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Abstract. A large part of the Pacific Arctic basin experiences ice-free conditions in summer as a result of sea ice cover steadily decreasing over the last decades. To evaluate the impact of sea ice retreat on the marine ecosystem, phytoplankton in situ observations were acquired over the Chukchi shelf and the Canadian basin in 2008, a year of high melting. Pigment analyses and taxonomy enumerations were used to characterise the distribution of main phytoplanktonic groups. Marked spatial variability of the phytoplankton distribution was observed in summer 2008. Comparison of eight phytoplankton functional groups and 3 size-classes (pico-, nano- and micro-phytoplankton) also showed significant differences in abundance, biomass and distribution between summer of low ice cover (2008) and heavy ice summer (1994). Environmental parameters such as freshening, stratification, light and nutrient availability are discussed as possible causes to explain the observed differences in phytoplankton community structure between 1994 and 2008.

1 Introduction

The Arctic Ocean is experiencing the fastest environmental changes, which are likely related to increasing CO₂ concentrations in the atmosphere. Both sea ice extent (10 % decrease per decade for 1979–2006; Comiso et al., 2008; Poliakov et al., 2010) and thickness (Kwok and Rothrock, 2009) have shown dramatic decreases in the recent years, reaching lowest values in 2007 and 2011 (Perovich, 2011). Simultaneously, river discharge increased (Peterson et al., 2006) leading to freshwater accumulation in the upper layer of the Arctic Ocean (Rabe et al., 2011), especially in the Canadian Basin. Environmental parameters driving phytoplankton growth and bloom occurrence such as light, stratification, temperature, freshening and nutrient availability have, thus, been modified with probable consequences on phytoplankton (Grebmeier, 2010; Wassmann et al., 2010).

Contradictory hypotheses have been proposed to explain the phytoplankton response to ice withdrawal. Increased primary production (PP) and total phytoplankton biomass in marginal Arctic seas is suggested by mathematical models (Zhang et al., 2010; Slagstad et al., 2011) and in situ data
from the Beaufort (Carmack and Chapman, 2003; Lee and Whitledge, 2005), Barents and Greenland seas (Rysgaard et al., 1999). Satellite observations also suggested increased PP in the Arctic basin as a result of deeper light penetration and a longer phytoplankton growth season (Arrigo et al., 2008; Pabi et al., 2008). Higher PP could also result from higher nutrient availability fed by wind-driven upwellings (Carmack et al., 2004; Yang et al., 2004), favouring the development of larger taxa such as diatoms (Babin et al., 2004). On the contrary, recent studies reported no PP increase in ice-free waters of the Canadian basin due to water column stratification restraining nutrient availability (Sundfjord et al., 2008; Lee et al., 2011; Joo et al., 2011). Li et al. (2009) documented a shift towards smaller sized phytoplankton. A northward displacement of sub-Arctic species such as coccolithophores in the Barents Sea (Hegseth and Sundfjord, 2008) and unprecedented recent blooms of the coccolithophorid Emiliania huxleyi have been observed and linked to changing climates regimes, including decreased mixed layer depth and increased surface temperature (Napp and Hunt, 2001). In the Ross Sea, deeper mixed layer would have favoured haptophyte Phaeocystis sp. growth, while shallower mixed layer along melting ice edges promoted diatom blooms (Arrigo et al., 1999). Finally, microbial DNA analyses pointed out a decrease of the bacterial communities during the 2007 minimum sea ice extent (Comeau et al., 2011), while ciliates became more common and stramenopiles less numerous.

Yet, the scarcity of data on Arctic phytoplankton strongly limits our understanding of the impact of ice melting on primary producers. Moreover, most of available phytoplankton observations are from shelf areas, i.e., the Barents and Chukchi Seas (Carmack and Wassmann, 2006; Grebmeier et al., 2006; Wassmann, 2006) and few of them relate to deep basins (Wassmann et al., 2010; Poulin et al., 2011). Better assessment of phytoplankton changes induced by ice melting at a regional scale is critical because of their consequences on carbon fixation and export to the deep sea (Sigman and Boyle, 2000), which ultimately affect the CO$_2$ Arctic sink (Bates et al., 2006; Anderson et al., 2010; Cai et al., 2010).

In this study, we report on the phytoplankton distribution in the Pacific Arctic and discuss their link to environmental parameters during 2008, a year of unusually low sea ice cover, compared with a high-ice year (1994).
Table 1. List of the pigments and their taxonomic significance (Wright and Jeffrey, 1997). The name of the twelve pigments used in CHEMTAX to distinguish between the eight phytoplankton classes are highlighted in bold. P: Picoplankton (< 2 µm), N: Nanoplankton (2–20 µm), M: Microplankton (> 20 µm).

<table>
<thead>
<tr>
<th>Pigments</th>
<th>Abbreviation</th>
<th>Size Classes</th>
<th>Taxonomic significance</th>
</tr>
</thead>
<tbody>
<tr>
<td>Fucoxanthin</td>
<td>Fuco</td>
<td>M</td>
<td>Diatoms, prymnesiophytes, some Dinoflagellates</td>
</tr>
<tr>
<td>Peridinin</td>
<td>Peri</td>
<td>M</td>
<td>Dinoflagellates</td>
</tr>
<tr>
<td>19′-Hexanoyloxyfucoxanthin</td>
<td>Hex</td>
<td>N</td>
<td>Prymnesiophytes</td>
</tr>
<tr>
<td>19′-Butanoyloxyfucoxanthin</td>
<td>But</td>
<td>N</td>
<td>Chrysophytes, Haptophytes</td>
</tr>
<tr>
<td>Chlorophyll c3</td>
<td>Chl c3</td>
<td>N</td>
<td>Prymnesiophytes, Chrysophytes</td>
</tr>
<tr>
<td>Alloxanthin</td>
<td>Allo</td>
<td>N</td>
<td>Cryptophytes</td>
</tr>
<tr>
<td>Prasinoxanthin</td>
<td>Pras</td>
<td>P</td>
<td>Prasinophytes</td>
</tr>
<tr>
<td>Neoxanthin</td>
<td>Neo</td>
<td>P</td>
<td>Chlorophytes, Prasinophytes</td>
</tr>
<tr>
<td>Chlorophyll b</td>
<td>Chl b</td>
<td>P</td>
<td>Chlorophytes, Prasinophytes</td>
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<tr>
<td>Violaxanthin</td>
<td>Viola</td>
<td>P</td>
<td>Chlorophytes, Prasinophytes</td>
</tr>
<tr>
<td>Lutein</td>
<td>Lut</td>
<td>P</td>
<td>Chlorophytes, Prasinophytes</td>
</tr>
<tr>
<td>Zeaxanthin</td>
<td>Zea</td>
<td>P</td>
<td>Cyanobacteria, Prochlorophytes, Chlorophytes, Chrysophytes</td>
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<tr>
<td>Divinyl Chlorophyll a</td>
<td>Dvchla</td>
<td>P</td>
<td>Prochlorophytes</td>
</tr>
<tr>
<td>Chlorophyll c2</td>
<td>Chl c2</td>
<td>All</td>
<td>Various</td>
</tr>
<tr>
<td>Diadinoxanthin</td>
<td>Diadino</td>
<td>All</td>
<td>Various</td>
</tr>
<tr>
<td>Diatoxanthin</td>
<td>Diato</td>
<td>All</td>
<td>Various</td>
</tr>
<tr>
<td>β-Carotenes</td>
<td>Car</td>
<td>All</td>
<td>Various</td>
</tr>
<tr>
<td>Chlorophyll a</td>
<td>Chl a</td>
<td>–</td>
<td>All – except Prochlorophytes</td>
</tr>
</tbody>
</table>

