

## Microzooplankton biomass distribution in Terra Nova Bay, Ross Sea (Antarctica)

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Received 15 October 1995; accepted 15 August 1997

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### Abstract

This work describes the spatial and vertical distribution of microzooplankton (20–200  $\mu\text{m}$ ) abundance and biomass of the upper layers (0–100 m), collected during the first oceanographic Italian expedition in Antarctica (1987/1988) in Terra Nova Bay (Ross Sea). Biomass was estimated by using biovolume calculations and literature conversion factors. Sampling was carried out at three depths, surface, 50 and 100 m. The dominant taxa were made up of tintinnid ciliates, ciliates other than tintinnids, larvae of micrometazoa and heterotrophic dinoflagellates. The abundance of the total microplankton fraction had its absolute maximum in the center of Terra Nova Bay at the surface with 31 042 ind.  $\text{dm}^{-3}$ . The areal and vertical distribution of heterotrophic microplankton biomass differs from that of abundance. On the basis of hydrological conditions, phytoplankton composition and biomass and microzooplankton biomass and structure it is possible to identify three groups of stations: 1—northern coastal stations (intermediate chlorophyll maxima, microphytoplankton prevalence, low microzooplankton biomass); 2—central stations (high surface chlorophyll, nanoplankton prevalence, high abundance of microzooplankton); 3—northern stations (deeper pycnocline, nanoplankton prevalence, high microzooplankton biomass at intermediate depths).

### Résumé

L'objet de cette étude a été la distribution spatiale et verticale de l'abondance et de la biomasse du microzooplancton échantillonné pendant la première expédition antarctique italienne (1987/1988) à Terra Nova Bay (Ross Sea). Les échantillons ont été prélevés à trois différentes profondeurs: surface, 50 et 100 m. Les taxa dominantes ont été les tintinnides ciliés, les ciliés autre que les tintinnides, les larves de micrometazoaire et les dinoflagelles hétérotrophes. La valeur maximale (31 042 ind.  $\text{dm}^{-3}$ ) d'abondance du microzooplancton total a été mesurée dans le niveau de surface de la zone centrale. La distribution spatiale et verticale de la biomasse du microzooplancton hétérotrophe est différente de celle de l'abondance. Trois groupes de stations ont été mis en évidence sur la base des conditions hydrologiques, de la composition et la biomasse du phytoplancton et la structure et la biomasse du microzooplancton: 1—stations côtières du nord (valeur maximal de chlorophylle dans les niveau intermédiaires, majorité de microphytolancton et faible biomasse du microzo-

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plankton); 2—stations centrales (haute valeur de chlorophylle de surface, majorité de nanoplancton et forte abondance de microzooplancton); 3—stations du nord (pycnocline plus profonde, majorité de nanoplancton, hautes valeurs de biomasse du microzooplancton dans les niveau intermédiaires). © 1998 Elsevier Science B.V. All rights reserved.

**Keywords:** microzooplankton; Terra Nova Bay; tintinnid ciliates

## 1. Introduction

The role of heterotrophic microzooplankton in the interpretation of the biological processes occurring in the Southern Ocean increased in the last decade due to the discovery of its importance as a key link in the transfer of newly formed organic matter by nano and picoplankton to mesozooplankton (Gifford, 1988; Alder and Boltovskoy, 1991; Burkill et al., 1993, 1995).

Pico- and nanoplanktonic producers often dominate the post-bloom successional stages of Antarctic phytoplankton accounting for up to 80–90% of total autotrophic production (Alder and Boltovskoy, 1993; Garrison et al., 1993). Microplanktonic heterotrophs, which almost alone are able to feed on nanoplankton and repackage the organic material into a size suitable for consumption at higher levels of the food web, can sustain in large part the high standing stock of Antarctic crustaceans. Among the heterotrophic microplankters, ciliates (including tintinnids), microcrustacea and various invertebrate larvae are all relevant in terms of biomass. Heterotrophic dinoflagellates also contribute significantly to the microheterotrophic biomass (Dodge and Priddle, 1987; Bjørnsen and Kuparinen, 1991; Elbrächter and Zölfel, 1993) sometimes accounting for more than 50% of the total heterotrophic biomass (Alder and Boltovskoy, 1993), and more than 60% in the 0–150 m depth range (Boltovskoy and Alder, 1992). Their distribution is affected by high heterogeneity, with patches of relatively high abundance related to some particular current pattern (Dodge and Priddle, 1987). The purpose of this work is to present the areal and vertical distributions (in the upper 100 m) of heterotrophic biomass in Terra Nova Bay (Ross Sea) during an austral summer (1987/1988) and their relationship with the water column structure and chlorophyll *a* distribution.

