

## Bacterivory of metazooplankton, ciliates and flagellates in a newly flooded reservoir

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**Abstract.** Bacterial consumption by metazoan zooplankton and phagotrophic protists was measured *in situ* during the period of thermal stratification in the epilimnion (1 m) and metalimnion (7 m) of a newly flooded reservoir (Sep reservoir, France). The mean bacterial consumption was  $2.53 \times 10^6$  bacteria  $l^{-1} h^{-1}$  at 1 m and  $0.97 \times 10^6$  bacteria  $l^{-1} h^{-1}$  at 7 m. The main consumers over the whole study period were the cladocerans *Daphnia longispina* and *Ceriodaphnia quadrangula*, accounting on average for 72% of the potential total predation of bacteria at 1 m and 56% at 7 m, especially during the months of May–June and August. Heterotrophic nanoflagellates (HNF), which accounted for 12% estimated total predation of bacteria at 1 m and 13% at 7 m, only exerted a limited predation, mainly by a *Monas*-type cell. Ciliates, dominated in terms of abundance by *Pelagohalteria viridis*, accounted for 4% of total predation in the epilimnion ( $0.00$ – $0.42 \times 10^6$  bacteria  $l^{-1} h^{-1}$ ). In a newly flooded reservoir, metazoan zooplankton appear to be the main consumers of bacteria. Predation of ciliates and HNF by zooplanktonic crustaceans could account for the low contribution of components of the microbial loop to bacterial consumption.

### Introduction

Heterotrophic bacteria form a large proportion of the biomass in pelagic ecosystems (Cho and Azam, 1988; Simon *et al.*, 1992). The small variations in bacterial abundance, despite the fact that bacterial generation times are of the order of a day in temperate regions (Pace, 1988), suggest that there is an effective regulatory system. Predation by protozoa (Sanders *et al.*, 1989), metazoan zooplankton (Güde, 1988; Arndt 1993; Jürgens, 1994), mortality caused by viruses (Proctor and Führman, 1992) and the quantity of dissolved organic carbon produced by algae (Cole *et al.*, 1988) are the most frequently cited regulatory mechanisms. Heterotrophic flagellates are generally thought to be the main consumers of bacteria in marine ecosystems (Fenchel, 1982) and fresh waters (Riemann, 1985; Güde, 1986), but phagotrophic phytoflagellates can also have an impact on bacteria and have grazing rates that are similar to those of heterotrophic flagellates (Bird and Kalff, 1986). In addition, ciliates can be important consumers of bacteria in freshwater (Simek *et al.*, 1995) and marine environments (Sherr *et al.*, 1989). Although protozoans, or more generally protists, are thought to be the main consumers of bacteria, some metazoans can nevertheless play an important role in regulating bacterial communities (e.g. Jürgens *et al.*, 1994). Among these, Cladocera, rather than Copepoda, are thought to be effective consumers of bacteria (Peterson *et al.*, 1978; Geller and Müller, 1981; Pace *et al.*, 1983; Güde, 1988; Jürgens, 1994), which can be an important food source during the summer (Pace *et al.*, 1983). Some field studies have highlighted their important predation activity (Børsheim and Olsen, 1984; Riemann, 1985; Bjørnson *et al.*, 1986), whereas others concluded that these crustaceans were only of minor importance in controlling bacterial populations (Toth, 1980;

Pedros-Alio and Brock, 1983). Although some rotifers do consume heterotrophic bacteria (Pourriot, 1977; Bogdan *et al.*, 1980; Ricci, 1984; Boon and Shiel, 1990; Ooms-Wilms *et al.*, 1995; Ooms-Wilms, 1997), their contribution to regulating the bacterial plankton is, however, modest (Sanders *et al.*, 1989; Pace *et al.*, 1990; Pernie *et al.*, 1990).

The literature therefore clearly shows that metazoans do consume bacteria, but little is known of the importance of this predation compared to that of heterotrophic and mixotrophic protists. Simultaneous observation of the succession of bacterial consumption by metazoans and protozoans has only rarely been attempted (Sanders *et al.*, 1989; Vaqué and Pace, 1992). Their role could also depend on the type of ecosystem being studied. For example, Paterson *et al.* (1997) suggested that metazoan zooplankton in a newly flooded reservoir could use organisms other than phytoplankton, such as bacteria, as an energy source.

In this study, we determined which taxa consumed bacteria-sized particles, estimated the filtration and ingestion rate of individual organisms on bacteria, and calculated the impact of grazing by protozoans and metazoans on the bacterial community. This study, therefore, provided information on the relative roles of the different predators in controlling the bacterial abundance in an ecosystem in a transitional stage.

## Method

The oligomesotrophic Sep reservoir (33 ha) situated in the Massif Central of France (46°2'N and 3°1'E) was first flooded in 1995 to provide irrigation water for crops. During the study period (April–August 1997), lake water was sampled every week using an 8 l Van Dorn-type bottle in the epilimnion (1 m) and metalimnion (7 m) at the deepest point in the lake. Estimations of grazing on bacteria by protozoans and metazooplankton were made every week from May to August. At the same time, the metazoan zooplankton in the water column were collected by taking three vertical hauls from the bottom to the surface using a Juday type net of 55 µm mesh size.

### *Abiotic and biotic variables*

The water temperature, dissolved oxygen content and pH were determined with a multiparameter probe (YSI GRANT 3800). Water transparency was estimated from the Secchi disc depth. Phosphorus (PO<sub>4</sub>-P), nitrates (NO<sub>3</sub>-N) and ammonium (NH<sub>4</sub>-N) contents were analysed using standard American Public Health Association (1992) methods. The chlorophyll *a* content was determined by spectrophotometry (Lorenzen, 1967; Strickland and Parsons, 1968). Dissolved organic matter was quantified after filtration through a 0.2 µm polycarbonate filter (Millipore). The dissolved protein concentration (DPROT) was determined using the micro BCA Protein Assay Reagent Kit (Pierce) and that of total dissolved carbohydrates (TDCHO) by Burney and Sieburth's (1977) and Johnson and Sieburth's (1977) method, after acid hydrolysis (1 N HCl; 100°C, 15 h).

*Sample storage*

The samples were fixed immediately after collection and stored at 4°C. Formaldehyde was used as fixative for bacteria (final concentration 2%), glutaraldehyde for flagellates (final concentration 1%), mercuric chloride for ciliates (final concentration 2.5%), and a mixture of formaldehyde and sucrose (final concentration 4%) for metazoans, the latter preventing release of eggs and physical deformation (Prepas, 1978).