2 Methods

2.1 Study area and environmental parameters

The R/V CHINARE 2008 oceanographic cruise was conducted onboard the Chinese icebreaker R/V Xuelong (Ice Dragoon) in late summer 2008 from 1 August to 8 September in the Pacific Arctic Ocean from latitudes 65 to 86° N (Fig. 1a). The study area covered the shallow Chukchi shelf (< 100 m) south of 74° N and the deep Canadian basin (100 to 4000 m) north of 74° N. Regions visited over the Canadian basin included the Canada Abyssal Plain, the Alpha Ridge, the Mendeleev Abyssal Plain (MAP) and the Chukchi Borderland.

To investigate the role of the environmental conditions on the phytoplankton growth and distribution, several indexes were calculated. Daily ice concentrations (%) were derived from the Special Sensor Microwave Imager (SSM/I) satellite data (level 2 products at 12.5 km spatial resolution). Temperature and salinity were acquired with a conductivity-temperature-depth system (CTD, Sea-Bird SBE 9). The euphotic depth (in m) was provided by two methods: ocean color satellite MODIS (http://oceancolor.gsfc.nasa.gov/) and PAR attenuation calculated at different depth from in situ measurements (Zhao et al., 2010). The stratification of the upper layer (100 m) was estimated by the stratification index (SI), in kilograms per cubic metre (kg m⁻³), defined as the density difference between the surface and 100 m depth. The amount of freshwater from meltwater and river discharges accumulated in the surface layer was quantified by the surface fresh layer (SFL), defined as the thickness (in m) of the layer above the isohaline 31. The reference value $S = 31$ was chosen to account for the impact of sea ice meltwater ($S = 4$) and rivers ($S = 0$) and exclude freshening due to the Pacific waters whose minimum salinity is 31 (Woodgate et al., 2005).

At each station, nutrient concentrations were determined at four to ten depths from the surface to the bottom of the ocean and determined onboard using the Scan ++ Continuous Flow Analyser (SKALAR). Nitrate (NO₃⁻) concentrations were measured using the method described by Wood et al. (1967) following the World Ocean Circulation Experiment (WOCE) protocol (Gordon et al., 1993) for the preparation of primary standards and reagents. The accuracy of the analytical system for nitrate concentrations in water samples is ±0.1 µm. In the water column, the nitracline depth was determined by the position of the inflection point of the nitrate concentration profile. This nutrient is considered as the most limiting in Arctic waters (Tremblay et al., 2006; Tremblay and Gagnon, 2009).

2.2 Phytoplankton

The phytoplankton distribution was determined by two methods: pigments analysed by high-performance liquid chromatography (HPLC, Jeffrey and Vesk, 1997) and microscopic enumerations of individual species. Until now, cross-comparison of pigments and taxonomy has not been undertaken in Arctic waters. The confrontation of these two methods provides a robust approach to fully describe the phytoplankton distribution in the Arctic Ocean. Light microscopy allows the identification of phytoplankton communities at the
Fig. 2. Concentrations of Chlorophyll \(a\) (Chl \(a\)) and major accessory phytoplankton pigments (in mg m\(^{-3}\)) (see Table 1 for pigment acronyms) over the Chukchi Shelf in (a) surface waters and (b) in the SCM. The miscellaneous group (Misc) refers to pigments present in proportions < 2 %, i.e., Chl \(c\), Diato, Lut, DVchla. The pie charts show the mean relative proportions of the major accessory pigments.

scale of the species. Yet, this method is time-consuming and the absence of discernable features hampers the identification of small size phytoplankton. In contrast, the reproducibility and rapidity of the HPLC allow for the analysis of a large number of samples. Photosynthetic pigments allow recognition of the smaller fraction of phytoplankton thanks to distinctive suites of marker pigments (Jeffrey and Vesik, 1997). However, many pigments are shared among different algal classes (Jeffrey et al., 1999) which imply possible misinterpretation in determining phytoplankton composition.

