

Sugar dynamics in large particles during *in vitro* incubation experiments

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ABSTRACT: Large particles (>60 µm) were collected using *in situ* pumps deployed at 30 and 200 m depth at 2 typical stations in the southern Indian Ocean (Polar Front Zone [PFZ] and Sub-Antarctic Area [SAr]). The samples were incubated *in vitro* with their own bacterial assemblages for 7 to 17 days in batches under oxic conditions in the dark. Particulate organic carbon (POC) and sugars were monitored over time. Particulate sugars (PCHO-C) accounted for 5 to 20% of POC in the SAr, while they represented 5 to 8% of POC in the PFZ station at the beginning of the experiments. Molecular level analysis indicated that at the time of collection, ribose was among the most abundant sugars (18 to 30 wt% of PCHO-C) at the PFZ station and was rapidly degraded ($k = 0.051$ to 0.058 d⁻¹) over the course of the experiment, while this sugar was below detection limits in the SAr. Our results also showed an increase in the relative abundance of deoxysugars (fucose and rhamnose), suggesting that these sugars have the potential to be used as indicators of the bacterial activity and evaluate the degradation status of POM in both areas. The kinetic study indicated that pentoses were degraded faster than hexoses, while deoxysugars exhibited the lowest degradation rates. This study demonstrated that total sugar degradation rates do not reflect the rates of all individual components, but rather a disparate collection of rates among sugars classes and individual sugars, which very likely can vary significantly in relation to the origin of particles.

KEY WORDS: Sinking particles · *In vitro* incubation experiments · Sugar degradation · Southern Indian Ocean

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INTRODUCTION

Understanding the fate of particulate organic matter (POM) during microbial decomposition is important because marine aggregates represent the primary mode of particle flux to the ocean interior (Asper et al. 1992, Asper & Smith 1999, Honjo et al. 2000), while most of marine primary production is recycled within the euphotic zone (Karl et al. 1988, Falkowski et al. 1998). During sedimentation, consumption of POM by bacteria produces degraded particulate organic carbon (POC) and dissolved organic carbon (DOC); the latter are further mineralized by free-living bacteria that release CO₂ to the marine environment through respiration (Cho & Azam 1988, Karner & Herndl 1992, Smith et al. 1992, Hoppe et al. 1993, Sempéré et al.

2000). Earlier investigations indicated the importance of carbohydrates and proteins as bacterial substrates in POM and seawater (Smith et al. 1992, Keil & Kirchman 1993, Skoog et al. 1999, Panagiotopoulos et al. 2002). Although the previous studies have provided some information on the fates of carbohydrate and protein, little quantitative molecular-level information is currently available on the alteration of carbohydrate in POM by marine bacteria. Furthermore, very few studies have provided data on the degradation rates of the total carbohydrate pool, carbohydrate classes (i.e. hexoses, pentoses) or individual monosaccharides.

Carbohydrates are generally considered as less labile compounds than amino acids and are, therefore, useful indicators to evaluate the degradation status of organic matter (Cowie & Hedges 1984,

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Tanoue & Handa 1987, Hedges et al. 1994, Hernes et al. 1996, Opshal & Benner 1999, Panagiotopoulos & Sempéré 2005a). For example, an increase in the weight percentage (wt%) of deoxysugars associated with a decrease in wt% of glucose is indicative of progressively degraded organic matter (Hedges et al. 1994). The latter observation was made while comparing different organic matter forms, including coarse (>63 μm) and fine (<63 μm) POM, and ultrafiltered dissolved organic matter (UDOM). Similar patterns were also observed during decomposition of vascular plants (Opshal & Benner 1999), polysaccharides excreted by phytoplanktonic species (Giroldo et al. 2003), and algal-derived DOM (Amon et al. 2001). Currently we do not know if these patterns are similar during decomposition of sinking POM because there is a dearth of molecular carbohydrate data of any kind in any marine environment. In a recent study based on sediment trapped material, Panagiotopoulos & Sempéré (2005a) showed that a high relative abundance of glucose (30 to 35 wt%) in sinking particles does not necessarily reflect the presence of fresh material.

