The role of cell-surface-bound phosphatases in species competition within natural phytoplankton assemblage: an *in situ* experiment

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ABSTRACT

Despite it is widely acknowledged that the ability to hydrolyze dissolved organic matter using extracellular phosphatases is diverse in freshwater phytoplankton, the competition within single species related to presence and quantity of cell-surface-bound phosphatases has not been examined in natural conditions yet. Here, we studied phytoplankton species competition in a freshwater reservoir during an in situ experiment. A natural plankton community, with the exclusion of large zooplankton, was enclosed in permeable dialysis bags inside two large containers of different bioavailable phosphate concentrations. Phytoplankton species biomass and the abundance of bacteria were determined in purpose to compare the development of enclosed microbial communities. Total and cell-surface-bound phosphatase activities in the phytoplankton were investigated using the Fluorescently Labelled Enzyme Activity (FLEA) technique that allows for direct microscopic detection of phosphatase-positive cells and, with image cytometry, enables quantification of phosphatase hydrolytic capacity. Production of extracellular phosphatases was not completely inhibited or stopped in the phosphate-enriched environment, phytoplankton cells only showed the activity less often. Under the phosphate-nonenriched conditions, the production of phosphatases was enhanced, but active species did not proliferate amongst phytoplankton assemblage. Further, specific growth rates of the phosphatase-positive species in the non-enriched environment were lower than the same phosphatase-positive species in phosphate-enriched environment. Interestingly, the phosphatase-positive cells of Ankyra ancora increased their size in both treatments equally, although the population in phosphate-enriched environment grew much faster and the cell-specific phosphatase activity was lower. We hypothesize that brand new daughter cells had sufficient phosphorus reserves and therefore did not employ extracellular phosphatases until they matured and needed extra bioavailable phosphorus to support their metabolism before cell division. Based on presented in situ experiment, we propose that the ability to hydrolyze organic polymers and particles with cell-surface-bound phosphatases is advantageous for longer persistence of given population in a phosphate-scarce environment; although phosphatase-positive species cannot dominate the reservoir phytoplankton solely because of specific phosphorus-scavenging strategy.

Key words: competition, ectoenzyme, ELF97 phosphate, image cytometry, species-specific activity

1. INTRODUCTION

Species diversity in phytoplankton assemblages have been intensively studied for decades because phytoplankton composition influences the aquatic food web as well as the water quality in terms of, e.g., algal blooms. The availability of nutrients is considered to be a crucial factor regulating species diversity. When the intracellular storage of nutrients is exhausted, growth rates are regulated by the external supply rate of the limiting nutrient (for review, see Reynolds 1997). The most often growth-limiting factor in freshwaters is phosphorus. Its bioavailable form, orthophosphate (P_i), can be depleted rapidly in freshwaters (Hecky & Kilham 1988). Many algal and cyanobacterial species synthesise extracellular phosphatases when they lack P_i, because it allows access to organically bound P-compounds (Siuda & Chróst 1987; Jansson et al. 1988). Moreover, Chróst (1991) suggested that cell-associated extracellular enzymes might allow the producer to dominate nutrientdepleted ecosystem through its nutritional advantage.

The occurrence of extracellular phosphatase activity could diagnose starvation of phytoplankton (Kuenzler & Perras 1965; Dyhrman et al. 2002); cells that produce phosphatases are most probably without polyphosphate storage and lack P_i. Phosphatase activity can be directly visualised with a fluorogenic substrate, ELF[®]97 phosphate (ELFP, Huang et al. 1992). A phosphate group is hydrolysed from the water-soluble ELFP molecule in the presence of either alkaline or acid phosphatases (Štrojsová & Vrba 2006). The resulting fluorescent product of the hydrolysis is ELF alcohol (ELFA), which precipitates at or near the site of enzyme activity and thus can distinguish the cells or populations within phytoplankton assemblages that are able to scavenge P. The Fluorescently Labelled Enzyme Activity (FLEA) assay (in earlier articles imprecisely called as ELF technique, as well) has been used to study enzymatic activities of marine phytoplankton and algal cultures (Gonzáles-Gil et al. 1998; Dyhrman & Palenik 1999; Lomas et al. 2004, Dyhrman & Ruttenberg 2006; Štrojsová & Dyhrman 2008), freshwater phytoplankton



Fig. 1. Diagram of the experimental design. Left: open polyethylene container submerged in the reservoir water with three dialysis bags assigned as RDB. Right: container with reservoir water enriched with P_i and three dialysis bags assigned as PDB. RDB and PDB contained filtered reservoir water without large zooplankton.

(Rengefors *et al.* 2003; Štrojsová *et al.* 2003; Dignum *et al.* 2004; Cao *et al.* 2005; Štrojsová & Vrba 2008) and planktonic rotifers (Štrojsová & Vrba 2005, 2007). ELFA fluorescence of chosen objects can be quantified using epifluorescence microscopy and image analysis (Nedoma *et al.* 2003; Štrojsová *et al.* 2005).

