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Diatom composition and fluxes over the Northwind Ridge, western Arctic Ocean: impact of marine surface circulation and sea ice distribution

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Abstract

Over the last decades the western Arctic Ocean has undergone unprecedented environmental changes. However, long-term marine phytoplankton *in situ* observations are still rare and therefore insufficient to fully characterize evolutionary trends. This study investigate diatom flux and composition in sediment trap material collected in the Northwind Ridge, western Arctic Ocean from August 2008 to September 2009. Our data show that *Chaetoceros* resting spores are the predominant species accounting for >40% of the diatom composition. The sea ice diatom group, which includes *Fossula arctica*, *Fragilariopsis cylindrus* and *F. oceanica*, dominates the rest of the assemblage throughout the observation period. While the diatom fluxes in winter are extremely low, higher values are found in summer, with summer 2009 flux values being twice as high as in 2008. High total mass and diatom fluxes in summer 2009 are attributed to the intertwined effect of a weakened Beaufort Gyre, strengthened Pacific Water Inflow (PWI) and distribution pattern of the sea ice. Enhanced values of coastal diatoms and terrigenous proxies in summer 2009 are in agreement with intensified PWI. Sea ice diatoms and sea ice biomarker IP₂₅ fluxes are both high during the sea ice melting season and significantly correlated ($r^2 = 0.64$, $p < 0.01$). Our data also suggest that sea ice diatoms are prone to selective dissolution in the water column and sediments, implying biases on diatom assemblages and subsequently on paleoceanographic reconstructions.

Key words: Diatoms, Sediment trap, Chukchi Sea, Northwind Ridge, Sea ice, Pacific water inflow, Beaufort Gyre

1. Introduction

Over the past decades, the Arctic Ocean has undergone unprecedented environmental changes with drastic impacts on the terrestrial (Post et al., 2013) and marine ecosystems (Carroll and Carroll, 2003; Grebmeier et al., 2006, 2012; Arrigo et al., 2008; Wassmann, 2011; Wassmann et al., 2011). Among other factors, the decline of sea ice has been shown to affect marine phytoplankton production and distribution patterns (e.g. Arrigo et al., 2008; McLaughlin et al., 2011; Coupel et al., 2012; Arrigo and van Dijken, 2015; Renaut et al., 2018). Blooms in autumn are now reported during the prolonged ice-free period in the Arctic (Ardyna et al., 2014) as well as significant changes in phytoplankton community due to enhanced freshening caused by global warming (Li et al., 2009; He et al., 2012; Coupel et al., 2015; Zhuang et al., 2018; Lee et al., 2019).

The highly productive Chukchi Sea in the western Arctic has experienced rapid sea ice retreat (Comiso, 2012; Serreze et al., 2016) leading to increased primary production and enhanced biological carbon uptake (Harada, 2016). The strengthening of the Pacific Water Inflow (PWI) bringing nutrients into the Chukchi Sea during the last decade (Woodgate et al., 2012; Woodgate, 2018) is another factor that may have altered the phytoplankton structure (Zhuang et al., 2016). The Chukchi Sea is thus a critical region for understanding oceanographic on-going and future changes in the Arctic Ocean and their consequences on polar ecosystems.

Apart from flagellates, diatoms appear to dominate the phytoplankton and bottom sea ice community in the Chukchi Sea (Booth and Horner, 1997; Sukhanova et al., 2009; Poulin et al., 2011; Joo et al., 2012). Because of diatom sensitivity to environment changes, their composition, abundance, distribution as well as size have been used as indicators of marine environmental conditions (Zernova et al., 2000; Smol and Stoermer, 2010; Romero and Armand, 2010). For instance, recent studies speculated that the occurrence of the endemic North Pacific diatom *Neodenticula seminae* in the North Atlantic might reflect stronger PWI driven by sea ice loss (Reid et al., 2007; Poulin et al., 2010; Bluhm et al., 2011). Better understanding of the relationship between diatoms and environmental variables today is also essential to produce robust paleoceanographic reconstructions based on fossil diatoms in Arctic sediments (e.g. Ren et al., 2009; Sha et al., 2016, 2017; Miettinen, 2018). Paleo-records are still very sparse in the western Arctic in part because of our poor knowledge of modern diatom ecology and the complex deposition pathways in this environment.

Sediment traps have been extensively deployed throughout the world ocean to study biological processes and carbon export (Romero and Armand, 2010). In the Arctic, sediment trap deployments have focused on the continental shelves and slopes (e.g. Wassmann et al., 2004 and references therein; O'Brien et al., 2006, 2011; Fahl and Nöthig, 2007; Forest et al.,

2007, 2008, 2011; Lalande et al., 2009; Fahl and Stein, 2012) as well as the deep basins (Honjo et al., 2010; Hwang et al., 2015; Lalande et al., 2019). Few studies have been undertaken in the Chukchi Sea. Analyses of the ice-tethered sediment traps over the Chukchi Rise revealed the heterogeneous patterns of biological processes and variety of particulate organic carbon sources (Honjo et al., 2010). Recent studies of the first multi-years mooring observations in the Northwind Abyssal Plain have documented seasonal variations of phyto- and zooplankton (Matsuno et al., 2014; Ikenoue et al., 2015; Onodera et al., 2015, 2016; Tokuhira et al., 2019) and also evidenced early winter high flux induced by cold eddies from the shelf-break (Watanabe et al., 2014, 2015).

In this study, we report on seasonal variations of diatom vertical flux obtained from the sediment trap mooring deployed over the Northwind Ridge, Chukchi Borderland from August 2008 to September 2009 as part of the Chinese National Arctic Research Expedition (CHINARE) program. These data are compared to the biomarker data of Bai et al. (2019) measured in the same samples and discussed in light of the overlying water mass characteristics, sea ice concentrations and atmospheric circulation in the western Arctic Ocean.

2. Oceanographic settings

Station DM is located at the southern Northwind Ridge, a steep ledge extending from the shallow Chukchi Shelf into the Arctic deep basin as a part of the Chukchi Borderland (Fig. 1). The surface hydrology of the region is influenced by the dynamic Beaufort Gyre (BG) circulation, the PWI entering the Arctic Ocean through the Bering Strait and the seasonal sea ice coverage.

The anticyclonic BG circulation is driven by the Beaufort High (Proshutinsky and Johnson, 1997). Over the last two decades, the stronger BG circulation has resulted in enhanced freshening in the Canadian Basin (Giles et al., 2012) with significant impact on the phytoplankton productivity and polar ecosystem (He et al., 2012; Coupel et al., 2015).

The PWI is primarily driven by a sea level difference between the Bering and Chukchi Seas (Coachman and Aagaard, 1966), leading to an average northward water transport of ~0.8 Sv (Roach et al., 1995). The PWI splits into three water masses in the Chukchi Sea, i.e. the saline and nutrient-rich Anadyr Water (AW) as the western branch, the fresher and nutrient-depleted Alaska Coastal Water (ACW) to the East, and in between the Bering Shelf Water (BSW) of intermediate salinity (Fig. 1; Woodgate et al., 2005a; Grebmeier et al., 2006). In the past two decades, long-term mooring observations in the Bering Strait have shown a gradual increase of the PWI volume transport (~70% from 2001 to 2014; Woodgate et al., 2012; Woodgate, 2018), entraining more heat and nutrients into the Arctic Ocean. In the western Chukchi Sea, the fresher Siberian Coastal Current (SCC) periodically interacts with the AW.

Sea ice occurs in the study area from November to July while it reaches a minimum over the northern Chukchi Sea in September (Fig. 1). Several studies have shown that sea ice reduction induced by PWI impacts on marine ecosystem (e.g. Shimada et al., 2006; Harada, 2016; Serreze et al., 2016). It has also been hypothesized that sea ice motion is also correlated, at least partially, with the Arctic Oscillation (AO) (Rigor et al, 2002; Shimada et al., 2006).

3. Material and methods

3.1 Sediment trap samples

A one year-round mooring was deployed at Station DM (DM hereafter) on the southern Northwind Ridge, in the vicinity of the Canada Basin, from August 2008 to September 2009 (Fig. 1; 74°24.0' N, 158°14.0' W, 1650 m water depth). This conical sediment trap (McLane PARFLUX Mark 78H-21) was equipped with 21 sampling cups and installed at ~870 m water depth. Samples were generally collected every two weeks (15 or 16 days, depending on the month) during the diatom production season (July to November) while during the low flux period in winter (December to June) sampling intervals were longer (28 to 31 days, depending on the month; Bai et al., 2019). The 21 sampling cups were filled with artificial seawater (salinity~35) and antiseptic HgCl₂ to preserve trap material from degradation.

After recovery, the wet samples were sieved using a 1-mm mesh nylon sieve to remove swimmers. The fine fractions (<1 mm) were split into aliquots with a McLane wet sample divider (WSD-10). An aliquot (1/4 or 1/8) was filtered on a polycarbonate filter (0.45 µm pore size, 47 mm diameter) for diatom and biochemical analyses. Samples were then dried in an oven at 45 °C for 72 h (Bai et al., 2019).

3.2 Diatom preparation and analysis

Quantitative diatom slides were prepared according to the standard method developed at the Alfred Wegener Institute, Germany (Gersonde and Zielinski, 2000). For each cup, 30 mg or 15 mg sub-sample was used. Although only 1 mg was available for cup 14# (May 2009), the sample was analyzed to keep the entire record intact. The subsamples were then treated with HCl (~36% w/w concentration) and H₂O₂ (~30% w/w concentration) to remove calcareous and organic material. After rinsing, a known aliquot of the residue was evenly dropped on a cover glass (24 mm × 24 mm). Permanent slides were mounted with Naphrax for diatom identification ($n_D \sim 1.7$; Fleming, 1954).