### 2.2.1 HPLC pigments

Samples for pigment analyses were collected at 65 stations (Fig. 1a). For each station two depths were sampled, i.e., at 3 m and in the sub-surface chlorophyll maximum (SCM) assumed to be the depth of maximum biomass determined from the peak of fluorescence. About 2 L of seawater were filtered through 25 mm Whatman GF/F filters (0.7 \(\mu\)m porosity), then stored in a freezer at \(-80^\circ\)C to avoid biological degradation. HPLC analyses were performed in SOA (Second Institute of Oceanography, Hangzhou, China) following the method developed by Van Heukelem and Thomas (2001). Pigments were extracted 1 h at \(-20^\circ\)C in methanol and placed in an ultra-sonic bath to disrupt cells. An internal standard, the DL-\(\alpha\) Tocopherol acetate, was added to the extraction solvent to correct pigment concentrations from recovery. Pigments were analysed using a Waters 600E HPLC and an Eclipse C8 column (150 x 4.6 mm, 3.5 \(\mu\)m) thermostated at 60 \(^\circ\)C at a flow rate of 1 m min\(^{-1}\). Every 30 samples, a standard mixture was analysed under the same conditions to avoid from HPLC deviation. Chlorophyll \(a\) (Chl \(a\)) and 17 accessory pigments were quantified based on their retention time (Table 1). Twelve pigments (in bold in Table 1) were used to distinguish the relative contribution between eight phytoplankton classes (diatoms, dinoflagellates, chrysophytes, prymnesiophytes, cryptophytes, prasinophytes chlorophytes and cyanobacteria) using the matrix factorization program CHEMTAX (CHEMical TAXonomy) running under MATLAB™ following Mackey et al. (1996) and using a conversion matrix ratios constructed for the southern polar ocean species by Wright et al. (1996). Pigments were gathered into three size classes: micro- (> 20 \(\mu\)m), nano- (2–20 \(\mu\)m) and picophytoplankton (< 2 \(\mu\)m), based on their major taxonomic significance in term of algal divisions and classes (Wright and Jeffrey, 1997). The micro-sized phytoplankton pigments include those produced by dinoflagellates and diatoms; nano-sized phytoplankton pigments are from prymnesiophytes, chrysophytes and cryptophytes and finally the pico-sized phytoplankton pigments comprise those synthesised by cyanobacteria, prochlorophytes, chlorophytes and prasinophytes.
Fig. 3. Concentrations of Chlorophyll \( a \) (Chl \( a \)) and major accessory phytoplankton pigment concentrations (in mg m\(^{-3}\)) (Table 1 for pigment acronyms) over the Chukchi Borderland and the Canada Abyssal Plain in (a) surface waters and (b) in the SCM. The miscellaneous group (Misc) refers to pigments present in proportions < 2%, i.e., Chl \( c2 \), Diato, Lut, DVchla. The pie charts show the average relative proportions of the major accessory pigments.

2.2.2 Light microscopy identification and counts of phytoplankton

The 27 stations underlined in Fig. 1a were sampled for taxonomic enumerations at the same two depths as for pigment analysis (surface and SCM). About 100 ml of water taken from Niskin bottles were used for microscopic identification and preserved with glutaraldehyde (final concentration 1%) before filtration through Gelman GN-6 Metricel filters (0.45 µm pore size, 25 mm diameter). The filters were set up, onboard, on microscope slides with water-soluble embedding media (2-hydroxypropyl methacrylate). In the laboratory, the slides were used to identify and count phytoplankton species following the procedure of Joo et al. (2011). At least 300 cells were counted under the microscope (BX51, Olympus) with a combination of light and epifluorescence microscopy at 400 times for microplankton and at 1000 times for pico- and nanoplankton. The carbon biomass associated to each phytoplanktonic group was estimated from specific species biovolumes according to the equations of Menden-Dauer and Lessard (2000). These equations correspond to the more recent upgrade for biovolume calculations. Biolume estimates of each species were based on cell dimensions measured by light microscopy using appropriate geometric shapes according to Sun and Liu (2003). As for the pigments, the phytoplankton species identified by microscopy were associated to three size classes. Micro-sized plankton includes diatoms and dinoflagellates. Unidentified nanoplankton, prymnesiophytes, dictyochophytes, chrysophytes and cryptophytes belong to nano-sized phytoplankton. Unidentified picoplankton and prasinophytes represent the pico-sized phytoplankton.

3 Results

3.1 Ice cover and spatial distribution of phytoplankton communities

Phytoplankton distribution at stations located in the shallow Chukchi shelf is described separately from those from the oligotrophic Canadian basin, because of significantly different pigments and taxonomic assemblages. Among the Canadian basin, three areas can be distinguished according to the ice conditions: (i) the southern part of the Canadian basin (74°–77° N) characterised by ice free condition (ice
<20 %) called the “ice-free basins” (IFB, Fig. 1a), (ii) the “marginal ice zone” (MIZ) over the Chukchi Borderland and the Canada Abyssal Plain associated with a partial ice cover (20–70 %), and (iii) the northern part of the basins called “heavy-ice basins” (HIB) where the ice cover was >70 %, reaching exceptionally 90 % or more at three stations north of 84° N. For each province, the surface and SCM data were compared.

3.1.1 Accessory pigments

**Chukchi shelf.** Pigment distributions of the Chukchi shelf are not significantly different in surface and SCM waters except for the photoprotective pigment carotene (Caro), slightly higher in surface layers (Table 1 in Supplement). The average concentration of accessory pigments was three times higher at the SCM (2.59±2.28 mg m⁻³, Fig. 2b) than at the surface (0.99±0.64 mg m⁻³, Fig. 2a) and varied in parallel to Chl a concentration (r² = 0.93, Table 2 in Supplement). Pigment assemblage at both depth were largely dominated by fucoxanthin mainly produced by diatoms (Fuco > 70 %, Fig. 2a, b). Other pigments such as prasinoxanthin (Prasino), chlorophyll b (Chl b), diadinoxanthin (Diadino) and Caro accounted for 3 to 5 % of the total pigment concentrations. The peridinin (Peri), primarily synthesized by dinoflagellates, represented less than 2 % of the accessory pigments.

Lowest pigment concentrations were observed along the west coast of Alaska (St. C31, R05, C23, Fig. 2b). While Fuco still prevailed in these areas (40 %), typical pigments of small-size species, such as Pras (10 %), Chl b (10–30 %), neoxanthin (Neo ∼ 5 %) and alloxeanthin (Allo ∼ 5 %) increased at both depths. Of note, the high concentrations of 19'-hexanoyloxyfucoxanthin (19HF, 1.69 mg m⁻³) found at 40 m depth in station R17, suggesting high concentrations of prymnesiophyte (coccolithophorids or Phaeocystis sp.).

