Nutrient cycling in a microflagellate food chain: I. Nitrogen dynamics*

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ABSTRACT: In a series of grazing experiments we found that the phagotrophic microflagellate $Paraphysomonas\ imperforata$ (Lucas) (7 to 12 μm dia.) fed equally well on the diatom $Phaeodactylum\ tricornutum$ and bacteria under batch conditions. Growth rates of the microflagellate were considerably higher than those of the phytoplankton prey. Regeneration of NH_4^+ was negligible in control cultures of phytoplankton alone, bacteria alone, or phytoplankton and bacteria. However, when the microflagellate was grazing on either phytoplankton or bacteria there was considerable regeneration of NH_4^+ This result was a clear indication that only the microflagellate was responsible for the excretion. Rates of NH_4^+ excretion were highest during exponential growth of the microflagellate and slackened off considerably with onset of the stationary phase when growth of the prey was not nutrient-limited. Regeneration efficiencies, however, were lowest during exponential growth (15 to 30 %), but increased to 50 % if calculated on the basis of the entire experiment which included the stationary phase. Regeneration efficiencies during exponential growth decreased to 8 % when the prey was grown under nitrogen-limitation. Nitrogen turnover rates per body weight are generally much higher in microflagellates than in macrozooplankton, but can be greatly influenced by many environmental factors including prey nutritional state.

INTRODUCTION

The pathways by which inorganic nutrients are recycled in surface waters of the open ocean after the first step of phytoplankton assimilation are poorly understood. The classical view has been that nutrients are recycled in the euphotic zone by bacterial mineralization of both dead phytoplankton and phytoplankton excretion products and by direct grazing on phytoplankton by macrozooplankton. In recent years there has been considerable evidence that frequently the bulk of nutrient regeneration occurs among plankton less than 100 μ m in size (summarized in Harrison 1980) and in some cases by microbes smaller than 10 μ m (Glibert 1982). Thus there is now some doubt that macrozooplankton grazing is an important process of nutrient regeneration in pelagic waters (Jackson 1980).

The role of bacteria in the nutrient regeneration

process is equally confusing. Bacteria are efficient scavengers of inorganic nutrients; they have been shown, for example, to compete successfully with phytoplankton for inorganic phosphorus in numerous studies (Rhee 1972, Faust & Correll 1976, Harrison et al. 1977). They also are very efficient in converting dissolved organic material (and presumably nutrients) to biomass (e.g. net growth yields between 50 and 70 % are commonly observed) (Calow 1977). Hence, the contemporary notion is that direct excretion of inorganic nutrients during active growth of bacteria is not a major mode of remineralization (Williams 1981, Joint & Morris 1982, Hagstrom & Larsson 1984). Additionally, bacterial hydrolysis of organic material from dead phytoplankton is probably minimal in the euphotic zone of pelagic waters for 2 reasons. First, the rate constants for such decomposition are known to be relatively low: 0.05 to 0.1 d⁻¹ for the most labile fractions of phytoplankton biomass (Harrison 1980). Second, there is no evidence for spontaneous death of phytoplankton populations in the open sea - the remains of intact phytoplankton cells are not often found in oceanic surface waters (Pomeroy 1984).

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The resulting scenario is that bacteria are not efficient remineralizers of nutrients assimilated by growing phytoplankton and may even compete with phytoplankton for nutrients in the open ocean. This leaves the perplexing but unanswered question of how inorganic nutrients assimilated by phytoplankton and converted to particulate organic material are then recycled back into the dissolved phase. The importance of this question is dramatized by the possibility that 80 to 90+% of nutrients in the euphotic zone of pelagic waters are recycled (Eppley & Peterson 1979).

It has now become abundantly clear that phagotrophic microflagellates are the main grazers of bacteria in the sea (Haas & Webb 1979, Fenchel 1982a, Azam et al. 1983, Davis & Sieburth 1984, Sieburth 1984). Although, their role in recycling nutrients assimilated by bacteria has been the subject of some debate (Johannes 1965, Barsdate et al. 1974, Fenchel & Harrison 1976), the importance of these voracious predators in this process now seems firmly established (Azam et al. 1983, Sherr et al. 1983).

Recently, one of us raised the possibility that phagotrophic microflagellates, being highly opportunistic, could play an important role in cropping both bacteria and phototrophs and in regenerating nutrients from all grazed microbes (Goldman 1984a,b). To test this hypothesis we performed a series of growth and nutrient regeneration studies under NH₄+, PO₄-3, and non-nutrient-limited conditions on a heterotrophic microflagellate that we found capable of grazing on both bacteria and a variety of phytoplankton species. General features of our experimental results are reported elsewhere (Goldman & Caron 1985). In this study we examine the details of nitrogen cycling by the microflagellate under the different nutritional conditions. In subsequent papers we report on population dynamics and carbon cycling (Caron et al. 1985) and on phosphorus cycling (Andersen et al. unpubl.).

