

Characterization of extracellular polymers of *Phaeocystis globosa* and *P. antarctica*

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ABSTRACT: Extracellular polymers of *Phaeocystis globosa* and *P. antarctica* were characterized by elemental analysis, X-ray photoelectron spectroscopy (XPS), and matrix-assisted laser desorption ionization – time of flight – mass spectrometry (MALDI-TOF-MS) analysis. The 2 species were grown in axenic and bacterized cultures to examine the effect of bacterial degradation on the composition of the extracellular polymers. The C:N ratios for extracellular polymers from *P. globosa* and *P. antarctica* grown under these conditions were lower (<9) than in previous reports (>12). *P. antarctica* polymer C:N was significantly lower than that of *P. globosa*. Contrary to expectations, the polymer C:N from bacterized cultures was slightly lower than from axenic cultures. XPS analysis indicated that the polymers are carbohydrate-rich, and MALDI-TOF-MS analysis showed that the nitrogen was not proteinaceous, but probably originated from low molecular weight compounds such as amino sugars. The low C:N ratios suggest that extracellular polymers may directly provide a substrate for bacteria and heterotrophic protists without the need for other sources of nitrogen.

KEY WORDS: *Phaeocystis* spp. · Carbohydrates · C:N ratio · Extracellular polymers

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INTRODUCTION

Phaeocystis spp., which are colony-forming prymnesiophyte algae, produce large amounts of extracellular polymers that can contribute to the high molecular weight dissolved organic matter pool in the ocean (Lancelot 1995, Biersmith & Benner 1998). The polymers produced during blooms can undergo sedimentation (Wassmann et al. 1990, Riebesell et al. 1995, Smith & Dunbar 1998), or microbial degradation by heterotrophic bacteria (Janse et al. 1999) or protists capable of ingesting high molecular weight compounds (Sherr 1988, Marchant & Scott 1993, Tranvik et al. 1993), but the biological and physical controls over these loss pathways are not well understood. Questions about the fate of these polymers are further raised by the fact that *Phaeocystis* blooms can result in either accumulation (i.e. North Sea: Lancelot et al. 1987) or no accumulation (i.e. Ross Sea: Carlson et al. 1998) of polymers in the dissolved organic carbon pool.

The differing degrees of microbial degradation of *Phaeocystis* extracellular polymers during blooms may depend on differences in the polymer chemical composition of species of *Phaeocystis* that thrive in distinct environments. The biodegradability of *Phaeocystis* polymers will in part depend on their ability to provide sufficient carbon and nitrogen to bacteria and protists. Extracellular polymers from most phytoplankton species consist primarily of carbohydrates (Guillard & Hellebust 1971, Mykkestad & Haug 1972, Painter 1983, Hoagland et al. 1993). If the polymers also contain glycoproteins or amino sugars, they may directly provide nitrogen to microheterotrophs. On the other hand, degradation of high carbon polymers may be limited by the availability of external sources of nitrogen such as NH_4^+ , NO_3^- or free amino acids from the surrounding seawater.

The molecular composition of extracellular polymers from *Phaeocystis*, as well as other phytoplankton, is not well defined. Most studies characterizing phytoplank-

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ton extracellular polymers have focused on assessing the carbohydrate fraction using bacterized cultures. The monosaccharide composition of *P. globosa* extracellular polymers consists primarily of arabinose, xylose, mannose, galactose, glucose, and rhamnose (Guillard & Hellebust 1971, Janse et al. 1996a,b, Aluwihare & Repeta 1999). Chitin has also been found in ejected filaments of *P. globosa* (Chretiennot-Dinet et al. 1997). Recently, various methods have been applied to assess the nitrogenous components of the polymers. Amide groups were detected in *P. globosa* colonial polymers (Hamm et al. 1999), suggesting that these have a protein or an amino sugar component.

Chemically distinct polymers may be produced by different *Phaeocystis* species. The C:N ratios of whole *P. pouchetti* colonies, consisting of both cells and extracellular polymers, vary from 4.5 to 20.5 (Verity et al. 1988, 1991, Riebesell et al. 1995, Hegarty & Villareal 1998), whereas those of *P. antarctica* colonies vary from 5.94 to 6.22 (Moisan & Mitchell 1999). Thus, the C:N ratio of *Phaeocystis* polymers may be highly variable, depending on species and growth conditions.

However, analyses of the composition of *Phaeocystis* extracellular polymers *without* the inclusion of cells are few (Biersmith & Benner 1998, Aluwihare 1999, Aluwihare & Repeta 1999), and are available for only 2 strains of *P. globosa* in bacterized cultures. In this study, we examined the chemical composition of axenic and bacterized extracellular polymers from another strain of *P. globosa* and from *P. antarctica*, which has not been previously characterized. We characterized polymers by X-ray photoelectron spectroscopy (XPS) and matrix-assisted laser desorption ionization–time of flight–mass spectrometry (MALDI-TOF-MS).