**Canadian basin.** The Canadian basin waters show statistically different pigment assemblage than in shelf waters (Fig. 3, Table 1 in Supplement). The oligotrophic surface waters (Chl a ∼ 0.14±0.08 mg m⁻³, Fig. 3a) of the Canadian basin were dominated by Fuco and Diadino, while 19HF was the third major pigment accounting for 5 % to 13 % of the pigment assemblage. Total pigment concentrations were up to 6× higher at SCM (Fig. 3b) than in surface waters, with more 19BF, 19HF, Neo, Pras but less Diadino in relative proportion (Table 1 in Supplement). The Fuco, Chl b, Prasino and 19HF prevailed and accounted for 80 % of the
Fig. 5. Phytoplankton abundance and carbon biomass derived from microscopic counts in surface (left panels) and SCM (right panels) over the Chukchi Borderland, Mendeleev Abyssal Plain (MAP) and the Canada Abyssal Plain. The four upper panels show the abundance (a–d) and carbon biomass (b–e) of the major taxa. Two bottom panels (c–f) show the abundances of the 14 dominant centric and penate diatoms.

3.1.2 Taxonomy

**Chukchi shelf.** Highest surface and SCM cell abundances were encountered in the northern part of the Chukchi shelf (71–73° N, Fig. 4a, d) partially ice covered (10 to 50%, top of Fig. 4). Unidentified nanoplanckton was dominant in abundance (52%, Fig. 4a) and biomass (60%, Fig. 4b) in the surface waters, while pennate diatoms *Fragilaria* sp. and *Fragilaritopsis* sp. and the centric diatom *Chaetoceros* spp. (Fig. 4f) dominated the biomass at the SCM (56%, Fig. 4e). In the southern shelf (67–69° N), centric diatoms *Chaetoceros* spp. and *Thalassiosira* spp. and the prymnesiophyte *Phaeocystis pouchetii* prevailed both at the surface (Fig. 4b, c) and in the SCM (Fig. 4e, f). The central shelf (69–70° N) had the lowest biomass mostly composed by the dinoflagellate *Gymnodinium* sp. (not shown) and the pennate diatom *Cylindrotheca* sp. in the SCM (Fig. 4e, f), and unidentified nanoplanckton in surface waters (Fig. 4b). Unidentified picoplankton (<2 µm) accounted for 36 and 16% of total cell abundances in surface (Fig. 4a) and SCM waters (Fig. 4d), respectively, and less than 1% of the total carbon biomass (Fig. 4b, d).

**Canadian basin.** Compared to the shelf, picoplankton abundances over the deep Canadian basin are higher by a factor two while nanoplanckton and diatoms are lower by a factor 5 and 10, respectively (Table 2). Surface and SCM phytoplankton abundances were dominated by picoplankton (55%) and nanoplanckton (40%), while diatoms amounted to only 5% of the cell number (Fig. 5a, d; Table 2). Conversely, diatoms and nanoplanckton dominated the Canadian
Fig. 6. Distribution of major groups of phytoplankton obtained by taxonomy (a, b, c, d) and pigments (e–f) in surface water (left panels) and in the SCM (right panels). Area charts show the abundance (a, b) and carbon biomass (c, d) of pico-, nano- and microplankton derived from taxonomy over the shelf, the ice-free basin (IFB), the marginal ice zone (MIZ) and heavy-ice basin (HIB). Area charts (e, f) show abundance of pico-, nano- and microplankton derived from pigment concentrations. Pie charts (a, b, c) and (d) show major phytoplankton groups derived from taxonomy counts. Pie charts (e) and (f) show the major phytoplankton groups calculated by the matrix factorization program CHEMTAX.

basin waters in term of carbon biomass while picoplankton accounted for only 5% (Fig. 5b, e; Table 2). Highest abundances and biomasses were found in the MIZ over the Chukchi Borderland (77–80° N) near the surface (5–20 m) and were dominated by diatoms type Nitzschia sp. (290 cells ml⁻¹, Fig. 5c), and to a lesser extent Fragilaropsis sp., Actinocyclus sp., and locally at greater depth by cryptophytes (St N01, Fig. 5e). In contrast, the IFB (75–77° N) largely represented by nanoplankton (90%) exhibited the lowest biomasses and abundances (∼5 mgC m⁻³, Fig. 5 and Table 2). Finally, phytoplankton biomasses over the HIB areas (> 80° N) were higher at SCM than surface waters and were dominated by nanoplankton, dinoflagellates type Gymnodinium sp. and Heterocapsa sp. between 80 and 83° N (Fig. 5b, e) and by the diatoms Minidiscus sp., Navicula sp. and Chaetoceros sp., north of 83° N (Fig. 5c, f). While the microplankton is well-identified by microscopy, the majority of the picoplankton and nanoplankton remained unidentified (> 90%). The few picoplankton cells identified were prasinophytes, type Micromonas sp., consistently with the observations published by Lovejoy et al. (2007) and for nanoplankton, the cryptophyte Cryptomonas sp. and the chrysophyte Dinobryon belgica, over the Alpha Ridge.

4 Discussion

4.1 Significance and comparison of pigments and taxonomic counts

R/V CHINARE 2008 uses two methods to characterise the phytoplankton distribution, i.e., pigments (HPLC) and taxonomy (microscopy), offering the possibility of cross-comparison and providing a useful test of the CHEMTAX algorithm for the Arctic Ocean. Pigments are useful indicators for offshore waters where more than 90% of the phytoplankton remains unidentified by microscopy. According to CHEMTAX, the picoplankton unidentified by cell counts (Fig. 6a, d) could be mainly prasinophytes (Fig. 6c, f), and unidentified nanoplankton could be chrysophytes and prymnesiophytes. Diatoms (r² = 0.93) and cryptophytes (r² = 0.61) pigments are correlated to their respective taxonomic abundances (Fig. 1a, b in the Supplement). Such correlation...
Table 2. Average abundances (cell ml\(^{-1}\)), carbon biomass (mgC m\(^{-3}\)) and relative contributions (%) derived from CHEMTAX for 4 phytoplankton groups in surface and SCM waters.

