Mixotrophic and heterotrophic nanoflagellate grazing in the convergence zone east of New Zealand

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ABSTRACT: Nanoflagellate grazing was investigated in the subtropical convergence region off the east coast of the South Island, New Zealand, in the summer of 1995. Clearance rates were estimated using 0.5 μm fluorescently labelled beads and fluorescently labelled bacteria to represent bacterial populations and 1.0 μm fluorescently labelled beads representing picophytoplankton populations. Nanoflagellate grazing by mixotrophs was on average lower than heterotrophic nanoflagellate clearance rates per individual for all prey types, and both heterotrophic and mixotrophic nanoflagellates showed a preference for picophytoplankton-sized particles over bacteria-sized particles when grazing on artificial prey. Despite lower clearance rates per individual, higher numbers of mixotrophic nanoflagellates meant that they contributed 57% of measured grazing impact on picophytoplankton-sized particles, 40% of grazing on bacteria-sized particles and 55% of grazing on stained bacteria per day. In addition to assessing grazing rates, by identifying the major genera involved we were able to distinguish the predominant grazers in 3 water masses and investigate how changes in species composition may be linked to grazing in this region.

KEY WORDS: Nanoflagellates · Mixotrophy · Grazing · Autotrophy · Heterotrophy · Microbial food web

INTRODUCTION

The flow of carbon within the microbial food web has been the focus of many studies since the introduction of the concept of the microbial loop by Azam et al. (1983). This concept has resulted in an upsurge of interest in the role of plankton <200 μm in size. Heterotrophic nanoflagellates (HNF) and ciliated protists are now considered the primary grazers of bacteria and picophytoplankton and are largely responsible for making energy from these sources available to higher trophic levels (Porter et al. 1985).

The importance of HNF within the microbial loop has led to further investigation into the role of nanoflagellates as a group. Nanoflagellates are flagellates that fall within the size range 2 to 20 μm and consist of 3 major groups; HNF, which graze on other living cells, autotrophic nanoflagellates (ANF), which are photosynthetic, and mixotrophic nanoflagellates (MNF), which are defined in this study as organisms which are capable of combining both photosynthesis and grazing (phagosing) particles (Sanders 1991). Reviews of mixotrophy in both freshwater and marine environments report highly variable clearance rates among different MNF populations (Borass et al. 1988, Sanders & Porter 1988, Sanders 1991, Jones 1994) but suggest that MNF can be important grazers on bacteria and picophytoplankton, especially in oligotrophic waters.

Sanders et al. (1990) and Jones (1994) both propose that nanoflagellates can occur anywhere on a continuum from complete heterotrophy to complete photo-otrophy, with the degree of either process depending on both the species and the environmental conditions. Nutrient limitation has been shown to stimulate mixotrophic grazing in some cases (Sibbald & Albright 1991, Rothhaupt 1996b). Facultative heterotrophy has

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also been reported within marine flagellates, and some photosynthetic species of nanoflagellates such as *Ochro-
monas* have also been shown to switch to heterotrophy when they are light limited (Sibbald & Albright 1991, 
Rothhaupt 1996a). Mixotrophic grazing on phospho-
rus-rich bacteria in a phosphorus-limited environment 
has even been found to induce nitrogen limitation in 
the mixotrophic species (Jansson et al. 1996). Despite 
such examples of facultative heterotrophy, in most 
cases the reason for a population to change its source 
of nutrition remains unclear, with mixotrophy provid-
ing different benefits for different species and groups 
(Jones 1994). Three main reasons for the develop-
ment of mixotrophy have been proposed. These suggest 
that heterotrophy may provide a source of; (1) carbon (Bird 
& Kalff 1986); (2) inorganic nutrients (Salonen & 
Jokinen 1988); and (3) organic nutrients essential for 
growth such as vitamins and essential amino acids 
(Caron et al. 1991). Overall, the occurrence of mixotro-
phy can be viewed as a competitive strategy to deal 
with the planktonic environment, which is subject to 
rapid temporal and spatial variations.

The impact of MNF grazing is recognised as impor-
tant and in some cases makes up a substantial part of 
the overall grazing on bacteria and picophytoplankton 
populations (Sanders et al. 1989, Berninger et al. 
1991a, Hall et al. 1993). Grazing by MNF has already 
been investigated in New Zealand waters during win-
ter and autumn periods (Hall et al. 1993, James et al. 
1996); these studies have indicated that the contribu-
tion by MNF can be significant. Hall et al. (1993) 
reported mean clearance rates by phytoflagellates of 
1.1 nl ind.$^{-1}$ h$^{-1}$ on bacteria-sized prey and 0.5 nl ind.$^{-1}$ 
h$^{-1}$ on picophytoplankton-sized prey off the west coast 
of the South Island in winter. James et al. (1996) 
reported mean clearance rates of 0.11 nl ind.$^{-1}$ h$^{-1}$ on 
bacteria-sized prey and 0.8 nl ind.$^{-1}$ h$^{-1}$ on picophyto-
plankton-sized prey during autumn in both west and 
east coast waters off the South Island. These studies 
indicate both spatial and seasonal variability in the 
clearance rates of MNF.

