

Photo- and bioreactivity of chromophoric dissolved organic matter produced by marine bacterioplankton

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ABSTRACT: The major source of oceanic dissolved organic matter (DOM) is organic compounds from phytoplankton released either directly during photosynthesis or via grazing. Bacterioplankton as a source of marine DOM have received considerably less attention. We determined the production of DOM during active bacterial growth in batch cultures with artificial seawater amended with glucose and inorganic nitrogen and phosphorus. During the initial growth of bacterioplankton, the fluorescence of bacterial-derived DOM increased steadily indicating release of chromophoric DOM. This bacterial-derived DOM was relatively rich in N and depleted in P as indicated by the C:N:P ratio of 147:29:1. Exposure of this bacterial-derived DOM to artificial solar radiation provoked a rapid decline in fluorescence indicating its photoreactivity. Re-growth experiments with natural bacterial assemblages inoculated into the solar radiation-exposed bacterial-derived DOM indicated that the bioavailability remained essentially unaltered, i.e. is refractory to bacterial utilization. Our findings indicate that bacterioplankton release chromophoric, refractory DOM during active growth and that this DOM, albeit being photoreactive does not stimulate bacterial growth activity upon exposure to solar radiation.

KEY WORDS: Bacterioplankton · Bacterial-derived DOM · UV radiation · Photoreactivity · Bioreactivity

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INTRODUCTION

The pool of oceanic dissolved organic matter (DOM) represents, besides soil humus, the largest reservoir of organic matter of the biosphere (Hedges 1992). The major fraction of this oceanic DOM pool is derived from phytoplankton, either directly, via extracellular release (Obernosterer & Herndl 1995, Teira et al. 2001, and references therein) or indirectly, through lysis and grazing (Bratbak et al. 1990, Lee & Fisher 1994, Sherr & Sherr 1994, Guixa-Boixareu et al. 1996).

Bacterioplankton as producers of marine DOM have received considerably less attention than phytoplankton. It has been shown, however, that bacteria produce copious amounts of largely refractory DOM (Brophy & Carlson 1989, Tranvik 1993, Heissenberger & Herndl 1994, Stoderegger & Herndl 1998). Ogawa et al. (2001)

showed in laboratory experiments that only 30% of the bacterial-derived DOM is hydrolyzable and, therefore, accessible for analysis on a molecular level.

Considerable information has accumulated over the past decade on the photochemical alteration and degradation of DOM. Photochemical degradation of DOM might enhance its overall availability for bacterioplankton (Lindell et al. 1995, Obernosterer et al. 1999). While this enhanced DOM bioreactivity upon exposure to solar radiation has been frequently reported (e.g. Kieber et al. 1989, Lindell et al. 1995, Obernosterer et al. 1999), DOM might also become less reactive upon exposure to solar radiation as compared to DOM held in the dark (Benner & Biddana 1998, Obernosterer et al. 1999, 2001a). It has been suggested by Obernosterer et al. (1999) that originally more labile DOM becomes more refractory while originally more

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refractory DOM becomes more labile upon exposure to solar radiation. Overall, it has been estimated that about 2 to 4% of the oceanic DOC is photochemically oxidized to CO₂ and CO per year (Moran & Zepp 1997, Obernosterer et al. 2001b).

In the present paper we tested the hypothesis that bacteria produce copious amounts of relatively refractory DOM from originally labile substrate during the exponential growth phase, and that this bacterial-derived DOM is photoreactive and becomes more bioavailable for bacterial utilization upon exposure to solar radiation. Previous studies have already shown that bacteria produce refractory DOM (Brophy & Carlson 1989, Tranvik 1993, Ogawa et al. 2001). Using short-term incubations of several days, Heissenberger & Herndl (1994) showed that bacterioplankton release about 6% of the incorporated leucine as DOM larger than 50 000 Da, while Stoderegger & Herndl (1998) demonstrated that the capsular envelope of bacteria is constantly renewed, releasing the older material into the ambient water. These authors also showed that this bacterial-derived capsular DOM is semi-labile to refractory. Since a substantial part of the released DOM is chromophoric (Tranvik 1993, this study), we expected that exposure to solar radiation would substantially increase its bioavailability.

