Vol. 39: 7-16, 2005

# Seasonal dynamics of bacterial growth efficiencies in relation to phytoplankton in the southern North Sea

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ABSTRACT: The main function of heterotrophic bacterioplankton in marine carbon cycling is the conversion of dissolved organic carbon (DOC) into biomass and  $CO_2$ . The relative importance of bacterial biomass production (BP) versus respiration (BR) is expressed by the bacterial growth efficiency (BGE = BP/(BP + BR) × 100). Studies on the dynamics of the BGE of bacterioplankton growing on natural DOC covering entire seasonal cycles are scarce. We measured BP and BR over a seasonal cycle in the southern North Sea at a total of 150 stations to determine seasonal variability in BGE. While BP varied over 1 order of magnitude over the seasonal cycle, BR varied only 2-fold. Cell-specific BP was related to primary production while BR was not. Mean BGE increased from  $6 \pm 3\%$  in the winter to  $25 \pm 9\%$  in the spring and summer. Depth-integrated BR was fairly stable over the seasonal cycle, averaging 57% of the particulate primary production. Based on the bacterioplankton respiration and the mean annual BGE of 20%, bacterioplankton organic carbon demand amounts to ~70% of the particulate primary production in the southern North Sea, suggesting that autochthonous organic matter production is sufficient to fuel bacterioplankton carbon demand.

KEY WORDS: Bacterioplankton  $\cdot$  Bacterial production  $\cdot$  Bacterial respiration  $\cdot$  Bacterial growth efficiency  $\cdot$  Primary production  $\cdot$  North Sea

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## INTRODUCTION

Bacterioplankton represent the largest living biomass in the world's ocean (Whitman et al. 1998). Their main ecological function is to sequester the dissolved organic carbon (DOC) pool by incorporating a fraction of it into biomass and remineralizing the major fraction of DOC to  $\rm CO_2$ . Depending on the trophic state of the environment, bacterioplankton contribute between 12 and >50 % to total community respiration (Robinson et al. 1999, 2002b, Søndergaard et al. 2000).

Whether high remineralization rates of the prokaryotic community render the vast open-ocean into a net heterotrophic system is currently under debate (Duarte & Agusti 1998, Williams 1998, Del Giorgio & Duarte 2002, Hansell et al. 2004). While bacterial production (BP) is routinely measured in marine ecosystem studies, bacterial respiration (BR) has only recently received adequate attention. Determining BR rates, however, is as important as BP to estimate the bacterial growth efficiency (BGE =  $BP/(BP + BR) \times$ 100) and the bacterial carbon demand. Reported BGEs span over a large range from <1 to 80%. The underlying reasons for this large variation are basically unknown (Del Giorgio & Cole 2000). Jahnke & Craven (1995) suggested that the variability in the reported BGEs is due to the limited data set on respiration and concomitantly performed production measurements. Del Giorgio et al. (1997) argued that BGEs should range from <10 to 25 % in most marine surface systems. Attempts to derive respiration rates from BP measurements are problematic because the correlation between BP and BR is usually weak (Del Giorgio & Cole 1998). Rivkin & Legendre (2001) found a significant relationship between BGEs and temperature using a data set across a large temperature

range; this makes it difficult to apply their model to specific locations with smaller temperature fluctuations (Robinson et al. 2002a).

Shelf seas such as the North Sea cover only around 7% of the global ocean surface but contribute  $\sim 30\%$  to the oceanic primary production (Gattuso et al. 1998). Most studies on BP and BR are single surveys and surprisingly few marine studies have been performed over a seasonal cycle (Sherry et al. 1999, Lemme et al. 2002).

In this study, we measured BP, BR and primary production during 6 cruises covering a full seasonal cycle in the non-stratified southern North Sea occupying a total of 150 stations. The ultimate aim of the study was to determine the seasonal relationship between BP and BR and to relate the BGE dynamics to primary production and other physico-chemical variables.

#### MATERIALS AND METHODS

**Study site.** The study was carried out during 7 cruises aboard the RV 'Pelagia' in the unstratified waters of the southern North Sea (Fig. 1), occupying at total of 150 stations between July 2000 and August 2002. The mean depth of the southern North Sea is around 30 m and strong tidal currents and relatively high turbidity are common (Otto & Zimmermann 1990).