*Counting organisms*

Heterotrophic bacteria were stained with DAPI (1 µg/l), then filtered through black polycarbonate membranes of 0.2 µm pore size (Millipore), using the protocol described by Porter and Feig (1980). After staining with primulin (final concentration 200 µg/ml) (Caron, 1983), flagellates were recovered on black polycarbonate filters of 0.8 µm pore size (Nuclepore). Both preparations were made within 24 h of sampling and were stored at -25°C to minimize losses of auto-fluorescence (Bloem *et al.*, 1986). Counts were made under an Olympus HBS epifluorescence microscope equipped with an epifluorescent HB2-RFL light source, an HBO-100 W mercury lamp and a neofluar 100/1.25 objective lens. Two types of light filters were used: UG-1, DM 400, L 435 (UV light) for heterotrophic bacteria and heterotrophic nanoflagellates (HNF), and BP 545, O 590 (blue light) for autotrophic nanoflagellates. In a preliminary series of triplicate counts of bacteria and flagellates (magnification ×1250) and ciliates (magnification ×500), the coefficient of variation for bacteria was <5%, when counting 500–800 bacteria in 30–60 fields, with an eyepiece fitted with a graticule delimiting the field. The coefficient of variation was 6% for pigmented flagellates (PF), 9% for HNF after counting 200–300 cells, and 12% for ciliates after counting 100–200 individuals. Large-sized and/or colonial ciliates and flagellates were counted using Utermöhl's (1958) method with a Leitz (Wild M40) inverted microscope. The entire counting field was examined at a magnification of ×500 for ciliates, within 2 months of sample fixation as recommended by Sime-Ngando and Grolière (1991). Ciliates were identified to genus or species level by reference to Kahl (1930–1935), Kudo (1966) and by consulting the works of Foissner (1994). The metazoan zooplankton were counted under a binocular microscope (Wild M3 Z) in a Dolfuss chamber. The dry weight of each taxon was calculated from the formulae of Bottrell *et al.* (1976).

*Estimating the predation rates of metazoan zooplankton and protozoans on the bacterial communities*

A stock solution of tracer particles 0.5 µm in diameter was prepared from a concentrated solution of Fluoresbrite Plain Microspheres (Polysciences). To ensure that the microspheres were well dispersed, the solution was treated with bovine serum albumin (BSA) at a concentration of 5 mg l<sup>-1</sup> (Pace and Bailif,

1987). The concentration of microspheres was estimated by epifluorescence microscopy, after filtration onto polycarbonate filters of 0.2 µm pore size.

The plankton from 1 and 7 m were acclimatized for 5 min in the lake in glass bottles with a capacity of 1 l for metazoan zooplankton (three replicates) and of 250 ml for protozoans (1–2 replicates). A concentration of microspheres of between 2 and 5% of the bacterial abundance in the lake (Ooms-Wilms *et al.*, 1995) was injected into each metazooplankton bottle, which was then turned upside down several times to mix. Preliminary experiments under laboratory conditions (20°C) showed that an incubation time of 5 min was less than the digestion time for the various metazoans. After incubation, each bottle was filtered through a nylon sieve of 55 µm pore size. To prevent any regurgitation, the metazooplankton were anaesthetized with carbonated water before fixation in a final concentration of 4% formaldehyde and 60 g l<sup>-1</sup> sucrose. For protozoans, a concentration of microspheres of between 8 and 12% (MacManus and Okubo, 1991; Carrias *et al.*, 1996) of the mean bacterial abundance in the lake was added to the 250 ml bottles. Preliminary time series experiments showed that bead uptake by protozoans was linear between 0 and 30–40 min. Thus, *in situ* incubations were stopped at 30 min by adding a cold solution of glutaraldehyde (2% final concentration) to prevent egestion of tracer particles (Sanders *et al.*, 1989). The plankton in the control bottles were fixed immediately after adding the tracer particles, to determine the concentration of particles adhering to organisms (background noise). The numbers of microspheres ingested by the metazooplankton, ciliates, microflagellates and colonial flagellates were determined under an epifluorescence microscope Leitz fluovert FU, filter A (513593, UV light) and in transmitted light after sedimenting 75–100 m of sample in counting cells. The counts of microspheres in protozoa were conducted using the same method as that described for ciliates. The numbers of microspheres ingested by metazooplankton were estimated by examining the entire alimentary tract at a magnification of ×125–250. The ingestion of tracer particles by phagotrophic nanoflagellates was determined after filtering two subsamples (10–25 ml) onto a black polycarbonate filter of 0.8 µm pore size. Two hundred to 300 cells were then examined using the method described above. Preliminary triplicate experiments showed that the coefficients of variation for the abundance of microspheres in protist organisms were 37% for HNF, 32% for pigmented flagellates and 26% for ciliates, whereas those in metazoans were 22% for *Daphnia longispina*, 8% for *Ceriodaphnia quadrangula*, 6% for *Bosmina longirostris* and 45% for rotifers.

The filtration [TF; µl individual (ind.)<sup>-1</sup> h<sup>-1</sup>], ingestion (TI; bacteria ind.<sup>-1</sup> h<sup>-1</sup>) and grazing rates (TGR; bacteria l<sup>-1</sup> h<sup>-1</sup>) for each taxon were calculated as follows:

$$TF = (M_t - M_0)/M \times T \text{ and } TI = TF \times (B + \text{microspheres})$$

$$TGR = TI \times \text{abundance of the taxon (l}^{-1}\text{)}$$

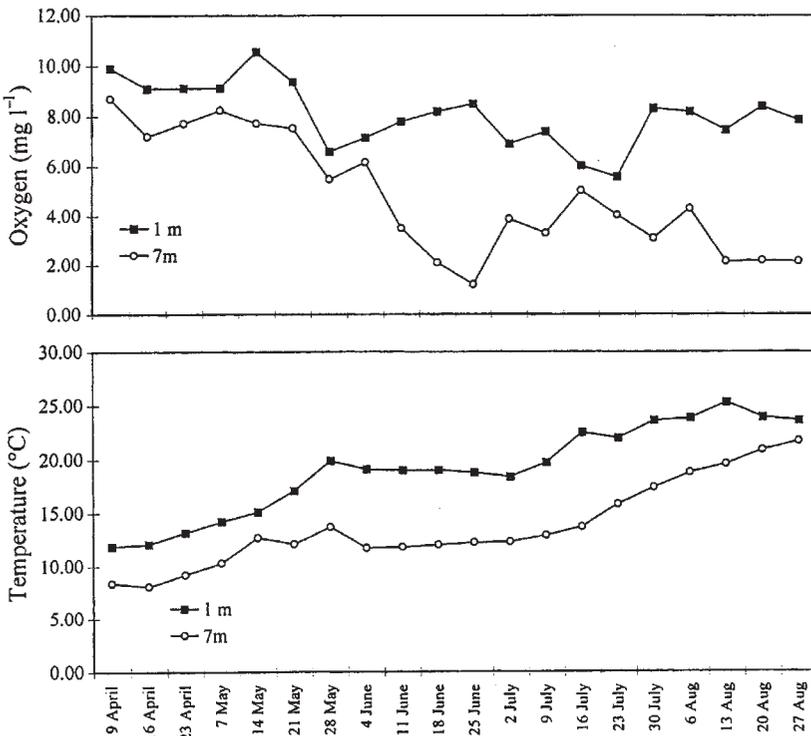
where  $M_t$  is the number of microspheres ingested per individual (microspheres ind.<sup>-1</sup>) at incubation time  $t$ ,  $M_0$  is the number of microspheres ingested per individual (microspheres ind.<sup>-1</sup>) at incubation time 0 (background noise),  $M$  is the

concentration of microspheres during incubation (microspheres  $\mu\text{l}^{-1}$ ),  $T$  is the incubation time (h) and  $B$  is the bacterial concentration during incubation (bacteria  $\mu\text{l}^{-1}$ ).