MATERIALS AND METHODS

Experimental approach to determine chromophoric DOM production by bacteria. The production of bacterial-derived chromophoric DOM was assessed in microcosm experiments. In total, 4 experiments each in duplicate were performed. Bacterioplankton of the coastal North Sea were collected from the NIOZ jetty at 0.5 m depth and immediately filtered through 0.8 µm filters (polycarbonate, 140 mm filter diameter). This filtrate (5 l) was concentrated with a 0.2 µm polycarbonate filter (Millipore, 47 mm) and resuspended in artificial seawater. The concentrated (~20 ml) bacterial community was inoculated in 2 l of 0.2 µm filtered artificial seawater, amended with 200 µM glucose-C, 20 µM NH₄⁺, and 2 µM PO₄³⁻. Bacteria were allowed to grow in the dark at 18°C for a period of 17 to 21 d. Artificial seawater without bacteria inoculated served as an abiotic control. In these seawater batch cultures and in the control flasks bacterial abundance, dissolved organic carbon (DOC), nitrogen (DON), and phosphorus (DOP) concentrations, as well as DOM fluorescence were monitored at regular intervals as described below.

Experimental methods to evaluate the photoreactivity of bacterial-derived chromophoric DOM and its bioavailability. After allowing the bacterioplankton

assemblage to grow for 17 to 21 d in nutrient amended artificial seawater, the bacteria were removed from the culture medium by filtering it twice through rinsed 0.2 µm filters (Millipore, polycarbonate). Subsequently, the 2 l filtrate was split. One half of the volume was exposed to artificial solar radiation in a quartz tube (2.8 cm inner diameter) for up to 24 h and the other half was held in the dark by wrapping the borosilicate tube in aluminum foil. During this exposure, the DOM fluorescence was monitored at regular intervals.

Artificial solar radiation was provided by 3 different types of light sources. Two HQI-T Powerstar (Osram) lamps provided photosynthetic active radiation (400 to 700 nm wavelength range), 2 TL 100W/10R fluorescent light tubes (Philips) were used to provide UV-A (320 to 400 nm) and 3 UVA-340 fluorescent light tubes (Q-Panel) supplied UV-A and UV-B (300–320 nm). The solar simulator was adjusted to 30–60% of the local maximum radiation intensity in late spring measured on a cloudless day (Pausz & Herndl 2002). Thus, the dose rate received by the bacterial-derived DOM in our exposure experiments is similar to that expected in the top surface layer of the water column. To maintain in situ water temperature (17 to 18°C) during the exposure to artificial solar radiation, the treatments were kept in a flow-through water bath connected to a temperature control unit (LAUDA RCS/RC-6).

After exposing the bacterial-derived chromophoric DOM to artificial solar radiation, a natural bacterioplankton community (prepared as described above) was inoculated into the solar radiation-exposed and into the dark treatment and allowed to grow in the dark at 18°C for 2 d. In order to check for growth limitation due to a lack of available C, N or P, different treatments were established enriched in these elements in 2 experiments (Expt II: unamended control, 200 µM glucose-C, 20 µM NH₄⁺, 2 µM PO₄³⁻; Expt III: unamended control, 5 µM NH₄⁺ and 1 µM PO₄³⁻).

Enumeration of bacteria. To determine the bacterial abundance, 1 to 5 ml of sample was fixed with 0.2 µm filtered (Acrodiscs, Gelman) 37% formaldehyde (4% final concentration), stained with 4',6-diamidino-2-phenylindole-dihydrochloride (DAPI, final concentration 10 µg ml⁻¹) and filtered onto a black 0.2 µm polycarbonate filter (Millipore, 25 mm filter diameter) for counting under the epifluorescence microscopy (Zeiss Axioplan, 1250× magnification; Porter & Feig 1980). Filters were stored in the dark at -20°C until analysis (within 3 months after collection). At least 30 fields or 300 bacteria per sample were counted.