MATERIALS AND METHODS

Culture conditions and isolation of extracellular polymers and polymer gels. Original stock cultures of *Phaeocystis globosa* (CCMP 629, isolated from the North Atlantic) and *P. antarctica* (isolated by E.J.L. from the Ross Sea, Antarctica) were transferred to f/2 media (Guillard & Ryther 1962) with antibiotics (Provasoli's antibiotic concentrated solution, Sigma, 0.5%) until axenic cultures were obtained. The absence of bacteria was confirmed by direct microscopic observation and the absence of bacterial growth on marine yeast-extract media (2216 marine broth, VWR Scientific Products). Axenic and bacterized cultures of *P. globosa* and *P. antarctica* were then grown in f/10 media (177 μM NO_3^- , N:P = 24:1; Guillard & Ryther 1962), transferred to 8 l carboys containing 4 l of f/10 media, and exposed to an irradiance of ca. 113 $\mu\text{mol photons m}^{-2} \text{s}^{-1}$ (ranging from 107 to 119 $\mu\text{mol photons m}^{-2} \text{s}^{-1}$ in the carboys) on a 14 h

light:10 h dark cycle at 18 and 4°C, respectively. Bacterial strains which had been isolated from cultures of each species were added to axenic cultures for the bacterized treatment (Strain 18-18 in *P. globosa* cultures, Strain 4-4 in *P. antarctica* cultures; bacteria isolated by E. Peele). Both bacterial strains were capable of growing on *Phaeocystis* polymers without added nitrogen or carbon sources (E. Peele & E. J. Lessard unpubl.). Samples of 10 ml were taken daily for determination of phytoplankton and bacterial cell numbers. Fixative (0.5% glutaraldehyde) and the fluorescent nuclear stain DAPI (5 $\mu\text{g ml}^{-1}$ final concentration) were added and filtered onto 0.2 μm black polycarbonate filters. The filters were placed on slides with low-fluorescing oil, and cells were enumerated using epifluorescence microscopy. The biomass of the algae and bacteria was estimated using 14.2 pg C cell^{-1} for *Phaeocystis* cells (Rousseau et al. 1990) and 13 fg C cell^{-1} for bacteria cells (Ducklow 2000).

During late exponential phase, after 9 to 15 d incubation, the cells were separated from polymers and polymer gels <0.2 μm in size by gravity-filtering through a SuporCap™ 0.2 μm capsule filter (Pall/Gelman Laboratories). Preliminary growth experiments under the same culture conditions indicated the onset of the stationary phase after 10 d in for *P. globosa* and after 16 d in for *P. antarctica*. The timing of polymer collection was designed to harvest both species of *Phaeocystis* at the same phase of growth under nonlimiting conditions when cells were healthy. The filtrate was allowed to sit for 3 d at culture temperature to allow dispersed polymers to assemble into gels. *Phaeocystis* polymer gels, which are hydrated polymer networks stabilized by ionic bridges, self-assemble from <0.2 μm -filtered dispersed polymers on a timescale of ca. 2 d (Chin et al. 1996, 1998). The polymers and gels in the filtrate were then concentrated from 2–3 l to 500 ml (concentration factor = 4 to 6) using a CH2L Amicon ultrafiltration system with a 1 nm pore size (1 kDa nominal molecular weight cut-off) regenerated cellulose spiral-wound cartridge (S1Y10), diafiltered with 2.4 l deionized water with 10 mM CaCl_2 , and lyophilized. To maximize retention of polymer gels, CaCl_2 was added during diafiltration to prevent the loss of Ca^+ , which is needed for the stability of *Phaeocystis* polymer gels (van Boekel 1992, Chin et al. 1998). The ultrafiltration was performed under sterile conditions (tubing and bottles were autoclaved, ultrafiltration unit and cartridge were gas-sterilized) under a laminar flow hood. Between uses, the cartridge was cleaned (0.01 N NaOH without pressure for 20 min and with pressure for 20 min, rinsed with autoclaved Nanopure water until the pH was the same as autoclaved Nanopure water) and preconditioned with the sample before concentration (after Dai et al. 1998).

Calibration of 1 kDa cartridge. We tested the retention efficiency of the 1 kDa-regenerated cellulose cartridge using protein (gramacidin S, 1.2 kDa; Sigma) and carbohydrate (maltoheptaose, 1.15 kDa; Sigma) standards. Initial standard concentrations were about 1 mg C l^{-1} in Nanopure water. Initial samples were taken of the standard before and after preconditioning. After concentration by a factor of 4, samples were taken from the retentate (concentrated sample) and permeate (filtrate) and measured with a total organic carbon analyzer for mass balance calculations. The retention efficiency of gramicidin and maltoheptaose was 14 and 53%, respectively. Most of the gramicidin and maltoheptaose was either lost to the permeate (35 and 17%) or absorbed by the cartridge membrane (52 and 29%).

Carbon and nitrogen analysis. Triplicate aliquots of the lyophilized concentrate from each culture were analyzed for carbon and nitrogen using a Leeman Labs Model CEC440 elemental analyzer. The concentrates were 0.10 to 0.17% C and 0.025 to 0.039% N by weight; the remaining material was CaCl_2 from the dialysate and other salts, including NO_3^- . To determine the contribution of NO_3^- -nitrogen to total nitrogen, triplicate aliquots of the lyophilized concentrates were resuspended in Nanopure water and analyzed for NO_3^- using a Technicon AA11 autoanalyzer. We corrected the C:N ratio of the polymers by subtracting the NO_3^- -N from the N value measured by the CHN analyzer for a known mass of sample.