Phosphorus enrichment bioassays have a long history in both fresh and marine waters (e.g., Wetzel 1981; Gonzáles 2000; Ojala *et al.* 2003; Rengefors *et al.* 2003). Orthophosphate is added and planktonic growth rates and biomass are compared relative to an unenriched control. The change in P_i concentration is typically accompanied by changes in phosphatase activity at the species level (Štrojsová *et al.* 2005). In most cases, enrichment results in increased biomass as well as a change in species composition (e.g., Arvola *et al.* 1996; Cottingham *et al.* 1998; Kobayashi & Church 2003).

In this *in situ* enclosure experiment, we examined effects of P_i enrichment on phytoplankton extracellular phosphatases, including bacteria, as important components of microbial community in the Římov Reservoir. We conducted the *in situ* experiment in late June, when P-limiting conditions usually develop. Alkaline phosphatase activity was detected and quantified using the FLEA assay and image cytometry. Two main objectives were addressed: (i) how does enrichment affect both plankton biomass dynamics and extracellular phosphatase activities at the species level? and (ii) are species that produce cell-associated phosphatases in advantage to the other, non-active species in natural phytoplankton assemblage, e.g., reaching extra P_i from dissolved organic phosphorus (DOP) compounds, could these species increase their specific growth rates?

2. MATERIAL AND METHODS

2.1. Study site and experimental design

The eutrophic Římov Reservoir is situated on the Malše River (Czech Republic; 48°83'N; 14°46'E). The main reservoir characteristics are: area 2.06 km², vol-

ume 34.5×10^6 m³, length 13.5 km, max./mean depth 43/16.5 m, mean retention time 100 days (Vrba *et al.* 1993). P-limiting conditions develop regularly during summer stratification in the euphotic zone.

The enclosure experiment was conducted on 18-25 June 2004 after the clear-water phase, which was characterised by the dominance of cryptomonad populations (Cryptomonas reflexa, C. erosa, and Rhodomonas *minuta*). Design of the experiment is presented in a diagram (Fig. 1). Water was sampled with a Friedinger sampler from a depth of 0.5 m at a sampling site located in the dam area. It was then filtered through a sieve (200 µm) to exclude large zooplankton and collected in a 50 L container. Three dialysis bags (diameter 75 mm, 1216 kDa, Poly Labo, France), were filled with 2.5 L of water from the container and placed vertically into a transparent polyethylene container (~80% of ambient light intensity, 50 L) that was supposed to simulate the reservoir (initial ambient concentration 7.7 μ g SRP L⁻¹) and was filled with unfiltered epilimnetic water amended by K_2 HPO₄ (final concentration 62 µg P L⁻¹). These P-enriched dialysis bags were designated as PDB. Three other dialysis bags (designated as RDB, no nutrient addition) were placed vertically in a bottomless control container with four triangular holes (~10 cm) on the sides to enable continuous reservoir-water exchange.

From each dialysis bag, samples were taken for plankton, chemical, and enzyme analyses on days 0, 3, 5 and 7 between 8.00 to 9.00 a.m. and transported immediately to the laboratory. All enzyme assays were processed within the next 3 hours. Water samples (~300 mL) were taken for particulate organic carbon, total P (neither analysed at day 5), soluble reactive P (SRP), and chlorophyll-*a* analyses. In addition, both containers were sampled and analysed for SRP and chlorophyll-*a*. Phytoplankton and rotifers were determined in the same samples (20 mL) preserved with Lugol's solution; other samples (20 mL) for bacteria determination were fixed with formaldehyde (2% final concentration).

2.2. Phosphatase assays

Three assays were used to determine phosphatase activity in the dialysis bags: (i) the MUF assay for measuring total phosphatase activity (PA_T) ; (ii) the FLEA assay for phosphatase activity detected on filters (PA_F) ; and (iii) the FLEA assay modified for measuring of cell-surface-bound phosphatase activity of single cells of phytoplankton (here *Ankyra ancora*, PA_A).

Total phosphatase activity (PA_T) was assayed with the common spectrofluorimetric method (Hoppe 1983) using 4-methyllumbelliferyl phosphate (MUFP, Glycosynth, UK). Duplicates of water samples were buffered with Tris/HCl buffer (pH 7.5; 10 mmol L^{-1} final concentration), supplemented with MUFP (100 μ mol L⁻¹ final concentration) and incubated for 2 h at 20 °C. The incubation was terminated by HgCl₂ addition (4 mmol L^{-1} final concentration). Parallel duplicates supplemented only with HgCl₂ were always incubated as blanks (Christian & Karl 1995). Prior to fluorescence reading, an alkaline solution (40 mmol L^{-1} of NaOH with 8 mmol L^{-1} of EDTA final concentrations, Vrba *et* al. 1993) was added. Fluorescence was measured using a spectrophotometer, with a fluorometric device (excitation at 365 nm and emission 465 nm) and corrected for the blank.