Analyses of DOM and inorganic nutrients. Samples for DOC, DON, DOP, and DOM fluorescence were filtered through combusted (450°C for 4 h) Whatman GF/F glass fiber filters. For DOC determination, 8 ml of

the filtrate was transferred into combusted glass ampoules, immediately acidified with 3 to 4 drops of 45% H_3PO_4 and then the ampoules were sealed. Samples were stored at -20°C until analysis.

For the determination of the inorganic nutrients, DON and DOP, the samples were collected in acid-rinsed polyethylene bottles and stored at -20°C until analysis. All analyses were done within 3 mo of collecting the samples.

DOC analysis was performed on quadruplicate samples using the high temperature combustion method on a modified Shimadzu TOC-5000A. The absorbance of the CO_2 evolving from the combusted DOC was detected with an external infrared cell (LiCor Model LI-6252) and quantified as peak area by the internal Shimadzu integrator. Ultra-pure CO_2 -free air was used as a carrier gas at a flow rate of 150 ml min^{-1} . Samples were automatically injected with a Shimadzu ASI-5000A autosampler on a platinumized aluminum catalyst (Elemental Microanalysis) held at 680°C (Benner & Strom 1993). Standards were prepared with potassium hydrogen phthalate (Baker) in Milli-Q water. The overall analytical precision was always better than 3%.

DON and DOP analyses were performed simultaneously following the method of Valderrama (1981). This procedure is based on an alkaline persulphate digestion (at 120°C in an autoclave for 90 min) over a wide pH range starting at pH 9 and ending at pH 4 using boric acid and sodium hydroxide. The resulting total nitrate and phosphate were measured using the autoanalyzer for inorganic nutrient determination as described below. The sum of the inorganic nitrogen species ($\text{NH}_4^+ + \text{NO}_3^- + \text{NO}_2^-$) and phosphate concentrations were subtracted from the corresponding total dissolved N and P concentrations, respectively, to calculate the concentrations of DON and DOP. The recovery efficiency of DON and DOP was checked using a cocktail of 10 different compounds containing organic N and/or P measured in parallel with the samples. The recovery was 80% of the total added N in DON and 94% of the total added P in PON.

Concentrations of NH_4^+ , NO_2^- , NO_3^- and PO_4^{3-} were determined in a TRAACS autoanalyzer system. NH_4^+ was detected with the indo-phenolblue-method (pH 10.5) at 630 nm wavelength (Helder & de Vries 1979). NO_2^- was detected after diazotization with sulphanilamide and *N*-(1-naphtyl)-ethylene diammonium-dichloride as the reddish purple dye complex at 540 nm (Parsons et al. 1984). NO_3^- was reduced in a copper cadmium coil to NO_2^- (using imidazole as a buffer) and then measured as NO_2^- . Phosphate was determined via the molybdenum blue complex at 880 nm according to Murphy & Riley (1962).

For the determination of the DOM fluorescence, 3 ml of the filtrate was immediately measured at an excita-

tion of 350 nm and an emission wavelength of 450 nm using a 1 cm quartz cuvette in a Hitachi F-2000 fluorometer. The fluorescence was standardized with a quinine sulfate solution and is given in QSU ($1\text{ QSU} = 1\text{ ppb quinine sulfate in } 0.05\text{ M H}_2\text{SO}_4$).

RESULTS

Production of chromophoric DOM by marine bacterioplankton

In the nutrient-amended artificial seawater batch cultures, bacterial abundance increased from initially $0.2 \times 10^6\text{ cells ml}^{-1}$ to about $10 \times 10^6\text{ cells ml}^{-1}$ within 5 d (Fig. 1A). Thereafter, the natural bacterial community remained in the stationary phase for 3 to 5 d and sub-

