

The Green Edge cruise: Investigating the Marginal Ice Zone processes during late spring / early summer to understand the fate of the Arctic phytoplankton bloom.

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Abstract. The Green Edge project was designed to investigate the onset, life, and fate of a phytoplankton spring bloom
75 (PSB) in the Arctic Ocean. The lengthening of the ice-free period and the warming of seawater, amongst other factors, have induced major changes in Arctic Ocean biology over the last decades. Because the PSB is at the base of the Arctic Ocean food chain, it is crucial to understand how changes in the Arctic environment will affect it. Green Edge was a large multidisciplinary collaborative project bringing researchers and technicians from 28 different institutions in seven countries together, aiming at understanding these changes and their impacts into the future. The fieldwork
80 for the Green Edge project took place over two years (2015 and 2016) and was carried out from both an ice-camp and a research vessel in Baffin Bay, in the Canadian Arctic. This paper describes the sampling strategy and the dataset obtained from the research cruise, which took place aboard the Canadian Coast Guard Ship (CCGS) *Amundsen* in late spring/early summer 2016. The sampling strategy was designed around the repetitive perpendicular crossing of the marginal ice zone (MIZ), using not only ship-based station discrete sampling, but also high-resolution measurements
85 from autonomous platforms (Gliders, BGC-Argo floats...) and under-way monitoring systems. The dataset is available at <https://doi.org/10.17882/86417> (Bruyant et al., 2022).

1 Introduction

The Arctic Ocean is currently experiencing unprecedented environmental changes. The increase of the summer ice-retreat lengthens the phytoplankton growing season, but also increases the area of the marginal ice zone (MIZ). If trends are maintained, all Arctic sea-ice may become seasonal as early as twenty years from now (Meredith et al., 2019), increasing the MIZ coverage even more. Ice edge blooms represent much of the annual phytoplankton primary production in the Arctic Ocean (Perrette et al., 2011; Ardyna et al., 2013), and their current phenology is relatively well known (Wassmann and Reigstad, 2011; Leu et al., 2015). However, we currently do not know how precisely primary production will respond to climate changes. The overarching goal of Green Edge was to understand the processes that control an arctic PSB as it expands northward, and to determine its fate through the investigation of related carbon fluxes (e.g., Trudnowska et al., 2021). This study was also motivated by the discovery that PSBs can and do occur underneath the ice (Arrigo et al., 2014) despite the limited amounts of under-ice available light (Mundy et al., 2009; Arrigo et al., 2014; Lowry et al., 2014; Assmy et al., 2017, Randelhoff et al., 2019). Field studies for the Green Edge project were carried out in 2015 and 2016 at an ice-camp located on landfast sea ice close to Qikiqtarjuaq (NU, Canada). Additionally, during late spring/early summer 2016, a cruise aboard CCGS *Amundsen* was conducted in Baffin Bay. As explained in Randelhoff et al. (2019), Baffin Bay is both relatively easy to access and represents an ideal framework for this study, because environmental conditions are representative of what is observed at the pan-arctic scale. Particularly, the warm Greenland current flowing north on the Greenland side, and colder waters on the Canadian side flowing south (Baffin Island current) induce an evenly retreating ice edge, allowing a straightforward sampling strategy. Here, we present an overview of the dataset gathered during this cruise.

2 Study area, sampling strategy and ship-based operations

For logistical reasons, the cruise was divided into two legs. Leg 1A started on June 3rd in Québec City, ended on June 23rd in Qikiqtarjuaq (NU), and included one week of transit to the study zone. Leg 1B started on June 23rd in Qikiqtarjuaq and ended in Iqaluit (NU) on July 14th. During the five-week period spent in Baffin Bay, the ship crossed the MIZ, from open waters (in the east) to sea ice-covered areas (in the west) and back again following latitudinal transects. A total of seven transects were covered between 68.0° N and 70.5° N (Fig. 1A). Three transects were covered during Leg 1A (68.5° N to 69.0° N) and four during Leg 1B (68.0° N and 69.5° N to 70.5° N). For each transect, stations were separated by six nautical miles (approximately 11 km) to obtain a relatively high spatial resolution. Fifteen to twenty-five stations were sampled within each transect, for a total of 144 stations visited during the campaign.

The activities conducted at each type of station are detailed in Table 1. Briefly, at so-called CTD (Current Temperature Depth) stations, rosette casts did not include seawater collection. The rosette, a Sea-Bird model 32 carousel equipped with twenty-two 12-L Niskin bottles, was geared with multiple sensors (see Table 2 for details). At NUT stations, seawater samples were additionally collected at several depths between 2 and 2000 m for nutrient analyses. At BASIC stations, the apparent optical properties of seawater were measured using underwater profiling optical instruments.

The variables measured at BASIC stations included the concentration of chlorophyll *a*, phytoplankton pigment, particulate carbon and nitrogen, and particulate absorption spectra. Finally, at FULL stations, a suite of measurements was made on seawater samples, and several underwater instruments were deployed as well, including a suite of optical sensors. Vertical plankton nets geared with a plankton imager, horizontal net trawls, benthic trawls and Ursnel box corers were also deployed at each FULL station. Note that the number of variables measured on seawater samples was significantly larger at FULL stations, where three rosette casts were necessary to cover the demand in water volume, compared with one rosette at BASIC stations. At least one FULL station was sampled in each of the three major domains covered by each transect, namely open waters, the MIZ and the ice-covered area. During Leg 1B a larger emphasis was placed on collecting ice-cores for analyses and measurements of light propagation through the ice and snow. FULL stations sampled in the ice-covered domain did not include trawling operations.

Our sampling strategy allowed successive crossings of different PSB stages: the early-bloom stage at the western end of transects covered in sea-ice, late- to post-bloom stages at the eastern end of transects in open waters, and full-bloom stage in the middle of transects around the ice edge (see Randelhoff et al., 2019). Between the stations, the ship-track water monitoring system (TSG SBE45 from Sea-Bird, WETStar fluorometer from WetLabs, LiCOR non-dispersive infra-red spectrometer (model Li-7000), Campbell Scientific CR1000 data logger) recorded temperature, salinity, chlorophyll-*a* fluorescence and pCO₂ at 7 m depth continuously when navigating outside the ice pack. A moving vessel profiler (MVP300-1700, AML Oceanographic, Victoria, BC Canada), equipped with a micro CTD (AML), a WetLabs C-Star transmissometer and a WetLabs Eco-FLRTD fluorometer; was deployed in open waters.

To significantly extend the monitoring of the PSB beyond the duration and space covered by the cruise, we deployed four BGC-Argo (Bio-Geo-Chemical-Argo) floats on July 9th. They collected data until late fall (Fig. 1B), performing a 0-1000 m profile each day. The ice-specific version of BGC-Argo floats that was deployed, the so-called PRO-ICE, is commercialized by NKE Electronics (France). These floats carry a typical biogeochemical payload (Sea-Bird 41 ARGO CTD, Aanderaa optode 4330 Oxygen sensor, Sea-Bird™ OCR-504 PAR (photosynthetic available radiation) and downwelling radiance sensor (380, 412 and 490 nm), Sea-Bird ECO-FLBBCD fluorescence chlorophyll-*a*, colored dissolved organic matter (CDOM), and backscattering sensor and Sea-Bird SUNA nitrate sensor). They also include a 2 way-directional Iridium communication Rudics for data transmission and a sea-ice detection system to protect the float from hitting sea ice on ascent (Le Traon et al., 2020; André et al., 2020). The 2016 data are available on the following website: <http://www.obs-vlfr.fr/proof/ftpv/greenedge/db/DATA/FLOATS/>.

Two SLOCUM G2 gliders (Teledyne Marine Inc.) were deployed during the cruise from the ship's zodiac either to revisit transects, or in areas that were not visited by the CCGS *Amundsen* (Fig. 1B). They both carried a similar scientific payload and communication system as the BGC-Argo floats, rendering the possibility of inter-calibration (Fig. 2). Gliders were primarily used in the MIZ where a 90-m icebreaker would disturb the fragile hydrological structure of shallow under-ice water masses. Gliders were deliberately directed through the same area as the BGC-Argo floats to compare CTD and optical data from both platforms. Results in Fig. 2 show a good agreement for the data. Gliders travel following a programmed sawtooth pattern, joining pre-defined waypoints. Data and instructions were transmitted both ways via iridium when the glider surfaced. Figure 3 shows an example of the level of detail

provided by the gliders when travelling through complex water structures. Representing chlorophyll fluorescence measured over a three-week journey covering almost 500 km, the data show that the glider(s) did travel through both surface blooms and a subsurface chlorophyll maximum between 20-50 m. The data presented in Fig. 3 represents 693,000 chlorophyll fluorescence measurements. These constitute robust results towards the validation of the use of multiple measurement platforms to investigate complex systems.

A schematic synopsis of all operations carried out during the campaign can be found in Fig. 4A. The use of multiple analysis platforms allowed the measurement of more than 150 parameters during the Green Edge cruise (Table 3), together they accurately describe the complexity of the MIZ trophic systems (Fig. 4B) and most of the stocks and processes involved in the development of the PSB.

3 Time-for-space formatting and data quality control

One of the challenging tasks, when assembling data from a large group of researchers, is to adopt a common frame for spatial and temporal tagging of samples. Geographic positions require a lot of attention and conversion of latitude and longitude into one format (we used decimal degrees east) to ensure data could be easily merged. The concomitant use of local time and Coordinated Universal Time (UTC) during the cruise also presented a challenge. For the Green Edge cruise, an operation logbook was created to keep track of all operations conducted on the ship in sequential order during each day (local time). Each operation was associated with a unique operation ID to which all other data could be referenced. The use of ordinal date (number of days since January first) was used to avoid confusion between European and American date writing conventions. Each cast within a given operation type (CTD/RO for Rosette, AOP for Apparent Optical Properties, etc.) was numbered sequentially, starting from 001, throughout the entire cruise. As a result, any given operation received a unique code which thereafter could be used to merge all the data acquired during that operation.

Different control procedures were adopted to ensure the quality of the data. First, the raw data were screened to identify and –where possible, eliminate errors originating from the measurement devices, including sensor (systematic or random) errors inherent to measurement procedures and methods. Instrumental pre- and post-calibration corrections were applied when necessary. Statistical summaries such as average, standard deviation and range were computed to detect and remove anomalous values in the data. Then, data were checked for duplicates and remaining outliers. Once raw measurements were cleaned, data were structured and gathered into single comma-separated values (CSV) files. Each of these files was constructed to gather variables of the same nature (e.g., nutrients). In each file, a minimum number of variables (columns) was always included to make dataset merging easier and accurate (Table 4).

4 Description of data collection: an overview

4.1 Meteorological, navigational and ice coverage data

Along the ship track, an Automated Voluntary Observing Ship (AVOS) system recorded all data related to navigation, including the position of the vessel and basic atmospheric meteorological data (including barometric pressure). A meteorological tower was also installed on the foredeck of the ship to measure additional variables (instantaneous wind speed and direction, air temperature and relative humidity, atmospheric partial pressure, and vertical fluxes of CO₂). Averaged wind speeds and directions over the entire Baffin Bay were retrieved from the Remote Sensing System website (<http://www.remss.com/measurements/ccmp/>) and were computed according to the Cross-Calibrated Multi-Platform (CCMP) wind vector analysis product (V2.0, Atlas et al., 2011) (see one example of a pattern calculated over the months of June and July in Fig. 5). A major change in wind patterns happened between June (light 1-2 m s⁻¹ southward winds) and July (4-5 m s⁻¹ northward winds), which impacted sea ice movements. Figure 6 shows sea-ice cover over four periods covering the total sampling time of the cruise (<https://nsidc.org/data/g02186>), NSIDC (U.S. National Ice Center and National Snow and Ice Data Center) (2010). The north-south general orientation of the ice edge is visible, along with, over time, the westward progression of the MIZ.

Ice cover history was compiled and expressed as Open Water Days (OWD) before sampling day (Fig. 7), calculated from the difference (in day number) between the date of sampling and the date at which the sea-ice concentration reached 10 (panel A), 50 (panel B) and 80% (panel C) in the geographical location under study. Ice concentration data was obtained from the Advanced Microwave Scanning Radiometer 2 (AMSR2) sea-ice concentration data on the 3.125 km grid (Spreen et al., 2008), downloaded from http://www.iup.uni-bremen.de:8084/amsr2data/asi_daygrid_swath/n3125/ (see Randelhoff et al., 2019, for details on the calculations). In Fig. 7, yellow and pale green colors on the Greenland side indicate a positive value between 20 and 40 OWD which reflects how long the open-water conditions had prevailed at those stations at the time of sampling. To the east, closer to the Canadian side, the colder colors and negative values indicate the presence of sea ice. Open water days is a useful metric/index and can be computed using different sea-ice cover (SIC) thresholds, depending on the goal. For example, in Randelhoff et al. (2019), the SIC value used for hydrological interpretation was 10% (as in Fig. 7 top panel). However, a biologist might want to consider a SIC of 80% (Fig.7 bottom panel) when looking for the onset of phytoplankton growth, as only 20% of open water surface already dramatically increases the amount of light available.

4.2 Physical data

4.2.1 Sea ice

During sea-ice sampling operations, snow depth, ice thickness and freeboard were measured at each ice coring site. Over the cruise, ice thickness varied between 32 and 108 cm, while freeboard varied between 10 and -8 cm (top of the ice under water). Several cores were retrieved at each site using a 9 cm diameter Mark II ice-corer (Kovacs Enterprises Inc., Roseburg, OR USA). Each ice core was sliced into 10 cm segments (from the bottom) after temperature was measured. Salinity was assessed after thawing and filtration using a salinometer (Guildline Autosol 8400B, Guildline

220 Instruments Ltd., Smith Falls, ON Canada). Snow density and granulometry were assessed opportunistically (Eicken
et al., 2009).

4.2.2 Water masses

Hydrological conditions during the cruise were determined using several tools. A moving vessel profiler (MVP) was
225 deployed, oscillating between 0 and 300 m depth while towed at an average speed of 12 knots, rendering a very high
spatial resolution (Fig. 8 bottom row, one profile every 2 km). Data obtained with the MVP matched the patterns
observed from the rosette data acquired on sampling stations (Fig. 8 middle row). Profiles of conductivity, temperature
and pressure were collected using a Sea-Bird SBE 911plus CTD system rigged on the rosette. The data were post-
processed according to the standard procedures recommended by the manufacturer and averaged over 0.2 m vertical
bins. While there was a sharp transition in SIC at the ice edge along transect 300, the change in SIC was less steep in
230 the more extensive MIZ of transect 500 (top row in Fig. 8). Nonetheless, both transects show similar patterns, with
100% SIC, colder (below $-1\text{ }^{\circ}\text{C}$) and fresher (salinity below 33.5 g kg^{-1}) waters close to the surface on the western
side, and 0% SIC, saltier (above 33.6 g kg^{-1}) and warmer (above $0\text{ }^{\circ}\text{C}$) waters within the first 50 m on the eastern side.
These observations were consistent with the northward inflow of Atlantic-origin waters along the Greenland shelf
break, and the southward outflow of Arctic/Pacific-origin waters along the Baffin Island shelf break.

235 Currents in the water column were measured using a hull-mounted 150 kHz Acoustic doppler current profiler (ADCP,
Teledyne RD Instruments Ocean Surveyor, California, USA), as well as two L-ADCP installed on the rosette structure
(RDI, WHM300-1-UG304) in a Master/Slave configuration. Vertical profiles of water turbulence were measured at
each FULL station using a self-contained autonomous microprofiler (SCAMP, Precision Measurement Engineering,
California, USA) deployed from the zodiac. A detailed description of the hydrological structures in the studied area
240 during the Green Edge cruise is presented in Randelhoff et al. (2019).

4.3 Chemistry

Partial pressure of CO_2 (pCO_2) was measured continuously (every two minutes) using a Li-7000 CO_2 analyzer
(LICOR, Lincoln NE, USA) coupled to a General Oceanics Underway System model 8050 (General Oceanics, Miami
FL, USA) connected to the ship-track water monitoring system. At each FULL, BASIC and NUT station, discrete
245 samples were collected using the Niskin bottles at 10 or more depths for seawater analysis (see Table 3 for the complete
list). To complement the pCO_2 data from the underway system and provide full profiles of the seawater CO_2 system,
total alkalinity and dissolved inorganic carbon (DIC) concentrations were determined on discrete samples according
to Dickson et al. (2007). Concentrations of the major macronutrients (nitrate, phosphate and orthosilicic acid) were
determined with a segmented flow AutoAnalyzer model 3 (Seal Analytical, Germany) using standard colorimetric
250 methods adapted from Grasshoff et al. (1999). Nitrate concentration in the water column was also determined during
each CTD cast using an in situ ultraviolet spectrophotometer (ISUS, Satlantic Inc., Halifax NS, Canada) mounted on
the rosette. Concentrations varied between 0 and non-limiting concentrations over the entire cruise, with
concentrations gradually increasing from the surface to bottom. Surface waters showed higher concentrations of

macronutrients in the western half of the transects than in the eastern half of the transects, indicating surface water
255 nutrients had been used by the developing PSB.

Organic elemental composition of total and dissolved matter (nitrogen, carbon, and phosphorus) was measured on
water samples taken from the Niskin bottles. Samples were immediately poisoned with sulfuric acid and brought back
to the lab for analysis using wet oxidation, as described by Raimbault et al. (1999). Subtracting signals obtained for
filtered samples (dissolved matter) from non-filtered samples (total matter) rendered calculated values for particulate
260 organic matter. Particulate organic carbon and nitrogen were also analysed on filtered samples (Whatman™ glass
fiber GF/F, GE Healthcare USA) using high-temperature oxidation combined with gas chromatography.