The protocol by Nedoma et al. (2003) for FLEA was followed for measuring both PA_F and PA_A. Water samples (4 mL) were incubated with ELF[®]97 phosphate (ELFP, Invitrogen – Molecular Probes; 20 μ mol L⁻¹ final concentration) at 20 °C. Samples were buffered with Tris/HCl buffer (pH 7.5; 10 mmol L⁻¹ final concentration) to ensure ELF alcohol (ELFA) precipitation. The incubation started within 1 hour after sampling. After a 2 h incubation, the samples were fixed with HgCl₂ (4 mmol L^{-1} final concentration) to prevent destruction of fragile flagellates and filtered over mild vacuum (<20 kPa) through a membrane filter (0.8 µm pore size). Filters with retained plankton were inspected for the presence of ELFA precipitates with an epifluorescence microscope (Olympus AX-70, Japan) with the UV-excitation filter set (excitation / emission: 360-370 nm / >420 nm). Both green ELFA fluorescence and red autofluorescence of chlorophyll-a were visible using this filter set.

A set of 40 images with ELFA fluorescence was taken from randomly selected areas $(470 \times 290 \ \mu\text{m}$ to quantify the phosphatase activity on a filter (PA_F) using the ELFA-specific filter set (excitation / emission: 360–370 nm / 520–540 nm). To quantify phosphatase activity of *Ankyra ancora* (PA_A), a set of images was taken to obtain typically 25 or more cells. The Lucia G/F 4.8 image analysis software (Laboratory Imaging, Czech Republic) was used for image processing and fluorescence quantification (more detailed description in Nedoma *et al.* 2003). Relative ELFA fluorescence in fluorescence units (FU) on a chosen area was calculated from the mean grey of the area, the mean grey of the

background, camera sensitivity, intensity of excitation light, and the objective-specific correction factor. ELFA (in fmol) was determined by multiplying the relative ELFA fluorescence by a conversion factor of 2.4 fmol FU^{-1} , determined by comparing the fluorescence of the same samples measured by image cytometry and in a spectrofluorimeter (Nedoma *et al.* 2003). The PA_A of *Ankyra ancora* was normalised per unit of plane projection area of each active cell measured.

2.3. Other methods

Soluble reactive phosphorus (SRP) was determined spectrophotometrically by the molybdate method (Murphy & Riley 1962). Total and dissolved phosphorus (TP and DP) were determined by perchloric acid digestion and the molybdate method according to Kopáček & Hejzlar (1993). Particulate phosphorus (PP) was determined as the difference between TP and DP and dissolved organic phosphorus (DOP) as the difference between DP and SRP. Particulate organic carbon (POC) was analysed by combustion of the glass-fibre filter with the retained particulate organic matter in a TOC 5000A analyser (Shimadzu, Kyoto, Japan). Molar C:P ratios were calculated from PP and POC. Concentration of chlorophyll-a (chl-a) was determined fluorimetrically after acetone extraction using the whole-water technique (Vyhnálek 1994). Unfiltered phytoplankton samples preserved with acid Lugol's solution were quantified in Utermöhl's sedimentation chambers on an inverted microscope after at least oneday sedimentation. Species-specific cell volumes, including flagellates (3–20 µm of cell diameter), were calculated from size measurements of living cells using geometric volume approximation. A conversion factor of 1 mg mm⁻³ of algal fresh mass was assumed. Specific growth rates of selected phytoplankton species were determined as the slope of the linear regression of the natural logarithm of cell density versus time. Bacteria were counted after DAPI staining using epifluorescence microscopy (Porter & Feig 1980). Bacterial biovolumes were measured with image analysis (Lucia G/F 4.8, Psenner 1993); cellular carbon content was calculated according to Norland (1993).

An unpaired *t*-Test was performed to compare triplicate means of PA_F and PA_A and phytoplankton biomasses determined from PDB and RDB. The relationships between PA_F or PA_A and TP, DOP or SRP were tested with a nonparametric Spearman rank correlation. Qualitative differences between the numbers of phosphatase-positive and non-positive taxa were tested using contingency tables and a χ^2 test. The above analyses were performed using Prism 4.0 (GraphPad, CA).

3. RESULTS

The proportions (abundance) of phytoplankton groups changed similarly in the non-enriched RDB and enriched PDB treatments during the first three days of



Fig. 2. Development of phytoplankton and rotifers during the *in situ* experiment: A: Phytoplankton biomass (as fresh mass, FM); and **B**: The most abundant rotifers. Bars represent means of triplicates in RDB (R) and PDB (P); S is the initial situation in the reservoir water.

the experiment. Initially, phytoplankton were dominated by cryptomonads, which were replaced by other populations, namely cyanobacteria, diatoms and flagellates, within three days (Fig. 2A). Overall, phytoplankton biomass increased more in the enriched PDB than in RDB by the day 7 (*t*-Test, P = 0.041). Moreover, fewer phytoplankton species were phosphatase-positive (Tab. 1) and specific growth rates of particular species were higher in PDB by the end of the *in situ* experiment (Tab. 2).