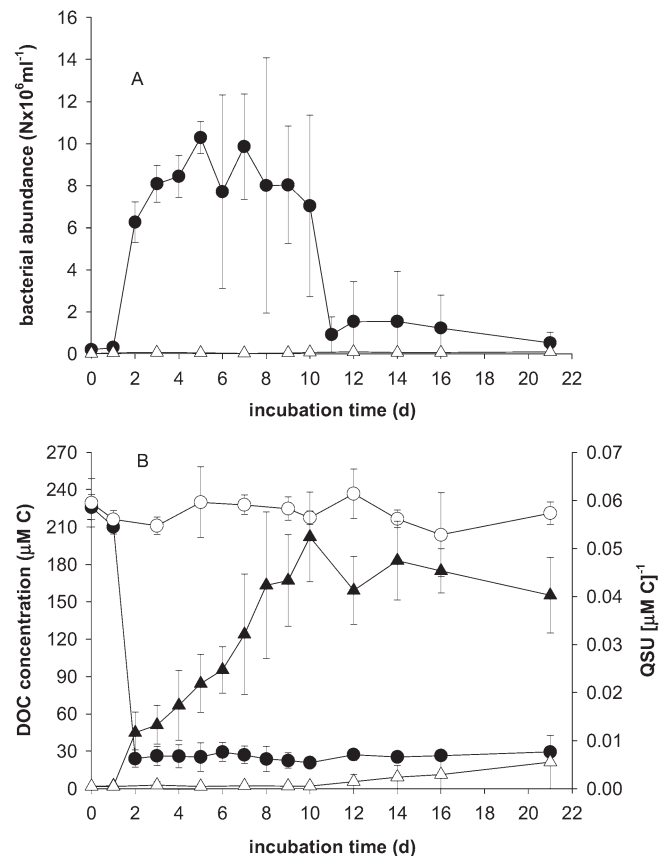


Fig. 1. Development of (A) the bacterial abundance and (B) the DOC concentrations and the DOC-normalized DOM fluorescence (in QSU [$\mu\text{M C}$] $^{-1}$) in the 4 batch culture experiments each done in duplicate with natural bacterial assemblages (solid symbols) and the corresponding control experiments where no bacteria were inoculated (open symbols). In (B), the DOC concentrations are indicated by circles and the DOC-normalized DOM fluorescence by triangles. The 3 data points for Days 14 to 21 of one of the abiotic control experiments were not included as bacterial growth was observed. Error bars indicate SD

Table 1. Development of the concentrations of the dissolved organic carbon (DOC), nitrogen (DON), and phosphorus (DOP) (all in μM) and the C:N, C:P and N:P ratios of the DOM in Expts II and III during the incubation of natural bacterial assemblages in batch cultures with artificial seawater amended with glucose and inorganic nutrients. The mean of duplicate incubations is given

Day of incubation	DOC ($\mu\text{M C}$)	DON ($\mu\text{M N}$)	DOP ($\mu\text{M P}$)	C:N	C:P	N:P
Expt II						
0	167	4.8	0.2	38	851	25
4	86	4.7	0.17	24	491	26
9	nd	7.4	0.18	nd	nd	43
17	46	7.3	0.29	6	168	27
Expt III						
0	212	3.8	0.03	66	6723	123
5	25	7.4	0.16	4	159	47
11	21	5.8	0.25	4	85	25
20	27	6.5	0.22	5	127	31

sequently, bacterial abundance decreased rapidly to about 1 to 2×10^6 cells ml^{-1} (Fig. 1A). The bacterial abundance in the abiotic controls remained below 0.2×10^6 cells ml^{-1} in all the experiments, except in one, where bacteria grew up in one of the duplicate flasks towards the end of this particular experiment (data not included in Fig. 1A).

Bacterioplankton were utilizing the $200 \mu\text{M}$ glucose-C added to the batch cultures within 2 d (Fig. 1B). On Day 2, the DOC concentrations were, on average, $24.1 \pm 6.9 \mu\text{M C}$ and varied subsequently between 20.8 and $29.5 \mu\text{M C}$. In the abiotic treatments without bacteria inoculation, the DOC concentrations remained stable over the entire incubation period (Fig. 1B), again with the exception of one experiment where bacterial growth was observed (data not included in Fig. 1B).

