Abundance, biomass and composition of spring ice algal and phytoplankton communities of the Laptev Sea (Arctic)

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Abstract Abundance, biomass and composition of the ice algal and phytoplankton communities were investigated in the southeastern Laptev Sea in spring 1999. Diatoms dominated the algal communities and pennate diatoms dominated the diatom population. 12 dominant algal species occurred within sea ice and underlying water column, including Fragilariopsis oceanica, F. cylindrus, Nitzschia frigida, N. promare, Achnanthes taeniata, Nitzschia neofrigida, Navicula pelagica, N. vanhoeffenii, N. septentrionalis, Melosira arctica, Clindrotheca closterium and Pyramimonas sp. The algal abundance of bottom 10 cm sea ice varied between 14.6 and 1562, 2×10^4 cells l^{-1} with an average of 639, 0×10^4 cells l^{-1} , and the algal biomass ranged from 7.89 to 2093.5 μg C l^{-1} with an average of 886. 9 µg C 1-1, which were generally one order of magnitude higher than those of sub-bottom ice and two orders of magnitude higher than those of underlying surface water. The integrated algal abundance and biomass of lowermost 20 cm ice column were averagely 7.7 and 12.2 times as those of upper 20 m water column, respectively, suggesting that the ice algae might play an important role in maintaining the coastal marine ecosystem before the thawing of sea ice. Ice algae influenced the phytoplankton community of the underlying water column. However, the "seeding" of ice algae for phytoplankton bloom was negligible because of the low phytoplankton biomass within the underlying water column.

Key words ice algae, phytoplankton, biomass, Laptev Sea, Arctic.

The Laptev Sea, an important shelf area of the Arctic Ocean, is significant for the fresh water balance and temperature and salinity structure of the Arctic Ocean (Steele et al. 1996; Johnson and Polyakov 2001). As an important new ice forming area, the Laptev Sea exports sea ice and influences the sea ice and planktonic community structure of the Arctic Ocean due to the existence of the Transpolar Drift, which transported the pack ice through the central Arctic Ocean to the Greenland Sea (Wollenburg 1993; Eicken et al. 1997; Dethleff et al. 2000).

The southeastern Laptev Sea is a shallow water area with the water depths of

less than 40 m in most area, and covers with sea ice between October and June or July in the next year. The water temperature near the sea bottom is below 0°C in the whole year. The fresh water influx from Lena and Yana rivers, brings abundant nutrients, organic matters and phytoplankton to this area (e. g. Sorokin and Sorokin 1996; Kattner et al. 1999; Matthiessen et al. 2000). Consequently, the algal community structure of this area was mostly dominantly controlled by the seasonal covering of sea ice and the influx of fresh water (Clark 1990).

Most researches of the spring ice algae and phytoplankton of the coastal seas were carried out in the Canadian Arctic seas (e. g. Hsiao 1980; Michel et al. 1993; Monti et al. 1996). In the Laptev Sea, several studies have been done mainly in summer and few in other seasons (e. g. Gran 1904; Kiselev 1932; Okolodkov 1992a, b; Heiskanen and Keck 1996; Soroki and Sorokin 1996; Cremer 1998). Juterzenka and Knickmeier (1999) and Tushling et al. (2000) reported the autumn chlorophyll a concentrations and micro-algal compositions within the sea ice and water column. Tushling (2000) compared the phytoplankton biomass and community structure of different seasons. No information was available for the spring ice algal community in this area.

The aim of this study is to describe the spring species composition, abundance and biomass both in the sea ice and underlying water column of Laptev Sea, to know further the ecological characteristic of ice algal community in this area, the relationship with phytoplankton and its significance to the marine ecosystem.

1 Materials and methods

1.1 Sampling stations and sample collection

Samples were collected from 10 stations in the Laptev Sea between April 17 and May 6, 1999, and most of the stations located in the fast ice zone (Fig. 1). Samples were collected twice at Stn. 01 and stn. 05 (marked as stn. 01—I/stn. 01—II and stn. 05—I/stn. 05—II, respectively). At stn. 03, new ice with 20 cm thickness was also collected (marked as stn. 03—II).