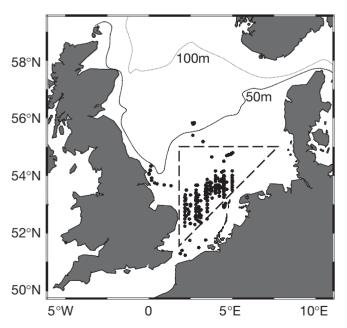


Fig. 1. Study area. Individual stations ( $\bullet$ ) occupied during the 7 cruises; only those stations enclosed by the dashed triangle were considered in the present analysis. Continuous and dotted lines mark 50 and 100 m isolines, respectively

Sample preparation for assessing bacterial parameters. For the parameters measured in unfiltered seawater, samples were taken at 5 and 15 m depth as well as 5 m above the bottom with 10 l NOEX bottles mounted on a CTD rosette. Part of the water collected was drawn under gentle vacuum (<-200 mbar) and filtered through 0.8 µm polycarbonate filters (Millipore ATTP; 47 mm diameter) to remove most of the nonbacterial particles. To avoid clogging, filters were changed as soon as the flow rate decreased to approximately half the initial rate. Subsequently, the 0.8 µm filtrate was used for bacterial production (BP) and respiration (BR) measurements and for bacterial abundance determination. Bacterial abundance and BP were also determined in unfiltered seawater. All sample handling was performed at in situ temperature (±2°C). Hydrographic properties of the study area are described in detail elsewhere (Baars et al. 2003). Additionally, samples were collected for chlorophyll a (chl a), phytoplankton production and dissolved organic carbon (DOC) as described below.

**Bacterial abundance.** We fixed 5 ml samples with 37% formaldehyde (4% final conc.), and subsequently determined bacterial abundance by 4′,6-diamidino-2-phenylindole (DAPI) staining and epifluorescence microscopy (Porter & Feig 1980) upon return to the laboratory within 1 wk. Since the abundance of Archaea in the coastal North Sea is generally low (Pernthaler et al. 2002), DAPI-stained cells with distinct cell boundaries were considered to be bacteria.

Bacterial production. BP for the unfiltered and 0.8 µm filtered seawater was measured by <sup>14</sup>C-leucine (leu) incorporation (specific activity 0.295 Ci mmol<sup>-1</sup>; 10 nmol l<sup>-1</sup> final conc.); 2 samples and 1 blank were incubated in the dark. The blank was fixed with formaldehyde (4% final conc., v/v) 10 min prior to adding the tracer. After incubating the samples and the blank at *in situ* temperature for 60 min, the samples were fixed with formaldehyde (4 % final conc.), filtered onto 0.2 µm nitrocellulose filters (Millipore HA; 25 mm diameter) and rinsed twice with 5 ml ice-cold 5% trichloroacetic acid (Sigma Chemicals) for 5 min. The filters were dissolved in 1 ml ethylacetate and, after 10 min, 8 ml scintillation cocktail (Insta-Gel Plus, Canberra Packard) was added. The radioactivity incorporated into bacterial cells was counted in a liquid scintillation counter (LKB Wallac, Model 1212). Leucine incorporated into bacterial biomass was converted to carbon production using the empirical conversion factor  $0.07 \times 10^{18}$  cells mol<sup>-1</sup> leu (Riemann et al. 1990) and assuming a bacterial carbon content of 20 fg C cell<sup>-1</sup> (Lee & Fuhrman 1987). The application of this conversion factor resulted in BP estimates similar to the theoretical factor of 1.55 kg C mol<sup>-1</sup> leu, assuming no isotope dilution (Simon & Azam 1989) (data not shown).

Bacterial respiration. Part of the 0.8 µm filtrate was carefully transferred to calibrated borosilicate glass BOD bottles with a nominal volume of 120 ml by a sipper system to avoid the formation of air bubbles. For the determination of the initial  $O_2$  concentration  $(t_0)$ , samples were fixed immediately with Winkler reagents and incubated together with the live samples in a water bath in the dark at in situ temperature (±1°C) for 12 to 24 h before the incubations were terminated  $(t_1)$ . Triplicate bottles were used for the determination of the initial and final O2 concentration. All glassware was washed with 10% HCl and thoroughly rinsed with Milli-Q water prior to use. Sample handling and fixation followed the recommendations of Carrit & Carpenter (1966). Oxygen concentrations of corresponding  $t_0$  and  $t_1$  bottles were measured in one run. The amount of total iodine was determined spectrophotometrically at a wavelength of 456 nm on a Hitachi U-1100 spectrophotometer using a 1 cm flowthrough cuvette at 20°C (Pai et al. 1993, Roland et al. 1999). To increase the sensitivity of the absorbance readings, a 4-digit voltmeter (Metex M4650) was connected to the spectrophotometer. Calibration was performed by standard additions of iodate to distilled water, resulting in an empirical coefficient of 0.54455 nM cm<sup>-1</sup> (Kraay pers. comm.). The samples were withdrawn from the BOD bottles with a Teflon tube and a peristaltic pump (Gilson Minipuls) and directly fed to the flow-through cuvette of the spectrophotometer. The end of the tube was placed near the bottom of the bottles to avoid loss of volatile iodine. The spectrophotometer was zeroed against Milli-Q water. The coefficient of variation of the oxygen determinations was < 0.5 %. To convert oxygen consumption into carbon we used a respiratory quotient of 1.

