Ammonium regeneration by microplankton in an upwelling environment

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ABSTRACT: Size-fractionated measurements of ammonium flux in the southern Benguela upwelling system provided evidence that regeneration by microplankton (<200 μm) was concentrated in the upper water column and generally supported the majority of phytoplankton demands. Based on the present study and literature data it was demonstrated that ammonium remineralization was dependent on the concentration of particulate nitrogen present when viewed on a broad scale. The < 15 μm size class, including bacteria and nanoflagellates, proved to be the predominant mineralizers accounting for 95% of total microplankton regeneration on average. Although ammonification by heterotrophic picoplankton (presumably mainly bacterial) was important in some instances, bacteriovorous nanoflagellates (1 to 15 μm) appeared to supply the bulk of microplankton excretion, i.e. a mean of 87% in surface waters and 55% near the base of the euphotic zone. Bacteria were relatively more active at depth than at the surface although in the majority of cases absolute rates were fastest in surface waters. The possibility exists that bacterial remineralization was underestimated in the < 1 μm fraction due to the absence of nanoflagellates during the incubation. Nevertheless, nanoflagellate feeding comprises an essential component of the remineralization process. Ammonium uptake by the different size classes was directly proportional to chlorophyll a concentrations. Results of this study are consistent with the proposed crucial role for small protozoans as intermediaries between bacteria and higher trophic levels and as mineralizers of nutrients for primary producers.

INTRODUCTION

Since the introduction of the concept of 'new' and 'regenerated' production in stratified surface waters (Dugdale & Goering 1967) it has become increasingly evident that a considerable proportion of primary production is based upon epipelagically excreted nutrients as opposed to nutrients introduced from deeper waters. Eppley & Petersen (1979) have demonstrated that broad oceanographic regions can be distinguished from each other by the proportion of 'regenerated' primary production occurring there. Whereas in the global ocean as much as 80% of primary production is estimated to be supported by 'regenerated' nutrients, in productive coastal upwelling environments this figure is about 50% or less. Nevertheless, absolute regeneration rates and primary production are likely to show synchronous trends. One can therefore expect a concurrent increase in the activities of the organisms responsible for regenerative flux with the increase in primary production due to elevated inputs of 'new' nutrients via upwelling (Harrison 1980). Coastal upwelling thus provides a favourable system for studying epipelagic nutrient regeneration and explicit trophodynamics especially considering the important role these play in extending the impact of the original input of 'new' nutrients.

The distinction between 'new' and 'regenerated' production is largely a distinction between physically and biologically mediated supplies of nutrients for primary production. Although there is a scarcity of information of the actual rates involved, the physical processes controlling the injection of nutrients to the photic layer are better understood than the roles of different planktonic organisms in nutrient recycling. Since the pioneering studies of Harris (1959) a considerable amount of information has accumulated on nutrient excretion by the larger zooplankton species which could effectively be isolated from the phytoplankton community (i.e. > 200 μm). The recent picture that is emerging, however, is that meso- and macrozooplankton are not as important in nutrient regeneration as was originally thought and rarely account for more than 30% of phytoplankton demands (see reviews by Harrison 1980, Williams 1981, Bidigare 1983). Johannes (1965), using phosphorus as an exam-
ple, was first to suggest that marine Protozoa were considerably more important than the macrofauna in nutrient excretion. Recently, with the application of the isotope dilution technique to marine plankton studies, it has become possible to perform direct measurements of nutrient excretion by natural assemblages of microorganisms. A number of these studies extending over a wide geographical range have demonstrated for both nitrogen (Harrison 1978, Caperton et al. 1979, Glibert 1982, Koike et al. 1982) and phosphorus (Harrison 1983) that microheterotrophs <200 µm, including protozoans (e.g. tintinnid ciliates, microflagellates), metazoans (e.g. developmental stages of copepods) and bacteria supply the major share of uptake requirements. As discussed by Goldman & Caron (1985) the high assimilation efficiencies (low regeneration efficiencies) typical of each grazing step at the microbial level necessitate a close interrelationship between nutrient regeneration and the structure of the food chain. Thus an ‘open’ system with a large input of ‘new’ nutrients will be characterised by a simpler and shorter food chain than one dependent largely on ‘regenerated’ nutrients and hence a number of regenerating steps.