4.4 Light Field and Bio-optics

The characteristics of the light field (quantity and quality) passing through snow and sea ice into the water column
were assessed, as light is the most important parameter triggering the PSB. From the top of the CCGS *Amundsen*
265 wheelhouse, the total solar downwelling radiation was measured using a pyranometer (0.3 to 300 μm wavelength) and
a radiometer (visible range 300 to 750 nm). Surface radiometry was performed using a HyperSAS (Hyperspectral
Surface Acquisition System, SeaBird Scientific USA) placed on the bow of the ship, including simultaneous
measurements of hyperspectral above-water downward irradiance ($E_d(0^+, \lambda)$), and sky and surface radiance ($L_{\text{sky}}(0^+, \lambda)$
or $L_{\text{tot}}(0^+, \lambda)$). Data were recorded at each FULL and BASIC station in open waters. Above-water hyperspectral
270 remote sensing reflectance ($R_{\text{rs}}(\lambda)$) was calculated from the radiometric quantities following the ocean optics protocols
of Mueller et al. (2003) and Mobley (1999). In-water vertical profiles of downward irradiance ($E_d(z)$) and upward
radiance ($L_u(z, \lambda)$ or irradiance ($E_u(z, \lambda)$) in the water column were measured using two different versions of the
Compact Optical Profiling System (C-OPS, Biospherical Instruments Inc.) radiometer. In open waters, the free-fall
version of the profiler was deployed from the ship (Antoine et al., 2013). The sea-ice version (ICE-Pro) was deployed
275 during ice sampling through an auger hole carefully filled with fresh snow to avoid, as much as possible, disturbing
the underwater light field. A reference sensor provided simultaneous measurements of downward irradiance in the air.
All measurements were made at 19 different wavelengths between 320 and 875 nm.

A profiling optical package was deployed at 28 stations to measure the inherent optical properties (IOPs) of seawater.
The measured properties (and sensors) included: fluorescence of chlorophyll-*a* and fluorescent dissolved organic
280 matter (FDOM) (WetLabs, Eco Triplets), spectral total non-water absorption coefficients between 360 and 764 nm
(Hobilabs, a-sphere), particle backscattering coefficient at six different wavelengths (Hobilabs, hydroscat-6, 394, 420,
470, 532, 620 and 700 nm) together with CTD data (Sea-Bird SBE 19plus attached to the package). In addition,
discrete water samples were taken at each FULL and BASIC station to measure in the lab the CDOM absorption
coefficient (a_{CDOM}) between 200 and 722 nm using an Ultrathin (World Precision Instruments), and the phytoplankton
285 and non-algal particle absorption coefficients between 200 and 860 nm determined from the “inside sphere” filter-pad
technique (Rottgers and Gehnke, 2012; Stramski et al., 2015) using a spectrophotometer equipped with a 155-mm
integrating sphere (Perkin Elmer Lambda 19). Note that a_{CDOM} , phytoplankton and non-algal particle absorption
coefficients were also measured on the bottom slice of thawed ice cores. The data revealed that the minimum light

amount required for net phytoplankton growth ($0.415 \text{ mol m}^{-2} \text{ d}^{-1}$; Letelier et al., 2004) can be reached deeper under the ice than expected (Randelhoff et al., 2019). Further details of the light field measurements can be found in Massicotte et al. (2020).

Three optical profilers were also attached to the rosette carousel and rendered 203 profiles of CDOM fluorescence (FluoCDOM Wetlabs USA) and chlorophyll concentration (estimated from in situ fluorescence, Seapoint fluorometer, USA), as well as 87 profiles of light transmittance (Wetlabs C-Star transmissometer, USA). Data is shown in Fig. 8 for transects 300 and 500. For both transects, the highest values of chlorophyll-*a* concentration and of the attenuation coefficient (both parameters being strong proxies for phytoplankton biomass) were observed close to the surface in the MIZ, and deeper at around 50 m in ice-free waters, showing a progression in the PSB development starting close to the surface along the ice edge and growing into a subsurface chlorophyll maximum (SCM) where surface waters were depleted in nutrients. Concentration of CDOM showed its lowest concentrations at the surface of open waters where SIC dropped below 50%, and where the surface waters had been depleted of nutrients by phytoplankton growth.

4.5 Biodiversity

The Green Edge project also aimed to understand the related potential impacts of evolving environmental conditions on Arctic food-webs in the context of climate change. Hence, great care was taken to sample the entire size spectrum of particulate matter and living organisms (Fig. 9) from the tiniest viruses and bacteria to demersal fishes, seabirds, and marine mammals. A wide variety of sampling techniques and analyses, from visual observation to highly automated underwater imaging systems, allowed us to ensure that almost all the levels of the trophic network were examined.

4.5.1 Viruses and bacteria

Abundance of viruses and bacteria was determined on fresh and preserved (glutaraldehyde 4% final concentration) water samples taken from the rosette at each FULL and BASIC station (10 depths) using two different flow cytometers. On board, fresh samples were counted using an Accuri™ C6 and preserved samples were counted back in the lab using a FACSCanto (both machines from Becton Dickinson Biosciences, San Jose, CA, USA). Samples were processed according to Marie et al. (2001). Bacteria (and viruses) are ubiquitous in the oceans, and in Baffin Bay we measured bacterial abundances up to $2.9 \times 10^6 \text{ cells ml}^{-1}$.

For bacterial diversity analysis, water samples were filtered sequentially onto 20 μm , 3 μm (both polycarbonate filters, Millipore) and 0.22 μm (Sterivex-GV, Millipore). The filters and Sterivex were stored at $-80 \text{ }^\circ\text{C}$ with RNAlater (Qiagen) until analyzed. The DNA/RNA co-extraction was carried out using the AllPrep DNA/RNA kit (Qiagen). The V4-V5 hypervariable region of the 16S rRNA gene was amplified by PCR using primers 515F-Y and 926R covering a broad spectrum of diversity, including Archaea and Bacteria (Parada et al., 2016). PCR, as well as sequencing settings and bioinformatic of sequence data, can be found in Dadaglio et al. (2018).

A comparison of the bacterial diversity as a function of geographic location and size fractions (free living bacteria, bacteria attached to particles smaller than 20 μm and larger than 20 μm) was made at a relatively broad taxonomic

level (Fig. 10). The samples from the different groups were mainly dominated by Bacteroidetes and Proteobacteria. In general, the proportion of Proteobacteria decreased from ice stations to open water stations featuring a more advanced stage of PSB, giving way to Bacteroidetes and more specifically Flavobacteriaceae.

4.5.2 Phytoplankton community

Flow cytometry was used (same protocols as for viruses and bacteria, Section 4.5.1) to count and differentiate the smallest cells (picophytoplankton, nanophytoplankton, cryptophytes and *Synechococcus*) according to their fluorescence and scattering properties at each FULL station. At the surface, picophytoplankton and nanophytoplankton cell concentrations could reach 60,000 (station 719 – transect 7 station 19) and 9,000 (station 515 –transect 5 station 15) cell ml⁻¹, respectively while cryptophytes were always below 360 cell ml⁻¹. *Synechococcus* cyanobacteria were never observed.

Some samples were used to start phytoplankton cultures, which were taken back to the Roscoff laboratory for purification using flow cytometry sorting, serial dilution, and single-cell pipetting. Pure cultures were characterized by microscopy and 18S rRNA gene sequencing. Most cultures isolated during the cruise belonged to diatoms, especially to the genera *Attheya* and *Chaetoceros* (Gérikas Ribeiro et al., 2020). All cultures were deposited in the Roscoff Culture Collection and are available for distribution (<http://www.roscoff-culture-collection.org/strains/shortlists/cruises/green-edge>).

To study the phytoplankton community composition, an Imaging FlowCytobot (IFCB, McLane Research Laboratories Inc., East Falmouth MA, USA) was used during Leg 1B. The IFCB is best used for the study and identification of cells between 1 and 150 µm. Fresh samples (5 ml) taken from the rosette at each FULL station (all depths) and some BASIC and NUT stations (2 to 7 depths) were analyzed, as well as samples from the 2 bottom-most slices of ice cores once melted. The IFCB takes pictures at a resolution of around 3.4 pixels per µm. Image descriptors/features were extracted with Matlab®, using scripts developed by Heidi Sosik (Sosik and Olson, 2007). Taxonomic determination was achieved using Ecotaxa (Picheral et al., 2017; <http://ecotaxa.obs-vlfr.fr>). Random forest algorithms were used for automatic classification. Reference set and validation of predictions were both done manually. Examples of specimens observed during the Green Edge campaigns can be found in Massicotte et al. (2020).

A total of 203 underwater vertical profiles were acquired using an underwater vision profiler (UVP, model 5-DEEP, Hydroptics France) installed on the frame of the rosette carousel. The UVP5 collects in-focus images in the small seawater volume lit by its light emitting diodes (LEDs), as it is lowered in the water column. An automated computer system (<https://ecotaxa.obs-vlfr.fr/>) was used to sub-sample images of individual objects and sort them into the appropriate category (marine snow or various taxa of zooplankton). The UVP has been mainly developed to count and identify particles larger than 100 µm. Figure 11 (left panels) show the average vertical profiles of particle concentration (ml⁻¹) over the top 350 m of the water column, for open-water (top) and under-ice (bottom) stations. The number of particles close to the surface in open waters coincides with the larger phytoplankton biomass observed there compared with ice-covered stations, consistent with lower primary and secondary production under sea ice. Deeper in the water column, around 300 m, the large particle concentrations observed at open-water stations likely reflect resuspension of

bottom sediments, because these observations were mostly made in the eastern part of Baffin Bay over the continental shelf, whereas under-ice stations were mostly located on the deeper Canadian side of the Bay (see the bathymetry in Fig. 1).

Samples for taxonomic analyses of micro-algae by microscopy were taken at each FULL and BASIC station at ten sampling depths. Half a litre of seawater was preserved with Lugol and kept at 4 °C until it was analyzed in the laboratory. Visual observation and taxonomic determination were done using an inverted microscope (Eclipse TS100, Nikon Instrument Inc.) according to the Utermöhl method (Utermöhl, 1958), using 25 or 50 ml columns. Three transects of 26 mm at 400x were systematically observed for identification and counting of Bacillariophyceae, Dinophyceae, flagellates and ciliates. Larger phytoplankton cells and colonies were observed in all chambers at 100x. Diatoms (Bacillariophyceae) were found at every station, primarily at the surface of the water column, along with flagellates (both at the surface and in the subsurface chlorophyll maximum, SCM) (Fig. 12). The most striking feature was the dominating presence of a *Phaeocystis* sp. (Prymnesiophyceae, blue bars in Fig. 12) at the SCM, reaching 60 to 90% of the cell counts (and to a lesser extent at the surface) across a wide range of ice cover conditions (OWD values between -12 and 12 days).

Phytoplankton and ice-algae pigments were measured to derive indices of micro-algae biomass and taxonomic composition, and to get information on processes such as photoacclimation, senescence and grazing activities (Roy et al., 2011). Rosette water samples were filtered onto GF/F filters (Whatman™, GE Healthcare Life Sciences) and quickly frozen in liquid nitrogen. Back in the land-based laboratory, samples were thawed and extracted in 100% methanol, separated, and identified by HPLC, as described by Ras et al. (2008). Twenty-five individual pigments or groups of pigments were identified and quantified at each FULL and BASIC station (10 depths sampled each time). Figure 13 shows the distribution of total chlorophyll-*a* and pheophorbide concentrations along transects 300 and 500. The high chlorophyll-*a* concentrations close to the surface in the MIZ, and deepening towards open waters in the east, confirmed the evolution of the PSB from an under-ice bloom to a SCM. Note that highest concentrations of pheophorbide were systematically found underneath the accumulation of chlorophyll *a*, indicating the sinking of degrading phytoplanktonic material.

4.5.3 Zooplankton and Fish

Zooplankton represents the second level of the food chain. The UVP5 and the Imaging FlowCytobot (see previous sections) both rendered valuable information on small zooplankton specimens (below 150 µm). Figure 11-right panels show copepod volumetric fraction ($\text{cm}^3 \text{m}^{-3}$) over the top 350 m of the water column, for open-water (top panel) and under-ice (bottom panel) stations. The total copepod volumetric fraction calculated based on automatic identification made using the Ecotaxa web application (<http://ecotaxa.obs-vlfr.fr>), shows subsurface peaks around 20 m at both open-water and ice-covered stations. A secondary peak right at the surface is present at ice-covered stations, where a subpopulation of copepods may stay close to the bottom of sea ice to feed on sympagic microalgae, small animals, and related detritus.

For bigger specimens, a series of vertical nets and trawls were deployed at each FULL station (no trawling operations took place when sea ice was present). An assembly of 4 nets of 3 different mesh sizes (50, 200 and 500 μm) coupled with a Lightframe Onsight Key species Investigation (LOKI) system rendered high-resolution pictures of individuals sampled along the water column together with actual specimens. The multi-net plankton sampler (Hydro-bios, Altenholz, Germany) uses a different sampling strategy. Composed of 9 identical nets (200 μm mesh size), it is hauled vertically in the water column with the nets opening sequentially at different depths, each net collecting a slice of the water column displaying the vertical distribution of the species sampled. Figure 13C shows abundance (ind m^{-3}) of zooplankton sampled using the vertical 200 μm mesh net along transects 300 and 500. Copepods represent the main zooplankton class observed during the cruise, and they were present at every sampling site. The particularly high abundance at station 507 (situated under the ice where a phytoplankton bloom had already disappeared) might be due to the relatively high abundance of copepod nauplii (25% of all copepod individuals compared to the usual 4% at other stations).

Ichthyoplankton was sampled using a double square net towed obliquely from the side of the ship at a speed of ca. 2-3 knots to a maximum depth of 90 m. A Star-Oddi® mini-CTD attached to the frame and flowmeters determined the real depth and volume of sampling. For fish sampling, an echo sounder (EK60, Simrad, Kongsberg Maritime, Norway) mounted on the hull was used to locate and determine the depth of fish aggregations along the ship track during the entire cruise. When pelagic juveniles and adult fish were present at the sampling stations, an Isaac-Kidd Midwater Trawl (IKMT, Filmar and Québec-Océan, Québec, Canada) was towed for 20 minutes at a speed of 2 to 3 knots. When demersal fish were detected, a Benthic Beam Trawl (BBT, Filmar and Québec-Océan, Québec, Canada) was used to sample bigger specimens (between 10 and 32 mm mesh size) living on the bottom sediment.

Samples of zooplankton were all processed in the same way. Swimmers (fish larvae and juveniles) were sorted out, measured, identified, and preserved in a mix of 95% ethanol and 1% glycerol (final concentrations) for later analysis, while zooplankton samples were preserved in 4% formaldehyde solution. Zooplankton abundance and diversity were determined using binoculars back at the laboratory. A few samples (hydro-bios) were analyzed using the zooscan and the ecotaxa identification tools (<https://ecotaxa.obs-vlfr.fr/prj/802>). Fish from IKMT and BBT sampling were sorted, counted, identified, and measured before preservation in a -20 °C freezer in case further analyses are needed in the future. Acoustic data from the EK60 were analyzed in Echoview® (see Geoffroy et al. (2016) for details).

4.5.4 Benthos sampling

Benthos sampling was performed to answer the following specific objectives to test if i) sea-ice cover is the primary environmental driver of contribution and geographic distribution of sympagic carbon on the seabed, ii) sympagic carbon is the most important baseline food source supporting benthic consumers during spring and summer in areas close to the MIZ, and iii) deep benthic food web dynamics and structural variability are directly linked with both depth and availability of food sources (Yunda-Guarin et al., 2020). The sampling was achieved using two different strategies; a total of 16 Agassiz trawling and 34 box coring operations (some down to more than 2000 m depth) were carried out during the cruise. The Agassiz trawl (KC Denmark a/s Research Equipment) is a medium-size dredge trawled behind

the ship, allowing sampling of the macrofauna living on the sediment surface. Once brought back onboard, the content of the net was immediately rinsed with seawater, manually sorted and identified to the lowest taxonomic level possible. Samples were brought back to the laboratory to be identified under a dissecting microscope when onboard identification was impossible. Figure 14 shows an example of the diversity of benthic organisms. More than 220 species of macrofauna were identified during the Green Edge cruise from more than 25 classes (Grant C. and Yunda-Guarin G. Unpublished data). A box corer was used to sample sediment from each FULL station. The sediment samples were then divided among the research teams for diverse analyses (see Table 3 for a complete list) including (but not limited to) identification of organisms living inside the sediment, incubations for respiration and nutrient utilization or chemical analysis.

4.5.5 Birds and marine mammals

During the entire cruise, a systematic bird and marine mammals survey was carried out from the ship's wheelhouse (see LeBlanc et al., 2019 for detailed methodology). A total of 20 different bird species and 8 different mammal species were identified. Northern fulmar, thick-billed murre and little auk were the most common bird species observed. Ringed seal, hooded seal and harp seal were the most common seals. The long-finned pilot whale was the most common whale species observed. A total of 10 polar bears were observed.

A total of 123 seabirds from 7 species were also collected from a zodiac deployed from the CCGS *Amundsen* in Greenland waters between June 10th and July 8th. This includes black-legged kittiwakes ($n=8$), glaucous gulls ($n=6$), great black-backed gull ($n=1$), little auks ($n=19$), northern fulmars ($n=42$) and thick-billed murres ($n=36$). Sampled birds were frozen at $-20\text{ }^{\circ}\text{C}$ until laboratory analyses. A first study aimed to investigate the co-distribution of seabirds and their fish prey along the MIZ (LeBlanc et al., 2019). To this end, stomach contents were examined for 74 birds (35 murres, 30 fulmars and 9 kittiwakes) under a dissecting microscope. Otoliths were retrieved and used to identify fish species, age and size. A second focus was recording of plastic in the stomachs. Plastic data are used in OSPAR monitoring and AMAP working groups on plastic pollution. A third study aimed to determine bird's association to sea ice and ice-derived resources by the combination of different trophic markers. Hence, liver, muscle, and blood (from the cardiac clot) samples were collected from a total of 52 bird carcasses (27 murres, 14 auks, 3 kittiwakes and 8 fulmars), on which Highly Branched Isoprenoids (HBIs), carbon and nitrogen stable isotopes and fatty acids were measured. Finally, a fourth study was looking at stable isotope data together with mercury (Hg) data for both muscle and liver.

5. Biological production and fluxes

5.1 Bacterial production, respiration, and viability

At each FULL station during the cruise, water samples were taken from 2-3 depths (surface, deep chlorophyll maximum (DCM) and below DCM) to determine bacterial respiration. Oxygen concentration was determined using the Winkler method on $1\text{-}\mu\text{m}$ filtered samples before and after a 5-day incubation in the dark at $1.5\text{ }^{\circ}\text{C}$. Bacterial

460 respiration varied overall between 0 and $1.63 \mu\text{mol O}_2 \text{ l}^{-1} \text{ d}^{-1}$, with a mean value of $0.35 \pm 0.41 \mu\text{mol O}_2 \text{ l}^{-1} \text{ d}^{-1}$. For
bacterial production determination, water was collected at each FULL station from 8-10 depths. Bacterial production
was measured by [^3H]-Leucine incorporation (Kirchman et al., 1985) modified for microcentrifugation (Smith and
Azam, 1992). Overall values varied between 0 and $1.51 \mu\text{gC l}^{-1} \text{ d}^{-1}$ around a mean value of $0.17 \pm 0.25 \mu\text{gC l}^{-1} \text{ d}^{-1}$.
The use of the propidium monoazide (PMA) method identified a high bacterial mortality in sea ice (up to 90%) and
465 in SPM material (up to 68%) collected in shallower waters at station 409 and station 418 (Burot et al., 2021).