Initial DOP and SRP concentrations in the dialysis bags were 19.8 μ g L⁻¹ and 7.5 μ g L⁻¹, respectively (Fig. 3A) whereas the ambient SRP concentration in the Pcontainer reflected the P addition. The concentration of SRP in the P-container decreased after 5 days to 6.25 μ g L⁻¹ and after 7 days reached 2.9 μ g L⁻¹, a similar value to that in the R-container (2.2 μ g L⁻¹; Fig. 3A). The initial addition of P_i did not change measurably SRP levels in the PDB; SRP even declined (2.6 μ g L⁻¹) similarly as in the non-enriched RDB (2.6 μ g L⁻¹) by day 3. On the other hand, TP increased in both treatments, but more in the PDB (55.7 μ g L⁻¹ and 79.2 μ g L⁻¹; Fig. 3A). Initially, the C:P ratio was 144 and decreased by the end of the experiment in both RDB and PDB (C:P = 132 and 107, respectively).

Initial chl-*a* concentration in the dialysis bags was $36.9 \ \mu g \ L^{-1}$, which then decreased in all variants by day

3 (9–12 µg L⁻¹, Fig. 3B). After that, the decline continued in the R-container, which equalled that of open water, being only 5.8 µg L⁻¹ at day 7. There was an increase in chl-*a* by day 7 (17.5 µg L⁻¹) in the RDB. Chl-*a* increased in the P-container four times since day 3 (47 µg L⁻¹), while its concentration increased in the PDB by day 5 and then decreased to the same level as in the RDB (Fig. 3B).

3.1. Plankton composition and development

Species composition of the phytoplankton is given in table 1. Cryptophyceae (*Cryptomonas erosa* and *C. reflexa*) dominated the phytoplankton at the beginning of the enclosure experiment (>85% of total biomass), while Cyanobacteria, Bacillariophyceae, Chlorophyceae and flagellates were rather negligible (altogether <15%; Fig. 2A). Cryptomonads decreased rapidly in the dialysis bags; they declined to <5% of total biomass in RDB by day 7 (from the initial 4.2 mg L⁻¹ to 0.08 mg L⁻¹; Fig. 2A). Flagellate biomass increased both in the RDB and PDB samples (from the initial 0.54 mg L⁻¹ to, respectively, 2.2 mg L⁻¹ and 3.4 mg L⁻¹); maximum of flagellates (3.8 mg L⁻¹) occurred in RDB on day 3 followed by a decrease (Fig. 2A).

Tab. 1. Summary of phytoplankton species/genera present in the Římov Reservoir (Start), and in non-enriched (RDB) and enriched (PDB) dialysis bags, and inspected for ELFA labelling. Symbols: (x) several cells of the population ELFA-labelled, (xx) >10% of the population ELFA-labelled, (o) present but not labelled.

	Start	RDB	PDB
Cvanobacteria			
Anabaena planctonica Brunnth.	Х	х	х
Merismopedia sp.	0	0	0
Planktothrix agardhii (Gom.) Anagn. et Kom.	0	0	0
Pseudanabaena limnetica (Lemm.) Kom.	0	0	0
Snowella litoralis Häyrén (Kom. et Hind.)	0	х	0
Woronichinia naegeliana (Unger) Elenk.	0	0	0
Chrysophyceae			
Synura sp.	0	0	0
Dinophyceae			
Amphidinium sp.	0	0	0
Gymnodinium uberrimum (Allman) Kofoid et Swezy	0	Х	0
Bacillariophyceae			
Asterionella formosa Hass.	0	х	0
Aulacoseira italica (Ehrenb.) Simonsen	0	Х	0
Cyclotella pseudostelligera Hust.	0	0	0
Fragilaria crotonensis Kitt.	0	Х	0
Nitzschia acicularis W. Smith	0	0	0
Stephanodiscus hantzschii Grun.	0	0	0
Cryptophyceae			
Cryptomonas erosa Ehrenb.	0	0	0
Cryptomonas reflexa Skuja	0	0	0
Rhodomonas minuta Skuja	0	0	0
Chlorophyceae			
Ankyra ancora (G.M. Smith) Fott	XX	XX	XX
Crucigeniella apiculata (Lemm.) Kom.	0	Х	Х
Dictyosphaerium sp.	0	Х	0
Elakatothrix genevensis (Reverd.) Hind.	0	0	0
<i>Eudorina elegans</i> Ehrenb.	Х	Х	0
Gonium sociale Warm.	0	Х	Х
Chlamydomonas spp.	0	0	0
Chlorogonium sp.	0	0	0
Micractinium pusillum Fres.	Х	Х	0
Monoraphidium contortum (Thur.) KomLegn.	0	0	0
Monoraphidium minutum (Nag.) KomLegn.	0	0	0
<i>Oocystis parva</i> W. et G.S. West	0	Х	х
Panaorina morum (O.F. Muller) Bory	0	х	Х
Pediastrum boryanum (Turp.) Menegh.	0	Х	х
Pediastrum auplex Meyen	X	0	X
Pealastrum tetras (Enrend.) Kalls	0	0	0
r iankiosphaeria gelalinosa (J.WI. Smith	0	X	X
Sceneuesmus sp. Totraadron ingus (Tail.) G.M. Smith	X	x	x
Tetrastrum sp	0	0	U
Terusrum sp.	0	0	A 1.1
I OTAL NO. OF ELFA-positive taxa	6	1/	11

Tab. 2. Mean species-specific growth rates for the phytoplankton species in both non-enriched (RDB) and enriched (PDB) dialysis bags; \pm standard error. The PDB:RDB ratio implies the difference between the treatments. Median of specific growth rates for all the species in R-dialysis bags was 0.11 day⁻¹ and 0.32 day⁻¹ for the same species in P-dialysis bags.