## 2. Materials and methods

During the 1987/1988 cruise hydrological profiles were performed at each station by means of a Meerestechnik Mod. KMS multiparametric (CTD) probe with a frequency of 8 readings/s (Boldrin and Stocchino, 1990).

Downwelling PAR irradiance and downwelling, upwelling and scalar underwater irradiance were

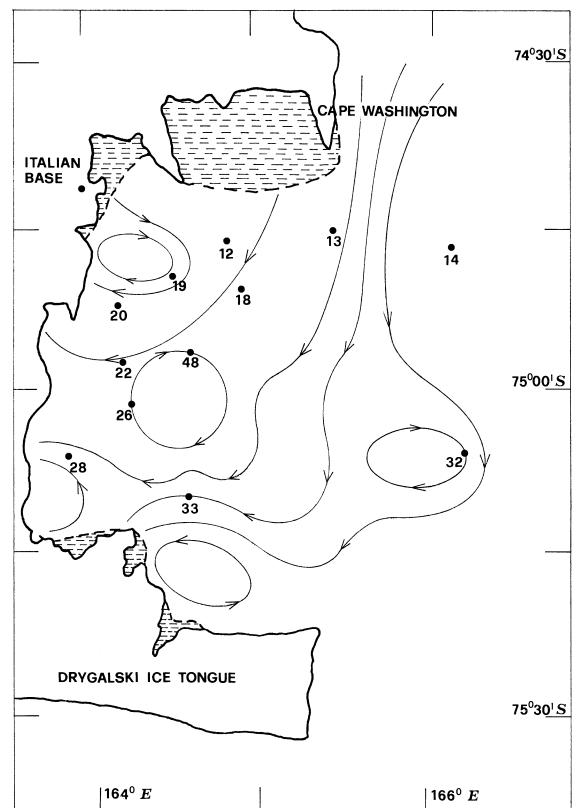


Fig. 1. Study area and sampling stations with the most important currents (redrawn from Stocchino and Manzella, 1991).

measured by means of LI-COR quantum sensor (Innamorati et al., 1990). Chlorophyll *a* concentrations were determined spectrophotometrically or fluorometrically, after extraction in 90% acetone (Innamorati et al., 1990).

From 9 January 1988 to 2 February 1988 water samples for microzooplankton analyses were obtained with sampling bottles at 3 depths (surface, 50 m, and 100 m) at 12 stations in Terra Nova Bay (Fig. 1). At each station water samples were simultaneously collected at the same depths for chlorophyll *a* concentration and nano—and micro—phytoplankton analyses. Sampling times varied between 10 AM and 8 PM. Each station was occupied once; stations 20, 19, 22, 26, 28 and 33 were sampled between 9 and 15 January; stations 12, 13, 14, 18, 48 and 32 between 24 January and 2 February. The farthest stations from the coast were stations 32 (42 miles) and 14 (33 miles); all other stations were within 15 miles from the coast.

For each sample, 2 l of water were collected, concentrated on 20  $\mu\text{m}$  mesh, and reduced to 250 cc. For dinoflagellate counting, unfiltered water was used. Samples were preserved with 4% formaldehyde solution buffered with sodium tetraborate. Subsamples (50–100 ml) were then examined in a settling chamber using an inverted microscope (magnitude  $\times 200$  and  $\times 320$ ), according to the method of Utermöhl (1958).

The biomass of microzooplankton organisms was estimated by measuring the linear dimension and equating shapes to standard geometric figures or combinations of them. Measurements were taken for each species (or groups of eggs, larvae, etc.) on more than 100 individuals (max. 394 for *Cymatocylis drygalskii*) for the abundant taxa and on the few present individuals for rare taxa. Conversion factors used to transform these volumes into organic carbon values were, dinoflagellates:  $\text{pg C} = \mu\text{m}^3 \times 0.13$  (Edler, 1979); tintinnids:  $\text{pg C} = \mu\text{m}^3 \times 0.053 + 444.5$  (Verity and Langdon, 1984); ciliates other than tintinnids:  $\text{pg C} = \mu\text{m}^3 \times 0.14$  (Putt and Stoecker, 1989); all other groups:  $\text{pg C} = \mu\text{m}^3 \times 0.08$  (Beers and Stewart, 1970).

The dinoflagellates included in our computation were all species belonging to the *Protoperidinium* genus (Dodge and Priddle, 1987; Elbrächter and Zölfel, 1993) or *Gyrodinium lachryma* (Lessard and

Swift, 1986) and *Gymnodinium* sp. (Bjørnsen and Kuparinen, 1991).

The microzooplankton composition at each station and at the three depths was organized in a matrix of 34 rows (taxa) and 36 columns (stations depths).

Cluster analysis, based on the complete linkage method, was computed by MATEDIT (Burba et al., 1992) on the matrix after calculating the correlation coefficient on species and the similarity ratio on stations' depths.