**Chlorophyll a determination.** We gently filtered 1 l water samples collected from 3 depths (5 m, 15 m, and 5 m above the bottom) through 47 mm Whatman GF/F filters and stored these at  $-60^{\circ}$ C until analysis (within 4 wk). Chl a was extracted in 10 ml of 90% acetone at  $-20^{\circ}$ C in the dark for 48 h. Subsequently, the filters were sonicated on ice for 1 min (Branson, Model 3200) and centrifuged to remove particles. The chl a concentration in the supernatant was determined fluorometrically with a Hitachi F-2000 fluorometer (Holm-Hansen et al. 1965).

**Primary production (PP) measurements.** For particulate PP measurements, the protocol of Gieskes et al. (1979) was followed. In brief, before sunrise, seawater was collected from 5 m and transferred into 250 ml polycarbonate bottles, and 10  $\mu$ Ci of <sup>14</sup>C bicarbonate was added to each sample. Subsequently, the samples were placed in 6 tubes of different light transmittance using neutral-density filters. The tubes were held at surface water temperature (±1°C) with a flow-through

seawater system. Simulated light intensities ranged from 85 to 0.6% of the surface irradiance. The incubations were run for 24 h; thereafter, the samples were filtered onto Whatman GF/F glass-fiber filters and fumed over HCl for 3 h. The filters were stored at –20°C and counted in an LKB Wallac liquid scintillation counter after adding 10 ml of Instagel II (Packard Canberra). Dark incorporation of <sup>14</sup>C bicarbonate was subtracted from the incubations in the light. Combining light-attenuation measurements of the water column with the primary production measurements performed under the different light regimens allowed us to calculate integrated PP over the water column.

DOC measurement. Samples for DOC were filtered through rinsed 0.2  $\mu$ m polycarbonate filters and sealed in pre-combusted (450°C for 4 h) glass ampoules after adding 50  $\mu$ l of 40% phosphoric acid. Subsequently, the samples were stored frozen at –20°C. DOC concentrations were determined by the high-temperature combustion method using a Shimadzu TOC-5000 analyzer (Benner & Strom 1993). Standards were prepared with potassium hydrogen phthalate (Nacalai Tesque). Ultrapure Milli-Q blanks were run before and after the sample analysis. The blank was on average 16.3  $\pm$  6.8  $\mu$ M, and the mean of triplicate injections was calculated for each sample. The average analytical precision of the instrument was <3%.

Calculations and statistical analysis. As no significant differences were detectable between the different stations occupied during the individual cruises, individual parameters from each cruise were pooled. To relate bacterial biomass to phytoplankton biomass, a C:chl a ratio of 30 was used (Banse 1977). Depth integration of PP, chl a, BP and BR was performed with the trapezoidal method. Areal PP was depth-integrated to the 1% light level while areal BP of unfiltered seawater and BR were integrated over the whole water column. Statistical analyses were done with the software package Statistica from Statsoft on logtransformed data whenever appropriate. For regressions, the ordinary least-squares (OLS) and reduced major axis (RMA) regressions were calculated. For comparison with other published empirical models OLS is presented, whereas RMA provides a better estimate of the true functional relationship (McArdle 2003) and references therein).