Therefore, in order to obtain more information from the carbohydrate perspective on sinking particle dynamics during microbial degradation, large particles were collected using *in situ* pumps (ISP) and were further subjected to biodegradation experiments as previously described (Sempéré et al. 2000, Panagiotopoulos et al. 2002). This sequel study of our work on biodegradation experiments in the southern Indian Ocean (Panagiotopoulos et al. 2002) aims to: (1) contrast combined neutral sugar composition in sinking POM collected from different marine environments, notably the Polar Front Zone (PFZ) and the Sub-Antarctic Area (SAr), both located in the southern Indian Ocean; (2) examine whether wt% of glucose and deoxysugars can be used as tracers of biological activities during

decomposition; and (3) provide and compare degradation rates for individual sugar constituents during decomposition.

MATERIALS AND METHODS

Large particles (>60 μm) were collected in January and February 1999 in the southern Indian Ocean at 2 stations: the Polar Front Zone (PFZ ~ 46° S, 62° E) and the Sub-Antarctic Area (SAr ~ 44° S, 63° E). General features of these sites are described in Panagiotopoulos et al. (2002). Detailed information about the study sites, including primary production, bacterial production, and total organic carbon (TOC) stocks, is presented in Table 1. Particles were collected at 30 and 200 m depth using ISP (Challenger Oceanics) equipped with 60 μm pore-size filters (Nitex, \varnothing : 142 mm). These sampling depths (within and beneath the euphotic zone) were chosen to represent freshly produced and slightly degraded organic material, respectively.

Seawater (up to 10 l) was also collected at the same depths and filtered through 0.2 μm pore-size filters (Millipore, \varnothing : 142 mm; pressure <50 mm Hg) in order to remove bacteria and grazers (Sempéré et al. 2000, Panagiotopoulos et al. 2002). This water was used to wash out particles from filters and for batch preparation. Extracted particles were likewise concentrated into a 2 l polycarbonate bottle and immediately transferred to 4 to 5 incubation bottles (500 ml each) using a peristaltic pump. Each bottle corresponds to the time-point of the incubation (Table 2). This material was incubated at *in situ* seawater temperature (5 to 14°C) under oxic conditions and in the dark (Table 1). The headspace-to-sample ratio inside the incubation bottles was about 1:1, which ensured oxic conditions over the course of the incubation (7 to 17 d; longer incuba-

Table 1. Summary of particle sampling conditions and *in situ* seawater characteristics. Large particles (>60 μm) were collected using *in situ* pumps (ISP) deployed at 30 and 200 m at 2 stations in the southern Indian Ocean: Polar Front Zone (PFZ; ~46° 01.22' S, 62° 56.70' E) and Sub-Antarctic Area (SAr; ~44° 10.92' S, 63° 23.37' E). Euphotic zone depth was 150 m at both stations. Primary production (PP), bacterial production (BP), and total organic carbon (TOC) are integrated values to 1% of light penetration

	Sampling depth (m)	ISP filtered volume (l)	Seawater temp. (°C)	PP (mmol C m ⁻² d ⁻¹) ^a	BP (mmol C m ⁻² d ⁻¹) ^{b,c}	TOC (mmol C m ⁻²) ^b
PFZ	30	7471	9	28	8	7474
	200	9906	5			
SAr	30	12068	14	47	13	9642
	200	8013	12			

^aLeblanc et al. (2002)
^bR. Sempéré (unpubl. data)
^cVan Wambeke (pers. comm.)

Table 2. Elemental carbohydrate compositions (weight percent of particulate combined aldoses) and particulate combined aldose (PCHO-C) concentrations during biodegradation of (>60 μm) particles collected with *in situ* pumps in the Polar Front Zone (PFZ) and Sub-Antarctic Area (SAr). The PCHO-C yields are the percentage of sugar-C to the POC pool (PCHO-C/POC). Fuc: fucose, Rha: rhamnose, Ara: arabinose, GlcN: glucosamine, Gal: galactose, Glc: glucose, Man: mannose, Xyl: xylose, Fru: fructose, Rib: ribose