<table>
<thead>
<tr>
<th></th>
<th>Diatoms (20–200 µm)</th>
<th>Dinoflagellates (10–100 µm)</th>
<th>Nanoplankton (2–20 µm)</th>
<th>Picoplankton (&lt; 2 µm)</th>
<th>Total</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>SURFACE</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Abundance (cell ml(^{-1}))</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>SHELF (68–74)</td>
<td>362.0 (12.2)</td>
<td>0.2 (0.0)</td>
<td>1542.8 (51.9)</td>
<td>1068.6 (35.9)</td>
<td>2973.6</td>
</tr>
<tr>
<td>BASINS (68–86)</td>
<td>32.9 (4.9)</td>
<td>2.0 (0.3)</td>
<td>245.9 (36.6)</td>
<td>390.5 (58.2)</td>
<td>671.3</td>
</tr>
<tr>
<td>IFB (75–77)</td>
<td>11.3 (4.1)</td>
<td>0.0 (0.0)</td>
<td>139.4 (50.0)</td>
<td>128.0 (45.9)</td>
<td>278.7</td>
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<tr>
<td>MIZ (78–82)</td>
<td>92.2 (8.7)</td>
<td>4.6 (0.4)</td>
<td>396.8 (37.3)</td>
<td>569.1 (53.6)</td>
<td>1062.6</td>
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<tr>
<td>HIB (83–86)</td>
<td>12.3 (2.9)</td>
<td>0.2 (0.0)</td>
<td>148.8 (35.0)</td>
<td>263.4 (62.0)</td>
<td>424.6</td>
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<tr>
<td>Carbon biomass (mgC m(^{-3}))</td>
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<tr>
<td>SHELF (68–74)</td>
<td>29.4 (37.8)</td>
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<td>47.2 (60.7)</td>
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<tr>
<td>BASINS (68–86)</td>
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<td>1.3 (7.9)</td>
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<td>MIZ (78–82)</td>
<td>14.4 (45.0)</td>
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<td>2.0 (6.3)</td>
<td>32.0</td>
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<td>HIB (83–86)</td>
<td>0.8 (13.9)</td>
<td>0.7 (11.5)</td>
<td>4.1 (66.8)</td>
<td>0.5 (7.6)</td>
<td>6.1</td>
</tr>
<tr>
<td>CHEMTAX (%)</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>SHELF (68–74)</td>
<td>92.6</td>
<td>0.0</td>
<td>4.6</td>
<td>2.8</td>
<td>1.19</td>
</tr>
<tr>
<td>BASINS (68–86)</td>
<td>49.9</td>
<td>0.2</td>
<td>19.0</td>
<td>30.9</td>
<td>0.14</td>
</tr>
<tr>
<td>IFB (75–77)</td>
<td>58.0</td>
<td>0.3</td>
<td>19.5</td>
<td>22.3</td>
<td>0.10</td>
</tr>
<tr>
<td>MIZ (78–82)</td>
<td>17.2</td>
<td>0.1</td>
<td>27.8</td>
<td>54.9</td>
<td>0.21</td>
</tr>
<tr>
<td>HIB (83–86)</td>
<td>99.4</td>
<td>0.2</td>
<td>0.2</td>
<td>0.3</td>
<td>0.04</td>
</tr>
<tr>
<td><strong>SCM</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Abundance (cell ml(^{-1}))</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>SHELF (68–74)</td>
<td>959.4 (40.6)</td>
<td>11.2 (0.5)</td>
<td>1009.6 (42.7)</td>
<td>385.5 (16.3)</td>
<td>2365.6</td>
</tr>
<tr>
<td>BASINS (68–86)</td>
<td>31.6 (3.2)</td>
<td>3.6 (3.7)</td>
<td>377.2 (38.7)</td>
<td>561.5 (57.6)</td>
<td>974.0</td>
</tr>
<tr>
<td>IFB (75–77)</td>
<td>0.4 (0.1)</td>
<td>0.0 (0.0)</td>
<td>160.4 (41.4)</td>
<td>226.8 (58.5)</td>
<td>387.6</td>
</tr>
<tr>
<td>MIZ (78–82)</td>
<td>22.9 (2.1)</td>
<td>3.3 (0.3)</td>
<td>556.8 (52.2)</td>
<td>484.4 (45.4)</td>
<td>1067.4</td>
</tr>
<tr>
<td>HIB (83–86)</td>
<td>72.7 (5.8)</td>
<td>6.7 (0.5)</td>
<td>240.1 (19.2)</td>
<td>929.1 (74.4)</td>
<td>1248.7</td>
</tr>
<tr>
<td>Carbon biomass (mgC m(^{-3}))</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>SHELF (68–74)</td>
<td>55.8 (55.6)</td>
<td>10.2 (10.2)</td>
<td>34.3 (34.2)</td>
<td>0.1 (0.1)</td>
<td>100.4</td>
</tr>
<tr>
<td>BASINS (68–86)</td>
<td>4.5 (21.3)</td>
<td>2.7 (12.8)</td>
<td>13.3 (63.0)</td>
<td>0.6 (2.8)</td>
<td>21.1</td>
</tr>
<tr>
<td>IFB (75–77)</td>
<td>0.1 (1.6)</td>
<td>0.0 (0.0)</td>
<td>4.7 (97.3)</td>
<td>0.0 (1.0)</td>
<td>4.9</td>
</tr>
<tr>
<td>MIZ (78–82)</td>
<td>3.7 (14.0)</td>
<td>2.9 (11.1)</td>
<td>19.7 (74.5)</td>
<td>0.1 (0.4)</td>
<td>26.4</td>
</tr>
<tr>
<td>HIB (83–86)</td>
<td>9.1 (38.5)</td>
<td>4.2 (17.6)</td>
<td>8.2 (34.5)</td>
<td>2.2 (9.3)</td>
<td>23.6</td>
</tr>
<tr>
<td>CHEMTAX (%)</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>SHELF (68–74)</td>
<td>88.1</td>
<td>0.2</td>
<td>4.9</td>
<td>6.8</td>
<td>2.16</td>
</tr>
<tr>
<td>BASINS (68–86)</td>
<td>10.8</td>
<td>0.7</td>
<td>35.1</td>
<td>53.4</td>
<td>0.50</td>
</tr>
<tr>
<td>IFB (75–77)</td>
<td>15.9</td>
<td>0.9</td>
<td>32.1</td>
<td>51.1</td>
<td>0.69</td>
</tr>
<tr>
<td>MIZ (78–82)</td>
<td>9.3</td>
<td>0.8</td>
<td>43.2</td>
<td>46.7</td>
<td>0.40</td>
</tr>
<tr>
<td>HIB (83–86)</td>
<td>2.5</td>
<td>0.0</td>
<td>23.4</td>
<td>74.1</td>
<td>0.28</td>
</tr>
</tbody>
</table>

In parenthesis are the given % abundance and % carbon biomass of the 4 phytoplanktonic groups for the different areas. Stations where taxonomic data are available were separated in Shelf and Basins. Basins are subdivided in three areas according to ice conditions: ice-free basins (IFB), marginal ice zone (MIZ) and the heavy ice basins (HIB). In the first column, the two numbers in parenthesis represent the range of latitude of each considered area.