### 3. Results

#### 3.1. Hydrology

The study area in Terra Nova Bay extended approximately between  $74^{\circ}30'$  and  $75^{\circ}30'S$  and  $164^{\circ}00'$  and  $166^{\circ}00'E$  (Fig. 1). The maximum depth ( $> 1100$  m) is in the central part of the Bay bordered by a high and rocky coast and characterized by an extremely steeply sloping sea-floor (Brambati et al., 1989). The hydrology of the study area, during the summer period is characterized by the presence of a stable pycnocline and high temperature and salinity gradients. The thick surface layer of Antarctic Surface Water (AASW) has temperatures exceeding  $0^{\circ}\text{C}$ . In the coastal areas the surface layer is characterized by a salinity of  $> 34.7$  psu and temperatures between  $1.9^{\circ}\text{C}$  to  $2.1^{\circ}\text{C}$ . This area is affected by prevailing surface thermal processes due to both atmospheric and orographic phenomena which can maintain the polynya over winter. Eastward, the offshore surface water is affected by dilution processes due to meteoric ice melting. Consequently, the salinity is  $< 34.6$  psu. Below the AASW a homogeneous water body of Ross Sea Shelf Water (RSSW) is present. Here the salinity increases slightly towards the bottom with an average temperature of  $-1.9^{\circ}\text{C}$  and salinity of 34.9 psu (Boldrin and Stocchino, 1990).

During the 1987/1988 cruise pycnocline deepened from 50–75 m depth of the first sampled coastal stations (19, 20, 22, 26, 28 and 33) to 100 m depth of the last northern stations (12, 13, 14 and 18); the easternmost station 32 and central station 48, sampled during the second period, still showed pycnocline at 50–75 m depth (Fig. 2); the latter was

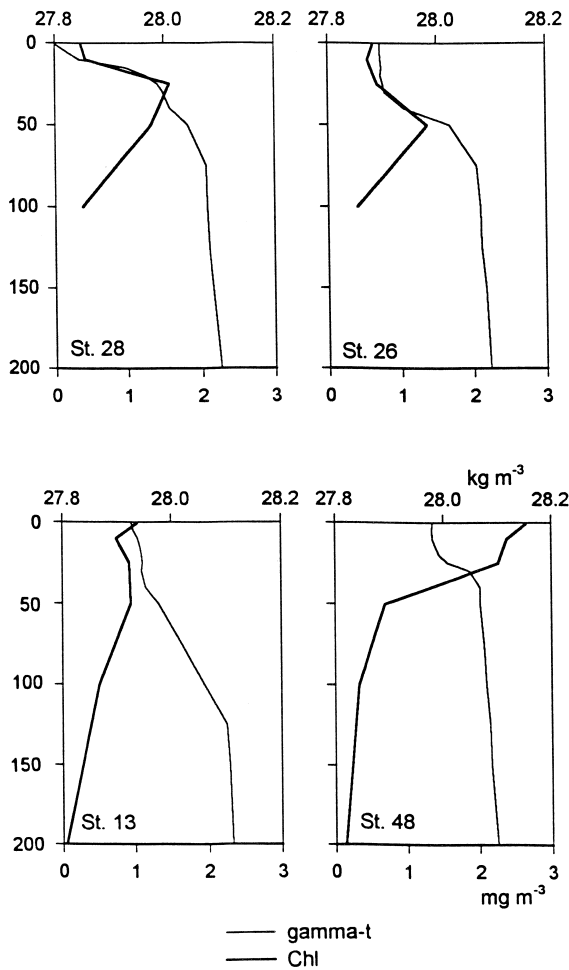


Fig. 2. Density and chlorophyll *a* typical profiles of the three main group of stations: stations 28 and 26 show profiles of coastal stations (19, 20, 22, 26, 28 and 33); station 13 shows profile of northern stations (12, 13, 14 and 18); station 48 shows profile of central stations 48 and 32.

characterized by the lowest temperatures (from  $-0.6^{\circ}\text{C}$  at surface to  $-2.0^{\circ}\text{C}$  at 300 m depth). At all other stations surface temperature was  $> 0.6^{\circ}\text{C}$ , with maxima at the most southern stations 28 and 33 ( $> 1.6^{\circ}\text{C}$ ). Only at station 20 surface temperature was  $< 0^{\circ}\text{C}$ . At the surface the lowest salinity was found at the easternmost stations 14 and 32 ( $< 34.6$  psu); while at all other stations it was  $> 34.75$  (Boldrin and Stocchino, 1990).

The surface currents are constituted by two branches (Fig. 1): the first originates from the NE in

the northernmost part of the Bay and the second from the E in the southernmost region south of  $75^{\circ}\text{S}$ . The NE current invades the central area and then turns N to become the coastal current. The second one, from the E, after reaching the Drygalski Ice Tongue, turns N to join the coastal current. Generally the surface circulation is characterized by a stable cyclonic gyre (Stocchino and Manzella, 1991).