#### **RESULTS**

## Total versus 0.8 $\mu m$ filtered bacterial abundance and BP

Bacterial abundance and BP obtained for the 0.8  $\mu m$  fraction closely correlated with the abundance and

production in unfiltered seawater. However, filtration through 0.8 µm filters to exclude non-bacterial particles from the BR measurements reduced BP by 40  $\pm$ 34% compared to the raw seawater, except in July, when BP in the 0.8 µm filtered fraction and unfiltered seawater was equal. With increasing abundance and production in the unfiltered seawater, the percentage of bacterial abundance and BP recovered in the  $0.8\,\mu m$ fraction decreased. The loss of cells due to filtration did not change significantly between different months and bacterial abundance recovered in the 0.8 µm fraction was  $63 \pm 21\%$  of that in the unfiltered seawater. We used the optical backscatter readings from the CTD to estimate the relative particle load of suspended matter in the water column. From April to August, turbidity was relatively constant and sharply increased in the fall. Total bacterial abundance was not correlated with turbidity, but BP measured in unfiltered seawater showed a weak negative correlation with increasing particle load (Spearman rank correlation: r = -0.45; p < 0.05; N = 108; data not shown). Below, BP is referred to as the production measured in 0.8 µm filtered seawater unless otherwise noted.

## Relations between bacterioplankton abundance, BP and BR

Bacterial abundance of 0.8  $\mu$ m filtered samples increased from 0.66  $\times$  10<sup>6</sup> cells ml<sup>-1</sup> in April to 1.67  $\times$  10<sup>6</sup> cells ml<sup>-1</sup> in August and declined again toward winter (Fig. 2). BP and BR were significantly correlated (Fig. 3, Table 1). BP and temperature explained 61% of the variation in BR by multiple regression analysis, while BP alone explained only 45% of the BR (Table 1). In contrast, BP could not be predicted reliably from BR and temperature (Table 1).

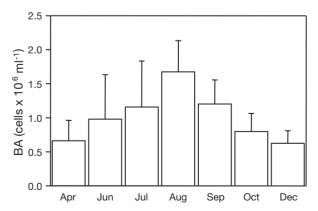


Fig. 2. Dynamics of bacterioplankton abundance (BA) in the southern North Sea (0.8  $\mu m$  pre-filtered). Data are averages per cruise, error bars: standard deviations of the mean; N = 11 to 28 for the different months

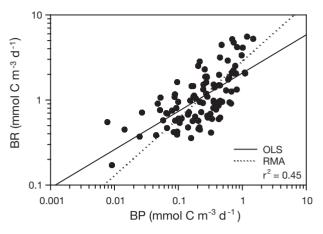


Fig. 3. Bacterial respiration (BR) as a function of bacterial production (BP) for 0.8  $\mu$ m filtered seawater. Ordinary least-squares model (OLS;  $r^2 = 0.45$ ) and reduced major-axis model (RMA;  $r^2 = 0.45$ ) are fitted to data. Model statistics in Table 1

Cell-specific BP declined steadily from 0.6 fmol C cell<sup>-1</sup> d<sup>-1</sup> in April to 0.06 fmol C cell<sup>-1</sup> d<sup>-1</sup> in December Cell-specific BR was fairly constant and varied only between 1 and 2 fmol C cell<sup>-1</sup> d<sup>-1</sup> (mean:  $1.4 \pm 0.71$  fmol C cell<sup>-1</sup> d<sup>-1</sup>) with a peak of 2.1 fmol C cell<sup>-1</sup> d<sup>-1</sup> in August (data not shown).

## Chlorophyll a and particulate primary production

Depth-integrated chl a concentrations were highest in April, with 96.1  $\pm$  78.8 mg chl a m $^{-2}$ , and declined toward December to 21.5  $\pm$  6.6 mg chl a m $^{-2}$  (Fig. 4). Particulate PP followed roughly the pattern of chl a (Fig. 4). The overall range of depth-integrated PP varied between 1.5 and 244.5 mmol C m $^{-2}$  d $^{-1}$  over all the stations, with an annual average of 62.0  $\pm$  59.8 mmol C m $^{-2}$  d $^{-1}$  (N = 59). Particulate PP was highest in April and July, with 133 and 143 mmol C m $^{-2}$  d $^{-1}$ , respectively, and lowest in December with an average of 5 mmol C m $^{2}$  d $^{-1}$  (Fig. 4).