5.2. Primary production and micronutrient cycling

To determine the fate of the phytoplankton spring bloom, one must first determine primary production. In situ
simulated incubations were carried out at each FULL station on water sampled from the rosette at 8 to 10 depths
determined as chosen percentages of surface photosynthetic available radiation (PAR, namely 100, 50, 25, 10, 6, 2.9,
470 1.2, 0.6 and 0.1%). The melted bottom-most slices of ice cores were also incubated, when available. After spiking the
water with a mix of $^{13}\text{C}/^{15}\text{N}$ tracers, samples were incubated on deck at simulated light levels identical to the sampling
light levels. The dissolved and particulate matter resulting from these incubations were analyzed by mass-spectrometry
resulting in detailed nitrogen assimilation and regeneration values (see Table 3 for a complete list of measurements),
as well as phytoplankton primary production (PP). Primary production varied between 0 and $88.13 \pm 3.0 \mu\text{gC l}^{-1} \text{ day}^{-1}$
475 over the entire cruise.

Photosynthetic parameters also allow calculation of primary production and provide insight into the efficiency and
characteristics of photosynthesis of a given sample. Onboard, photosynthetic parameters were determined by the P vs
E curves method using $\text{NaH}^{14}\text{CO}_3^-$ spiked incubations of water samples (Lewis and Smith, 1983). Changes in the
saturation parameter E_k ($\mu\text{mol Quanta m}^{-2} \text{ s}^{-1}$) in the surface waters of the transect show clear variations in light level
480 acclimation, increasing from lower values around $39 \mu\text{mol Quanta m}^{-2} \text{ s}^{-1}$ at the western under-ice stations to $217 \mu\text{mol}$
 $\text{Quanta m}^{-2} \text{ s}^{-1}$ at the eastern open-water stations (transect 700 values given as an example). Great emphasis was placed
on the contribution of diatoms to primary production, as diatoms are the main phytoplankton group present during the
PSB. Experiments on silica production and dissolution were performed throughout the cruise to locate the actively
growing diatoms. These experiments confirmed the occurrence of active silicification beneath the sea ice, where both
485 centric and pennate diatoms were observed (see details in Lafond et al. (2019)).

5.3 Fate of the phytoplankton spring bloom

Some organic matter produced by the PSB was exported down the water column, as algal cells aggregated and sank,
or were grazed upon by vertically migrating zooplankton. One ambitious experiment was conducted during the cruise
to monitor the export of the PSB at a high temporal resolution. A sequential sediment trap (Technicap PPS4 France;
490 12 sampling cups) was anchored to an ice floe and deployed 25 m under the ice from June 15th to July 9th, 2016.
Sediment trap collection cups were filled with filtered seawater adjusted to a salinity of 38 psu with NaCl and a
formalin concentration of 4% to preserve samples during deployment and after recovery. The carousel holding the
sampling cups was programmed to rotate every 2 days. The sediment trap was deployed in the marginal ice zone along

495 transect 200 (Fig. 1), eventually drifted south with the ice, and was recovered on the way back to Iqaluit. The sediment trap was no longer anchored to its floe at recovery, but sea ice was still present in the region. Taxonomic identification of the algal cells collected showed a constant export of diatoms (~ 50 million cells $m^{-2} d^{-1}$) from June 15th to early July, when a 6-fold increase in diatom fluxes was observed from July 5th to July 7th (~ 300 million cells $m^{-2} d^{-1}$) along with a peak in chlorophyll-*a* fluxes. More than half of the cells exported during the peak in algal fluxes were identified as the ice-associated pennate diatom *Navicula* spp.. Fluxes of the ice-obligate pennate diatom *Nitzschia frigida*, among 500 the first species to be consistently exported from the melting sea ice in the Arctic Ocean (Lalande et al., 2019; Dezutter et al., 2021; Nadaï et al., 2021), peaked from June 23rd to June 25th, probably indicating the onset of sea ice melt. Fluxes of copepod fecal pellets collected in the sediment trap were higher prior to June 27th, suggesting under-ice grazing of ice algae until the ice melted.

5.4 Benthic processes

505 Taken from the box corer, portions of the sediment were incubated at in situ simulated conditions of temperature and light to assess the consumption of oxygen and nutrients by endofauna. Oxygen use in the sediment cores allowed calculation of the benthic carbon demand ($mgC m^{-2} d^{-1}$) which was found to be especially high at stations in open waters where the PSB had already reached senescence and sinking organic matter had reached the bottom.

5.5 Other data

510 The exhaustive list of parameters measured during the cruise is presented in Table 3 along with the responsible principal investigators (PIs) name.

6. Data availability

Making such a large and diverse dataset available to others requires the use of many different platforms. 515 Administrative rules and previous habits and commitments explain why our dataset is hosted by various websites with some of it in more than one place (see in Table 3, the link and file information, when applicable for each parameter acquired during the cruise). Some funding Agencies require data to be deposited in a specific database as a deliverable. In our case, the “Les Enveloppes Fluides et l’Environnement-Cycles Biogéochimiques Environnement et Ressources” (LEFE-CYBER) repository is our main host: http://www.obs-vlfr.fr/proof/php/GREENEDGE/x_dataлист_1.php?xxop=greenedge&xxcamp=amundsen. This is where ALL data and 520 associated metadata can be found for the Green Edge cruise. Particularly, detailed metadata files associated with each variable contain the principal investigator’s contact information. For specific questions, the PI associated with the data should be contacted directly. The LEFE-CYBER platform, however, does not deliver DOI, which is a very important feature for visibility of data. To obtain a DOI for the Green Edge cruise dataset, we uploaded the available formatted data on SEANOE (SEA scieNtific Open data Edition) under the CC-BY license: <https://www.seanoe.org/data/00752/86417>. All data hosted on the SEANOE website have been formatted as described

in section 3, but not all data original format allows transformation into the “.csv” file type. It is therefore important to keep a repository up to date where one can find all raw data.

530 Major long-term research programs often have their own repository / database available which has been used since the onset of their research. While the data of the BGC-Argo floats we deployed during the Green Edge cruise are hosted on the LEFE-CYBER repository, they also have been made available, together with the entire BGC-Argo data, from the biogeochemical Argo database: <https://biogeochemical-argo.org/data-access.php>

535 Some more specific data acquired during the cruise are also available on dedicated websites. For example, dissolved inorganic carbon (DIC), alkalinity and ¹⁸O data are also archived with the Ocean Carbon and acidification Data System (OCADS): <https://doi.org/10.25921/719e-qr37>.

Geographical specificity might also be a motivation for cross-uploading of data. Since the *Amundsen* has been used to conduct polar research, all navigation, AVOS, ADCP, MVP and CTD data are systematically uploaded to the Polar Data Catalog (PDC): <https://www.polardata.ca/>.

540 Please note that in Table 3 only one address for each parameter is provided while most of them are also available from other sources.

7. Lessons learned

As for any scientific cruise, a large amount of data was acquired by many people. Even though guidelines had been suggested ahead of time for data formatting, merging and storage, a tremendous amount of effort was necessary to collect, assemble and standardize the data. It is important that a clear and streamlined data management plan be established ahead of time to avoid errors or loss of data in the merging process. For oceanographic CTD-rosette sampling-based cruises, depth of sampling necessitates special attention, as it is crucial to use Niskin bottle number instead of nominal depth to correctly merge the data. Furthermore, we cannot emphasize enough that data management specialists must be involved from the beginning of such large-scale projects, to ensure that data is properly documented, to render the best quality dataset possible and avoid loss of both valuable time and data.

550 8. Conclusion

The Green Edge cruise was of typical oceanographic design. In terms of goal achievement, the cruise was extremely successful and generated an impressive dataset over a diverse set of disciplines, providing a global picture of the explored environment and of all the processes fuelling the Arctic food-web. Figure 4B represents all interactions existing and/or measured during the cruise between compartments of the various trophic levels. The generated dataset contains a much larger number of parameters than those presented in this paper. All data can be obtained from the data repository and provide an excellent opportunity for re-use and comparison with other Arctic datasets. A special issue of the *Elementa: Science of the Anthropocene* journal entitled “Green Edge –The phytoplankton spring bloom in the Arctic Ocean: past, present and future response to climate variations, and impact on carbon fluxes and the marine food web” contains a collection of research papers referring to this cruise. A complete list of peer reviewed journal

560 publications presenting data from either or both the Green Edge Ice Camp or Green Edge cruise can be found in Table
5.

9. Authors contribution

MB designed the Green Edge project, including the scientific objectives and sampling strategy. MB, KL, FB, TL, PG,
GJ, TB, MP, CM, KC, DM, MT, GD, GF, LD, JD, CL, ML, GN, NM, MC, MP, K-ML, HJ-W, ER, AV, LBdF, DD,
565 NG, HC, AS, BQ, SH, GB, SEG, CG, P-LG, JET, EB, CS, MGT, JR, AB, RA, CB, BM, JL, EL, PB and PC were
onboard the ship and took part in sampling and on-board analysis. FB, PG, GJ, TB, MP, CM, KC, DM, MT, GD, LD,
GF, LD, JD, CL, LM, GN, NM, NP, K-ML, HJ-W, ER, AV, LBdF, DD, NG, HC, AS, BG, SH, GB, GC, P-LG, M-
NH, CS, MGT, JR, AB, RA, CB, BM, JL, EL, BS-B, PC, PA, LA, SB, DC, VC-B, FC, MD, CD, BE, IE, JF, LF, MG,
CGR, CG, PG, CG, SH, RH, FJ, AL, PL, FLG, KL, JL, AL, ML, ALdS, GM, AM, LAM, P-IM, AM, AM, CP, MP,
570 PR, J-FR, RS, JT, A-PT, DV and CN took part in processing and analyzing of the samples and in generating data. M-
PA, and PM cleaned, merged, and assembled the dataset. CS maintains the Les Enveloppes Fluides et
l'Environnement-Cycles Biogéochimiques Environnement et Ressources (LEFE-CYBER) repository it is stored in.
NS, SM, JS, LRL, TP and PB oversaw communication and outreach. MHF, JF, JL and FB oversaw logistics. FB wrote
the manuscript.

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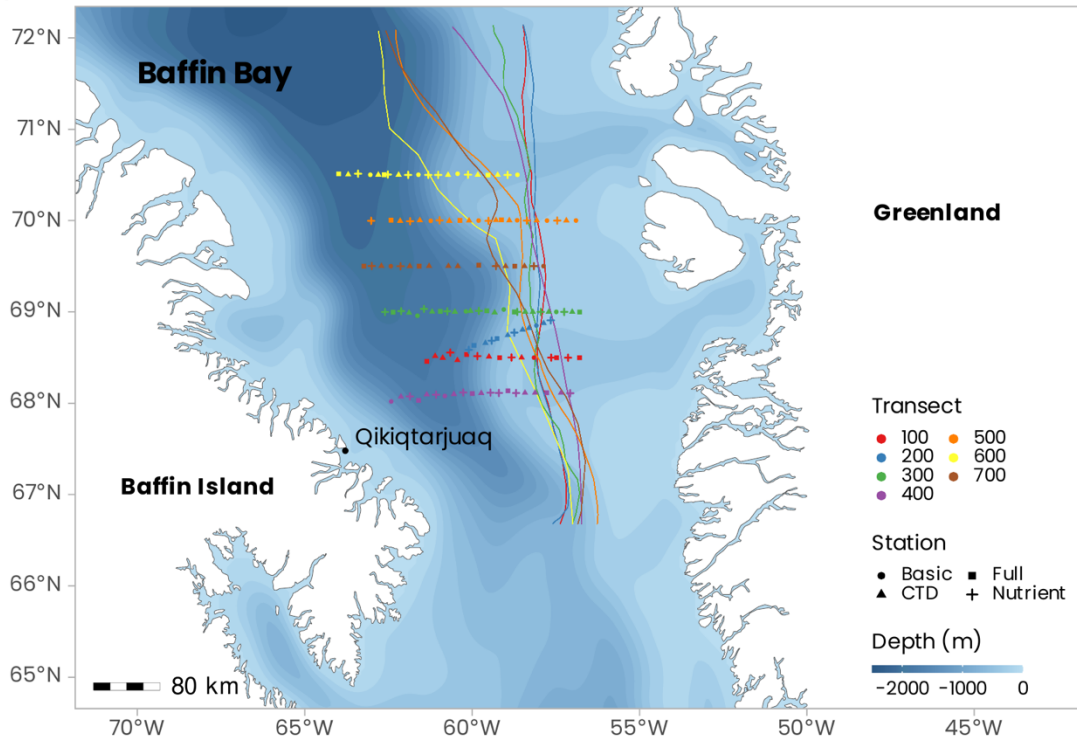
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FIGURES

A



B

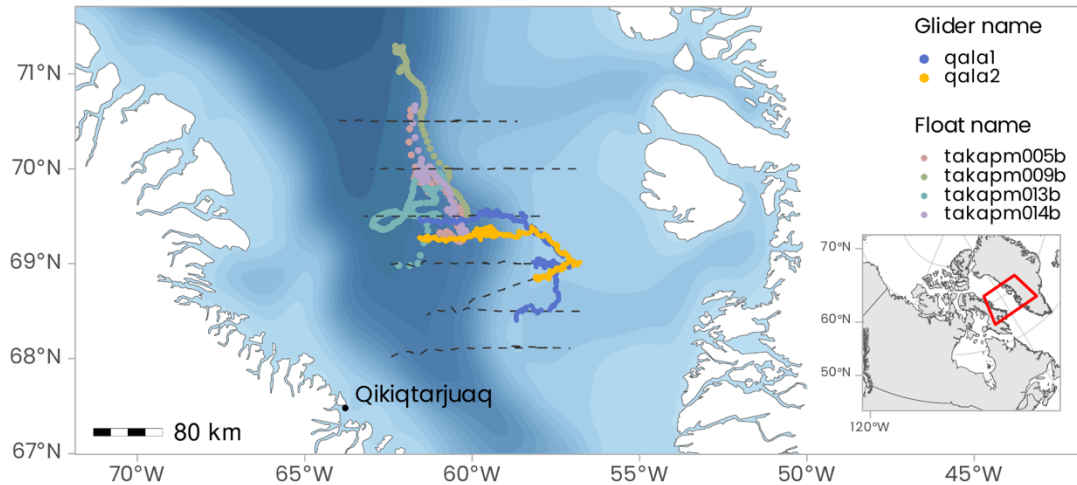


Figure 1: Cruise maps showing A. the seven transects and all stations with shape distinction between station type (FULL, BASIC, NUT and CTD, see Table 1 for description). Colored lines indicate the position of the ice edge (sea ice concentration at 80%) on each transect at date of sampling; ice cover persisted on the western side of Baffin Bay, while the eastern side cleared earlier. Starting date of each transect sampling is as follows: June 9th (100), June 14th (200), June 17th (300), June 24th (400), June 29th (500), July 3rd (600) and July 7th (700). B. Tracks of the 4 BGC-Argo floats deployed during the Green Edge cruise over their first 101 days of life, and the journey of the two Slocum gliders (blue and orange). The dashed line represents the ship track along the seven transects.

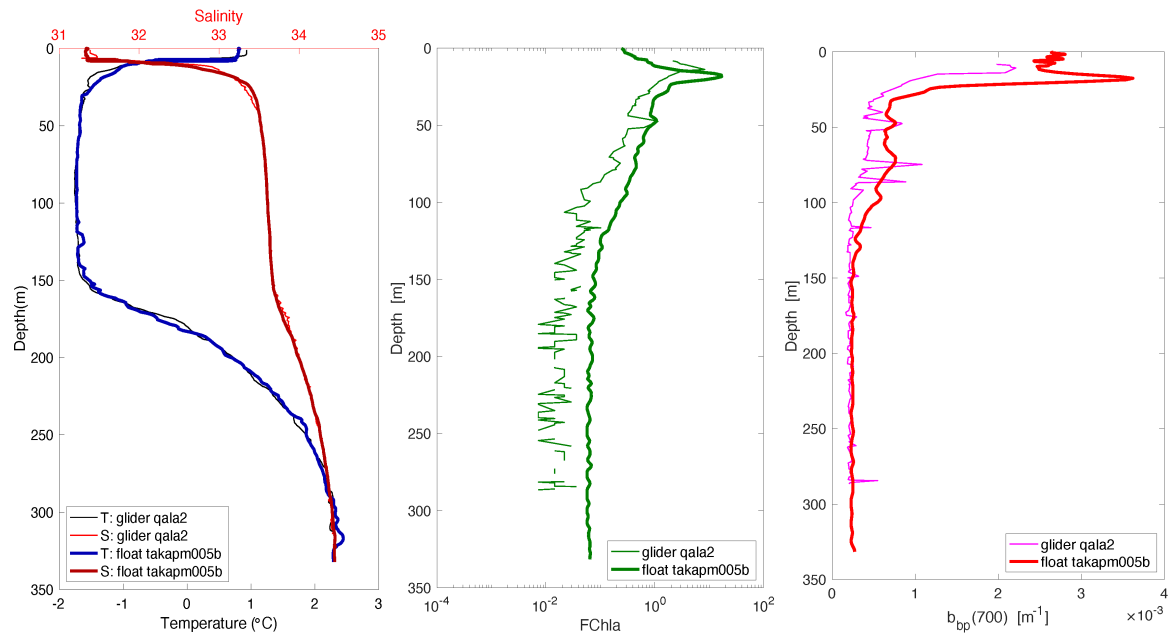


Figure 2: Comparison between salinity (psu), temperature (°C), chlorophyll fluorescence (relative units) and backscattering at 700 nm ($b_{bp}(700)$, m^{-1}) data from the qala2 glider and the takapm005b BioArgo float at their closest common position (Fig. 1B).