The aim of this study is to establish the vertical distribution of ammonium regeneration in the upper water column and to identify important classes of organisms involved using size fractionation techniques. The size fractions are treated as inclusive classes rather than size intervals. The predominant heterotrophs in the size fractions are considered to be bacteria in the <1 µm size class, bacteria and nanoflagellates in the <3 µm and <15 µm size classes, and bacteria, nanoflagellates, dinoflagellates and large ciliates in the <200 µm size class.

**METHODS**

**Sampling.** Ammonium regeneration and uptake were measured on 4 cruises during December 1983, October 1984, March 1985 and July 1985 aboard the research vessels ‘Africana’ and ‘Sardinops’ off the west coast of South Africa. All stations sampled were over the shelf area with the exception of Stn 1 which was located in deeper oceanic waters (Fig. 1). Samples were taken using plastic Niskin bottles or, in the case of surface water, a bucket. Water from the surface and either the 1% light level (3 × Secchi depth) or the thermocline region were used in the size-fractionation experiments. These samples were incubated in simulated in situ deck boxes at either 100 or 1% surface irradiance. Samples for measurement of depth profiles of ammonium regeneration were incubated in the dark. All incubation bottles were cooled by flowing surface seawater. Temperature data were obtained from continuous profiles using the ship’s CTD system aboard the ‘Africana’.

**Analytical.** Particulate matter for each size fraction was concentrated onto glass fibre filters (Whatman GF/F filters were used throughout) and frozen for later analyses. Chlorophyll a was measured according to Jeffrey & Humphrey (1975) and particulate nitrogen on a Heraeus CHN analyser. Ammonium concentrations were determined in triplicate according to the manual method described in Grasshoff (1976) scaled down to 5 ml. With the exception of Stns 3 and 4 ammonium concentrations were determined immediately after addition of the 15N tracer. Samples for these 2 stations were frozen and analysed about a week later.

**Phytoplankton and microheterotroph counts.** Samples for counting were preserved in 1% Lugol’s iodine. Bacteria were counted in thiosulphate-cleared samples using the fluorochrome acridine orange (Hobbie et al. 1977). Six different classes of bacteria were identified according to size and shape, and total bacterial biomass was calculated from the cell volumes of each class (Painting et al. 1985) using a specific gravity of 1.1 g cm⁻³ (Doetsch & Cook 1973). Phytoplankton,
tintinnid ciliates and nanoflagellates were counted using an inverted microscope. Ciliates other than tintinnids were not frequently observed. No distinction was made between chlorotic and apochlorotic nanoflagellates. Recent evidence indicates that the presence of an active chloroplast in oceanic flagellates does not necessarily preclude phagotrophy (Estep et al. 1986).

Ammonium flux experiments. All water used in the 15N experiments was pre-screened through a 200 μm plankton net. Further size fractionation was through either a 15 μm net, a 3 μm Nuclepore filter or a 1 μm Nuclepore filter. Following this 2 l was measured out and spiked with 0.2 μmol 15NH4Cl (99.7%) and the sample split. One half was decanted into a 1 l glass bottle and placed immediately in an incubator. The other half was filtered under vacuum and a portion of the filtrate used for ammonium analyses. Exactly 900 ml of the filtrate was spiked with 10 μmol NH4Cl to satisfy the mass requirements for atomic spectroscopy and stored frozen. Incubations lasted 4 to 8 h and were generally centred around midday.