XPS. XPS can analyze several nanometers of a surface, give information on all elements and molecular environment (e.g. oxidation state), and identify organic groups (Ratner & Castner 1997). XPS analyses were made according to Arnarson & Keil (2001) using a Surface Science Instruments (SSI) S-Probe ESCA instrument. An aluminum K_{α} 1,2 monochromatized X-ray source was used to stimulate photoemission, and an electron flood gun set at 4 eV was used to minimize surface charging of the samples. The energy of the emitted electrons was measured with a hemispherical energy analyzer at pass energies of either 150 eV (overview spectra for elemental composition) or 50 eV (high-resolution spectra for chemical composition of carbon and nitrogen). SSI data analysis software was used to calculate the elemental compositions from the peak areas of the overview spectra and to peak-fit the high-resolution spectra with Gaussian functions. The overview spectra that included salts (data not shown), was narrowed down to 280–300 eV to examine the C_{1s} (electron) spectra. The accuracy of the elemental compositions is limited to 10% relative error due to uncertainty about the sensitivity factors used (Arnarson & Keil 2001).

TOF-MALDI-MS. TOF-MALDI-MS can give information about polymers in their intact state, while other

techniques break the polymers into smaller fragments (Holcombe et al. in press). The same polymers analyzed by XPS were run on a Bruker Reflex III TOF mass spectrometer (Bruker-Franzen) with a single probe inlet, equipped with a UV-nitrogen laser. Matrix-assisted laser desorption and ionization time of flight mass-spectrometry analyses were conducted as described by Holcombe et al. (in press) using 2,5-dihydroxybenzoic acid (DHB, Aldrich) as the matrix. All spectra were acquired in the positive-ion linear reflector mode. Typically, 100 to 500 laser shots were accumulated in the resulting spectra and the attenuation setting was approximately 40%. The first scan was from 100 to 100 000 m/z, while the subsequent scans were in smaller groups (50 000 to 100 000, 20 000 to 100 000, 7000 to 500 000, followed by 100 to 1000 m/z). Mass calibration was performed by using singly and doubly protonated molecular ion signals of chicken egg-white lysozyme protein standard (molecular wt = 14 264).

Statistical analysis. Data were analyzed by mixed-model analysis-of-variance techniques (ANOVA; SAS 2001) which contain both fixed and random factors. The 3 factors were cell/polymer, species, and treatment. The mixed ANOVA also considered the 2- and 3-factor interactions. The 2 assumptions of ANOVA techniques, homogeneity of variance and normality, were tested. The assumption of homogeneity of variance was examined using goodness-of-fit statistics and residual variance was found to be greater for *Phaeocystis globosa* than for *P. antarctica*. Mixed ANOVA techniques were then used to partition the residual variance into 2 residuals, 1 for each species, solving the heterogeneity of variance problem. The assumption of normality was examined and found satisfactory within species. Means of *Phaeocystis* species cell and polymer C:N ratios were compared by planned contrasts.

RESULTS

Growth of *Phaeocystis* species cultures

Phaeocystis globosa incubated at 18°C grew at 0.59 d^{-1} , and *P. antarctica* incubated at 4°C grew at 0.35 d^{-1} (Table 1, Fig. 1). The growth rates of both *Phaeocystis* species were slightly higher in the presence of bacteria. Bacteria (Strain 18-18) in the *P. globosa* culture grew at 1.73 d^{-1} from an initial standing stock of $7 \times 10^3 \text{ cells ml}^{-1}$ while bacteria (Strain 4-4) in the *P. antarctica* cultures grew more slowly, at 0.44 d^{-1} , from an initial standing stock of $7 \times 10^2 \text{ cells ml}^{-1}$ (Fig. 1, Table 1). In both bacterized *P. globosa* and *P. antarctica* cultures, bacterial biomass was <1% of the total biomass when polymers were harvested.

Table 1. *Phaeocystis globosa* and *P. antarctica*. Growth rates in axenic and bacterized cultures. Growth rate of Bacterial Strains 18-18 and 4-4 in the cultures are also shown

Species	Temp. (°C)	<i>Phaeocystis</i> growth rate (d ⁻¹)	Bacteria growth rate (d ⁻¹)
<i>P. globosa</i>	18	0.59	
<i>P. globosa</i> with Bacterial Strain 18-18	18	0.60	1.73
<i>P. antarctica</i>	4	0.35	
<i>P. antarctica</i> with Bacterial Strain 4-4	4	0.37	0.44