METHODS

Microbial species. A heterotrophic microflagellate, identified as Paraphysomonas imperforata (Lucas) (7 to 12 μm in diameter) (see Goldman & Caron 1985), was isolated from a tidal channel connecting Vineyard Sound, Massachusetts with an estuarine pond near the Woods Hole Oceanographic Institution (WHOI). A clonal culture of the microflagellate was started from the second stage of a 2-stage continuous culture system. The first stage was a culture of the marine diatom Phaeodactylum tricornutum (clone TFX-1) grown on synthetic seawater medium (Goldman & McCarthy 1978). To start an enrichment culture in the second stage a few ml of a centrifuged concentrate of 35 μm

filtered seawater from the tidal channel was added and a diverse population of protozoa developed. The microflagellate, chosen because it grazed directly on the diatom, was then isolated and made bacteria-free by micropipetting individual cells into an axenic culture of *P. tricornutum* containing antibiotics. The microflagellate culture was maintained by making approximate weekly transfers to fresh cultures of *P. tricornutum*. Routine tests were made for bacterial contamination.

The culture of *Phaeodactylum tricornutum*, used both to isolate the microflagellate and for the grazing and nutrient regeneration experiments, came from the culture collection of R. R. L. Guillard. Bacterial populations used in the grazing and regeneration experiments were either a mixed bacterial population grown on 0.8 μm filtered (Nuclepore) seawater from Vineyard Sound enriched with 0.01% yeast extract, or a pure culture of *Pseudomonas halodurans* obtained from H. W. Jannasch at WHOI and maintained on nutrient seawater agar.

Experimental protocols. Three sets of experiments were performed, 1 at 24°C (Experiment A) and the other 2 at 20 °C (Experiments B & C) (Table 1). Experiment A involved all possible combinations of the alga, bacteria (mixed population), and microflagellate plus control experiments with the alga and bacteria alone. It was intended to make the prey cultures N-limited in Experiment A by adjusting the N:P ratio of synthetic seawater media (Goldman & McCarthy 1978) to 10 (100 μ g-at l^{-1} NH_4^+ and 10 μ g-at l^{-1} PO_4^{-3}). However, as will be discussed in the 'Results' section, the grazing phase of the experiments was commenced while the phytoplankton or bacterial cultures were still in the exponential phases of growth; thus we did not attain clear-cut N-limited conditions. Experiment B was similar to Experiment A, but with the prey grown on a Plimited growth medium with an N:P ratio of 60 $(150 \,\mu g\text{-at l}^{-1} \text{ of } NH_4^+ \text{ and } 2.5 \,\mu g\text{-at l}^{-1} \text{ of } PO_4^{-3}).$ Phosphorus or nitrogen limitation was ensured in the prey cultures (Phaeodactylum tricornutum and Pseudomonas halodurans) of Experiment C by using the appropriate N: P ratios in the medium and allowing the cultures to enter the stationary phase before adding the microflagellate (Table 1).

All experiments were performed with 7 l of media in 8 l glass carboys, except for the bacterial control in Experiment A which was performed with 900 ml of medium in a 1 l flask. The bacterial medium was supplemented with 36 mg l^{-1} glucose. Sterile growth media were inoculated with either *Phaeodactylum tricornutum* or bacteria from stock cultures grown to late exponential phase in order to minimize carry-over of nutrients. The cultures were mixed with a magnetic stirring bar and with bubbled air (filter-sterilized). The

Experi- ment	Tempera- ture (°C)	Microbes involved	N:Pratio in medium (by atoms)	Prey cell quota (pg N cell ⁻¹)	Microflagellat μ ± SE (d ⁻¹)	
A 24		Diatom Bacteria* Bacteria** Diatom + bacteria*	10:1	2.0 0.03 0.02 2.5	- - - -	
		Diatom + microflagellate Diatom + bacteria* + microflagellate Bacteria* + microflagellate		2.2 2.0 0.03	3.48 ± 0.189 3.41 ± 0.225 3.50 ± 0.221	
В	20	Diatom + bacteria** Diatom + microflagellate Diatom + bacteria** + microflagellate	60 : 1	1.7 1.5 1.3	2.55 ± 0.199 2.52 ± 0.021	
С	20	Bacteria · · + microflagellate Diatom + microflagellate Diatom + microflagellate	50 : 1 60 : 1 10 : 1	0.05 1.0 0.8	2.47 ± 0.101 2.10 ± 0.075 2.31 ± 0.077	

Table 1. Summary of experimental conditions for grazing in the dark by the phagotrophic microflagellate *Paraphysomonas* imperforata on the marine diatom *Phaeodactylum tricornutum* and on bacteria

••• Pseudomonas halodurans

experiments were performed in a walk-in incubator maintained on a 12 h light-12 h dark cycle. Air bubbling was stopped when the cultures reached designated points in the growth curves and inocula of the microflagellate and/or bacteria were added. Cultures were then darkened and time series measurements of a variety of biological and chemical parameters were made in approximate 6 h intervals over a 6 to 7 d period. Although a constant dark period is unrealistic, we wanted to avoid nutrient uptake and growth of *P. tricornutum* during grazing. Such a condition would have seriously complicated our attempts to calculate grazing and nutrient regeneration rates of the microflagellate.