Sampling depth (m)	Incubation period (d)	Fuc	Rha	Ara	GlcN	Gal	Glc	Man	Xyl	Fru	Rib	PCHO-C (mg l ⁻¹)	PCHO-C/POC (%)
PFZ													
30	0	3.59	2.05	2.38	0.98	19.9	22.6	5.38	8.78	4.92	29.5	0.49	4.64
	1	3.67	2.38	2.24	1.75	18.1	19.0	5.92	8.40	3.95	34.6	0.38	5.61
	2	3.36	3.18	0.87	2.27	16.9	22.0	6.07	6.85	4.84	33.6	0.40	4.55
	4	3.48	3.82	1.09	3.82	17.1	16.0	3.95	7.59	4.98	38.2	0.30	3.40
	7	5.63	3.97	2.30	3.15	21.0	24.5	5.62	8.76	5.30	19.7	0.40	4.96
	13	5.34	6.67	0.57	2.10	22.0	24.2	6.05	10.7	7.66	14.7	0.39	6.61
200	0	3.04	1.26	–	10.5	16.2	31.9	7.17	7.72	4.11	18.1	0.33	7.62
	1	3.58	1.66	–	8.47	19.8	28.3	6.30	13.1	8.86	9.98	0.31	8.89
	4	4.74	2.85	–	14.8	19.4	23.0	8.47	8.77	5.13	12.8	0.36	9.01
	9	7.10	5.69	–	12.1	24.6	34.3	6.86	9.34	–	–	0.25	6.32
	17	10.0	5.80	–	9.08	24.2	26.6	9.95	14.4	–	–	0.23	7.72
SAr													
30	0	4.27	3.52	1.02	0.83	21.6	54.5	5.64	8.62	–	–	1.10	19.9
	1	3.98	3.40	1.09	1.68	21.7	51.0	7.11	10.1	–	–	1.39	23.8
	2	4.76	3.49	1.12	1.43	23.9	48.8	6.56	9.97	–	–	0.81	16.4
	4	5.23	4.58	1.33	2.55	26.6	39.4	8.20	12.1	–	–	1.05	20.1
	7	5.74	5.88	1.01	2.03	25.7	42.6	6.80	10.3	–	–	0.62	13.8
	14	8.17	7.25	0.46	0.69	26.2	37.4	7.32	12.6	–	–	0.50	12.0
200	0	4.25	3.44	1.56	0.99	12.9	57.4	6.62	12.9	–	–	0.101	5.08
	1	5.07	1.96	0.88	2.36	13.3	56.6	6.56	13.2	–	–	0.079	5.88
	2	4.04	3.16	1.53	2.78	12.8	60.5	4.48	10.8	–	–	0.098	8.61
	4	3.12	2.66	1.48	2.50	9.75	69.4	4.00	7.13	–	–	0.117	9.10
	7	5.76	4.37	2.52	2.16	14.7	55.9	4.05	10.5	–	–	0.065	4.19

tion times would probably induce anoxic conditions). Incubation bottles were also gently and regularly swirled upside down to homogenize their contents and avoid particle deposition on the walls.

Biodegradation experiments were only conducted once at each depth due to a limited amount of particulate material. Sub-sampling of incubations was performed regularly by collecting aliquots for POC and particulate sugars (PCHO-C) in precombusted glass bottles (450°C, 6 h). Particles were collected on GF/F filters (nominative retention, 0.7 μm) and stored in the dark at –17°C. Incubation sub-samples were analyzed using previously described methods for POC (Tan & Strain 1979) and neutral sugars (Panagiotopoulos et al. 2001, Panagiotopoulos & Sempéré 2005a,b). At the beginning of the incubation, the standard deviation for particle distribution was 5% (POC measurements in 5 replicates). During the course of the incubations experimental errors, based on duplicate analysis of sub-samples, were estimated to be about 10% for the concentration of each chemical variable (POC and PCHO). A 2-fold replicate analysis of individual monosaccharides had a standard deviation of <8% for all sugars except ribose (15%).