Experiments were terminated by filtration (Whatman GF/F) and the filters frozen after rinsing with 50 ml 0.2 μm filtered seawater. The filtrate was used for ammonium determinations and 900 ml for determination of the aqueous ammonium enrichment was stored frozen as before. Aqueous ammonium was recovered by diffusion. Sufficient MgO was added to the thawed sample to raise the pH above 9. A 25 mm glass fibre filter wetted with 0.05 ml 6N H2SO4 was then suspended above the sample and the bottle capped tightly. The bottles were left at room temperature for about 2 wk during which time generally in excess of 50% of the aqueous ammonium was recovered on the filter. The speed and efficiency of recovery can be greatly improved by continual stirring throughout the recovery period. The filters were shaken in 5 ml double-distilled-deionised water and an aliquot removed for colorimetric determination of ammonium content. Enrichment with 15N was determined by atomic emission spectroscopy according to a Rittenberg oxidation procedure (Fielder & Proksch 1976). Particulate 15N enrichment was determined in a similar manner following Kjeldahl digestion and distillation.

Ammonium uptake rates (u) were calculated using the following equation in which correction is made for isotopic dilution (Glibert et al. 1982a):

\[ u = \frac{PE \times PN}{R \times T} \]

where \( PE \) = percent 15N enrichment of the particulate fraction in excess of the natural abundance; \( PN \) = particulate nitrogen concentration (μmol l⁻¹); \( T \) = duration of the incubation; and \( R \) = exponential average aqueous 15N enrichment: \( R = \frac{Ro}{kT} (1 - e^{-kT}) \)

where \( k = -\frac{\ln(Rt/Ro)}{T} \) and \( Ro \) and \( Rt \) are the measured values of the time course of an experiment (measured concentrations within 0.02 μmol l⁻¹ at ammonium concentrations <1 μmol l⁻¹) regeneration rates were calculated from the equation of Laws (1984):

\[ r = \frac{\ln(Ro/Rt)}{\ln(St/So)} \left( \frac{So-St}{T} \right) \]

Statistical. Curves were computer fitted to untransformed data by a least squares iterative process (STAT-PRO statistical package). The best fit of either an exponential or power function was selected on the basis of minimizing the residual sum of squares. The power function proved to give the better fit of the two. Goodness of fit (r²) was calculated from a semilogarithmic plot of the data. Analysis of covariance was used in the test for coincidental linear regressions.

RESULTS

Microscopical examination

Inverted microscope counts of the numerically important taxa at 4 of the stations sampled for the size fractionation studies are shown in Table 1. The net-plankton fraction (200 to 15 μm) was dominated by the diatoms Nitzschia spp., Chaetoceros spp., and Asterionella glacialis. Heterotrophic dinoflagellates and tintinnids were also common in this size class. Copepod nauplii were observed very infrequently. The nanoplankton (<15 μm) was comprised mainly of numerically dominant flagellates. On average 69 ± 7% (95% confidence limits, n = 14) of these nanoflagellates were <5 μm. This ratio was apparently not affected by sample depth.

There was a general tendency for bacterial biomass to be greater in surface waters (Fig. 2), in agreement with previous studies off the southeast coast of England (Joint & Pomroy 1983, Holligan et al. 1984a) and the southern Benguela region (Lucas et al. 1986). Bacterial biomass at the oceanic site (Stn 1) was smaller than over the shelf.
Table 1. Densities (cells ml"1) of the numerically important taxa at 4 of the stations in the southern Benguela region