Sampling was started within 8 h of placing the cultures in the dark. The bacterial inoculum added to the algal cultures came from the 7 l bacterial culture and the microflagellate inoculum came from a stock culture grown on *Phaeodactylum tricornutum*. Aseptic sampling was done with a gravity fed siphon clamped at the end. Before each sampling a volume of liquid exceeding the volume of the siphon tube was wasted; approximately 400 ml of culture were then collected and used for all analyses required in this study and in the companion studies of Caron et al. (1985) and Andersen et al. (unpubl.).

Analytical methods. Particulate nitrogen (PN) was measured with a Perkin Elmer 240 Elemental Analyzer on 25 ml samples retained on pre-combusted Whatman GF/F glass-fiber filters. Filtration pressure differentials were 25 mm Hg. Sufficient filtrate was collected to measure $\mathrm{NH_4}^+$ (McCarthy & Kamykowski 1972) and

urea (McCarthy 1970) immediately and to freeze a portion for later analyses of NO₃⁻ (Wood et al. 1967). Cells of *Phaeodactylum tricornutum* were counted in Experiment A with a Spencer Bright-line hemacytometer on samples preserved in Lugol's solution. Epifluorescence microscopy with acridine orange staining (Watson et al. 1977, Davis & Sieburth 1982) was used to count bacteria and *Paraphysomonas imperforata* in Experiment A and all microbes in Experiments B and C. Samples were preserved in 1% glutaraldehyde in filtered seawater.

Data analyses. The microflagellate growth and grazing curves were divided into 5 convenient intervals to facilitate analyses and interpretation of the data. Interval I defined the portion of each growth curve where the initial decrease in prey cell number was so small that it was impossible to calculate a nitrogen ingestion or excretion rate. Interval II described the exponential phase of microflagellate growth which was concomitant with pronounced decreases in prey cell number and increases in NH₄⁺ excretion. Interval III was the transition phase between exponential growth and the stationary phase where both the rates of grazing and nitrogen excretion decreased markedly. Finally, Intervals IV and V defined the early and late stationary phases, respectively. Each interval was designated by visual inspection of the experimental curves.

Specific growth rates μ (d⁻¹) of the heterotrophic microflagellate were determined from linear regression analyses of the plots of the natural log of cell count *versus* time for the exponential phase of growth (Intervals I and II). Both the average nitrogen ingestion and

excretion rates per microflagellate cell for each interval were determined by the expressions:

 $I_{N} = Q_{N_{D}}(P_{1} - P_{2}) \cdot (\overline{F} \cdot \Delta t)^{-1}$ (1)

and

$$E_{N} = (\Sigma N_{2} - \Sigma N_{1}) \cdot (\overline{F} \cdot \Delta t)^{-1}$$
 (2)

where I_N and E_N = nitrogen ingestion and excretion rates, respectively (pg N cell⁻¹ d⁻¹); $Q_{Np} = nitrogen$ cell quota for Phaeodactylum tricornutum or bacteria at the time the microflagellate was introduced into the culture (pg N cell⁻¹); P_1 and P_2 = prey cell concentration (cells ml^{-1}) at the start and end, respectively, of each interval Δt (d); ΣN_2 and ΣN_1 = sum of excreted NH₄⁺ and urea at the end and beginning, respectively, of each interval (pg N l^{-1}); $\overline{F} = \ln$ average concentration of microflagellates (cells ml-1) during each interval. The equation $(F_2-F_1) \cdot (\ln F_2 - \ln F_1)^{-1}$ of Heinbokel (1978) was used to determine \overline{F} , where F_2 and F_1 = microflagellate cell number (cells ml^{-1}) at the end and start respectively of each interval. Nitrogen regeneration efficiencies (R_N in %) were calculated either as $(E_N \cdot I_N^{-1}) \cdot 100$ for each interval or the sum of Intervals I to III or I to IV by the expression:

$$R_{N} = [(\Sigma N_{t} - \Sigma N_{O}) \cdot P N_{O}^{-1}] \cdot 100$$
 (3)

where ΣN_O and PN_O = concentrations of ΣN and particulate nitrogen respectively at the start of the dark phase; ΣN_t = concentration of ΣN at the end of the designated summed interval.

Estimates of the microflagellate nitrogen cell quota at each sampling were made from the following expression:

$$Q_{Nf} = (PN_{t} - Q_{Np}P_{t}) \cdot F_{t}^{-1}$$
 (4)

where Q_{N_f} and PN_t = flagellate nitrogen cell quota (pg

N cell⁻¹) and particulate nitrogen concentration respectively at each sampling; P_t and F_t = cell concentrations of the prey organisms and the microflagel-late respectively at each sampling.