## Results

### *Abiotic and biotic variables*

Vertical oxygen and temperature profiles were measured at every metre. The values presented are those obtained at the depths of the experiments (Figure 1). The mean temperature in the epilimnion was  $19.2 \pm 4.3^\circ\text{C}$  and that of the metalimnion  $14.0 \pm 4.2^\circ\text{C}$ . The dissolved oxygen concentration at 1 m depth only changed slightly with time, with a mean value of  $8.1 \pm 1.3 \text{ mg l}^{-1}$ . At 7 m, the oxygen content declined from the start of measurements to reach a minimum value of  $1.20 \text{ mg l}^{-1}$  on 25 June; it then varied between 2.20 and  $5.03 \text{ mg l}^{-1}$  (mean  $3.4 \pm 1.0 \text{ mg l}^{-1}$ ). The mean concentrations of nitrates, ammonium and orthophosphates were  $1.59 \text{ mg N l}^{-1}$  ( $0.87\text{--}2.3 \text{ mg N l}^{-1}$ ),  $0.006 \text{ mg N l}^{-1}$  ( $0.003\text{--}0.009 \text{ mg N l}^{-1}$ ) and  $0.030 \text{ mg P l}^{-1}$  ( $0.015\text{--}0.047 \text{ mg P l}^{-1}$ ), respectively. The mean chlorophyll  $a$  concentration was  $1.9 \mu\text{g l}^{-1}$ , and fluctuated during the study period between 0.6 and  $10.5 \mu\text{g l}^{-1}$  (Figure 2). The DPROT and TDCHO concentrations varied



**Fig. 1.** Temporal changes in dissolved oxygen ( $\text{mg l}^{-1}$ ) and temperature ( $^\circ\text{C}$ ) at 1 and 7 m depths.

between 4.11 and 7.64 mg l<sup>-1</sup> (mean 5.88 mg l<sup>-1</sup>) and from 1.21 to 3.13 mg l<sup>-1</sup> (mean 2.04 mg l<sup>-1</sup>), respectively.

### Plankton community succession

**Bacteria.** The abundance of heterotrophic bacteria (Figure 2) varied from 1.4 to 4.5 × 10<sup>6</sup> cells ml<sup>-1</sup> (mean = 2.7 × 10<sup>6</sup> cells ml<sup>-1</sup>). The abundance was slightly higher at 1 m than at 7 m, except in April. Most of the bacteria were in the form of cocci, with a size between 0.4 and 0.6 µm.

**Flagellate and ciliate protists.** The various taxa (HNF, pigmented flagellates and ciliates) encountered in the Sep reservoir are shown in Table I. The mean

**Table I.** Size, biovolume, filtration rate and ingestion rate of the various taxa of protists

Protists	Mean length (µm)	Biovolume (µm <sup>3</sup> )	Bead uptake	Clearance (10 <sup>-3</sup> µl ind. <sup>-1</sup> h <sup>-1</sup> )			Ingestion (bacteria ind. <sup>-1</sup> h <sup>-1</sup> )		
				Min.	Max.	Mean	Min.	Max.	Mean
Ciliates									
Oligotrichida									
<i>Pelagohalteria viridis</i>	29.0	3450	Y	3.1	47.5	26.9	10.1	236.7	125.1
<i>Halteria</i> sp.	28.0	2520	Y	0.0	24.7	9.9	0.0	118.8	46.8
<i>Strobilidium caudatum</i>	21.5	3111	N						
<i>Strombidium viride</i>	51.0	23 459	Y			4.0			13.6
Prostomatida									
<i>Urotricha furcata</i>	20.5	1251	N						
<i>Balanion</i> sp.	13.5	1294	N						
Scuticociliatida									
<i>Cyclidium</i> sp.	13.0	618	N						
Undetermined Scuticociliatida	37.5	1193	N						
Haptorida									
<i>Askenasia volvox</i>	30.0	7650	N						
<i>Didinium</i> sp.	42.5	12 750	N						
<i>Paradileptus elephantinus</i>	280.0	714 737	N						
Peritrichida									
<i>Vorticella</i> sp.	55.5	17 028	Y	35.1	52.3	43.7	192.1	195.1	193.5
Suctorida	50.1	11 780	N						
Undetermined	35.0	22 444	N						
Pigmented flagellates									
<i>Chrysococcus</i> sp.	10.0	524	N						
<i>Chrysidalis</i> sp.	4.5	31	N						
<i>Cryptomonas ovata</i>	24.5	1856	Y	0.0	4.3	0.7	0.0	14.0	2.6
<i>Dinobryon cylindricum</i>	11.5	220	Y	28.4	51.6	31.8	63.2	137.6	103.8
<i>Mallomonas</i> sp.	20.0	1097	N						
<i>Pandorina morum</i>	12.5	1023	N						
<i>Peridinium volzii</i>	50.5	65 449	N						
<i>Rhodomonas minuta</i>	9.5	173	N						
<i>Synura</i> sp.	13.0	1050	N						
Heterotrophic nanoflagellates									
<i>Kathablepharis</i> sp.	5.8	65	N						
Monas-like cells	6.0	72	Y	0.0	3.3	0.7	0.0	14.1	2.5
Chrysomonadine (2–5 µm)	3.0	17	N						
Undetermined	4	26	N						