	RDB (day ⁻¹)		PDB (day ⁻¹)		PDB:RDB
Anabaena planctonica	0.11 ± 0.19	+	0.39 ± 0.08	+	3.5
Asterionella formosa	0.22 ± 0.06	+	0.31 ± 0.06	-	1.4
Aulacoseira italica	0.09 ± 0.09	+	0.17 ± 0.21	-	1.9
Nitzschia acicularis	0.40 ± 0.08	-	0.47 ± 0.21	-	1.2
Ankyra ancora	0.11 ± 0.06	+	0.36 ± 0.16	+	3.3
Elakatothrix genevensis	0.17 ± 0.08	-	0.11 ± 0.02	-	0.6
Micractinium pusillum	0.03 ± 0.17	+	0.29 ± 0.04	-	9.7
Monoraphidium contortum	0.11 ± 0.11	-	0.14 ± 0.10	-	1.3



Fig. 3. Selected parameters measured in RDB (open symbols) and PDB (solid symbols) or in containers during the *in situ* experiment: **A**: Soluble reactive phosphorus (SRP, left, note logarithmic scale) and dissolved organic phosphorus (DOP, right scale) concentrations. **B**: Chlorophyll-*a* concentration (chl-*a*). **C**: Bacterial biomass (BB, left scale) and abundance (BA, right scale) in dialysis bags. **D**: Proportion of bacteria >5 μ m in bacterial biomass (as μ g C L⁻¹). *t*-Test results showed significant differences between RDB and PDB at day 7, probability level: *, *P* <0.05. All measurements in the dialysis bags were means of triplicates, error bars show standard error.



Fig. 4. The most abundant phytoplankton species during the *in situ* experiment: comparison of their proportions in the total phytoplankton biomass in non-enriched (RDB) and enriched (PDB) dialysis bags at day 7. Bars represent means of triplicates; *: significantly different values (*t*-Test, P < 0.05).

Cyanobacteria, namely *Anabaena planctonica*, initially formed <1% of total biomass (0.1 mg L⁻¹); by the end of the study, *A. planctonica* was more successful in PDB, where formed 22% of total biomass (1.8 mg L⁻¹), compared to 6.7% in RDB (0.28 mg L⁻¹; *t*-Test, P = 0.023; Fig. 4). The difference in Cyanobacteria biomass between RDB and PDB was significant by day 7 (*t*-Test, P = 0.023). Diatoms *Asterionella formosa*, *Aulacoseira*

italica and *Nitzschia crotonensis* formed initially less than 5% of total biomass (Fig. 2A). All three species were relatively more successful in the RDB treatment by day 7 (17%, 3%, and 8.5% of total biomass, respectively); however, the differences were nonsignificant (Fig. 4). Therefore, diatoms were the second most abundant group in RDB after flagellates at the end of the experiment, while cyanobacteria, chlorophytes and

flagellates were more abundant in the PDB samples (Fig. 2A). Chlorophyceae had the largest species diversity (20 taxa), although their biomass was not high during the whole experiment (initial 0.32 mg L^{-1} ; final 0.36mg L^{-1} in RDB and 1.08 mg L^{-1} in PDB). Ankyra ancora, Micractinium pusillum, and Monoraphidium contortum were the most abundant green algae in both treatments by day 7, while the first two reached higher biomass in PDB (Fig. 4). The difference in biomass was significant only for *M. pusillum* (*t*-Test, P = 0.045). Bacterial biomass grew very similarly in both RDB and PDB; it increased from an initial 79 μ g C L⁻¹ to a final 102 μ g C L⁻¹ (Fig. 3C). While bacterial number decreased twice during the experiment (Fig. 3C), the proportion of large bacteria (>5 µm) increased from 2.3% to 27% in RDB and to 17% in PDB (Fig. 3D).

than in RDB by day 5, while the opposite was the case for PA_F. No significant differences were found between triplicate means of PA_F determined from RDB and PDB (Fig. 5A). The PA_T of PDB significantly differed from that of RDB on days 5 and 7 (P = 0.0498 and 0.007, respectively). TP levels were correlated significantly with both PA_T (r = 0.857, P = 0.014) and PA_F (r =0.964, P = 0.001), on the other hand, DOP was not significantly linked to either PA_T or PA_F.