RESULTS

Prey physiological state before grazing

We found no evidence of N or P limitation in any of the prey cultures, except when purposely kept in the growth phase for an extended period (Experiment C). At the end of the growth phase of Experiment A not only was there a slight residual of NH₄⁺ remaining in all cultures, but the particulate C:N ratios (by atoms) were about 5 to 6 in the phytoplankton cultures and about 4 in the bacterial culture (data not shown). For the most part, any remaining NH₄+ in these experiments was rapidly consumed at the beginning of the dark phase with a concomitant rapid but small increase in phytoplankton or bacterial cell numbers (Fig. 1 to 3). When the phytoplankton and bacterial cultures were grown on medium with an N:P ratio of 50 to 60 (Experiments B & C) there was a very large residual of NH₄⁺ at the end of the growth phase (Fig. 4 D & 5 C) and the particulate N: P ratios were < 20 (data not shown). In contrast, the particulate N:P ratio rose to 23 to 30 in the P-limited cultures of Experiment C. Also, particulate alkaline phosphatase in these cultures was significantly higher than in any of the others (Anderson et al. unpubl.). The particulate C:N ratio was about 9 in the N-limited culture of Experiment C as opposed to ratios of 5 to 6 in the non-nutrientlimited cultures (data not shown).

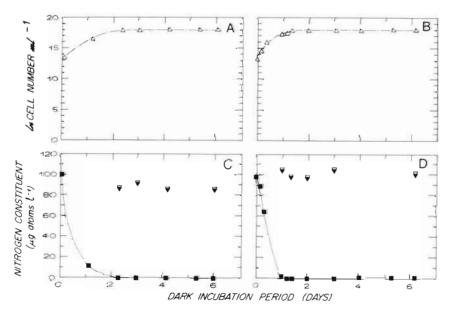


Fig. 1 Control bacterial experiments. Cell number (Δ) of (A) Pseudomonas halodurans, (B) mixed bacterial assemblage, grown under non-nitrogen- and non-phosphorus-limiting conditions in the dark. Corresponding plots of nitrogen mass balance are shown in (C) P. halodurans, (D) mixed bacterial population. (\blacksquare) Σ N (urea plus NH₄+); (\blacktriangledown) particulate N; (\triangledown) total N (PN + urea + NH₄+). Time zero indicates when initial bacterial inoculum was added to medium

Fig. 2. Control phytoplankton experiments. Cell number (\square) of *Phaeodacty-lum tricornutum* first grown under non-nutrient-limiting conditions (Experiment A) and then placed in the dark in (A) absence or (B) presence of bacteria (\triangle). Corresponding plots of nitrogen mass balance are shown in (C) without bacteria, (D) with bacteria. (\blacksquare) ΣN ; (\triangledown) particulate N; (\triangledown) total N. Time zero indicates when cultures had grown to mature populations. Same for Fig. 3 to 5

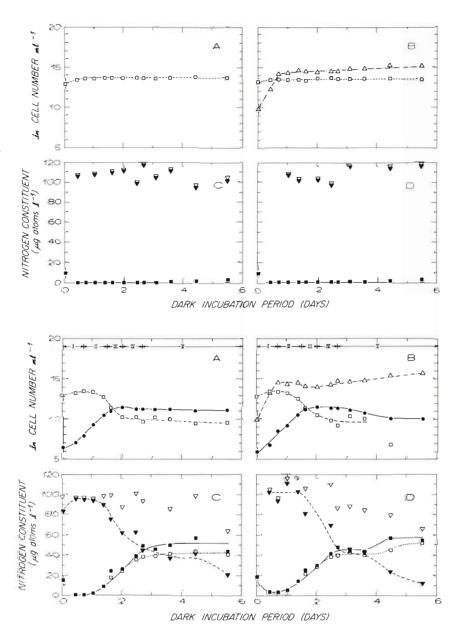


Fig. 3. Grazing of Phaeodactylum tricornutum (\square) grown under non-nutrient-limiting conditions (Experiment A) by the phagotrophic microflagellate Paraphysomonas imperforata (\bullet) in (A) absence or (B) presence of bacteria (\triangle). Intervals I to V define intervals of grazing as described in text. Corresponding plots of nitrogen mass balance are shown in (C) without bacteria, (D) with bacteria. (\blacksquare) ΣN ; (\square) NH_4^+ ; (\blacktriangledown) particulate N; (\triangledown) total N

Phytoplankton and bacterial control experiments

After the initial changes described above, we found no further changes over time in cell number or particulate nitrogen in the control cultures of Experiment A consisting of a mixed bacterial population alone (Fig. 1B,D), *Pseudomonas halodurans* alone (Fig. 1A,C), or *Phaeodactylum tricornutum* either alone (Fig. 2A, C) or in combination with a mixed bacterial population (Fig. 2B, D). There was no evidence of NH_4^+ or urea excretion in the bacterial control cultures during the course of the incubation (Fig. 1C, D), but there was a slight ($< 3 \,\mu g$ -at l^{-1}) build up of NH_4^+ in the phytoplankton control cultures by Day 5.5 (Fig. 2C, D). We obtained similar results with the control incubations of

Experiment B involving P. tricornutum (medium N:P ratio = 60) plus bacteria, although in these experiments there were large residuals of NH_4^+ remaining at the time the dark phase was started. Because all the results of Experiment B were virtually identical to those of Experiment A, we have, with exception of growth rate data (Table 1), excluded them from presentation.