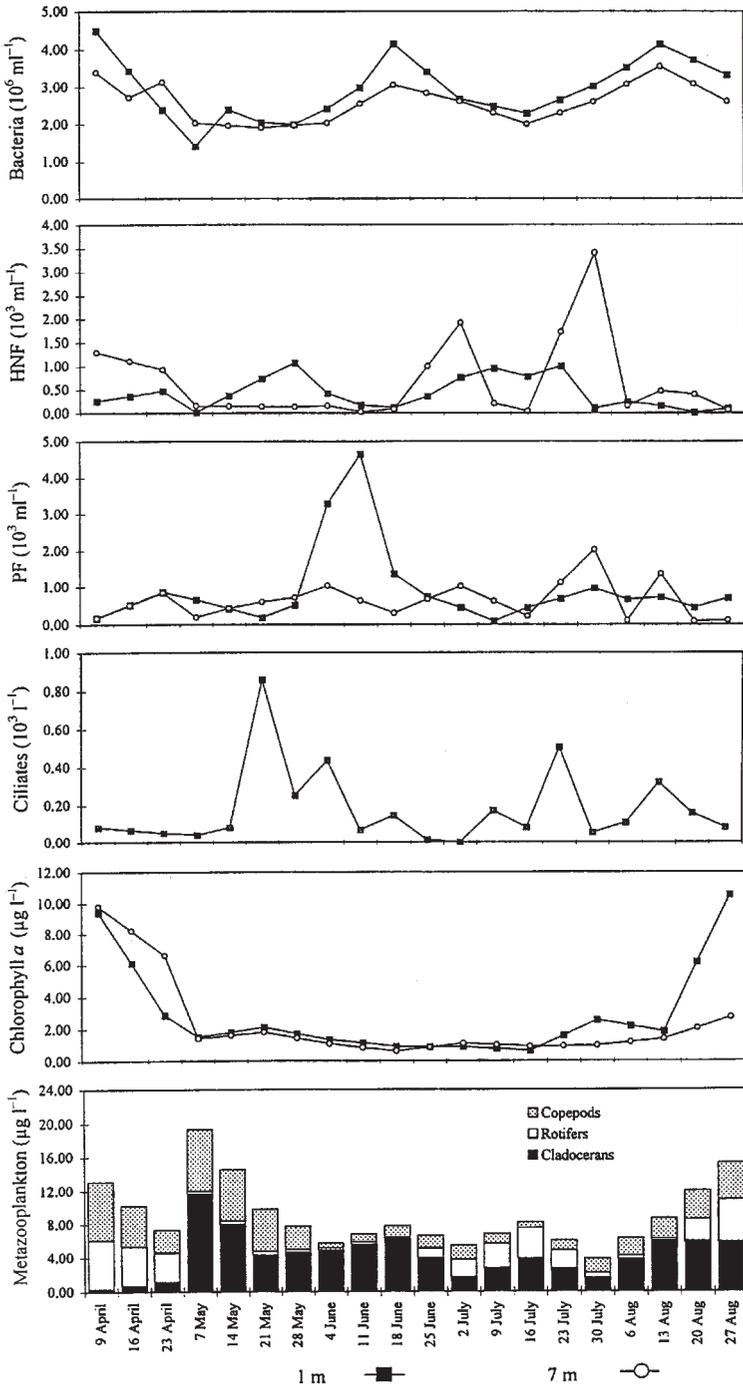
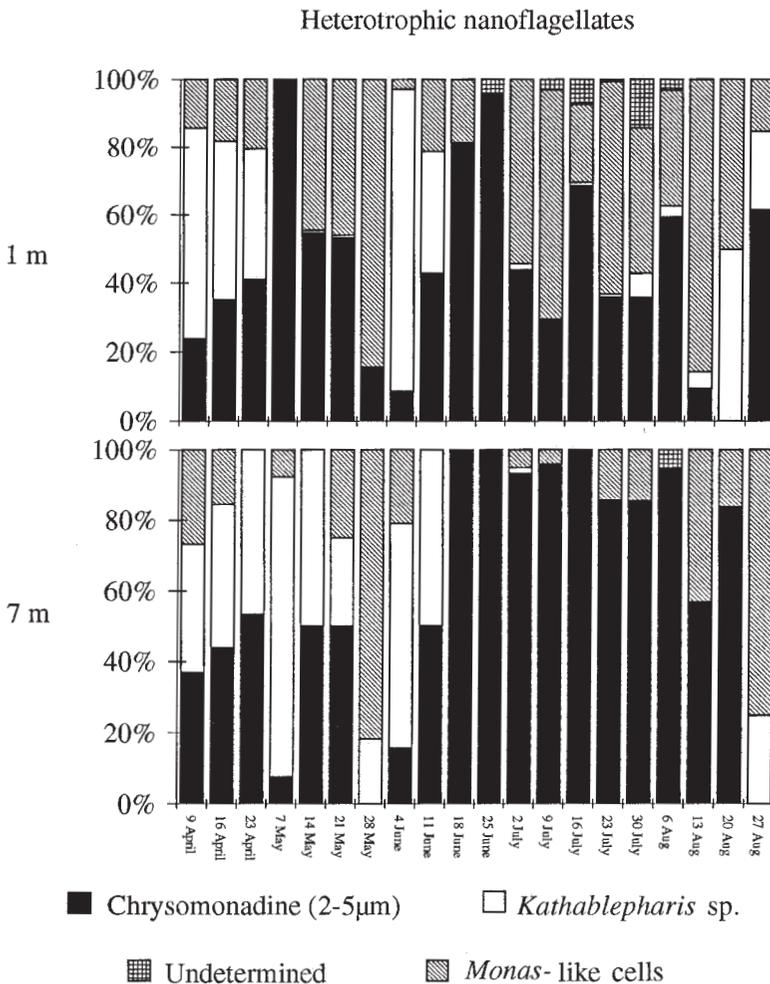


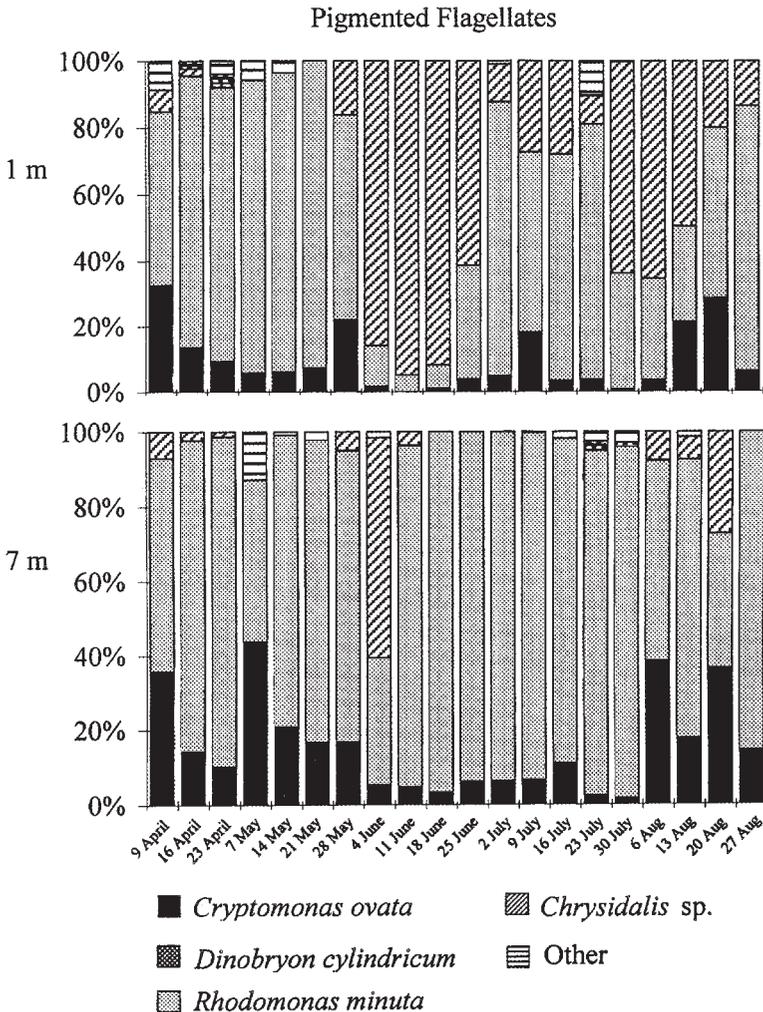
Fig. 2. Abundances of bacteria, heterotrophic nanoflagellates (HNF), pigmented flagellates (PF) and ciliates, and chlorophyll *a* at 1 and 7 m, and metazooplankton biomass in the water column.

