



Carbon Export in the Seasonal Sea Ice Zone North of Svalbard From Winter to Late Summer

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Phytoplankton blooms in the Arctic Ocean's seasonal sea ice zone are expected to start earlier and occur further north with retreating and thinning sea ice cover. The current study is the first compilation of phytoplankton bloom development and fate in the seasonally variable sea ice zone north of Svalbard from winter to late summer, using short-term sediment trap deployments. Clear seasonal patterns were discovered, with low winter and pre-bloom phytoplankton standing stocks and export fluxes, a short and intense productive season in May and June, and low Chl *a* standing stocks but moderate carbon export fluxes in the autumn post-bloom conditions. We observed intense phytoplankton blooms with Chl *a* standing stocks of >350 mg m⁻² below consolidated sea ice cover, dominated by the prymnesiophyte *Phaeocystis pouchetii*. The largest vertical organic carbon export fluxes to 100 m, of up to 513 mg C m⁻² day⁻¹, were recorded at stations dominated by diatoms, while those dominated by *P. pouchetii* recorded carbon export fluxes up to 310 mg C m⁻² day⁻¹. Fecal pellets from krill and copepods contributed a substantial fraction to carbon export in certain areas, especially where blooms of *P. pouchetii* dominated and Atlantic water advection was prominent. The interplay between the taxonomic composition of protist assemblages, large grazers, distance to open water, and Atlantic water advection was found to be crucial in determining the fate of the blooms and the magnitude of organic carbon exported out of the surface water column. Previously, the marginal ice zone was considered the most productive region in the area, but our study reveals intense blooms and high export events in ice-covered waters. This is the first comprehensive study on carbon export fluxes for under-ice phytoplankton blooms, a phenomenon suggested to have increased in importance under the new Arctic sea ice regime.

Keywords: vertical carbon export, sea ice, phytoplankton, seasonality, Arctic ocean, under-ice bloom

INTRODUCTION

Vertical carbon export plays a key role in the biological carbon pump by contributing to the sequestration of CO₂ from the surface to the deep ocean, as well as providing a food source for mesopelagic, deep-sea and benthic ecosystems (Ducklow et al., 2001; Boyd and Trull, 2007). In the Arctic Ocean, primary productivity and pelagic–benthic coupling are primarily driven by the distribution, thickness, and melting of sea ice, controlling the light transmittance and stratification in the waters below (Wassmann and Reigstad, 2011; Lalande et al., 2014). The proportion of multiyear ice (MYI) has declined by 90% since 1979 (Stroeve and Notz, 2018; IPCC report, 2019), and the Arctic Ocean is expected to be ice free in summer by the end of the century or earlier (Massonnet et al., 2012; Wang and Overland, 2012). Thus, the whole Arctic Ocean may eventually transition into a seasonally ice-covered ocean. As a result of the shorter, more transparent, and dynamic sea ice cover, large under-ice phytoplankton blooms in the Arctic have been reported (Fortier et al., 2002; Arrigo et al., 2012; Mundy et al., 2014; Assmy et al., 2017; Johnsen et al., 2018; Ardyna and Arrigo, 2020; Ardyna et al., 2020). Recent studies have described the phytoplankton dynamics and environmental drivers of these under-ice blooms (Ardyna et al., 2020), but the fate of the carbon produced by these blooms is yet to be explored (Ardyna and Arrigo, 2020).

As a consequence of the changing Arctic sea ice regime, blooms of phytoplankton are now beginning earlier in the seasonal sea ice zone (SIZ) (Kahru et al., 2011). The SIZ is defined as the area that encompasses the annual minimum and maximum sea ice extents as well as the ice margin significantly influenced by ocean (Wadhams, 1986). It has been predicted that the northern portions of the SIZ will experience the largest changes and that the earlier phytoplankton blooms would result in earlier and an overall increase in carbon export (Wassmann and Reigstad, 2011). Additionally, Wassmann and Reigstad (2011) predicted an extended ice-free period that would lead to a longer productive season, especially in the form of regenerated production dominated by heterotrophic processes once nutrients are limiting. However, little quantitative information exists from the northernmost regions of today's SIZ. Contrary to the Barents Sea, the areas further north have previously been logistically challenging to access, and thus, the current SIZ of the Eurasian Arctic has been understudied, especially across seasons.

Daily measurements of vertical export can reveal central processes occurring in the upper water column, including the timing and synchrony between phytoplankton blooms and zooplankton grazers, giving us insight into the partitioning of organic carbon between pelagic and benthic ecosystems. Since a large portion of pelagic–benthic coupling is determined by the gravitational flux of particles [but see Boyd et al. (2019) for other mechanisms], the protist community composition plays an important role. Protists are unicellular or unicellular-colonial eukaryotes, including heterotrophic, phototrophic, and mixotrophic organisms. Diatoms, large (generally >10 μm) unicellular phytoplankton encased in silicified shells, are considered to be important contributors to vertical carbon export under certain conditions. Once diatom blooms terminate their

growth phase and nutrients become limited, mass sedimentation events of ungrazed cells are often observed (Kjørboe et al., 1996; Olli et al., 2002; Wassmann et al., 2006b; Bauerfeind et al., 2009; Juul-Pedersen et al., 2010; Lalande et al., 2014; Roca-Martí et al., 2017; 2019). Some diatom species form large chains that can aggregate or entrap other cells in sticky mucilage and sometimes spines, producing rapidly sinking flocs of cells (Assmy et al., 2019). On the other hand, several studies revealed limited export of *Phaeocystis* spp.-dominated blooms (Riebesell et al., 1995; Reigstad and Wassmann, 2007; Wolf et al., 2016), while others have reported significant contribution of this prymnesiophyte to export (Lalande et al., 2011; Moigne et al., 2015; Wollenburg et al., 2018), mainly as a consequence of deep mixing, aggregation, ballasting, or heavy grazing by large zooplankton who mediate export through the production of fast-sinking fecal pellets (Hamm et al., 2001; Reigstad and Wassmann, 2007; Boyd et al., 2019). This is consistent with a recent study, partly based on vertical export data presented herein, showing that the carbon flux attenuation coefficient is negatively correlated with the dominance of diatoms compared to non-ballasted *Phaeocystis*, and therefore, the relative contribution of diatoms is a good predictor for carbon flux attenuation in the upper 200 m of the water column, especially in the cold waters of the Arctic (Wiedmann et al., 2020).

As the Arctic is warming and stratification is strengthening as a result of freshening, generally reducing nutrient supply, the flagellated species, such as *Phaeocystis* spp., have been predicted to become more dominant, because their high surface-area-to-volume ratio makes them more efficient at acquiring nutrients, and their size and colonies embedded in polysaccharide gel matrixes make them hydrodynamically resistant to sinking (Li et al., 2009; Peter and Sommer, 2012; Metfies et al., 2016). A recent study has found bio-optical evidence for increasing dominance of *Phaeocystis pouchetii* in the Barents Sea (Orkney et al., 2020). This shift toward smaller cells has further been modeled to increase the length of the food web and, consequently, retain phytoplankton carbon and to be respired in the upper water column (Vernet et al., 2017). In addition, the decline in silicic acid concentrations, an essential nutrient for diatom growth, in the Nordic seas and Barents Sea (Hátún et al., 2017) seems to have led to the increased dominance of *Phaeocystis* in the Barents Sea (Orkney et al., 2020). Shifts in bloom-forming protists' composition, as well as their seasonal timing in relation to heterotrophic consumers, will greatly influence vertical export.

In order to measure the pelagic processes and associated export fluxes, synchronized and daily sampling is necessary. In our study, we investigate these processes seasonally under generally dense sea ice cover, from a collection of 22 short-term sediment trap deployments in the SIZ north of Svalbard. The study focuses on how the magnitude and attenuation of vertical export in the epipelagic zone is determined by seasonal variability in sea ice cover, protist community composition, and the composition of the exported organic material. With a seasonal coverage from late January to August in a region previously hard to access, this study provides new insight to the export efficiencies in the changing Arctic Ocean, including under-ice blooms.

MATERIALS AND METHODS

Study Area

Vertical carbon export was studied during three field campaigns in the SIZ north of Svalbard: CarbonBridge (Bridging productivity regimes in the Arctic Ocean, with R/V *Helmer Hanssen*), N-ICE2015 (Norwegian young sea ICE Cruise, with R/V *Lance*), and TRANSSIZ (Transitions in the Arctic Seasonal Sea Ice Zone, PS92, ARK XXIX/1, with R/V *Polarstern*). In total, areas represented by 22 stations were sampled in May and August 2014 and from late January to late June 2015. The ice cover in the visited areas ranged from 0 to 100% in sea ice cover and was influenced to various degrees by Atlantic water transported by the West Spitsbergen Current (WSC) and by overlaying Arctic surface waters (ArSW) advected in the opposite direction (Rudels et al., 2000; Fer et al., 2017; Meyer et al., 2017; Crews et al., 2018; Menze et al., 2019) (**Figure 1**).

Sampling and Vertical Export Measurements

Short-term sediment trap arrays [KC Demark, parallel cylinders mounted on a gimbaled frame, with an aspect ratio (height:diameter) of 6.25 to avoid turbulent mixing in the cylinders during sampling] were deployed at the edge of or through a hole in a floe of drifting sea ice, at depths ranging from 5 to 200 m for 21 to 71.5 h, with longer deployments during periods of low flux (**Table 1**). The trap cylinders are weighted and mounted on a gimbaled frame equipped with a vane and will remain vertical and perpendicular to the current directions. No fixatives or poisons were added to the traps during deployment, but the trap cylinders were filled with a saturated NaCl solution to reduce microbial activity during the deployments of N-ICE2015, which were longer.

Upon retrieval, the contents of the two trap cylinders from each depth were pooled, homogenized carefully, and subsampled for measurements of particulate organic carbon (POC) and nitrogen (PON) and chlorophyll *a* (Chl *a*) and for counts of protists and fecal pellets (FP) from large zooplankton. The 100-ml protist samples for CarbonBridge and TRANSSIZ were fixed with glutaraldehyde–Lugol mixture (2% final concentration), and the 200-ml fecal pellet samples with formaldehyde (2% final concentration). For N-ICE2015, samples for protist cell counts (100 ml) were fixed with an aldehyde mixture (0.1% glutaraldehyde and 1% hexamethylenetetramine-buffered formaldehyde final concentrations). All samples were stored refrigerated (4°C) until microscopy analyses were performed.

Water samples for concentrations of suspended POC, PON, and Chl *a*, as well as for protists (during CarbonBridge and N-ICE2015), algal marker pigments determined by high-performance liquid chromatography (HPLC) (during TRANSSIZ), and inorganic nutrients [nitrates (nitrate and nitrite combined), phosphate, and silicic acid] analyses, were collected from discrete depths between 0 and 200 m, with Niskin bottles mounted on a Sea-Bird rosette water sampler equipped with a conductivity, temperature, and depth (CTD) probe (SBE911+). During CarbonBridge and TRANSSIZ, the inorganic nutrient samples were collected in acid-washed 100 ml

plastic bottles and immediately frozen at -20°C , while during N-ICE2015, the samples were immediately fixed with 0.2 ml chloroform and refrigerated, until analysis could be performed in a land-based lab.

The POC, PON, and Chl *a* samples from both the suspended and exported profiles were filtered (300–500 ml) onto GF/F filters (Whatman), precombusted at 500°C for the POC/PON samples, and immediately frozen at -20°C , or immediately dried (at 60°C) and stored on PALL filter slides (N-ICE2015 samples), and analyzed later. For HPLC pigment analyses, water samples (1–2 L) were filtered on GFF filters and immediately frozen in liquid nitrogen. Samples were thereafter stored at -80°C until analyses. The samples were then processed according to the procedures described in Kiliyas et al. (2013).

