for the analyses of δ13C-DIC.

647  **Author contribution**

648  M.B., M.A. and G.Ci. conceived and wrote the main part of the article. M.B., M.A., G.Ci., M.G.,
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650  G.M., C.T., L.P. F.S., F.D., C.C. contributed to the collection and/or processing of the data,
651  preparation of figures, and to the discussion of the results. M.A. led the CASSANDRA project. All
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