densities of HNF measured during the study periods at 7 m ( $0.7 \times 10^3$  cells ml<sup>-1</sup>) were slightly higher than those at 1 m ( $0.4 \times 10^3$  cells ml<sup>-1</sup>). The highest densities of HNF in the epilimnion were recorded in May and July (Figure 2). In the metalimnion, the highest density was recorded at the end of July. This community (Figure 3) was dominated by cells of the chrysomonadine, *Monas*-like cells (46% at 1 m and 14% at 7 m) and a small unidentified heterotrophic flagellate (unidentified sp. 2) with a size of between 2 and 5 µm (41% at 1 m and 73% at 7 m). *Kathablepharis* sp. was mainly present in spring when it accounted for 13% of total HNF abundance (Figure 3). The density of pigmented flagellates (Figure 2), which fluctuated from 0.09 to  $4.6 \times 10^3$  cells ml<sup>-1</sup>, was higher at 1 m (mean  $1.0 \times$



**Fig. 3.** Relative abundance of heterotrophic nanoflagellates (HNF) at 1 and 7 m depths.

$10^3$  cells  $\text{ml}^{-1}$ ) than at 7 m (mean  $0.7 \times 10^3$  cells  $\text{ml}^{-1}$ ). The highest density was recorded in June in the epilimnion and in July in the metalimnion. *Chrysidalis* sp. (57% of abundance at 1 m and 6% at 7 m) and *Rhodomonas minuta* (36% of abundance at 1 m and 82% at 7 m) accounted for most of the total abundance of pigmented flagellates. *Cryptomonas ovata* was especially abundant in April–May and at the end of summer (Figure 4). Densities of ciliates varied from 0.04 to  $0.86 \times 10^3$  cells  $\text{l}^{-1}$  (mean  $0.18 \times 10^3$  cells  $\text{l}^{-1}$ ) (Figure 2), the highest densities being recorded in May. These protozoans were dominated in terms of density by a mixotrophic oligotrich, *Pelagohalteria viridis*, which on average accounted for 67% of cell abundance during the study period.



**Fig. 4.** Relative abundance of pigmented flagellates (PF) at 1 and 7 m depths.

*Metazooplankton.* The various taxa (crustaceans and rotifers) counted in the Sep reservoir are shown in Table II. The total biomass of metazoan zooplankton in the water column (Figure 2) varied from 3.9 to 19.3  $\mu\text{g l}^{-1}$  (mean 9.10  $\mu\text{g l}^{-1}$ ). In April, copepods (*Eudiaptomus gracilis* and *Cyclops vicinus*) and the rotifer *Polyarthra* sp. dominated the community in terms of biomass. Over the whole year, copepods consisted mainly of juvenile stages (nauplii and copepodites). The calanoid *E.gracilis* was the most abundant copepod from May onward, but most of the metazooplankton biomass was composed of Cladocera and especially *D.longispina*, which accounted for 99.7% of the biomass of these crustaceans from May to June. In July and August, *Hexarthra mira* and *Polyarthra* sp. were the dominant species of rotifers. *Ceriodaphnia quadrangula*, which only occurred in August, accounted for 36% of the cladoceran biomass at this time. When chlorophyll *a* reached values of  $<2 \mu\text{g l}^{-1}$  (Figure 2), from the end of May to July, the metazooplankton biomass remained low. This period coincided with the peaks in the abundance of HNF, ciliates, PF and bacteria. At the start of the study, the decrease in bacterial abundance also coincided with an increase in the biomass of *D.longispina*.

*Estimated consumption of bacteria by flagellate and ciliate protists and metazooplankton*

The range and mean values for filtration and ingestion rates for each taxon of flagellate and ciliate protists and metazooplankton are shown in Tables I and II. Among the heterotrophic nanoflagellates, only cells of the *Monas* type ingested

**Table II.** Filtration and ingestion rate of the various taxa of metazooplankton

Metazooplankton	Bead uptake	Clearance ( $\mu\text{l ind.}^{-1} \text{h}^{-1}$ )			Ingestion ( $\times 10^3$ bacteria $\text{ind.}^{-1} \text{h}^{-1}$ )		
		Min.	Max.	Mean	Min.	Max.	Mean
Crustaceans							
Cladocerans							
<i>Bosmina longirostris</i>	Y	1.0	17.0	9.0	3.3	138.0	28.0
<i>Ceriodaphnia quadrangula</i>	Y	7.0	144.0	76.0	2.9	456.2	175.8
<i>Daphnia longispina</i>	Y	10.0	290.0	90.0	10.6	704.8	229.1
Copepods							
<i>Cyclops vicinus</i>	N						
<i>Acanthocyclops robustus</i>	N						
<i>Eudiaptomus gracilis</i>	N						
Rotifers							
<i>Asplanchna priodonta</i>	N						
<i>Conochilus unicornis</i>	Y	0.8	2.3	2.2	2.2	14.2	6.5
<i>Filinia terminalis</i>	N						
<i>Hexarthra mira</i>	Y	11.2	17.2	14.2	30.2	46.1	38.1
<i>Kellicottia longispina</i>	N						
<i>Keratella cochlearis</i>	N						
<i>Keratella quadrata</i>	N						
<i>Polyarthra</i> sp.	N						
<i>Synchaeta</i> sp.	N						

bacteria-sized particles, whereas, among pigmented flagellates, only *Cryptomonas ovata* and *Dinobryon cylindricum* ingested bacteria-sized particles (Table I). The ciliates consuming bacteria-sized particles were Oligotrichida and Peritrichida. Cladocera all consumed bacteria-sized particles, whereas *Polyarthra* sp., which accounted for most of the rotifer biomass, and Copepoda never ingested microspheres.

Estimated total consumption of bacteria in the epilimnion varied between 0.08 and  $10.27 \times 10^6$  bacteria  $l^{-1} h^{-1}$  (Figure 5A). Predation of bacteria-sized particles by heterotrophic and mixotrophic flagellates averaged  $0.64 \times 10^6$  bacteria  $l^{-1} h^{-1}$ . The highest grazing rates by protists were recorded on 13 and 20 August ( $2.54$  and  $2.47 \times 10^6$  bacteria  $l^{-1} h^{-1}$ ), when they were caused by a heterotrophic flagellate (*Monas*-like cells) and then by a pigmented flagellate (*Cryptomonas ovata*)