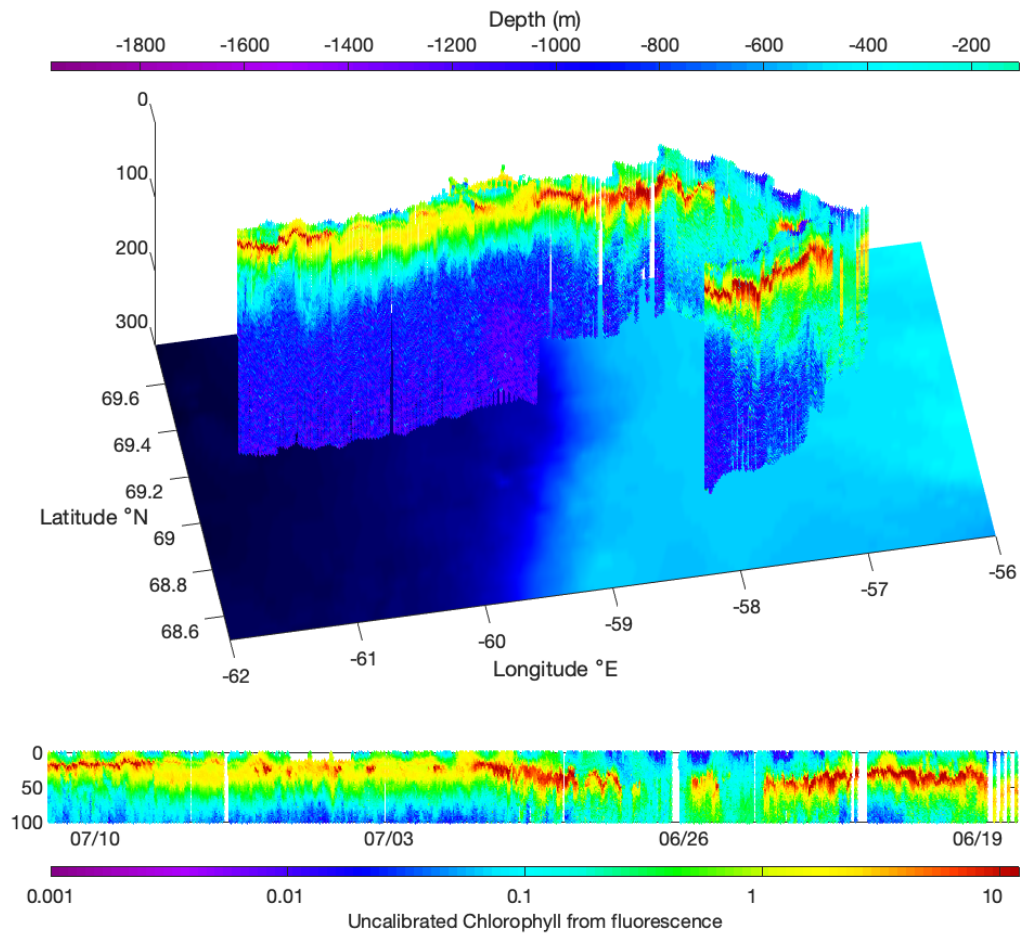


Figure 3: Chlorophyll fluorescence measured by the Glider qala2 during its 23-day journey (Fig. 1B).

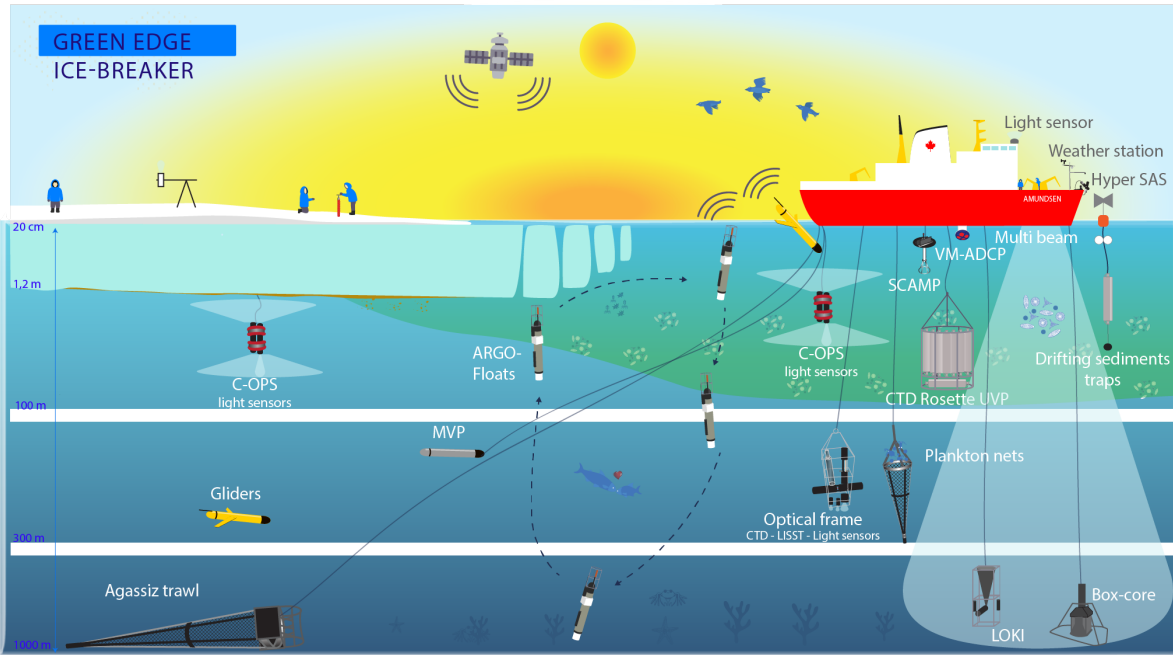


Figure 4A: Representation of all operations carried out during the Green Edge campaign.

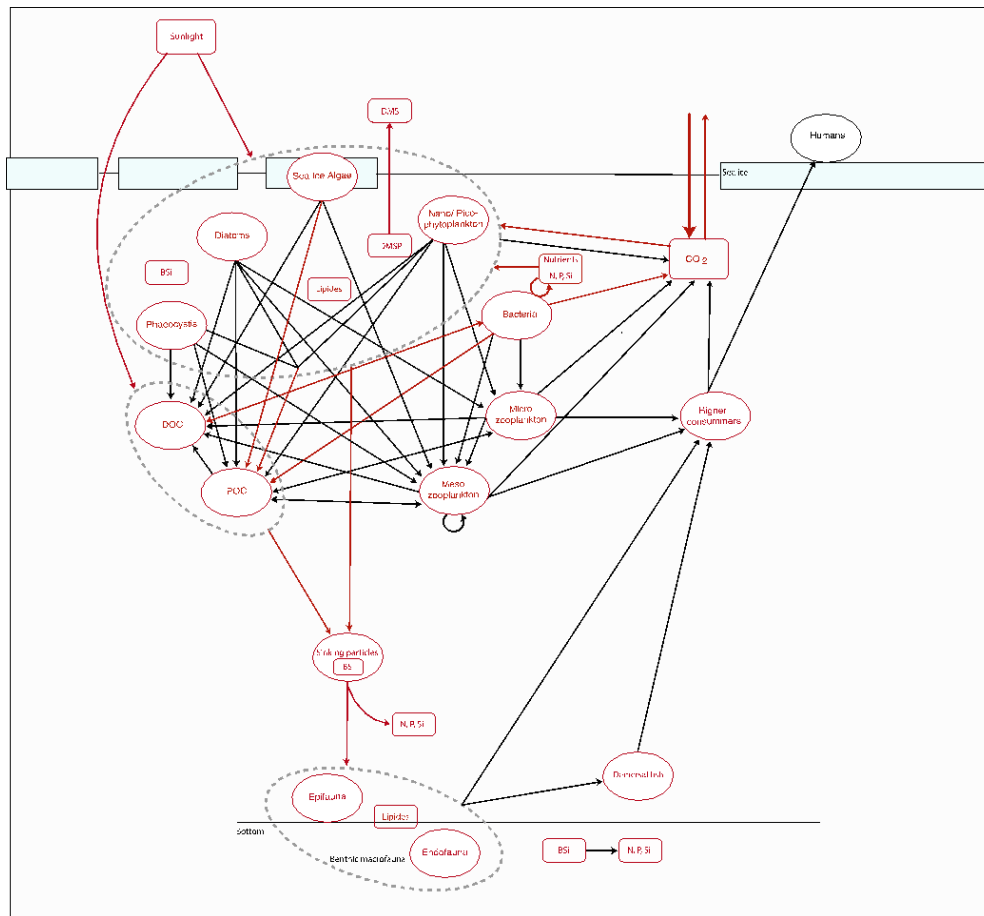


Figure 4B: Schematic representation of all stocks (bubbles) and fluxes (arrows) measured during the Green Edge cruise as part of the MIZ trophic system. Measured and calculated variables are represented in red.

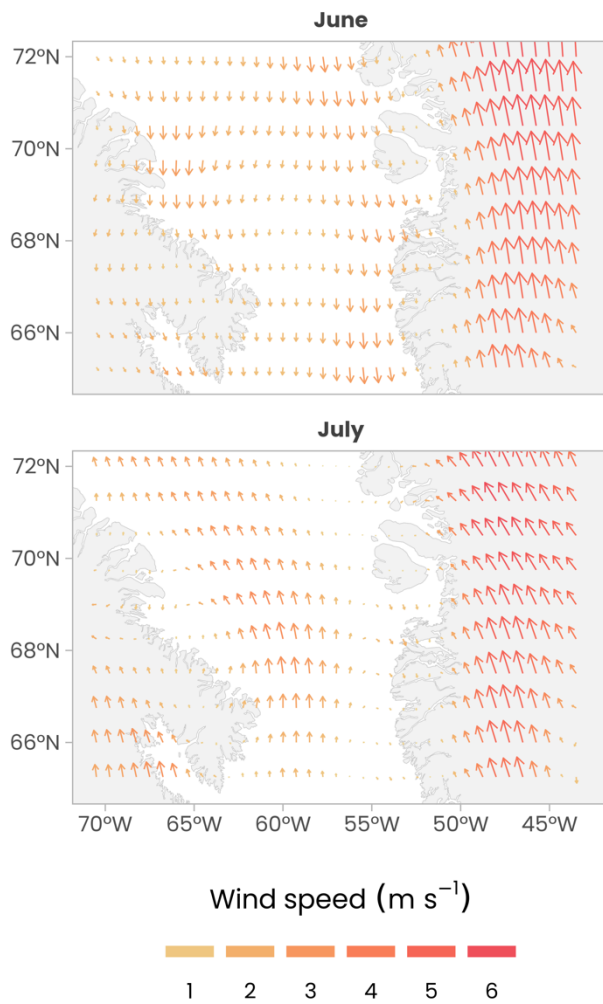


Figure 5: Average wind direction and speed (arrow length and color) over Baffin Bay in June 2016 (top panel) and July 2016 (bottom panel); CCMP wind vector analysis product (V2.0, Atlas et al., (2011)).

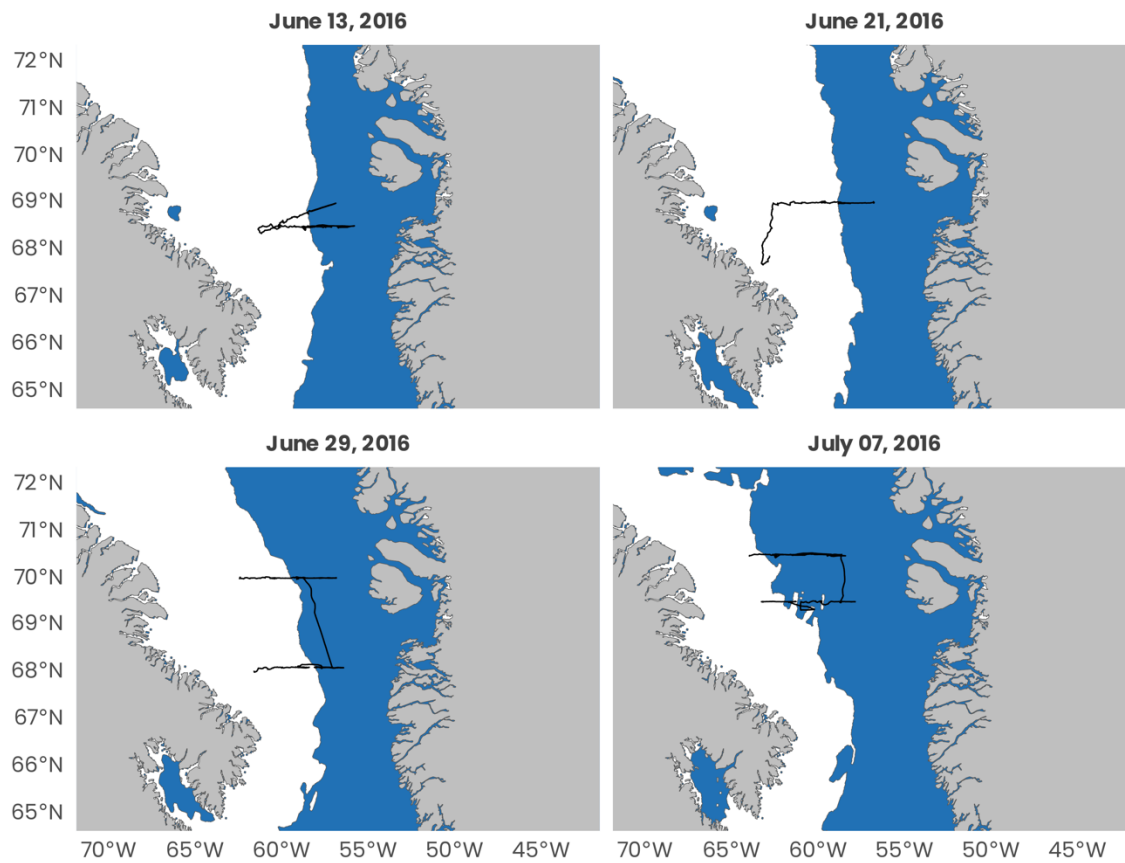


Figure 6: Average weekly sea-ice extent for each of the four 8-day periods during the Green Edge cruise. Date is the median date of each 8-day sampling period. The white areas represent sea ice and the blue areas the open water. Black lines represent the ship route during said 8-day period.

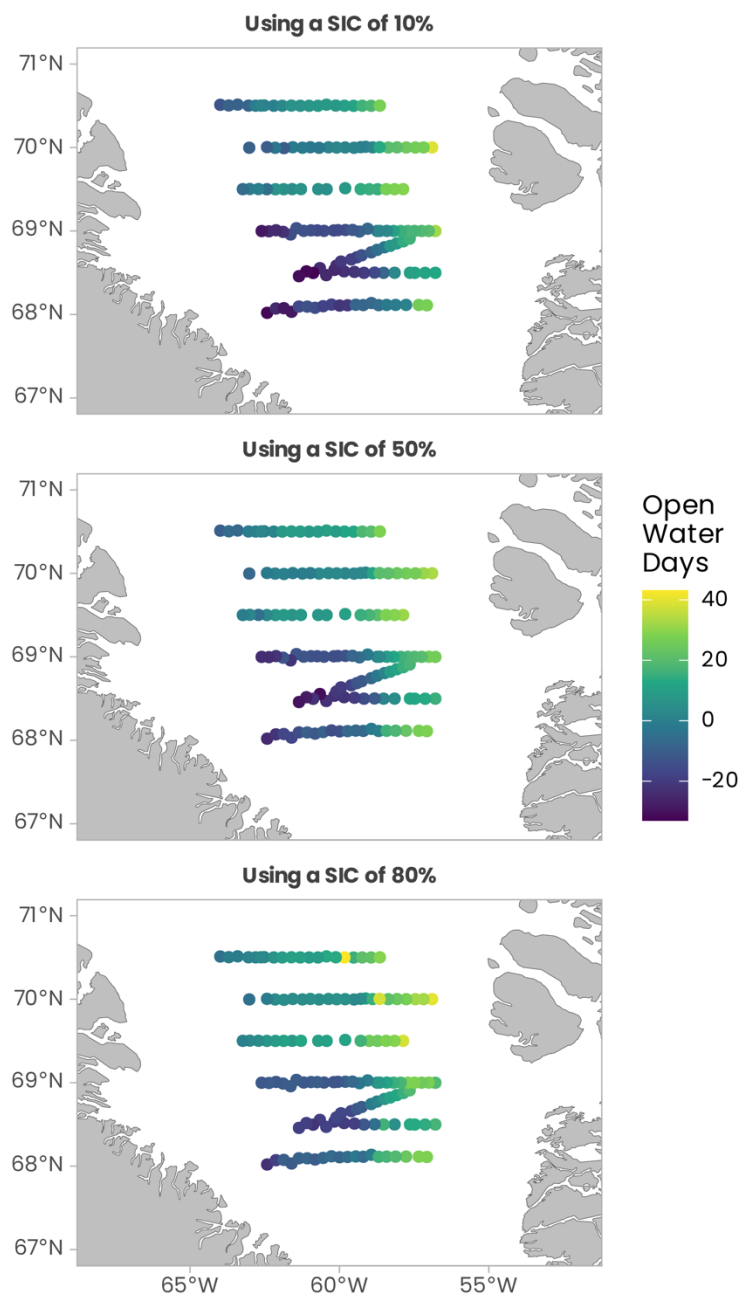


Figure 7: Open water days (OWD) before sampling (values in days). The yellow colour corresponds to positive values meaning the water was already free of ice on the day of sampling. Bluer darker values correspond to stations that were still covered in ice at sampling date (negative values). Values of OWD can be computed using different SIC (sea-ice coverage) values: top panel SIC = 10%, middle panel SIC = 50% and bottom panel SIC = 80%.