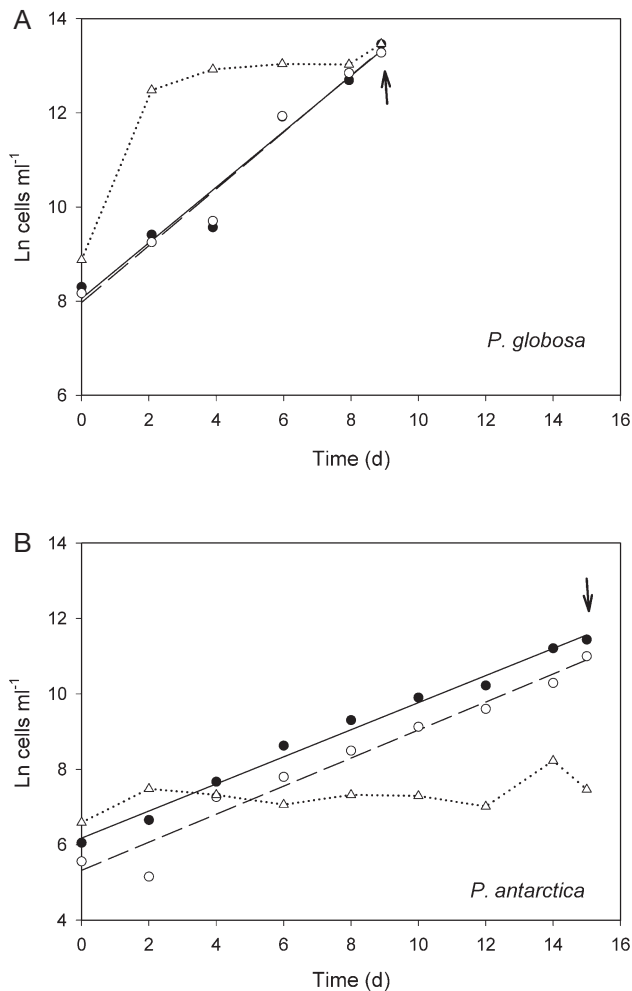


Fig. 1. *Phaeocystis globosa* and *P. antarctica*. Growth curves. (A) *P. globosa* (●) and *P. globosa* with Bacterial Strain 18-18 (○); growth curve for Bacterial Strain 18-18 is also shown (Δ). (B) *P. antarctica* (●) and *P. antarctica* with Bacterial Strain 4-4 (○); growth curve for Bacterial Strain 4-4 (Δ) is also shown. Arrows denote time when *Phaeocystis* polymers and polymer gels were collected

Comparison of C:N ratios of *Phaeocystis globosa* and *P. antarctica* extracellular polymers and cells

The C:N ratios of extracellular polymers from the 2 *Phaeocystis* species ranged from 6.20 to 8.32 (Fig. 2). The C:N ratio of *P. globosa* polymers from axenic culture was higher than those from bacterized culture (mean \pm SE: 8.32 ± 0.28 vs. 7.40 ± 0.03), but the difference was significant only at $p = 0.07$. The C:N ratios of polymers from axenic *P. antarctica* were significantly higher than those from bacterized cultures, but the difference (6.86 ± 0.03 vs. 6.20 ± 0.04 , $p < 0.01$) was less than for *P. globosa*. *P. globosa* polymers had a significantly higher C:N than *P. antarctica* polymers under both axenic and bacterized conditions ($p < 0.05$). The C:N ratios of *Phaeocystis* cells (range 4.56 to 6.05) were lower than the C:N ratios of polymers. In all cultures except in the bacterized *P. antarctica* treatment, cell C:N was significantly lower than polymer C:N ($p < 0.05$, *P. antarctica*, $p = 0.2$). The *P. antarctica* cell C:N of axenic and bacterized culture was significantly different (5.22 ± 0.10 vs. 6.05 ± 0.09 , $p < 0.01$), while *P. globosa* cell C:N was not (4.46 ± 0.44 vs. 4.99 ± 0.32 , $p = 0.36$). The interspecific difference between cellular C:N was significant only for the bacterized cultures ($p < 0.05$).

XPS and MALDI-TOF-MS

XPS analysis indicated very little lipid (CH_x) in the samples, and the spectra had a general C_{1s} binding energy distribution consistent with the hypothesis that the extracellular polymers were composed

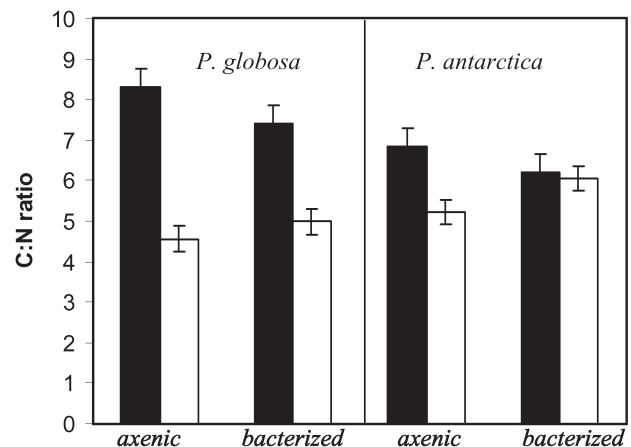


Fig. 2. *Phaeocystis globosa* and *P. antarctica*. Comparison of C:N ratios of axenic and bacterized cultures. Black bars: polymer C:N ratios; white bars: cell C:N ratios; error bars: standard error ($n = 3$)