Particulate sugar degradation rates. Particulate sugar degradation rates (k) were calculated according to the formula (Amon & Benner 1996):

$$k = (C_{\text{init}} \times C_{\text{fin}}) \times (t)^{-1} \times (C_{\text{init}})^{-1} \quad (1)$$

where C_{init} and C_{fin} are the initial and final concentration ($\mu\text{g l}^{-1}$) of any constituent, respectively, and t is the incubation time (days). To analyze the relative proportion of the particulate fraction of each individual sugar that is available for bacterial consumption, all rates were normalized to the initial concentration of the incubation (Panagiotopoulos et al. 2002).

RESULTS

In situ PCHO-C concentrations

In situ PCHO-C concentrations in large (>60 μm) particles from 30 m depth at both PFZ and SAr stations ranged from 14 to 18 ng C l^{-1} and decreased to a range of 6 to 8 ng C l^{-1} at 200 m. These values are roughly 6 orders of magnitude lower than those measured at the beginning of the biodegradation experiment (Table 2).

Table 3. Degradation rates (k) of particulate combined aldoses (PCHO-C), hexoses, pentoses, deoxysugars, and individual monosaccharides calculated according to Eq. (1) (see text). Units in $(\mu\text{mol C l}^{-1} \text{d}^{-1})/(\mu\text{mol C}_{\text{init}} \text{l}^{-1})$. Abbreviations of sugars as in Table 2

Sampling depth (m)	$k_{\text{PCHO-C}}$	k_{hexoses}	k_{pentoses}	$k_{\text{deoxysugars}}$	k_{Fuc}	k_{Rha}	k_{Ara}	k_{GlcN}	k_{Gal}	k_{Glc}	k_{Man}	k_{Xyl}	k_{Fru}	k_{Rib}
PFZ														
30	0.016	0.021	0.044	-0.032	0.001	-0.090	0.065	0.023	0.020	0.022	0.019	0.015	0.018	0.051
200	0.018	0.016	0.037	-0.084	-0.070	-0.120	-	0.025	0.001	0.026	0.005	0.025	0.057	0.058
SAr														
30	0.038	0.041	0.024	0.008	0.004	0.014	0.055	0.042	0.029	0.047	0.026	0.020	-	-
200	0.049	0.047	0.054	0.013	0.009	0.018	0.001	0.028	0.030	0.047	0.083	0.063	-	-

Assuming a sinking velocity of 100 m d^{-1} (Alldredge & Silver 1988), these *in situ* PCHO-C concentrations correspond to a theoretical sugar flux at 200 m of $0.83 \text{ mg C m}^{-2} \text{ d}^{-1}$ for SAr and $0.64 \text{ mg C m}^{-2} \text{ d}^{-1}$ for PFZ, which is in the same order of magnitude as the fluxes estimated from the sediment traps (Panagiotopoulos & Sempéré 2005a).

Concentrations, relative abundances, and degradation rates of PCHO-C

In the PFZ biodegradation experiments, PCHO-C concentrations decreased by 20 and 30% from their initial concentration at 30 and 200 m, respectively, while the corresponding decreases at SAr were 54 and 36% at 30 and 200 m (Table 2). The PCHO-C yields (defined as PCHO-C/POC ratios) increased slightly for PFZ particles from 5 to 7% at 30 m and remained constant (~8%) at 200 m over the course of the experiment (Table 2). In contrast, in the SAr experiment, PCHO-C yields at 30 and 200 m depths decreased from 20 to 12% and from 5 to 4%, respectively, over the course of the experiment.

The initial sugar composition at SAr varied only slightly between the 2 depths, with the exception of galactose (Table 2). At SAr, the collected particulate material was characterized by the dominance of glucose (55 and 57 wt% at 30 and 200 m depth, respectively) followed by galactose (22 and 13 wt% at 30 and 200 m) and xylose (9 and 13 wt% at 30 and 200 m). The remaining sugars contributed <7 wt% to the carbohydrate pool. It is important to note that ribose and fructose were below detection limits, indicating the absence of storage or intracellular polysaccharides in SAr particles at $t = 0$. In contrast, PFZ particles at 30 m at $t = 0$ were characterized by the dominance of ribose (30 wt%) followed by glucose (23 wt%), galactose (20 wt%), and xylose (9 wt%). For PFZ particles at 200 m, glucose was initially the major aldose (~32 wt%), followed by ribose (18 wt%), galactose (16 wt%), and xylose (8 wt%) (Table 2).