Concurrently with the sharp decline in DOC due to the uptake of the added glucose until Day 2, the DOC-normalized DOM fluorescence increased from undetectable levels on Day 0 to $0.012 \pm 0.004 \text{ QSU } [\mu\text{M C}]^{-1}$ on Day 2 (Fig. 1B). Subsequently, DOC-normalized fluorescence increased steadily until Day 9 when the highest DOM fluorescence was reached ($0.052 \pm 0.009 \text{ QSU } [\mu\text{M C}]^{-1}$). Thereafter, the DOC-normalized DOM fluorescence decreased slightly and remained more or less constant thereafter until the end of the experiment (averaging $0.043 \pm 0.006 \text{ QSU } [\mu\text{M C}]^{-1}$) between Day 10 and Day 21. In the abiotic treatments, the DOC-normalized DOM fluorescence ranged from 0 to $0.001 \text{ QSU } [\mu\text{M C}]^{-1}$, again with the exception of one experiment where bacteria were growing up (data not included in Fig. 1B).

The dynamics of DON and DOP produced by bacterioplankton growing on inorganic N and P were

monitored as well in 2 duplicate experiments (Expts II and III) and are shown in Table 1. Over the course of the 21 d incubation period, the DON concentrations increased from about 4 to about $7 \mu\text{M N}$ in both experiments and DOP concentrations from 0.20 to $0.29 \mu\text{M P}$ in Expt II and in Expt III from 0.03 to $0.22 \mu\text{M P}$ (Table 1).

Generally, the C:N:P ratios of the bacterial-derived DOM pool remained remarkably stable over the course of the incubation period after the initial decrease from Day 0 to Day 4 due to the uptake of glucose (Table 1). At the end of the incubation, the C:N ratio of the bacterial-derived DOM pool was 6 and 5 in Expts II and III, respectively. The C:P ratio of the DOM pool at the end of the experiments was 168 and 127 in Expts II and III, respectively, and the N:P ratios were 27 and 31 in Expts II and III, respectively (Table 1).

Photoreactivity of the bacterial-derived chromophoric DOM

Exposure of the bacterioplankton-derived chromophoric DOM to solar radiation resulted in a rapid decline in the DOM fluorescence, while the DOM fluorescence in the dark treatments remained constant (Fig. 2). In Expt II, mean DOM fluorescence decreased from 0.069 to $0.020 \text{ QSU } [\mu\text{M C}]^{-1}$ within the initial 6 h of exposure to solar radiation. Thereafter, DOM fluorescence declined only slightly in Expt II reaching $0.016 \text{ QSU } [\mu\text{M C}]^{-1}$ at the end of the 24 h exposure period. In Expt III, however, DOM fluorescence decreased steadily throughout the entire exposure period reaching also a DOM fluorescence value of $0.016 \text{ QSU } [\mu\text{M C}]^{-1}$ after 11 h (Fig. 2). The decline in the DOC-normalized DOM fluorescence upon exposure to solar radiation indicates that bacterial-derived DOM is photoreactive although no significant decrease in DOC concentrations during the exposure to solar radiation was discernable (data not shown). The DOM fluorescence of the bacterial-derived DOM held in the dark remained essentially constant throughout the exposure period (Fig. 2).

Utilization of bacterial-derived DOM by marine bacterioplankton

Subsequent inoculation of natural bacterioplankton consortia with solar radiation-exposed bacterial-

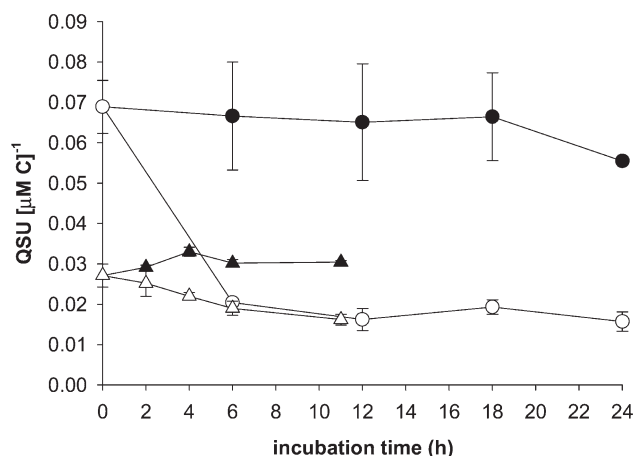


Fig. 2. Dynamics of the DOC-normalized DOM fluorescence during the exposure of bacterial-derived DOC to artificial solar radiation in Expts II (solid circles) and III (triangles). Open symbols represent solar radiation exposed and solid symbols DOM held in the dark. Symbols represent means of duplicate treatments; vertical lines indicate the range