In winter or autumn, nutrients are rarely considered 
limiting to phytoplankton growth; however, during the 
summer, inorganic nutrients are often low and are 
likely to be limiting to growth. This study was aimed at 
assessing the role of nanoflagellate grazing during this 
period with a special focus on the role of mixotrophic 
species. In addition, we aimed to identify the major 
genera and predominant grazers in the 3 water masses 
and investigate how changes in species composition 
may be linked to grazing in this region. The study was 
conducted in the subtropical, convergence and sub-
antarctic waters of the subtropical convergence region 
off the east coast of the South Island (Fig. 1).

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**Fig. 1.** Location map identifying the 3 major water masses sampled and the approximate location of sampling sites (■).
METHODS

General sampling. Samples for grazing experiments were collected at 30 stations from the RV "Gilanes" during voyage 3024 across the Chatham Rise in late summer of 1995. The Chatham Rise was selected because within this relatively small geographic zone, we were able to assess populations and clearance rates in subtropical, convergence and subantarctic water masses (Fig. 1). The convergence zone is a region where cool, nutrient-rich, subantarctic waters of the Southern Ocean meet warmer, nutrient-poor, subtropical waters (Jeath 1985).

Eight stations were sampled in the subtropical and convergence water masses and 7 in the subantarctic water mass on 2 transects across the convergence zone between 19 February and 5 March 1995 (Fig. 1). Water samples for biological and chemical analysis were collected at selected depths, in and below the mixed layer with 5 and 10 l Niskin bottles. At each station conductivity, temperature and density profiles were recorded using a Seabird 9/11 CTD profiler. Conductivity was calibrated using a Guildline Autosal Laboratory Salinometer Model 8400A.

Subsamples of water from 6 to 8 depths were filtered through acid washed Whatman GF/F filters for nutrient analysis. Ammonia (NH4-N), nitrate (NO3-N) and dissolved reactive phosphorus (DRP) concentrations were measured using the analytical methods described in Vincent et al. (1991). For measurement of chlorophyll a (chl a), 500 ml subsamples were filtered onto Whatman GF/F filters which were frozen in liquid nitrogen and kept stored frozen until analysed. Chl a was subsequently extracted using 90% acetone, with fluorescence being measured by a Perkin Elmer LS 50 B spectrofluorometer (Strickland & Parsons 1972).

Planktonic abundance. Bacterial and picophytoplankton numbers were determined by direct counts at all sampling depths. Twenty millilitre subsamples for bacterial enumeration were fixed with 1 ml of formalin, and refrigerated in the dark for a maximum of 24 h. A 5 ml subsample was stained with acridine orange for 5 min, and filtered onto prestained 0.2 μm Nuclepore filters (Hobbie et al. 1977). Bacteria were counted under blue light excitation using a Leitz compound microscope (BP 450 to 490 nm excitation, LP 515 barrier filter, RPK 510 dichromatic beam splitter). Samples for picophytoplankton enumeration were passed through a 2 μm Nuclepore pre-filter before being fixed with 1 ml of paraformaldehyde (0.2% final concentration) for at least 1 h. Duplicate 50 ml subsamples were then filtered onto pre-dyed 0.2 μm Nuclepore filters which were then mounted in a gelatin/glycerol mix, sealed onto glass slides and frozen (Hall 1991). Enumeration of eukaryotic picophytoplankton was conducted under blue light excitation (BP 450 to 490 nm excitation, LP 515 barrier filter, RPK 510 dichromatic beam splitter), resulting in a red fluorescence from chl a. Enumeration of prokaryotic picophytoplankton was conducted under green light excitation (BP 530 to 560 nm excitation, LP 580 barrier filter, RPK 580 dichromatic beam splitter), resulting in an orange fluorescence from phycoerythrin.

Samples collected for nanoflagellate enumeration were size fractionated through a 20 μm nylon mesh. The filtrate was then fixed 1:1 with ice-cold glutaraldehyde (2% final concentration) for 1 h (Sanders et al. 1989). Fixed samples were filtered onto pre-stained 0.8 μm black Nuclepore filters, stained for 5 min with 2 ml primulin, rinsed with 2 ml Tris HCl, mounted on slides and stored frozen (Bloem et al. 1986). Nanoflagellates were counted under UV excitation using a Leica compound microscope (BP 450 to 490 nm excitation, LP 520 barrier filter, FT 510 dichromatic beam splitter). Forty randomly selected fields were counted per filter. Nanoflagellate biovolumes were calculated using dimensions and approximated geometric shape (Chang 1988). Biovolumes were calculated from measurements on a minimum of 200 cells, collected at 10 m from all stations. Biovolumes were then combined and averaged within each water mass to give an average biovolume per water mass. Samples for ciliate enumeration were preserved in Lugol’s iodine (1% final concentration) and enumerated in Utermöhl chambers with a Leica inverted microscope (James & Hall 1998).

Cell carbon estimates were based on those used by Li et al. (1992), for bacteria 20 fg C cell−1, and for picophytoplankton 250 fg C cell−1. A carbon to chl a conversion factor of 50 was used to convert chl a biomass to carbon (Banse 1982).

Bacterial productivity was measured by the incorporation of [methyl-3H] thymidine into bacterial DNA following the method of Wicks & Robarts (1987). Size-fractionated primary production was assessed during simulated on-deck 24 h incubations using 14C as described in Hawes et al. (1997).