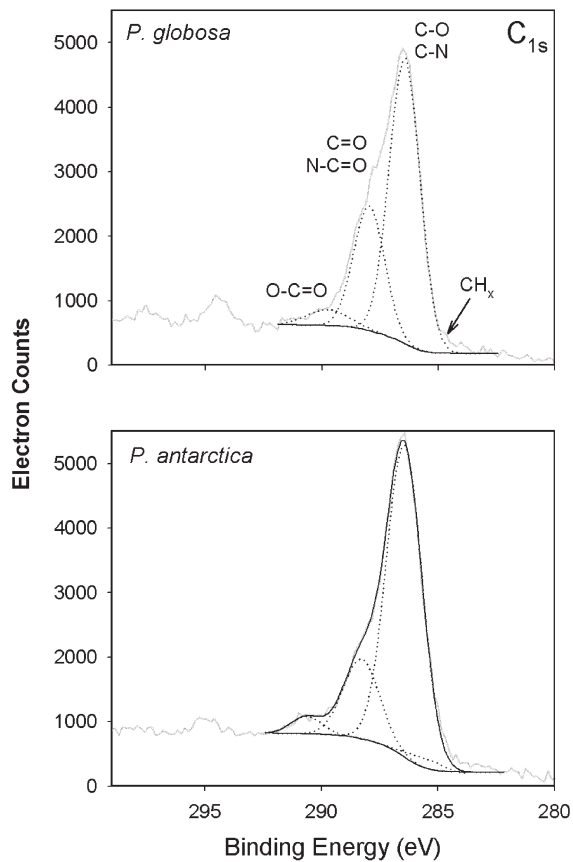


Fig. 3. *Phaeocystis globosa* and *P. antarctica*. X-ray photoelectron spectroscopy analysis of C_{1s} electron peak from 280 to 300 eV for extracellular polymers. CH_x (lipid) peak set at 285.0 V is reference point for binding-energy scale. Chemical shift values for functional groups are as follows: C–N (286.0), C–O (286.5), N–C=O (288.2), C=O (288.0), and O–C=O (289.0) (Ratner & Castner 1997)

largely of carbohydrates (Fig. 3). Integration of the peaks broadly assigned to the C–O, C=O and O–C=O functionalities resulted in a ratio of 12:6:1. The ratio for a simple sugar (e.g. an aldose) is 5:1:0, the ratio for an amino acid is roughly 1:0:1, and the ratio for amino acids in peptides (amide bonds) is 0:1:1. Given a C:N ratio of roughly 7 and assuming that the nitrogen is contained within amide bonds (as observed by Hamm et al. 1999), possible fits for the XPS spectra included mucopolysaccharides, aldoseamines, or a small glycopeptide such as *n*-glycoside.

The MALDI-TOF-MS data were consistent with the presence of carbohydrates, not proteins, in the sample (e.g. Harvey 1999). Analysis of peak distributions were done using available databases (PepFind, etc.: Holcombe et al. in press). Compounds larger than