### 3.2. Chlorophyll *a* and phytoplankton distribution

Absolute surface chlorophyll maxima ( $> 2.3$   $\text{mg m}^{-3}$ ) were recorded at stations 32 and 48; at stations 12, 13, 14 and 18 less intense ( $< 1.3$   $\text{mg m}^{-3}$ ) surface or subsurface maxima were registered (Fig. 3). At all other stations, the chlorophyll maximum occurred at depths ranging from 25 to 50 m, corresponding to the different water mass structures (Fig. 2). At stations 32 and 48 the euphotic layer depth ( $Z_{\text{eu}}$ ) was particularly thin; where 1% of surface PAR recorded at 21 and 22 m depths, respectively.  $Z_{\text{eu}}$  was maximum at station 14 (43 m). Integrated over 100 m depth chlorophyll (Fig. 4) was higher in the coastal northern area, followed by the central part of the grid.

Phytoplankton abundance (Fig. 5) was maximum at station 48 (25 m depth), reaching  $963 \times 10^3$   $\text{cell l}^{-1}$ ; at this depth 90.77% of total phytoplankton was constituted by nanoplankton. This station was the richest also at surface and 10 m depth, and still nanoplankton was the dominant fraction, as well as at stations 14, 18, 20, 32 and in the upper layers of 12 and 13. Microphytoplankton prevailed at stations 19, 22, 26, 28 and 33; nanoplankton generally increased below 50 m depth. Total phytoplankton abundance at these stations ranged between  $2 \times 10^3$  and  $166 \times 10^3$   $\text{cell l}^{-1}$ . Generally, the highest total abundances of phytoplankton corresponded to the highest percentages of nanoplankton.

### 3.3. Microzooplankton abundance and biomass

Abundance of the total microplankton fraction (including dinoflagellates, tintinnids, aloricate ciliates, larvae and eggs of invertebrates) is maximum (Fig. 6) at station 48 at the surface ( $31042$   $\text{ind. dm}^{-3}$ ), due to the conspicuous contribution of *Pro-*

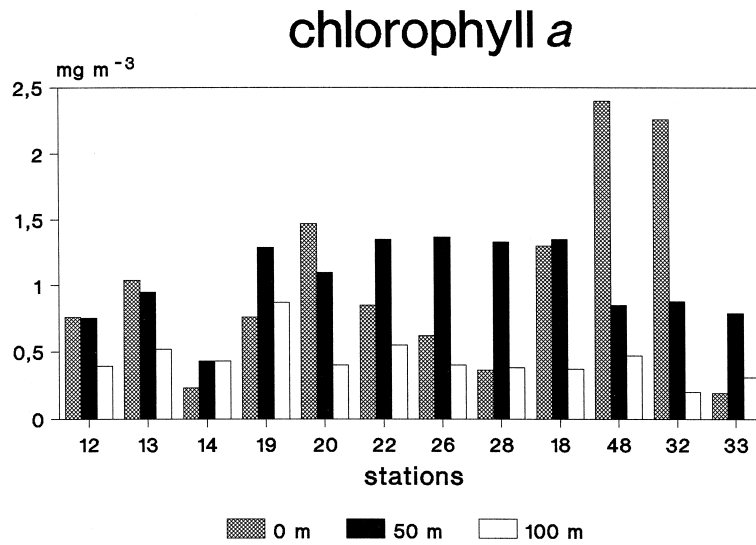


Fig. 3. Chlorophyll *a* values ( $\text{mg m}^{-3}$ ) at surface, 50 and 100 m.

*tooperidinium defectum*, *P. mediocre* and *Gymnodinium* sp. At the surface at all other stations, values are lower and variable. The contribution by dinoflagellates is considerable everywhere. At intermediate depths numbers were more uniform than in surface waters and dinoflagellate contribution to the total abundance was of the same order as at the surface

(Fig. 7a and b). At 100 m, the maximum, although less than  $20\,000 \text{ ind. dm}^{-3}$ , appears at station 13; at all other stations values are low, particularly when dinoflagellates are absent (Fig. 7c).

In Table 1, where individual biovolume and carbon content calculated for each organism is reported, it is noticeable that the biomass values of some dinoflagellates, in particular *Protoperidinium antarcticum* and *P. pseudoantarcticum*, are in the same range as tintinnid carbon values. The other dinoflagellates are 10–20 times less abundant in terms of carbon biomass.

The spatial and vertical distribution of heterotrophic microplankton biomass shows a quite different pattern from that of abundance (Fig. 8). At the surface, the richest station was 13, while at station 48, biomass is relatively high but of the same order of magnitude as at all other stations. At intermediate depths, stations 13 and 14 exhibited biomass maxima more than 4 times higher than at all other stations. Only at 100 m depth at stations 14 and 13 was there a good correspondence between abundance and biomass distribution.