derived DOM and that held in the dark resulted in a maximum bacterial abundance not significantly different between the 2 treatments regardless whether additional NH_4^+ and/or PO_4^{3-} were added (Kruskal-Wallis, abundance data from 7 time points per treatment used for analysis, $p > 0.05$) (Fig. 3). Only the addition of glucose-C in Expt II resulted in about twice the maximum bacterial abundance in both, the solar radiation-exposed and the corresponding dark treatment (Fig. 3) indicating that the bioavailability of C was limiting bacterioplankton growth.

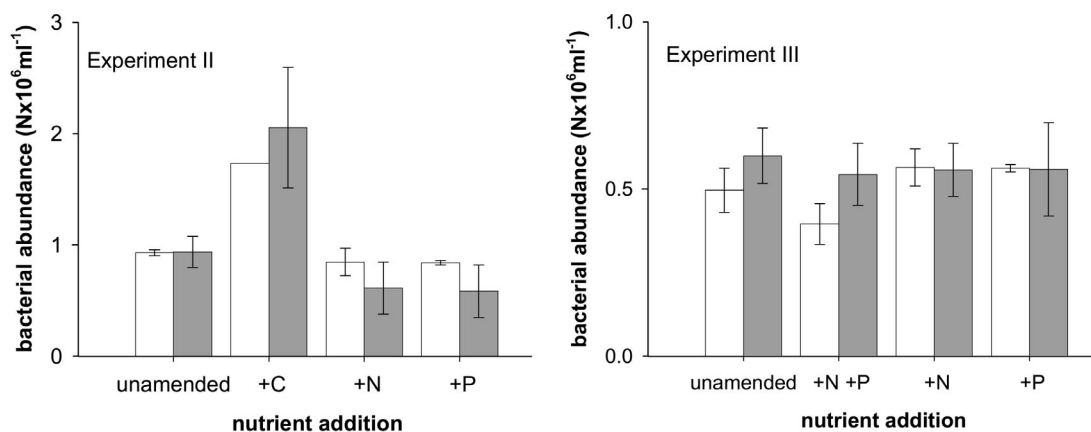


Fig. 3. Maximal bacterial abundance in the re-growth experiments after exposing bacterial-derived DOM to solar radiation or held in the dark and then inoculating a natural bacterial community after amending the bacterial derived DOM with different nutrients. The nutrient additions for Expt II were 200 μM glucose-C (+C), 20 μM NH_4^+ (+N), 2 μM PO_4^{3-} (+P), and an unamended control (unamended); for Expt III the nutrient additions were 5 μM NH_4^+ and 1 μM PO_4^{3-} (+N +P), 5 μM NH_4^+ (+N), 1 μM PO_4^{3-} (+P), and an unamended control (unamended). White bars represent solar radiation exposed treatments; gray bars represent treatments held in the dark prior to inoculating bacteria. Average values and the range of duplicate experiments are given. Maximum bacterial abundance was reached after an incubation of 30 to 48 h

DISCUSSION

In the batch cultures with artificial seawater amended with nutrients, bacterioplankton were rapidly transforming labile organic matter (here in the form of glucose) into chromophoric DOM, as indicated by the steady increase of the DOM fluorescence in the treatments inoculated with a natural bacterial community (Fig. 1B). Correcting our DOC concentrations for the background DOC concentration of the artificial seawater ($\sim 20 \mu\text{M}$ DOC), bacterioplankton consumed essentially all the added glucose-C within 2 d. Using a similar experimental approach, Ogawa et al. (2001) found that 85% of the added labile C was taken up by the bacteria within 2 d. After 10 d, DOC concentrations amounted to 9% of the initial concentration. Similarly, in the experiments of Ogawa et al. (2001), DOC concentrations decreased to 8% of the initial DOC after 7 d.