<table>
<thead>
<tr>
<th>Species</th>
<th>Stn 5 0 m</th>
<th>Stn 5 27 m</th>
<th>Stn 6 0 m</th>
<th>Stn 6 12 m</th>
<th>Stn 7 0 m</th>
<th>Stn 7 28 m</th>
<th>Stn 9 0 m</th>
<th>Stn 9 35 m</th>
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</thead>
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<tr>
<td>Diatoms</td>
<td></td>
<td></td>
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<td></td>
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<tr>
<td>Nitzschia seriata</td>
<td>1.5</td>
<td>1.2</td>
<td>1068.6</td>
<td>814.3</td>
<td>7.5</td>
<td>0.1</td>
<td></td>
<td></td>
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<tr>
<td>N. closterum</td>
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<td></td>
<td></td>
<td></td>
<td>2.3</td>
<td>4.8</td>
<td></td>
<td></td>
</tr>
<tr>
<td>N. delicatissima</td>
<td></td>
<td>3.0</td>
<td></td>
<td></td>
<td>133.2</td>
<td>91.9</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Asterionella glacialis</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td>388.0</td>
<td>507.1</td>
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<td>Bacteriastrum lyaliun</td>
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<td></td>
<td></td>
<td></td>
<td></td>
<td>67.2</td>
<td>17.6</td>
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<td>Chaetoceros spp.</td>
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<td></td>
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<td></td>
<td>5.8</td>
<td>8.6</td>
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<td>Corethron criophilum</td>
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<td></td>
<td>0.4</td>
<td>0.2</td>
<td></td>
<td>20.8</td>
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<td></td>
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<tr>
<td>Dinoflagellates</td>
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<td></td>
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<td></td>
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<td></td>
</tr>
<tr>
<td>Protoperidium pallidum</td>
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<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
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<td></td>
</tr>
<tr>
<td>Gymnodinium spp.</td>
<td>3.0</td>
<td>2.3</td>
<td>0.8</td>
<td>1.1</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Scripsiella spp.</td>
<td>8.7</td>
<td>3.7</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Small (&lt; 15 μm) unknown</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td>99.8</td>
<td></td>
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<tr>
<td>Other</td>
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<td></td>
<td></td>
<td></td>
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<td></td>
</tr>
<tr>
<td>Nanoflagellates</td>
<td>1227.7</td>
<td>1015.5</td>
<td>1905.2</td>
<td>2610.5</td>
<td>387.7</td>
<td>207.7</td>
<td>1776.6</td>
<td>1041.6</td>
</tr>
<tr>
<td>Tintinnids</td>
<td>6.1</td>
<td>6.9</td>
<td>1.4</td>
<td>2.7</td>
<td></td>
<td></td>
<td>7.0</td>
<td>5.1</td>
</tr>
</tbody>
</table>

Fig. 2. Combined depth profile of bacterial biomass (B) measured at all stations. Open symbols: oceanic station

Fig. 3. Size-fractionated ammonium uptake (u) at the surface, the 1% light level (1%), thermocline (T), and the aphotic zone

Ammonium uptake

Ammonium uptake rates for the 4 size fractions are shown in Fig. 3. Data for Stns 5 and 6 are not available as the filters for 15N determinations were damaged during storage. Uptake rates were higher at the surface than at greater depth. It is clear that the <15 μm size class frequently accounts for the majority of ammonium uptake; being 72% on average. In fact a considerable proportion of uptake activity was in the <3 μm size class. Uptake below the euphotic zone appeared to be largely bacterial (<1 μm). Ammonium uptake rates for each size class generally followed the chlorophyll a values (Fig. 4). There was no statistical difference between the slopes and elevations of the 4 regression equations for the different size classes (p > 0.25). The common relation is described by the equation: \[ u = 0.008 \text{ Chl} \, a - 0.001 \] \( (r^2 = 0.88, p < 0.001, n = 40) \). Eppley et al. (1979) also demons-
Ammonium regeneration

The majority of ammonium regeneration by microplankton at Stns 1, 8 and 10 occurred in the upper water column above the thermocline (Fig. 5). Maximum regeneration rates at Stns 2 and 7 occurred somewhat deeper and were an order of magnitude slower than for the other stations. Ammonium concentrations, however, reached a maximum at the thermocline (Fig. 5). Since an increase in microplankton regeneration at the thermocline was only evident at the oceanic Stn 2, a likely explanation for the accumulation of ammonium at the thermocline is a reduced demand through light limitation of photosynthesis and/or utilization of alternative nitrogen sources (e.g. nitrate) by primary producers. The present findings, however, do not preclude meso- and macro-zooplankton excretion as contributing significantly to ammonium regeneration in some instances.