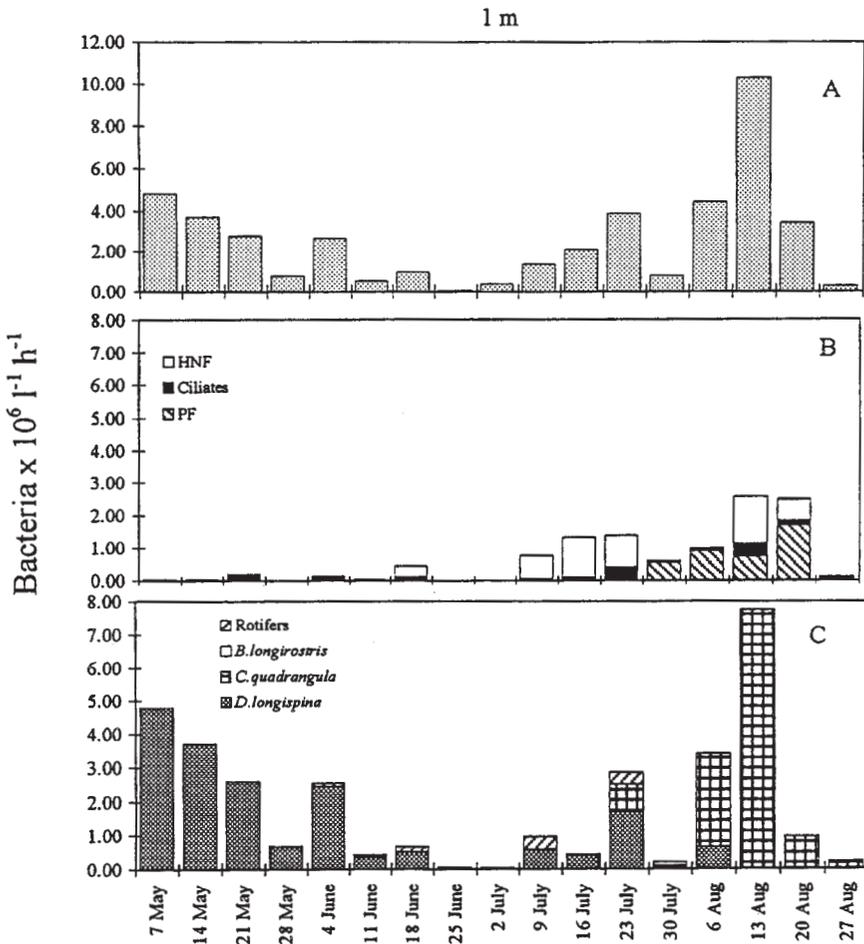


Fig. 5. Temporal changes in total bacterial consumption (A), by phagotrophic protists (B) and by metazoan zooplankton (C) at 1 m depths.

(Figure 5B). There was no consumption of bacteria-sized particles by protists on 25 June and 2 July. The mean consumption of bacteria-sized particles by ciliates was  $0.10 \times 10^6$  bacteria  $l^{-1} h^{-1}$ , accounting for 15% of the total grazing by protists for all the measurements combined. Two oligotrich species, *Pelagohalteria viridis* and *Halteria* sp., were responsible for 95% of the ingestion of microspheres. *Vorticella* were only encountered on a single date, on 4 June, when their potential predation of bacteria was  $0.022 \times 10^6$  bacteria  $l^{-1} h^{-1}$ . The impact of scuticociliates was negligible because of their low density throughout the study.

The average predation by metazooplankton was  $1.89 \times 10^6$  bacteria  $l^{-1} h^{-1}$ . In spring, *D.longispina* was the main bacterial consumer, whereas *Ceriodaphnia quadrangula* became the main consumer in summer. These two species accounted for 57 and 39% of bacterial predation by metazoans, respectively, whereas *B.longirostris*, which was only counted at 1 m, only accounted for 2% of total metazoan activity. *Hexarthra mira* and *Conochilus unicornis*, with mean filtration rates of 14.2 and 2.2  $\mu l$  ind. $^{-1} h^{-1}$ , respectively, were the two rotifer species that ingested microspheres in June and July.

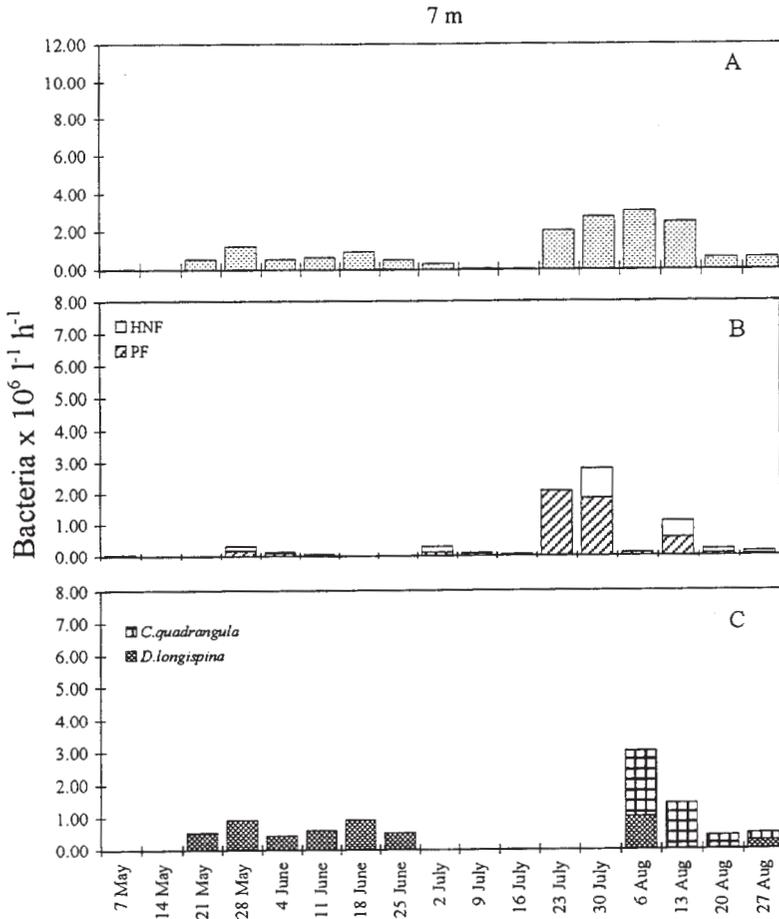
Bacterial consumption in the metalimnion fluctuated between 0 and  $3.07 \times 10^6$  bacteria  $l^{-1} h^{-1}$  (mean  $0.97 \times 10^6$  bacteria  $l^{-1} h^{-1}$ ) (Figure 6A). The mean grazing rate of ciliates and total flagellates decreased slightly with depth and was only  $0.43 \times 10^6$  bacteria  $l^{-1} h^{-1}$  at 7 m (Figure 6B). The numbers of ciliates counted in each replicate throughout the study period, and of crustaceans and rotifers in early May and in July, were too low in the metalimnion to calculate the grazing impact.

From the start of the study until 16 July, bacterial consumption by flagellates was  $<0.31 \times 10^6$  bacteria  $l^{-1} h^{-1}$ . Pigmented phagotrophic flagellates (*Dinobryon cylindricum* and *Cryptomonas ovata*), which only accounted for 12% of consumption at 1 m, had a higher contribution at 7 m (32%). This was due to their high consumption rates on 23 and 30 July ( $2.06$  and  $1.82 \times 10^6$   $l^{-1} h^{-1}$ ) (Figure 6B). The grazing impact of protists was highest on 30 July ( $2.78 \times 10^6$  bacteria  $l^{-1} h^{-1}$ ) and was mainly due to the pigmented flagellate *Dinobryon cylindricum* whose consumption accounted for 75% of grazing on bacteria-sized particles by protists at this depth. As in the epilimnion, most of the bacterial consumption by metazooplankton in the spring was by *D.longispina* and mainly by *Ceriodaphnia quadrangula* in summer (Figure 6C).