In parenthesis does not exist for the 4 others main phytoplankton groups (picoplankton, nanoplankton, dinoflagellates and prasinophytes, Fig. 1c, d, e, f in the Supplement). CHEMTAX seems to overestimate the diatom contribution and to minimise the importance of nanoplankton as compared to taxonomic observations (Table 2). For example, in shelf surface waters dominated by Fuco, CHEMTAX concludes to a large dominance of diatoms (92.6 %, Fig. 6c) while taxonomic counts indicate dominant nanoplankton cell abundances (51.9 %, Fig. 6a) and carbon biomass (60.7 %, Fig. 6b). The systematic assignment of Fuco to diatoms by CHEMTAX, while this pigment is also a major pigment in nanoplankton like prymnesiophytes or dinoflagellates, may be responsible for this apparent discrepancy (Jeffrey et al., 1997; Rodriguez et al., 2002). In the Canadian basin and Chukchi shelf, pigments combined to cell counts suggest that Fuco is produced by nanoplankton rather than by diatoms. Similar conclusions have been achieved in the oligotrophic gyres of the
subtropical Atlantic Ocean by Hirata et al. (2008). Furthermore, Chl \(b\) associated here to pico-sized prasinophytes has been attributed by Hirata et al. (2011) to nano-sized phytoplankton. While CHEMTAX reflects relatively well the phytoplankton abundance in Antarctic waters (Wright et al., 1996; Rodriguez et al., 2002), a calibration would improve its performance in Arctic waters.

Taxonomic data indicate that picoplankton (\(< 2 \mu m\)) accounted for 25% of the cell abundance, but contributed for less than 1% of the total phytoplankton carbon biomass over the shelf (Table 2). On the contrary, diatoms with a 1000 times larger biovolume than picoplankton contribute for a large part to the total carbon biomass over the Canadian basin, despite low cell numbers. The pigment cell content increases with depth in response to adaptation to low light availability (Henriksen et al., 2002). Pigment biomasses over the Canadian basin are 5 times higher in the deep SCM (60 m) than in surface (Fig. 6c, f), despite no significant increase of cells abundance (Fig. 6a, d and Table 2) and carbon biomass (Fig. 6b, e). Pearson test (Table 2, Supplement) indicate that Chl \(a\) better correlates with carbon biomass \((r = 0.83)\) than cell abundances \((r = 0.58)\) excepted for the SCM where Chl \(a\) fit better with cell abundances than carbon biomass. No significant correlation was found between Chl \(a\) and the cell abundance or carbon biomass over the Canadian basin. Finally, low pigment concentrations in ultra-oligotrophic surface waters of the IFB and HIB \((< 0.05 \text{mg Chl } a^{-1})\) could hamper the detection of minor pigments by the HPLC. We estimated that a limit of detection for minor pigments occurs at total pigments concentration lower than 0.05 \(\text{mg m}^{-3}\).

The cross-comparison of pigments and taxonomic data highlight the different information provided by the two approaches to diagnose phytoplankton populations. Improvement of the relationship between taxonomic enumeration and pigment fingerprints would benefit from both additional field and in vitro experiments from the Arctic Ocean.

### 4.2 Comparison with previous campaigns in Arctic Ocean

Arctic main phytoplankton distribution differs widely between the marginal shelves and the deep central basins and depends strongly on sea ice conditions (Poulin et al., 2011). High biomasses and abundances observed in 2008 over the Chukchi shelf agree with the high productivity commonly observed in summer across the shelf (Hameedi, 1978; Cota et al., 1996; Sakshaug, 2004; Hill and Cota, 2005). The high contribution of large cells, such as diatoms *Fragilaria* sp., *Cylindrotheca* sp. and dinoflagellates during the low sea ice summer 2008 are comparable to those of more icy years in the marginal Barents and Chukchi seas dominated by the centric diatoms *Chaetoceros furcellatus*, *Thalassiosira* sp., the pennate diatoms *Cylindrotheca closterium*, *Fragilariopsis oceanica* and dinoflagellates (*Hill et al., 2005; Sukhanova et al., 2009; Poulin et al., 2011*). Dominance of nanoplanckton in the surface shelf waters in 2008, likely reflect post-bloom conditions as also reported by Hodal and Kristiansen (2008) in the Barents Sea.

In contrast to the shelf, phytoplankton assemblages in summer 2008 in the oligotrophic Canadian basin point to high abundances of prasinophytes such as *Micromonas* sp. Previous phytoplankton observations during more icy years highlighted abundant prasinophytes better adapted to low nutrient concentrations (Lovejoy et al., 2006, 2007). Also remarkable is the dominating nanoplanckton biomass in all of the three offshore ice areas (IFB, MIZ and HIB). Another notable feature is the abundance of pelagic diatoms (*Minidiscus* sp., *Navicula* sp. and *Chaetoceros* sp.) in summer 2008 at the highest more icy years in the marginal Barents and Chukchi Sea, at higher latitudes of the Canadian basin \((> 80^\circ N)\). High diatom abundances have been previously reported close to the pole covered by ice throughout the year, mainly as sea ice-associated algae such as the centric diatoms *Melosira arctica* (Booth and Horner, 1997; Gradinger, 1999; Melnikov et al., 2002). During summer 2008 and probably because of the sea-ice shrinking close to the pole, taxonomic observations indicate that sea-ice associated diatom species could have been replaced in part by pelagic diatoms.