Ice cores were collected using a CRREL ice auger, bottom 10 cm (lowermost 0~10 cm) and sub-bottom (10–20 cm) fragments of sea ice were cut down for algal observation. Each fragment was melted in seawater pre-filtered through a 0.2 μ m membrane to avoid osmotic stress (Garrison and Buck 1986). Underlying water samples were collected with a water sampler at 0 m, 5 m, 10 m and 20 m depths. 250 ml subsamples were collected using brown glass bottles, fixed with 0.2 μ m filtered borax-buffered formalin (1.0% final concentration, v/v), preserved avoiding light and at 4°C until algal counting.

1.2 Counting and biomass conversion

50 ml of each fixed water sample was taken and algal cells were settled using a Hydro-bios sediment chamber according to the method of Utermöhl (1931), and

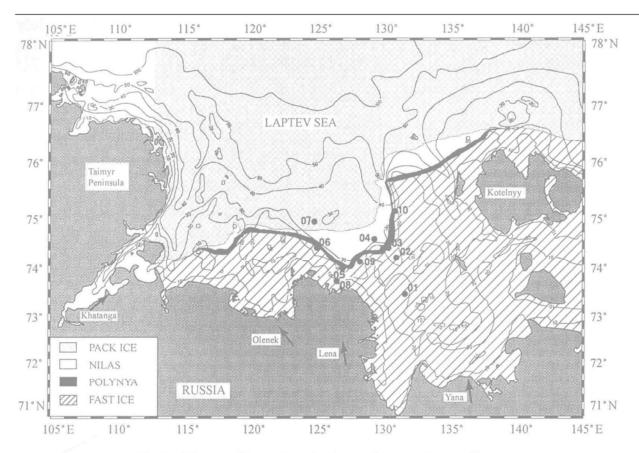


Fig. 1 The sampling stations in the southeastern Laptev Sea.

counted under a Zeiss Axiovert 135. One or two lines of algal cells were counted under a $40\times$ objective lens and the total counting cells were not less than 400 per samples. Several lines, or even whole chamber of algal cells were counted for the phytoplankton samples because of the low abundance of phytoplankton. The cell size of protists was measured using a Porton G12 ocular micrometer (Graticiles Ltd. UK), and the bio-volume was calculated according to the method described by HELCOM (1989). Algal biomass was estimated by the bio-volume with a conversion factor of 0.11 pg C μ m⁻³, and for the armored dinoflagellates, a conversion factor of 0.13 pg C μ m⁻³ was used (HELCOM 1989). Bray-Curtis similarity factor was used to describe the community similarity between those of bottom and sub-bottom ice, and of bottom ice and underlying surface water. The similarity factor was calculated according to Bray and Curtis (1957) with

$$C_N = 2J_N/(aN+bN)$$

Where aN and bN are species numbers of sample a and b, respectively. JN is the common species number occurred in both samples. Considering the considerable variation of algal abundance occurred among micro-habitats (bottom ice, sub-bottom ice and underlying surface water), algal species number other than algal abundance was adopted in the formula for the similarity estimation.

The species with the abundance percentage higher than 10% was regarded as dominant species.

2 Results

2.1 Ice algal abundance and biomass

The ice algal abundance and biomass of the bottom and sub-bottom ice at each station were summarized in Table 1. Algal abundance of bottom ice ranged from 14. 6 to 1562.2×10^4 cells l^{-1} with an average of 639.0×10^4 cells l^{-1} , and algal abundance of sub-bottom ice varied between 5.0 and 1170.1×10^4 cells l^{-1} , with an average of 143.2×10^4 cells l^{-1} . Biomass of bottom ice ranged from 7.9 and 2093.5 μ g C l^{-1} with an average of 886.9 μ g C l^{-1} , and biomass of sub-bottom ice ranged from 5.0 and 594.5 μ g C l^{-1} , with an average of 132.4 μ g C l^{-1} .

Except for stn. 03—II, the ice algal abundance and biomass of the bottom ice were higher than those of the sub-bottom ice, or even 2 orders of magnitude higher at some stations. At stn. 03—II, the total thickness of sea ice (new ice) was 20 cm, and the abundance and biomass of upper 10 cm ice section were 80 and 75 times higher than those of the downside 10 cm ice section, respectively. Stn. 07 was located in the pack ice zone, and no considerable variation was observed between stn. 07 and the stations located in the fast ice zone.