For taxonomic evaluation of the nanoflagellate population, slides prepared for the assessment of grazing were examined using epifluorescence. Six stations within each water mass were selected at random. Nanoflagellates were identified by a number of characteristic features and then grouped into classes based on these criteria. Characteristic features included criteria such as: cell size, shape; presence or absence of chloroplasts, chloroplast colour (fluorescence), num-
ber, size, type and position of flagellum; presence or absence of haptonema. Where possible, identification was taken to species level using the criteria described in Fenchel (1982a), Sournia (1986), Patterson & Larson (1991), Thronsden (1993). The methodology was not designed to be comprehensive, given that preserved cells are difficult to identify, due to a lack of motility, distortions in shape and cell damage, including loss of flagellum from some individuals. For these reasons, the use of epifluorescence combined with compound microscopy could only provide an indicative analysis of nanoflagellate composition.

**Grazing experiments.** Fluorescent microspheres, 0.5 and 1.0 μm in diameter (Polysciences Inc., Warrington, PA), were used to simulate bacteria- and picophytoplankton-sized particles, respectively. Under blue light excitation (BP 450 to 490 nm excitation, LP 515 barrier filter, RPK 510 dichromatic beam splitter), 0.5 μm microspheres fluoresced a bright yellow-green and 1.0 μm microspheres fluoresced bright red. Microspheres were added to the sample to give bacteria- and picophytoplankton-sized particle concentrations of $5 \times 10^{5}$ and $1 \times 10^{5}$ ml$^{-1}$, respectively. This resulted in an average bacteria to bead ratio of $\sim 5.0$, which corresponds to an average tracer concentration of 19% of the bacterial population. Although higher than reported in some studies (Christoffersen 1994), this is lower or similar to the tracer concentration used in other studies (Sanders et al. 1989, Marrase et al. 1992, Hall et al. 1993). Due to an initial overestimate of picophytoplankton numbers, the concentration of picophytoplankton-sized particles was, however, on average 5 times higher than the observed natural picophytoplankton populations. This induced a prey-saturated environment as described by Fenchel (1982b). In such an environment, prey uptake rate becomes independent of population density and is limited by the ingestion rate (Fenchel 1982b). Although this is not ideal and may lead to an overestimate of the uptake rate in the natural environment, it still provides a useful measure of the grazing potential of nanoflagellates on picophytoplankton populations in these waters.

To reduce clumping and minimise selection by nanoflagellates against artificial particles, all microspheres were soaked overnight in bovine serum albumin (10 mg ml$^{-1}$) (James et al. 1996). Samples were sonicated at 30 W power level for four, 2 s bursts using a Misonix XL2020 sonicator (Pace & Bailiff 1987).

Fluorescently labelled bacteria (FLB) were also used in grazing experiments. Monospecific cultures of *Escherichia coli* were grown in sterile media and then centrifuged in a Sorvall RC26 centrifuge at 15,000 rpm (22,000 × g) for 20 min, and the pellet was then suspended in 10 ml of phosphate-buffered saline solution (0.05 M Na$_2$HPO$_4$, at ambient salinity adjusted to pH 7.6). Two milligrams of 5-(4, 6-dichlorotiazin-2-yl) amino fluorescein (DTAF) (Sigma Chemical Co., St. Louis, MO) were added to the cell suspension which was then incubated in a waterbath for 2 h at 60°C. After incubation, the cells were centrifuged and washed 3 times with the phosphate-buffered saline solution to remove excess DTAF fluorescein. After the final wash, the cells were resuspended in 20 ml of 0.02 M tetrasodium pyrophosphate (PP)–NaCl solution. These solutions were stored frozen until used as described by Sherr et al. (1987). The concentration of bacteria in the solution was determined using the bacterial enumeration methods described earlier. Subsamples of the solution were thawed and sonicated at 30 W power level for four, 2 s bursts prior to use. FLB were added to give a final concentration of $5 \times 10^{5}$ ml$^{-1}$.

FLB varied in size from 0.3 to 2.0 μm (~0.8 μm average size); the natural bacterial population was less variable in size and was on average ~0.5 μm in diameter. It is likely that this size difference between FLB and the natural bacterial population would have influenced clearance rates through size selectivity (Gonzales et al. 1990). However, other factors such as chemosensory responses (Sherr et al. 1987, Léndy et al. 1991) and predator size (Havskum & Hansen 1997) also influence selectivity. Taking these factors into consideration and recognising that the size of FLB did not represent either type of fluorescently labelled bead, it was decided that clearance rates on FLB related more closely to the bacterial population and therefore were compared to clearance on 0.5 μm bacteria-sized fluorescently labelled beads.

Grazing experiments using FLB and both sizes of artificial prey were conducted at 7 stations within each water mass, from 3 selected depths above and below the deep chl a maxima. Subsamples of 300 ml were poured gently into 500 ml polycarbonate bottles and then placed in an on-deck incubator under shade cloth to approximate irradiance at the depth of the sample for 30 min to allow the assemblage to recover from handling. Solutions of either microspheres or FLB were added, and 75 ml subsamples for evaluation of nanoflagellate grazing taken at 0 and 20 min. Subsamples were fixed 1:1 with ice-cold glutaraldehyde (2% final concentration) for 1 h (Sanders et al. 1989). Duplicate samples were filtered onto 0.2 μm Nuclepore filters and the filters were stained as previously described for nanoflagellate enumerations. Ingested beads were counted under blue light excitation with a minimum of 100 consumed beads per filter being enumerated. Phototrophic nanoflagellates were distinguished from HNF by the chl a fluorescence under blue light excitation. The mean number of beads phagosed per individual for both HNF and MNF populations was calculated. This result was in turn used to estimate total.
grazing rates for both HNF and MNF, taking into account the ambient prey populations.