Microflagellate grazing experiments

There was a fairly consistent pattern of grazing by Paraphysomonas imperforata regardless of whether the prey was Phaeodactylum tricornutum only

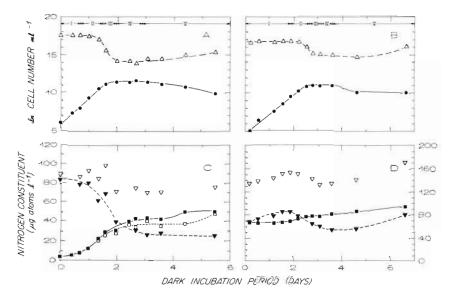


Fig. 4. Same as Fig. 3 except that prey are (A) a mixed bacterial assemblage (Experiment A) or (B) P-limited *Pseudomonas halodurans* (Experiment B)

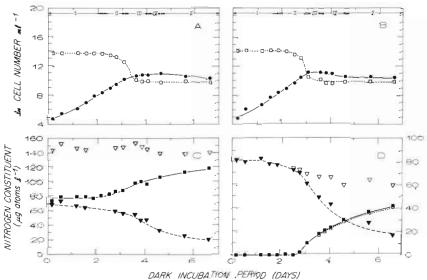


Fig. 5. Same as Fig. 3 except that prey are *Phaeodactylum tricornutum* that are (A) P-limited or (B) N-limited (Experiment C)

(Experiment A: Fig. 3A; Experiment C: Fig. 5A, B), *P. tricornutum* plus bacteria (Experiment A: Fig. 3B), or bacteria only (Experiments A & C: Fig. 4A, B). Microflagellate numbers in each experiment increased exponentially for 1 to 2 d (Intervals I & II) with corresponding reductions in prey population size. Transition from exponential growth to stationary phase (Interval III) always occurred rapidly. Prey threshold numbers corresponding to the onset of the stationary phase were about 10^4 cells ml⁻¹ for *P. tricornutum* (Fig. 3A, B & 5A, B) and about 1 to 4×10^6 cells ml⁻¹ for bacteria (Fig. 4A, B).

There was little change in cell numbers of both the microflagellate and *Phaeodactylum tricornutum* in the absence of bacteria after the stationary phase was reached (Intervals IV & V) (Fig. 3A & 5A, B). Visible aggregation of particulate material and wall growth

developed, however, in all cultures containing the mixed bacterial assemblage (Fig. 3B & 4A, B) during late stationary phase (Interval V). This aggregation was coincident with a general decrease in measured microflage. Late numbers and an increase in bacterial numbers for the duration of the incubations. Similar wall growth and aggregation was not observed in cultures containing *Pseudomonas halodurans*.

Nitrogen turnover and mass balance

Peak values of the nitrogen ingestion rate of the microflagellate (I_N) occurred during Interval II of Experiment A. Values of I_N ranged from 80 pg N cell⁻¹ d⁻¹ with only the alga present (based on curve in Fig. 3A) to 63 pg N cell⁻¹ d⁻¹ when bacteria were

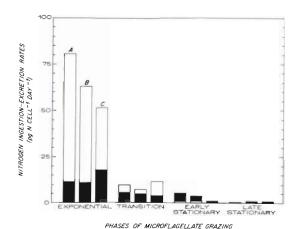
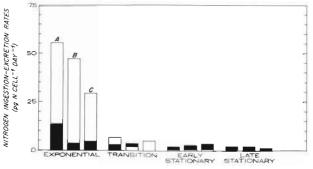


Fig. 6. Paraphysomonas imperforata. Rates of nitrogen ingestion (open bars) and excretion (solid bars) during different intervals of grazing on non-nutrient-limited cultures (Experiment A) of: Bar A: Phaeodactylum tricornutum without bacteria present; Bar B: P. tricornutum with bacteria present; Bar C: mixed bacteria alone

included with the alga (based on curve in Fig. 3B) and to $50 \text{ pg N cell}^{-1} \text{ d}^{-1}$ when only bacteria were present (based on curve in Fig. 3A) (Fig. 6). The nitrogen ingestion rate then dropped off dramatically in the transition stage in all 3 cultures and was negligible by the early stationary phase.

Ammonium excretion occurred concomitant with the onset of grazing in experiments where the prey were either not nutrient-limited (Fig. 3C,D & 4C,D) or were P-limited (Fig. 5C). The rate of $\mathrm{NH_4}^+$ excretion per cell was fairly constant through the exponential growth phase (about 11 to 17 pg N cell⁻¹ d⁻¹ depending on the experiment) but then decreased through the transition and early stationary phases until excretion was negligible by late stationary phase (Fig. 6 & 7). Urea excretion occurred during the transition or early stationary phases, but only in the cultures that were



PHASES OF MICROFLAGELLATE GRAZING

Fig. 7. Same as Fig. 6 except that Bar A: P-limited *Phaeodactylum tricornutum*; Bar B: N-limited *P. tricornutum* (Experiment C); Bar C: P-limited *Pseudomonas halodurans* (Experiment C)

originally grown on media with an N:P ratio of 10, and it never exceeded 15 % of the total N excreted (Fig. 3 C,D & 4C). Ammonium excretion in the N-limited culture of Experiment C was delayed until the late exponential phase. A low but slowly decreasing excretion rate (4 to 2 pg N cells⁻¹ d⁻¹) followed through the late stationary phase (Fig. 5D & 7). Nitrate concentrations always were below detection limits (< 0.03 μ g-at l⁻¹).