DON derived from bacterioplankton increased in concentration during the course of the incubation by 52 and 70% in Expts II and III, respectively, representing a net production of 2.5 and 2.7 μM DON, respectively (Table 1). The initial inorganic N concentrations (data not shown) decreased by 32% in Expt II and by 36% in Expt III, resulting in a net uptake of 7.5 and 8.8 μM N, respectively, during the incubation. Thus, in our experiments, the bacterial community transformed a net percentage of around 30% of the inorganic N taken up into semilabile to refractory DON within 17 to 20 d. Heissenberger & Herndl (1994) found that after ~ 6 d, 60 to 70% of the added ^{14}C -leucine was transformed into high molecular weight DOM (>1000 Da).

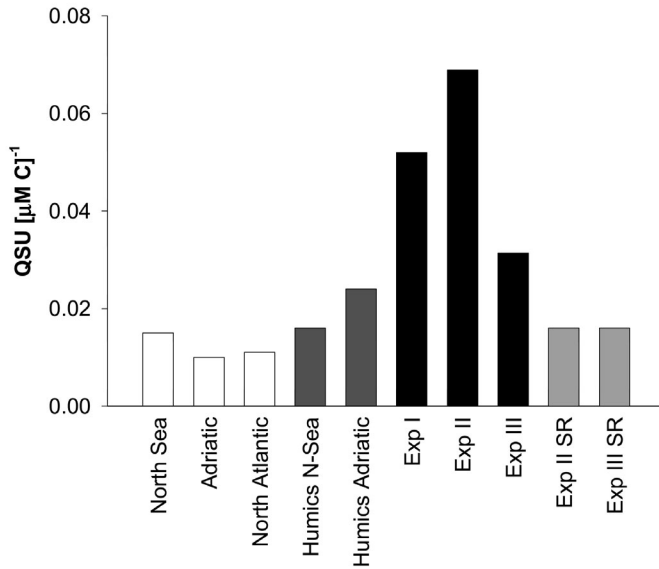


Fig. 4. Comparison of the DOC-normalized fluorescence of marine bulk DOM and its humic fraction obtained from the North Sea and the Adriatic with that of bacterial-derived DOM obtained in the present experiments. Expt II SR and Expt III SR are average values of duplicate treatments after the exposure to artificial solar radiation from Expts II and III. North Sea and Adriatic data are from Obernosterer & Herndl (2000); North Atlantic data from Obernosterer et al. (2001b)

At the end of the incubation period of 10 d, however, only 6% of the ^{14}C -leucine was in the $>50\,000$ Da fraction of the DOM (Heissenberger & Herndl 1994). Ogawa et al. (2001) measured bacterial-derived total hydrolyzable dissolved amino acid concentrations of 0.6 to 1.0 $\mu\text{M N}$ representing only $\sim 30\%$ of the total DON pool. Thus, taking this 30% recovery of DON into account, the bacterial-derived DON concentrations were 2 to 3 $\mu\text{M DON}$, which is similar to the average DON concentration we determined (2.6 μM).

After 20 d of incubation, the C:N:P ratio of the DOM produced by bacterioplankton was 168:27:1 in Expt II and 127:31:1 in Expt III. These C:N:P ratios of the bacterial-derived DOM are substantially lower than for bulk marine surface (C:N:P = 300:22:1, C:N = 17.8) and deep water DOM (C:N:P = 444:25:1, C:N = 13.6) compiled in Benner (2002). The low C:N ratios of 6 and 5 for Expts II and III, respectively, and the high N:P ratios indicate that bacterial-derived DOM is relatively rich in N but depleted in P as compared to the bulk oceanic DOM and the overall elemental composition of bacteria (Fagerbakke et al. 1996). This indicates that bacterioplankton are retaining P in the cell while releasing N-rich compounds into the water. This also suggests that the viral lysis probably played only a minor role in releasing intracellular material into the ambient water since the P-content of the DOM would then reflect

more closely the elemental composition of the bacterioplankton. Especially in the initial growth phase of bacteria in the batch culture (Days 4 to 5) viral lysis certainly played only a negligible role and because the bacterial-derived DOM already exhibited high C:P and N:P ratios (Table 1).