Around 400 diatom valves (max. 604.5, min. 372) were counted for each sample with a Motic BA400E microscope at ×1000 magnification. For cup 14# only 215 diatom valves could be counted. Diatom valves were counted according to the method of Schrader and Gersonde (1978). Diatoms were identified to species level following Medlin and Priddle (1990), Hasle

and Syvertsen (1997), Suto et al. (2004), Katsuki et al. (2009), Obrezkova et al. (2014) and Tsoy and Obrezkova (2017). Because resting spores of genus *Chaetoceros* can hardly be identified at a species level by light microscopy, they were combined and labeled as *Chaetoceros* resting spores (RS).

The diatom counts were converted into relative abundances with respect to the total diatom assemblage. The diatom concentration (valves per gram dry sample) was estimated using the following formula (Esper et al., 2010):

$$\text{Diatom concentration} = (1/dw) \times (csa/ta) \times (sv/split) \times (vn/tn) \quad (1)$$

in which, dw is the dried sample weight in milligrams, csa is the area of the cover slide (576 mm²), ta is the area of one counted traverse (6 mm²), sv is the volume of processed diatom suspension (6-15 ml), $split$ is the volume of aliquot dropped onto the cover slide (from 27 to 81 ×10⁻² ml), vn is the total counted diatom valves and the tn is the number of fully counted traverses.

The diatom fluxes were calculated by the following equation:

$$\begin{aligned} \text{Diatom flux [valves } m^{-2} d^{-1}] \\ = \text{Total mass flux [mg } m^{-2} d^{-1}] \times \text{Diatom concentration [valves mg}^{-1}] \end{aligned} \quad (2)$$

where the diatom concentration is derived from equation (1).

Diatom taxonomic diversity was evaluated by the H-index based on the Shannon-Weaver formula (Shannon and Weaver, 1949):

$$H = -\sum P_i \log_2 P_i \quad (3)$$

where H is the diversity index and P_i is the relative abundance of species i .

3.3 Biomarker analysis

The concentrations of IP₂₅ and brassicasterol (24-methylcholesta-5,22E-dien-3β-ol) published by Bai et al. (2019) are used for comparison with our diatoms fluxes to discuss sea ice conditions. Campesterol (24-methylcholest-5-en-β-ol) and 24-ethylcholest-5-en-3β-ol are also used to assess higher plant inputs. All sterols were silylated with 100 μL BSTFA (bis-trimethylsilyl-trifluoroacet-amide) (80 °C, 1 h) prior to the gas chromatography (GC) analysis. The GC analyses were carried out on a Varian 3300 with a septum programmable injector and a flame ionization detector (Bai et al., 2019).

3.4 Carbonate and lithogenic matter

The CaCO₃ content of sinking material was calculated from particulate inorganic carbon (PIC) values determined by the difference between total carbon (TC) and particulate organic

carbon (POC; Bai et al., 2019). The lithogenic matter (LM) was estimated as the difference between the total mass (TM; from Bai et al., 2019) and the biogenic material composed of CaCO_3 , opal (Wu et al., unpublished data) and organic matter (OM) that conventionally convert from POC by a factor of 1.8 (Müller et al., 1986):

$$LM = TM - (CaCO_3 + Opal + OM) \quad (4)$$

The lithogenic matter percentage was then calculated by the following equation:

$$LM (\%) = LM / TM \quad (5)$$

3.5 Environmental data

Daily environmental data over the study period at DM, including shortwave radiation, snow thickness, sea ice cover and sea ice thickness, were from the National Centers for Environmental Prediction (NCEP)/Climate Forecast System Reanalysis (CFSR) 6-hourly dataset (Saha et al., 2010, 2014).

Monthly Pacific inflow data, e.g. volume transport, temperature and salinity, were acquired from the “Bering Strait: Pacific Gateway to the Arctic” project (Woodgate et al., 2015; Woodgate, 2018).

Regional sea ice concentrations were from Nimbus-7 SMMR and DMSP SSM/I-SSMIS Passive Microwave Data with a grid resolution of 25×25 km (Cavalieri et al., 1996).

Summer (July, August, September) monthly averaged sea surface chlorophyll-*a* concentrations for a $2^\circ \times 2^\circ$ area around DM were retrieved from the Moderate Resolution Imaging Spectroradiometer (MODIS) with a 4×4 km resolution and calculated by algorithm of Gohin et al (2002) collected by GlobColour data (<http://globcolour.info>).

4. Results

4.1 Environmental conditions

Station DM experiences polar night from November to mid-February and receives maximum solar irradiance of $\sim 369 \text{ W m}^{-2}$ in summer, from mid-April to mid-July (Fig. 2a). Seasonal sea ice was present from late October 2008 to mid-August 2009. The ice-free season (sea ice concentration $< 15\%$) is approximately one month longer in 2008 than in 2009 (mid-July to late October vs. mid-August to early October, respectively; Fig. 2b). Winter sea ice thickness over the station is 2.19 m on average with a maximum thickness of 2.79 m recorded in March 2009 (Fig. 2c). Similarly, snow on ice appears in mid-October 2008 and nearly has thawed out early June 2009 (Fig. 2d). The snow accumulation is over 25 cm in winter.

4.2 Total mass flux

The seasonal pattern of the total mass fluxes (TMF) earlier described by Bai et al. (2019), show higher values in August and September, both in 2008 and 2009 (Fig. 2e). However, the TMF in September 2009 is ca. eightfold larger than in 2008. Particulate organic carbon (POC) fluxes exhibit similar fluctuations as TMF, with extremely low values in winter and early spring, from November 2008 to June 2009 (Fig. 2f). One exception to this is the POC flux value of 3.51 mg m⁻² d⁻¹ in April 2009, representing approximately 12% of the TMF as compared to an annual mean value of 4% (Bai et al., 2019).

4.3 Lithogenic matter flux

In few samples, the PIC content is negative as a result of POC values exceeding the TC values (PIC = TC - POC). These numbers reflect the very low PIC content of these samples (<1%), and uncertainties on TC and POC measurements. Thus the CaCO₃ contents based on these PIC values became negative. In general, CaCO₃ is less than 10% of the total mass and has thus no significant impact on the lithogenic matter content. Accordingly, we revised the negative carbonate values to zero. The derived lithogenic matter fluxes reach 225 mg m⁻² d⁻¹ and ~600 mg m⁻² d⁻¹ in summer of 2008 and 2009, respectively, while low values (<50 mg m⁻² d⁻¹) are recorded in winter (Fig. 2e). On average, the lithogenic matter accounts for >70% of the total mass with no clear seasonal pattern (Fig. 2e).

4.4 Terrigenous sterol flux

Fluxes of 24-ethylcholest-5-en-3β-ol show similar variations as terrigenous campesterol suggesting a primary terrestrial source of this sterol, thus most likely mainly represented by β-sitosterol (Fig. 2g). Campesterol and β-sitosterol fluxes reach ~1 and 2.8 μg m⁻² d⁻¹ in August 2008, respectively. Both decrease to less than 0.5 μg m⁻² d⁻¹ in the following winter (Fig. 2g) and increase to highest fluxes in summer 2009 peaking at 9.6 μg m⁻² d⁻¹ in July for β-sitosterol, and to 8.4 μg m⁻² d⁻¹ in September for campesterol.

4.5 Diatom flux and species composition

Total diatom fluxes are strongly seasonal (Fig. 2h). Two peak values occur early August 2008 (~10×10⁶ valves m⁻² d⁻¹) and late July to September 2009 (>14×10⁶ valves m⁻² d⁻¹), respectively. The diatom fluxes in winter are two orders of magnitude lower (<0.5×10⁶ valves m⁻² d⁻¹). They rise abruptly late July 2019 when solar irradiance is nearly at maximum and sea ice starts retreating (Fig. 2a, b, h).

In total 93 diatom species or species groups were identified in the sediment trap samples. Apart from *Chaetoceros* RS, diatoms were categorized into 4 groups based on their ecological preferences (Medlin and Priddle, 1990; Hasle and Syvertsen, 1997; von Quillfeldt, 1997, 2000; Kohly, 1998; von Quillfeldt et al., 2003; Onodera et al., 2015). The sea ice group mainly

consists of *Fossula arctica*, *Fragilariopsis cylindrus*, *F. oceanica*, *Nitzschia frigida*, *Pauliella taeniata* and certain species of genera *Diploneis*, *Navicula*, *Nitzschia*, *Pinnularia*. The cold-water diatom group is mostly composed of *Thalassiosira antarctica* var. *borealis* RS, *T. nordenskiöldii* and, in minor amounts, *Bacterosira bathyomphala* RS and *T. bulbosa*. The cosmopolitan diatom group includes *Chaetoceros atlanticus*, *Thalassionema nitzschioides* and *Thalassiosira eccentrica*, while *Paralia sulcata* and species of genera *Cocconeis*, *Cyclotella*, *Cymbella*, *Delphineis* and *Eunotia* form the coastal diatom group. The remaining diatom species make up the other group.

Chaetoceros RS is the predominant group over the entire observation period (Fig. 3a). It increases from 40% in August 2008 to ~80% in late September 2009. In contrast, the sea ice diatom group shows a broad decline since the deployment of the trap (Fig. 3b-e). Within this group, the relative abundances of *F. arctica*, *F. cylindrus* and *F. oceanica* are two- to three-fold higher in summer 2008 than in 2009, whereas *P. taeniata* peaks in November 2008. The cold-water diatom group, however, remains low throughout the study period (Fig. 3f-g). Unlike the sea ice diatom species, the cold-water *T. antarctica* var. *borealis* RS, *T. bulbosa* and *T. nordenskiöldii* are more abundant in summer 2009 than in summer 2008. Interestingly, *T. nitzschioides*, the major representative of the cosmopolitan diatom group, is more frequently observed in winter than in summer (Fig. 3h). Similarly, the coastal diatom group, dominated by *P. sulcata*, is also found in low relative abundances with slightly higher values from October 2008 to April 2009 when sea ice prevails (Fig. 3i).