The high values at stations 13 and 14 are due to the conspicuous presence of dinoflagellates (Fig. 9), which account for more than 97% of the total (station 14; 100 m). At these stations, a bloom of the very large *P. antarcticum*, whose carbon content is

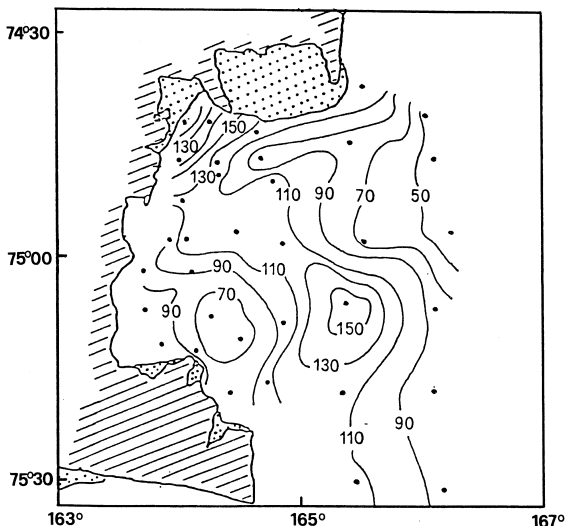
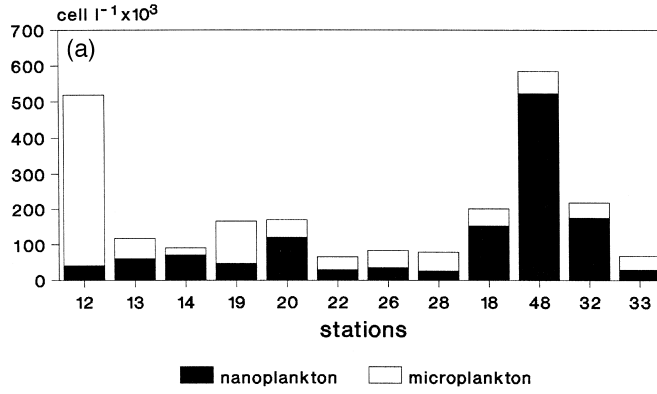
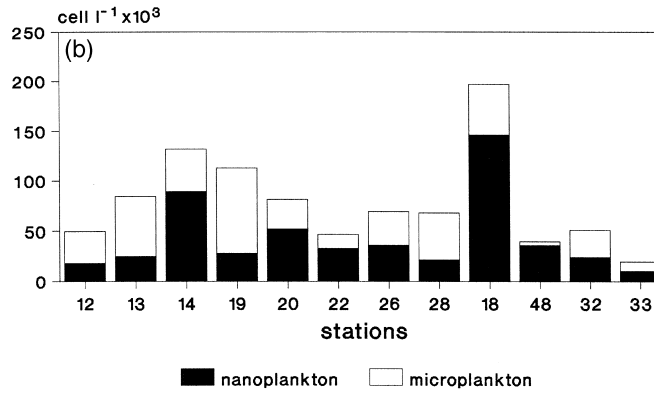


Fig. 4. Contour map of integrated over 100 m depth chlorophyll *a* (from Innamorati et al., 1992).