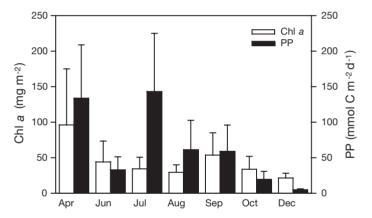
## Relation between phytoplankton and bacterial biomass and activity

Over the seasonal cycle, depth-integrated total bacterial biomass (BB) was similar to depth-integrated phytoplankton biomass (PB), indicated by a BB:PB ratio of ~1 (data not shown). However, in April and June, algal biomass dominated over bacterial biomass, whereas bacterial biomass was almost twice as high as phytoplankton biomass in August. During the remaining months, the ratio of total bacterial biomass versus phytoplankton biomass was not different from 1 (Stu-

Table 1. Regression statistics for volumetric and depth-integrated relationships of bacterial production (BP), bacterial respiration (BR), primary production (PP) and temperature (T). All variables were  $\log_{10}$ -transformed except for temperature. Reduced major axis (RMA) is more appropriate than ordinary least-squares (OLS) when x is measured with error (McArdle 2003). A conversion factor (CF) must be used when converting log-transformed to arithmetic scale (Sprugel 1983). Volumetric units of BP and BR in mmol C m<sup>-3</sup> d<sup>-1</sup>, depth-integrated units in mmol C m<sup>-2</sup> d<sup>-1</sup> and T in °C. Regressions follow form  $\log y = \log b_0 + x_1 \log b_1 (+x_2 b_2)$ ; SE: standard error; SEE: standard error of estimate used to calculate CF; 95% CI: upper and lower (95%) confidence limit of estimated parameters;  $r^2$ : coefficient of determination; p: significance level; N: number of measurements; \*not significant

Model	y	<i>x</i> <sub>1</sub>	<i>x</i> <sub>2</sub>	$b_0 \pm SE$	$b_1 \pm SE$	$b_2 \pm SE$	95%	6 CI	$\mathbf{r}^2$	p	SEE	CF	N
							$b_0$	$b_1$					
Volum	etri	rel	atior	ships									
OLS	BR	BP		$0.32 \pm 0.04$	$0.45 \pm 0.05$		0.24 0.39	0.35 0.55	0.45	< 0.001	0.23	1.15	102
RMA	BR	BP		$0.46 \pm 0.04$	$0.67 \pm 0.05$		0.38 - 0.54	0.57 0.77	0.45				102
OLS	BR	BP	T	$-0.32 \pm 0.11$	$0.32 \pm 0.05$	$0.04 \pm 0.01$			0.61	< 0.001	0.19		102
OLS	BP	BR	$T^*$	$-0.70 \pm 0.18$	$0.98 \pm 0.15$	$0.002 \pm 0.01$			0.45	< 0.001	0.34		102
Depth-	inte	grat	ed re	elationships									
OLS	BP	PP		$-0.24 \pm 0.16$	$0.66 \pm 0.09$		-0.56 $0.09$	0.48 0.85	0.50	< 0.001	0.35	1.38	55
RMA	BP	PP		$-0.70 \pm 0.16$	$0.94 \pm 0.09$		-1.03 -0.38	0.76 1.12	0.50				55
OLS	BR	PP		$1.29 \pm 0.12$	$0.18 \pm 0.07$		1.04 1.54	0.04 0.33	0.15	< 0.01	0.23	1.16	39
RMA	BR	PP		$0.82 \pm 0.12$	$0.48 \pm 0.07$		0.57 1.07	0.33 0.62	0.15				39

3.0



2.5 - Oite 2.0 - Oite 1.5 - Oite 1.5 - Oite 1.0 Oite 1.5 - Oite 1.0 Oite 1.

Fig. 4. Monthly averages (+SD) of depth-integrated chlorophyll a (chl a) and primary production (PP) measured in the southern North Sea. N = 4 to 21 for PP and 11 to 31 for chl a for the different months

Fig. 5. Monthly mean (+SD) ratio of depth-integrated total bacterial biomass (BB) and depth-integrated phytoplankton biomass (PB). Dashed line indicates unity

dent's t-test for single means; p < 0.001) (Fig. 5). Depthintegrated BP in unfiltered seawater averaged over the different months ranged from 1 to 26 mmol C m $^{-2}$  d $^{-1}$ . Thus, while particulate PP varied over 2 orders of magnitude, total BP was much less variable (Fig. 6a). Total BP as a percentage of particulate PP was highest after the bloom period in June and August, amounting for 49% of particulate PP. Over the annual cycle, total BP amounted to  $16 \pm 9\%$  of PP. BR did not exhibit any particular trend with depth in the well-mixed study area (data not shown). As for BP, BR was much less variable than phytoplankton production. Depth-integrated PP explained only about 15% of the variation in depthintegrated BR (Fig. 6b, Table 1). About 40% of the depth-integrated BR measurements were higher than

the areal PP estimates. On a volumetric basis, cell-specific BP of the 0.8  $\mu m$  filtered fraction significantly correlated with particulate PP (Spearman rank correlation: r = 0.50; p < 0.05; N = 54), while cell-specific BR was not related to particulate PP (Fig. 7).