Among the sea ice diatoms, *Haslea crucigeroides* and *Pleurosigma stuxbergii* var. *rhomboides*, that are known producers of sea ice biomarker C₂₅ monounsaturated hydrocarbon IP₂₅ (Brown et al., 2014), are detected though in low relative abundance (<1%), in both August 2008 after sea ice just retreated, and in July - early August 2009 at the onset of sea ice thaw (not shown). Although widely observed in the Arctic Ocean, the sea ice bottom tethered diatom *Melosira arctica* is only found sporadically over this one-year experiment (Abelmann, 1992; Zernova et al., 2000; von Quillfeldt et al., 2003; Boetius et al., 2013; Lalande et al., 2014, 2019).

The North Pacific predominant and endemic taxa *Neodenticula seminae* (Jousé et al., 1971; Sancetta, 1982; Onodera and Takahashi, 2009; Ren et al., 2014), which has been recently reported in the North Atlantic water column and surface sediment (Starr et al., 2002; Reid et al., 2007; Miettinen et al., 2013), is also found once in sinking particles at DM, in line with the data obtained at the nearby sediment trap NAP site (Onodera et al., 2015).

4.6 Diatom taxonomic diversity

The diatom diversity index, H-index, decreases from a mean of ~2.6 in 2008 to ~2.3 in 2009 with an outlier value of ~3.9 in early July 2009 (Fig. 3j). The summer 2009 diversity minimum

is reached at maximum TMF and diatom fluxes (Fig. 2e, h, 3j). The H-index of summer 2009 is lower than found in summer 2008 (H-index: 1.8 vs 2.7) most probably due to the high abundance of *Chaetoceros* RS and decrease of other species.

5. Discussion

5.1 Interannual variability of diatom fluxes: link to surface circulation and sea ice

Our sediment trap time-series shows high TMF and diatom flux values during the summer season. However, in summer 2009 and particularly late September, fluxes are two- to three-fold higher than measured in 2008 (Fig. 2e, h; Bai et al., 2019). Diatom fluxes in summer 2008 ($\sim 10 \times 10^6$ valves $\text{m}^{-2} \text{d}^{-1}$) are comparable to those found during the same season in the nearby station NAP at 180/260 m depth in 2011 ($\sim 11 \times 10^6$ valves $\text{m}^{-2} \text{d}^{-1}$), regardless of the mooring depths and collection years of each site (Onodera et al., 2015). Interestingly, the high diatom flux values in summer 2009 at DM (14 to 27×10^6 valves $\text{m}^{-2} \text{d}^{-1}$) are of similar magnitude as winter high flux events at NAP attributed to shelf-break eddies (Watanabe et al., 2014; Onodera et al., 2015).

Lower diatom fluxes in 2008 than 2009 may suggest that the summer bloom was not captured (Fig. 2-3) and that these low values reflect post-bloom production. Indeed, an early bloom was reported in May 2008 in the Amundsen Gulf, East of the Beaufort Sea (Brown et al., 2011; Belt et al., 2013) although this does not imply that the Northwind Ridge experienced similar production growth. By contrast, a deep sediment trap in the southwestern Canadian Basin indicates high POC values in August and September 2008, which based on a sinking velocity of 85 m d^{-1} , corresponds to high surface production and export in July - August (Station A in Fig. 1; Hwang et al., 2015; Onodera et al., 2015). Furthermore, ice-ocean-ecosystem coupled model simulations point out peaking chlorophyll-*a* values in the Chukchi Sea in August 2008 (Wang et al., 2013). Therefore, we can reasonably infer that the diatom flux measured in August 2008 represents the blooming season locally and that it was thus lower than in 2009. This assumption is further supported by MODIS summer sea surface chlorophyll-*a* concentrations of the area indicating $\sim 36\%$ higher values in 2009 ($0.382 \mu\text{g l}^{-1}$) than 2008 ($0.282 \mu\text{g l}^{-1}$).

Biogenic production at DM is influenced by the PWI, the oligotrophic waters of the BG and the sea ice distribution during the blooming season (Fig. 1). The Arctic Oscillation (AO) plays an important role on sea ice distribution and surface ocean circulation (Rigor et al., 2002; Serreze et al., 2003; Steele et al., 2004).

Since 1997, the BG circulation is driven by an anticyclonic wind circulation resulting in enhanced freshening of the BG waters (Giles et al., 2012; Rabe et al., 2014; Proshutinsky et al., 2015; Wang et al., 2018). The subsequent stratification prevents nutrient to replenish the upper

ocean and thus stalls phytoplankton growth (McLaughlin and Carmack, 2010). The strength of the BG is linked to the Arctic Ocean Oscillation (AOO) index, which varied annually and notably shifted from a strong BG (AOO index = 2.9) in 2008 to a weaker BG in 2009 (AOO index = 1.9; Fig. 4a) (Proshutinsky et al., 1999, 2015). We can thus hypothesize that phytoplankton productivity at DM in 2008 was limited by stratified conditions maintaining oligotrophic conditions in the upper ocean due to freshening under intensified BG. Likewise, weakened and reduced BG in 2009 provided more favorable conditions for refueling surface waters in nutrients to sustain higher production. This is consistent with fresher surface waters reported near DM in summer 2008 than summer 2009 (Zhang, 2009; Kikuchi, 2009). Model results also show prevailing west-northwestward surface flow in the Chukchi Borderland in summer 2008 carrying nutrient-depleted waters from the Beaufort Sea to DM, whereas in summer 2009 the surface ocean circulation was nearly northward bringing nutrient-rich PWI waters to our site (Onodera et al., 2015). These findings highlight the key role of atmospheric conditions in driving the surface current and nutrients transport at the BG boundary and their impact on diatom flux.

The PWI is regarded as a pivotal nutrient source for the Arctic Ocean ecosystem (Walsh et al., 1997). The mean annual transport of the PWI through the Bering Strait (~ 0.8 Sv) fluctuates seasonally as well as annually (Roach et al., 1995; Woodgate et al., 2015) around values that regularly increased since 1990 (Woodgate et al., 2012; Woodgate, 2018). The PWI in summer 2009 (~ 1.7 Sv) was stronger than in 2008 (~ 1.35 Sv) (Fig. 4b). Furthermore, during its maximum transport period (April to July) this flow was ~ 0.2 Sv higher in 2009 than 2008. Based on the distance between the Bering Strait and DM and current velocities in the Chukchi Sea (Weingartner et al., 2005; Woodgate et al., 2005b), we can estimate that ~ 4 months is needed for the PWI to reach our mooring site, which is in agreement with the time lag between the PWI peak and high biogenic production in summer 2009 (Fig. 4). It is thus very reasonable to conclude that stronger PWI by entraining more nutrients into the Arctic Ocean likely triggered enhanced phytoplankton production in 2009.

The sea ice distribution during summer also shows different patterns in 2008 and 2009 (Fig. 5). As expected from AO and subsequent ocean circulation, in 2008 sea ice started retreating westwards from the East, favoring the inflow of freshened and oligotrophic BG waters. Meanwhile, the presence of sea ice in the South likely prevented nutrient-rich PWI waters from advecting to our site. In contrast, sea ice melting in 2009 began from the South allowing the PWI to flow to DM and influence primary production. This is further supported by surface wind fields shown in Fig. S1. Easterly winds in July 2008 favored transport of freshened and oligotrophic waters from the BG towards to the northwest or North of DM thereby limiting biogenic production. In August 2008, southward winds further contributed to counteract the

northward advection of PWI, whereas in summer 2009 (July and August) Northerly winds enhanced transport of nutrient-rich PWI to our mooring site, favoring diatom blooming (Fig. S1)

In summary, our results provide consistent evidences on the impact of contrasting atmospheric and oceanic circulation, notably the BG circulation, PWI nutrient transport as well as sea ice retreat pattern on diatoms fluxes in the Chukchi Sea, between 2008 and 2009 (Fig. 5). They also underline that the southern Northwind Ridge is an area of strong interactions between water masses from the Canadian Basin, the Pacific and even coastal Siberia, where seasonal sea ice drift and motion makes it a sensitive region to ongoing climate change.

5.2 Lateral transport of diatoms

Coastal diatoms, mostly dwelling in the continental shelf area, are indicative of lateral advection from the nearby Chukchi Sea shelf waters. Previous studies have shown that while *Paralia sulcata* prevails in the sediments of the Chukchi Sea Shelf, this species is rarely found in the Chukchi Borderland (Ran et al., 2013). At DM, the coastal diatom fluxes are also dominated by *P. sulcata* peaking in September 2009, pointing out a significant influence of water masses from the shelf area under stronger PWI conditions (Fig. 6b; See 5.1; Woodgate, 2018). Apart from this episode, values were low and stable (Fig. 6b).

Chaetoceros RS is also considered as an indicator of allochthonous material (Onodera et al., 2015). In our dataset *Chaetoceros* RS displays similar seasonal pattern as terrigenous biomarker campesterol and β -sitosterol (Fig. 6c-d). The flux of these biomarkers produced by higher plants (Huang and Meinschein, 1976) were found to also parallel lithogenic matter fluxes in both shallow and deep sediment traps of the central Arctic Ocean (Fahl and Nöthig, 2007). The co-eval temporal evolution of terrigenous sterols and lithogenic material at DM is consistent with lateral advection from the shelf in summer of 2009 (Fig. 6d-e). In contrast, the lithogenic matter content (in %) drops to its lowest values in summer 2009 witnessing a reduced contribution of allochthonous material (Fig. 6b-d, f). This discrepancy cannot be explained by sea ice transport of coastal diatoms and biomarkers in winter and later settling under ice-free conditions in September 2009 (Fig. 6a). Besides, this mechanism cannot account either for the concomitant decrease of lithogenic matter content in summer 2009. Another explanation is that coastal diatoms and terrigenous material might be entrained by PWI. Enhanced PWI would thus be responsible for higher phytoplankton production in summer 2009 and the advection of terrigenous matter (See 5.1). This is further supported by the decrease of the C/N ratio (bulk organic carbon over bulk organic nitrogen mole ratio; modified from Bai et al, 2019) to 6.4 in summer 2009. This value, close to the Redfield ratio of 6.6, points out a major marine origin of settling material (Fig. 6g). Boosted blooming production and export of organic matter likely

explain enhanced contribution of marine constituents to the overall flux and the subsequent decline of lithogenic content (Fig. 6f).