Table 1 summarises phosphatase activity for 38 phytoplankton species/taxa determined during the enrichment experiment. The ELFA labelling was rare among the phytoplankton, with only several cells of particular populations (except for *Ankyra ancora*) being found on each inspected filter (i.e., in 4 mL of water sample). At the beginning, *Anabaena planctonica* and



Fig. 5. Phosphatase activity measured in RDB (open symbols) and PDB (solid symbols) during the *in situ* experiment. **A**: Extracellular phosphatase activities PA_T , measured with the MUF assay, and PA_F , measured with the ELF assay. *t*-Test found significant differences between PA_T in RDB and PDB treatments at days 5 and 7, probability levels: *, P < 0.05; **, P < 0.01. **B**: Extracellular phosphatase activity of *Ankyra ancora* population (PA_A, left scale) and mean projection area of active cells (right scale). *t*-Test found significant differences between PA_A in RDB and PDB treatments at days 3, 5, and 7, probability levels: **, P < 0.01; ***, P < 0.001. All values are means of triplicates, error bars show standard error.

Rotifers formed significant part of the enclosed plankton. The most abundant taxa were *Polyarthra* spp., *Synchaeta* spp., *Keratella cochlearis*, and *Trichocerca cylindrica* (Fig. 2B). Initially, 8,000 individuals L^{-1} were in the dialysis bags. *Synchaeta* populations grew by more than six times by day 3 in both RDB and PDB treatments (17,300 and 21,000 individuals L^{-1} , respectively), further followed by their rapid decline (Fig. 2B). Other rotifer species continued growing; more rotifers were finally in RDB (21,000 individuals L^{-1}) than in the enriched PDB (14,000 individuals L^{-1}). Total rotifer numbers in RDB and PDB were not significantly different at day 7 (*t*-Test, P = 0.078), only *T. cylindrica* numbers differed significantly (*t*-Test, P = 0.011).

3.2. Extracellular phosphatase activity

The initial PA_T measured with the MUF assay and PA_F measured with the FLEA assay on filters were comparable (30–38 nmol L⁻¹ h⁻¹; Fig. 5A). Both PA_T and PA_F showed a similar pattern of continuous increase until day 5; PA_T then decreased in both variants, the decrease being greater in RDB. PA_T was higher in PDB

Micractinium pusillum were ELFA-labelled and were relatively abundant compared to the other labelled species. Populations of A. planctonica were partly and rarely ELFA-labelled both in RDB and PDB. Micractinium pusillum cells were sometimes ELFAlabelled in RDB only. Phosphatase-positive cells were observed among a number of chlorophytes (A. ancora, Crucigenia apiculata, Gonium sociale, Oocystis parva, etc.). The cells of A. ancora were labelled intensively and often; therefore it was possible to calculate PA_A (Fig. 5B). The highest PA_A was at day 3 in RDB (2.56 fmol $\mu m^{-2} h^{-1}$), which then declined by more than half by day 7 (1.16 fmol $\mu m^{-2} h^{-1}$). Ankyra ancora was less often ELFA-labelled in PDB, with the PA_A decreasing by day 5 (0.56 fmol μm^{-2} h⁻¹); however, it slightly increased afterwards (0.8 fmol $\mu m^{-2} h^{-1}$). The PA_A in RDB and PDB differed significantly on days 3, 5 and 7 (P = 0.0005, 0.001 and 0.009, respectively). The area of ELFA-labelled cells of A. ancora almost doubled in both treatments by day 3 and then remained stable (Fig. 5B). Cryptomonads, the most abundant phytoplankton group at the beginning of the study, were never ELFA-

labelled. Among the diatoms, Asterionella formosa, Aulacoseira italica and Fragilaria crotonensis were ELFA-labelled in RDB only. Finally, there were 17 ELFA-labelled taxa in RDB, while 11 taxa were labelled in PDB. The difference between initial and final numbers of ELFA-labelled taxa was significant in RDB ($\chi^2 = 7.54$, P = 0.003) but nonsignificant in PDB ($\chi^2 = 1.89$, P = 0.08). The difference in the number of ELFA-labelled taxa between RDB and PDB was nonsignificant ($\chi^2 = 2.04$, P = 0.077)

3.3. Specific growth rates

Specific growth rates of several phytoplankton species were significantly greater (P = 0.029) in PDB (median = 0.32 day⁻¹) than in RDB (median = 0.11 day⁻¹; Tab. 2). In all but one case (*Elakatothrix genevensis*), the growth rates were higher in enriched PDB than in RDB; however, the differences within a particular species population were all nonsignificant. Species that grew faster in PDB bags were either not producing phosphatases or the phosphatase activity was lower, as documented for *Ankyra ancora* (Fig. 5B). This was not proven for *Anabaena planctonica* since it was not possible to measure cell-specific phosphatase activity on its dense filaments.

4. DISCUSSION

Different ability to produce cell-surface-bound phosphatases could have fundamental implication for the adaptive biology of phytoplankton species. We investigated to what extent the production of extracellular phosphatases favoured single species in natural environment and, moreover, if this ability could help the species to proliferate amongst natural phytoplankton assemblage. Further, we compared the species-specific growth rates in two different bioavailable phosphate concentrations.