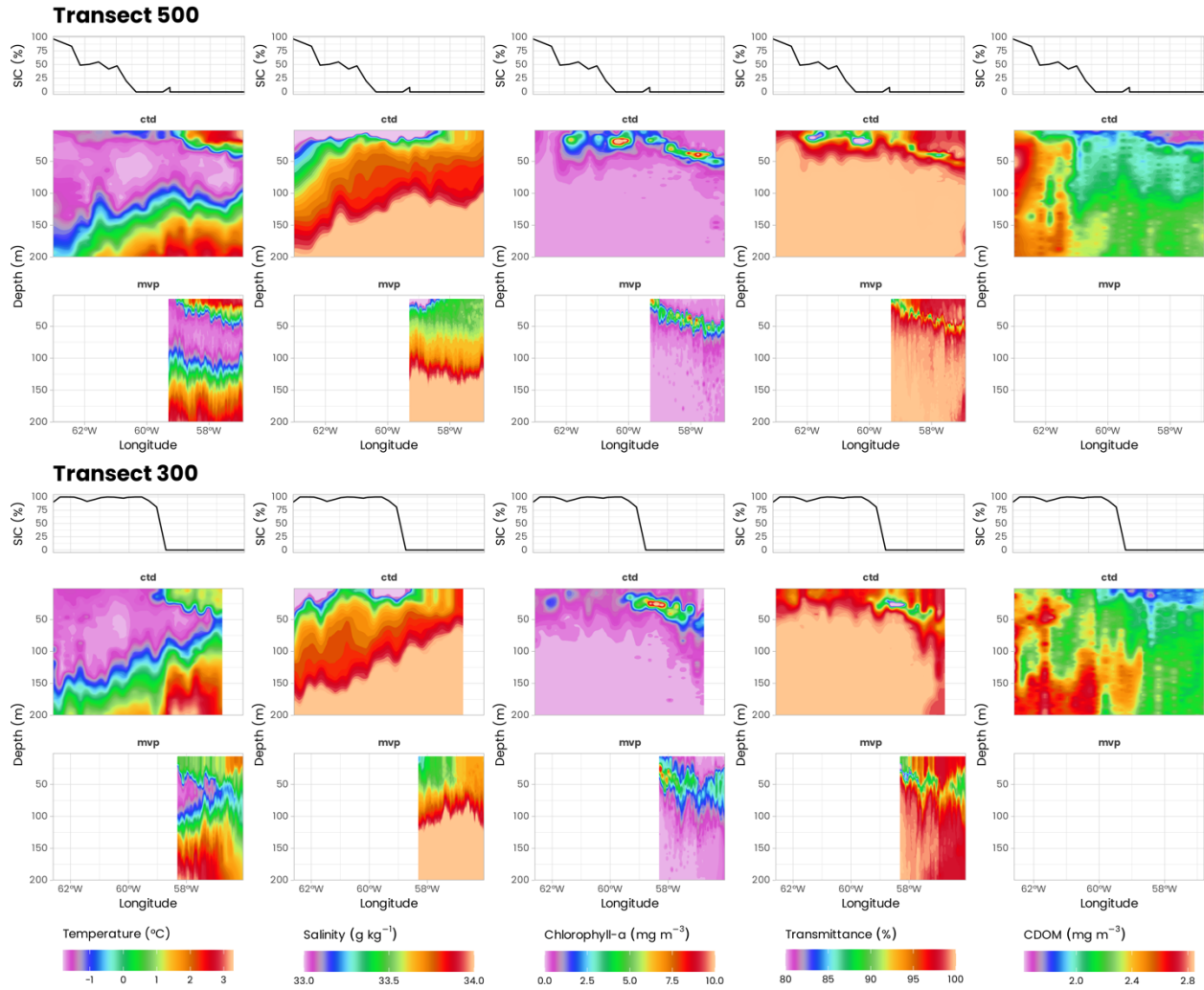


Figure 8: Physical properties of seawater along transects 500 (70° North) and 300 (69° North). Middle and bottom panels show data recorded by sensors deployed on the rosette and on the MVP, respectively. From left to right: temperature (°C), salinity (g/Kg), chlorophyll-*a* concentration (mg m⁻³), transmittance (%) and CDOM concentration (mg m⁻³) as a function of depth (m, y axis) and longitude (degrees West, x axis). Topmost panels show ice coverage (%) at corresponding longitudes. Note that the MVP did not carry a CDOM sensor.

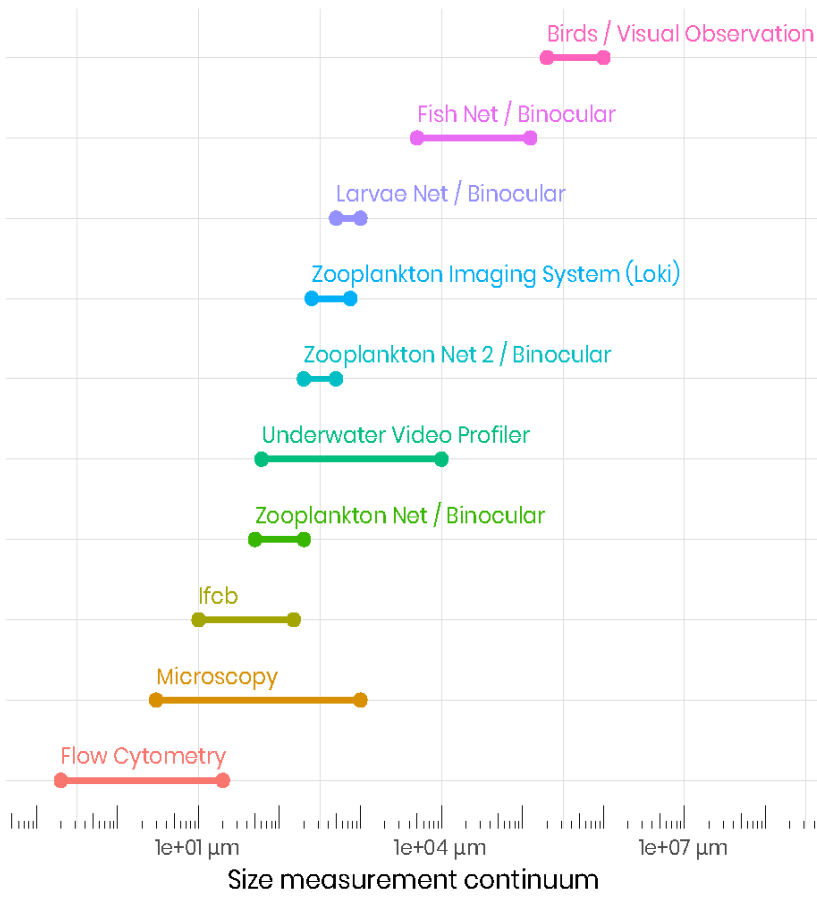


Figure 9: Schematic of the biological sampling size continuum across the various methods and tools used during the Green Edge cruise.

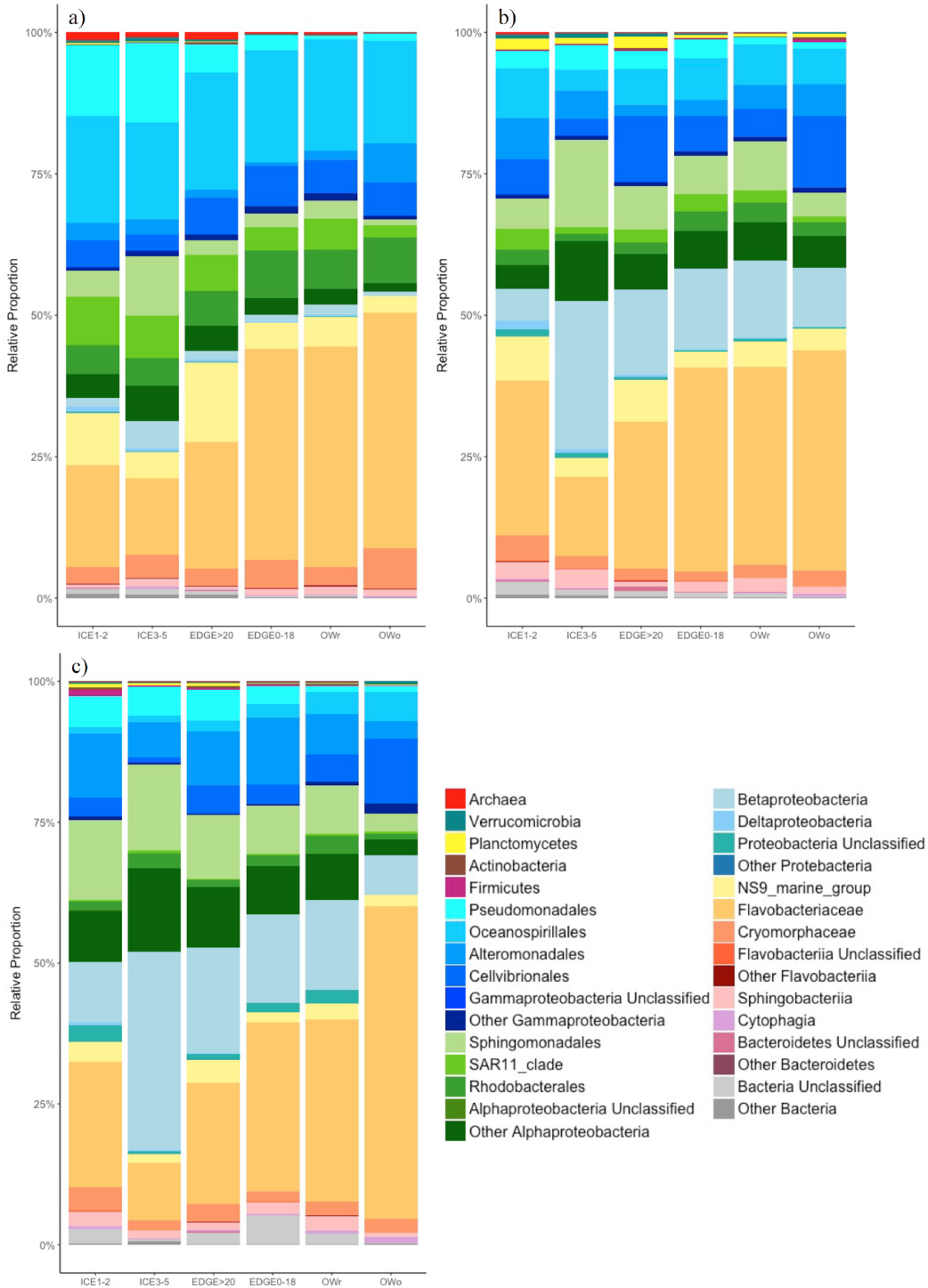


Figure 10: Relative proportions of the different microbial taxa within the different groups of samples, a) for the 0.2-3 μm size fraction (free-living bacteria), b) for the 3-20 μm size fraction (particle-attached bacteria) and c) for the size fraction > 20 μm (particle-attached bacteria). ICE1-2: Ice stations in the transects 100 and 200; ICE3-5: Ice stations in transects 300 and 500; EDGE0-18: Samples between 0 and 18 m depth from the edge stations (stations 107, 204, and 312); EDGE> 20: Samples greater than 20 m deep from the edge stations (stations 107, 204, and 312); OWr: Open water stations where the ice had receded between 2 and 8 days previously; OWo: Open water stations where the ice has receded more than 15 days previously.

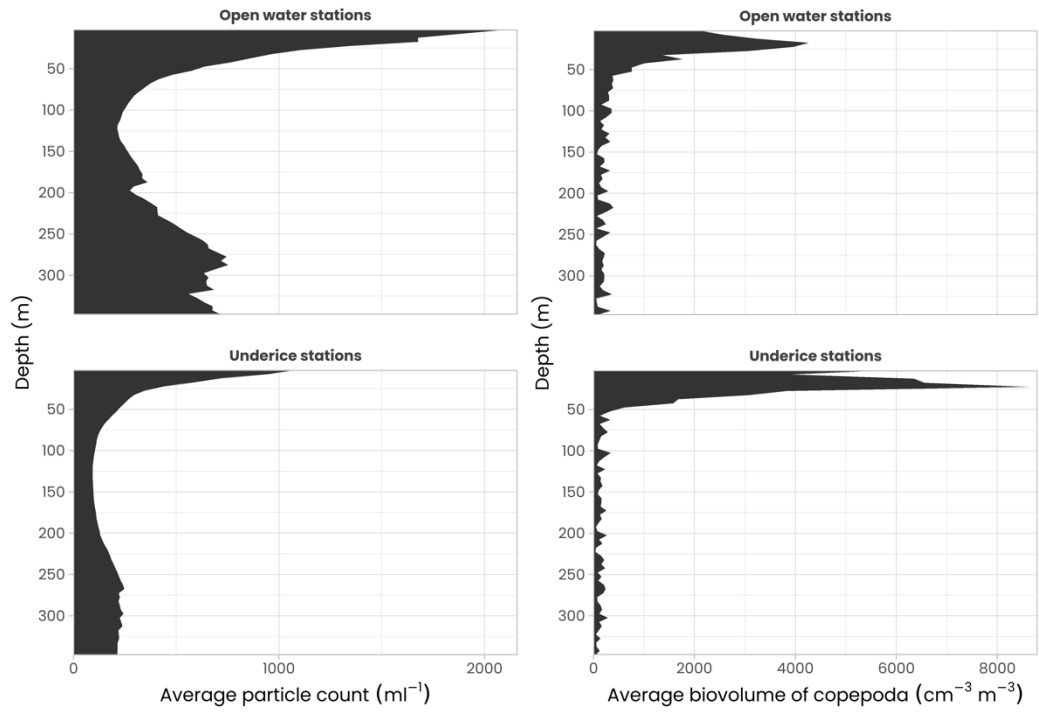


Figure 11: Underwater Vision Profiler data. Average vertical profiles of particle concentration (per ml; left panels) and copepod volumetric fraction (cm³ m⁻³; right panels) at open-water (top panels) and ice-covered (bottom panels) stations over the top 350 m of the water column.



Figure 12: Relative abundance (%) of main phytoplankton groups (colored bars) at the surface (left) and at the subsurface chlorophyll maximum (SCM; right) for all stations analyzed. Stations are sorted according to their OWD value (Fig. 7). Top panel represents the chlorophyll *a* concentration (mg m^{-3}) at the relevant depths for each station.

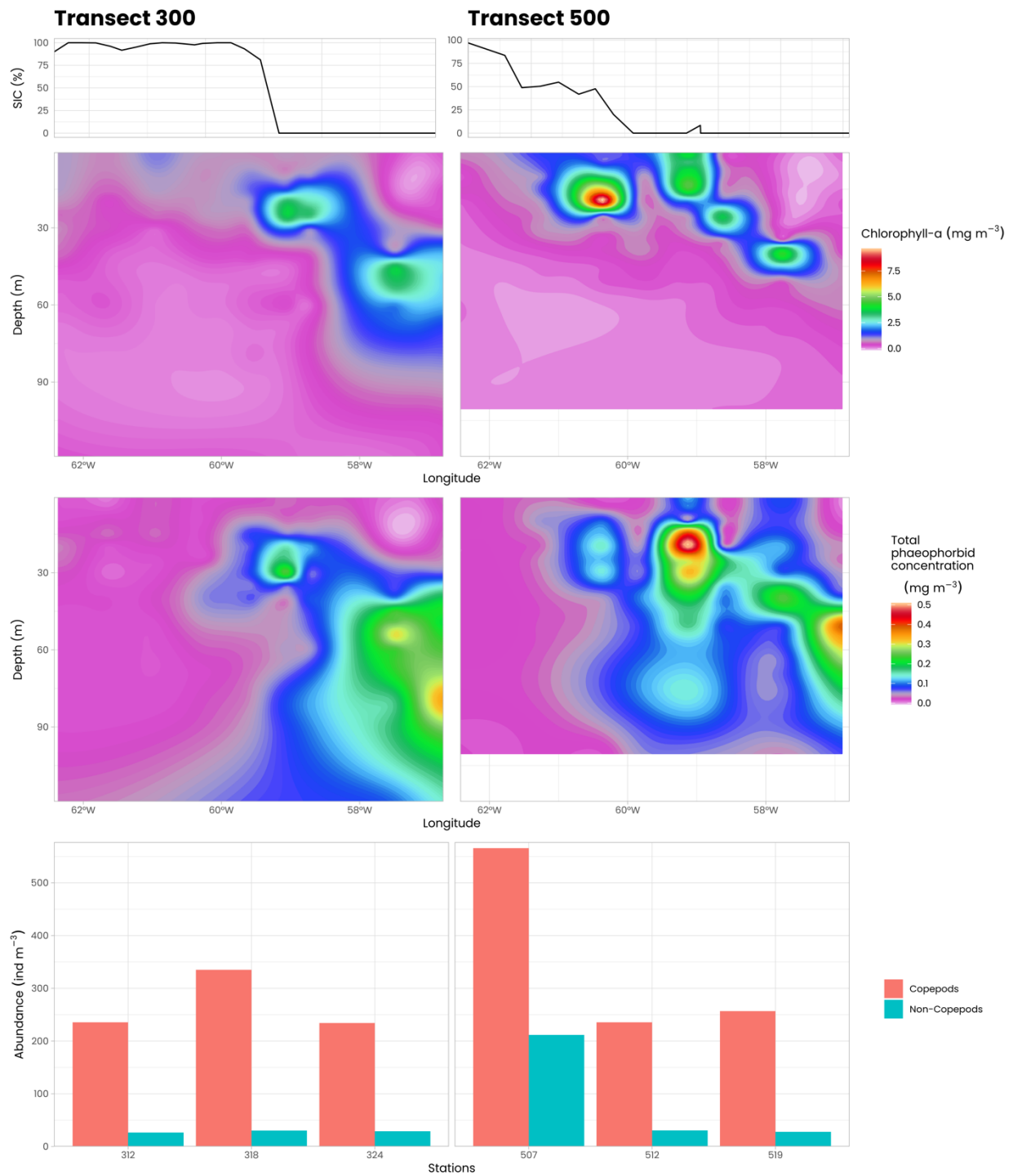


Figure 13: Water column chlorophyll *a* concentration (mg m^{-3} , top panels), and total phaeophorbide *a* concentration (mg m^{-3} , bottom panels) along transects 300 (69° North, left panels) and 500 (70° North, right panels), as measured by HPLC on discrete samples obtained from the Niskin bottles at every FULL and BASIC station. Bottom panels, Abundance of zooplankton (ind m^{-3}), copepods in red and non-copepods in blue, at each of the three FULL stations sampled on each transect (lower station numbers towards the east). The top graphs indicates the SIC at each station at the time of sampling.