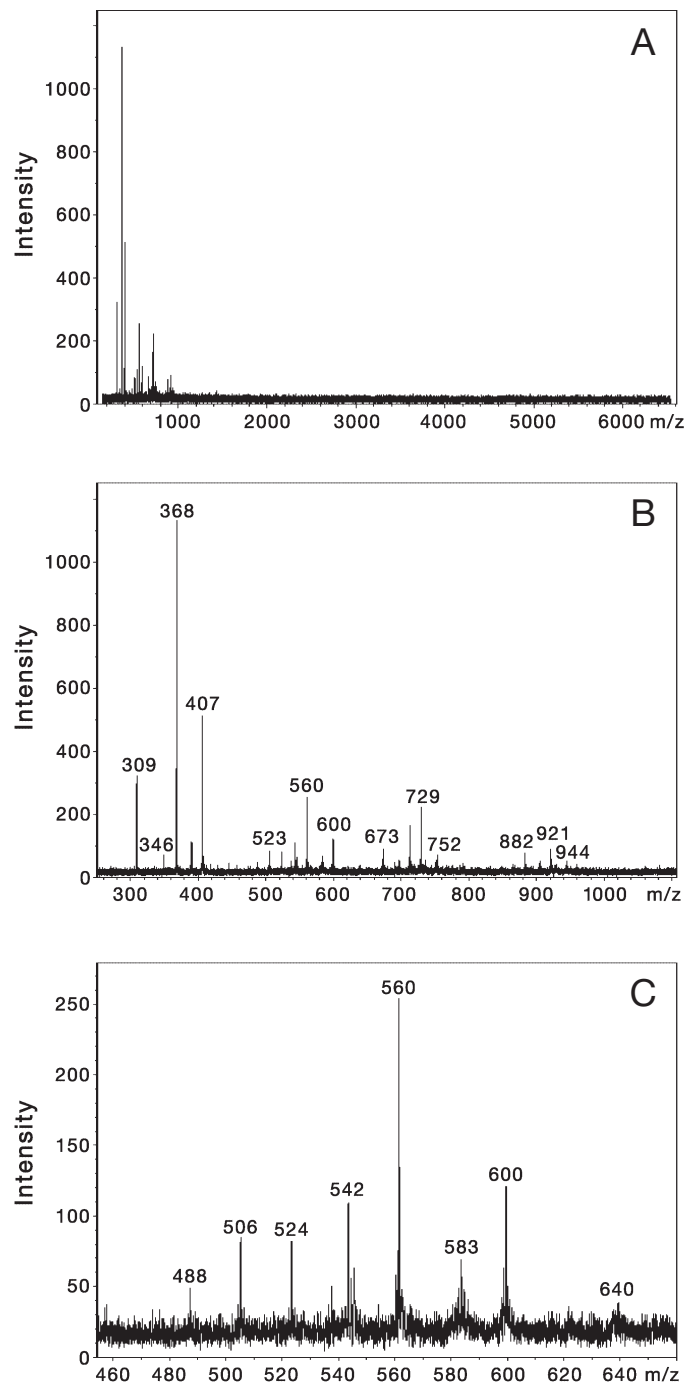


Fig. 4. *Phaeocystis globosa*. Matrix-assisted laser desorption/ionization–time of flight–mass spectrometry spectra for extracellular polymers isolated from axenic cultures. (A) Full spectra from 100 to 7000 m/z showing components only in the <1000 m/z range; spread of the 4 major clusters of peaks in the overview spectra is indicative of macromolecules with repeating structural units (i.e. carbohydrates: Harvey 1999). (B) Spectra from 250 to 1000 m/z , illustrating general pattern of approximately ~ 192 m/z between clusters of peaks (e.g. Peaks 368–560–752–944). (C) Spectra from 450 to 650 m/z illustrating progressive loss of water (18 m/z) from m/z 560 peak, sodium (23 m/z) from m/z 583 peak, and calcium (40 m/z) from m/z 600 peak

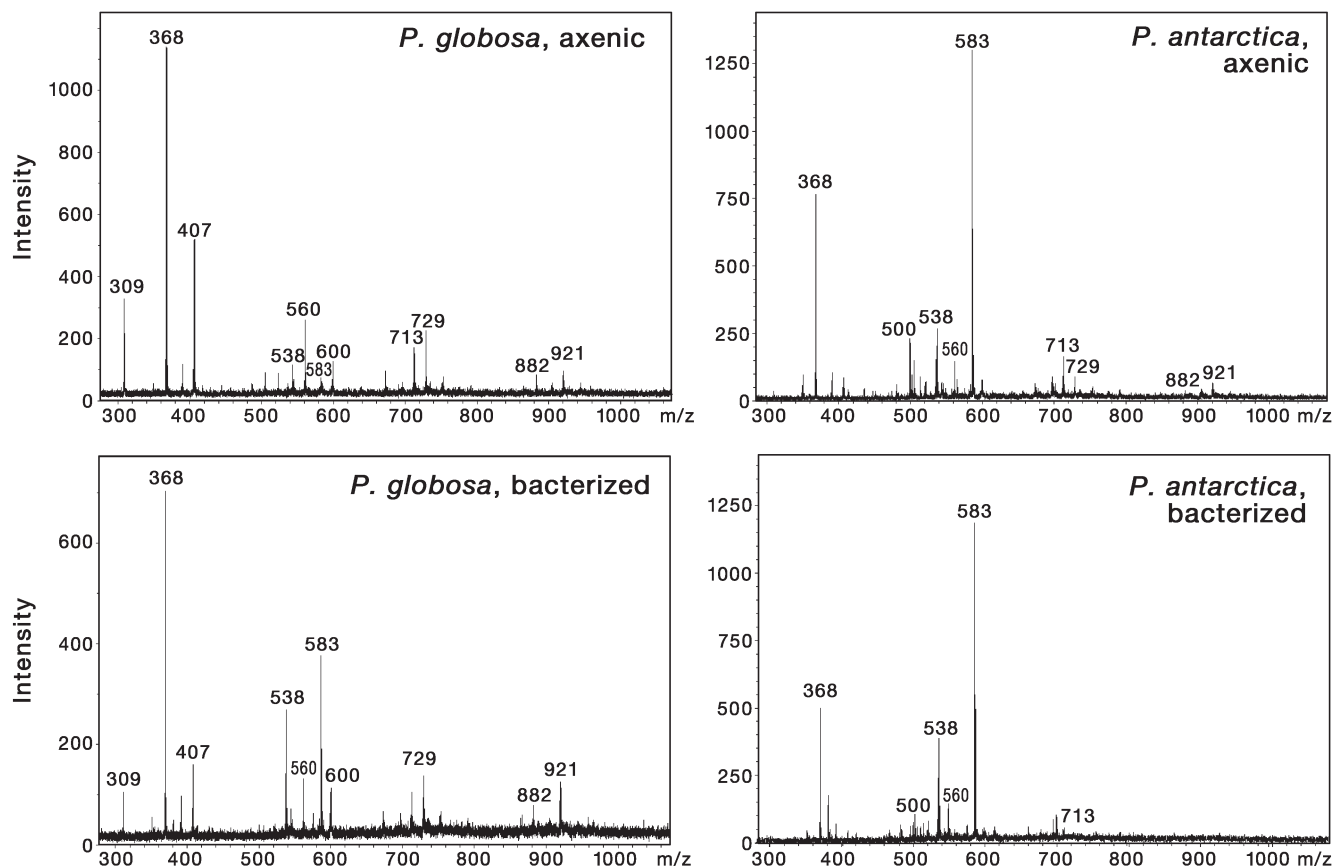


Fig. 5. *Phaeocystis globosa* and *P. antarctica*. Mass spectra in m/z range 250 to 1100 for algae with and without bacteria present during growth