4.1. Dynamics of phytoplankton community

In the beginning of the experiment, similar trends were recorded in both the enriched PDB and nonenriched RDB samples: a rapid decline in cryptomonads and increases in flagellates and cvanobacteria during the first three days. Interestingly, the chl-a concentration in the reservoir water (i.e., in the open R-container) decreased similarly as in the dialysis bags, which could imply uniform development of the phytoplankton assemblage in the reservoir as well as in the enclosures. On the other hand, enclosures, even if being permeable dialysis bags, most probably generated different conditions inside as compared to the external environment within next few days. Still, our aim was to compare the development of the same initial phytoplankton assemblage split and exposed to different P_i concentrations, but under comparable environmental conditions. Extra P_i was added into the P-container and not directly into the dialysis bags, to study plankton response to a gradual input rather than to a single pulse of P. However, exchange through the permeable membrane was slow and bacteria and phytoplankton possibly took up majority of P_i already in the container and the microbial community inside the PDB utilised only a limited amount. On the other hand, it is very likely that some additional P_i could gradually enter the PDB due to accelerated P regeneration by the microbial community in the container. Thus, the plankton community in PDB could pump inward the P_i available from the P-container. As a result, phytoplankton biomass increased more in the PDB samples and also less species were phosphatase-positive.

Cryptophyceae were dominant at the beginning of the experiment. Cryptomonads occasionally form the majority of phytoplankton biomass during the clearwater phase in the Římov Reservoir (Komárková *et al.* 2003; Štrojsová *et al.* 2003). Regular sampling in July 2004 showed that cryptomonads formed a peak in the reservoir, which lasted even during the experiment (data of the Institute of Hydrobiology, České Budějovice). The rapid decrease in the abundant cryptomonad population could not be explained only by, e.g., rotifer grazing. Lethal enclosure effect on cryptomonads has been observed in several Czech reservoirs (P. Znachor, personal communication), which could be the main reason of the cryptomonad decline in our experiment.

Cyanobacteria, specifically Anabaena planctonica, formed a higher proportion of the total biomass in enriched PDB than in RDB. The N₂-fixing A. planctonica population might be favoured when Pi is amended and the N:P ratio is low (Levine & Schindler 1999). Although the N:P ratio was not determined in our study, the P-enrichment might shift the ratio to lower level, which might play a role in the success of A. planctonica. Relatively low cyanobacterial biomass in RDB compared to higher biomass of chlorophytes and diatoms might support this explanation. Large diatoms (Asterionella formosa, Aulacoseira italica, and Nitzschia acicularis) increased in abundance in both RDB and PDB. However, diatoms formed a higher proportion of the total biomass in RDB, especially A. formosa, which is tolerant of lower irradiance (Reynolds 1997), e.g., inside the container. Thingstad et al. (2005) suggested that the success of diatoms could be based on their "Winnie-the-Pooh" strategy: a high nutrient affinity relative to a large cell size that allows escape from predation. In this experiment, large zooplankton were removed and therefore diatoms were probably even more favoured. Diatoms were relatively less successful in PDB, probably because of stronger competition from the abundant cyanobacterium A. planctonica.

Planktonic rotifers in the Římov Reservoir have never reached such a high abundance (cf. Devetter & Sed'a 2003) as in the dialysis bags. Rotifers probably profited from their ability of fast adaptation to varying conditions (Herzig 1987) and high rate of reproduction (Cajander 1983). Moreover, ample food supplies along with no predation pressure by fish and large zooplankton favoured their increase. Increase in rotifer abundance (except for *Synchaeta* by day 7) in all dialysis bags implied rather tolerance to the enclosures as was previously observed by, e.g., Gonzáles (2000). Rotifers probably controlled the phytoplankton assemblage, because – especially in dialysis bags with higher rotifer numbers (PDB at day 3 and RDB at days 5 and 7) – phytoplankton biomass was lower. Despite this relationship was not significant, abundant rotifer populations most probably intensified phytoplankton biomass turnover and nutrient regeneration in the dialysis bags.

4.2. Phosphatase activity

The current concept of extracellular phosphatase activity regulation predicts an inverse relationship between phosphatase activity and phytoplankton growth rate (Healey & Hendzel 1979; Olsen *et al.* 1983) or cellular P (Gage & Gorham 1985). Low phosphatase activity is assumed to be a consequence of the repression of phosphatase synthesis caused by high cellular P content (e.g., Chróst 1991) and high cellular P content leads (if microbial growth is P-limited) to a high growth rate (Droop 1973). Conversely, high phosphatase activity should occur at low cellular P content, which also implies a low growth rate (Vadstein *et al.* 1988).

Based on the above assumptions, extracellular phosphatase activity has been proposed as an indicator of Plimited phytoplankton; for such indication, bulk phosphatase activity (e.g., PA_T) used to be normalised per phytoplankton biomass (e.g., Healey & Hendzel 1980; Gage & Gorham 1985; Chróst 1991). Superficial interpretation of the above, however, may lead to an incorrect assumption that the whole P-limited phytoplankton react this way and plankton ecologists should be aware of this oversimplification. Recent FLEA studies have suggested very diverse species-specific reactions in the same ambient P_i concentrations (Rengefors et al. 2003; Cao et al. 2005; Štrojsová et al. 2003, 2005; Dyhrman & Ruttenberg 2006). Moreover, the activity of cell-surface-bound phosphatases, produced by particular algal population, could fluctuate on the time scale of several hours (Štrojsová & Vrba 2008). Thus, phytoplankton could cause explicit short-term variation in the bulk activity (PA_T) . Therefore, we would like to warn that PA_T alone could be misleadingly interpreted as an indicator of P deficiency/sufficiency. We suggest more frequent sampling for phosphatase analysis or to use the FLEA assay to evaluate an actual proportion of phosphatase-positive phytoplankton (or bacterioplankton) species.