The abundance and flux of lithogenic matter, mostly composed of silt and clay, are higher at our site than at nearby mooring station NAP and LOMO2 (Fig. 1) in the central Arctic, despite a difference in water depths. Lithogenic particles might be derived from a proximate branch of the Northwind Ridge, lying at ~900 m water depth (Poore et al., 1994), close to the depth of our sediment trap (~870 m). Benthic nepheloid layer developing hundreds meters above the seafloor is another possible mechanism able to remobilize sediments from the Northwind Ridge and transport them to the sediment trap location (Rutgers van der Loeff et al., 2002).

5.3 Sea ice diatoms and IP_{25}

Fluxes of sea ice diatoms and IP_{25} at DM display quite similar seasonal patterns (Fig. 7a). Since IP_{25} is produced by specific sea ice diatoms (Brown et al., 2014), some correlation is expected between the two fluxes ($n = 21$; $r^2 = 0.64$; $p < 0.01$; Fig. 7b; Table 2). This correlation is statistically unchanged after discarding low flux values ($IP_{25} < 0.5 \text{ ng m}^{-2}\text{d}^{-1}$; sea ice diatom $< 1 \times 10^5 \text{ valves m}^{-2} \text{ d}^{-1}$) ($n = 13$; $r^2 = 0.61$; $p < 0.01$). Both fluxes are high during the sea ice melting season for both years. Although sea ice reconstructions based on micropaleontological fossils or biomarkers preserved in the sediment cores have been successfully carried out in the Atlantic and Arctic Ocean (e.g. diatoms: Sha et al., 2016, 2017; IP_{25} : Massé et al., 2008; Müller et al., 2009; Vare et al., 2009; Müller and Stein, 2014; Belt et al., 2015; Xiao et al., 2015; Cabedo-Sanz et al., 2016; Hörner et al., 2016, 2017; Stein et al., 2016, 2017; Kolling et al., 2017; Clotten et al., 2018; Kremer et al., 2018), they have rarely been produced and compared in the same core. Few available studies show poor agreement between sea ice diatoms and IP_{25} , a result that might be attributed to the specificity of IP_{25} producers as opposed to the broader habitat of sea ice diatoms (Weckström et al., 2013; Sha et al., 2015). Our sediment trap record, however, indicates consistent and synchronous fluctuations of the sea ice diatom group and IP_{25} fluxes with respect to sea ice distribution (Fig. 7a). Departures of diatom versus biomarker signals in sediments can, to some extent, be attributed to dissolution and degradation (Belt and Müller, 2013). Previous studies in the Arctic have shown that IP_{25} is biosynthesized by sea ice diatoms *Haslea crucigeroides* and *Pleurosigma stuxbergii* var. *rhomboides* (Belt et al., 2013; Brown et al., 2014). These species are present in our sediment trap in summer mainly during high IP_{25} fluxes (Fig. 7a). Although these species have been frequently reported at the bottom sea ice and, to a lesser extent, in the water column in the Arctic (e.g. von Quillfeldt, 1997; von Quillfeldt et al., 2003; Katsuki et al., 2009), they were identified in only a few summer samples at ~870 m, possibly because of dissolution during settling. These species have seldomly been found in Arctic sediments (e.g. Ran et al., 2013; Obrezkova et al., 2014; Astakhov et al., 2015;

Limoges et al., 2018). The non-IP₂₅ producing sea ice diatoms, e.g. *Fossula arctica*, *Fragilariopsis cylindrus*, *F. oceanica*, also suffer from dissolution. Sediment trap data from the Nordic Seas have shown that weakly silicified diatoms strongly dissolve in the sediments while the robust ones are preserved (Kohly, 1998). Selective degradation of diatom frustules was also demonstrated by comparing diatoms from shallow and deep traps at the nearby mooring site NAP (Onodera et al., 2015). The relative abundance of *Rhizosolenia* spp., for instance, decreased remarkably in the deep trap, whereas some diatom taxa increased in the abyssal material (Onodera et al., 2015). In another study in the Chukchi Sea, the diatom assemblages in sea ice and underlying sediments showed significant differences (von Quillfeldt et al., 2003). Despite the fact that the diatom composition and species diversity of sediment trap at DM are comparable to those of the sediments in the northern Chukchi Sea, weakly silicified diatoms, *Chaetoceros* RS and sea ice diatoms among others, became less abundant in the sediments while diatoms with stronger frustules such as *Thalassiosira antarctica* var. *borealis* RS increased (Table 1; Ran et al., 2013). This finding supports the idea of selective diatom dissolution during export to deep sediments, which pleads for cautious interpretation of paleo-reconstructions. In contrast, given its low photo-reactivity and wide distribution in the Arctic sediments, IP₂₅ should introduce less bias due to diagenesis (Brown and Belt, 2012; Belt and Müller, 2013). Although a possible biogeochemical degradation of IP₂₅ was suggested in the central Arctic based on shallow and deep sediment traps, the IP₂₅ seasonal patterns remained similar from 150 to 1550 m water depth (Fahl and Stein, 2012). Nevertheless, investigations are needed to explore IP₂₅ degradation and its possible impact on past sea ice reconstruction.

Major phytosterols, namely brassicasterol and dinosterol, show a high correlation with cold-water diatoms ($r^2 = 0.83$ and 0.77 , respectively; $p < 0.01$; Table 2). From this result we can infer that both phytosterol producers and cold-water diatoms thrive in ice-free conditions (Bai et al., 2019). Cold-water diatoms, however, have been also reported underneath consolidated sea ice (Arrigo et al., 2012, 2014). In this case, it is expected that both cold-water and sea ice diatoms sink synchronously and result in a significant correlation between their respective fluxes ($r^2 = 0.64$; $p < 0.01$; Table 2). Interestingly, the tri-unsaturated highly branched isoprenoid alkene (HBI-III), presumably associated with marginal ice zone (Belt et al., 2015; Smik et al., 2016; Smik and Belt, 2017; Bai et al., 2019), shows a weak correlation with the sea ice diatom flux ($r^2 = 0.48$; $p < 0.01$; Table 2). Although HBI-III producers have not yet been fully recognized, the hypothesized producers are within diatom genera *Pleurosigma* and *Rhizosolenia* (Belt, 2018; Belt et al., 2000, 2015; Rowland et al., 2001). In this area diatom species belonging to *Pleurosigma* are mainly related to sea ice while *Rhizosolenia* is found in the ice-free waters (von Quillfeldt et al., 2003; Onodera et al., 2015). The weak correlation between HBI-III and sea ice diatoms rather hints a sea ice source or sea ice edge conditions, but more investigation

is needed to be more conclusive.

Diatom sinking velocity of $>85 \text{ m d}^{-1}$ in summer has been recently estimated based on the time lag between the shallow and deep trap diatom fluxes at NAP (Onodera et al., 2015). Using this value, the summer diatom flux at $\sim 870 \text{ m}$ water depth at our site should represent the export of material from surface waters or the sea ice produced about 10 days earlier. Hence, the diatom bloom of summer 2009 might have started early July when the sea ice was about to retreat but still compact (Fig. 2, 5). Incidentally, massive under sea ice phytoplankton blooms have been observed in the Beaufort Sea in summer 2008 (Mundy et al., 2009) and in the Chukchi Sea in summer 2011, maximizing at $>100 \text{ km}$ into the sea ice cover boundary (Arrigo et al., 2012, 2014). This phenomenon was attributable to intensified light penetration through sea ice into the water column due to more melt ponds (Arrigo et al., 2014; Mundy et al., 2014). Thawing snow over sea ice as well as leads in the ice pack might also have played a role in promoting under ice bloom (Ambrose et al., 2005; Róžańska et al., 2009; Arrigo, 2017; Assmy et al., 2017; Lalande et al., 2019). Although no melt ponds data exist at DM for summer 2009, on-site model estimates indicate that the snow cover had almost vanished and the sea ice thickness decreased from $\sim 2.8 \text{ m}$ to $\sim 1.5 \text{ m}$, at that time of maximum solar irradiance (Fig. 2a-d), leading to similar sea surface state in summer 2009 as in 2011, allowing light penetration to trigger under sea ice bloom.

6. Conclusions

The one-year long sediment trap time-series (August 2008 - September 2009, $\sim 870 \text{ m}$ water depths) from the southern Northwind Ridge in the western Arctic Ocean highlights the strong seasonality of diatom abundances and fluxes. *Chaetoceros* RS and, to a lesser extent, the sea ice diatoms, e.g. *Fossula arctica*, *Fragilariopsis cylindrus*, *F. oceanica*, dominate the diatom composition.

Diatom fluxes are two- to three-fold higher in summer 2009 than in summer 2008, a result that is explained by a weaker BG circulation, an intensified PWI bringing nutrients into the Arctic Ocean and the earlier retreat of sea ice from the South in 2009. Our findings underline the sensitivity of the southern Northwind Ridge export fluxes to the spatio-temporal pattern of sea ice melting and upper ocean circulation regime.

High fluxes of coastal diatoms, terrigenous biomarkers and lithogenic matter show indicate lateral advection in summer 2009 likely triggered by enhanced PWI rather than by drifting sea ice.