1000 m/z (Da) were not present, and there was a repeating pattern in the spectra (Figs. 4 & 5). The repeating unit of ~ 192 m/z (Da) was close to the molecular weight of the amino sugar glucosamine (187 Da) and the initial mass of 309 was similar to that of sialic acid. Although speculative, the repeat of ~ 190 indicates that the polymer might not be a glycopeptide, since common bacterial glycopeptides have molecular weights between 400 and 700 and no simple way of generating repeated fragments with a m/z of ~ 190 . The peaks of the size ~ 560 m/z (Da) could represent a small polymer of 3 amino sugars ($3 \times 187 = 561$). The MALDI-TOF-MS data also showed consistent signals of Ca^{2+} (40 m/z), Na^{2+} (23 m/z), and H_2O (18 m/z) from the seawater and CaCl_2 additions during isolation of the polymers. This is further evidence that each of the clusters of peaks separated by ~ 190 units represents a carbohydrate. Polymers were present in similar compositions in both axenic and bacterized cultures of both species, although the bacterized sample of *Phaeocystis antarctica* did not show the presence of any peaks larger than ~ 720 m/z .

DISCUSSION

Extracellular polymer composition

The C:N ratios of the 2 *Phaeocystis* species extracellular polymers obtained in this study were much lower than those previously reported for 2 other strains of *P. globosa* (19.3 and 12: Biersmith & Benner 1998, Aluwihare & Repeta 1999, respectively). Our relatively low C:N ratios suggest that the extracellular polymers may contain significant amounts of protein or amino groups, and do not entirely consist of carbohydrates as suggested by many authors (Guillard & Hellebust 1971, Painter 1983). Our results are somewhat surprising, since monosaccharides have been found to be major components of polymers by independent means such as gas chromatography, ^1H NMR (proton nuclear magnetic resonance), and the MBTH method (Biersmith & Benner 1998, Aluwihare & Repeta 1999).

Could protein potentially be an important component of *Phaeocystis* extracellular polymers? Amino ($-\text{NH}_2$) groups have been shown to be prevalent on the

Phaeocystis colonial skin using the fluorescent probe TAMRA-SE (5-and 6-carboxytetramethylrhodamine succinimidyl ester), and hypothesized to belong to proteins (Hamm et al. 1999). In some studies of *Phaeocystis* extracellular polymers, the total sugar measured did not match the total carbon measured, suggesting that the unaccounted carbon may have been protein-C or glycoprotein-C (M. van Rijssel pers. comm.). Protein has also been inferred to occur in *Phaeocystis* extracellular polymers based on ^1H MNR spectra (Aluwihare & Repeta 1999), whereby *Phaeocystis globosa* (CCMP 628) had small resonances from between 2.8 to 1.5 ppm, reflecting the N-H bond (Aluwihare et al. 1997). However, our MALDI-TOF-MS data suggest that the nitrogen in *Phaeocystis* extracellular polymers is not from proteins or peptides but from small nitrogen-containing compounds such as amino sugars.

The MALDI-TOF-MS data showed that the extracellular polymers are present as molecules <1000 Da in size. This is consistent with data of Aluwihare et al. (1997), who found that a major component of natural macromolecular dissolved organic matter consisted of a chain of 7 neutral sugars of approximately 1000 Da. The molecular weights found in the MALDI-TOF-MS data may reflect glycoproteins that are cleaved at glycosidic or cross-ring bonds (Harvey 1999). The compounds most likely to produce the XPS and MALDI spectra we obtained are amino sugars, gangliosides, or sialic acid. One of the larger fragments (i.e. 921 m/z) may be part of a chain of 7 neutral sugars, as suggested by Aluwihare et al. (1997), whereas the other fragments may be chains of acetylated sugars and amino sugars. Therefore, nitrogen is contained in amino groups located on the sugars rather than in proteins.

Possible factors contributing to differences in reported C:N ratios

Numerous differences between the present study and those of Biersmith & Benner (1998) and Aluwihare & Repeta (1999) may have contributed to the wide range of apparent C:N ratios of *Phaeocystis* extracellular polymers. Possible confounding factors are the use of different *Phaeocystis* strains or species, different light levels for growth, different growth stages, and the presence of bacteria. Biersmith & Benner (1998) grew *P. globosa* Strain CCMP 627 (from the Gulf of Mexico) at 20°C and ca. 100 $\mu\text{mol photons m}^{-2} \text{s}^{-1}$ in bacterized culture and measured a C:N ratio of 19 for extracellular polymers harvested during the stationary phase. Aluwihare & Repeta (1999) grew *P. globosa* Strain CCMP 628 (from Vineyard Sound, Massachusetts) at 26°C in bacterized culture at an unspecified light level and obtained a C:N ratio of 12 for polymers in a late

exponential culture. In our study, *P. globosa* Strain CCMP 629 (from the Gulf Stream) and *P. antarctica* (from the Ross Sea) were grown at 18 and 4°C, respectively, at ca. 113 $\mu\text{mol photons m}^{-2} \text{s}^{-1}$, and were harvested at late exponential stage; their C:N ratios were 6 to 8. Therefore, the discrepancy between the extracellular polymer C:N ratios recorded in different studies may be due to the use of different strains or species of *Phaeocystis* at different light levels or growth stages.