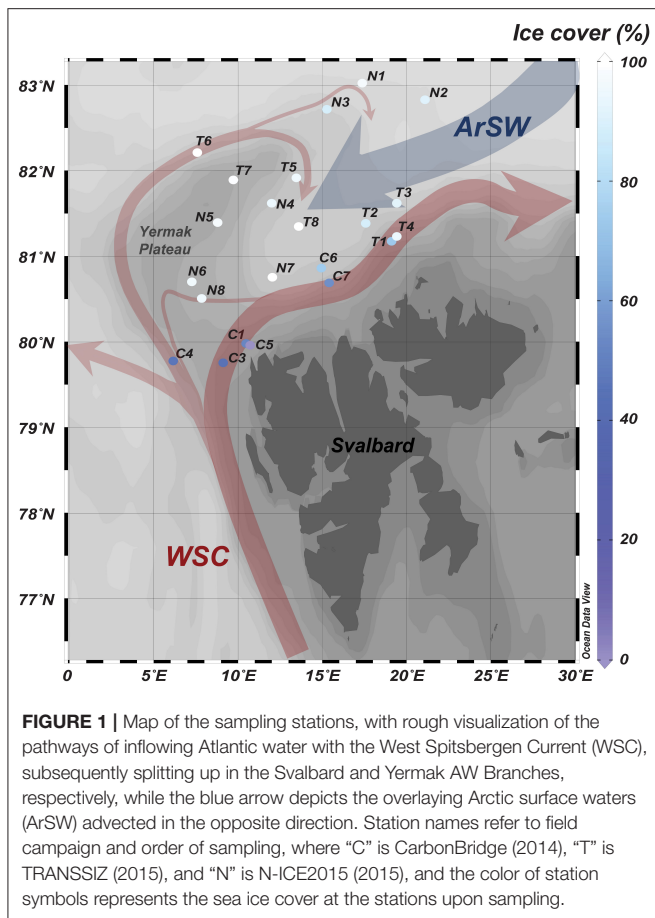
Chl *a*, POC/PON, and Nutrient Analyses

The frozen POC/PON filters were dried at 60°C for 24 h and subsequently placed in an acid fume bath (concentrated HCl) for 24 h to remove all inorganic carbon. The filters were then replaced into the 60°C drying oven for an additional 24 h and finally packed into nickel capsules. The samples were analyzed using an Exeter Analytical CE440 CHN elemental analyzer. Samples for POC/PON from N-ICE2015 were analyzed with continuous-flow mass spectrometry (CF-IMRS) carried out with a RoboPrep/Tracermass mass spectrometer (Europa Scientific, UK). Chl *a* was extracted from filters for roughly 12 h in the dark, in 100% methanol at $4-5^{\circ}\text{C}$, and measured fluorometrically using a pre-calibrated (Sigma, C6144) Turner Design AU-10 fluorometer. The samples were also measured fluorometrically for the degradation product phaeopigment by the addition of HCl (5% concentration) following Holm-Hansen and Riemann (1978).

The CarbonBridge and TRANSSIZ inorganic nutrient samples were analyzed for nitrate, nitrite, phosphate, and silicic acid following standard seawater methods, using a Flow Solution IV analyzer from O.I. Analytical, USA. The analyzer was calibrated using reference seawater from Oceanic Scientific International Ltd., UK. The N-ICE2015 samples were measured spectrometrically in a modified Skalar Autoanalyzer. The measurement uncertainty for nitrite is 0.06 mmol m^{-3} and 10% or less for nitrate, phosphate, and silicic acid.

Protist and Fecal Pellet Analyses

For N-ICE2015 and CarbonBridge, samples for protist cell counts were settled in Utermöhl sedimentation chambers (HYDRO-BIOS[®], Kiel, Germany) for 48 h. Protists, here including phototrophic, heterotrophic, and mixotrophic unicellular eukaryotes, were identified and enumerated at 100–600 \times magnification using an inverted Nikon Ti-S light and epifluorescence microscope. The organisms were identified to the lowest possible taxonomic level, ideally to the species level, otherwise to the genus level or grouped into size classes. Cell carbon was estimated according to Menden-Deuer and Lessard (2000). For TRANSSIZ, the taxonomic structure of the protist community was deduced from the marker pigments measured by HPLC and applying the CHEMTAX program (Mackey et al., 1996), which takes into account that several marker pigments



are not unique to one phytoplankton group. Pigment ratios used in the CHEMTAX program were constrained as suggested by Higgins et al. (2011) based on microscopic examination of representative samples during the cruise and applying the input matrix published by Fragoso et al. (2017). The resulting protist group composition was adjusted according to Chl *a* concentrations. For comparability between field campaigns, the relative biomass of protist groups in the upper water column (0–30 m) was used. Comparable protist data between field campaigns were not available from the sediment traps.

Depending on the concentration of pellets, 25–100 ml of the zooplankton fecal pellet subsamples were left for 24 h to settle in an Utermöhl sedimentation column and, subsequently, enumerated using a Leica inverted microscope. The length and width of each fecal pellet was measured, and its condition noted (intact, end-pieces or mid-pieces), and pellet volumes were calculated according to the shape of the pellets. Long cylindrical pellets were classified as calanoid copepod pellets, larger fecal strings with cut ends (filiform) were classified as euphausiid/krill pellets, and ellipsoid pellets were attributed to appendicularians and amphipods depending on their size (González, 2000; Wexels Riser et al., 2006). Fecal pellet volumes were converted into fecal pellet carbon (FPC) using a volumetric carbon conversion factor of 0.0943 mg C mm⁻³ for copepod pellets, 0.1861 mg

C mm⁻³ for carnivorous amphipod pellets, 0.0451 mg C mm⁻³ for krill pellets, and 0.025 mg C mm⁻³ for appendicularians pellets (Wexels Riser et al., 2006).

Calculations: Depth-Integrated Standing Stocks, Loss Rate, and Phytoplankton Carbon

Standing stock was calculated for the upper 100 m of the water column by integrating the suspended Chl *a* or POC biomass samples. The 100-m depth horizon was chosen as previous studies have found that vertical flux at around 100 m explains >85% of the variation in biogenic matter on the surface sediment of the sea floor (Renaud et al., 2008). The fraction of the standing stock exported daily to 100 m is termed the loss rate and measured in % per day (Reigstad et al., 2008). For the stations where sediment traps were positioned at 90 m, the export flux was extrapolated to 100 m based on the flux curve (Boyd and Trull, 2007).

Estimates of the contribution of phytoplankton-derived POC (PPC) were derived from an estimation of the C:Chl *a* ratio, using the slope of the linear regression of POC against Chl *a* taken from the measurements of all the suspended samples at all stations, here 40.4 (w:w). The slope of the relationship will be the average C:Chl *a* of that organic matter which is statistically associated to the Chl *a* pigment (Banse, 1977), and the ratio of 40.4 found by the current study is in the range provided by previous literature, especially for biomasses including diatoms and prymnesiophytes (Sathyendranath et al., 2009).

The detritus fraction of the exported organic carbon was estimated by subtracting the PPC and FPC fractions from the total POC in the sediment trap samples.

Seasonal and Bloom Stage Categorization

The stations were separated according to the stage of the bloom they were in, based on the following parameters: sea ice cover, hydrographical characteristics and mixed layer depth, nutrient profiles, distinction and depth of the Chl *a* maximum, Chl *a* standing stocks, and protist community composition (Table 2). These parameters were chosen according to Hodal and Kristiansen (2008), Kubiszyn et al. (2017), Leu et al. (2015), Reigstad et al. (2002, 2008), Sundfjord et al. (2008), Tremblay et al. (2006), and Wassmann et al. (1999). Assessment of categorization was further supported by other studies from the field campaigns (Peecken, 2016; Assmy et al., 2017; Olsen et al., 2017; Massicotte et al., 2019; Sanz-Martín et al., 2019; Svensen et al., 2019). Depending on the focus of the study, the parameters may vary. Winter stations, ranging from late January to late April, were further separated from the pre-bloom stage according to sea ice cover and thicknesses (data found in Kowalczyk et al., 2017), nutrient profiles, and especially the standing stocks of Chl *a* and protist biomass. For further details, see Table 3.

Sea Ice Cover, Distance to Open Water, and Water Mass Characteristics

Sea ice cover for all stations were derived from satellite data provided by the Center for Satellite Exploitation and Research

TABLE 1 | Sampling station, cruise ID, sediment trap deployment date, location, water depth, sea ice cover, distance to open water, deployment period, and deployment depths.

Field campaign	Station name	Cruise ID	Deployment date	Lat (°N) Long (°E)	Water depth (m)	Sea ice cover (%)	Distance to open water (km)	Deployment period (h)	Deployment depths (m)
CarbonBridge	C1	P1	19.05.14	79.974 10.654	397	52	27	22.1	20, 30, 40, 50, 60, 90, 120, 200
	C3	P3	23.05.14	79.756 9.138	417	44	35	22.8	20, 30, 40, 50, 60, 90, 120, 200
	C4	P4	25.05.14	79.778 6.163	1,051	46	57	27.1	20, 30, 40, 50, 60, 90, 120, 200
	C5	P5	09.08.14	79.968 10.737	341	0	-23*	21.0	20, 30, 40, 50, 60, 90, 120, 200
	C6	P6	12.08.14	80.861 14.946	993	74	74	22.3	20, 30, 40, 50, 60, 90, 120, 200
	C7	P7	14.08.14	80.691 15.418	622	55	59	26.7	20, 30, 40, 50, 60, 90, 120, 200
	N-ICE2015	N1	Floe1	28.01.15	83.023 17.384	3,955	97	167	48.5
N2		Floe2	14.03.15	82.831 21.101	-	90	289	36.0	5, 25, 50, 100
N3		Floe3	24.04.15	82.720 15.247	1,932	89	265	48.0	5, 25, 50, 100
N4		Floe3	08.05.15	81.621 12.001	1,932	94	136	44.8	5, 25, 50, 100
N5		Floe3	15.05.15	81.392 8.808	1,127	97	118	71.5	5, 25, 50, 100
N6		Floe3	28.05.15	80.704 7.275	866	96	49	50.0	5, 25, 50, 100
N7		Floe4	10.06.15	80.756 12.051	1,461	100	132	49.0	5, 25, 50, 100
N8		Floe4	14.06.15	80.508 7.847	823	95	61	48.0	5, 25, 50, 100
TRANSSIZ	T1	PS92/0019-6	28.05.15	81.174 19.135	377	73	52	24.0	30, 40, 60, 90
	T2	PS92/0027-2	31.05.15	81.386 17.587	876	89	110	24.0	30, 40, 60, 90, 200
	T3	PS92/0031-2	03.06.15	81.620 19.427	1,963	91	251	24.0	30, 40, 60, 90, 200
	T4	PS92/0032-4	06.06.15	81.233 19.431	481	94	98	16.0	30, 40, 60, 90, 200
	T5	PS92/0039-6	11.06.15	81.917 13.459	1,589	95	234	23.0	30, 40, 60, 90, 200
	T6	PS92/0043-4	15.06.15	82.211 7.588	804	100	240	24.0	30, 40, 60, 90, 200
	T7	PS92/0046-1	17.06.15	81.891 9.728	906	99	193	22.0	30, 40, 60, 90, 200
	T8	PS92/0047-3	19.06.15	81.347 13.609	2,171	100	92	24.0	30, 40, 60, 90, 200

*Negative distance value represents an open water station, where the value represents the distance until the ice edge begins.

[CERSAT, Ezraty et al. (2007)]. The product is available on a 12.5 × 12.5-km grid based on 85-GHz SSM/I brightness temperatures, using the ARTIST Sea Ice (ASI) algorithm. The distance to open water was calculated by measuring the distance to the closest area with ice areal cover of <15% as an ice edge indicator.

Mixed layer depths (MLD) were taken as the average depth of the mixed layer during the duration of the sediment trap

deployment, based on existing datasets from the field campaigns. During CarbonBridge and TRANSSIZ, the MLD is defined as the depth where the potential density crosses 20% of the density difference between a surface layer density (3–5 m) and deeper (50–60 m) values (Nikolopoulos et al., 2016; Randelhoff et al., 2018). For N-ICE2015, the MLD was defined as the depth in each profile where the potential density first exceeded the density at

TABLE 2 | Conditions for bloom stage categorization.