Sea ice diatom fluxes are coherent with IP_{25} fluxes, both showing high values during the sea ice melting period reflecting marginal ice zone conditions. However, sea ice diatoms might be prone to selective dissolution in the water column that is likely to alter diatom assemblage

530 in sediments and introduce bias in sea ice reconstructions. This result reinforces the need for
531 multi-proxy approaches to assess proxy biases and improve past sea ice estimates.

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548 **Author contributions**

549 J. R. designed the study and wrote the manuscript with contribution of M.-A. S., J. C., H.
550 L., J. Z., Y. B., Y. Z and B. W. J. R. carried out the diatom identification. Y. B. and Y. L.
551 contributed the biomarker analyses (campesterol and β -sitosterol), J. Z. performed the
552 determination of particulate inorganic carbon (PIC) and calculated the CaCO₃ content. Z. J.
553 and H. J. contributed the particulate organic carbon (POC) and B. W. the opal flux. Z. Y. and L.
554 L. retrieved the environmental data from different database. All authors contributed to the final
555 version of the manuscript.

556 **Declarations of interest:** None

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Diatom composition and fluxes over the Northwind Ridge, western Arctic Ocean: impact of marine surface circulation and sea ice distribution

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Figures captions

Figure 1. a) Location of Station DM (red circle) and surface currents in the study area (grey arrows). The white dash line represents the multi-year sea ice minimum for 1979-2017 (20% of sea ice concentration) (from Cavalieri et al., 1996). Two other sites for discussion are also presented (yellow circles). Currents system: PWI- Pacific Water Inflow; ACW-Alaska Coastal Water; AW-Anadyr Water; BSW-Bering Slope Water; BG-Beaufort Gyre; SCC-Siberia Coastal Current. Topography: NR-Northwind Ridge; NAP-Northwind Abyssal Plain; CR-Chukchi Rise; CAP-Chukchi Abyssal Plain; MR-Mendeleev Ridge. b) Location of the sediment traps discussed in the text. The red dot shows Station DM (this study), the yellow dots show the other sites from previous studies.

Figure 2. Time series of daily environment variables, diatoms and biogeochemical tracers measured in the sediment trap moored at Station DM from August 2008 to September 2009. a) Shortwave radiation (W m^{-2}); b) Sea ice concentration (%); c) Sea ice thickness (m); d) Snow thickness (cm). a-d) are derived from the National Centers for Environmental Prediction (NCEP)/Climate Forecast System Reanalysis (CFSR) 6-hourly dataset (Saha et al., 2010, 2014). e) Total mass flux in $\text{mg m}^{-2} \text{d}^{-1}$ (dark bar; from Bai et al., 2019), lithogenic matter flux in $\text{mg m}^{-2} \text{d}^{-1}$ (grey bar) and percent weight of lithogenic matter (brown line); f) Particulate organic carbon (POC) flux in $\text{mg m}^{-2} \text{d}^{-1}$ (Bai et al., 2019); g) Campesterol and β -sitosterol flux in $\mu\text{g m}^{-2} \text{d}^{-1}$ represented by orange and brown bars, respectively; h) Stacked flux of different diatom groups in $10^6 \text{ valves m}^{-2} \text{d}^{-1}$ (see text). Note that the fluxes in e) and g) are not stacked.

Figure 3. a-i) Diatom fluxes (in $10^5 \text{ valves m}^{-2} \text{d}^{-1}$) and relative abundances (in %) of the main species and species groups (see text) and j) diatom species diversity (H-index). Main species of *Chaetoceros* RS group in green (a), sea ice group in blue (b-e), cold-water group in purple (f-g), cosmopolitan group in red (h) and coastal group in yellow (i).

Figure 4. Time series of environment data and fluxes measured at Station DM. a) Arctic Ocean Oscillation index (Proshutinsky et al., 2015) which indicates the strength of Beaufort Gyre; b) Transport in Sv of the Pacific water inflow (Woodgate et al., 2015; Woodgate 2018); c) Sea ice concentration in % (Saha et al., 2010, 2014); d) Total diatom flux in $10^6 \text{ valves m}^{-2} \text{d}^{-1}$; e) Total mass flux in $\text{mg m}^{-2} \text{d}^{-1}$ (Bai et al., 2019).

Figure 5. Representation of the surface currents and sea ice conditions in the study area in July and August of 2008 and 2009. PWI: Pacific water inflow, BG: Beaufort Gyre. Blocked currents by sea ice are indicated by the dashed lines. Sea ice data were retrieved from Nimbus-7 SMMR and DMSP SSM/I-SSMIS Passive Microwave Data with a grid resolution of $25 \times 25 \text{ km}$ (Cavalieri et al., 1996).

Figure 6. a) Sea ice concentration (in %) and terrigenous related fluxes: b) Coastal diatom flux in $10^5 \text{ valves m}^{-2} \text{d}^{-1}$; c) *Chaetoceros* resting spore flux in $10^5 \text{ valves m}^{-2} \text{d}^{-1}$; d) Total flux of campesterol and β -sitosterol in $\mu\text{g m}^{-2} \text{d}^{-1}$; e) Lithogenic matter flux in $\text{mg m}^{-2} \text{d}^{-1}$ (grey bars) and lithogenic matter in % weight (red line). g) Bulk organic C/N ratio is also shown (blue line; Bai et al., 2019) and compared to the Redfield ratio as a reference (grey dash line).

Figure 7. a) Fluxes of sea ice diatom in 10^5 valves $\text{m}^{-2} \text{d}^{-1}$ and IP_{25} in $\text{ng m}^{-2} \text{d}^{-1}$ (relative to the internal standard). The occurrence of sea ice diatoms that produces IP_{25} are indicated by squares. b) Correlation plot between sea ice diatom flux and IP_{25} flux ($r^2 = 0.64$; $p < 0.01$).

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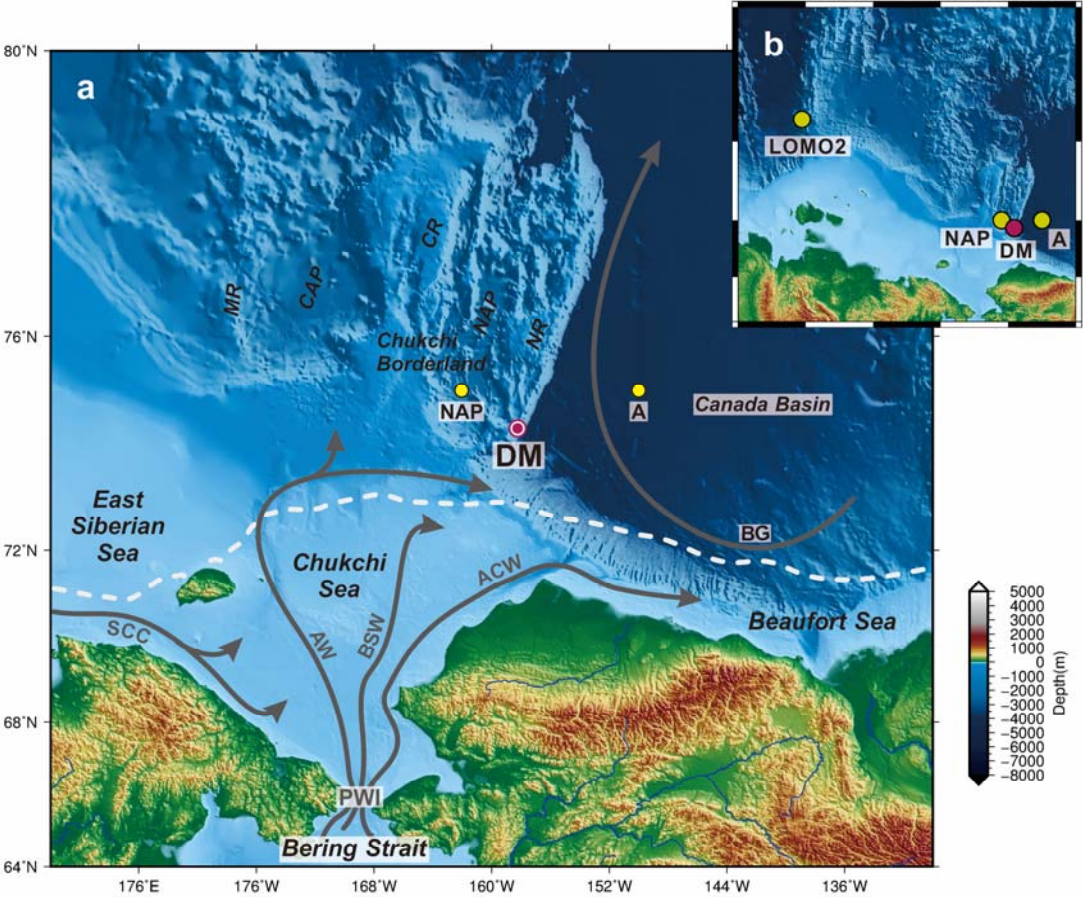
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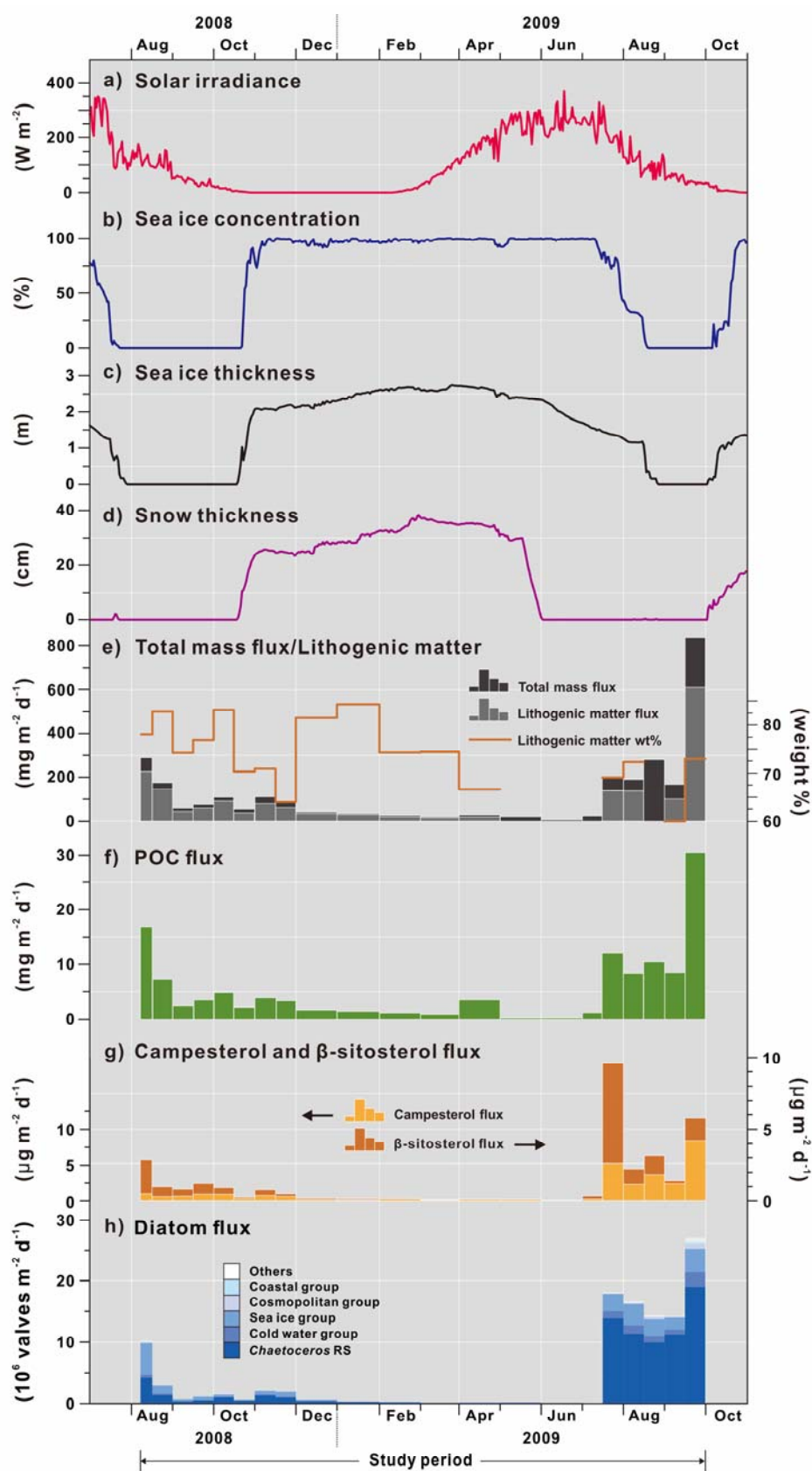
9 Figures