In all experiments for both stations, major decreases in the concentration ($\mu\text{g C l}^{-1}$) of pentoses (arabinose, xylose, and ribose) and hexoses (mannose, glucose, and galactose) were observed. Conversely, the concentrations of deoxysugars (fucose and rhamnose) increased at PFZ and remained stable at SAr. The relative abundances of mannose, arabinose, xylose, fucose, and rhamnose increased or remained constant, while those of glucose and ribose always decreased over time at both stations, suggesting a selective extraction of these sugars from POM.

Calculated values of k for PFZ and SAr stations at both depths were 0.016 to 0.049 d^{-1} for PCHO-C and 0.001 to 0.083 d^{-1} for individual monosaccharides. At the PFZ station, negative degradation values were recorded for deoxysugars due to the fact that concentrations of these sugars (sum of fucose and rhamnose) increased over time (Table 3). This result was surprising since Amon et al. (2001) found that fucose and rhamnose generally remained constant (or increased slightly) during a short incubation period. Therefore, for the remainder of this investigation the negative degradation values for rhamnose, fucose, and deoxysugars at PFZ (30 and 200 m) were excluded.

DISCUSSION

PCHO-C yields and relative abundance of combined aldoses in ISP

At the beginning of the biodegradation experiments, particulate combined aldoses comprised <8% of the POC pool (except SAr particles at 30 m; Table 2). These results are in agreement with earlier investigations carried out in sediment traps at other locations including the central equatorial Pacific Ocean (5 to 18%, Hernes et al. 1996), North Pacific (7 to 10%, Tanoue & Handa 1987), Sargasso Sea (3 to 6%, Ittekkot et al. 1984), southern Indian Ocean (3 to 4%, Panagiotopoulos & Sempéré 2005a), and Mediterranean Sea (4 to 16%, Kerhervé et al. 1999,

Panagiotopoulos & Sempéré 2005a). It is important to note that at $t = 0$, amino acids were predominant in PFZ particles, accounting for ~38% of POC, while they represented only ~15% of POC in SAR particles (Panagiotopoulos et al. 2002). Since amino acids are indicative of fresh algal detritus and are more rapidly degraded than other biochemical classes (Harvey et al. 1995, Harvey & Mannino 2001), their high relative abundance at PFZ suggests that particles initially collected were less degraded than in SAR waters. Therefore, the low initial PCHO-C yields measured at PFZ (5%) compared with those at SAR (20%) imply the presence of fresh material at the PFZ station (Table 2).

Our results also showed that at $t = 0$ ribose was the major aldose at PFZ at 30 m, but was second in abundance at PFZ at 200 m. These results are comparable with those reported in sinking particles in the same area using sediment traps (Panagiotopoulos & Sempéré 2005a). The difference in the relative abundance of sugars collected in sediment traps (Panagiotopoulos & Sempéré 2005a) and ISP (present study) may be due to the fact that ISP collect a wide spectrum of particle sizes (small and large), while sediment traps collect only fast sinking particles. It is also likely that small particles absorbed in the *in situ* pump filters are more degraded than large sinking particles collected with sediment traps. However, the high abundance of ribose in PFZ particles at $t = 0$ indicates the presence of non-structural and labile compounds, which might be further degraded rapidly by bacteria (Table 2). This result corroborates our previous observations suggesting that sinking particles in PFZ waters were less degraded than those in SAR waters. Finally, glucose was the major aldose (>32 wt%) at SAR station followed by galactose and xylose (Table 2). Similar results have also been reported for sediment trap samples in other oceanic areas (Tanoue & Handa 1987, Hernes et al. 1996, Panagiotopoulos & Sempéré 2005a).