As with uptake, ammonium regeneration in the aphotic zone was largely bacterial (Fig. 6). No obvious generalizations can be made about the size structure of the ammonium-regenerating community at the surface and at or near the base of the euphotic zone, although bacteria did appear to be relatively more active with depth. At the surface the <1 µm fraction contributed an average of 11 % (range 1 to 35 %) to total regeneration while near the base of the euphotic zone it supplied 32 % (range 0 to 79 %). Absolute regeneration rates for each size class, however, were faster in surface waters, with the exception of Stns 5 and 6. One unifying trend that does emerge from the data is that the bulk (93 %) of microplankton ammonium regeneration was by microbes that passed a 15 µm screen. Of this some 87 % (range 66 to 99 %) of the activity was in the 15 to 1 µm size interval at the surface and 55 % (range 19 to 90 %) near the base of the euphotic zone. It is apparent that the nanoflagellate-dominated size class was responsible for a substantial proportion of total microplankton ammonium excretion. Total nanoflagellate counts (Table 1) proved to be closely correlated with ammonium regeneration rates ($r^2 = 0.69$, $p < 0.01$, $n = 8$).

Since the majority of nanoflagellates were close to 3 µm size in diameter the apparent enhancement of
regeneration for the <3 μm size class for Stn 6 may have resulted from the stimulation of nanoflagellate bacterivory in the absence of their predators after passing through the membrane filter. Alternatively, Ferguson et al. (1984) have shown that filtration of seawater through a 3 μm Nuclepore filter resulted in a significant increase in the concentration of dissolved primary amines, presumably as a result of cell rupture. Such an enrichment with dissolved organic matter is likely to stimulate bacterial decomposition and nutrient release.

Assimilation/regeneration ratios

The relation between ammonium assimilation and regeneration for the various microplankton size classes is shown in Fig. 7. Although appreciable deviations were evident at low rates, assimilation and regeneration were roughly in balance for the entire microplankton community (<200 μm). Size-related trends were not obvious in these data.

DISCUSSION

Ammonium assimilation

This study supports earlier conclusions that the majority of microplankton ammonium assimilation during non-bloom conditions is by nanoplankton (<15 μm) (Glibert et al. 1982b, Furnas 1983, Probyn 1985). This is illustrated in the seasonal study in Narragansett Bay (Rhode Island, USA: Furnas 1983) where the total phytoplankton community ammonium uptake was dominated by large chain-forming diatoms in the winter-spring and by nanoplankton (<10 μm) during the summer. Field measurements of nitrogen resource partitioning indicate ammonium may or may not constitute a similar proportion of the total nitrogen ration for both nanoplankton and the intact phytoplankton community (Furnas 1983, Nalewajko & Gar-side 1983, Probyn 1985). However, relative preference indices for ammonium, nitrate and urea suggest that there is no obvious size-related preference for either reduced or oxidised nitrogen by netplankton and nanoplankton (Furnas 1983, Probyn 1985). Picoplankton, however, do utilise proportionally more urea and ammonium than nitrate compared to netplankton and nanoplankton (Probyn 1985). There is evidence to suggest that picoplankton also utilise reduced nitrogen more efficiently (per unit phytoplankton biomass) than
larger phytoplankton species. Studies in Hawaiian surface waters (Bienfang & Takahashi 1983), the equatorial Atlantic (Le Bouteiller 1986) and Agulhas Bank waters (Probyn & Lucas 1987) have demonstrated that ammonium uptake rates as a function of chlorophyll a concentration were faster for picoplankton than nanoplanckton or netplankton. This did not appear to be the case in the present study where ammonium uptake rates and chlorophyll a concentrations for the different size fractions shared a common regression line (Fig. 4).