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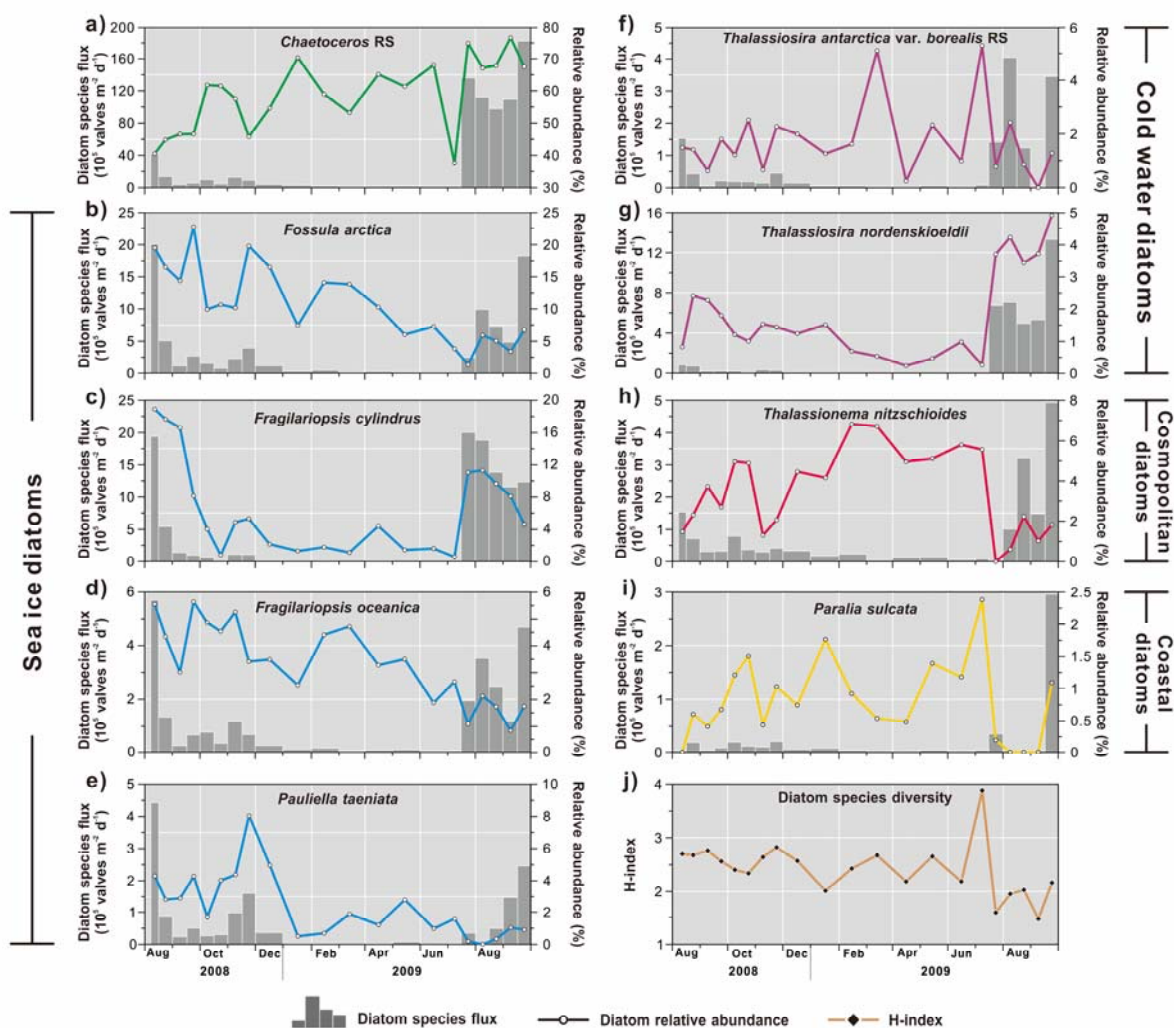
11 Figure 1-7