	Winter	Pre-bloom	Bloom	Post-bloom
Sea ice cover	89–97%	94–100%	44–100%	0–74%
Mixed layer depth	58–79 m	43–72 m	7–31 m	5–15 m
Surface nitrates	4.6–6.8 mmol m ⁻³	6.0–9.4 mmol m ⁻³	0.0–5.0 mmol m ⁻³	0.0–1.8 mmol m ⁻³
Chl <i>a</i> maximum	None	Indistinct	Distinct; in upper 50 m	Somewhat distinct; shallow (>30 m)
Chl <i>a</i> standing stock	3–4 mg m ⁻²	15–32 mg m ⁻²	69–532 mg m ⁻²	55–92 mg m ⁻²
Protist community	Ciliates, dinoflagellates	Dinoflagellates, ciliates, prasinophytes, diatoms	Diatoms/ <i>Phaeocystis pouchetii</i>	Small flagellates (≤10 μm), coccolithophores

TABLE 3 | Physical and biological properties during the different bloom stages at the 22 sampling stations in 2014 and 2015.

	Winter	Pre-bloom	Bloom	Post-bloom
Date	28 st Jan–24 th April	08 th May–17 th June	19 th May–19 th June	9 th –14 th August
Sea ice cover	89–97%	94–100%	44–100%	0–74%
Distance to open water	167–298 km	119–240 km	26–251 km	–23*–74 km
Mixed layer depth	58–79 m	43–72 m	7–31 m	5–15 m
Chl <i>a</i> standing stock	3–4 mg m ⁻²	15–32 mg m ⁻²	69–532 mg m ⁻²	55–92 mg m ⁻²
POC standing stock	1.9–3.6 g m ⁻²	3.0–5.0 g m ⁻²	8.4–23.4 g m ⁻²	11.9–14.1 g m ⁻²
Nitrates at Chl <i>a</i> maximum	4.6–6.8 mmol m ⁻³	6.0–9.4 mmol m ⁻³	0.0–5.0 mmol m ⁻³	0.0–1.8 mmol m ⁻³
Chl <i>a</i> max depth	N/A	Indistinctive	1–50 m	1–25 m
POC export 100 m	19–57 mg C m ⁻² day ⁻¹	31–154 mg C m ⁻² day ⁻¹	74–513 mg C m ⁻² day ⁻¹	108–215 mg C m ⁻² day ⁻¹
POC loss rate 100 m	1.0–2.0% day ⁻¹	0.8–3.0% day ⁻¹	0.4–3.1% day ⁻¹	0.8–1.6% day ⁻¹
Avg. suspended C:N (full range)	6.0 (2.4–8.1)	6.5 (2.0–10.1)	7.4 (1.3–15.3)	7.1 (5.5–10.3)
Avg. exported C:N (full range)	10.7 (6.9–16.4)	8.6 (5.5–11.9)	7.4 (5.5–16.9)	7.0 (6.2–7.8)
Biomass-dominating groups of protists	Ciliates, dinoflagellates	Dinoflagellates, ciliates, prasinophytes, diatoms	Diatoms/ <i>Phaeocystis pouchetii</i>	Small flagellates (≤10 μm), coccolithophores
PPC/POC export contribution 100 m	0.6–2%	0.8–16%	21–79%	19–32%
FPC/POC export contribution 100 m	Unknown	0–3%**	3–30%**	6–15%
POC:Chl <i>a</i> (<i>n</i>)	143.3 (9)	36.4 (34)	39.8 (101)	82.3 (39)
Sampling stations	N1, N2, N3	N4, N5, T5, T6, T7	C1, C3, C4, N6, N7, N8, T1, T2, T3, T4, T8	C5, C6, C7

*Negative distance value represents an open water station; here, the distance until the ice edge begins.

**FPC data unavailable from N-ICE2015 stations.

20 m depth by 0.01 kg m⁻³ in winter and the near-surface value by 0.003 kg m⁻³ in spring (Meyer et al., 2017).

Statistical Analyses

To assess the strength of trends in the data, linear regressions were performed on log-transformed Chl *a* standing stocks against the stations' distances to open water and against the mixed layer depth of each station. Standing stocks for the stations were log-transformed as they increased in an exponential manner as blooms developed.

RESULTS

Environmental and Bloom Settings

All stations were located in ice-covered waters (44–100%), apart from one open water station (C5) sampled in August (Figure 1, Table 1). During N-ICE2015 and TRANSSIZ, ice thickness ranged from 0.90 to 1.45 m and snow thickness from

0.10 to 0.54 m, with the thickest snow and ice in early May over the northern part of the Yermak Plateau and thinnest in mid-June over the Sofia Basin and southern Yermak Plateau [for full details, see Kowalczyk et al. (2017) and Massicotte et al. (2019)]. Ice and snow thickness were not measured during the CarbonBridge campaign. Unfortunately, light measurements were not available for all campaigns, but see Pavlov et al. (2017) and Katlein et al. (2019) for the N-ICE2015 and TRANSSIZ/PS92 stations, respectively.

Winter and Pre-bloom

Typical winter conditions were found at the deep Nansen Basin stations sampled from late January to late April and characterized by dense ice cover and large distance (>160 km) to open water. The mixed layer depth at these stations ranged between 58 and 79 m within which water temperatures were close to the freezing point (on average –1.7°C) and salinities around 34.3 psu (Figures 2, 3B). In the upper 50 m, Chl *a* and POC

standing stocks were very low (3–4 mg m⁻², the lowest observed during the study), and nitrates (nitrate+nitrite) and silicic acid concentrations were between 5 and 7 mmol m⁻³ and between 2 and 3 mmol m⁻³, respectively (see **Table 3**, **Figures 2, 3A**).

Five stations over the northern Yermak Plateau sampled in May and June were categorized as pre-bloom stations based on the dense sea ice cover, cold and saline surface conditions (**Figure 2**), low but increasing Chl *a* standing stocks, and high nutrient concentrations (6–9 mmol m⁻³ for nitrates and 2.5–4 mmol m⁻³ for silicic acid, **Figure 3A**) in the upper 100 m (**Table 3**). The pre-bloom stations had mixed layer depths between 43 and 72 m (**Figure 3B**).

Bloom

The highest Chl *a* standing stocks and low nutrient concentrations were recorded at 11 stations sampled in May and June (in 2014 and 2015). They were classified as bloom stations. In surface waters, the concentrations of nitrates ranged from 0 to 5 mmol m⁻³ (**Figure 3A**) and silicic acid from 0.2 to 4 mmol m⁻³ (**Figure 2**). For additional profiles of nitrates, silicic acid, and phosphate during the diatom and *Phaeocystis* blooms, see the **Supplementary Materials**. The areas over the Yermak Plateau (sampling stations N6–N8 and T8) and the shelf break (stations T1–T4) showed cold Arctic surface waters (<0°C) above the warmer modified or core Atlantic waters (Rudels et al., 2000), while the areas further south (stations C1–C4) showed warmer and more saline water masses throughout the water column (**Figure 2**). All stations had core Atlantic water (>2°C) below 100 m depth (with the exception of station T1) and mixed layer depths between 7 and 31 m (**Figure 3B**). For further details of water mass characteristics for these areas, see Meyer et al. (2017), Peeken (2016), and Randelhoff et al. (2018).

Post-bloom

The areas sampled in August were in a post-bloom stage (stations C5–C7), and nutrient concentrations in surface waters were very low (0–2 mmol m⁻³ for nitrates and 0.5–1 mmol m⁻³ for silicic acid), displaying warm and saline unstratified water column at station C5 and an advanced state of sea ice melt with a cold and fresh surface layer overlaying the warm and saline waters at depth at stations C6 and C7 (**Figures 2, 3**). These stations had lowered Chl *a* standing stocks (55–92 mg m⁻²) and mixed layer depths between 5 and 15 m (**Figure 3B**).

Seasonal Patterns

Standing Stocks

The standing stocks of Chl *a* and POC displayed large seasonal variation, with the broadest variation in May and June (**Figure 4**). During winter, standing stocks remained below ~3 mg Chl *a* m⁻² and 5 g C m⁻² (**Table 3**). Pre-bloom POC standing stocks were similar to winter conditions, but Chl *a* standing stocks were 10-fold higher (32 mg m⁻²).

During the bloom stage in May and June, Chl *a* and POC standing stocks ranged between 69 and 532 mg m⁻² and between 8 and 23 g m⁻², respectively. In August, the post-bloom stage, Chl *a* standing stocks had decreased to 55–92 mg m⁻², while

those of POC remained at intermediate levels of 12–14 g C m⁻², twice as high as during the pre-bloom stages (**Figure 4**).