### DISCUSSION

Generally, the southern North Sea is a highly dynamic system with strong tidal forces and a permanently well-mixed water column (Postma & Zijlstra 1988). This probably led to the low spatial variability in phyto- and bacterioplankton biomass and activity we recorded.

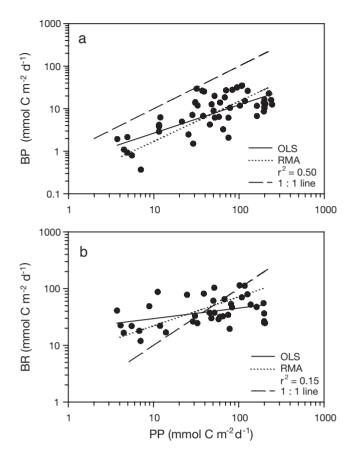


Fig. 6. (a) Depth-integrated bacterial production (BP) of unfiltered seawater and (b) depth-integrated bacterial respiration (BR) as a function of primary production (PP). Ordinary least-squares model (OLS) and reduced major-axis model (RMA) are fitted to data. 1:1 line: unity of parameters. Model statistics in Table 1

It is well known that bottle confinement, such as in BOD bottles, can bias bacterioplankton respiration estimates. An increase in bacterial abundance and community shifts during the course of incubations has been reported by several authors (Pomerov et al. 1994, Gattuso et al. 2002). However, Williams (1981) found that despite an increase in bacterial abundance during the incubation, the respiration rate is usually linear over incubation times up to 48 h. In our study, the average bacterial turnover rate was  $0.2 \pm 0.3 \,\mathrm{d}^{-1}$ , and most of our incubations for respiration measurements were between 12 and 18 h and never longer than 24 h. Thus, it is unlikely that bacterial abundance increased substantially during our incubations, and consequently shifts in the community composition should also be minimal. Moreover, our regression of BR on BP (Fig. 3) was very similar to that obtained by Del Giorgio & Cole (1998), further indicating that BR in the southern North Sea is in the range of published respiration rates.

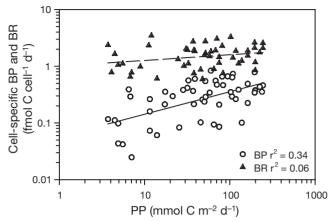


Fig. 7. Cell-specific bacterial production (BP) and respiration (BR) for 0.8 µm-filtered seawater as a function of particulate primary production (PP)

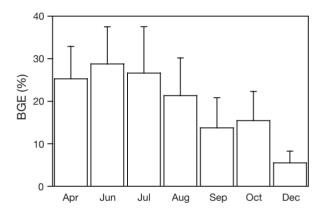


Fig. 8. Seasonal dynamics of bacterial growth efficiency (BGE) calculated as BGE =  $BP/(BP + BR) \times 100$  from April to December. Means (+SD); N = 9 to 21 estimates for the different months

Determining BGE depends on the choice of the conversion factor used in BP estimates and on the respiratory quotient (RQ) used. While RQ values are assumed to center around 1 and are considered a minor source of error (Del Giorgio & Cole 1998), conversion factors for BP generally vary over a much larger range (Simon & Azam 1989, Riemann et al. 1990, Ducklow et al. 1992, Kirchman 1992). We scaled the bacterial production measurements with a conversion factor typical for coastal systems (Riemann et al. 1990); however, choosing a higher conversion factor of (e.g.) 3.1 kg C mol<sup>-1</sup> leu (Simon & Azam 1989) would increase our BGEs 1.7 times. We found a clear seasonal pattern in BGE, with a median value of 25% in the spring and summer (mean  $25 \pm 9\%$ ; N = 60), decreasing to a median BGE of 14% in the fall (mean  $15 \pm 7\%$ ; N = 33) and 5% in the winter (mean  $6 \pm 3\%$ ; N = 9) (Fig. 8). The annual median BGE in the southern North Sea was 19% (mean  $20 \pm 11\%$ ; N = 102), which is similar to the overall BGE of 20% obtained by Del Giorgio & Cole (2000) compiling available information from marine surface waters and Carlson et al. (1999) from a similar high-latitude system.