**Ammonium assimilation/regeneration**

There is conflicting experimental evidence regarding the time scales over which ammonium fluxes through the microplankton community are coupled. Experiments in the Southern Californian Bight and CEPEX enclosures, British Columbia, demonstrated that ammonium regeneration generally exceeds assimilation over a 24 h period (Harrison 1978). Similarly, short-term (1 to 4 h) daytime excretion measurements in such diverse regions as the Scotia Sea, Vineyard Sound, Massachusetts, USA (Glibert 1982) and Bedford Basin, Nova Scotia, Canada (La Roche 1983) indicate, in agreement with the present study, that microplankton regeneration generally equals or exceeds assimilation. It has also been shown experimentally that a steady state can exist between ammonium efflux and uptake even over a period of minutes (Glibert 1982). By way of contrast ammonium assimilation during daylight hours was shown to exceed microplankton regeneration in the Oslofjord, Norway (Paasche & Kristiansen 1982) and the Mid-Atlantic Bight (Harrison et al. 1983). These findings, however, can be reconciled with the other studies when one considers that regeneration is likely to exceed assimilation during darkness. This has been demonstrated in Kaneohe Bay, Hawaii (Caperon et al. 1979) and the Sargasso Sea (Glibert 1982). These authors demonstrated that microplankton regeneration and uptake were only in balance when integrated over 24 h due to the mismatch of fluxes over the light/dark cycle.

Harrison (1978) has shown that both ammonium assimilation and regeneration rates are faster in the euphotic zone than in deeper waters. As was the case in the present study (Fig. 5) assimilation decreased relative to regeneration with depth resulting in an accumulation of ammonium below the 1 % light level. Such a depth-related variation in assimilation/regeneration ratios might result in a marked imbalance in ammonium fluxes at discrete depths. Nevertheless, when integrated over the upper mixed layer, uptake and regeneration will be in balance (Harrison et al. 1983). These findings emphasize that when both the appropriate time and space scales are considered, microplankton excretion probably accounts for most of ammonium demand in the euphotic zone.

**Ammonium regeneration**

Large scale spatial variation in remineralization generally follows gradients in primary production (Harrison 1980) and consequently, gradients in particulate matter. Koike et al. (1982) have demonstrated a positive correlation between particulate nitrogen and ammonification in enclosure experiments. Fig. 8 shows a compilation of the present data and literature data on particulate nitrogen concentrations and ammonium regeneration rates (Harrison 1978, Caperon et al. 1979, Gilbert 1982, Gilbert et al. 1982a). The relation between the 2 variables can be described by the power equation: 

$$ r = 0.0008 \cdot PN^{2.153} $$

$$ r^2 = 0.52, p < 0.001, n = 91. $$

The nature of the relation indicates that the turnover rate of particulate nitrogen as ammonium
increases with increasing particulate concentration. It is noteworthy that the fastest regeneration rates reported in the literature (>0.2 \( \mu \text{mole } l^{-1} \text{ h}^{-1} \)) were associated with estuarine environments such as Kaneohe Bay (Caperon et al. 1979), Chesapeake Bay (Gilbert 1982) and Bedford Basin (La Roche 1983). The present range of regeneration rates for the shelf region of the southern Benguela are intermediate between those reported for coastal and oligotrophic waters.

This study confirms previous findings in very diverse oceanographic regions that the smaller microplankton (heterotrophic nanoplankton and picoplankton) are generally most important in remineralization (Gilbert 1982). The extent to which bacterial decomposition contributes directly to nutrient regeneration in the epipelagic is a contentious issue. Decomposition has been demonstrated to be greatly increased in the presence of protozoan grazers (Barsdate et al. 1974, Fenchel & Harrison 1976) which through their bacterivorous pursuits, enhance the labile dissolved organic matter pool and thereby stimulate bacterial growth and mineral turnover (Sieburth & Davis 1982, Taylor et al. 1985). An additional explanation is that bacterivores maintain bacterial populations in a state of physiological youth thereby increasing the assimilation of organic matter (Fenchel & Harrison 1976). Other laboratory studies, however, suggest that the importance of bacteria in the upper water column is not in remineralization directly but rather as packagers of labile dissolved organic matter into edible particles (Johannes 1965, Azam & Hodson 1977, Goldman & Caron 1985). This latter view is particularly appealing considering that bacteria are very efficient immobilisers of dissolved nutrients and have been demonstrated to compete successfully with phytoplankton for this resource (Rhee 1972, Krempin et al. 1981). Field studies employing \(^{15}\text{N}\), however, have shown that microplankton ammonium remineralization may at certain times, such as towards the end of a phytoplankton bloom, be dominated by organisms passing a 1 \( \mu \text{m} \) filter (Harrison 1978, Gilbert 1982, Harrison et al. 1983). Bacterial catabolism of dissolved free amino acids has been shown to be accompanied by a substantial release of ammonium amounting to some 60% of the total nitrogen regeneration in enclosure experiments (Hollibaugh et al. 1980). For at least one station in the present study, Stn 5, it appears that bacteria may be very effective nitrogen remineralizers on both an absolute and proportional basis. Generally any heterotrophic activity associated with the <1 \( \mu \text{m} \) size class is regarded as being primarily bacterial (Azam & Hodson 1977, Harrison et al. 1977) although the recent demonstration that active eukaryotic bacterivores may be <0.6 \( \mu \text{m} \) suggests this might not always be the case (Fuhrman & McManus 1984, Cynar et al. 1985).