RESULTS

General features of water masses

In the waters off the east coast of the South Island of New Zealand, the late-summer mixed layer depth was similar for subantarctic and subtropical waters, with means of 41 and 40 m, respectively. The mixed layer in the convergence waters was shallower, with a mean of 34 m. Nitrate concentrations in the mixed layer of the subtropical and convergence waters were low in contrast to the subantarctic waters (Table 1). Chl a concentrations were similar in all 3 water masses, ranging from 0.06 to 0.86 mg m⁻³ (Table 1). In subtropical waters, 50 to 60% of the chl a in the mixed layer was in the size fraction 0.2 to 2 μm i.e. picophytoplankton; in subantarctic waters, 30 to 40%, and in convergence waters the contribution ranged between 30 and 60%.

Planktonic abundance

Bacterial numbers were relatively similar in all water masses, ranging from 0.5 × 10⁶ to 2.8 × 10⁶ cells ml⁻¹ in subantarctic waters, 0.6 × 10⁶ to 4.0 × 10⁶ cells ml⁻¹ in convergence waters and 0.5 × 10⁶ to 5.6 × 10⁶ cells ml⁻¹ in subtropical waters (Table 1, Fig. 2). Bacterial carbon dominated the carbon pool with ratios of bacterial carbon to phytoplankton carbon at 10 m of 4.5:1 in convergence waters, 2.8:1 in subantarctic waters and 2.7:1 in subtropical waters. Bacterial productivity measured by [methyl-³H] thymidine produced daily mean growth rates of 9% in subantarctic, 12% in convergence and 16% in subtropical waters.

Prokaryotic picophytoplankton numbers were more variable across the 3 water masses and were approximately 2 orders of magnitude lower than bacterial numbers. Prokaryotic picophytoplankton numbers ranged from 0.2 × 10⁴ to 4.9 × 10⁴ cells ml⁻¹ in subantarctic waters, 0.2 × 10⁴ to 10.5 × 10⁴ cells ml⁻¹ in convergence waters and 0.4 × 10⁴ to 9.1 × 10⁴ cells ml⁻¹ in subtropical waters (Table 1, Fig. 2). Primary productivity measured by ¹⁴C uptake in the <2 μm size fraction produced daily mean growth rates of 54% in subantarctic, 58% in convergence and 70% in subtropical waters. Eukaryotic picophytoplankton numbers were not reported as they represented only a small proportion of <2 μm biomass and were near or below the detection limits of the enumeration method.

HNF numbers ranged from 71 to 857 cells ml⁻¹ in subantarctic waters, 138 to 782 cells ml⁻¹ in convergence waters and 153 to 957 cells ml⁻¹ in subtropical waters (Table 1, Fig. 2). Phototrophic nanoflagellate numbers were, however, more variable within each water mass, ranging from 336 to 1728 cells ml⁻¹. In subantarctic waters all genera identified were found to possess the ability to graze. For this reason, although not every individual within these genera grazed, we decided to regard all phototrophic nanoflagellates in these water masses as MNF for analysis purposes. Within subantarctic and convergence waters all genera identified were found to possess the ability to graze. For this reason, although not every individual within these genera grazed, we decided to regard all phototrophic nanoflagellates in these water masses as MNF for analysis purposes. Within subantarctic waters, however, 3 genera representing ~10% of the phototrophic nanoflagellate population were not observed to ingest prey during our experiments. In this water mass we regarded 90% of the phototrophic nanoflagellate population as MNF.

HNF biovolume was approximately twice MNF biovolume in all water masses, which was opposite to the general trend reported previously in this region for winter and spring seasons (James & Hall 1996, Safi & Hall 1997). HNF biovolumes were similar between convergence and subtropical waters, with a mean biovolume at 10 m of 37.7 and

Table 1. Means and standard errors for biological and chemical variables at 10 m within each water mass. MNF: mixotrophic nanoflagellates, HNF: heterotrophic nanoflagellates, DRP: dissolved reactive phosphorus