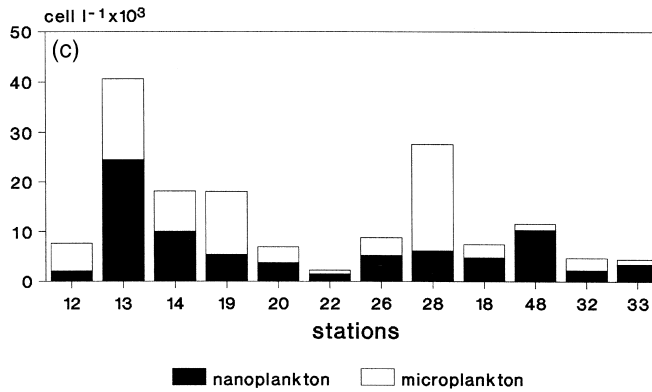
# Phytoplankton abundance surface



## 50 m



## 100 m



## Microzooplankton abundance

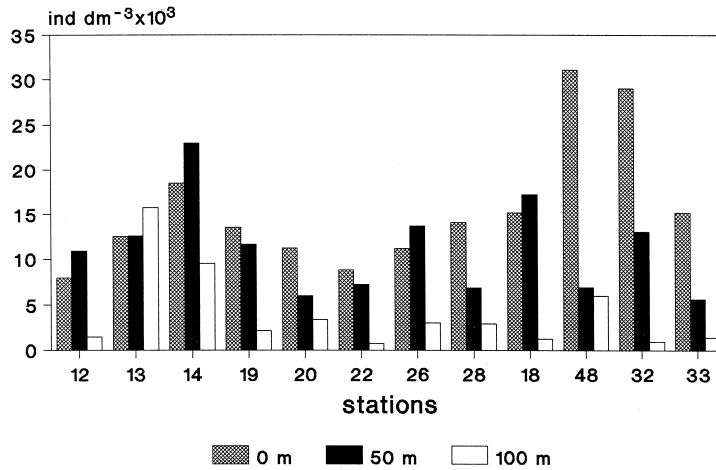


Fig. 6. Total microzooplankton abundance (ind. dm<sup>-3</sup>) at surface, 50 and 100 m depth.

60930 pg C (Table 1), was particularly developed at 50 m depth.

At all other stations the contribution of dinoflagellates to the total biomass is generally higher at the surface. The differences observed between abundance and biomass distribution are mainly due to the different composition of the heterotrophic dinoflagellate community at the various stations: e.g., at station 48 (surface) *P. defectum* and *P. mediocre* dominated, characterized by a small biovolume and consequently low carbon content (Table 1).

The contribution to the total biomass of ciliates other than tintinnids is again more important at the surface with a maximum of 26% at station 22 and generally it is more than 10%.

Tintinnids show generally the highest percentage at intermediate and bottom layers with a maximum of 82% at station 22 at 100 m.

The biomass of eggs and larval forms of invertebrates, among which the contribution of copepod nauplii is not very significant, was more abundant in the northern coastal stations (12, 18, 19 and 20),

reaching a maximum of 31% of the total biomass at stations 19 at 50 and 100 m.

### 3.4. Community structure

Cluster analysis defined 2 main stations' depth groups: 1 and 2 (Fig. 10); group 2 is comprised by 100 m samples of stations 19, 20, 22, 12, 33 and 26.

Group 1 can be divided into six subgroups: 1.1 and 1.2 identified surface samples of stations 13, 22, 12, 33, 14, 19, 28 and 50 m samples of station 19. Group 1.3 included surface samples of stations 18, 26 and 32 and 100 m samples of stations 28 and 32. Group 1.4 comprehended 50 m samples of stations 18, 28, 26, 32, 22, 33 and 12; group 1.5 samples at 50 and 100 m depths of stations 13 and 14. In group 1.6 are included all the depths of station 48, surface and 50 m of station 20 and the 100 m sample of station 18.

Cluster analysis applied on species identified six groups of species (Fig. 11): only group C is more or less homogeneously distributed in all the stations' depth groups (Table 2), being constituted primarily