PCHO-C dynamics during bacterial degradation

With the exception of PFZ particles at 200 m, in which we observed an accumulation of sugars in the total dissolved fraction, dissolved sugar (DCHO-C) concentrations closely followed the decrease of PCHO-C, implying a rapid bacterial assimilation of dissolved monosaccharides or oligosaccharides produced during POC decomposition. We did not analyze sugars in the dissolved fraction by chromatography, only the total amount of dissolved sugars is measured by colorimetric techniques (see Panagiotopoulos et al. 2002). Nevertheless, our results indicated comparable de-

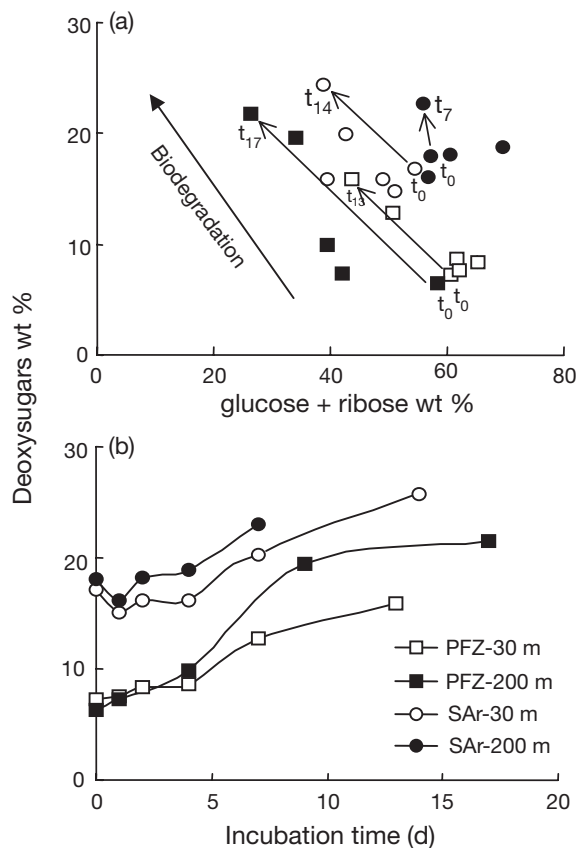


Fig. 1. (a) Plot of weight percentage (wt%) of deoxysugars (fucose plus rhamnose) vs. wt% of glucose plus ribose and (b) wt% of deoxysugars during biodegradation experiments. Weight percentages of deoxysugars and ribose were calculated on a glucose-free basis

creases in both fractions (PFZ at 30 m, 21% for PCHO-C and 17% for DCHO-C; SAR at 30 m, 54% for PCHO-C and 43% for DCHO-C; SAR at 200 m, 34% for PCHO-C and 37% for DCHO-C), indicating that PCHO-C concentrations in batches are likely to better represent the decay of sugars in POM.

The 2 most striking changes in composition of neutral sugars during biodegradation were the rapid decrease in the wt% of glucose and ribose, and the increase in wt% of deoxysugars (fucose and rhamnose; Fig. 1a). Glucose is considered to be a labile compound of storage polysaccharides (β 1-3 glucanes), while ribose is a common component of RNA and nucleotides (Haug & Myklestad 1976, Cowie & Hedges 1984, Hicks et al. 1994). The high decrease of glucose and ribose and the relative increase of mannose, xylose, and galactose over time clearly suggests a selective extraction of these labile sugars (glucose and ribose) from the PCHO-C pool and is in agreement with previous investigations (Tanoue & Handa 1987, Hernes et al. 1996).