12 **Figure 1**

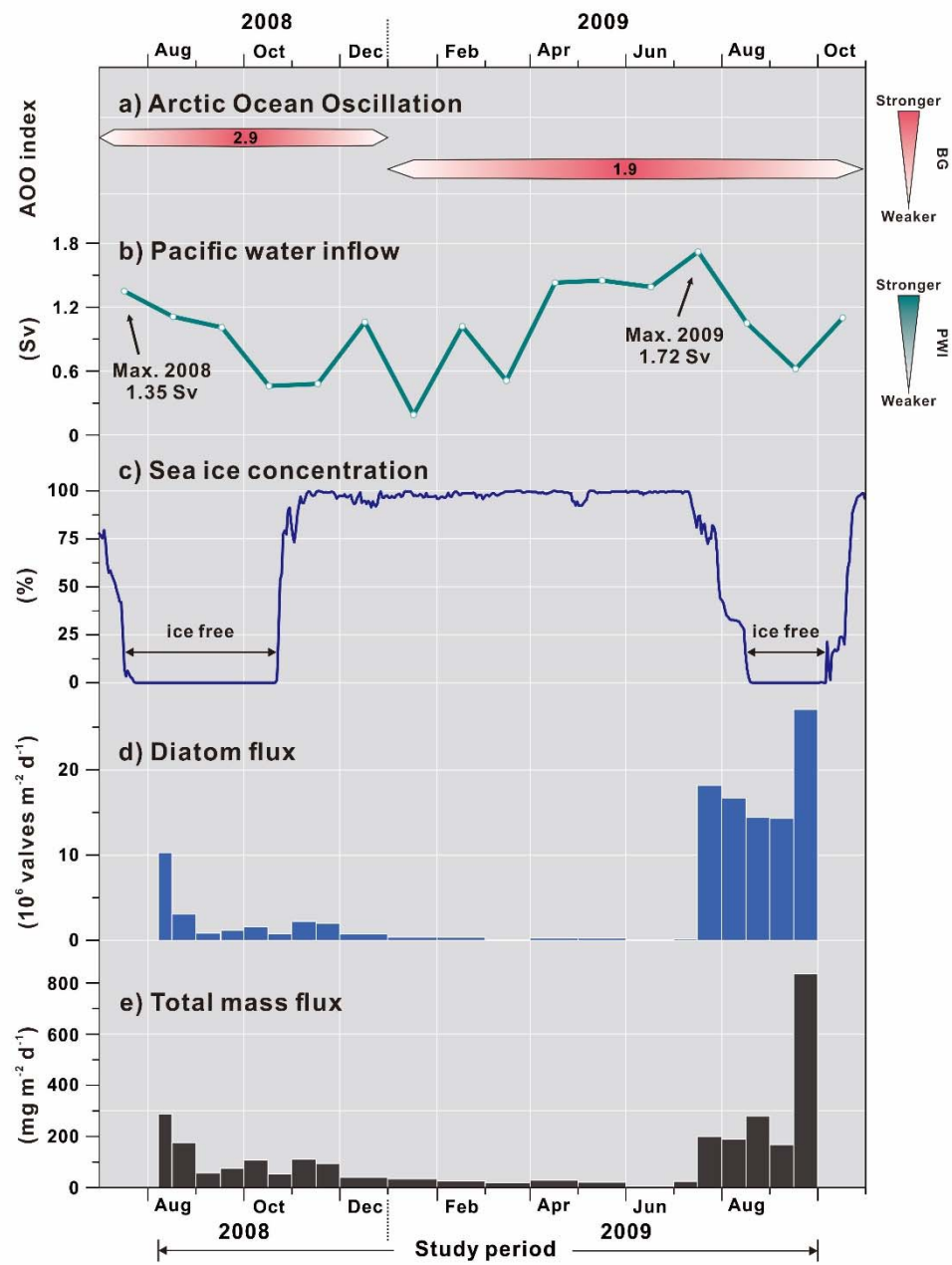




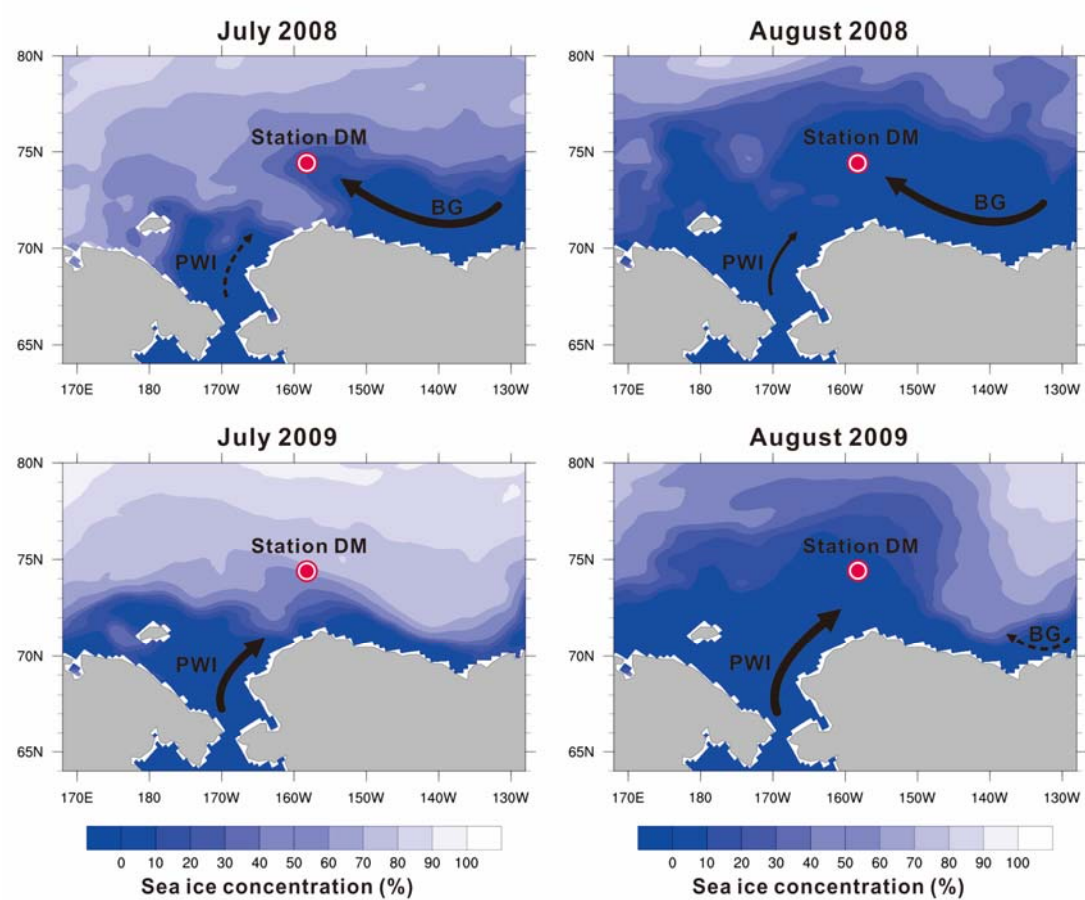
14 Figure 3



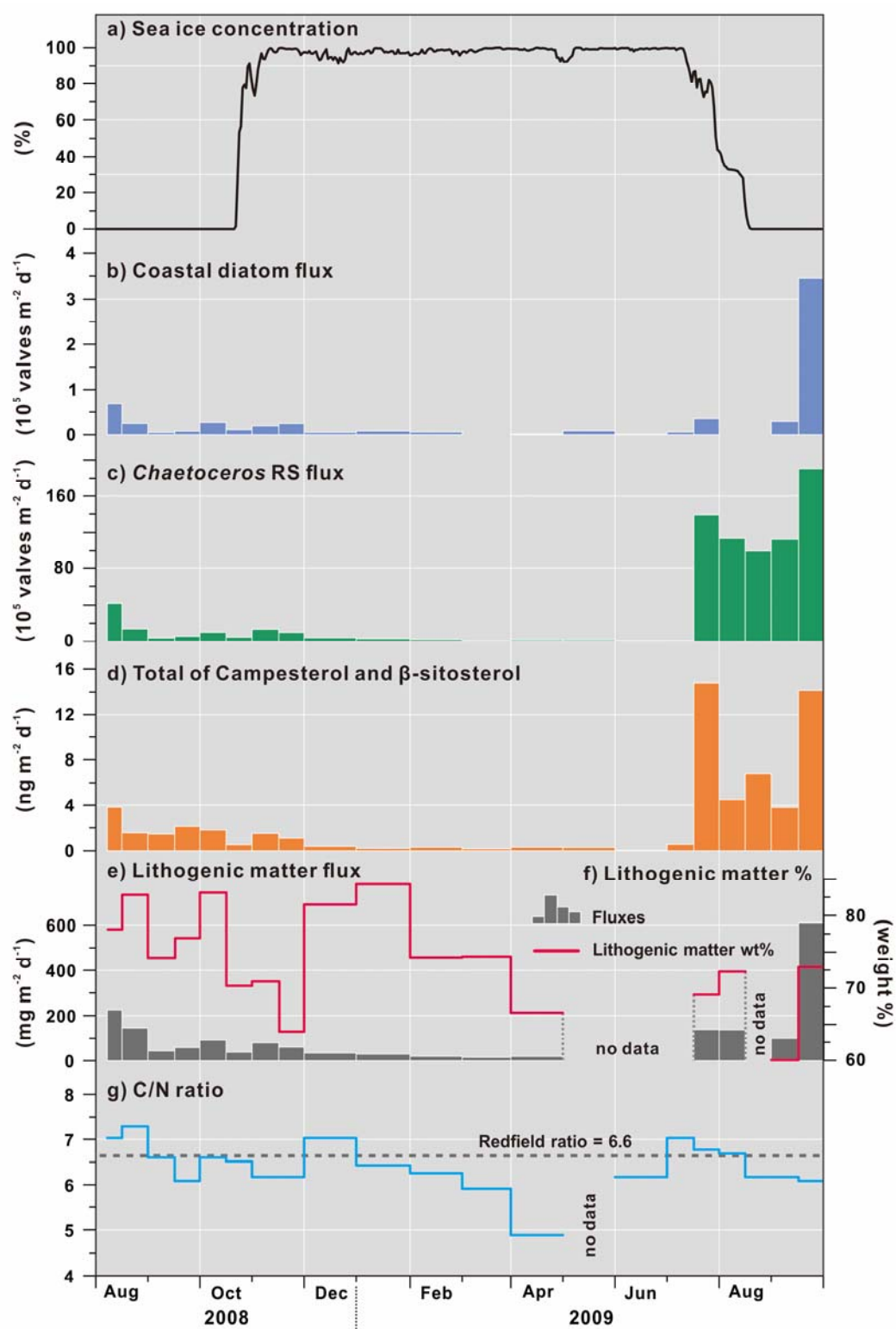
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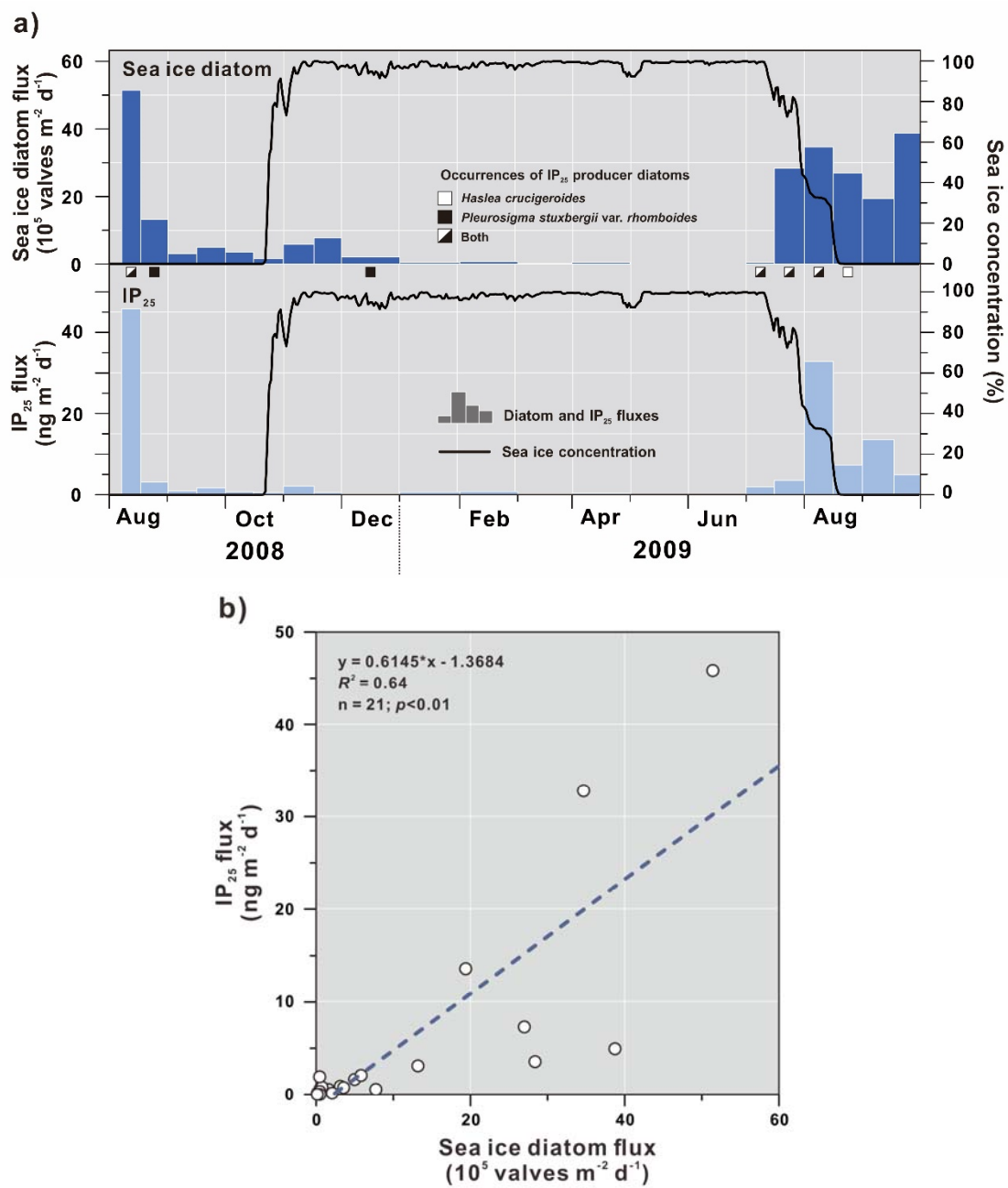
16 **Figure 5**