The seasonality in BGE is mainly driven by the changes in BP, which vary 6-fold from spring to winter. BR was generally rather uniform over the seasonal cycles, although there was a slight increase in cell-specific BR toward the warmer summer months. A peak in respiration coinciding with the annual temperature maximum was reported previously (Smith & Kemp 1995 and references therein), but in our study this increase was too small to significantly influence the pattern of BGE. While Sherry et al. (1999) found that the seasonal dynamics in BGE were influenced more by bacterial respiration rather than production along an E–W transect in the NE Pacific, our finding is in agreement with that of Del Giorgio & Cole (2000), who concluded that BP was mainly determining BGEs.

Reports on temperature-dependence of bacterioplankton metabolism are contradictory. White et al. (1991) used a multiple regression including bacterial abundance and temperature to explain the variability in BP. Pomeroy & Wiebe (2001) argued that temperature and substrate availability are both limiting factors for heterotrophic bacteria, while Del Giorgio & Cole (1998) concluded that temperature was not an important factor controlling bacterioplankton activity. Over the temperature range of our study (9 to 18°C; Table 2), temperature explained only around 32% of the variability in BP. Thus, in the southern North Sea temperature does not explain the seasonality in BP, and the model from Rivkin & Legendre (2001) to derive BGE from BP and temperature cannot be applied here, as they used a much larger temperature range. In a study in the northern North Sea, Robinson et al. (2002b) calculated BR from the model of Rivkin & Legendre (2001), and the resulting estimate grossly underestimated the bacterial contribution to total respiration compared to the model of Del Giorgio & Cole (1998)

Table 2. Monthly averages of mixed-layer depth (m), salinity (S, PSU), temperature (T,  $^{\circ}$ C) and DOC concentration ( $\mu$ M)

Month	Depth	S	T	DOC	N
Apr	$40 \pm 20$	$34.01 \pm 1.19$	$8.8 \pm 0.8$	$72.7 \pm 20.5$	32
Jun	$29 \pm 7$	$34.12 \pm 0.29$	$12.2 \pm 0.3$	$139.8 \pm 34.7$	17
Jul	$28 \pm 6$	$34.63 \pm 0.28$	$14.9 \pm 0.2$	$86.5 \pm 9.6$	20
Aug	$33 \pm 7$	$34.28 \pm 0.30$	$18.4 \pm 0.5$	$265.3 \pm 89.0$	23
Sep	$28 \pm 6$	$34.31 \pm 0.21$	$16.6 \pm 1.0$	$87.1 \pm 19.9$	17
Oct	$31 \pm 8$	$34.48 \pm 0.23$	$13.1 \pm 1.2$	$107.6 \pm 19.2$	31
Dec	$30 \pm 8$	$34.55 \pm 0.48$	$10.3 \pm 0.9$	$152.2 \pm 43.5$	11

which gave more a reasonable value of  ${\sim}60\,\%$  of total respiration.

Other factors causing seasonal fluctuations in BP are changes in the concentrations of readily utilizable dissolved organic matter (DOM) due to variable extracellular release of phytoplankton, grazing activity and/or allochthonous input of organic material via rivers. In our study, phytoplankton biomass (measured as chl a) was not related to bacterial abundance; however, a positive relationship was found between PP and specific BP (Fig. 7).

The lowest mean DOC concentrations of around 73 µM were measured in the spring and DOC peaked in August with an average of 265 µM, but no clear seasonal pattern was apparent (Table 2). Søndergaard & Middelboe (1995) estimated that in marine environments around 19% of the bulk DOC is labile and used by prokaryotes within 1 and 2 wk. The more labile components of the DOC pool are directly or indirectly fueled by the release from phytoplankton, and this DOM is remineralized within hours (Fuhrman 1987, Rich et al. 1997). Natural DOC consists of a continuum of size classes of differing diagenetic state (Amon & Benner 1996), which makes it difficult to directly relate bulk DOC measurements to BP and growth. As a measure of the availability of labile DOC we used the ratio of BP to bulk DOC (Obernosterer et al. 1999). In the spring and summer, relatively more labile DOC is available, as indicated by a higher ratio between BP and bulk DOC concentration than in the winter (Fig. 9a). The ratio of BP to DOC, as a function of phytoplankton production, indicates that the increase in the relative availability of DOC is related to increased pp (Fig. 9b). Therefore, we conclude that the BP in the southern North Sea is mainly coupled to the seasonal dynamics in primary production and that BGE is directly linked to the bioavailability of DOC and indirectly to PP (Fig. 9c). While such a conclusion would be expected for open-ocean systems, for a shelf sea such as the North Sea this conclusion is not so obvious, considering the large input of terrigenous material via rivers.