The fluorescence of the bacterial-derived DOM was up to 7 times higher than that of marine bulk DOM (Fig. 4) and at least twice as high as the fluorescence of the humic fraction of marine DOM extracted with XAD-8 resin (Obernosterer & Herndl 2000) (Fig. 4). This comparison indicates that bacterial-derived DOM consists of highly chromophoric DOM. Chromophoric DOM has been shown to be highly photoreactive (Obernosterer & Herndl 2000). Upon solar radiation cleavage products are formed from the parent humic substances which are efficiently utilized by bacterioplankton leading to an overall higher bioavailability of solar radiation-exposed humic substances (Obernosterer & Herndl 2000). It has been shown that ammonia and amino acids are formed from N-containing humic matter upon exposure to solar radiation (Bushaw et al. 1996). Thus, we assumed that the N-rich bacterial-derived DOM might become at least partially available to bacterioplankton utilization after exposure to solar radiation.

Upon exposure to artificial solar radiation, the fluorescence of the bacterial-derived DOM decreased during the initial 6 h, thereafter, only small changes were detectable (Fig. 2). Although the initial DOC-normalized DOM fluorescence in Expt II was more than twice as high than the initial DOM fluorescence in Expt III, the final DOM fluorescence was similar in both experiments (~ 0.016 QSU [$\mu\text{M C}^{-1}$]). Thus, bacterial-derived DOM is photoreactive (Fig. 4). Bulk marine DOM is less photoreactive than bacterial-derived DOM since Obernosterer et al. (1999), exposing bulk marine DOM to similar radiation levels, report a decrease of DOC-normalized DOM fluorescence by ~ 50 and 22% for mesopelagic and surface waters of the Mediterranean Sea, respectively. For the coastal North Sea, the site from where the water used in this study originated, a decrease of DOC-normalized DOM fluorescence of only 17% was measured (see Table 2 in Obernosterer & Herndl 2000). This is substantially lower than the decrease in fluorescence of bacterial-derived DOM upon exposure to solar radiation measured in this study (78 and 40%, Fig. 4).

Re-growth experiments of bacterioplankton did not indicate enhanced bioavailability of solar radiation-exposed bacterial-derived DOM (Fig. 3). Only in the carbon amended cultures could a significant growth of the bacterioplankton be observed independent of the exposure to solar radiation. Contrasting effects of solar radiation on the bioavailability of marine DOM have

been previously reported (Benner & Biddanda 1998, Obernosterer et al. 1999, 2001a). It was found that originally more labile DOM becomes more refractory upon solar radiation, while more refractory parent DOM becomes more labile (Obernosterer et al. 1999). Extrapolating these findings to our experiment, we would have expected a significant increase in the bioavailability of solar radiation-exposed bacterial-derived DOM in the re-growth experiments. The lack of any significant changes in the bioavailability of bacterial-derived DOM upon solar radiation as compared to the DOM held in the dark indicates that either the production of growth stimulating and inhibiting photo-products were balanced or, alternatively, the photo-products formed were not accessible for the bacterial community. At present we cannot resolve the underlying processes leading to this lack in growth response of bacteria to solar radiation-exposed bacterial-derived DOM but similar observations have been made with bovine serum albumin exposed to solar radiation (Obernosterer et al. 1999).

In summary, we found that bacterioplankton are producing refractory DOM and that this DOM is photo-reactive as indicated by the rapid loss in fluorescence upon exposure to solar radiation. This photoreactivity, however, does not lead to changes in the bioavailability of bacterial-derived DOM. Taken together, our findings indicate that bacterioplankton metabolism might be a significant source of refractory DOM in the ocean. The fate of this DOM in the oceanic water column, however, remains unclear.

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