<table>
<thead>
<tr>
<th></th>
<th>Subantarctic</th>
<th>Convergence</th>
<th>Subtropical</th>
</tr>
</thead>
<tbody>
<tr>
<td>MNF (mean cells ml⁻¹)</td>
<td>1232 ± 200</td>
<td>680 ± 98</td>
<td>528 ± 56</td>
</tr>
<tr>
<td>HNF (mean cells ml⁻¹)</td>
<td>689 ± 56</td>
<td>486 ± 96</td>
<td>295 ± 33</td>
</tr>
<tr>
<td>MNF mean biovolume (μm cell⁻¹)</td>
<td>28.7 ± 6</td>
<td>14.7 ± 4</td>
<td>17.1 ± 5</td>
</tr>
<tr>
<td>HNF mean biovolume (μm cell⁻¹)</td>
<td>57.4 ± 3</td>
<td>37.7 ± 8</td>
<td>37.4 ± 9</td>
</tr>
<tr>
<td>Picophytoplankton (cells ml⁻¹ × 10⁹)</td>
<td>1.1 ± 0.1</td>
<td>3.2 ± 0.4</td>
<td>1.6 ± 0.2</td>
</tr>
<tr>
<td>Bacteria (cells ml⁻¹ × 10⁹)</td>
<td>2.8 ± 0.3</td>
<td>3.0 ± 0.3</td>
<td>1.5 ± 0.1</td>
</tr>
<tr>
<td>Chl a total (mg m⁻³)</td>
<td>0.18 ± 0.8</td>
<td>0.31 ± 0.11</td>
<td>0.29 ± 0.08</td>
</tr>
<tr>
<td>Chl a &lt;2μm (mg m⁻³)</td>
<td>0.12 ± 0.01</td>
<td>0.10 ± 0.05</td>
<td>0.08 ± 0.01</td>
</tr>
<tr>
<td>NH₄ (μmol⁻¹)</td>
<td>0.74 ± 0.36</td>
<td>0.20 ± 0.17</td>
<td>0.15 ± 0.03</td>
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<tr>
<td>NO₃ (μmol⁻¹)</td>
<td>7.5 ± 2.5</td>
<td>0.45 ± 0.02</td>
<td>0.22 ± 0.01</td>
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<tr>
<td>DRP (μmol⁻¹)</td>
<td>0.6 ± 0.19</td>
<td>0.08 ± 0.02</td>
<td>0.03 ± 0.01</td>
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Fig. 2. Depth profiles of micro-organism abundance (ml⁻¹) in (a) subtropical waters, (b) convergence waters, and (c) subantarctic waters.

37.4 μm³, respectively. Similar genera were observed in these 2 water masses, with Kinotoplastidea and Chrysophyceae dominating and Choanoflagellidea also present (Thordens 1993, Patterson & Larson 1991). In subantarctic waters, however, HNF biovolumes were considerably larger, with a mean biovolume of 57.4 μm³. This was due to the presence of larger colourless dinoflagellates in addition to genera of Kinotoplastidea and Chrysophyceae.

A similar trend in biovolume was found for MNF populations in all 3 water masses with pyrmenesioflagellates, Chrysochromulina sp. and Imantonia sp. (Chretiennot-Dinet et al. 1990, Thordens 1993) identified as the dominant genera. An average biovolume for MNF of 14.7 and 17.1 μm³ was reported at 10 m in convergence waters and subtropical waters, respectively. Subtropical waters were distinguished by the presence of euglenophyceans including Eutreptiella sp. (Chretiennot-Dinet et al. 1990, Thordens 1993) and to a lesser extent Raphidophyceae including Fibrocapsa sp. (Thordens 1993), neither of which were observed in the convergence or subantarctic waters. In subantarctic waters the mean biovolume was significantly larger at 28.7 μm³, due to the presence of Dinoflagellates, Oxytoxum sp., Protop eridinium sp. and Gymnodinium sp. (Sournia 1986).

Ciliate numbers were highest in subantarctic waters with a mean of 1144 cells l⁻¹, lower numbers were observed in convergence waters with a mean of 737 cells l⁻¹ and the lowest numbers of 542 cells l⁻¹ were observed in subtropical waters. Ciliate to HNF ratios were calculated based on 10 m. Results varied between the 3 water masses; the highest mean ratio was observed in the convergence waters at 659:1, with subtropical waters at 559:1 and subantarctic waters having the lowest ratio at 404:1.

Grazing experiments

All phototrophic algal classes identified in this study, with the exception of Raphidophyceae and euglenophyceans, are reported to have members capable of phagotrophy (Borass et al. 1988). The genera belonging to these 2 classes were only found in subantarctic waters and consisted of Heterosigma, Fibrocapsa (Raphidophyceae) and Eutreptiella (Euglenophyceae). No individuals from these genera were found to exhibit phagotrophy, which concurs with the findings of Thordens (1993) and Sanders (1991).

MNF clearance rates per individual in convergence and subantarctic waters were calculated using the total phototrophic algal population. This calculation may have been an underestimation of mixotrophic grazing on a per individual basis as uptake rate per individual was averaged over both individuals that grazed and those who did not. This calculation was based on the assumption that individuals of the same species, when found in a relatively homogeneous environment, should express a similar mixotrophic capacity. We therefore calculated a mean uptake rate per individual based on the premise that all genera that exhibited mixotrophy were mixotrophic. The non-mixotrophic genera in subtropical waters were estimated to represent ~10% of the phototrophic algae present in this water mass, so MNF clearance
rates were calculated using only 90% of the total phototrophic algal population.

Despite their lower clearance rates per individual, the MNF populations contributed 55% of measured grazing on FLB, 40% of grazing on bacteria-sized artificial particles and 57% of grazing on picophytoplankton-sized particles due to their greater abundance than HNF.

Nanoflagellate uptake of FLB, and 0.5 and 1.0 μm fluorescently labelled beads occurred at all stations. Grazing by both HNF and MNF populations was highest on FLB, with mean clearance rates across all water masses of 4.8 and 3.2 nl ind. \(^{-1}\) h\(^{-1}\), respectively (Table 2). HNF uptake rates per individual on FLB were significantly higher (ANOVA \(t = 4.0, p > 0.001\)) than MNF. Bacteria-sized particles were grazed at 2.5 and 0.9 nl ind. \(^{-1}\) h\(^{-1}\) for HNF and MNF, respectively. HNF uptake rates for bacteria were also significantly higher (\(t = 3.9, p > 0.001\)) than MNF uptake rates per individual (Table 2). Both HNF and MNF had higher clearance rates on picophytoplankton-sized particles than bacteria-sized particles when grazing on artificial prey, with clearance rates on picophytoplankton-sized particles of 4.2 and 3.5 nl ind. \(^{-1}\) h\(^{-1}\), respectively. HNF uptake rate was again significantly higher (\(t = 3.9, p > 0.03\)) than MNF uptake rates per individual. We observed a high degree of variability in clearance rates between depths both within and across water masses. Consequently, we found no statistically significant changes in clearance rates between depths.