Table 1. Comparison of algal abundance and biomass among ice bottom (0-10 cm), sub-bottom (10-20 cm) and underling water surface

Stations		stn. 01 —	I -II	stn. 02	stn. 03-	I - II	stn. 04 s	tn. 05-	I —II_	stn. 06	stn. 07	stn. 08	stn. 09	stn. 10
Abundance	Sub-bottom ice	14.6	15.5	5.0	9.6	1170.1	18.1	55.4	113.6	34.3	32.2	8. 5	340.1	44.1
	Bottom ice	177.2	884.4	127.3	1239.4	14.6	906.1	74.6	1273.5	168.5	595.1	643.5	641.1	1562.2
	Surface water	0.1	1.0	1.0	0.8	2.3	1.9	1.7	26.4	10.6	1.5	0.4	24.8	5.26
Biomass	Sub-bottom ice	33.4	35.0	5.0	24.9	594.5	35.8	46.0	237.7	20.0	52.8	10.5	579.8	46.2
	Bottom ice	292.4	2093.5	155.4	1387.0	7.9	1448.8	117.6	1642.8	104.2	911.2	971.7	890.3	1506.4
	Surface water	0.1	1.5	1.1	0.3	3.0	1.2	1.0	160.1	4.3	2.2	0.5	36.0	9.48

^{*} The unit of abundance: 104 cells l-1; the unit of biomass: µg C l-1.

The integrated biomass within bottom 20 cm ice column ranged from 13.0 to 160.6×10^7 cells m⁻², with an average of 79.9 \times 10⁷ cells m⁻². The integrated biomass varied between 12.4 and 212.8 mg C m⁻², with an average of 112.0 mg C m⁻² (Table 2).

Table 2. Comparison of integrated abundance and biomass of bottom 20 cm ice and underlying 20 m water column

Stations	,	stn. 01 - I	I stn. 02	stn. 03	stn. 04	stn. 05 —	1 -11	stn. 06	stn. 07	stn. 08	stn. 09	stn. 10	Average
Abundance	Sea ice	90.0	13. 2	124.9	92.4	13.0	138. 7	20.3	62.7	65. 2	98. 1	160.6	79.9
	Water column	6.7	8.3	6.6	14.8	13.1	263.5	72.5	32.1	1.7	308.7	63.0	71.9
	Ratio	13.4	1.6	18.9	6. 2	1.0	0.5	0.3	2.0	38.4	0.3	2.5	7.7
Biomass	Sea ice	212.8	16.0	141.2	148.5	16.4	188.0	12.4	96.4	98. 2	147.0	155.3	112.0
	Water column	10.1	8.5	4.0	8. 7	6.7	1781.7	25.2	34.3	1.9	270.4	193.1	213.1
	Ratio	21.1	1.9	35.3	17.1	2.4	0.1	0.5	2.8	51.7	0.5	0.8	12. 2

^{*} The unit of abundance: 107 cells m⁻²; the unit of biomass: mg C m⁻².

2.2 Phytoplankton abundance and biomass

Phytoplankton abundance ranged from 0.02 and 26.4 \times 10⁴ cells l⁻¹, and the bi-

omass was between 0.03 and 324.9 μ g C l⁻¹. The maximum occurred at 5 m depth of stn.05—II, and the minimum occurred at 20 m depths of stn.02. The phytoplankton abundance and biomass decreased with the depth except for few stations, at which the maximum occurred at 5 m depth.

The phytoplankton abundance of underlying surface water ranged from 0. 1 to 26.4×10^4 cells l^{-1} with an average of 6.0×10^4 cells l^{-1} . The algal abundance varied between 0. 1 and 160. 1 μ g C l^{-1} with an average of 17.0 μ g C l^{-1} (Table 1). The ratios of algal abundance and biomass between bottom ice and underlying surface water had high variability among stations. The minimum was at stn. 03—II (6. 3 and 2. 7, respectively), and the maximum was at stn. 03 with the ratios of 1609. 6 and 4203. 0, respectively. In general, the algal abundance and biomass of the underlying surface water were 2 orders of magnitude lower than those of bottom ice, and one order of magnitude lower than those of sub-bottom ice.

The integrated abundance of upper 20 m water column was between 1. 7 and 263.5×10^7 cells m⁻² with an average of 71. 9×10^7 cells m⁻², the integrated biomass ranged from 1. 9 to 1781. 7 mg C m⁻², with an average of 213. 1 mg C m⁻², suggesting that the integrated biomass of upper 20 m water column was even lower than that of the lowermost 20 cm ice column (Table 2).