In icier past summers, highest production occurred in spring as “ice edge blooms” over the shelves (Sakshaug and Skodjal, 1989; Luchetta et al., 2000; Tremblay et al., 2006; Perrette et al., 2011) and was mainly composed of centric diatom genera *Chaetoceros* sp., *Thalassiosira* sp. and prymnesiophyte *Phaeocystis pachetii* (Wassmann et al., 1999). In 2008, high abundance and biomass of centric diatoms *Chaetoceros* and *Fragilaria* sp. and typical polar species *Phaeocystis pachetii* were found at the ice edge over the shelf. The genus *Thalassiosira* sp. was mainly observed south of the shelf, an area influenced by the Pacific inflow in 2008. High biomasses dominated by diatoms (*Nitzschia* sp. and *Fragilariopsis* sp.) and haptophytes were also observed in the marginal ice zone (MIZ) of the Canadian basin at higher latitudes \((77–80^\circ N)\) than during icier years during which the MIZ was never observed further north than the continental shelves.

Overall, in summer 2008, the phytoplankton species and main functional groups are similar to those described earlier in the Pacific Arctic but their distributions are somewhat different. Because of lack of data, the comparison is weak and limited in the Arctic Ocean, especially in the central Arctic basins. There was no previous pigment data over the Canadian Arctic basin. The only taxonomic data from this region, before the extensive melting occurring since 2007, have been produced from the Arctic Ocean Section (R/V *AOS*) cruise in July–August 1994 (Booth et al., 1997; Gosselin et al., 1997). We, thus, compared phytoplankton distributions from the Chukchi Borderland and the Mendeleev Abyssal Plain during a summer of intense ice melting 2008 (R/V *CHINARE*) and the icier summer 1994 (R/V *AOS*). The 1994 and 2008 cruises have similar ship tracks between
Fig. 7. Comparison of the taxonomic distribution of phytoplankton obtained during the R/V CHINARE 2008 (left panels) and R/V AOS 1994 (right panels) cruises. The phytoplankton abundance (a, b, g, h) and biomass (c, d, i, j) of three size classes: flagellates (< 2 µm), flagellates (> 2 µm) and diatoms and the abundances of the main seven planktonic taxa (e, f, k, l) are averaged over four latitude ranges in surface and at the SCM.

Sampling was performed west of the Mendeleev Abyssal Plain (~ 175° W) in 1994 and east of the Mendeleev Abyssal Plain during the 2008 cruise. The Chl a concentration maximum in the Arctic follows the sea ice retreat and is typically observed between May and June over the Chukchi shelf (Longhurst, 1995; Wang et al., 2005) and probably later over the Canadian basin. Both R/V AOS and R/V CHINARE cruises were carried out during post-bloom conditions.

Phytoplankton communities in 2008 show differences in terms of geographical distribution, abundance and species dominance as compared to 1994. At all stations considered, the average standing stock of diatoms and two size classes of flagellates (< 2 µm and > 2 µm) was drastically lower in 2008 (Fig. 7a, b) than in 1994 (Fig. 7g, h). Differences in cell numbers were mainly attributed to a ten times lower abundance of picoplankton in surface and sub-surface waters in 2008. While in 1994, picoplankton accounted for 96% of the total abundance (Fig. 7g, h) and 20% of the total biomass (Fig. 7i, j); it represented only 42% of total abundances (Fig. 7a, b) and 5% of the total biomass in 2008 (Fig. 7c, d). The drastic picoplankton decrease in 2008 could result from the sensitivity of this group to photo-inhibition (Finkel et al., 2010; Key et al., 2010) caused by sea ice thinning and removal of snow deposited on sea ice. Moreover, the dinophyceae Gymnodinium sp., abundant in 1994 (~30 cell ml⁻¹; Fig. 7k, l) was an order of magnitude lower in 2008 (Fig. 7e, f).

Despite reduced total abundances in 2008, regions such as the MIZ over the Canadian basin (77–80° N) showed phytoplankton biomasses (Fig. 7c, d) three times higher than in 1994 (Fig. 7l, j) due to a higher abundances of unidentified nanoplanckton, cryptophytes and diatoms in 2008 (Fig. 7e, f) than in 1994 (Fig. 7k, l). In contrast, in 2008 the IFB
(75–77° N) was lower in phytoplankton abundance, biomass and diversity than during 1994 ice-covered conditions. In 1994, phytoplankton biomass at this latitude (75–77° N) was mainly composed of diatoms (45%), large flagellates (30%) and picoplankton (25%). In contrast in 2008, the biomass was 2–3 times lower and largely dominated by unidentified nanoflagellates (> 90%) both in surface and SCM while diatoms and dinoflagellates abundances was 10 and 20 times lower than in 1994. In the northern part of the Canadian basin (83–86° N) diatom abundances were surprisingly high (~ 100 cells ml⁻¹) as well in SCM in 2008 (Fig. 7f) than in surface and SCM in 1994 (Fig. 7k, l). The nutrient sources sustaining these large taxa are still unknown and further studies on the river or multi-year sea ice nutrient budget in the central Arctic should provide some answers.

Differences between phytoplankton distributions between 1994 and 2008 could come in part from the microscopic countings. Settling chambers and inverted light microscopy (Utermöhl, 1931) were used for cell enumerations for the 1994 samples while normal light microscopy was used in 2008. Moreover, different transfer functions were used to derive carbon biomass from phytoplankton biovolumes. Equations were based on Strathmann (1967) for 1994 samples and on Mender-Deuen and Lessard (2000) for those collected in 2008. Comparison of both methods on 2008 samples revealed that picoplankton is overestimated by a factor 6 and nanoplankton underestimated by 20% with the Strathmann method. Nevertheless, the bias resulting from the selected method affects the biomass calculations, but cannot explain the difference in cell abundances between the two years.

Therefore, despite methodological biases, sea ice retreat and melting could be a determinant parameter in the different phytoplankton pattern between 1994 and 2008. Sea ice retreat over the Canadian basin could locally increased phytoplankton abundances due to the presence of an offshore MIZ. But, the total disappearance of the sea ice cover in the south of the Canadian basin likely explain the phytoplankton impoverishment and decrease in phytoplankton abundance and Chl a biomass of less large cells (diatoms and dinoflagellates) and picoplankton and higher production of nanoplankton (chrysophytes and prymnesiophytes).