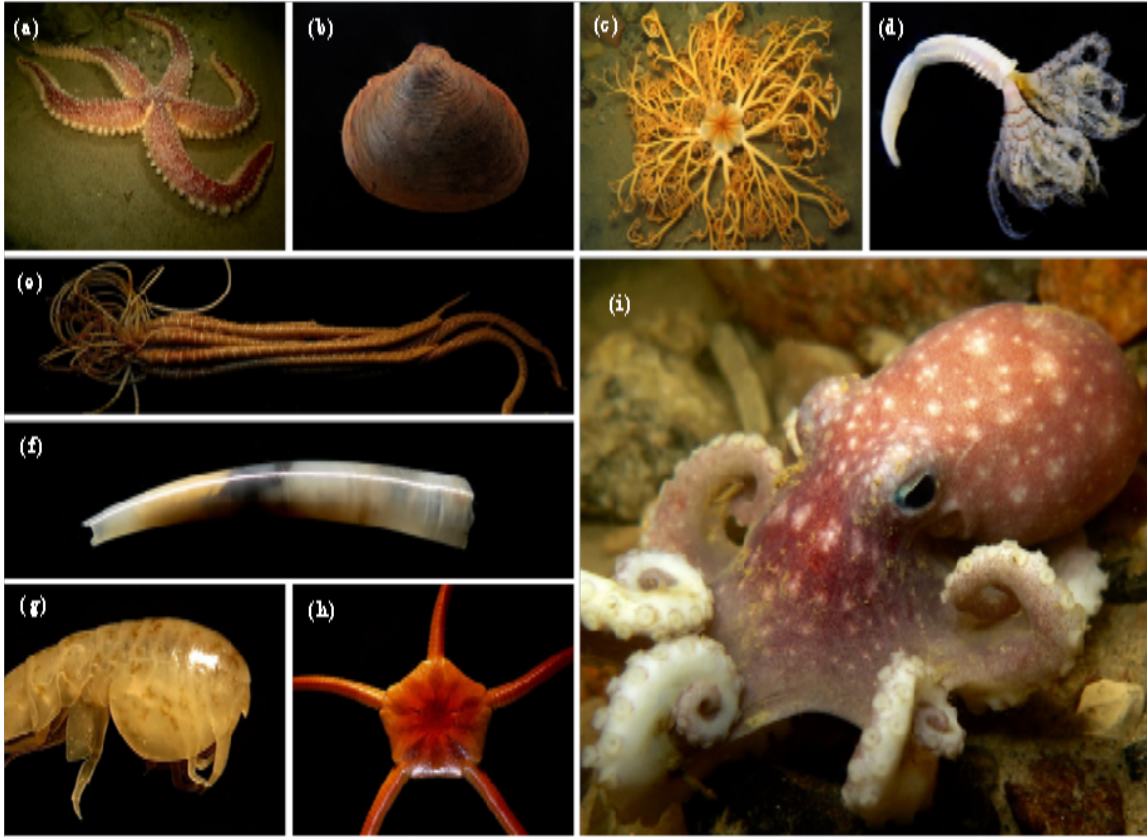


Figure 14: Examples of benthic organisms. (a) *Urasterias lincki*, (b) *Astarte borealis*, (c) *Gorgonocephalus eucnemis*, (e) *Brachiomma* sp., (f) *Heliometra glacialis*, (g) *Siphonodentalium lobatum*, (h) *Stegocephalus inflatus*, (i) *Ophiopleura borealis*, *Bathypolypus* sp. (pictures by benthic ecology lab Université Laval, Gonzalo Bravo).

Table 1: List of operations carried out during the Green Edge cruise at each different station type. [(+) opportunistic sampling only]. CTD conductivity temperature depth, AOP apparent optical properties, C-OPS compact optical profiling system, IOP inherent optical properties, BB9 backscattering meter, CDOM colored dissolved organic matter, SCAMP self-contained autonomous microprofiler, LOKI lightframe on-site key species investigation.

OPERATION TYPE	FULL	BASIC	NUT	CTD
<i>CTD cast (see Table 2 for sensors list)</i>	+	+	+	+
<i>Water sampling (niskin/rosette)</i>	+	+	+	
<i>AOP profile (C-OPS)</i>	+	+		
<i>IOP profile (CTD, BB9, a-Sphere, CDOM and EcoTriplet fluorometers)</i>	+			
<i>SCAMP profile</i>	+			
<i>LOKI and vertical integrated Plankton nets</i>	+			
<i>Mesozoic plankton trawl net</i>	+			
<i>Hydro-bios MultiNet Maxi</i>	+			
<i>Beam trawl</i>	+	(+)		
<i>Agassiz trawl</i>	+			
<i>Box core</i>	+	(+)		
<i>Ice sampling (when ice covered)</i>	+			
<i>Sediment traps</i>	<i>Deployed only once for long term sampling</i>			
TOTAL	26	18	49	51

Table 2: List of the sensors attached to the rosette carousel. CDOM colored dissolved organic matter, PAR photosynthetic available radiation, ADCP Acoustic doppler current profiler, UVP underwater vision profiler.

SENSOR	TYPE	UNIT	SERIAL #	CALIBRATION	# CASTS
<i>Temperature</i>	SBE 3plus	ITS-90 deg C	03P4318	12/2015	203
<i>Conductivity</i>	SBE 4	mS/cm	42696	12/2015	203
<i>Pressure</i>	Paroscientific Digiquartz®	db	679	01/2016	203
<i>Oxygen</i>	SBE 43	mL / L	430427	12/2015	203
<i>CDOM</i>	WetLabs	mg / m ³	2344	01/2016	203
<i>Fluorescence</i>	Seapoint	µg / L	3120	01/2016	203
<i>Transmissometer</i>	WetLabs C-Star	%	CST671DR	01/2016	87
<i>Sonar Altimeter</i>	Benthos PSA-916	m	1065	02/2016	203
<i>PAR / Irradiance</i>	QCP-2300 Biospherical Instr.	µEm ⁻² s ⁻¹	4664	01/2014	203
<i>Nitrate</i>	MBARI-ISUS Satlantic	µM	132	05/2016	104
<i>L-ADCP</i>	WHM300-RDI	NA		03/2016	203
<i>Particles</i>	UVP5 Hydroptic	NA		NA	203

Table 3. List of all the variables sampled during the Green Edge cruise.

VARIABLE	METHOD	SAMPLING METHOD	ACCESS TO DATASET	NAME OF FILE	PI
<i>¹⁵N-Nitrate assimilation</i>	¹⁵ N spiking - incubation - mass-spectrometry	Rosette Deck incubations	https://www.seanoe.org/data/00752/86417/	Primary Production incubations experiments	Tremblay J. E.
<i>¹⁵N-Nitrate primary production (¹³C)</i>	¹⁵ N spiking - incubation - mass-spectrometry	Rosette Deck incubations	https://www.seanoe.org/data/00752/86417/	Primary Production incubations experiments	Tremblay J. E.
<i>¹⁵N-Urea assimilation</i>	¹⁵ N spiking - incubation - mass-spectrometry	Rosette Deck incubations	https://www.seanoe.org/data/00752/86417/	Primary Production incubations experiments	Raimbault P. / Garcia N.
<i>Above water Reflectance (Rrs(0⁺))</i>	C-OPS Biospherical Instr.	Profile mode	https://www.seanoe.org/data/00752/86417/	C-OPS data	Belanger S.
<i>Above water Reflectance (Rrs(0⁺))</i>	Radiometer (Satlantic HyperSAS)	Above-water sensor	https://www.seanoe.org/data/00752/86417/	Remote Sensing Reflectance (Rrs) measured by the HyperSas	Belanger S.
<i>Absorption (non-algal particles)</i>	Spectrophotometer (filters)	Rosette water sample	https://www.seanoe.org/data/00752/86417/	Particulate absorption	Bricaud A. / Sciandra A. / Matsuoka A.
<i>Absorption (particulate matter)</i>	Spectrophotometer (filters)	Rosette water sample	https://www.seanoe.org/data/00752/86417/	Particulate absorption	Bricaud A. / Sciandra A. / Matsuoka A.
<i>Absorption coefficient (total)</i>	HOBILabs a-sphere	In-water profiler	http://www.obs-vlfr.fr/proof/ftpv/greenedge/db/DATA/AMUNDSEN/IOP/	GreenEdge_hydroscat_Asp here	Belanger S.
<i>Acoustic determination of fish presence</i>	Echosounder	Continuous on way	https://www.polardata.ca/pdcsearch/?doi_id=12841	Multiple files available. Data search required	Fortier L.
<i>Air Relative Humidity</i>	Humidity Sensor	Meteorological Tower	https://www.seanoe.org/data/00752/86417/	Meteorological Tower data	Else B. / Burgers T.
<i>Air Temperature</i>	Temperature probe	Meteorological Tower	https://www.seanoe.org/data/00752/86417/	Meteorological Tower data	Else B. / Burgers T.

VARIABLE	METHOD	SAMPLING METHOD	ACCESS TO DATASET	NAME OF FILE	PI
<i>Alkalinity total (TA)</i>	Potentiometry	Rosette water sample	https://www.seanoe.org/data/00752/86417/	Bottle salinity, DIC concentration and Delta 180	Else B. / Miller L.
<i>Ammonium (assimilation)</i>	Isotopic dilution ¹⁵ N	Rosette Deck incubations	https://www.seanoe.org/data/00752/86417/	Primary Production incubations experiments	Raimbault P. / Garcia N.
<i>Ammonium (regeneration)</i>	Isotopic dilution ¹⁵ N	Rosette Deck incubations	https://www.seanoe.org/data/00752/86417/	Primary Production incubations experiments	Raimbault P. / Garcia N.
<i>Ammonium concentration</i>	Fluorescence	Rosette water sample	https://www.seanoe.org/data/00752/86417/	Water column nutrient concentrations	Tremblay J. E.
<i>Backscattering coefficient</i>	BGC-Argo profiler	In-water profiler	http://www.obs- vlfr.fr/proof/php/GREENEDGE/greenedge autonomous.php	multiple	Babin M.
<i>Backscattering coefficient (6 wavelength)</i>	Hydroscat-6	In-water profiler	http://www.obs- vlfr.fr/proof/ftpv/greenedge/db/DATA /AMUNDSEN/IOP/	GreenEdge_hydroscat_Asp here	Belanger S.
<i>Bacterial abundance</i>	Flow Cytometry	Rosette water sample	https://www.seanoe.org/data/00752/86417/	photosynthetic and non-photosynthetic eukaryotes and prokaryotes concentration (flow cytometry)	Vaulot D.
<i>Bacteria infected by virus</i>	Electron Microscopy	Rosette water sample	http://www.obs- vlfr.fr/proof/ftpv/greenedge/db/DATA /AMUNDSEN/EM/	direct	Joux F.
<i>Bacterial diversity</i>	16S rRNA Illumina sequencing	Rosette water sample	https://www.seanoe.org/data/00752/86417/	Bacterial production and respiration	Joux F.
<i>Bacterial production</i>	Leucine- ³ H incorporation	Rosette water sample	https://www.seanoe.org/data/00752/86417/	bacterial production and respiration	Joux F.
<i>Bacterial respiration (whole community)</i>	O ₂ consumption - Winkler - Incubations	Rosette water sample	https://www.seanoe.org/data/00752/86417/	bacterial production and respiration	Joux F.

VARIABLE	METHOD	SAMPLING METHOD	ACCESS TO DATASET	NAME OF FILE	PI
<i>Bacterial salinity- and light-induced biomarkers</i>	GC/MS	Rosette water sample	http://www.obs-vlfr.fr/proof/ftp/greenedge/db/DATA/AMUNDSEN/Bact_Viab/	direct	Rontani J.-F. / Amiraux R. / Burot C.
<i>Bacterial viability</i>	Incubation	Rosette water sample	http://www.obs-vlfr.fr/proof/ftp/greenedge/db/DATA/AMUNDSEN/Bact_Viab/	direct	Rontani J.-F. / Amiraux R. / Burot C.
<i>Bathymetry</i>	Kongsberg EM302 multi-beam echosounder	Continuous horizontal	https://geoapp.bibl.ulaval.ca/Home/Ind ex	search	Lajeunesse P. / Joyal G. / Brouard E.
<i>Benthic ammonium flux</i>	Incubations - Colorimetry	Box corer	https://www.seano.e.org/data/00752/86417/	Sediment incubations data	Morata N.
<i>Benthic carbon content</i>	CN analyser	Box corer	https://www.seano.e.org/data/00752/86417/	Sediment incubations data	Morata N.
<i>Benthic Macrofauna abundance</i>	Microscopy	Box corer	https://www.seano.e.org/data/00752/86417/	Abundance and weight of benthic organisms	Archambault P.
<i>Benthic Macrofauna biomass</i>	Wet weight	Box corer	https://www.seano.e.org/data/00752/86417/	Abundance and weight of benthic organisms	Archambault P.
<i>Benthic Macrofauna diversity</i>	Microscopy	Box corer	https://www.seano.e.org/data/00752/86417/	Abundance and weight of benthic organisms	Archambault P.
<i>Benthic nitrate flux</i>	Incubations - Colorimetry- Autoanalyzer	Box corer	https://www.seano.e.org/data/00752/86417/	Sediment incubations data	Morata N.
<i>Benthic nitrite flux</i>	Incubations - Colorimetry- Autoanalyzer	Box corer	https://www.seano.e.org/data/00752/86417/	Sediment incubations data	Morata N.
<i>Benthic nitrogen content</i>	CN analyser	Box corer	https://www.seano.e.org/data/00752/86417/	Sediment incubations data	Morata N.
<i>Benthic phosphate flux</i>	Incubations - Colorimetry- Autoanalyzer	Box corer	https://www.seano.e.org/data/00752/86417/	Sediment incubations data	Morata N.

VARIABLE	METHOD	SAMPLING METHOD	ACCESS TO DATASET	NAME OF FILE	PI
<i>Benthic silicic acid flux</i>	Incubations - Colorimetry - Autoanalyzer	Box corer	https://www.seanoe.org/data/00752/86417/	Sediment incubations data	Morata N.
<i>Chlorophyll a and Phaeopigments concentration</i>	Fluorimetry	Sea ice core	https://www.seanoe.org/data/00752/86417/	Chlorophyll and Phaeopigment concentration (fluorescence technique)	Bruyant F. / Babin M.
<i>Chlorophyll a and Phaeopigments concentration</i>	Fluorimetry	Rosette water sample	https://www.seanoe.org/data/00752/86417/	Chlorophyll and Phaeopigment concentration (fluorescence technique)	Bruyant F. / Babin M.
<i>Chlorophyll a and Phaeopigments concentration (benthic)</i>	Fluorometric analysis	Box corer	http://www.obs-vlfr.fr/proof/ftpv/greenedge/db/DATA/AMUNDSEN/SEDIMENT_BC/	GE_Amundsen_Sediment_BoxCore.csv	Archambault P.
<i>Chlorophyll a and Phaeopigments concentration (benthic)</i>	Fluorometric analysis	Box corer	https://www.seanoe.org/data/00752/86417/	Sediment incubations data	Morata N.
<i>Chlorophyll a fluorescence</i>	Seapoint fluorometer	In-water profiler	https://www.seanoe.org/data/00752/86417/	CTD data 2.0 m resolution with fluorescence	Guillot P. / Gombault C.
<i>Chlorophyll a fluorescence</i>	Fluorescence (Wetlabs)	In-water profiler	http://www.obs-vlfr.fr/proof/ftpv/greenedge/db/DATA/AMUNDSEN/IOP/	GreenEdge_AMundsen_CTD_IOPs	Belanger S.
<i>Chlorophyll a fluorescence</i>	Fluorometer	Continuous horizontal	https://www.seanoe.org/data/00752/86417/	CTD data 2.0m resolution with fluorescence	Guillot P. / Gombault C.
<i>Chlorophyll a fluorescence</i>	BGC-Argo profiler	In-water profiler	http://www.obs-vlfr.fr/proof/php/GREENEDGE/greenedge_autonomous.php	multiple	Babin M.
<i>Chlorophyll a fluorescence</i>	Wetlabs FLRTD	Moving Vessel Profiler	https://www.seanoe.org/data/00752/86417/	Moving vessel profiler data	Morisset S. / Gombault C.
<i>Chlorophyll-a concentration</i>	Fluorometer	Drifting sediment trap	https://www.seanoe.org/data/00752/86417/	drifting traps data (25m depth)	Lalande C.

VARIABLE	METHOD	SAMPLING METHOD	ACCESS TO DATASET	NAME OF FILE	PI
<i>Chromophoric Dissolved Organic Matter absorption</i>	Ultrapath	Rosette water sample	https://www.seanoe.org/data/00752/86417/	Colored Dissolved Organic Matter (CDOM) absorption data	Matsuoka A.
<i>Chromophoric Dissolved Organic Matter fluorescence</i>	Fluorescence (Wetlabs)	In-water profiler	http://www.obs-vlfr.fr/proof/ftpv/greenedge/db/DATA/AMUNDSEN/IOP/	GreenEdge_AMundsen_CT D_IOPs	Belanger S.
<i>Chromophoric Dissolved Organic Matter fluorescence</i>	Fluorescence (Wetlabs, rosette)	In-water profiler	https://www.seanoe.org/data/00752/86417/	CTD data 2.0m resolution with fluorescence	Guillot P. / Gombault C.
<i>Chromophoric Dissolved Organic Matter fluorescence</i>	BGC-Argo profiler	In-water profiler	http://www.obs-vlfr.fr/proof/php/GREENEDGE/greenedge_autonomous.php	multiple	Babin M.
<i>CO₂ partial pressure (pCO₂)</i>	Licor	Continuous horizontal	https://www.seanoe.org/data/00752/86417/	Continuous pCO ₂ and salinity	Else B. / Burgers T.
<i>Conductivity, Temperature and depth (CTD)</i>	Seabird (IOP optical frame)	In-water profiler	http://www.obs-vlfr.fr/proof/ftpv/greenedge/db/DATA/AMUNDSEN/IOP/	GreenEdge_AMundsen_CT D_IOPs	Belanger S.
<i>Conductivity, Temperature and depth (CTD)</i>	Seabird (rosette)	In-water profiler	https://www.seanoe.org/data/00752/86417/	CTD data 2.0m resolution with fluorescence	Guillot P. / Gombault C.
<i>Conductivity, Temperature and depth (CTD)</i>	Seabird	Continuous horizontal	https://www.seanoe.org/data/00752/86417/	CTD data 2.0m resolution with fluorescence	Guillot P. / Gombault C.
<i>Conductivity, Temperature and depth (CTD)</i>	BGC-Argo profiler	In-water profiler	http://www.obs-vlfr.fr/proof/php/GREENEDGE/greenedge_autonomous.php	multiple	Babin M.
<i>Conductivity, Temperature and depth (CTD)</i>	AML micro CTD	Moving Vessel Profiler	https://www.seanoe.org/data/00752/86417/	Moving vessel profiler data	Morisset S. / Gombault C.