To assess the relative importance of autotrophic versus heterotrophic processes, the ratio of phytoplankton production to bacterial carbon demand (BCD) based on the total BP measurements was calculated. We used the depth-integrated values of PP and BR because, as Williams (1998) pointed out, PP should be considered on an areal basis to account for the solar radiation dependency of PP. A considerable part of the phytoplankton production may be lost to the dissolved phase through extracellular release of DOC (Kaltenböck & Herndl 1992). Thus our PP measurements are likely to underestimate actual total phytoplankton production (Teira et al. 2001). The PP:BCD ratio was significantly

higher than 1 in April and July while in August and December the southern North Sea was heterotrophic (Student's t-test for single means; p < 0.001) (Fig. 10). The remaining months exhibited roughly a metabolic balance, i.e. the ratio was not significantly different

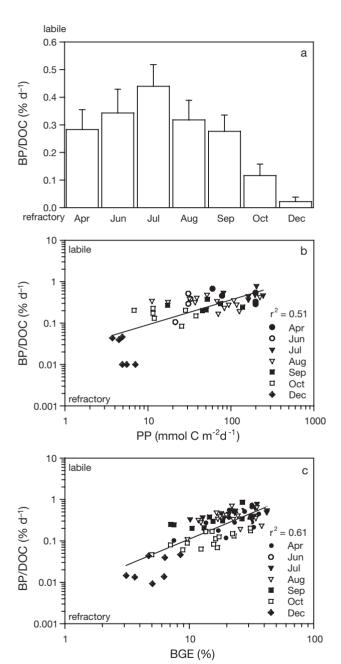


Fig. 9. Total bacterial production (BP; mmol C  $\rm m^{-3}$  d<sup>-1</sup>) normalized to dissolved organic carbon (DOC; mmol C  $\rm m^{-3}$ ) to indicate relative DOC availability (BP/DOC  $\times$  100). (a) Distribution of DOC available over the seasonal cycle, showing ratio of monthly averages; means (+SD). (b) Relation of depthintegrated primary production and available DOC. (c) Dependence of bacterial growth efficiency (BGE) on DOC availability

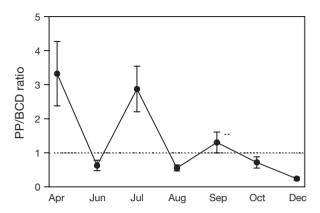


Fig. 10. Ratio between particulate primary production (PP) and total bacterial carbon demand (BCD) over seasonal cycle. Means  $(\pm SD)$  of 10 estimates

from 1 (Fig. 10). This pattern generally agrees with a recent study of Thomas et al. (2004), who measured fluxes of  $CO_2$  in the North Sea on a seasonal scale. They found that the southern part was slightly heterotrophic throughout the year, except in spring when  $CO_2$  uptake from the atmosphere prevailed.

Recently, evidence has been presented that openocean systems are slightly heterotrophic (Del Giorgio et al. 1997, Duarte & Agusti 1998, Williams 1998). In marginal seas such as the North Sea, short-term trophic imbalances might be compensated by terrestrial input via rivers and terrestrial runoff. In the southern North Sea, the rivers Humber, Thames and the Rhine provide considerable amounts of DOM (A. V. Borges et al. unpubl.); thus, potentially, even long periods of net system heterotrophy would be possible. Decreasing PP toward the winter was paralleled by decreasing BGE. However, in the spring and early summer the supply of autochthonous DOM supports a relatively high BGE until most of the labile DOM is depleted in August.

In summary, we have shown that seasonality in BP is mainly responsible for the dynamics in BGE which, in turn, is influenced by the bioavailability of DOC. Remineralization of organic carbon is the main function of prokaryotes in carbon cycling and, as such, is relatively stable over the annual cycle.

Acknowledgements. We thank J. Hegeman, S. Oosterhuis, A. Noordeloos and S. Gonzalez for the primary production, chlorophyll a and DOC data. Special thanks to G. van Noort and C. Winter for help with analysis of the bacterial abundance data and to M. Baars for his excellent job as chief scientist during the Bloom and Plume project and to the captain and crew of the RV 'Pelagia'. This work was supported by the Dutch Science Foundation (NWO-ALW) project 811.33.002 and the European Commission through the COMET project to G.J.H. This work is in partial fulfillment of the requirements for a PhD degree from the University of Groningen by T.R.

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Editorial responsibility: Frede Thingstad, Bergen, Norway

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Submitted: June 21, 2004; Accepted: January 10, 2005 Proofs received from author(s): March 11, 2005