Relationships Between Chl *a* Standing Stock and Sea Ice Conditions

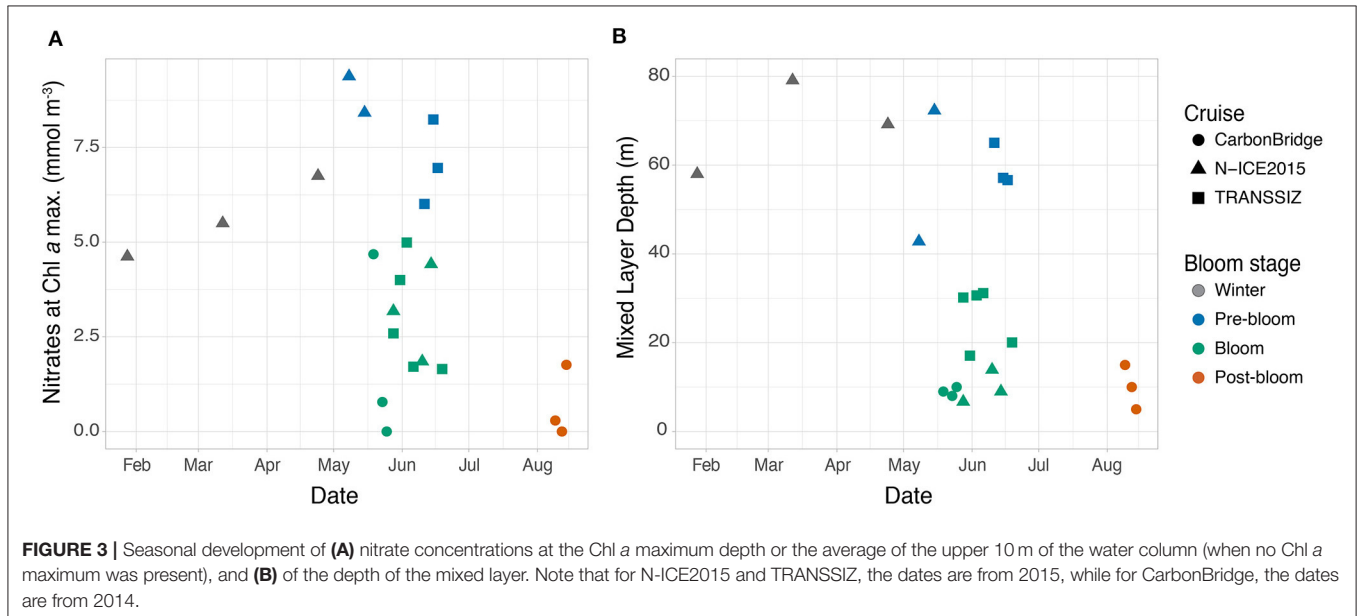
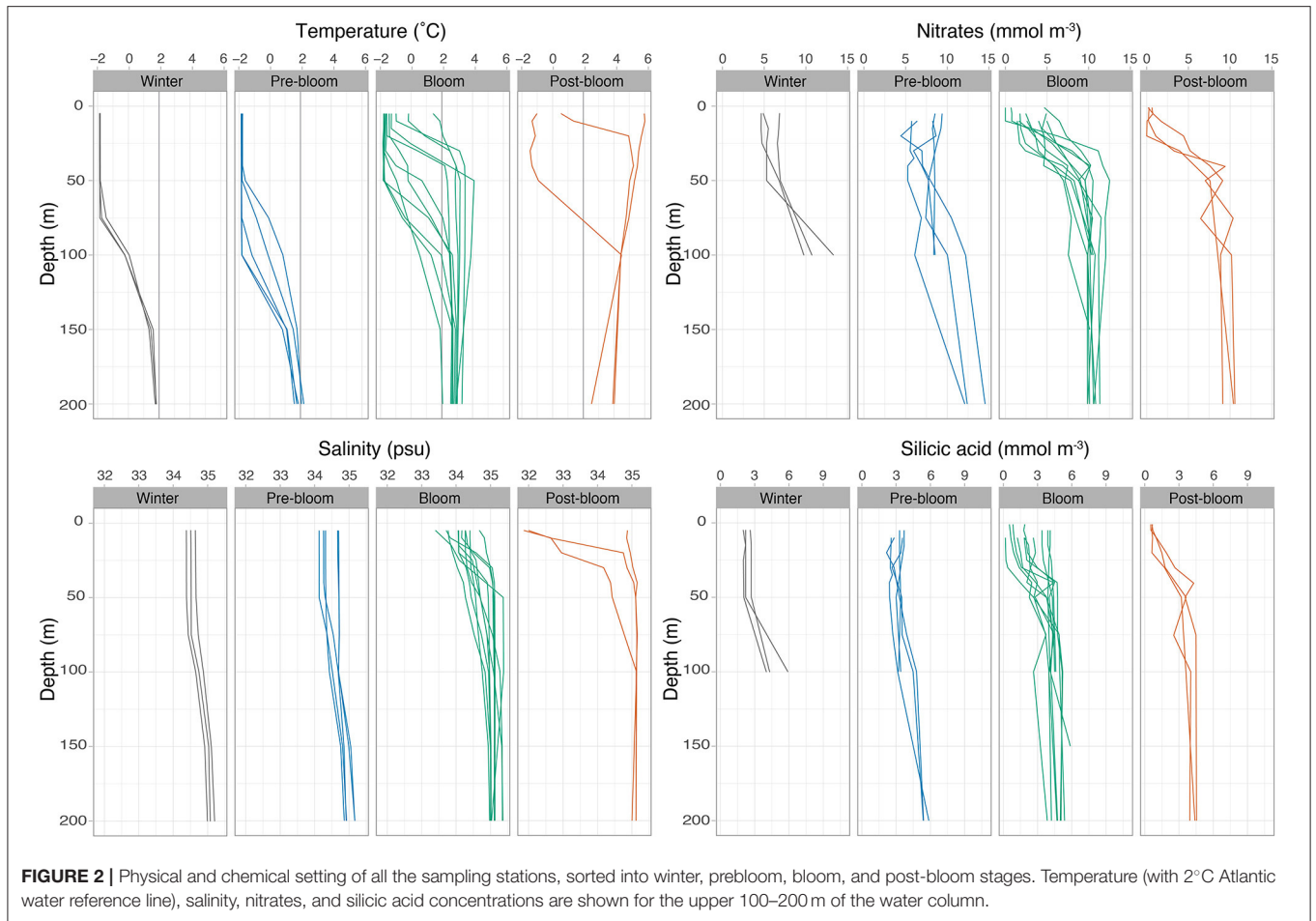
Strong negative correlations were found between the standing stock of Chl *a* and the distance to open water ($r^2 = 0.54$, $P = 0.00006$; **Figure 5A**) and between Chl *a* standing stock and the mixed layer depth ($r^2 = 0.69$, $P = 0.000001$; **Supplementary Material**). The sampling areas located closest to open water (<15% sea ice cover) were generally characterized by the highest standing stocks of Chl *a*, while those farthest into the ice had lower stocks. Notably, the sampling station in open water had lower Chl *a* standing stock than those situated between 27 and 109 km into the ice pack. The highest standing stock (532 mg Chl *a* m⁻²) was recorded at a station located 35 km into the ice. The sampling stations with the shallowest mixed layer depth (<25 m) generally had the highest Chl *a* standing stock, while areas with deeper mixed layers were mainly associated with lower standing stocks. No significant correlation was found between Chl *a* standing stock and ice cover ($r^2 = 0.06$, $P = 0.142$) as the 15 stations where ice cover ≥89% showed almost the full range of integrated standing stocks (**Figure 5B**). It is important to note that these linear models span over many seasons and were not weighted accordingly.

Surface Water Protist Community

During the winter over the Nansen Basin, ciliates and dinoflagellates dominated in terms of biomass in the upper 30 m of the water column (**Figure 6**). Ciliates, which accounted for 32–84% of the total biomass, were mainly composed of *Mesodinium rubrum*, *Uronema marinum*, and unidentified cysts, while dinoflagellates, accounting for 12–69% of the biomass, were mainly of *Gymnodinium* spp. and also *Alexandrium* sp. at station N2.

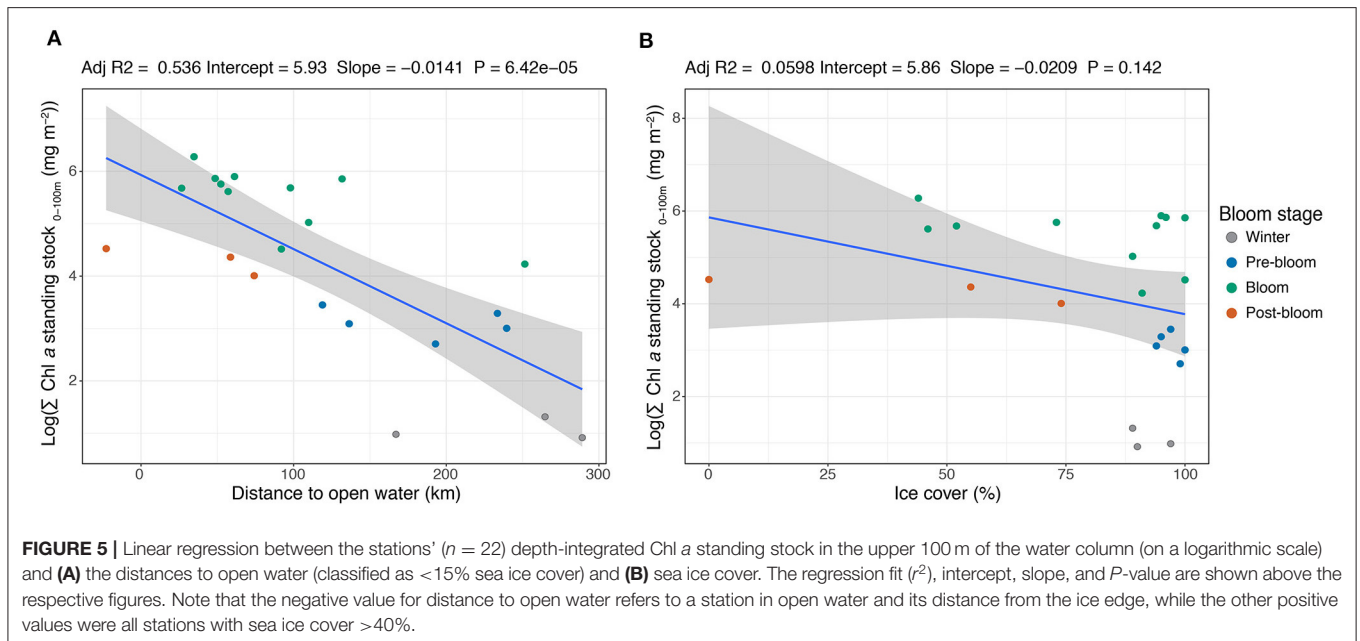
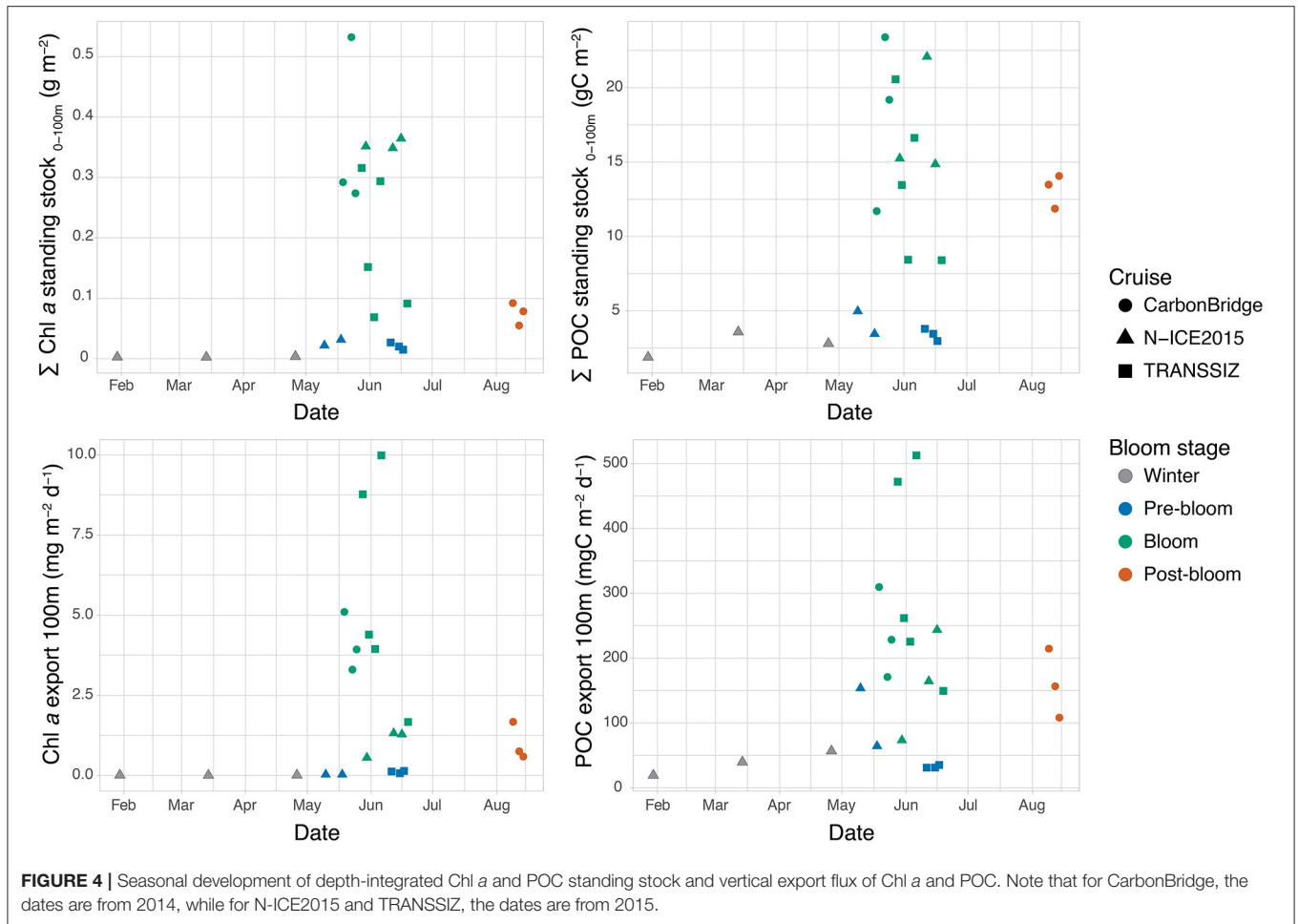
For the stations in a pre-bloom state over northern Yermak Plateau, the composition of the protist community in the upper 30 m differed between the N-ICE2015 and TRANSSIZ campaigns, which can be largely attributed to differences in timing. The N-ICE2015 stations in mid-May (N4–N5) had a 33–93% dominance of ciliates (mainly *Mesodinium rubrum*) and at station N4 of dinoflagellates (38%, *Gymnodinium* spp.) (**Figure 6**). The TRANSSIZ stations in mid-June 2015 (T5–T7) were dominated by prasinophytes and diatoms (24–48% and 3–27%, respectively). The relative biomass contributions from TRANSSIZ were based on HPLC analyses, and thus, further taxonomic resolution was not possible.