Total N (PN + Σ N) generally was conserved in all the experiments, at least through several days incubation: increases in Σ N were balanced by losses of PN. We attributed losses in total N towards the end of particular experiments (e.g. Fig. 3D & 4D) generally to bacterially-mediated aggregation and wall growth. Apparent losses of total N in the N-limited culture not containing bacteria (Fig. 5D) are not easily explainable, but may have been due to insufficient mixing that led to some observed settling of particulate material.

The nitrogen cell quota (Q_{Np}) for *Phaeodactylum* tricornutum under non nutrient-limited conditions ranged from 2.0 to 2.5 pg N cell⁻¹ (Experiment A) to 1.3 to 1.7 pg N cell⁻¹ (Experiment B). Under P-limitation Q_{Np} was 1.0 pg N cell⁻¹ and under N limitation it was $0.8 \text{ pg N cell}^{-1}$ (Experiment C) (Table 1). The nitrogen cell quota for bacteria was 0.02 to 0.03 pg N cell-1 in Experiment A and 0.05 pg N cell-1 in Experiment C (Table 1). In calculating the nitrogen cell quota for the microflagellate (Q_{Nf}) we assumed that Q_{Np} remained constant through each incubation period. During exponential growth (Interval II) Q_{Nf} was maximal at about 11 to 19 pg N cell-1, but then decreased to < 6 to 8 pg N cell^{-1} as the food supply became exhausted and starvation set in (Table 2). The magnitude of QNf seemed not to be affected by the source or nutritional state of the prey.

Nitrogen regeneration efficiency

Total nitrogen excretion rates of the microflagellate (E_N) , as discussed previously, decreased during the course of the incubations, but at a much slower rate than the ingestion rate (I_N) . In fact, some excretion of ΣN continued through the stationary phase in each experiment, reflected in the diminishing value of Q_{Nf} (Table 2). However, because the magnitude of E_N during exponential growth of the microflagellate was small relative to I_N – only 5 to 10 pg N cell⁻¹ d⁻¹ (Fig. 6) – the actual efficiency of N regeneration (R_N) during this phase of growth was very low, varying from 14.5 % when only *Phaeodactylum tricornutum* was the prey to 34 % when only bacteria were grazed (Table 2). Although R_N increased dramatically during each subsequent interval (Fig. 5; Table 2), total N

Growth interval	Di	atom	Diat	iment A om + eria*	Bad	cteria*	•	iment B eria**		mited atom	ment C N-lin diat	
	15	(15)	18	(17)	19	(34)	11	(17)	17	(24)	13	(8)
III	8	(54)	8	(66)	16	(28)	11	(0)	13	(56)	9 (>	100
ΓV	8 (> 100)	7 (>	> 100)	3	(100)	9 (>	> 100)	7 (3	> 100)	9 (>	100
V	8 (> 100)	ş (>	> 100)	Ş	(100)	ş (>	> 100)	8 (2	> 100)	6 (>	100
I-III • • •		(24)		(30)		(30)		(8)		(37)		(22)
I–IV•••		(46)		(49)		(41)		(16)		(46)		(37)
• Mixed bacterial	popula	` ,		(49)		(41)		(10)		(40)		(S

Table 2. Paraphysomonas imperforata. Nitrogen cell quota (Q_{Ni}: pg N cell⁻¹) and, in parentheses, nitrogen regeneration efficiency (R_N: %) during different intervals of grazing

regeneration through Interval IV never exceeded 49 % of the initial prey nitrogen (as PN) that was present when the microflagellate was introduced (Table 2).

Patterns of N regeneration in all 3 experiments (Table 2) were, for the most part, virtually the same although absolute values of I_N and E_N in Experiment B and C were somewhat lower than in Experiment A. However, the delay in N excretion by the microflagellate when grazing on nitrogen-limited Phaeodactylum tricornutum (Fig. 5 B, D) led to a very low (8 %) regeneration efficiency during exponential growth (Table 2). Nonetheless, because ΣN regeneration continued at a relatively high rate through the stationary phase (Fig. 5D & 7), the summed ΣN regeneration efficiencies in this experiment were comparable to those in the other incubations (Table 2).

Microflagellate growth rates

Specific growth rates (µ) of the microflagellate during Interval II appeared to be dependent on temperature during each set of experiments and not on the prey source or nutritional state: μ varied from 3.41 to 3.50 d^{-1} in Experiment A at 24 °C, to 2.10 to 2.55 d^{-1} in Experiments B and C at 20°C (Table 1).