The proportion of the standing crop of bacteria removed per day by the total nanoflagellate population was highest using FLB as a tracer. Across all water masses FLB represented a removal rate of 11.0% \(d^{-1}\) of standing crop (Table 3). Stained bacteria were removed by HNF at rates ranging from 3.5 to 11.6% of the standing crop per day in subantarctic waters, 4.3 to 12.2% in convergence waters and 1.3 to 8.9% in subtropical waters. The proportion of the standing crop of bacteria removed per day by the MNF population, using FLB, was similar to HNF, with removal rates ranging from 4.5 to 13.9% in subantarctic waters, 2.8 to 13.4% in convergence waters and 1.62 to 5.9% in subtropical waters (Table 3).

The mean removal rate of standing crop using bacteria-sized (0.5 μm) fluorescently labelled beads per day by the total nanoflagellate population was much lower than observed using FLB at 4.2%. Bacteria-sized fluorescently labelled beads were removed by HNF at rates ranging from 1.0 to 6.7% of the standing crop per day in subantarctic waters, 0.8 to 6.4% in convergence waters and 0.1 to 3.0% in subtropical waters (Table 3). The proportion of the standing crop of bacteria removed per day by the MNF population was lower, with rates ranging from 0.7 to 5.1% in subantarctic waters, 0.3 to 4.3% in convergence waters and 0.1 to 1.6% in subtropical waters (Table 3).

The mean removal rate of picophytoplankton standing crop per day across all 3 water masses by the total nanoflagellate population using 1.0 μm fluorescently labelled beads was 9.5%. The proportion of the standing crop of picophytoplankton removed per day by the HNF population was double the removal of bacteria-sized (0.5 μm) fluorescently labelled beads, with rates of removal ranging from 1.2 to 7.5% in subantarctic waters, 0.9 to 7.6% in convergence waters and 0.4 to 5.9% in subtropical waters. The proportion of the standing crop of picophytoplankton removed per day by the MNF population was higher, with rates of removal ranging from 0.1 to 12.4% in subantarctic waters, 1.4 to 19.5% in convergence waters and 0.6 to 7.6% in subtropical waters (Table 3).

**DISCUSSION**

**Role of the microbial food web in the convergence zone**

The microbial food web has been reported to have a significant, but variable, role in the 3 water masses which make up the convergence zone off the east coast of New Zealand. Previous investigations in winter and spring reported that during both seasons in the subantarctic waters, and in the subtropical waters during spring, the microbial food web was the dominant pathway for carbon and energy flows (James et al. 1996, Hall et al. 1999, in this issue). During these periods the majority of phytoplankton biomass was found to be in the <20 μm size class and bacterial biomass dominated phytoplankton carbon (Hall et al. 1999). In this late summer study chl a was generally low, reflecting oligotrophic conditions (Table 1), and the majority of phyto-

<table>
<thead>
<tr>
<th>Water Mass</th>
<th>Subantarctic</th>
<th>Convergence</th>
<th>Subtropical</th>
</tr>
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<tbody>
<tr>
<td>MNF 1.0 bead</td>
<td>2.25 ± 0.29</td>
<td>3.43 ± 0.99</td>
<td>2.58 ± 0.33</td>
</tr>
<tr>
<td>HNF 1.0 bead</td>
<td>3.23 ± 0.29</td>
<td>5.23 ± 0.80</td>
<td>4.05 ± 0.55</td>
</tr>
<tr>
<td>MNF 0.5 bead</td>
<td>0.92 ± 0.09</td>
<td>1.26 ± 0.22</td>
<td>0.57 ± 0.09</td>
</tr>
<tr>
<td>HNF 0.5 bead</td>
<td>2.37 ± 0.29</td>
<td>2.62 ± 0.61</td>
<td>2.52 ± 0.27</td>
</tr>
<tr>
<td>MNF FLB</td>
<td>3.15 ± 0.29</td>
<td>3.53 ± 0.55</td>
<td>2.66 ± 0.31</td>
</tr>
<tr>
<td>HNF FLB</td>
<td>6.03 ± 0.47</td>
<td>4.45 ± 0.57</td>
<td>3.99 ± 0.54</td>
</tr>
</tbody>
</table>
plankton biomass was in the <20 μm size class in all 3 water masses. In addition, between 30 and 70% of total phytoplankton biomass in the different water masses was contributed by picophytoplankton.

The ratio of bacterial carbon to phytoplankton carbon at a depth of 10 m was 4.5:1 in convergence waters, 2.8:1 in subantarctic waters and 2.7:1 in subtropical waters. Bacterial numbers across all 3 water masses were higher than previously reported in this region (Safi & Hall 1997, Smith & Hall 1997) and ranged from 1.6 × 10^6 to 2.9 × 10^6 (Table 1). These observations are similar to Cho & Azam (1990), who report that, in low chlorophyll environments, bacterial carbon can often exceed phytoplankton carbon, commonly contributing 2 to 3 times the carbon of phytoplankton.