During the blooms when Chl *a* standing stocks were high, protist biomass in the upper water column was either dominated by diatoms or the prymnesiophyte *P. pouchetii* (**Figure 6**). The exception was N-ICE2015 station N8, sampled over the southern Yermak Plateau in June, where ciliates accounted for 33% (*Leegaardiella sol* and several unidentified species), dinoflagellates for 29% (*Gymnodinium* spp.), and prymnesiophytes for 27%, which were mainly unidentified coccolithophores, in the upper 30 m. However, for station N8, at 50 m and below, the prymnesiophyte *P. pouchetii* dominated in terms



of biomass (data not shown). The stations sampled over the shelf break north of Svalbard (T1–T4) were dominated by diatoms, 73–91% (Figure 6), mainly *Thalassiosira antarctica*

var. *borealis* and *Thalassiosira hyalina* (pers. comm. Eva-Maria Nöthig). In the southernmost stations (C1–C4), the southern Yermak Plateau station N6, and the deeper Sofia Basin stations



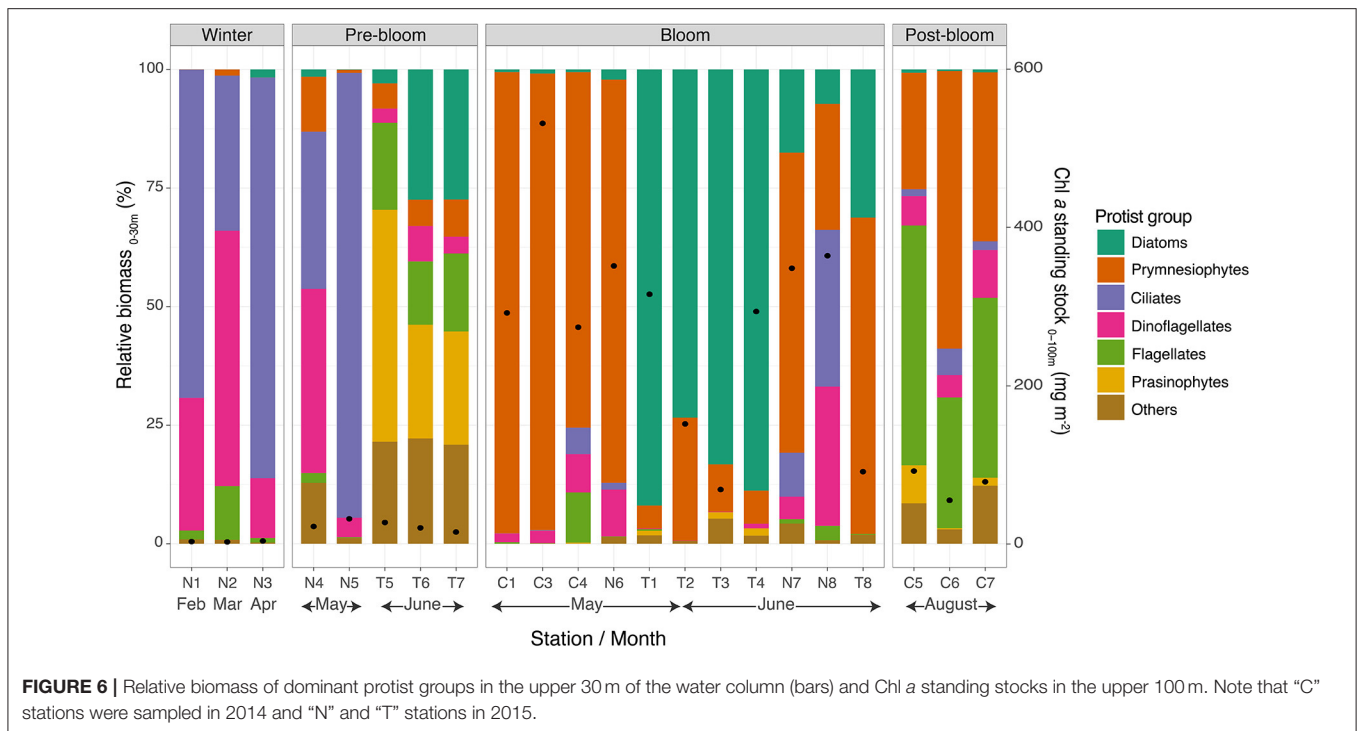


FIGURE 6 | Relative biomass of dominant protist groups in the upper 30 m of the water column (bars) and Chl *a* standing stocks in the upper 100 m. Note that “C” stations were sampled in 2014 and “N” and “T” stations in 2015.

(N7 and T8), the relative protist biomass in the upper 30 m was dominated by *P. pouchetii*, 63–97% (Figure 6), mainly in the form of colonies.

The southernmost stations in August in post-bloom states (C5–C7) were found to be dominated by small unidentified flagellates (28–51%, mainly <10 μm in size) and by unidentified coccolithophores (24–59%) (Figure 6).

Vertical Export Fluxes

Similar seasonal patterns in Chl *a* and POC export fluxes were observed at 100 m depth (Figure 4). Export fluxes of Chl *a* were low during winter and the pre-bloom stage, between 0.005 and 0.14 $\text{mg m}^{-2} \text{day}^{-1}$, and generally low for POC, between 19 and 65 $\text{mg C m}^{-2} \text{day}^{-1}$, with the exception of the N-ICE2015 station N4 in May, which had a distinctly higher export flux of 154 $\text{mg C m}^{-2} \text{day}^{-1}$ despite its low POC standing stock (4.9 g C m^{-2}). During the blooms, the export fluxes varied greatly, with fluxes generally 10-fold higher than during the pre-bloom period, ranging from 1.3 to 10 $\text{mg m}^{-2} \text{day}^{-1}$ and from 150 to 513 $\text{mg C m}^{-2} \text{day}^{-1}$ for Chl *a* and POC, respectively. The exception was N-ICE2015 station N6, where export fluxes remained low (0.5 $\text{mg Chl a m}^{-2} \text{day}^{-1}$ and 74 $\text{mg C m}^{-2} \text{day}^{-1}$) while standing stocks were high (351 mg Chl a m^{-2} and 15 g C m^{-2}). The post-bloom August conditions had relatively low Chl *a* export flux (0.6–1.7 $\text{mg m}^{-2} \text{day}^{-1}$) but elevated POC export flux (108–215 $\text{mg C m}^{-2} \text{day}^{-1}$), consistent with the higher POC standing stocks relative to Chl *a* standing stocks at those stations in August. Notably, the highest Chl *a* and POC standing stocks were found at the southernmost stations and over the southern Yermak Plateau (CarbonBridge and N-ICE2015 stations), but the highest export fluxes were measured

over the shelf break north of Svalbard (TRANSSIZ stations) (Figure 4).

Vertical Depth Profiles of POC Export

Vertical depth resolution of POC export fluxes revealed that not only the magnitude of the exported material but also the vertical profiles changed over the seasons sampled. During winter, pre-bloom, and post-bloom conditions, the vertical profiles of POC fluxes remained low and relatively constant from the surface down to 100–200 m depth (Figure 7). For the bloom stations, the vertical export flux profiles showed a sharp decrease from the upper water column to depths of 100–200 m (Figure 7). The strongest attenuation of POC export was observed at bloom stations T1, T4, and N7, with very high export fluxes of 964–1,165 $\text{mg C m}^{-2} \text{day}^{-1}$ at 25–30 m decreasing to 165–513 $\text{mg C m}^{-2} \text{day}^{-1}$ at 100 m.

Loss Rates

During winter, vertical POC export fluxes were low (19–57 $\text{mg C m}^{-2} \text{day}^{-1}$), equivalent to a loss of 0.6–2% day^{-1} of the POC standing stock, but increased with time (Figure 8). During the pre-bloom phase, POC fluxes were generally lower, but noteworthy differences were observed between the N-ICE2015 and TRANSSIZ stations, all of which were sampled over the Yermak Plateau (Figure 8). The TRANSSIZ sampling stations (T5, T6, and T7), located in dense ice cover in June, had low vertical export fluxes at 100 m ($\sim 30 \text{ mg C m}^{-2} \text{day}^{-1}$), representing a daily loss of 0.5–0.9% day^{-1} of the POC standing stock. The N-ICE2015 stations (N4, N5), sampled in mid-May,

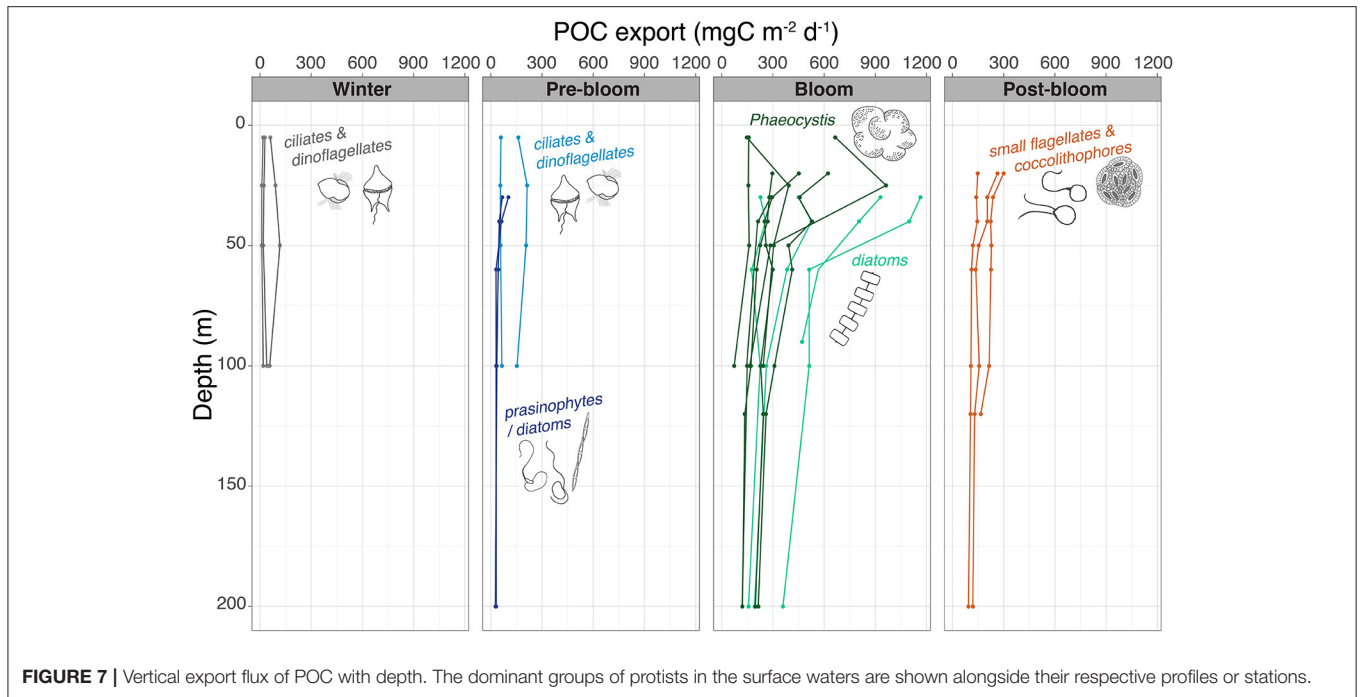


FIGURE 7 | Vertical export flux of POC with depth. The dominant groups of protists in the surface waters are shown alongside their respective profiles or stations.