DISCUSSION

Nitrogen turnover by microflagellates

As emphasized by Fenchel (1982a), there are important restrictions to studying grazing dynamics in batch cultures. Most important, balanced growth is never

attained. Hence, it is extremely difficult to calculate grazing rates (and concomitantly, nutrient incorporation rates by the grazer) during exponential growth when predator and prey numbers are changing rapidly in opposite directions. In this study, estimates of nitrogen ingestion per microflagellate cell (I_N) during Interval II had to be based on subjective fits by eye of the curves of prey disappearance over a relatively short period when orders of magnitude decreases in prey occurred (see Fig. 3A,B). Thus for Interval II the differences in I_N between experiments at a given temperature (Fig. 6 & 7) probably were due as much to subjective errors in curve fitting by eye as they were to any real physiological differences resulting from the different experimental treatments. The generally higher values of I_N at 24 °C - 52 to 81 pg N cell⁻¹ d⁻¹ (Fig. 6) than at $20\,^{\circ}\text{C}$ – 28 to 55 pg N cell⁻¹ d⁻¹ (Fig. 7) – most likely were related to differences in growth rate which clearly were temperature-dependent.

In spite of the difficulty in accurately estimating I_{N_1} our results show unequivocally that, relative to bacterial and phytoplankton activity, the bulk of NH_4^+ regeneration was performed by Paraphysomonas imperforata. A complicating factor in previous experiments on comparative nutrient regeneration by bacteria and protozoa has been the difficulty in separating the contribution of each microbial component when bacteria were the sole prey (Johannes 1965, Barsdate et al. 1974). In our study, because phytoplankton and bacteria both were prey, we were able to measure regeneration of nitrogen from grazed phytoplankton in the absence and presence of bacteria (Fig. 3C, D). Under these conditions we saw no appreciable difference in both the excretion rate per microflagellate during exponential growth and the pattern of regeneration over the full incubation period between the 2

microbial systems (Fig. 6). The lack of nitrogen regeneration in the bacterial control cultures (Fig. 1C, D) and the small release of $\mathrm{NH_4}^+$ by *Phaeodactylum tricornutum* only during late stationary phase (Fig. 2C, D) is clear confirmation that the microflagellate solely was responsible for the net excretion of nitrogen we observed.

Still another indication that bacteria played no role in the regeneration of nitrogen is that the rate of NH₄⁺ regeneration in the culture containing the microflagellate grazing only on bacteria was comparable to that when the prey was Phaeodactylum tricornutum (Fig. 6 & 7). Cryptic growth of bacteria occurred once the microflagellate entered the stationary phase (Interval V), most likely on excreted organic carbon (see Caron et al. 1985), both when bacteria were added to the culture of P. tricornutum (Fig. 3 B) and when bacteria were the sole prey (Fig. 4A, B). This bacterial growth occurred without any concomitant increase in nitrogen regeneration above that found in the culture containing only P. tricornutum as the prey. From these results it would appear that regeneration of NH_4^+ also was minimal during active bacterial growth, a conclusion consistent with the contemporary view that bacteria are consumers rather than regenerators of nutrients (Williams 1981, Azam et al. 1983, Hagstrom & Larsson 1984). Overall, our findings are in line with the view of Johannes (1965) and Pomeroy (1984) that protozoa rather than bacteria are the main regenerators of nutrients at the microbial level, at least in systems in which primary productivity is in reasonable balance with grazing (e.g. pelagic surface waters).

It is not surprising that NH_{Δ}^{+} was the main excretory product of the microflagellate since marine zooplankton and protozoa are considered to be primarily ammonotelic (Stout 1980, Bidigare 1983). Similarly, our finding that urea excretion was about 15 % of ΣN excreted is consistent with contemporary evidence that urea is a secondary nitrogenous excretion product of macrozooplankton. Although the magnitude of urea excretion is variable, typically it constitutes about 10 to 20 % of the nitrogen excreted by marine zooplankton (Bidigare 1983). Amino acids also can contribute 10 to 15 % of excreted N in macrozooplankton (Bidigare 1983). In our study it was possible to achieve closure of our nitrogen mass balance by summing only NH₄⁺, urea and PN; thus excretion of amino acids must have occurred at very low levels, if at all. Anderssen et al. (1985), in fact, found that of the bacterial nitrogen ingested by bacterivorous microflagellates, only 0.02 % was excreted as amino acids, principally serine, aspartic acid, glutamic acid, and ornithine. Other potential N excretion products such as purines (Soldo et al. 1978) would likewise be undetectable by our mass balance approach.

Nitrogen regeneration efficiency

Our second major finding was that during exponential growth (Interval II) the microflagellate was extremely efficient in conserving not only nitrogen (the regeneration efficiency for nitrogen was only about 15 to 25 % when the prey food was not nutrient-limited), but also phosphorus (Andersen et al. unpubl.) and carbon (Caron et al. 1985). These results are comparable to those of Anderssen et al. (1985), who found a nitrogen regeneration efficiency (R_N) of 13 % when bacteria were consumed by microflagellates. The ability to conserve considerable nitrogen even when the food source is exhausted (R_N over Intervals I to IV never exceeded 50 %) raises important questions about how nutrients are cycled in the microbial food chain.

Ammonium regeneration rates for Paraphysomonas imperforata, when normalized on a per unit dry weight basis, are comparable to those obtained for several other microflagellates grazing on bacteria and are considerably higher than those reported for at least one large ciliate and numerous macrozooplankton species (Table 3). As seen from our own study (Fig. 6 & 7), and also from the companion study on phosphorus dynamics (Andersen et al. unpubl.), the nitrogen and phosphorus regeneration rates for a given species can vary greatly depending on environmental factors such as temperature, predator nutritional state, and quality of the prey food. Thus, while we agree with the conclusion of Johannes (1964) that small protozoa have a much higher nutrient turnover potential per unit of body nutrient than macrozooplankton, we caution that for a given species these rates, like respiration rates (Fenchel & Finlay 1983, Caron et al. 1985), can be highly variable.