Over the entire study, cladocerans accounted on average for 72% of predation at 1 m and 56% at 7 m, and were the main consumers of bacteria-sized particles, especially in May, June and August. HNF accounted for 12% of bacterial consumption at 1 m and 13% at 7 m, and therefore only had a limited predation mainly due to a *Monas*-type cell. Ciliates and rotifers accounted for 4 and 2% of total predation, and were the organisms with the lowest grazing impact.

## Discussion

Various methods have been used to quantify bacterial consumption by individual planktonic organisms in the field. The use of fluorescent microspheres is one of the standard methods used for estimating the ingestion rate of protists (e.g. MacManus and F urhman, 1988), rotifers (e.g. Ooms-Wilms *et al.*, 1995; Ooms-Wilms, 1997)



**Fig. 6.** Temporal changes in total bacterial consumption (A), by phagotrophic protists (B) and by metazoan zooplankton (C) at 7 m depths.

and cladocerans (Børshiem and Andersen, 1987; Wiedner and Vareschi, 1995). Protozoans and copepods are organisms that are capable of selecting their food (Monger and Landry, 1992; Gonzales *et al.*, 1993), in contrast to rotifers (Arndt, 1993) and cladocerans (DeMott, 1986; Lampert, 1987) which are thought to be relatively unselective in terms of the bacterial particle size. Various studies have shown that using fluorescent microspheres underestimates predation by some ciliates and flagellates that preferentially ingest bacteria labelled with fluochrome (FLB) rather than the microspheres (Sherr *et al.*, 1987; Simek and Straskrabova, 1992), whereas other studies have shown that there are no significant differences in predation between the different types of particles (e.g. Sanders *et al.*, 1989; Jones and Rees, 1994).

For metazoans, the method used has an influence on the calculation of the ingestion rates (Ooms-Wilms *et al.*, 1995). For example, the rapidly digested FLB

tend to underestimate grazing by Cladocera. Furthermore, although some authors have determined bacterial consumption in rotifers using FLB (Ooms-Wilms, 1991; Turner and Tester, 1992), Boon and Shield (1990) demonstrated that there are methodological problems. Copepods are capable of selecting their prey (DeMott, 1988) and could, therefore, be sensitive to the method used. However, to our knowledge, there are no experimental data to support such a hypothesis. In fact, the work on food selection by these organisms was conducted using particles of a size between 2 and 20  $\mu\text{m}$  (DeMott, 1988; Kerfoot and Kirk, 1991), which is much greater than the length of bacteria occurring in lake ecosystems. Fluorescent microspheres also have the advantage of being easy to see in the alimentary tract of the different metazoans, in contrast to labelled bacteria (FLB or minicells).

The use of microspheres would, therefore, appear to be the best compromise for estimating simultaneous consumption of bacteria by both protists and metazooplankton. The predation on bacteria-sized particles measured in this study is obviously only representative of the two zones that were studied, and especially in the case of metazoans that can migrate over long distances through the water column.

Metazoans appeared to be the main consumers of bacteria in the epilimnion and metalimnion in this study. The measurements of predation at 1 and 7 m, and the highly significant negative correlation ( $P < 0.001$ ) between the abundance of the cladoceran *D.longispina* in the water column and the mean density of bacteria in the zones studied, demonstrate the predator-prey relationships that exist between these two communities. The consumption, which at maximum reached 7% of the standing stock of bacteria, was similar to that recorded by Vaqué and Pace (1992) in a dystrophic lake. The potential bacterial consumption is low and cannot at first sight satisfactorily explain the control of bacterial production, even if we accept a very conservative estimate of one doubling of total bacteria every 5 days. However, in this ecosystem, few of the bacteria are active (0.2–3.2%) (Jugnia *et al.*, 1998) and only these are capable of dividing (Roszak and Calwell, 1987). It should also be noted that the experiments were conducted during daylight hours, which certainly led to an underestimate of the impact of organisms on bacteria, since the metazooplankton populations undertake vertical migrations and are usually more concentrated at the surface during the night (Lampert, 1989). Our results, in general, confirm studies conducted in the natural environment, which emphasize the importance of the bacterial grazing activity of Cladocera (Jürgens, 1994). *Daphnia longispina* and *Ceriodaphnia quadrangula*, which were of small average size in this ecosystem ( $720 \pm 220$  and  $381 \pm 78 \mu\text{m}$ , respectively), have a filter mesh size of  $<0.5 \mu\text{m}$  (Geller and Müller, 1981; Brendelberger and Geller, 1985) which makes them efficient at retaining small-sized particles (Jürgens, 1994; Toth and Kato, 1997) and suggests that they are effective consumers of bacteria (Peterson *et al.*, 1978; Geller and Müller, 1981; Pace *et al.*, 1983; Güde, 1988; Jürgens, 1994). These results are confirmed by grazing measurements which showed that *D.longispina* was the species with the highest bacterial consumption activity. The calculated filtration rate values (Table II) for this species are close to those measured by Børshem and Andersen (1987) using

microspheres and by Kankaala (1988) using bacteria labelled with tritiated thymidine. *Ceriodaphnia quadrangula* was the main bacterial consumer in August. The filtration rate measured for this species ( $7\text{--}144 \mu\text{l ind.}^{-1} \text{h}^{-1}$ ; mean  $76 \mu\text{l ind.}^{-1} \text{h}^{-1}$ ) was higher than that recorded by Bern (1987) with radioactively labelled bacteria, but the incubation temperature ( $14.0^\circ\text{C}$ ) was lower than in this study ( $22.5 \pm 2.1^\circ\text{C}$ ). The filtration measured for *B.longirostris* (between  $1$  and  $17 \mu\text{l ind.}^{-1} \text{h}^{-1}$ ; mean  $9.0 \mu\text{l ind.}^{-1} \text{h}^{-1}$ ) was close to that found by Toth and Kato (1997) ( $13.4 \pm 4.9 \mu\text{l ind.}^{-1} \text{h}^{-1}$ ), and was the lowest of all the species of Cladocera recorded in this study. This result confirms the many studies that have shown that Bosminidae usually have a low bacterial grazing activity (Bogdan and Gilbert, 1982, 1984; DeMott and Kerfoot, 1982; Schoenberg and Maccubin, 1985; Ross and Munawar, 1987; Gulati *et al.*, 1991).

Few studies have been conducted on the consumption of bacteria by rotifers under natural conditions (Bogdan and Gilbert, 1982; Sanders *et al.*, 1989; Ooms-Wilms, 1997). Heterotrophic bacteria could represent a food source for these organisms which are relatively non-specific micro filter feeders over a size range of between  $0.5$  and  $2.0 \mu\text{m}$  (Pourriot, 1977; Rothhaupt, 1990), but rotifers only accounted for 2% of total predation on bacteria. This is in agreement with studies which have shown that their contribution to total bacterial consumption is generally  $<10\%$  (Sanders *et al.*, 1989; Pace *et al.*, 1990; Pernie *et al.*, 1990). Some of the species present in the lake (e.g. *Asplanchna priodonta*, *Polyarthra* sp. and *Synchaeta* sp.) never ingested the fluorescent particles and probably do not consume bacteria (Bogdan *et al.*, 1980; Boon and Shiel, 1990; Ooms-Wilms *et al.*, 1995). Other species, such as *Hexarthra mira* and *Conochilus* sp., had a similar bacterial consumption activity to that measured by Sanders *et al.* (1989) and Ooms-Wilms *et al.* (1995). In agreement with the studies of Brett *et al.* (1994) and Sanders *et al.* (1989), this study also showed no evidence of bacterial consumption by copepods, in contrast to that of Roff *et al.* (1995) in the marine environment.