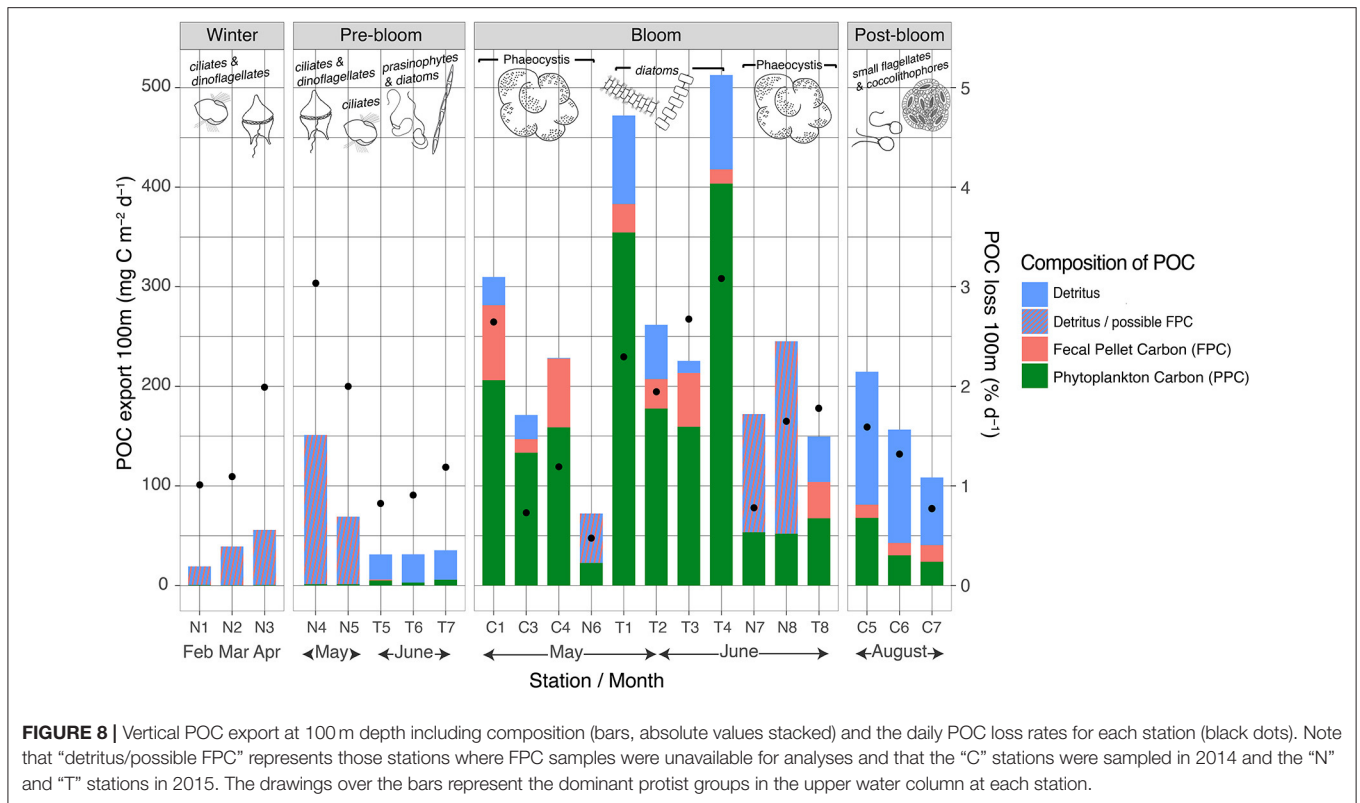


FIGURE 8 | Vertical POC export at 100 m depth including composition (bars, absolute values stacked) and the daily POC loss rates for each station (black dots). Note that “detritus/possible FPC” represents those stations where FPC samples were unavailable for analyses and that the “C” stations were sampled in 2014 and the “N” and “T” stations in 2015. The drawings over the bars represent the dominant protist groups in the upper water column at each station.

had higher 100 m export fluxes. At station N4, these fluxes translated into a high daily POC loss rate of 3% day⁻¹.

A large variation was found within and between the stations sampled during blooms, with POC fluxes at 100 m ranging from

74 to 513 mg C m⁻² day⁻¹, translating into daily loss rates of 0.4–3% day⁻¹ at 100 m (Figure 8). Furthermore, the strongest attenuation was at station N7 (Figure 7). At this station, vertical export rate at 100 m was only 17% of the flux at 25 m, while

at stations T1–T4, 100 m export fluxes were 44–51% of that at 30 m (Figure 7). The highest 100-m vertical export was found at sampling stations T1 and T4, sampled in late May and early June over the shelf break north of Svalbard (Figure 1). Here, the POC loss rates were 2–3% of the standing stock (Figure 8). Notably, the four shelf-break stations had consistently higher 100 m loss rates for Chl *a* than for POC (3–6% day⁻¹ compared to 2–3% day⁻¹). For the southernmost stations sampled in May (C1–C4, Figure 1), the POC export and loss rates at 100 m varied greatly (Figure 7), between 74 and 310 mg C m⁻² day⁻¹ and between 0.4 and 2.6% day⁻¹, respectively (Figure 8). Station C1 had a higher loss rate of POC (2.6% day⁻¹) than for Chl *a* (1.7% day⁻¹). Notably, the highest depth-integrated standing stocks of Chl *a* and POC (532 mg m⁻² and 23.4 g m⁻², respectively) were found at station C3, yet with low loss rates (0.6 and 0.7% day⁻¹, respectively), and POC fluxes of 171 mg C m⁻² day⁻¹ at 100 m. Likewise, at sampling stations over the Yermak Plateau (N6, N7, and N8), Chl *a* standing stocks of ~350 mg m⁻² were recorded, which were some of the highest values observed in the study, yet had Chl *a* loss rates of <0.4% day⁻¹ at 100 m.

Finally, the stations in a post-bloom stage (C5, C6, and C7), sampled in August on the shelf north of Svalbard, found POC export fluxes of 108–215 mg C m⁻² day⁻¹ and POC loss rates of 0.8–1.6% day⁻¹ (Figure 8).

Vertical Export Composition

During winter, the composition of the material exported at 100 m was largely unidentifiable particles classified as detritus (Figure 8). The calculated contribution of PPC to POC flux was ≤2%.

During the pre-bloom stage, the exported material was mainly comprised of detritus, 0–3% being FPC, and with a PPC contribution of 9–16% (Figure 8). The N-ICE2015 stations (N4 and N5), sampled in mid-May over the Yermak Plateau, had higher POC fluxes (64–153 mg C m⁻² day⁻¹) but with a lower contribution of PPC (<2%) (FPC contribution was unknown).

At the bloom-classified stations T1, T2, T3, and T4, sampled in late May and early June over the shelf break north of Svalbard (Figure 1), the highest 100-m vertical export fluxes were found and were composed of 75–79% PPC and 2.8–6.0% FPC (calanoid copepod and euphausiid pellets) (Figure 8). Of the southernmost stations in bloom (C1, C3, and C4) and those over the Yermak Plateau or Sofia Basin (N6, N7, N8, and T8), the highest fluxes were found where the exported POC had a larger contribution of FPC and/or detrital material (stations C1, C4, N8, and T8). For example, station C1, sampled in May at the southernmost area (Figure 1), had a 24% FPC contribution to the POC exported at 100 m, mainly produced by large calanoid copepods and euphausiids. Stations N6, N7, and N8 had Chl *a* standing stocks of ~350 mg m⁻², some of the highest values observed in the current study, yet the contribution of PPC to exported POC was only 21–32% (Figure 8). FPC data were unavailable for analysis for these N-ICE2015 stations.

Of the POC exported during the post-bloom stages, 19.4–31.6% was found to be PPC derived and 6.1–15.4% to be FPC derived (calanoid copepod and euphausiid pellets) (Figure 8).

Similar to winter and pre-bloom conditions, most of the exported POC was of detrital origin.

DISCUSSION

The current study is the first seasonal compilation of the fate of phytoplankton blooms in the SIZ north of Svalbard, including under-ice blooms. Despite the relatively large area covered during this study, clear seasonal patterns were found: low winter, pre-bloom, and post-bloom biogenic standing stocks and export fluxes and a short and intense productive season in May and June, with the exception of the northern Yermak Plateau. The results revealed how high export fluxes, up to 513 mg C m⁻² day⁻¹ at 100 m, can occur even in areas densely covered by thinning sea ice. These under-ice blooms were aided by favorable environmental conditions including stratification, winter-accumulated nutrients, a sufficient phytoplankton inoculum, and light transmission to the surface water column through melt ponds or leads (Assmy et al., 2017; Randelhoff et al., 2018; Katlein et al., 2019; Kauko et al., 2019). The distance to open water was a better predictor of standing stocks and carbon export than sea ice cover at the stations. Furthermore, despite spatial and temporal heterogeneity, two types of blooms could be distinguished: one dominated by the prymnesiophyte *P. pouchetii* and one dominated by diatoms. The largest vertical export fluxes of POC and Chl *a* were found at bloom stations dominated by diatoms, while the largest standing stocks were found at stations dominated by *P. pouchetii*. Previously, the ice edge (marginal ice zone) was considered the most productive region in the area, but our study recorded intense blooms and high export events under ice-covered waters. The results reveal large variations in carbon export fluxes during different blooms and highlight the roles of phytoplankton community composition, the synchrony with zooplankton grazers, and the impact of advection in determining the fate of blooms in ice-covered waters.

Earlier Phytoplankton Blooms Trigger Earlier Vertical Export Pulses

As the sea ice cover in the Arctic Ocean is shrinking and thinning, blooms of phytoplankton are now beginning earlier and already partly below the consolidated ice cover (Kahru et al., 2011; Ardyna and Arrigo, 2020). Through a series of conceptual models, Wassmann and Reigstad (2011) suggested that the earlier onset of blooms would be reflected in the export of carbon, with the largest changes occurring in the northern portions of the SIZ. The study area north of Svalbard has experienced dramatic changes in sea ice conditions in the last decade with increased fragmentation of sea ice. No study has yet been able to quantify the vertical export rates under the new sea ice regime. During the current study, early and intense phytoplankton blooms were recorded in May and June in both 2014 and 2015, with Chl *a* and POC standing stocks reaching up to 0.5 and 23 g m⁻², respectively, and POC fluxes exceeding 500 mg C m⁻² day⁻¹ at 100 m (Figure 4). This is a substantial increase when compared to a study performed in the same region in the summer of 1991 and

1997, when the ice was much thicker (classified as multiyear ice), where very low POC export fluxes were found in July ($17\text{--}26\text{ mg C m}^{-2}\text{ day}^{-1}$ at 100 m, $\sim 20\text{--}50\text{ mg C m}^{-2}\text{ day}^{-1}$ at 20–25 m) (Andreassen et al., 1996). As Andreassen et al. (1996) describe pre-bloom conditions in July, with surface nitrate concentrations of 4–6 mM, we strongly suggest a trend that phytoplankton blooms are occurring earlier in the season under the new sea ice regime. The earlier shift in carbon export fluxes has been found in the Central Arctic (Lalande et al., 2019), although the flux is mainly governed by ice algae, which play a negligible role in this study (Olsen et al., 2017).

Very little is known about winter and early spring vertical export flux, when access to the northern SIZ is logistically challenging. Our study reveals that during winter and the pre-bloom phase, under very dense sea ice cover, generally low POC export fluxes prevailed ($19\text{--}57\text{ mg C m}^{-2}\text{ day}^{-1}$) (Figure 4). These fluxes are comparable to observations in the Barents Sea in March 1998 ($20\text{--}70\text{ mg C m}^{-2}\text{ day}^{-1}$, with lower values in Arctic water masses and higher values in Atlantic water masses) (Olli et al., 2002). The ice cover during the current study was mostly composed of first-year ice (FYI) and second-year ice (SYI) (Granskog et al., 2017; Katlein et al., 2019), and high vertical export fluxes between $72\text{--}513\text{ mg C m}^{-2}\text{ day}^{-1}$ and Chl *a* standing stocks of $69\text{--}352\text{ mg m}^{-2}$ were found below sea ice cover of 91–100%, similar high values used to be typical only in the SIZ of the highly productive Barents Sea (Wassmann and Slagstad, 1993; Wassmann et al., 2006a; Reigstad et al., 2008). Our study shows one of the largest and earliest phytoplankton blooms recorded for the European Arctic north of 80°N and allows for the first time the evaluation of the strong seasonality in daily vertical carbon export of this previously inaccessible region of the European Arctic.