Virtually nothing is known of the nutritional requirements and metabolic patterns of food digestion in phagotrophic microflagellates. From a physiological standpoint it is difficult to explain why there was a lag in NH₄⁺ excretion (Fig. 5D), but not phosphorus release (Andersen et al. in prep.) only when the phytoplankton food was N-limited. A similar retention of phosphorus, but not NH₄+, occurred under P-limiting conditions (Andersen et al. unpubl.). Most likely, there are fairly rigid stoichiometric nutrient requirements of the microflagellatte while growing exponentially. Thus when there is an imbalance in the N:P ratio of the food ration the nutrient in shortest supply (in this case nitrogen) seems to be greatly conserved until exhaustion of the available food. Under N-limiting conditions synthesis of protein must dominate over the catabolic process of amino acid transamination and concomitant NH₄⁺ excretion. Just how this process is regulated biochemically is poorly understood. Conceivably, synthesis of GDH (glutamate dehydrogen-

Table 3. Comparison of NH ₄ excretion rates of marine microzooplankton and macrozooplankto	n
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Species	Temperature (°C)	NH_4^+ excretion rate ($\mu g N mg dry wt^{-1} h^{-1}$)	Sources
Microzooplankton*			
Ochromonas sp.	RT	28	Anderssen et al. (1985)
Monas sp.	20	10-17	Sherr et al. (1983)
Paraphysomonas imperforata	20	310	This study
	24	9-20	*
Euplotes vannus	25	3	Gast & Horstmann (1983)
Macrozooplankton***			
Temora stylifera	16	4.9	Debs (1984)
Eucalanus pileatus	20	1.2	Gardner & Paffenhöfer (1982
Acartia clausi	15	2.2	Mayzaud (1973)
Tigriopus brevicornis	23	1.3	Harris (1973)
Euphausia superba	1	1.5	Hirche (1983)
Natural populations (102 to 223 µm)	16-18	0.8	Smith & Whitledge (1977)

RT Room temperature

- $^{\bullet}$ Dry weight taken as 0.2 pg μm^{-3} (Sherr et al. 1983)
- ** Nitrogen-limited Phaeodactylum tricornutum
- *** Selected high excretion rates; numerous lower values are summarized in Ikeda (1974), Ikeda et al. (1982), George & Fields (1984)

ase), the enzyme thought to be responsible for regulating NH_4^+ excretion, is suppressed by production of GTP (guanosine triphosphate), a key factor in protein synthesis (Bidigare & King 1981). This possibility is in line with our observation that respiration rates of the microflagellate under these conditions were about half those that occurred when the food supply was nutritionally balanced (Caron et al. 1985); yet, interestingly, μ of the microflagellate at a given temperature remained relatively unchanged (Table 1) under all growth conditions.

Ecological implications of low nutrient regeneration

The high nutrient conversion efficiencies we observed are consistent with the well-established fact that protozoa along with bacteria have gross energy conversion efficiencies that can reach 50 to 70 % (Calow 1977, Fenchel 1982a, Sherr et al. 1983, Caron 1984). Yet, this important point has never been reconciled with the contemporary view that 80 to 90+ % of nutrients are recycled in the euphotic zone of nutrient impoverished waters (Eppley & Peterson 1979) by plankton that frequently are < 100 µm in size (Harrison 1980) and sometimes smaller than 10 μm (Glibert 1982). If our results are typical then, at most, about a 50 % regeneration efficiency is possible in a simple microbial food chain consisting of a phagotrophic microflagellate grazing on phytoplankton and/or bacteria. Moreover, this relatively high regeneration efficiency

can be achieved only when the predator has exhausted its food supply and goes through a starvation phase. This mode of nutrient regeneration would be characteristic of the feast or famine existence attributed to oceanic protozoa (Goldman 1984b, Sieburth 1984).

To reconcile these opposing views requires that the microbial food chain be more complicated then even currently envisioned (Azam et al. 1983) with, perhaps, numerous feeding steps at the microbial level. For example, inclusion of a microflagellate predator such as a ciliate with a similar nutrient regeneration efficiency of 50 % would raise the total nutrient regeneration efficiency to 75 %. An additional grazing step would raise R_N to almost 90 %. Other ways in which regeneration efficiency could increase might be through cannibalism among individual protozoan species (Fenchel 1982b) and/or by a hierarchy of sizedependent grazing among different microflagellates. Given that even more nitrogen or phosphorus is conserved by the microflagellates during active grazing when the prey is N- or P-limited, it follows that the microbial food chain would have to be extraordinarily complicated to achieve 80 to 90 % regeneration when phytoplankton growth rates were nutrient-limited. Alternatively, overall nutrient regeneration efficiencies in the euphotic zone may be lower than envisioned (Jenkins & Goldman 1985).

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