The biomass of the smaller prey (bacteria and picophytoplankton) in summer represented ~90% of the total combined bacterial and phytoplankton carbon, suggesting that the microbial food web dominates biological processes at this time. HNF, MNF and ciliates were included in these estimates. However, the biomass of the smaller prey in subantarctic waters was lower than in subtropical and convergence waters. This difference may be due to differences in the environment, such as temperature and salinity, which can affect bacterial and phytoplankton growth.

**Table 3. Mean removal rates and standard errors for standing stock (SS) removal per day by HNF and MNF on bacterial and picophytoplankton populations calculated using clearance rates for FLB in 3 water masses**

<table>
<thead>
<tr>
<th>Picophytoplankton</th>
<th>Subantarctic</th>
<th>Convergence</th>
<th>Subtropical</th>
</tr>
</thead>
<tbody>
<tr>
<td>MNF 1.0 bead</td>
<td>5.4 ± 0.7</td>
<td>6.1 ± 1.7</td>
<td>4.7 ± 0.6</td>
</tr>
<tr>
<td>HNF 1.0 bead</td>
<td>3.3 ± 0.3</td>
<td>5.0 ± 0.8</td>
<td>3.7 ± 0.5</td>
</tr>
<tr>
<td>Bacteria</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>MNF 0.5 bead</td>
<td>2.2 ± 0.5</td>
<td>2.2 ± 0.5</td>
<td>1.1 ± 0.2</td>
</tr>
<tr>
<td>HNF 0.5 bead</td>
<td>2.4 ± 0.3</td>
<td>2.5 ± 0.6</td>
<td>2.3 ± 0.4</td>
</tr>
<tr>
<td>MNF FLB</td>
<td>7.6 ± 0.7</td>
<td>6.3 ± 1.0</td>
<td>5.0 ± 0.6</td>
</tr>
<tr>
<td>HNF FLB</td>
<td>6.2 ± 0.5</td>
<td>4.3 ± 0.6</td>
<td>3.7 ± 0.5</td>
</tr>
</tbody>
</table>

**HNF grazing**

Clearance rates by HNF measured on bacteria-sized artificial prey (0.5 μm) across all water masses ranged from 0.1 to 12.3 nl ind.−1 h−1 (mean 2.5 nl ind.−1 h−1), a larger range than previously reported in these waters during autumn by James et al. (1996) (0.03 to 7.9 nl ind.−1 h−1, mean 0.32 nl ind.−1 h−1) but similar to other experiments carried out by Hall et al. (1993), Cynar & Sieburth (1986), Pace & Baliff (1987) and Sieraki et al. (1987).

Clearance rates measured for dead FLB in this study were on average much higher than those using 0.5 μm fluorescently labelled beads, with a mean of 4.8 nl ind.−1 h−1 for FLB compared to 2.5 nl ind.−1 h−1 for bacteria-sized fluorescently labelled beads. The cause of this selectivity is likely to be have been 2-fold; the variable size of FLB meant that, on average, they were larger than the natural population of bacteria, which could have led to selectivity by size as reported by Gonzales et al. (1990), Sherr et al. (1987) and Landry et al. (1991), however, reported significant selectivity between dead FLB and fluorescently labelled beads of the same size, with Sherr et al. (1987) reporting a maximum selectivity of 10 for dead FLB over fluorescently labelled beads and Landry et al. (1991) reporting a maximum of 20 for FLB over fluorescently labelled beads. Selectivity in the present study calculated by comparing the ratio of HNF clearance rates on 0.5 μm fluorescently labelled beads to grazing on dead FLB resulted in a selectivity of 15, mid-way between that reported by Sherr et al. (1987) and Landry et al. (1991). These results, although inconclusive, highlight the importance of prey selectivity.

For the first time in this region we identified the dominant HNF populations and found that Kinetoplastida and Chrysophyceae classes dominated grazing in all 3 water masses. In subantarctic waters, high numbers of heterotrophic dinophyceans occurred (Table 1). Dinophyceans often have a preference for prey equal to their own size as opposed to other HNF, which have a preferred predator to prey ratio size of 3:1 (Hansen et al. 1994). Due to the different predator to prey ratio exhibited by dinoflagellates, for the first time we have evidence of how HNF species composition may influence grazing across the convergence zone.

**MNF grazing**

Photosynthetic nanoflagellate populations in all 3 water masses were dominated by the pyrmenesioflagellate genera *Chrysochromulina* and *Imantonia* (Chretiennot-Dinet et al. 1990, Throndsen 1993), both of which exhibited the ability to ingest prey during our experiments. In addition, dinoflagellates which represented a significant proportion of the photosynthetic nanoflagellate population in subantarctic waters were also observed to ingest prey. The dominance of these 2 classes in all 3 water masses indicates that the majority of photosynthetic nanoflagellate genera were capable of grazing at this time. These data highlight the importance of MNF during summer in terms of both their contributions to nanoflagellate populations and their grazing on bacteria and picophytoplankton. Despite the similarities between water masses, some clear distinctions could be observed. Subantarctic waters had the highest mean biovolume for MNF populations due
to the presence of mixotrophic dinophyceans. Subtropical waters could also be clearly distinguished by the occurrence of non-mixotrophic genera from euglenophycean and raphidophycean classes. Taking this into account, MNF was still found to dominate nanoflagellate numbers, comprising 69.9, 65.1 and 63.3% of the total nanoflagellate population in subantarctic, convergence and subtropical waters, respectively.