VARIABLE	METHOD	SAMPLING METHOD	ACCESS TO DATASET	NAME OF FILE	PI
<i>Cryptophytes (abundance)</i>	Flow cytometry	Rosette water sample	https://www.seanoe.org/data/00752/86417/	photosynthetic and non-photosynthetic eukaryotes and prokaryotes concentration (flow cytometry)	Vaulot D.
<i>Current speed and direction</i>	ADCP (150kHz)	Continuous horizontal	https://www.seanoe.org/data/00752/86417/	Acoustic Doppler Current Profiler (ADCP)	Guillot P. / Gombault C.
<i>Current speed and direction</i>	ADCP (LADCP)	In-water profiler	https://www.seanoe.org/data/00752/86417/	Acoustic Doppler Current Profiler (ADCP)	Guillot P. / Gombault C.
<i>delta ¹⁸O - water</i>	Mass Spectrometry	Rosette water sample	https://www.seanoe.org/data/00752/86417/	Bottle salinity, DIC concentration and Delta ¹⁸ O	Else B. / Mucci A.
<i>Demersal fish diversity</i>	Beam trawl	Fish trawl	https://www.seanoe.org/data/00752/86417/	Demersal fish abundance and sizes (Beam trawl sampling)	Fortier L.
<i>Diacids (Aerosol)</i>	HPAEC-PAD	Atmosphere	http://www.obs- vlfr.fr/proof/ftpv/greenedge/db/DATA/AMUNDSEN/AEROSOLS/	direct	Panagiotopou los C. / Sempere R.
<i>Diatom (abundance)</i>	Microscopy	Rosette water sample	https://www.seanoe.org/data/00752/86417/	Taxonomy data Diatoms abundance inverted microscope	Lafond A.
<i>Diatom frustules (abundance)</i>	Ludox/colloidal silica extraction	Box corer	https://www.seanoe.org/data/00752/86417/	Sediment incubations data	Morata N.
<i>Diatoms (bacilliarophyta) abundance</i>	Inverted microscopy	Rosette water sample	https://www.seanoe.org/data/00752/86417/	Taxonomy data Diatoms abundance inverted microscope	Leblanc K. / Quéguiner B./Cornet V.
<i>Diatoms (bacilliarophyta) taxonomy</i>	Inverted microscopy	Rosette water sample	https://www.seanoe.org/data/00752/86417/	Taxonomy data Diatoms abundance inverted microscope	Leblanc K. / Quéguiner B./Cornet V.
<i>Diffuse attenuation coefficient (K_d)</i>	C-OPS	In-water profiler	https://www.seanoe.org/data/00752/86417/	C-OPS data	Belanger S.

VARIABLE	METHOD	SAMPLING METHOD	ACCESS TO DATASET	NAME OF FILE	PI
<i>Dimethyl sulfide (DMS)</i>	Gas Chromatography-Mass Spectrometry	Rosette water sample	https://www.seanoe.org/data/00752/86417/	DMS and DMSP concentration	Masse G. / Galí M.
<i>Dimethyl sulfide (sea-air flux)</i>	Gas Chromatography-Mass Spectrometry	Rosette water sample	https://www.seanoe.org/data/00752/86417/	DMS and DMSP concentration	Masse G. / Galí M.
<i>Dimethylsulfopropionate (DMSP)</i>	Gas Chromatography-Mass Spectrometry	Rosette water sample	https://www.seanoe.org/data/00752/86417/	DMS and DMSP concentration	Masse G. / Galí M. / Lizotte M. / Hussherr R.
<i>Dissolved inorganic Carbon (DIC)</i>	Coulometry	Rosette water sample	https://www.seanoe.org/data/00752/86417/	Bottle salinity, DIC concentration and Delta ¹⁸ O	Else B. / Miller L.
<i>Dissolved Organic Carbon (DOC)</i>	Wet oxidation	Rosette Deck incubations	https://www.seanoe.org/data/00752/86417/	Dissolved Inorganic and Organic matter concentrations	Raimbault P. / Garcia N.
<i>Dissolved Organic Nitrogen</i>	Wet oxidation	Rosette Deck incubations	https://www.seanoe.org/data/00752/86417/	Dissolved Inorganic and Organic matter concentrations	Raimbault P. / Garcia N.
<i>Dissolved Organic Nitrogen (release)</i>	Isotopic procedure	Rosette Deck incubations	https://www.seanoe.org/data/00752/86417/	Primary Production incubations experiments	Raimbault P. / Garcia N.
<i>Dissolved Oxygen concentration</i>	Seabird 43	In-water profiler	https://www.seanoe.org/data/00752/86417/	CTD data 2.0m resolution with fluorescence	Guillot P. / Gombault C.
<i>Dissolved Oxygen concentration</i>	BGC-Argo profiler	In-water profiler	http://www.obs-vlfr.fr/proof/php/GREENEDGE/greenedge_autonomous.php	multiple	Babin M.
<i>Downward longwave radiation</i>	Pyrgeometer	Atmosphere	https://www.seanoe.org/data/00752/86417/	Downwelling radiation (pyrgeometer)	Else B. / Burgers T.
<i>Downward shortwave radiation</i>	Pyranometer	Atmosphere	https://www.seanoe.org/data/00752/86417/	Downwelling radiation (pyrgeometer)	Else B. / Burgers T.
<i>Downwelling Irradiance (E_d(z))</i>	C-OPS	In-water profiler	https://www.seanoe.org/data/00752/86417/	C-OPS data	Belanger S.

VARIABLE	METHOD	SAMPLING METHOD	ACCESS TO DATASET	NAME OF FILE	PI
<i>Downwelling irradiance above the surface ($E_d(0^+)$)</i>	Radiometer (Satlantic HyperSAS)	Above-water sensor	https://www.seanoe.org/data/00752/86417/	Remote Sensing Reflectance (R_{rs}) measured by the HyperSas	Belanger S.
<i>Downwelling Irradiance above the surface ($E_d(0^+)$)</i>	SBDART	Surface mode	http://www.obs- vlfr.fr/proof/ftpv/greenedge/db/DATA /SBDART/AM2016/	AM2016_SBDART_AllCasts. zip	Babin M. / Galí M.
<i>Downwelling Irradiance above the surface ($E_d(0^+)$)</i>	C-OPS	In-water profiler	https://www.seanoe.org/data/00752/86417/	C-OPS data	Belanger S.
<i>Downwelling Radiance ($E_d(z)$)</i>	BGC-Argo profiler	In-water profiler	<a href="http://www.obs-
vlfr.fr/proof/php/GREENEDGE/greened
ge_autonomous.php">http://www.obs- vlfr.fr/proof/php/GREENEDGE/greened ge_autonomous.php	multiple	Babin M.
<i>Epibenthic fauna abundance</i>	Microscopy	Agassiz trawl	https://www.seanoe.org/data/00752/86417/	Benthic organisms' identification and abundance	Archambault P.
<i>Epibenthic fauna biomass</i>	Wet weight	Agassiz trawl	https://www.seanoe.org/data/00752/86417/	Benthic organisms' identification and abundance	Archambault P.
<i>Epibenthic fauna diversity</i>	Microscopy	Agassiz trawl	https://www.seanoe.org/data/00752/86417/	Benthic organisms' identification and abundance	Archambault P.
<i>Eukaryotic diversity</i>	Metabarcoding	Rosette water sample	https://www.seanoe.org/data/00752/86417/	photosynthetic and non-photosynthetic eukaryotes and prokaryotes concentration (flow cytometry)	Vaulot D.
<i>Fish abundance (Midwater)</i>	IKMT trawl	Fish trawl	https://www.seanoe.org/data/00752/86417/	Pelagic fish abundance and sizes (IKMT sampling)	Fortier L.
<i>Lipid biomarkers concentrations</i>	GC/MS	Collected Organisms (bird)	https://www.seanoe.org/data/00752/86417/	Lipid biomarkers in benthic and sediment fauna	Mosbech A. / Fort J.

VARIABLE	METHOD	SAMPLING METHOD	ACCESS TO DATASET	NAME OF FILE	PI
<i>Microturbulence</i>	SCAMP profiler	In-water profiler	http://www.obs-vlfr.fr/proof/ftp/greenedge/db/DATA/AMUNDSEN/SCAMP/		Vladoiu A. / Dumont D.
<i>Nanoeukaryotes (abundance)</i>	Flow cytometry	Rosette water sample	https://www.seanoe.org/data/00752/86417/	photosynthetic and non-photosynthetic eukaryotes and prokaryotes concentration (flow cytometry)	Vaulot D.
<i>Nitrate (NO₃⁻) assimilation</i>	Isotopic dilution 15N	Deck incubations	https://www.seanoe.org/data/00752/86417/	Primary Production incubations experiments	Raimbault P. / Garcia N.
<i>Nitrate concentration (NO₃⁻)</i>	Colorimetry/Autoanalyser	Rosette water sample	https://www.seanoe.org/data/00752/86417/	Water column nutrient concentrations	Tremblay J. E.
<i>Nitrate concentration (NO₃⁻)</i>	Colorimetry/Autoanalyser	Deck incubations	https://www.seanoe.org/data/00752/86417/	Primary production incubations experiments	Garcia N.
<i>Nitrate concentration (NO₃⁻)</i>	BGC-Argo profiler	In-water profiler	http://www.obs-vlfr.fr/proof/php/GREENEDGE/greenedge_autonomous.php	multiple	Babin M.
<i>Nitrate concentration (NO₃⁻)</i>	MBARI-ISUS Satlantic	In-water profiler	https://www.seanoe.org/data/00752/86417/	CTD data 2.0m resolution with fluorescence	Guillot P. / Gombault C.
<i>Nitrification</i>	¹⁵ N labeling	Deck incubations	https://www.seanoe.org/data/00752/86417/	Primary Production incubations experiments	Raimbault P. / Garcia N.
<i>Nitrite concentration (NO₂⁻)</i>	Colorimetry/Autoanalyser	Rosette water sample	https://www.seanoe.org/data/00752/86417/	Water column nutrient concentrations	Tremblay J. E.
<i>Nitrite concentration (NO₂⁻)</i>	Colorimetry/Autoanalyser	Deck incubations	https://www.seanoe.org/data/00752/86417/	Primary production incubations experiments	Garcia N.
<i>Open Water Days</i>	AMSR	Satellite	https://www.seanoe.org/data/00752/86417/	Days of Open Water (DOW)	Massicotte P. / Ferland J.
<i>Orthosilicic acid (uptake rate)</i>	³² Si absorption	Rosette water sample	https://www.seanoe.org/data/00752/86417/	Silicate uptake rate	Leynaert A. / Quéguiner B./Gallinari M.

VARIABLE	METHOD	SAMPLING METHOD	ACCESS TO DATASET	NAME OF FILE	PI
<i>Orthosilicic acid concentration (Si(OH)₄)</i>	Colorimetry/Autoanalyser	Rosette water sample	https://www.seanoe.org/data/00752/86417/	Water column nutrient concentrations	Tremblay J. E.
<i>Orthosilicic acid concentration (Si(OH)₄)</i>	Colorimetry/Autoanalyser	Deck incubations	https://www.seanoe.org/data/00752/86417/	Primary production incubations experiments	Garcia N.
<i>Orthosilicic acid concentration Si(OH)₄</i>	Technicon	Rosette water sample	https://www.seanoe.org/data/00752/86417/	Silicate absorption kinetics experiments	Leynaert A. / Moriceau B./Gallinari M.
<i>Orthosilicic acid Si(OH)₄ - uptake kinetics</i>	³² Si absorption - incubation	Rosette water sample	https://www.seanoe.org/data/00752/86417/	Silicate absorption kinetics experiments	Leynaert A./ Quéguiner B./Gallinari M.
<i>Particle Size Distribution</i>	UVP-5	In-water profiler	https://www.seanoe.org/data/00752/86417/	Underwater Vision Profiler (UVP) Particles	Picheral M.
<i>Particulate mass</i>	Dry Weight	Drifting sediment trap	https://www.seanoe.org/data/00752/86417/	drifting traps data (25m depth)	Lalande C.
<i>Particulate Nitrogen content</i>	CHN	Drifting sediment trap	https://www.seanoe.org/data/00752/86417/	drifting traps data (25m depth)	Lalande C.
<i>Particulate Organic Carbon (POC)</i>	CHN	Rosette water sample	https://www.seanoe.org/data/00752/86417/	Dry weight, Particulate Carbon and Nitrogen (CHN)	Bruyant F. / Lariviere J. / Babin M.
<i>Particulate Organic Carbon (POC)</i>	High combustion	Deck incubations	https://www.seanoe.org/data/00752/86417/	Dissolved Inorganic and Organic matter concentrations	Raimbault P. / Garcia N.
<i>Particulate Organic Carbon (POC)</i>	CHN	Drifting sediment trap	https://www.seanoe.org/data/00752/86417/	drifting traps data (25m depth)	Lalande C.
<i>Particulate Organic Nitrogen (PON)</i>	CHN	Rosette water sample	https://www.seanoe.org/data/00752/86417/	Dry weight, Particulate Carbon and Nitrogen (CHN)	Bruyant F. / Lariviere J. / Babin M.

VARIABLE	METHOD	SAMPLING METHOD	ACCESS TO DATASET	NAME OF FILE	PI
<i>Particulate Organic Nitrogen (PON)</i>	High combustion	Deck incubations	https://www.seanoe.org/data/00752/86417/	Dissolved Inorganic and Organic matter concentrations	Raimbault P. / Garcia N.
<i>PDMPO uptake</i>	Spectrophotometry/Spectrofluorometry	Rosette water sample	https://www.seanoe.org/data/00752/86417/	Silicification of Diatoms	Leblanc K. / Quéguiner B./Cornet V.
<i>Phosphate concentration (PO₄)³⁻</i>	Colorimetry/Autoanalyser	Rosette water sample	https://www.seanoe.org/data/00752/86417/	Water column nutrient concentrations	Tremblay J. E.
<i>Phosphate concentration (PO₄)³⁻</i>	Colorimetry/Autoanalyser	Deck incubations	https://www.seanoe.org/data/00752/86417/	Primary production incubations experiments	Garcia N.
<i>Photo Eukaryotes (morphology)</i>	Scanning Electron Microscopy	Rosette water sample	https://www.seanoe.org/data/00752/86417/	photosynthetic and non-photosynthetic eukaryotes and prokaryotes concentration (flow cytometry)	Vaulot D.
<i>Photo Eukaryotes Sorted (Morphology)</i>	Scanning Electron Microscopy	Rosette water sample	https://www.seanoe.org/data/00752/86417/	photosynthetic and non-photosynthetic eukaryotes and prokaryotes concentration (flow cytometry)	Vaulot D.
<i>Photosynthetic Available Radiation (PAR)</i>	Biospherical Instrument QCP-2300	In-water profiler	https://www.seanoe.org/data/00752/86417/	CTD data 2.0m resolution with fluorescence	Guillot P. / Gombault C.
<i>Photosynthetic Available Radiation (PAR)</i>	C-OPS	In-water profiler	https://www.seanoe.org/data/00752/86417/	C-OPS data	Belanger S.
<i>Photosynthetic Available Radiation (PAR)</i>	SBDART	Surface mode	http://www.obs-vlfr.fr/proof/ftpv/greenedge/db/DATA/SBDART/AM2016/	AM2016_SBDART_AllCasts.zip	Babin M. / Galí M.
<i>Photosynthetic Available Radiation (PAR)</i>	Radiometer	Atmosphere	https://www.seanoe.org/data/00752/86417/	Downwelling radiation (pyrgeometer)	Else B. / Burgers T.
<i>Photosynthetic Available Radiation (PAR)</i>	BGC-Argo profiler	In-water profiler	http://www.obs-vlfr.fr/proof/php/GREENEDGE/greenedge_autonomous.php	multiple	Babin M.

VARIABLE	METHOD	SAMPLING METHOD	ACCESS TO DATASET	NAME OF FILE	PI
<i>Photosynthetic parameters</i>	¹⁴ C P vs. E curve	Rosette water sample	https://www.seanoe.org/data/00752/86417/	Photosynthetic parameters	Lewis K.
<i>Phytoplankton (taxonomy)</i>	Inverted Microscopy	Rosette water sample		Phytoplankton taxonomy (microscopy)	Babin M.
<i>Phytoplankton (Micro-taxonomy)</i>	Microscopy	Drifting sediment trap	https://www.seanoe.org/data/00752/86417/	drifting traps data (25m depth)	Lalande C.
<i>Phytoplankton (taxonomy)</i>	Imaging flowcytobot	Sea ice core	http://www.obs-vlfr.fr/proof/ftpv/greenedge/db/DATA/AMUNDSEN/IFCB/	multiple	Bruyant F. / Grondin P.L. / Babin M.
<i>Phytoplankton (taxonomy)</i>	Imaging flowcytobot	Rosette water sample	http://www.obs-vlfr.fr/proof/ftpv/greenedge/db/DATA/AMUNDSEN/IFCB/	multiple	Bruyant F. / Grondin P.L. / Babin M.
<i>Phytoplankton cultures</i>	Sorted by flow cytometry, serial dilution, single cell pipetting	Rosette water sample	https://www.seanoe.org/data/00752/86417/	photosynthetic and non-photosynthetic eukaryotes and prokaryotes concentration (flow cytometry)	Vaulot D.
<i>Picoeukaryotes (abundance)</i>	Flow cytometry	Rosette water sample	https://www.seanoe.org/data/00752/86417/	photosynthetic and non-photosynthetic eukaryotes and prokaryotes concentration (flow cytometry)	Vaulot D.
<i>Pigments concentration</i>	HPLC	Rosette water sample	https://www.seanoe.org/data/00752/86417/	phytoplankton pigments concentration (HPLC)	Ras J. / Claustre H. / Dimier C.
<i>Pigments concentration</i>	HPLC	Sea ice core	https://www.seanoe.org/data/00752/86417/	phytoplankton pigments concentration (HPLC)	Ras J. / Claustre H. / Dimier C.
<i>Plankton taxonomy</i>	UVP-5	In-water profiler	https://www.seanoe.org/data/00752/86417/	Underwater Vision Profiler (UVP) zooplankton	Picheral M.
<i>Primary production</i>	¹³ C labeling	Deck incubations	https://www.seanoe.org/data/00752/86417/	Primary Production incubations experiments	Raimbault P. / Garcia N.