The membrane material of the ultrafiltration cartridge used for polymer concentration may also have affected the C:N ratios. We used regenerated cellulose membrane cartridges, whereas Biersmith & Benner (1998) and Aluwihare & Repeta (1999) used a polysulfone membrane cartridge (no longer manufactured). The different materials may have different absorption characteristics for carbohydrates, proteins and amino sugars. In general, polysulfone membrane absorbs more protein than regenerated cellulose (Cheryan 1998), and this could account for the higher C:N found in the previous studies. However, calibration studies with ultrafiltration cartridges using standard compounds do not portray a consistent picture about the absorption differences between the materials of the 2 cartridges. Dai et al. (1998) compared regenerated cellulose and polysulfone and found carbohydrate loss to the cartridge to be undetectable and 35%, respectively, whereas protein loss was 40 to 70% and 40%, respectively. Other calibration studies with 1000 Da polysulfone cartridges (Guo & Santschi 1996) have found lower absorption losses and better retention characteristics for Vitamin B-12 (80%) and for various proteins (between 93 and 97%). Aluwihare (1999) reported retention of $80 \pm 10\%$ for carbohydrate and 50% for protein. Retention of carbohydrate and protein of the cartridge used in our study was 53 and 14%, respectively, suggesting that more protein than carbohydrate was lost to the cartridge membrane. Calibration studies among polysulfone and regenerated cellulose cartridges may not be consistent but, overall, regenerated cellulose cartridges seem to absorb more protein than polysulfone cartridges. Thus, the lower C:N found in our study is not likely to have been due to preferential absorption of carbohydrates by the regenerated cellulose cartridge.

Influence of bacterial degradation on C:N ratios of extracellular polymers

Phaeocystis spp. extracellular polymers have been shown to effectively broken down by bacteria, but some fraction remains undegraded (Janse et al. 1999, Orellana et al. 2001). Since bacteria can degrade

Phaeocystis extracellular polymers, they could affect the C:N ratio of the recovered polymers during culture growth. We hypothesized that the high C:N ratios of *P. globosa* extracellular polymers found in previous studies (Biersmith & Benner 1998, Aluwihare & Repeta 1999) may have been due to the presence of bacteria in the experimental cultures. Even if only a low percentage of total dissolved organic carbon (DOC) is degraded, bacterial degradation could result in an increase in the C:N ratio of residual organic matter if relatively higher amounts of dissolved organic nitrogen are utilized. It is possible that the relatively high C:N ratios (>12) of *Phaeocystis* polymers reported in the previous studies may have been due to preferential dissolved organic nitrogen (DON) uptake by bacteria, even though bacterial biomass was <2% of the total DOC.

To test this, we compared the C:N ratios of recovered polymers in axenic and bacterized cultures. Although we expected to see a higher C:N in bacterized cultures, the C:N ratios were actually somewhat lower in the presence of bacteria. The greater decline in C:N for *Phaeocystis globosa* than for *P. antarctica* (8.3 to 7.3 vs. 6.8 to 6.2) may have been due to the larger standing stock and higher biomass production of bacteria. Bacterial degradation of diatom extracellular polymers has also reported to result in lowered C:N ratios (as low as 5.6: Aluwihare & Repeta 1999). How might bacteria decrease the C:N ratio? One possibility is that bacterial extracellular enzymes such as proteases and beta-glucosidases (Martinez et al. 1996, Vetter & Deming 1999) may be released during growth. Another possibility is that bacteria may have utilized the residual nitrate in the cultures and had a net excretion of DON. In our experiments, there was excess nitrate at the time of harvest, so it is unlikely that nitrate was limiting. In the Biersmith & Benner (1998) study, total DIN reached almost zero towards the end of the experiment (Biddanda & Benner 1997), so bacteria may have had a net uptake of DON, increasing the C:N ratios of the *Phaeocystis* spp. polymers. The contribution of bacteria to altering the C:N ratio of *Phaeocystis* extracellular polymers needs to be further investigated.

Conclusion

The C:N ratios of extracellular polymers produced by *Phaeocystis globosa* (CCMP 629) and *P. antarctica* during late exponential growth were relatively low, ranging from 6 to 8. XPS and MALDI-TOF-MS analyses showed that the nitrogenous component was not due to proteins but possibly to small nitrogen-containing compounds such as amino sugars. Other studies have also shown that amino acids comprise a small percent-

age of dissolved organic nitrogen in the high molecular weight pool (Mannino & Harvey 2000).

Surprisingly, the C:N ratio of *Phaeocystis* extracellular polymers in bacterized cultures was lower than that in axenic cultures. It is possible that released bacterial enzymes or excretion of DON by bacteria contribute to the organic nitrogen fraction of the recovered polymers. *P. antarctica* polymers are more nitrogen-rich than those of *P. globosa*, which may make them more biodegradable and may be a contributing reason to the low accumulation of DOC in the Ross Sea compared to the North Sea. In conclusion, freshly produced dissolved organic matter from phytoplankton may be a source of nitrogen as well as carbon for heterotrophic bacteria and protists. The nitrogenous component of low and high molecular weight DOM as well as the bacterial degradation of polymers need to be further investigated.

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