Among the phagotrophic protists, pigmented flagellates were the most important consumers, followed by HNF and ciliates. Pigmented phagotrophic flagellates accounted for 32% of grazing on bacteria at 7 m. Among these organisms, *Cryptomonas ovata* and *Dinobryon cylindricum* showed a similar activity to that recorded by Tranvik *et al.* (1989) and Bird and Kalf (1986). The estimated filtration rate for *D.cylindricum* was similar to that recorded by Carrias *et al.* (1996), whereas that obtained for *C.ovata* (mean  $0.7 \times 10^{-3} \mu\text{l ind.}^{-1} \text{h}^{-1}$ ) was slightly higher than the value determined by Tranvik *et al.* (1989) using FLB in a humic mesotrophic lake ( $0.27\text{--}0.55 \times 10^{-3} \mu\text{l ind.}^{-1} \text{h}^{-1}$ ).

The low density of ciliates encountered in the epilimnion of the Sep reservoir was similar to that reported from oligotrophic environments (e.g. Laybourn-Parry, 1994). The community was mainly dominated by a mixotrophic oligotrich *Pelagohalteria viridis*, which is frequently found in lake plankton (Carrias *et al.*, 1996). On average, ciliates accounted for 15% of total bacterial consumption by protists (4% of total grazing). This result is similar to that reported for the eutrophic Lake Oglethorpe (Sanders *et al.*, 1989) and for the oligomesotrophic Lake Pavin (Carrias *et al.*, 1996). *Pelagohalteria viridis*, which contains symbiotic

algae (Foissner *et al.*, 1988), paradoxically had a higher mean ingestion rate than the heterotrophic oligotrich *Halteria* sp.

The structure of the HNF community was similar to that encountered in lakes of oligomesotrophic to mesotrophic trophic status (Salbrechter and Arndt, 1994; Mathes and Arndt, 1995; Carrias *et al.*, 1998). Among these microorganisms, only the *Monas*-like cells ingested microspheres, with a filtration rate ( $0.4\text{--}3 \times 10^{-3} \mu\text{l ind.}^{-1} \text{h}^{-1}$ ) similar to that obtained by Sanders *et al.* (1989) in the eutrophic Lake Oglethorpe and close to that of Carrias *et al.* (1996) in the oligomesotrophic Lake Pavin. The ingestion rates that were measured for these HNF ( $0\text{--}14$  bacteria  $\text{ind.}^{-1} \text{h}^{-1}$ ) were similar to those obtained by Bloem *et al.* (1989) in a eutrophic lake ( $2\text{--}17$  bacteria  $\text{ind.}^{-1} \text{h}^{-1}$ ).

This study showed that metazoans, and particularly Cladocera, were partly responsible for controlling the abundance of bacteria in a newly flooded lake. However, predation by *Ceriodaphnia quadrangula* on bacteria in August paradoxically coincided with an increase in bacterial abundance. It seems likely that bacterial production, which is strongly temperature dependent (White *et al.*, 1991), exceeded predation at this time. At times of low chlorophyll *a* concentrations, bacteria could be an important food supplement for some metazoans (Pace *et al.*, 1983), but they are poorly assimilated relative to algae (Hessen *et al.*, 1989) and they lack the long-chain unsaturated fatty acids that may be crucial for zooplankton nutrition (Muller-Navarra, 1995).

Furthermore, as shown by Thouvenot *et al.* (1999) in field experiments in the same ecosystem, metazoans could exert a heavy predation on some protozoans, which would result in a reduction in the overall predation on bacteria by these organisms. This predation by metazoans could be exerted on ciliates of the size measured in this study, which was frequently  $<30 \mu\text{m}$ , and which correspond with metazooplanktonic nutritional requirements (Wiackowski *et al.*, 1994; Rabette *et al.*, 1998). The dominance of *Pelagohalteria viridis* could then be attributed to its ability to move by leaps (Tamar, 1979), enabling it to escape from predators and be one of the few ciliates that is capable of existing in the presence of Cladocera (Jürgens *et al.*, 1994). The inverse relationship ( $P < 0.01$ ) between the biomass of copepods and rotifers in the water column and of PF in the euphotic zone suggests that there is a predator–prey relationship between these metazoans and pigmented flagellate protists.

Several facts also tend to indicate that HNF were subjected to heavy predation. Firstly, the density of heterotrophic flagellate protists ( $0.5\text{--}3.6 \times 10^3$  cells  $\text{ml}^{-1}$ ) was slightly lower than the values generally reported for oligotrophic or mesotrophic environments ( $0.2\text{--}5.6 \times 10^3$  cells  $\text{ml}^{-1}$ ) (Pick and Hamilton, 1994; Tzaras and Pick, 1994) and, secondly, there was a negative correlation ( $P < 0.01$ ) between the abundance of the cladoceran *D.longispina* in the water column and these flagellates. The ratios of bacterial abundance to HNF abundance (7026:1 at 1 m and 3835:1 at 7 m) were greater than those recorded by Amblard *et al.* (1995) (2590:1) and Simek *et al.* (1995) (1000:1), and could be interpreted as the result of predation. The low bacterial consumption by HNF could also be explained by the structure of this community. *Kathablepharis ovalis* is a species that in fact prefers larger sized particles such as nanophytoplankton (Carrias *et al.*, 1996). In

addition, the chryomonadine (2–5 µm), which was difficult to identify because of its small size (Finlay *et al.*, 1988) and which dominated the HNF community, never consumed bacteria and we were never able to detect autotrophic picoplankton in its cells. Osmotrophic activity of HNF (Sherr and Sherr, 1987; Marchant and Scott, 1993) could have been favoured by the high DOM concentrations that were greater than those usually found in lakes of this trophic level (Striquer-Soares and Chevolut, 1996). These findings could explain why protists, and particularly HNF, which are generally considered to be the main microorganisms capable of transferring picoplankton production towards higher trophic levels (Güde, 1986; Cole *et al.*, 1988; Sherr and Sherr, 1988; Sanders *et al.*, 1989; Carrias *et al.*, 1996), only played a limited role in regulating the bacterial communities in this type of ecosystem.

The low abundance of phagotrophic protists in this newly flooded reservoir and the low proportion of HNF capable of consuming bacteria have therefore resulted in the metazoan zooplankton becoming the main consumers of bacteria. These results, combined with those of Thouvenot *et al.* (1999), demonstrate that these organisms have a high predation impact on bacteria, HNF and ciliates, thus confirming the hypothesis of Paterson *et al.* (1997), who suggested that some components of the microbial loop could be an important source of food for metazoans in a newly flooded reservoir. Nevertheless, the proportion of carbon coming from these organisms compared to that from strict autotrophs remains to be determined.

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