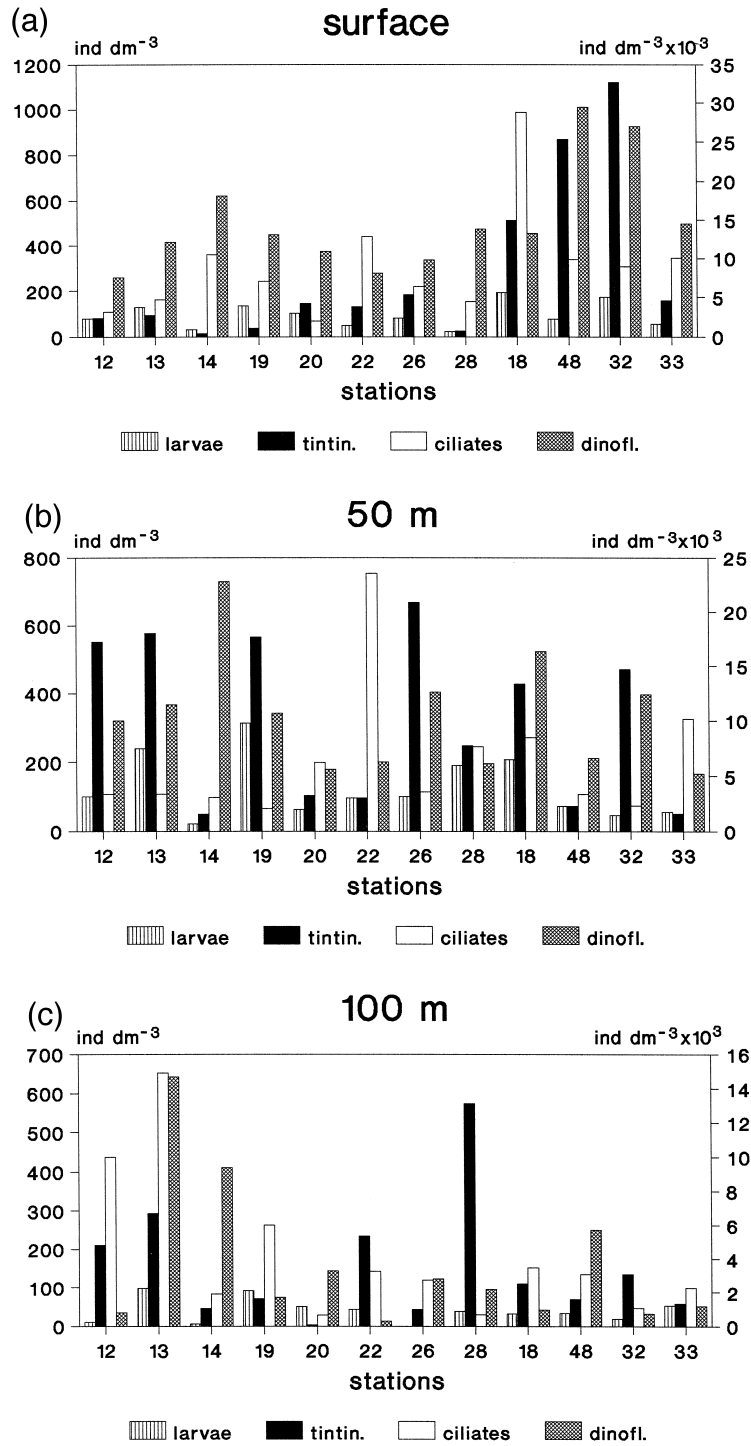


Fig. 7. Microzooplankton abundance (ind.  $dm^{-3}$ ) of dinoflagellates, tintinnids, ciliates and larvae and eggs, at surface (a), 50 m (b) and 100 m (c). Second scale is used to show dinoflagellate abundances.



Table 1  
Individual biovolume and carbon content

	$\mu\text{m}^3$	pg C
<b>Dinoflagellates</b>		
<i>Gymnodinium</i> sp.	4320	561
<i>Gyrodinium lachryma</i>	3470	4520
<i>Protoperidinium adeliense</i>	7600	990
<i>P. antarcticum</i>	468700	60930
<i>P. applanatum</i>	11970	1560
<i>P. defectum</i>	6790	880
<i>P. pseudoantarcticum</i>	278220	36170
<i>P. mediocre</i>	14930	1940
<i>P. unipes</i>	24870	3230
<b>Tintinnids</b>		
<i>Cymatocylis drygalskii</i>	467440	25220
<i>C. flava</i>	800220	42860
<i>Cymatocylis</i> sp.	513980	27680
<i>Laackmanniella naviculaefera</i>	384200	20810
<i>L. prolongata</i>	312870	17030
<i>Laackmanniella</i> sp.	247100	13540
<i>Codonellopsis gaussi</i>	98770	5680
<i>C. glacialis</i>	57270	3480
<i>Codonellopsis</i> sp.	116000	6590
<i>Salpingella</i> sp.	251200	13760
<i>Coxiella frigida</i>	72350	4280
<i>Ptychocylis</i> sp.	108090	57730
<b>Ciliates</b>		
<i>Ciliates</i> A	36250	5070
<i>Enchelys</i> sp.	61790	8650
<i>Prorodon</i> sp.	101900	14260
<i>Lacrymaria</i> sp.	9650	1350
<i>Halteria</i> sp.	127510	17850
<i>Strombidium</i> sp.	21040	2940
<i>Patronella</i> sp.	100970	14140
<i>Leegardiella</i> sp.	27450	3840
<i>Lohmaniella</i> sp.	25960	3630
<i>Oligotricha</i> sp.	84780	11870
<b>Larvae and eggs</b>		
Nauplii	367500	29400
<i>Limacina veliger</i>	569890	45590
Larvae unid.	145680	11650
Eggs unid.	112530	9000

by *Cymatocylis* sp. and rare species such as *Sticholonche zanclea* and *Laackmaniella naviculifera*.

Group A was constituted by heterotrophic dinoflagellates, is typical of 50 and 100 m depths of northern offshore stations (13 and 14).

Groups B and D species characterized the intermediate depths of group 1.4 (above). In group B, *P. defectum* and *Laackmaniella prolongata* prevailed,

in group D, *Lohmaniella* sp., *Leegardiella* sp. and unidentified ciliates as well as *Cymatocylis flava* prevailed.

Group E species in which *Cymatocylis drigalskii* prevailed, characterized the upper layer of the coastal stations (group 1.1).

Group F species of which the most abundant is *Codonellopsis gaussi* characterized the upper layers at stations 18, 26 and 28 and the 100 m depth of stations 28 and 32.

Samples from coastal stations at 100 m depth of (group 1.6) were characterized by the absence or scarcity of all groups of species.