In their model of the euphotic zone and thermocline region Moloney et al. (1986) and Newell et al. (1987) describe a succession of organisms following a pulse of 'new' nutrients, starting with bacteria and leading through nanoflagellates to larger Protozoa and mesozooplankton. Direct remineralization of nutrients by bacteria is postulated to be more important at the thermocline than in surface waters (Newell et al. 1987). The present size fractionation studies provide some support for these ideas on a relative rate basis. Ammonification in the <1 \( \mu \text{m} \) fraction contributed on average 3 times more towards total microplankton ammonium regeneration at the base of the euphotic zone and the thermocline than in surface waters. Furthermore, the absolute rates of bacterial ammonification at depth for Stns 5 and 6 were faster than at the surface. Such findings may be related to the breakdown of particulate detritus at the thermocline and dissolved organic matter at the surface as depicted in the 2-layered model of Newell et al. (in press).

Bacterial biomass and ammonium regeneration in this study were similarly distributed through the water column being highest in the surface waters. The correlation between these 2 variables (Fig. 9) can be described by the power curve: \( r = 0.1528 B^{1.334} \) \((r^2 = 0.78, p < 0.001, n = 28)\). Newell et al. (1987) have been able to predict ammonium regeneration at open sea bacterial concentrations using a linear fit to the data shown in Fig. 9 with some degree of success. The shortcomings of such a linear fit would become evident...
at the high bacterial biomass typical of coastal waters. This relation, although of limited quantitative significance, does imply that bacterial biomass is at least indirectly important to ammonium regeneration.

The present study, in demonstrating that microbes in the 1 to 15 µm size interval were the predominant remineralizers, provides support for the growing body of evidence regarding the crucial role of Protozoa in nutrient recycling in the epipelagic zone. Phagotrophic flagellates appear to dominate in the heterotrophic nanoplankton (Sieburth et al. 1978, Sorokin 1979, Sherr et al. 1984). However, small aloricate ciliates may prove to be more important in the future with the application of improved preservation and counting techniques (Sherr et al. 1986). The metabolic potential for the excretion of significant quantities of inorganic and organic nutrients by nanoflagellates has been clearly demonstrated in a number of microcosm studies (Sherr et al. 1983, Andersson et al. 1985, Goldman & Caron 1985, Goldman et al. 1985). Phagotrophic nanoflagellates are regarded as the most voracious grazers of planktonic bacteria in the sea and as such provide a critical link in energy and nutrient transfer from the smallest living component of the food chain to higher trophic levels (Haas & Webb 1979, Fenchel 1982a, Azam et al. 1983, Davis & Sieburth 1984, Sherr et al. 1984, Andersen & Fenchel 1985, McManus & Fuhrman 1986). If, as discussed by Goldman & Caron (1985), flexible feeding strategies and omnivory are common amongst nanoflagellates, they might play an important additional role in recycling nutrients incorporated by photosynthetic nanoplankton, the most active primary producers in the global ocean (Malone 1980). However, the general dependence of remineralization on bacterial biomass (Fig. 9) and the positive relation between heterotrophic bacteria and heterotrophic flagellates (Sieburth & Davis 1982, Sherr et al. 1984) provides indirect evidence for a predominantly bacterial diet for these microbes.