MNF showed significantly higher clearance rates on FLB and 1.0 μm fluorescently labelled beads compared to 0.5 μm fluorescently labelled beads, with clearance rates on FLB ranging from 1.1 to 7.2 nl ind.⁻¹ h⁻¹ (mean 3.1 nl ind.⁻¹ h⁻¹). These results were lower than HNF but comparable to results from other experiments (Sanders 1991, Jones 1994). Despite this, the greater abundance of MNF meant they dominated the grazing impact on bacteria in all 3 water masses. MNF contributed 55% of the nanoflagellate grazing on FLB across all stations, 57% of the grazing on picophytoplankton-sized fluorescently labelled beads and 48% of the grazing on bacteria-sized fluorescently labelled beads. These results show the important link that MNF represent in the transfer of carbon and nutrients through the microbial food web in the convergence zone.

Grazing by the photosynthetically capable nanoflagellate populations is recognized to be energetically inefficient (Thingstad et al. 1996), which raises the question of why grazing occurred in the 3 different water masses investigated in this study. Salonen & Jokinen (1988) suggested that MNF may be seeking inorganic nutrients, and more recent studies in the northern hemisphere have also shown that nutrient limitation stimulates mixotrophic grazing (Sibbald & Albright 1991, Rothhaupt 1996b). Nutrients in both convergence and subtropical waters were at levels which could be considered as limiting to growth (NO₃-N 0.5 and 0.2 μmol, respectively), and in subantarctic waters it has been suggested that micronutrients may control phytoplankton growth (Banse 1996, Hawes et al. 1997). We concede that other factors may be influencing the switch to grazing, but from the results of this study we suggest that MNF are likely to be seeking inorganic nutrients in these waters.

**Impact of nanoflagellate grazing on picophytoplankton standing stocks**

Picophytoplankton in this study represent the second most abundant source of prey for nanoflagellates and appear to play an important role in the microbial food web in all water masses. HNF and MNF clearance rates were higher on picophytoplankton-sized fluorescently labelled beads than on bacteria-sized fluorescently labelled beads in all water masses, suggesting that picophytoplankton-sized fluorescently labelled beads were being selectively grazed (Table 2). This result agrees with James et al. (1996), who also reported a preference for picophytoplankton-sized fluorescently labelled beads over bacteria-sized fluorescently labelled beads in these waters. However, clearance rates calculated on 1.0 μm particles are also likely to have been influenced by the higher tracer density used in these experiments, thus making direct comparisons difficult to interpret. In the present study the removal of standing crop by the total
nanoflagellate population was 9.4%. This rate was higher but within the range observed by James et al. (1996), who reported clearance rates from 0.2 to 15.7% d⁻¹, and was also similar to removal rates reported in similar studies (Sanders et al. 1989, Berninger et al. 1991b, Kuosa 1991, Christoffersen 1994).

When we consider the individual water masses, we find that the observed removal rates using 1.0 µm fluorescently labelled beads account for only a small proportion of the prokaryotic picophytoplankton production. The specific growth rate of the <2 µm size fraction of chl a in subtropical waters was 70% d⁻¹, with ~8.7% of standing stock per day grazed by nanoflagellates (Table 3). Convergence waters had a specific growth rate of 58% d⁻¹ and a grazing rate of 8.4% (Table 3), and subantarctic waters had a specific growth rate of 54% d⁻¹ and a grazing rate of 10.1% (Table 3). These results are within the range reported by James et al. (1996) but lower than results from dilution experiments conducted by James & Hall (1998). The results indicate that the highest grazing pressure on standing stock occurred in subantarctic waters, where we also found the highest nanoflagellate numbers (Table 1). Overall, however, these results indicate that, unlike bacterial grazing, grazing by nanoflagellates has only a moderate impact on the potential daily production of prokaryotic picophytoplankton. Given that stained bacteria is reported to be selected over artificial bacteria-sized particles (Sherr et al. 1987, Landry et al. 1991), it is also likely that the use of artificial picophytoplankton-sized particles may result in an underestimation of grazing rates on picophytoplankton.

This study has identified both HNF and MNF as key grazers of bacteria and picophytoplankton populations in summer. The results also indicate that both HNF and MNF are key components in the transfer of bacterial and picophytoplankton biomass to higher trophic levels, with the observed clearance rates accounting for a high percentage of the bacterial production in all 3 water masses. The identification of MNF as having similar if not greater grazing potential to HNF indicates that both forms must be considered as being important in the transfer of carbon and nutrients in the microbial food web. This study also identifies for the first time the key nanoflagellate genera responsible for grazing in this region and begins the important process of linking composition to grazing rates. Overall, this work has highlighted the need to reassess our concepts of food web structure to allow a more flexible approach to organisms that have more than 1 potential mode of nutrition.

Acknowledgements. We thank the master and crew of the RV 'Gilijanes', and S. Pickmere and G. Payne for their assistance with analytical work. We also thank M. James and C. Kemp for their helpful comments on the manuscript.

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Submitted: August 14, 1998; Accepted: May 7, 1999

Proofs received from author(s): October 26, 1999