Initial PA_F and PA_T were low, as is usual during spring and/or early summer in the Římov Reservoir (Vrba *et al.* 1993; Štrojsová *et al.* 2003). PA_F was higher in RDB by day 3, as was expected; however, the difference was nonsignificant. PA_T was very similar in both variants. PA_F and PA_T increased in both treatments by day 7, however, significantly more in PDB. This could be explained by the equally low SRP in all dialysis bags starting at day 3, whereas phytoplankton biomass was higher in PDB. The more abundant phytoplankton in PDB produced less phosphatase per biomass unit. No direct relationships between PA_T , PA_F and DOP or SRP were found, suggesting that the rate of enzymatic hydrolysis did not depend on ambient SRP concentration or that P_i concentration was sufficient for the majority of phytoplankton, as implied by the observed C:P ratios. Therefore, the internal microbial P pool probably regulated the production of phosphatases (Caruso *et al.* 2005).

Chrysophyceae and Dinophyceae were not phosphatase-positive except in one case (*Gymnodinium uberrimum*). That could be caused by their different sensitivity to P availability and likewise their mixotrophic capacities (Reynolds 1997). Three diatom species (*Asterionella formosa, Aulacoseira italica,* and *Fragilaria crotonensis*) were phosphatase-positive in RDB only and grew slower than in enriched PDB, where P supply strongly supported their development (Buzzi 2002).

Ankyra ancora reduced its phosphatase activity in enriched PDB after Pi addition. When this extra ambient P_i was exhausted, the PA_A slightly increased again. Such an effect shows the flexibility and adaptability of A. ancora. A comparable influence of P concentration on phosphatase activity in populations of A. ancora, two cyanobacteria Microcystis aeruginosa and Aphanizomenon flos-aquae, and the diatom Fragilaria crotonensis was previously found in Římov Reservoir (Štrojsová et al. 2005). Interestingly, the phosphatase-positive cells of A. ancora increased their size in both treatments equally, although the population in PDB grew much faster and the PA_A was lower. We hypothesize that brand new daughter cells had sufficient phosphorus reserves and therefore did not employ extracellular phosphatases until they matured and needed extra bioavailable phosphorus to support their metabolism before cell division.

Lower PA_F, compared to PA_T, was already found in our earlier study (Štrojsová *et al.* 2005). This phenomenon can be caused by (i) different ELFP and MUFP hydrolysis rates and (ii) the ELFP concentration (20 μ mol L⁻¹) used. The ELFP hydrolysis rate only slightly exceeded 50% of the maximum velocity at this concentration (Nedoma *et al.* 2003, 2007). The chosen ELFP concentration was a compromise between assay sensitivity and affordability. A lag phase before an increase in ELFA fluorescence occurred (Nedoma *et al.* 2003). Thus, some amount of ELFA molecules could pass through the filter when not precipitating. Nevertheless, differences in the *A. ancora* cell-associated ELFA fluorescence measured in this study reflect actual differences in PA_A.

P_i addition caused increased phytoplankton growth and partial depression of phosphatase production. In the non-enriched environment, phytoplankton species produced extracellular phosphatases more often and also their specific growth rates were lower; growth was constrained by lower P_i supply and probably the extra energy invested in synthesis of these enzymes.

The difference in extracellular phosphatase production was shown within different algal groups, for instance, Cyanobacteria or Chlorophyceae were phosphatase-positive more often than Chrysophyceae or Cryptophyceae in three consecutive years (Štrojsová et al. 2003). Contrasting growth strategies (survivalists vs bloomers) and/or distinct resource co-limitation of the phytoplankton species (Arrigo 2005), and also physiological history of individual cell, internal P pool, and/or the need of extra P_i (e.g., before cell division) may reflect the heterogeneity in phosphatase production. Nevertheless, extracellular phosphatases advantage their algal or cyanobacterial producer in competition for resources in a situation when internal reserves or ambient P_i is scarce, because only the cells that can produce these enzymes will be able to cleave P_i from DOP, and therefore survive or accomplish more cell divisions, even though their growth rates may be suboptimal. Based on our data, we assume that the threshold concentration of P_i under which cell activates the production of the enzyme is distinct among phytoplankton species and would be the most interesting topic for future laboratory studies. Here we have shown that, under conditions where P_i was present in lower concentrations, the production of phosphatases was enhanced, but these active species did not proliferate amongst phytoplankton assemblage. The regulation of the cell-surface-bound enzyme activity and the proportion to which the enzyme is present in different phytoplankton species populations needs particular attention and further investigation.

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