“The Under-Ice Productivity Belt” – Blooms and Accompanying Export Found up to 250 km Into the SIZ

As the SIZ of the European Arctic has shifted northwards from the Barents Sea to the area north of Svalbard, it has moved into a heterogeneous area. North of Svalbard, the continental slope encompasses sharp changes in bathymetry, which influences the behavior and location of the topographically steered AW inflow and the degree of mixing with surrounding water masses (Crews et al., 2018; Renner et al., 2018). The effects of AW advection are discussed in the *Export in a Dynamic Advective Environment* section below. Additionally, the thinner sea ice is moving and fragmenting more rapidly with the Transpolar Drift (Spreen et al., 2011; Krumpen et al., 2019). Situated in a dense but variable ice-covered area, our results show that the location in relation to the ice edge and open water was generally more important in determining the phytoplankton bloom magnitude (in terms of Chl *a* standing stocks) than the percentage ice cover itself (Figure 5). Closer to the ice edge, the sea ice is more likely to be fragmented and characterized by leads (Willmes and Heinemann, 2016), allowing for more light to penetrate into the water column (Assmy et al., 2017), which may be especially important before melt pond formation. We suggest that the distance to open water

be considered as an important predictor of blooms in sea ice-covered ecosystems. The most intense bloom ($532\text{ mg Chl } a\text{ m}^{-2}$, standing stock at station C3) was found in waters covered by 44% sea ice, roughly 35 km from open water (where sea ice cover was <15%). This bloom was classified as an ice edge bloom, which is often extensive as long as growth conditions are favorable (Wassmann et al., 1999; Tremblay et al., 2006), and contributes a substantial export of organic carbon to deeper waters ($170\text{ mg C m}^{-2}\text{ day}^{-1}$, 100 m at station C3).

In contrast, considerable standing stocks between 300 and $370\text{ mg Chl } a\text{ m}^{-2}$ in the upper 100 m were found at stations with >95% ice cover, and 50–130 km into the ice pack, illustrating that blooms can develop under densely ice-covered conditions. Early stages of a bloom (moderate Chl *a* standing stocks and high surface nitrates and silicic acid) were even found at 250 km from open water. These under-ice blooms are comparable to large ice edge blooms in the productive Barents and Beaufort Seas (Reigstad et al., 2008; Mundy et al., 2009) and show a similar or even higher daily vertical fluxes between $74\text{--}513\text{ mg C m}^{-2}\text{ day}^{-1}$ at 100 m depth. The blooms of the current study are likely locally produced blooms under the ice for the stations over the Yermak Plateau [see Assmy et al. (2017)] and possibly advected under the ice for those on the shelf break north of Svalbard where the AW boundary current is prominent [such as in Johnsen et al. (2018)]. Observations of under-ice blooms date back to the 1950s (Apollonio, 1961), but an increasing number of under-ice blooms have been reported during the last two decades, either produced locally under the ice (Fortier et al., 2002; Arrigo et al., 2012; Lowry et al., 2014; Mundy et al., 2014; Assmy et al., 2017; Ardyna et al., 2020) or advected below the ice cover (Metfies et al., 2016; Johnsen et al., 2018). More frequent observations of under-ice blooms can be partly attributed to the increased research efforts in the Arctic but largely to the changing sea ice conditions (Ardyna and Arrigo 2020). The current study area may be especially prone to large under-ice blooms in the future, with declining sea ice thickness causing changes in light climate (Nicolaus et al., 2012; Oziel et al., 2017; Katlein et al., 2019) and being positioned at the principal gateway for Atlantic water masses entering the Arctic Ocean (Aagaard et al., 1987; Crews et al., 2018; Renner et al., 2018). Although the environmental drivers of under-ice blooms have been examined [Ardyna et al. (2020) and studies therein], no study could yet provide any estimate of the carbon export potential of these under-ice blooms (Ardyna and Arrigo, 2020). Our study can reveal that if these blooms provide between $150\text{--}500\text{ mg m}^{-2}\text{ day}^{-1}$ of organic carbon to the deeper waters, they could certainly increase the productivity of mesopelagic and benthic habitats and must be included in productivity and carbon flow models of the Arctic Ocean.

Our results further highlight the importance of surface stratification for primary production and subsequent vertical export fluxes to increase in the SIZ. A clear seasonal trend with a sharp reduction in surface nitrate concentrations and a shallowing of the mixed layer depth was observed from winter to spring and summer (Figure 3). Notably, mixed layer depths of $\leq 30\text{ m}$ were always observed during blooms, suggesting that a shallow mixed layer (associated with melting sea ice) may aid

locally produced blooms to occur under the ice [supported by Strass and Nöthig (1996), M. Fortier et al. (2002), Tremblay et al. (2006), Leu et al. (2015), and Oziel et al. (2019)]. This is further supported by the strong negative correlation between mixed layer depth and Chl *a* standing stocks ($P = 0.000001$, **Supplementary Material**).

Winter surface nitrate concentrations were lower in late January (4.6 mmol m^{-3}) and increasing to a maximum of 9.4 mmol m^{-3} during pre-bloom conditions in May. The winter stations sampled during the N-ICE expedition were positioned further north in the Nansen Basin and northern Yermak Plateau where surface nutrients are generally lower (Bluhm et al., 2015), which likely explains this apparent increase. Continuous mixing throughout winter will further contribute to increasing surface nutrient concentrations toward early spring (Randelhoff and Guthrie, 2016). Blooms that occur earlier in the SIZ (i.e., in April) and further north in the AO where nitrate concentrations are lower and mixing is limited (Tremblay et al., 2015) would be subject to lower surface nutrient concentrations at the initiation of the bloom, potentially decreasing the bloom magnitude and the strength of pelagic–benthic coupling. On the other hand, surface nutrient concentrations have been shown to be well-mixed in the immediate Atlantic inflow using observations based on moorings equipped with nitrate sensors, deployed from 2012 to 2013 on the shelf break north of Svalbard (Randelhoff et al., 2015).

Seasonal Changes in Phytoplankton Community and Export Composition Are Reflected in the Magnitude of Export

Very few seasonal studies of daily export flux of organic material exist in the SIZ, especially during winter and for short-term sediment trap deployments [for other Arctic regions, see Juul-Pedersen et al. (2008), Olli et al. (2002), and Szymanski and Gradinger (2016)]. Samples collected using short-term sediment traps represent snapshots in time, but as they can be combined with *in situ* suspended water and community samples, they can examine central processes occurring in the upper water column, including the synchrony between phytoplankton blooms and zooplankton grazers, illustrating the partitioning of organic carbon in the ecosystems of the pelagic and benthic realms. Our results demonstrate how the seasonal and regional variability in the pelagic community and the composition of the exported material affect the magnitude and depth of exported POC in the SIZ north of Svalbard.

The flat winter profiles with low POC export fluxes and moderate loss rates indicate that the low amounts of organic matter were evenly spread through the water column (**Figure 7**). Pre-bloom POC standing stocks were similar to winter conditions, but Chl *a* standing stocks were 10-fold higher, which is noteworthy as it suggests a decoupling of phytoplankton from the heterotrophic community in the winter to pre-bloom transition at high latitudes (Behrenfeld and Boss, 2014). Furthermore, during the period preceding the bloom, one station in early May over the northern Yermak Plateau (station N4) had, despite low Chl *a* and POC standing stocks, elevated

POC loss to 100 m ($>3\% \text{ day}^{-1}$ of POC standing stock, **Figure 8**). The vertical export at this station was strongly dominated by the ciliate *M. rubrum* and coincided with the collapse of a dense *M. rubrum* bloom at the ice–water interface. The bloom collapse likely resulted in the downward migration of this species and interception by the deeper sediment traps (Olsen et al., 2019), which represents active export of organic matter as opposed to passive settling of particles.

During phytoplankton blooms, high biomass will accumulate as long as vertical export is low. This was the case at station C3, which had relatively low vertical export accounting for loss rates of only $0.7\% \text{ POC day}^{-1}$ and $0.6\% \text{ Chl } a \text{ day}^{-1}$ at 100 m, allowing for the buildup of the highest Chl *a* (532 mg m^{-2}) and POC (23 g C m^{-2}) standing stocks observed in the study. On the other hand, the highest vertical export fluxes are generally associated with intense blooms at the end of their growth phase when nutrients are exhausted (Agustí et al., 2020), grazing pressure is low, and the community composition allows for aggregation or mass flux of ungrazed cells (Wassmann et al., 2006a; Henson et al., 2019). The vertical export and loss rates varied greatly between the stations sampled during blooms, which highlights the need to examine the composition and vertical profiles of the exported material in order to outline the processes involved in the sedimentation of the blooms.

Diatoms are ballasted by silica frustules and often form long chains, while the prymnesiophyte *P. pouchetii* is a small flagellate ($\sim 5 \mu\text{m}$ in diameter) that often forms large polysaccharide-embedded colonies during blooms (Wassmann et al., 1999). Once diatoms terminate their growth phase, they tend to aggregate and ungrazed cells or resting spores can sink out of the surface layers at high rates, often leading to mass export events (Smetacek, 1985; Henson et al., 2006; Assmy et al., 2019). *Phaeocystis*, on the other hand, may form large blooms, but their contribution to carbon export is generally considered to be low (Reigstad and Wassmann, 2007; Wolf et al., 2016), unless ballasted by cryogenic gypsum (Wollenburg et al., 2018) or deep-mixed by eddies (e.g., Lalande et al., 2011). Recent findings also conclude that ballasted and sticky diatoms are generally poorly attenuated in the upper water column compared to non-ballasted *Phaeocystis*, and therefore, the relative contribution of diatoms is a good predictor for carbon flux attenuation, especially in the cold waters of the Arctic (Wiedmann et al., 2020). The current study found that the largest blooms were formed by *Phaeocystis*, but with loss rates $<2\% \text{ day}^{-1}$ of POC standing stock and with average POC flux of $190 \text{ mg C m}^{-2} \text{ day}^{-1}$ at 100 m. The diatom bloom on the shelf break revealed very high POC export fluxes out of the epipelagic (**Figure 8**), with loss rates $>2.5\% \text{ day}^{-1}$ and flux $>450 \text{ mg C m}^{-2} \text{ day}^{-1}$, despite having some nitrates and silicic acid available in the upper water column and a strongly stratified layer below the Chl *a* maximum. These POC export fluxes are comparable and higher than many previous investigations in the highly productive Barents Sea (Wassmann and Slagstad, 1993; Olli et al., 2002; Reigstad et al., 2008) and higher than export estimates for the future SIZ (Wassmann and Reigstad, 2011). Furthermore, even when organic carbon was strongly attenuated in the surface waters in diatom-dominated systems (**Figure 7**), their contribution to carbon export at 100 m was

higher than during *Phaeocystis* blooms. This demonstrates that diatom contribution to pelagic–benthic coupling is substantial and will very likely play a role in the fate of these blooms including the eventual carbon sequestration on the sea floor.

Furthermore, the locally produced under-ice *Phaeocystis* blooms over the Yermak Plateau had the lowest contribution of PPC to total POC export at 100 m of the bloom stations (Figure 8). Station N7, considered to be at peak bloom, showed great attenuation of organic carbon in surface layers, where POC export at 100 m was only 17% of that at 25 m. This demonstrates that at its peak (usually as growth conditions become unfavorable), where pelagic–benthic coupling is generally highest, the under-ice *Phaeocystis* bloom was attenuated and remineralized or grazed in the surface layers. On the other hand, two of the *Phaeocystis*-dominated sampling stations had notably higher export fluxes and loss rates of POC (Figure 8). At these stations, the contribution of fecal pellets to the carbon exported was quite high (25–30%), produced by large calanoid copepods and krill. Our results cannot confirm that these grazers fed directly on *P. pouchetii*, but nevertheless, it is likely that they facilitated much of the export in a *Phaeocystis*-dominated system [as found by Hamm et al. (2001)]. It is significant to note that out of the seven stations where *Phaeocystis* blooms were found in the current study, only two had depleted silicic acid stocks in the surface (Supplementary Material). Previous studies have suggested that diatoms will generally dominate blooms as long as silicic acid is not limiting (Ardyna et al., 2020). This was not the case in our study, which suggests that the light climate was possibly the decisive factor in the succession of these blooms (Assmy et al., 2017).