Paradoxically, recent laboratory experiments indicate that nanoflagellates might generally have low regeneration efficiencies. Nitrogen regeneration efficiencies of 13 to 25% have been reported for nanoflagellates fed on bacteria and phytoplankton (Andersson et al. 1985, Goldman et al. 1985). As discussed by Goldman et al. (1985) it is unlikely that a single protozoan grazing step characterized by such low regeneration efficiencies could recycle sufficient nutrients to satisfy phytoplankton requirements, or at least our present conception of the crucial importance of regenerated nutrients to phytoplankton production (Epplly & Petersen 1979). A logical conclusion of these findings is that a number of grazing steps would be required to boost the overall protozoan regeneration efficiency. The present results indicate that such a nanoflagellate predator would be in the same size range as bacterivorous nanoflagellates.

Netplankton ciliates (<15 µm) constitute an important component of the microzooplankton community which can contribute significantly to mineral recycling in the euphotic zone. The hypotrich ciliate Euplotes vannus has been demonstrated to rapidly excrete ammonium following either an algal or bacterial meal (Gast & Horstmann 1983). Verity (1985) has calculated for Narragansett Bay that ammonium excretion by tintinnid ciliates could supply about 18% of phytoplankton demands over the annual cycle with a maximum of 29% during the late summer. Field measurements in the Oslofjord, however, indicate that even during periods of peak abundance tintinnids accounted for <10% of microzooplankton excretion (Paasche & Kristiansen 1982). The accumulating evidence from laboratory experiments suggests that although weight-specific excretion rates of ciliates are considerably faster than marine crustaceans they, in turn, are much slower than for heterotrophic nanoflagellates. The clearance rates of bacterivorous ciliates are similarly 10 to 50 times slower than for phagotrophic nanoflagellates (Fenchel 1982b). Considering that microflagellates are the most dominant protozoans in the marine plankton both in terms of numbers and biomass (Sieburth et al. 1978, Davis 1982, cited in Davis & Sieburth 1984) it appears that the contribution of large ciliates to nutrient regeneration in the field is relatively unimportant. Confirmation of this conclusion is provided by the present study in which the 15 to 200 µm size interval generally contributed only a small amount to the total microplankton ammonium regeneration.

In conclusion, the present findings demonstrate that microplankton ammonium regeneration is a broad spectrum of marine environments is regulated by the concentration of particulate matter present. Remineralization increases geometrically with an increase in particulate nitrogen and decreases in an offshore direction. Generally sufficient ammonium was produced by heterotrophic microplankton to supply phytoplankton needs. Although this study did not address the larger mesozooplankton their relative contribution to community regeneration is likely to vary temporally as depicted in the model of Moloney et al. (1986), and spatially, as has been demonstrated in the western English Channel (Holligan et al. 1984b). Heterotrophic nanoplankton (mainly flagellates) appear to be the most active remineralizers in the shelf epipelagic microplankton community of the southern Benguela. It is possible that ammonium excretion by the nanoplankton fraction is the result of multiple feeding strategies amongst and predation on nanoflagellates. Direct ammonification by bacteria may be significant, espe-
cially at depth, although generally, their more important role in the remineralization process is regarded as nourishment for bacterivorous nanoplagellates. The present size fractionation techniques, however, do not pre-empt the possibility of an enhancement of bacterial decomposition rate (and ammonium regeneration) in the presence of bacterivorous Protozoa (Barsdate et al. 1974, Fenchel & Harrison 1976, Taylor et al. 1985). Results of this study are consistent with the new conceptual model of trophodynamics in the surface ocean in which a considerable portion of primary production is channelled through bacterioplankton into heterotrophic nanoplagellates.

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LITERATURE CITED


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