2.3 Species composition of ice algae

Table 3 lists the spring dominant species of ice algae in the Laptev Sea. The dominant species included Achnanthes taeniata, Clindrotheca closterium, Fragilariopsis cylindrus, Fragilariopsis oceanica, Melosira arctica, Navicula pelagica, Navicula septentrionalis, Navicula vanhoeffenii, Nitzschia frigida, Nitzschia neofrigida and Nitzschia promarme. Among which, Fragilariopsis cylindrus, Fragilariopsis oceanica and Nitzschia frigida were dominated in few station, and Melosira arctica, N. septentrionalis and Clindrotheca closterium dominated only at one station, respectively. The dominant species composition was variable among stations (Table 3).

The similarity factor of ice algal community between bottom and sub-bottom ice varied between 0.37 and 0.64 with an average of 0.52, suggesting that similarity, in some extent, existed between these two communities.

2.4 Species composition of phytoplankton

The dominant species of phytoplankton included Achnanthes taeniata, Fragilariopsis cylindrus, Fragilariopsis oceanica, Melosira arctica, Navicula pelagica, Navicula septentrionalis, Nitzschia frigida, and Nitzschia promarme. Among which, Fragilariopsis cylindrus, Fragilariopsis oceanica, Nitzschia frigida and Pyramimonas sp., as dominant species, occurred only in one station. Nitzschia neofrigida, Navicula vanhoef fenii and Clindrotheca closterium, which were the dominate species.

nant species within the ice, were not the dominant species in the water column. By contraries, *Pyramimonas* sp. dominated only in the water column (surface water at stn. 10).

The diatoms generally dominated the phytoplankton of the whole water column, except for stn. 10. In the surface water at stn. 10, phytoflagellates such as *Pyramimonas* sp. constituted 61% of total algal abundance and 14.4% of total biomass.

The similarity factor of algal community between underlying surface water and bottom ice ranged from 0. 22 and 0. 59 with an average of 0. 41, which was lower than that between bottom and sub-bottom ice.

Table 3. List of dominant algal species (percentage abundance>10%) in bottom ice, sub-bottom ice and underlying surface water

	underlying surface water				
Stations	Sub-bottom ice	Bottom ice	Underlying surface water		
Stn. 01 - I	N. frigida(23.1), N. neofrigida(19. 2), F. cylidrus(17.3), N. vanhoeffenii(15.8)	C. closterium(13.0) N. noefrigida(10	A. taeniata(67.6), N. frigida(16.9), F. oceanica(11.3)		
-II	F. cylindrus (50. 3), N. frigida (22. 7)	 N. frigida(24.4), A. taenicata(16.9) F. cylindrus(16.0), N. neofrigida(13 1) 	, A. taeniata(22,7), N. frigida(14,6), F. cylindrus(12,7) N. promare(11,2),		
		• /	F. oceanica(10.4)		
Stn. 02	F. cylindrus(40.3), F. oceanica(15.2),	F. cylindrus(43.1), N. frigida(11.8)	F. cylindrus(57.1)		
	N. arctica(10. 2)				
Stn. 03 -1	F. cylindrus(23, 0), M. arctica(21, 5)	F. oceanica (30. 4), F. cylindrus (22. 8),	F. oceanica(92, 0)		
-11	F. oceanica(57, 9), N. septentrionalis(20,7),	N. promare(22. 2) F. oceanica (51. 7), F. cylindrus (15. 5).	N. septentrionalis(33.5), M. arctica(26.8),		
Stn. 04	A. taeniata(11.2) F. oceanica(11.5) F. frigida(29.2), F. cylindrus(18.2), N. plagica(13.7), S. radisonii(13.0)	F. cylindrus(11.9) N. frigida(36.8), N. plagica(24.3)	F. cylindrus(41.7), F. oceanica(23.2)		
Stn. 05 - I	N. cylindrus(52, 6)	N. frigida(25.9), N. pelgaica(22.2)	F. cylindrus(56, 4), F. oceanica(13, 9)		
-11	N. frigida(48.1), N. promare(24.0)	N. frigida(39.4), N. promare(39.1)	M. arctica(71.1), F. cylindrus(12.3)		
Stn. 06	F. oceanica(89, 4)	F. oceanica(86. 6)	F. oceanica(64.4), F. cylindrus(14.8)		
Stn. 07	N. frigida(49.7)	N. frigida(65.1)	N. frigida(52.8), N. plagica(13.0),		
			F. cylidrus(10, 1)		
Stn. 08	N. frigida(29.5), F. oceanica(10.8)	N. frigida(37.5), F. cylindrus(15.0)	N. frigida(26.9), A. taenicata(10.8)		
Stn. 09	N. frigida (20, 8), F. oceanica (20, 8), A. taeniata (18, 3), N. promera (11, 8),	N. frigida(28.5), N. promare(26.6), F. oceanica(44.3), N. frigida(12.5), N. vanhoe fenii (17.8), M. arctica (10.9)			
Stn. 10	M. arctica(11, 4) F. cylindrus(51, 2), N. promare(27, 8)		Pyramimonas sp. (26.8), M. arctica(19.3)		