4.3 Role of the environmental conditions on the 1994 and 2008 phytoplankton distributions

What environmental parameters could explain such differences in phytoplankton spatial distributions between 1994 and 2008? The more remarkable environmental change between the two years was the sea ice coverage (Fig. 8a, Table 3). In 1994, all offshore stations were sampled under thick ice cover (> 90%) while in 2008, the ice retreat above the Canadian basin reached 77° N and allow for the determination of three zones, the IFB (74–77° N), the MIZ (77–80° N) and the HIB (80–86° N). This exceptional ice melting resulted in the a surface salinity decrease by 3 units (Fig. 8c, Table 3) and a thickening of the surface fresh layer (SFL) by a factor 2 (Fig. 8d) over the Canadian basin as compared to year 1994 (Fig. 8d, Table 3). The freshening of the surface layer leads to a stronger stratification as shown by the correlation between the salinity and the stratification index (SI) (R_{Salinity, SI} = −0.87). Stronger stratification prevents the supply of deep-water nutrients while the establishment of a thick SFL deepens the nitracline (Fig. 8e; R_{SFL, Nitracline} = 0.67). The observed deepening of the nitracline agrees with previous observations from the Canadian Basin (Mc Laughlin and Carmack, 2010). The decrease of the nutrient availability in surface waters subsequent to a deeper nitracline led to lower surface Chl a concentration (R_{Nitracline, Surf Chl a} = −0.62) and a deepening of the SCM (Fig. 8f; R_{Nitracline, DepthSCM} = 0.67). Since the euphotic depth did not change much between 1994 and 2008 (Table 3), deepening of the SCM in 2008 is likely responsible for less productive phytoplankton communities due to light attenuation. The strong freshening and associated deep nitracline observed in the IFB in 2008 (Fig. 8c, d, e) was associated with lower abundances of diatoms and dinoflagellates (Fig. 7c, f) and a deeper SCM (Fig. 8f, Table 3) than in the same areas under heavy ice conditions in 1994 (Fig. 7k, l). The nanoplankton dominated the phytoplankton assemblages in the IFB suggesting a better adaptation to the low nutrients and light occurring in this region. Even if the vertical stratification resulting from freshening reduces surface nutrient availability, part of the nutrient depletion observed in the IFB might be attributed to consumption by phytoplankton.

In contrast to IFB, a weak freshening (Fig. 8c, d) and a shallow nitracline (Fig. 8e) associated with the formation of an offshore MIZ due to extensive withdrawal of sea ice in 2008, likely provided optimal conditions for diatoms and nanoplankton growth (Fig. 7c, d, e, f). These high levels of phytoplanktonic biomass in the MIZ were comparable to shelf values and could be assimilated to sea ice.
edge blooms found in marginal Arctic seas (Luchetta et al., 2000; Hill et al., 2005; Tremblay et al., 2006). However, species inhabiting these “offshore ice edge blooms” are composed of pennate diatoms* Nitzschia *sp. and *Fragilariopsis *sp., and the nanoplanckton classes dictyochoyphyes and crysophytes, thus, different from species living in sea ice edge over the shelf, usually dominated by the diatoms *Chaetoceros *spp., *Fragilaria *sp. and *Fragilariopsis *sp. and by unidentified nanoplanckton and *Phaeocystis puchetti* (Alexander and Niebauer, 1981; Sukhanova et al., 2009; Seegreva et al., 2010).

Interannual variability in the hydrography, the circulation pattern and river inputs could be other causes for 1994–2008 phytoplankton differences. Recent studies highlighted the weak influence of McKenzie River on the pool of nutrients in the Canadian basin (Emerton et al., 2008; Simpson et al., 2008), while Russian rivers (Holmes et al., 2000; Hessen et al., 2010) should impact the phytoplankton distribution in this basin. Changes in oceanic circulation and stratification of the upper water layers associated with both sea-ice and continental ice cover reduction would be key drivers of nutrient availability and phytoplankton patterns in the illuminated surface layer.

5 Conclusions

In situ data from the RV CHINARE 2008 cruise provide new pigment and taxonomy data in a poorly documented area of the deep central basin of the Pacific Arctic, after extreme sea ice melting in summer 2007. These results highlight significantly different phytoplankton distribution between an unusually low sea ice covered year 2008, and an icier year 1994 which can be summarised by lower abundances of large cells (diatoms and dinoflagellates) and picoplankton (prasinophytes) and more abundant nanoplanckton probably due to chrysophytes and prymnesiophytes increased.

Earlier studies suggested that increase light availability due to sea ice retreat should result in increased primary production and biomass. We propose that freshening, by deepening the nutricline and reinforcing the stratification, would reduce light and nutrients availability for phytoplankton growth. The strong freshening observed in 2008 over the “ice free basin” was associated with lowest biomass and likely promoted nanoflagellates while picoplankton, less adapted to higher and longer exposure to UV, declined in surface waters. In contrast, appearance of offshore marginal sea ice zone stimulates the production of pelagic diatoms. Finally, the strong freshening observed in HIB would be responsible for the deepening of phytoplankton communities. However, both in 1994 and 2008, relatively high diatom abundances were found in the ice covered highest latitude (> 83° N), but the origin of the nutrients source feeding these large taxa is still an open question.

Our in situ phytoplankton data suggest that the deep Arctic basins may evolve towards lower phytoplankton biomass and production. With the northern extension of the ice-free areas and enhanced freshening subsequent to predicted increase of sea ice melting and river discharges, impoverishment of the ice-free basins would extend northward. We can, thus, anticipate that carbon production and export might decrease in the ice-free basins and probably increase in the offshore marginal ice zone, as a consequence of changes in the phytoplankton abundance and size structure.

This study points out discrepancy between information derived from pigment analyses and microscopic counts. High fucoxanthin concentrations over the Chukchi shelf and in the surface waters of the Canadian basin suggest that besides diatoms, nanoflagellates may have contributed to the production of this pigment. Our findings also underline that pigments represent an efficient tool for the description of phytoplankton over Arctic deep basins, which are dominated by small phytoplankton poorly identified by light microscopy. Future improvement of CHEMTAX calibration for Arctic pytoplankton using in vitro experiments and in situ
observation combining HPLC and microscopic taxonomy are needed to fully exploit the pigments data.

The R/V CHINARE cruises planned every two years in the Arctic should provide additional biogeochemical data to improve our understanding of the response of phytoplankton to on-going climate changes and get a more comprehensive picture of the Arctic ecosystem evolution in relation to ice and sea ice melting and subsequent environmental changes.

Supplementary material related to this article is available online at: http://www.biogeosciences.net/9/4835/2012/bg-9-4835-2012-supplement.pdf.

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References