VARIABLE	METHOD	SAMPLING METHOD	ACCESS TO DATASET	NAME OF FILE	PI
<i>Quantum Efficiency of PSII (Fv/Fm) (Phytoplankton)</i>	Benchtop PAM - PhytoPAM	Rosette water sample	http://www.obs- vlfr.fr/proof/ftpv/greenedge/db/DATA /AMUNDSEN/PAM/	direct	Joy-Warren H.
<i>Sea ice concentration</i>	Satellite	Surface mode	https://www.seanoe.org/data/00752/8 6417/	Sea ice history	Massicotte P.
<i>Sediment grain size</i>	Laser	Box corer	http://www.obs- vlfr.fr/proof/ftpv/greenedge/db/DATA /AMUNDSEN/SEDIMENT_BC/	GE_Amundsen_Sediment_BoxCore.csv	Archambault P.
<i>Sediment organic content</i>	Lost by ignition	Box corer	http://www.obs- vlfr.fr/proof/ftpv/greenedge/db/DATA /AMUNDSEN/SEDIMENT_BC/	GE_Amundsen_Sediment_BoxCore.csv	Archambault P.
<i>Silica Biogenic (BSi)</i>	BSi extraction	Rosette water sample	https://www.seanoe.org/data/00752/8 6417/	Biogenic and Lithogenic silica concentration	Leblanc K./Quéguiner B./Leynaert A./Moriceau B./Legras J./Gallinari M.
<i>Silica Biogenic (BSi) dissolution rate</i>	dissolution in filtered sea water	Rosette water sample	https://www.seanoe.org/data/00752/8 6417/	Biogenic and Lithogenic silica concentration	Moriceau B./Gallinari M.
<i>Silica Biogenic concentration</i>	BSi extraction	Box corer	https://www.seanoe.org/data/00752/8 6417/	Sediment incubations data	Morata N./Gallinari M.
<i>Silica Lithogenic (LSi)</i>	LSi extraction	Rosette water sample	https://www.seanoe.org/data/00752/8 6417/	Biogenic and Lithogenic silica concentration	Leblanc K./Quéguiner B./Leynaert A./Moriceau B./Legras J./Gallinari M.
<i>Sub-bottom profiles</i>	Knudsen 320 sub-bottom echosounder	Continuous horizontal	https://geoapp.bibl.ulaval.ca/Home/Ind ex	search	Lajeunesse P. / Joyal G. / Brouard E.

VARIABLE	METHOD	SAMPLING METHOD	ACCESS TO DATASET	NAME OF FILE	PI
<i>Sugars (Aerosol)</i>	HPAEC-PAD	Atmosphere	http://www.obs-vlfr.fr/proof/ftpv/greenedge/db/DATA/AMUNDSEN/AEROSOLS/	direct	Panagiotopoulos C. / Sempere R.
<i>Suspended Particulate Material (SPM)</i>	Particle dry weight (gravimetry)	Rosette water sample	https://www.seanoe.org/data/00752/86417/	Dry weight, Particulate Carbon and Nitrogen (CHN)	Bruyant F. / Lariviere J. / Babin M.
<i>Synechococcus (abundance)</i>	Flow cytometry	Rosette water sample	https://www.seanoe.org/data/00752/86417/	photosynthetic and non-photosynthetic eukaryotes and prokaryotes concentration (flow cytometry)	Vaulot D.
<i>Total Organic Carbon (TOC)</i>	Wet oxidation	Rosette Deck incubations	https://www.seanoe.org/data/00752/86417/	Dissolved Inorganic and Organic matter concentrations	Raimbault P. / Garcia N.
<i>Total Organic Nitrogen (TON)</i>	Wet oxidation	Rosette Deck incubations	https://www.seanoe.org/data/00752/86417/	Dissolved Inorganic and Organic matter concentrations	Raimbault P. / Garcia N.
<i>Total Organic Phosphorus (TOP)</i>	Wet oxidation	Rosette Deck incubations	https://www.seanoe.org/data/00752/86417/	Dissolved Inorganic and Organic matter concentrations	Raimbault P. / Garcia N.
<i>Transmittance (of light in water)</i>	CST Wetlabs	Moving Vessel Profiler	https://www.seanoe.org/data/00752/86417/	Moving vessel profiler data	Morisset S. / Gombault C.
<i>Transmittance (of light in water)</i>	C-Star	In-water profiler	https://www.seanoe.org/data/00752/86417/	CTD data 2.0m resolution with fluorescence	Guillot P. / Gombault C.
<i>Upwelling Irradiance ($E_u(z)$)</i>	C-OPS	In-water profiler	https://www.seanoe.org/data/00752/86417/	C-OPS data	Belanger S.
<i>Upwelling radiance (L_u)</i>	C-OPS	In-water profiler	https://www.seanoe.org/data/00752/86417/	C-OPS data	Belanger S.
<i>Variable fluorescence and Rapid Light Curves parameters (Phytoplankton)</i>	Benchtop PAM	Rosette water sample	http://www.obs-vlfr.fr/proof/ftpv/greenedge/db/DATA/AMUNDSEN/PAM/	direct	Joy-Warren H.

VARIABLE	METHOD	SAMPLING METHOD	ACCESS TO DATASET	NAME OF FILE	PI
<i>Weather and navigation</i>	On Board	AVOS	https://www.seanoe.org/data/00752/86417/	Automated Volunteer Observing Ship (AVOS) weather data	Morisset S. / Gombault C.
<i>Wind Direction</i>	Wind monitor	Meteorological Tower	https://www.seanoe.org/data/00752/86417/	Meteorological Tower data	Else B. / Burgers T.
<i>Wind Speed</i>	Wind monitor	Meteorological Tower	https://www.seanoe.org/data/00752/86417/	Meteorological Tower data	Else B. / Burgers T.
<i>Wind speed</i>	Cross Calibrated Multi Platform (CCMP)	Atmosphere	http://www.obs- vlfr.fr/proof/ftpv/greenedge//db/DATA/WIND/DATA/	direct	Massicotte P.
<i>zooplankton (abundance)</i>	Multi nets (Hydro-bios)	Plankton net	https://www.seanoe.org/data/00752/86417/	Zooplankton abundance and diversity (vertical nets)	Fortier L.
<i>Zooplankton (Abundances)</i>	Vertical plankton net 200 µm	Plankton net	https://www.seanoe.org/data/00752/86417/	Zooplankton abundance and diversity (vertical nets)	Fortier L.
<i>Zooplankton (Meso-) (abundance)</i>	Tucker net	Plankton net	https://www.seanoe.org/data/00752/86417/	Ichthyoplankton vertical nets sampling	Fortier L.
<i>Zooplankton (Meso-) (taxonomy)</i>	Tucker net	Plankton net	https://www.seanoe.org/data/00752/86417/	Ichthyoplankton vertical nets sampling	Fortier L.
<i>Zooplankton (taxonomy)</i>	Multi nets (Hydro-bios)	Plankton net	https://www.seanoe.org/data/00752/86417/	Zooplankton abundance and diversity (vertical nets)	Fortier L.
<i>Zooplankton (Taxonomy)</i>	Vertical plankton net 200 µm	Plankton net	https://www.seanoe.org/data/00752/86417/	Zooplankton abundance and diversity (vertical nets)	Fortier L.
<i>Zooplankton (taxonomy)</i>	Microscopy	Drifting sediment trap	https://www.seanoe.org/data/00752/86417/	drifting traps data (25m depth)	Lalande C.
<i>Zooplankton fecal pellets</i>	Microscopy	Drifting sediment trap	https://www.seanoe.org/data/00752/86417/	drifting traps data (25m depth)	Lalande C.

Table 4. Name and description of variables systematically included in datasets (i.e., in each .csv file)

Variable	Description
<i>mission</i>	mission identifier (“ice_camp_2016”, “Amundsen_2016”, ...)
<i>date</i>	sampling date (UTC)
<i>latitude</i>	latitude of sampling location (degree north decimals)
<i>longitude</i>	longitude of sampling (degree east decimals)
<i>station</i>	station name (ex: G206)
<i>operation code</i>	GE_0008-1 (unique operation identifier)
<i>PI</i>	name(s) of the principal investigator(s) responsible of the measured (calculated) variable

Table 5. List of all peer-reviewed journal articles published so far using Green Edge cruise and/or Green Edge Ice Camp data. Only the first 4 authors are indicated in this table.

Authors	Year	Title	Journal	DOI	Cruise	Ice Camp
<i>Amiriaux, R., Jeanthon, C., Vaultier, F., Rontani, J.-F.</i>	2016	Paradoxical effects of temperature and solar irradiance on the photodegradation state of killed phytoplankton	Journal of phycology	https://doi.org/10.1111/jpy.12410	n	y
<i>Rontani, J.-F., Belt, S.-T., Brown, T.-A., Amiriaux, R., et al.</i>	2016	Monitoring abiotic degradation in sinking versus suspended Arctic sea ice algae during a spring ice melt using specific lipid oxidation tracers	Organic Geochemistry	https://doi.org/10.1016/j.orggeochem.2016.05.016	n	y
<i>Amiriaux, R., Belt, S.-T., Vaultier, F., Galindo, V., et al.</i>	2017	Monitoring photo-oxidative and salinity-induced bacterial stress in the Canadian Arctic using specific lipid tracers	Marine chemistry	https://doi.org/10.1016/j.marchem.2017.05.006	n	y
<i>Rontani, J.-F., Galeron, M.-A., Amiriaux, R., Artigue, L., et al.</i>	2017	Identification of di- and triterpenoid lipid tracers confirms the significant role of autoxidation in the degradation of terrestrial vascular plant material in the Canadian Arctic	Organic Geochemistry	https://doi.org/10.1016/j.orggeochem.2017.03.011	n	y
<i>Dadaglio, L., Dinasquet, J., Obernosterer, I., Joux, F.</i>	2018	Differential responses of bacteria to diatom-derived dissolved organic matter in the Arctic Ocean	Aquatic Microbial Ecology	https://doi.org/10.3354/ame01883	y	n
<i>Goyens, C., Marty, S., Leymarie, E., Antoine, D., et al.</i>	2018	High Angular Resolution Measurements of the Anisotropy of Reflectance of Sea Ice and Snow	Earth and Space Science	https://doi.org/10.1002/2017EA000332	n	y
<i>Massicotte, P., Bécu, G., Lambert-Girard, S., Leymarie, E., et al.</i>	2018	Estimating underwater light regime under spatially heterogeneous sea ice in the Arctic	Applied Sciences	https://doi.org/10.3390/app8122693	n	y
<i>Rontani, J.-F., Amiriaux, R., Lalande, C., Babin, M et al.</i>	2018	Use of palmitoleic acid and its oxidation products for monitoring the degradation of ice algae in Arctic waters and bottom sediments	Organic geochemistry	https://doi.org/10.1016/j.orggeochem.2018.06.002	n	y
<i>Rontani, J.-F., Belt, S.-T., Amiriaux, R.</i>	2018	Biotic and abiotic degradation of the sea ice diatom biomarker IP 25 and selected algal sterols in near-surface Arctic sediments	Organic Geochemistry	https://doi.org/10.1016/j.orggeochem.2018.01.003	n	y
<i>Lafond, A., Leblanc, K., Quéguiner, B., Moriceau, B., et al.</i>	2019	Late spring bloom development of pelagic diatoms in Baffin Bay	Elementa	http://doi.org/10.1525/elementa.382	y	n
<i>LeBlanc, M., Gauthier, S., Garbus, S. E., Mosbech, A., et al.</i>	2019	The co-distribution of Arctic cod and its seabird predators across the marginal ice zone in Baffin Bay	Elementa	http://doi.org/10.1525/elementa.339	y	n
<i>Randelhoff, A., Oziel, L., Massicotte, P., Bécu, G., et al.</i>	2019	The evolution of light and vertical mixing across a phytoplankton ice-edge bloom.	Elementa	https://doi.org/10.1525/elementa.357	y	n
<i>Amiriaux, R., Smik, L., Köseoğlu, D., Rontani, J.-F., et al.</i>	2019	Temporal evolution of IP25 and other highly branched isoprenoid lipids in sea ice and the underlying water column during an Arctic melting season.	Elementa	https://doi.org/10.1525/elementa.377	n	y
<i>Else, B. G. T. T., Whitehead, J. J., Galindo, V., Ferland, J., Mundy, C. J., Gonski, S. F., et al.</i>	2019	Response of the Arctic marine inorganic carbon system to ice algae and under-ice phytoplankton blooms: A case study along the fast-ice edge of Baffin Bay.	Journal of Geophysical Research - Oceans	https://doi.org/10.1029/2018JC013899	n	y
<i>Gourdal, M., Crabeck, O., Lizotte, M., Galindo, V., et al.</i>	2019	Upward transport of bottom-ice dimethyl sulfide during advanced melting of arctic first-year sea ice.	Elementa	https://doi.org/10.1525/elementa.370	n	y
<i>Matthes, L. C., Ehn, J. K., L.-Girard, S., Pogorzelec, N. M., et al.</i>	2019	Average cosine coefficient and spectral distribution of the light field under sea ice: Implications for primary production.	Elementa	https://doi.org/10.1525/elementa.363	n	y
<i>Oziel, L., Massicotte, P., Randelhoff, A., Ferland, J et al.</i>	2019	Environmental factors influencing the seasonal dynamics of spring algal blooms in and beneath sea ice in western Baffin Bay.	Elementa	https://doi.org/10.1525/elementa.372	n	y

Authors	Year	Title	Journal	DOI	Cruise	Ice Camp
Sampei, M.	2019	An estimation of the quantitative impacts of copepod grazing on an under sea-ice spring phytoplankton bloom in western Baffin Bay, Canadian Arctic	Elementa	https://doi.org/10.1525/elementa.2019.00092	n	y
Sansoulet, J., Pangrazi, J.-J., Sardet, N., Mirshak, S., et al.	2019	Green Edge Outreach Project: a large-scale public outreach and educational initiative	polar record	https://doi.org/10.1017/S0032247419000123	n	y
Burgers, T. M., Tremblay, J.-É., Else, B. G. T., & Papakyriakou, T. N.	2020	Estimates of net community production from multiple approaches surrounding the spring ice-edge bloom in Baffin Bay	Elementa	https://doi.org/10.1525/elementa.013	y	n
Randelhoff, A., Lacour, L., Marec, C., Leymarie, E., et al.	2020	Arctic mid-winter phytoplankton growth revealed by autonomous profilers	Science Advances	https://doi.org/10.1126/sciadv.ab02678	y	n
Saint-Béat, B., Fath, B. D., Aubry, C., Colombet, J., et al.	2020	Contrasting pelagic ecosystem functioning in eastern and western Baffin Bay revealed by trophic network modeling	Elementa	https://doi.org/10.1525/elementa.397	y	n
Yunda-Guarin, G., Brown, T. A., Michel, L. N., Saint-Béat, B., et al.	2020	Reliance of deep-sea benthic macrofauna on ice-derived organic matter highlighted by multiple trophic markers during spring in Baffin Bay, Canadian Arctic	Elementa	https://doi.org/10.1525/elementa.2020.047	y	n
Amiriaux, R., Burot, C., Bonin, P., Massé, G., et al.	2020	Stress factors resulting from the Arctic vernal sea ice melt: impact on the viability of the bacterial communities associated to sympagic algae	Elementa	https://doi.org/10.1525/elementa.076	n	y
Else, B. G. T. T., Whitehead, J. J., Galindo, V., Ferland, J., et al.	2020	Green Edge ice camp campaigns: understanding the processes controlling the under-ice Arctic phytoplankton spring bloom.	Earth System Science Data	https://doi.org/10.5194/essd-12-151-2020	n	y
Matthes, L. C., Mundy, C. J., L-Girard, S., Babin, M et al.	2020	Spatial Heterogeneity as a Key Variable Influencing Spring-Summer Progression in UVR and PAR Transmission Through Arctic Sea Ice	Frontiers in Marine Science	https://doi.org/10.3389/fmars.2020.00183	n	y
Sansoulet, J., Therrien, M., Delgove, J., Pouxviel, G., et al.	2020	An update on Inuit perceptions of their changing environment, Qikiqtaaluk (Baffin Island, Nunavut)	Elementa	https://doi.org/10.1525/elementa.025	n	y
Ardyna, M., & Arrigo, K. R.	2020	Phytoplankton dynamics in a changing Arctic Ocean.	Nature Clim. Change	https://doi.org/10.1038/s41558-020-0905-y	y	y
Ardyna, M., Mundy, C. J., Mills, M. M., Oziel, L., et al.	2020	Environmental drivers of under-ice phytoplankton bloom dynamics in the Arctic Ocean	Elementa	http://doi.org/10.1525/elementa.430	y	y
Gérikas Ribeiro, C., dos Santos, A. L., Probert, I., Vaultot, D., et al.	2020	Taxonomic reassignment of <i>Pseudohaptolina birgeri</i> comb. nov. (Haptophyta)	Journal of Phycology	https://doi.org/10.1080/00318884.2020.1830255	y	y
Ribeiro, C. G., Dos Santos, A. L., Gourvil, P., Le Gall, F., et al.	2020	Culturable diversity of Arctic phytoplankton during pack ice melting.	Elementa	http://doi.org/10.1525/elementa.401	y	y
Tisserand, L., Dadaglio, L., Intertaglia, L., Catala, P., et al.	2020	Use of organic exudates from two polar diatoms by bacterial isolates from the Arctic Ocean	Phil. Trans. R. Soc. A.	https://doi.org/10.1098/rsta.2019.0356	y	y
Yau, S., Lopes dos Santos, A., Eikrem, W., Gérikas Ribeiro, C., et al.	2020	<i>Mantoniella beaufortii</i> and <i>Mantoniella baffinensis</i> sp. nov. (Mamiellales, Mamiellophyceae), two new green algal species from the high arctic	Journal of Phycology	https://doi.org/10.1111/jpy.12932	y	y
Toullec, J., Moriceau, B., Vincent, D., Guidi, L., et al.	2021	Processes controlling aggregate formation and distribution during the Arctic phytoplankton spring bloom in Baffin Bay	Elementa	https://doi.org/10.1525/elementa.2021.00001	y	n
Vilgrain, L., Maps, F., Picheral, M., Babin, M., et al.	2021	Trait-based approach on zooplankton in situ images reveals contrasted ecological patterns along ice melt dynamics	Limnology & Oceanography	https://doi.org/10.1002/lno.11672	y	n
Amiriaux, R., Rontani, J.-F., Armougom, F., Frouin, E., et al.	2021	Bacterial diversity and lipid biomarkers in sea ice and sinking particulate organic material during the melt season in the Canadian Arctic	Elementa	https://doi.org/10.1525/elementa.2019.040	n	y

Authors	Year	Title	Journal	DOI	Cruise	Ice Camp
<i>Gali, M., Lizotte, M., Kieber, D.J., Randelhoff, A., et al.</i>	2021	DMS emissions from the Arctic Ocean marginal ice zone	Elementa	https://doi.org/10.1525/elementa.2020.00113	y	n
<i>Laliberté, J., Rehm, E., Hamre, B., Goyens, C et al.</i>	2022	A method to derive satellite PAR albedo time series over first-year sea ice in the Arctic Ocean	Elementa	https://doi.org/10.1525/elementa.2020.00080	n	y