3 Discussion

3.1 Spatial distribution

The variation of algal communities among stations was considerable. For the bottom or sub-bottom ice, the maximum of variation between stations could be 2 orders of magnitude higher, and for the surface water, even 4 orders of magnitude higher. Tuschling et al. (2000) also reported the high biomass variations of new ice, surface water or water with 5 m depths at different stations. Previous researches suggested that in the Arctic estuary area, covering snow on the sea ice and the salinity of

the ice-water interface were two crucial factors controling the ice algal abundance of the bottom ice (e. g. Monti et al. 1996).

The abundance and biomass within ice were high in contrast to those in the water column. Table 2 shows that the average ratios of integrated abundance and biomass between 20 cm of bottom ice and 20 m of underlying water column were 7.7 and 12. 2. respectively, suggesting that in the southeastern Laptev Sea with the water depth of less than 40 m in most area, the algal biomass within ice column might be higher than those in the whole water column, and served as an important food source for a variety of the zooplankton in spring. Horner and Schrader (1982) reported that in the Beaufort Sea, ice algae contributed about 2/3 of the spring primary production in this area. No information was available for the grazing activity of zooplankton of spring Laptev Sea. However, Werner and Martinez Arbizu (1999) reported that in summer Laptev Sea, nauplii aggregated at the ice-water interface, and the abundance maximum could be 23911 individuals m⁻³.

3.2 Relationship between ice algae and underlying phytoplankton

Our present study shows that the similarity factor of algal community between underlying surface water and bottom ice ranges from 0. 22 and 0. 59 with an average of 0. 41. It was not high. However, species such as N. frigida and F. cylindrus, the typical ice algae, were common in the water column, suggesting that at least some of the dominant species in the water column came from bottom ice. The continuance and extent of the influence of spring ice algae to the underlying phytoplankton community was depended on the release velocity of ice algae. Fortier et al. (2002) proposed that covering snow was crucial for the ice algae release. Ice algae would release in a short period, if the rapid warming or precipitation led to the quick disappearance of covering snow.

Several researches in the Arctic coastal seas showed that ice algae could play as "seeding" of the spring phytoplankton bloom (Schandelmeir and Alexander 1981; Michel et al. 1993). Present study showed that the dominant species such as F. cylindrus, F. oceanica and N. frigida were all algal aggregates. When released from sea ice, these aggregates would be grazed easily by zooplankton or settled quickly to the sea floor (Sancetta 1981; Cremer 1999; Fortier et al. 2002), and therefore the role of these aggregated to the "seeding" was negligible. It has been reported that the summer phytoplankton community in the Laptev Sea affected considerably by the influx of fresh water (Clark 1990). However, in present study the phytoplankton biomass was low and most of the species were marine species, suggesting that the influence of fresh water algal species to the phytoplankton community did not emerge yet during the investigation period.

The algal community in the surface water at stn. 10 dominated with phytoflagellates other than diatom, which might be caused by the spring phytoflagellate bloom in the ice-water interface. Gradinger (1996) reported a summer *Pyramimonas* sp. bloom in the downside melting pond under pack ice of the Greenland Sea, and the abundance could be as high as 19.1×10^6 cells l^{-1} . Some fresh water species such as Aulacosira sp. were observed in the bottom ice of stn. 03-I, suggesting that the bottom ice community might be influenced, in some extent, by the phytoplankton.