The post-bloom settings in August were associated with warmer and more saline waters, with shallow mixed layer depths, small flagellates and coccolithophores, and intermediate carbon export fluxes at 100 m, though with a lowered contribution of PPC (Figure 8). Likewise, the standing stocks of POC remained considerable (12–14 g C m⁻²), while those of Chl *a* were low (Figure 4). The FPC contribution to carbon export was also relatively low during the post-bloom stages (6–15% of total POC), possibly indicating a more detritus-derived carbon source, where grazing by large zooplankton was low (Rivkin et al., 1996). Small heterotrophs were more prominent during the post-bloom period (Lavrentyev et al., 2019), which would likely increase recycling and remineralization in the upper water column, preventing organic matter from sinking out of the epipelagic zone (Olli et al., 2002) and allowing high POC standing stocks to remain suspended. These findings agree with previous observations and predictions for the future SIZ, where a longer ice-free period late in the summer would be characterized by regenerated production, where the quality of the exported material would be increasingly degraded and recycled (Wassmann and Reigstad, 2011; Flores et al., 2019).

Export in a Dynamic Advective Environment

Since the early 2000s, the area investigated in this study has been subject to “Atlantification,” and as a result, it is characterized

by declining sea ice cover and weakening stratification between the Atlantic water and polar surface waters above it (Polyakov et al., 2017; Lind et al., 2018). Due to their positions at the inflow of Atlantic water into the Arctic Ocean, the stations over the Yermak Plateau are likely experiencing a different water mass stability than those north and northwest of Svalbard (Figure 1; Fer et al., 2017; Crews et al., 2018; Menze et al., 2019). Therefore, it is likely that spatial variability in blooms and subsequent export observed in the current study is to a large extent driven by the variability in Atlantic water advection.

The sampling stations in May and June were characterized by large spatial variability, ranging from pre-bloom to bloom conditions in a relatively small area. Pre-bloom conditions were found in June over the northern Yermak Plateau, possibly as a result of inadequate light [see Massicotte et al. (2019)], great distance to open water (≥ 200 km), a deep-mixed layer depth, and possibly weaker Atlantic water advection (pers. comm. Sebastian Menze). Over the southern Yermak Plateau, slightly closer to open water (50–130 km), the large *P. pouchetii* blooms were likely locally produced under the sea ice, as Atlantic water advection was not found to be prominent but the algal cells were adapted to the low but variable light conditions penetrating through refrozen leads (Assmy et al., 2017; Pavlov et al., 2017; Kauko et al., 2019). On the other hand, the *P. pouchetii* blooms at the southernmost stations closer to Fram Strait in May 2014 were likely ice edge blooms due to the reduced ice cover (44–52% cover) and seeded by the West Spitsbergen Current (Johnsen et al., 2018; Paulsen et al., 2018; Sanz-Martín et al., 2019; Vernet et al., 2019). The vertical export fluxes and relative contribution of PPC differed between these two types of *Phaeocystis* blooms. The locally produced bloom generally had lower POC export fluxes with a smaller contribution of PPC to total POC exported, while the advected ice edge *Phaeocystis* bloom had a larger contribution of both PPC and FPC to the POC export at 100 m (Figure 8). This might suggest that the latter advected bloom consisted of phytoplankton cells that were brought into less favorable conditions (nutrient or light limited) and that large grazers advected into the area and facilitated higher POC export fluxes out of the surface waters (Svensen et al., 2019; Vernet et al., 2019; Wassmann et al., 2019), as opposed to the under-ice bloom which was still growing under favorable conditions with limited grazing (Kauko et al., 2019).

The current study cannot ascertain whether the diatom blooms on the shelf break were locally grown or advected underneath the sea ice cover, but we suspect the latter. Model simulations suggest that the area along the shelf break receives up to 50 times more advected gross primary production than what is locally produced (Vernet et al., 2019). On the other hand, no ice edge blooms were observed in the area, using satellite imagery, in the days prior to these blooms. The blooms may nevertheless have been seeded by advection of deeper Atlantic water masses (found below 40–170 m at these stations). Very high vertical POC export fluxes were found in this area (>500 mg C m⁻² day⁻¹), mostly consisting of PPC (~70%). Diatoms have been suggested to require stable light conditions to sustain growth rates (Hoppe et al., 2015; Assmy et al., 2017) and tend to sink faster under limited silicic acid conditions (Gross, 1948; Krause

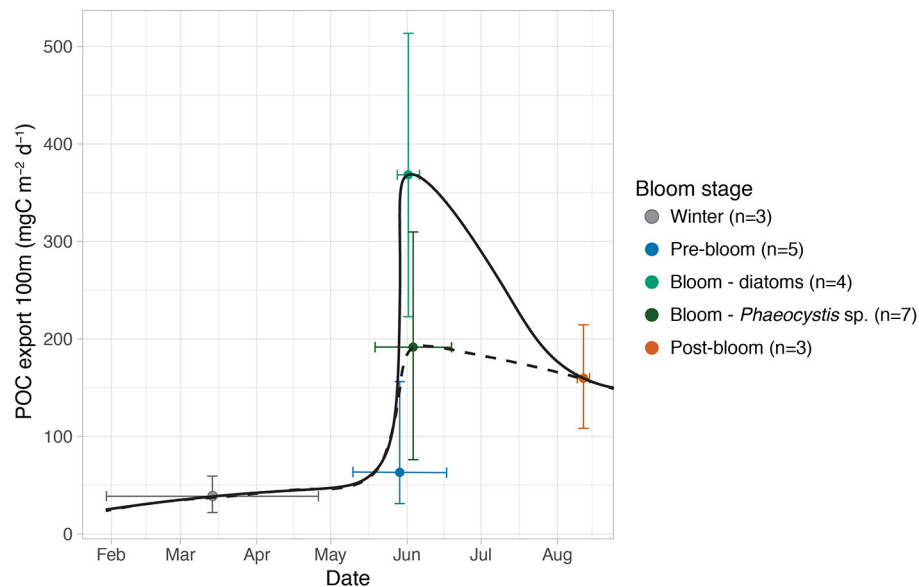


FIGURE 9 | General seasonal progression of vertical POC export fluxes at 100 m in the region north of Svalbard (mean mg POC m⁻² day⁻¹, +/- full range). The export fluxes are separated into bloom conditions and dominant phytoplankton groups during the blooms (n, number of measurements). Black lines: rough visual interpolation of seasonal trend using diatoms (solid) and Phaeocystis (dashed) blooms.

et al., 2018; Agustí et al., 2020). The increase in export fluxes from stations T1 to T4 (Figure 8), which were separated in time but similar in location, may suggest that once silicic acid became limiting (Figure 2), the growth conditions for these diatoms were reduced and subsequently increased their sinking velocity (Agustí et al., 2020). This is further supported by the deepening of the Chl *a* maxima from 10 to 25 m during the 8 days between sampling at stations T1 and T4, a reduction in surface silicic acid, and a very low mean light transmittance of 0.4 and 0.2% at these stations, respectively (Katlein et al., 2019). If the diatom bloom was advected under the pack ice, the change in light conditions may have contributed to the increased export flux, but unfortunately, our study cannot provide further evidence for the origin of this bloom.

The variability in FPC contribution to total POC export during the blooms (Figure 8) may also have been a result of the variable presence of Atlantic water and time of year. Large copepods (such as *Calanus* spp.) and krill were generally the main contributor to FPC export (data not shown). The southernmost region closer to Fram Strait and the WSC, where FPC export was the highest, is likely to receive larger advective biomass of *Calanus* earlier in the spring than areas further northeast along the AW boundary current, where they arrive or develop later in July and August (Basedow et al., 2018; Wassmann et al., 2019). This suggests that the reduced FPC contribution to POC export on the shelf break north of Svalbard may be a consequence of mismatch in timing between the phytoplankton bloom and the presence of large grazers (Wexels Riser et al., 2008).

Finally, advective variability highlights three concerns in the sampling methodology: (1) sampling of suspended biomass and calculations of depth-integrated standing stocks represent

a snapshot in time, and therefore, subsequent comparisons to exported material (integrating over ≥ 24 h) and loss rates can be over- or underestimated; (2) sediment traps that drift with the ice instead of the water column may not follow even a semi-Lagrangian drift, an assumption upon which the calculations of loss rates and export fluxes are based (Olli et al., 2007); and (3) confounding of spatial and temporal trends by geographic spread of the stations. These considerations are especially important in advective areas, where production and sedimentation can occur many kilometers apart. Although these concerns add uncertainty to the results, the ice-tethered sediment traps are currently the only feasible method for collecting direct measurements of daily vertical export flux in ice-covered waters in combination with suspended particle stocks, and their long history of use makes them instrumental for comparisons.

CONCLUSION

Through extrapolation of the short-term measurements of vertical POC export from the current study, rough estimations of annual POC flux at 100 m for the area were found to be in the range of 24–34.5 g C m⁻² year⁻¹, depending on the conservatism of the estimate (for details, see **Supplementary Materials**). This calculation, although rough, is the first estimate of annual carbon fluxes in the new SIZ north of Svalbard, including under-ice blooms. Notably, it is comparable to sediment trap estimates from the southern Barents Sea (32–97 g C m⁻² year⁻¹) (Slagstad and Wassmann, 1996; Reigstad et al., 2008), but much higher than long-term sediment trap deployment estimates for other parts of the Arctic

(0.1–10 g C m⁻² year⁻¹), with the lowest values from the central Arctic Ocean, summarized in Stein and Macdonald (2004).

Moreover, the present export fluxes were higher and occurred earlier in the year than previously observed in the same region (Andreassen et al., 1996; Lalande et al., 2014). Our results indicate that the thinner and more deteriorated sea ice in the area (Itkin et al., 2017) has extended the SIZ further north. Previously, the marginal ice zone was considered the most productive region in the area. Our study found intense phytoplankton biomasses and high carbon export fluxes, of up to 513 mg C m⁻² day⁻¹, to be found up to 250 km from open water, demonstrating stronger pelagic–benthic coupling in ice-covered waters than formerly reported.

The vertical export fluxes of POC followed the seasonal bloom progression, but the mass export of organic matter was influenced by the bloom composition and synchrony with grazers. Whether diatoms or *P. pouchetii* dominated the bloom had an impact on the carbon export, as the highest export fluxes were found in stations dominated by diatoms (Figure 9). If *Phaeocystis* blooms become more common in the Arctic (Orkney et al., 2020), it is likely to reduce absolute mass flux and thus limit pelagic–benthic coupling in the future Arctic SIZ, especially if no special circumstances are present to facilitate their export (e.g., Wollenburg et al., 2018). The presence of advected Atlantic water masses likely affected the magnitude of export fluxes by changing the growth conditions of phytoplankton blooms and their synchrony with the large zooplankton grazers. We emphasize the need to monitor seasonal successions of flux compositions, the timing and growth conditions of blooming phytoplankton under the ice, and their interactions with heterotrophic communities in order to make accurate predictions about the fate of productivity and the partitioning of organic carbon in the ecosystems of the pelagic and benthic realms in the future Arctic seasonal sea ice zone.

DATA AVAILABILITY STATEMENT

The datasets generated for this study are available on request to the corresponding author. All data are made available to the public; see PANGEA (<https://www.pangae.de>) and the Norwegian Polar Data Center (<https://data.npolar.no>).

AUTHOR CONTRIBUTIONS

PA, MR, IP, and AN planned parts of the various field campaigns. CD, PA, LO, AR, MR, and TK collected the samples

and conducted field measurements. AR and AN conducted oceanographic measurements. TK contributed to the sea ice data and performed sea ice analyses from satellite data. CD, AT, and JW did the microscopy analyses. IP performed the HPLC analysis. CD did the data analysis and wrote the first draft of the manuscript, and all co-authors contributed to the final version. All authors contributed to the article and approved the submitted version.

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fmars.2020.525800/full#supplementary-material>

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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