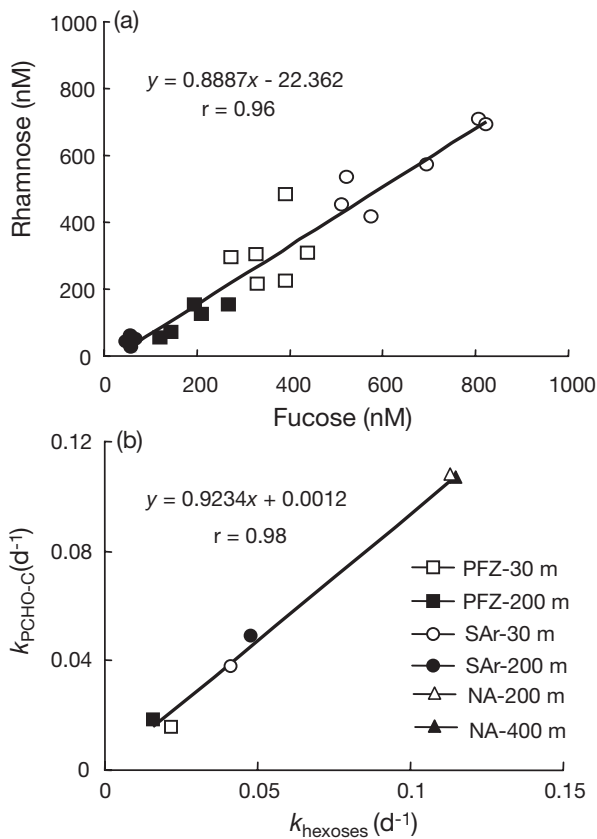


Fig. 2. (a) Correlation between fucose and rhamnose at 30 and 200 m depth, for the Polar Front Zone (PFZ) and Sub-Antarctic (SAR) stations during biodegradation experiments. (b) Correlation between degradation rates (k) of PCHO-C and hexoses. We included some of our unpublished data from other biodegradation experiments in the north Atlantic Ocean (NA at 200 and 400 m depth) performed in the same way as the present experiments

The relative increase of deoxysugars during decomposition has already been reported for several organic matter types and environments (Cowie & Hedges 1984, Hedges et al. 1994, Opsahl & Benner 1999, Amon et al. 2001, Girollo et al. 2003) and is often attributed to the high abundance of fucose and rhamnose in bacteria (Ogier et al. 2001). Our results indicated that particles at the PFZ station were initially rich in amino acids (Panagiotopoulos et al. 2002) and ribose, and relatively poor in deoxysugars (6.3 to 7.3 wt%), while the opposite observation was found at SAR (17.1 to 18.0 wt% deoxysugars; Fig. 1b). This result reinforces the idea that elevated relative abundance of deoxysugars is found in highly degraded material. As Fig. 2b indicates, after 13 to 17 d of decomposition deoxysugars wt% values in PFZ particles reached those of SAR particles (SAR at $t = 0$), which also suggests a difference in degradation status between PFZ and SAR particles. These results illustrate that deoxysugars have the

potential to be used as indicators of the bacterial activity and to evaluate the degradation status of POM. The high abundance of fucose and rhamnose in bacteria, and therefore in POM during decomposition, is further confirmed by a highly significant correlation ($r = 0.96$; $p < 0.01$; $n = 22$) between these deoxysugars in the collected particles (Fig. 2a).

Kinetics of PCHO-C and individual monosaccharides during bacterial degradation

Although the literature contains a plethora of values on degradation rates of bulk organic matter, including sinking particles, sediments, and phytoplankton, few studies provide data on PCHO-C degradation rates (Westrich & Berner 1984, Harvey et al. 1995, Rabouille et al. 2000, Sempéré et al. 2000, Panagiotopoulos et al. 2002). Furthermore, little data exist for the degradation rates of sugar classes and individual monosaccharides. In this study, degradation rates of PCHO-C ranged from 0.016 to 0.049 d⁻¹, resulting in a residence time of 20 to 63 d. Using a 1G model in a flow-through system during phytoplankton decay Harvey et al. (1995) reported degradation rates of PCHO-C ranging from 0.092 to 0.094 d⁻¹ for an incubation period of 45 to 77 d. However, if these rates were calculated according to Eq. (1) they would range from 0.013 to 0.021 d⁻¹, which is in the same order of magnitude as the present study. Similarly, Girollo et al. (2003) applied a 1G model during microbial degradation of extracellular polysaccharide and observed sugar degradation rates in the order of 0.050 d⁻¹ for an incubation period of 35 d. However, if their results were calculated in the same manner as our study they would correspond to a degradation rate of 0.025 d⁻¹, again in agreement with this study.

Therefore, during decomposition of fresh organic matter (sinking particles, phytoplankton, extracellular polysaccharides) particulate sugars have a residence time of 20 to 77 d, or roughly 1 to 3 mo, indicating that these compounds are major components of semilabile DOC. It is worth noting that these residence time values were always higher than those of amino acids, but comparable with or lower than lipids, which is consistent with the initial chemical composition reported for phytoplankton and fast-sinking particles (20 to 60% amino acids, 10 to 15% lipids, and 2 to 15% sugars; Harvey et al. 1995, Wakeham et al. 1997, Panagiotopoulos et al. 2002, Panagiotopoulos & Sempéré 2005a). This further suggests that the mechanisms that control particle degradation also depend on their initial chemical composition (Wakeham et al. 1997, Panagiotopoulos et al. 2002).