3.3 Seasonal variation

No information for autumn ice algal abundance is available in the Laptev Sea because of the high sediments within ice and difficulty of cell counting under microscope (Tuschling 2000). Comparison of ice algal communities between of spring and autumn suggest that seasonal variation of species composition occurred (Tushling 2000; Krember and Angle 2001). This variation might be caused by physical or biological processes. In summer, influenced by fresh water influx, summer phytoplankton was dominated by fresh-brackish water species or brackish water-marine species (Cremer 1998; 1999). In autumn, in the processes of ice formation, ice crystal harvested phytoplankton from the upper water column (Ackley 1982, Garrison et al. 1983, He et al. 2003). Consequently, the autumn ice algal community would influenced by the phytoplankton community of the upper water column. Our present study showed that the spring phytoplankton abundance was low, suggesting that the high biomass of ice algae was caused by the biological process, the in situ growth of ice algae.

Marked seasonal variations of phytoplankton abundance and biomass were observed in the Laptev Sea. The spring phytoplankton abundance of the surface water ranged from 0.1 to 26.4 \times 10⁴ cells l⁻¹, and the biomass was between 0.1 and 160.1 μ g C l⁻¹, which was higher than that from spring 1996 (<0.1-88.4 μ g C l⁻¹, Tushling 2000), and much higher than that from autumn (351-33660 cells l⁻¹, 0.1 \sim 5.3 μ g C l⁻¹, Tushling *et al.* 2000). The researches have shown that the maximum of algal biomass occurred in summer (Tushling 2000). Zernova *et al.* (2000) also reported that in the northern Laptev Sea, the vertical algal flux in summer were several hundreds higher than that in other seasons.

3.4 Comparison to other Arctic seas

The dominant species of spring Laptev Sea resembled to those of the coastal seas of Beaufort and Siberia Seas, and of Canadian Arctic seas. F. cylindrus, F. oceanica and N. frigida were common spring dominant species of Arctic coastal seas (Cross 1982; Horner and Schrader 1982; Okolodkov 1992a, 1993; Michel et al. 1993). The ice algal community had similarity, in some extent, with that of the first-year pack ice. However, there was obvious variation with that of multi-year pack ice in the high Arctic (Syvertsen 1990; Krember and Engel 2001). The percentage abundance of Melosira arctica in the bottom ice at stn. 09 was 10.9%. This species was commonly a dominant ice algal species of the multi-year pack ice of high Arctic, and the abundance maximum could be 11.57×106 cells 1-1 in summer (Booth and Horner 1997). However, traditional view suggested that Melosira arctica was not important in the community of the fast ice (Hsiao 1980; Okolodkov 1992a), and has not been de-

scribed as a dominant ice algal species of the fast ice before.

The spring phytoplankton abundance of underlying surface water was higher than that of Beaufort Sea, but lower than that of Canadian Arctic seas and Chukchi Sea (Hsiao 1980; Horner and Schrader 1982; Okolodkov 1992a).

The results from re-sampling station of stn. 01 and stn. 05 showed that the biomass of the bottom ice and underlying water column was increased during the investigation period. However, the observation of ice algal samples showed there were many diatom frustules, and most of dominant species had attached algae such as Synedra hyperborean, Attheya septentrionalis and Pseudogomphonema arctica, suggesting that the ice algae community was not healthy and the peak of algal biomass might not occur in the late May to early June, just before the thawing of sea ice, which usually occurred in other Arctic fast ice zone (Hsiao 1980; Horner and Schrader 1982).

4 Conclusion

There were some characteristics of the spring algal community of the southeastern Laptev Sea:

- 1) Algae aggregated mostly in the bottom of sea ice. In general, the abundance and biomass of the lowermost 10 cm sea ice were one order of magnitude higher than those of the sub-bottom 10 cm sea ice, and two orders of magnitude higher than those of the underlying surface water.
- 2) The variation of algal spatial distribution was considerable. The maximum of algal abundance and biomass could be 2 to 4 orders of magnitude as the minimum.
- 3) Ice algae community would influence the phytoplankton community during the investigation period, but the "seeding" role of ice algae to phytoplankton was negligible.
- 4) Although the spring ice algal biomass was lower than that of summer phytoplankton, the spring ice algal bloom occurred in the bottom of sea ice suggests that, ice algal community might be significant in the ecosystem at least in the special period when the phytoplankton in the water column was low.

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