17 Figure 6



18 **Figure 7**



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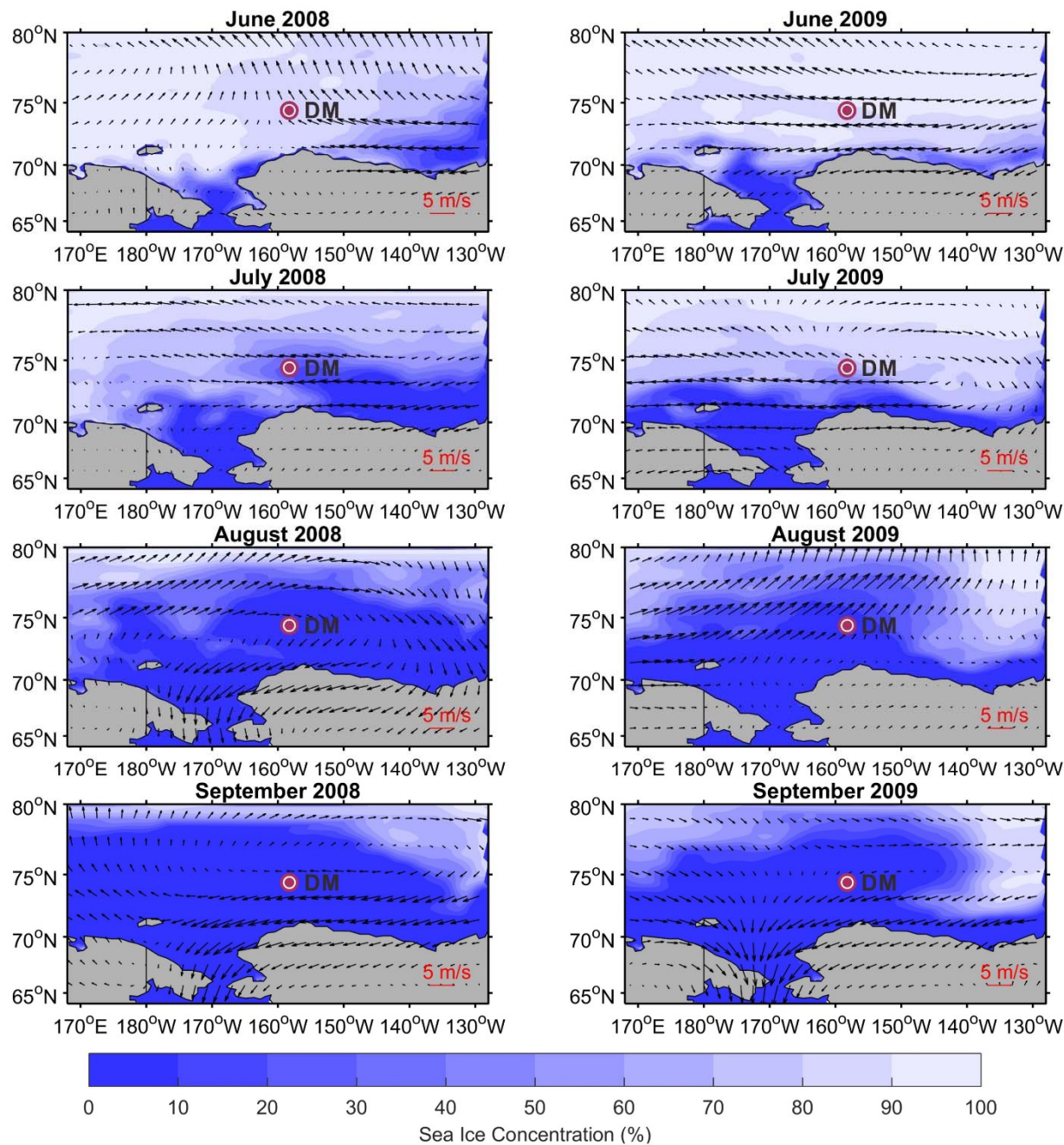
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9 Supplementary Figure

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11 Figure S1

12 **Figure S1**



13 **Figure S1.** Wind field of the study area in June, July, August and September of 2008 and 2009.
14 The wind data were retrieved from the NCEP/ NCAR atmospheric reanalysis data of daily 10
15 m-height wind speed (Kistler et al., 2001), with a horizontal resolution of 1.875°. Station DM
16 and sea ice concentration are also shown.

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9 Table 1-2

10 **Table 1**

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12 **Table 1. Comparison of the diatom diversity and relative abundances from the sediment trap sinking particles at Station DM and nearby surface**
 13 **sediments**

	Nr. of Species	H-index	<i>Chaetoceros</i> RS (%)	<i>Fossula arctica</i> (%)	<i>Fragilariopsis cylindrus</i> and <i>F. oceanica</i> (%)	<i>Thalassiosira nordenskiöldii</i> (%)	<i>Thalassiosira antarctica</i> var. <i>borealis</i> RS (%)
Station DM ^a	93	2.30	64.3 ± 11.3	8.2 ± 6.0	11.5 ± 6.0	2.9 ± 1.4	1.3 ± 1.4
Surface sediments ^b	78	2.46	63.3 ± 8.3	5.5 ± 2.9	11.3 ± 7.6	1.9 ± 1.1	3.3 ± 2.1

14 ^a Annual values based on merged seasonal samples

15 ^b Based on 10 surface sediments from northern Chukchi Sea (Ran et al., 2013)

Table 2

Table 2. Coefficient of determination (r^2) between fluxes and relative abundances (with and without *Chaetoceros* RS, respectively) of sea ice diatoms and cold water diatom group, fluxes of brassicasterol and dinosterol and fluxes of IP₂₅ and HBI-III from August 2008 to September 2009 at Station DM. Significant correlations are shown in bold and underlined, while weak correlations are in bold and italic.

	Sea ice diatom flux ^a	Cold-water diatom flux ^a	Sea ice diatom %	Cold-water diatom %	Sea ice diatom % (without <i>Chaetoceros</i> RS)	Cold-water diatom % (without <i>Chaetoceros</i> RS)	Brassicasterol flux ^b	Dinosterol flux ^b	IP ₂₅ Flux ^c	HBI-III flux ^c
Sea ice diatom flux ^a	1	-	-	-	-	-	-	-	-	-
Cold-water diatom flux ^a	<u>0.64</u> [*]	1	-	-	-	-	-	-	-	-
Sea ice diatom %	0.02	0.09	1	-	-	-	-	-	-	-
Cold-water diatom %	0.02	0.09	0.002	1	-	-	-	-	-	-
Sea ice diatom % (without <i>Chaetoceros</i> RS)	0.11	0.01	0.65 [*]	0.05	1	-	-	-	-	-
Cold-water diatom % (without <i>Chaetoceros</i> RS)	0.14	<u>0.54</u> [*]	0.43 [*]	0.281 ^{**}	0.21 ^{**}	1	-	-	-	-
Brassicasterol flux ^b	0.44 [*]	<u>0.83</u> [*]	0.04	0.09	0.01	0.35 [*]	1	-	-	-
Dinosterol flux ^b	<u>0.56</u> [*]	<u>0.77</u> [*]	0.06	0.05	0.002	0.45 [*]	<u>0.67</u> [*]	1	-	-
IP ₂₅ flux ^c	<u>0.64</u> [*]	0.16	0.09	<0.0001	0.17	0.002	0.03	0.08	1	-
HBI-III flux ^c	<u>0.48</u> [*]	0.03	0.24 ^{**}	0.02	0.21 ^{**}	0.06	0.01	0.02	<u>0.75</u> [*]	1

^a in 10⁵ valves m⁻² d⁻¹

^b in µg m⁻² d⁻¹

^c in ng m⁻² d⁻¹

* p -value < 0.01

** p -value < 0.05