Degradation rates of sugar classes (hexoses, pentoses, and deoxysugars) ranged from 0.008 to 0.054 d⁻¹,

resulting in a residence time of 19 to 125 d. The highest residence time corresponded with deoxysugar degradation (SAr at 30 and 200 m), while the lowest time with that of hexoses (mannose and glucose) and some pentoses (ribose). As previously indicated little current data exist for sugar classes and individual monosaccharides degradation rates in the literature. Therefore, the present data cannot be compared with values existing for sugar degradation rates. However, we can note from literature that degradation rates of some lipid classes (free fatty acids, sterols and triacylglycerols) during fresh algal decomposition fall in the range of 0.026 to 0.092 d⁻¹ (Harvey & Macko 1997, Caradec et al. 2004). Again if these values were calculated according to Eq. (1) they would range from 0.011 to 0.022 d⁻¹, which is in agreement with the present study (Table 3).

Pentoses also degraded faster than hexoses, and deoxysugars exhibited the slowest degradation rates (Table 3). This result was somewhat surprising, since we observed a strong correlation ($r = 0.98$; $p < 0.01$; $n = 6$) between degradation rates of PCHO-C and hexoses. This indicates that sugar degradation rates are controlled mainly by the degradation of hexoses, which are generally more abundant than pentoses in sinking particles (Hernes et al. 1996, Panagiotopoulos & Sempéré 2005a). A weaker correlation ($r = 0.88$; $p = 0.01$; $n = 6$) was observed between PCHO-C and pentoses degradation rates. Therefore, it seems likely that degradation rates of very labile compounds such as ribose, which in some cases may be dominant sugars in POM (e.g. PFZ particles at 30 m), are not directly reflected in the total carbohydrate pool degradation rate.

Individual sugar degradation rates were higher at the SAr station than at the PFZ, which corroborates the bulk degradation PCHO-C rates (Table 3). This further suggests that sugars were more efficiently degraded in SAr than in PFZ sinking particles, in agreement with their initial higher relative abundance at SAr. With the exception of SAr particles at 200 m, glucose degradation rates were always higher than those of galactose and mannose. Galactose and mannose are derived from structural polysaccharides (i.e. galactans, mannan, galactomannans), while glucose comes from both structural and storage compounds of marine phytoplankton (Hecky et al. 1973, Haug & Myklesstad 1976, Tanoue & Handa 1987). The higher degradation rates of glucose compared with mannose and galactose clearly suggest the preferential use of glucose storage compounds and is in agreement with previous investigations (Hernes et al. 1996, Panagiotopoulos & Sempéré 2005a).

The present data clearly demonstrate that total sugar degradation rates do not reflect the rates of all individ-

ual components but rather a disparate collection of rates among sugars classes and individual sugars, which can vary significantly. Additional studies are needed to examine the bacterial degradation of POM, specifically to focus on other sugar classes or individual compounds such as amino sugars and uronic acids.

Acknowledgements. We thank the captain and crew of the RV 'Marion Dufresne' for excellent service at sea as well as J. Le Fèvre (leader of the Antares project) and M. Denis (chief scientist of Antares-4 oceanographic cruise). We thank C. Jeandel, C. Bournot-Marec, and L. Coppola for assistance in sample collection and batch preparation. We also thank R. Lafont, L. Ziolkowski, and the 2 anonymous reviewers for valuable comments and suggestions. This research was supported by France-PROOF Antares-4 project. Financial support for C.P. came from the Alexander S. Onassis Public Benefit Foundation, Greece.

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Editorial responsibility: Otto Kinne (Editor-in-Chief), Oldendorf/Luhe, Germany

*Submitted: October 25, 2005; Accepted: June 21, 2006
Proofs received from author(s): January 10, 2007*