#### 4. Discussion and conclusion

On the basis of the water column structure it is possible to identify 3 main groups of stations: Group 1: coastal stations 19, 20, 22, 26, 28 and 33, sampled at the end of microphytoplankton bloom, with a pycnocline located between 50 and 75 m depths, maxima of chlorophyll located just above the pycnocline and intermediate values of  $Z_{eu}$ ; Group 2: stations 32 and 48, sampled during the second part of the cruise, with a pycnocline still between 50 and 75 m depths, high chlorophyll maximum at surface and the shallowest euphotic layer; Group 3: northern stations 12, 13, 14 and 18, sampled during the second part of the cruise at the beginning of nanoplankton bloom, with a deeper pycnocline at, about 100 m depth, less intense surface maxima of chlorophyll and deeper  $Z_{eu}$ .

In group 1 total phytoplankton abundance was lower with a higher percentage of the microphytoplankton fraction in the upper layer and an increase of the nanoplankton fraction in the deeper layers, with the sole exception of station 20 which exhibited features similar to group 3 (see below).

In group 2, phytoplankton abundance was the highest with maxima in the upper layers constituted almost exclusively by nanoplankton.

In group 3, phytoplankton abundance was still high, characterized by high percentages of nanoplankton and maxima at 25–50 m depths.

Group 1 was characterized by low microzooplanktonic biomass, which was slightly higher at interme-

## Heterotrophic biomass

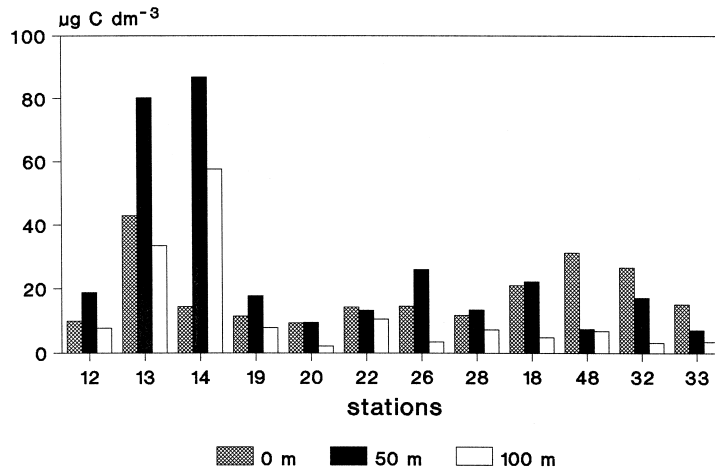


Fig. 8. Total heterotrophic biomass ( $\mu\text{g C dm}^{-3}$ ) at surface, 50 and 100 m.

diated depths. Microzooplankton abundance showed the same pattern.

Group 2 was characterized by intermediate values of heterotrophic biomass with surface maxima, mostly due to tintinnids. Here too microzooplankton abundance was the highest with surface maxima.

In group 3, two stations (13 and 14) showed a particularly high microzooplankton biomass at 50 and 100 m depths while all other stations belonging to this group showed high abundances at intermediate depths.

Cluster analysis clearly separated stations 13 and 14 at 50 and 100 m depths from the others, due to the high percentage of some species of heterotrophic dinoflagellates such as *Protoperdinium antarcticum* and *P. mediocre*. These stations are the most northern and offshore and they were sampled at the end of the cruise.

Stations 48 and 20 at all the depths were also separate from the others due to the prevalence of *P. defectum* at surface and to the presence of *Laackmaniella* spp., Pteropoda larvae and the richness of unidentified ciliates at all depths.

Surface samples of stations 18, 26 and 32 were grouped together with 100 m samples of stations 32 and 28 due to the dominance of *Codonellopsis gaussi*.

For all other stations, cluster analysis generally separated the water column into three discrete layers

(surface, intermediate and bottom), indicating that depth impose structure on microzooplankton communities more than other variables. This is true also for phytoplankton communities (Nuccio et al., 1992) which showed well defined structures at each layers, particularly at 50 m depth where a *Phaeocystis* bloom was recorded at all the stations of the grid.

Water mass structure changed during the cruise which covered firstly coastal stations and later off-shore stations: the pycnocline deepened from the former to the latter, reaching 100 m depths. Chlorophyll distribution changed from maxima at intermediate depths (25–50 m) to surface maxima and phytoplankton composition shifted from microplankton to nanoplankton dominance. In summary, it would appear that the first leg of the cruise encountered the end of the initial phytoplankton bloom (Innamorati et al., 1992; Nuccio et al., 1992), characterized by high percentages of microplankton while the second leg encountered the onset of a secondary bloom, mostly due to nanoplankton.

The contribution of heterotrophic biomass (10–200  $\mu\text{m}$ ) appears more significant in the lower layers and in particular at 50 m depth, which corresponds to the pycnocline and to the base of the photic zone during the first cruise leg. This particular vertical distribution might be explained by the predator/prey relationships characteristic for microplankton con-

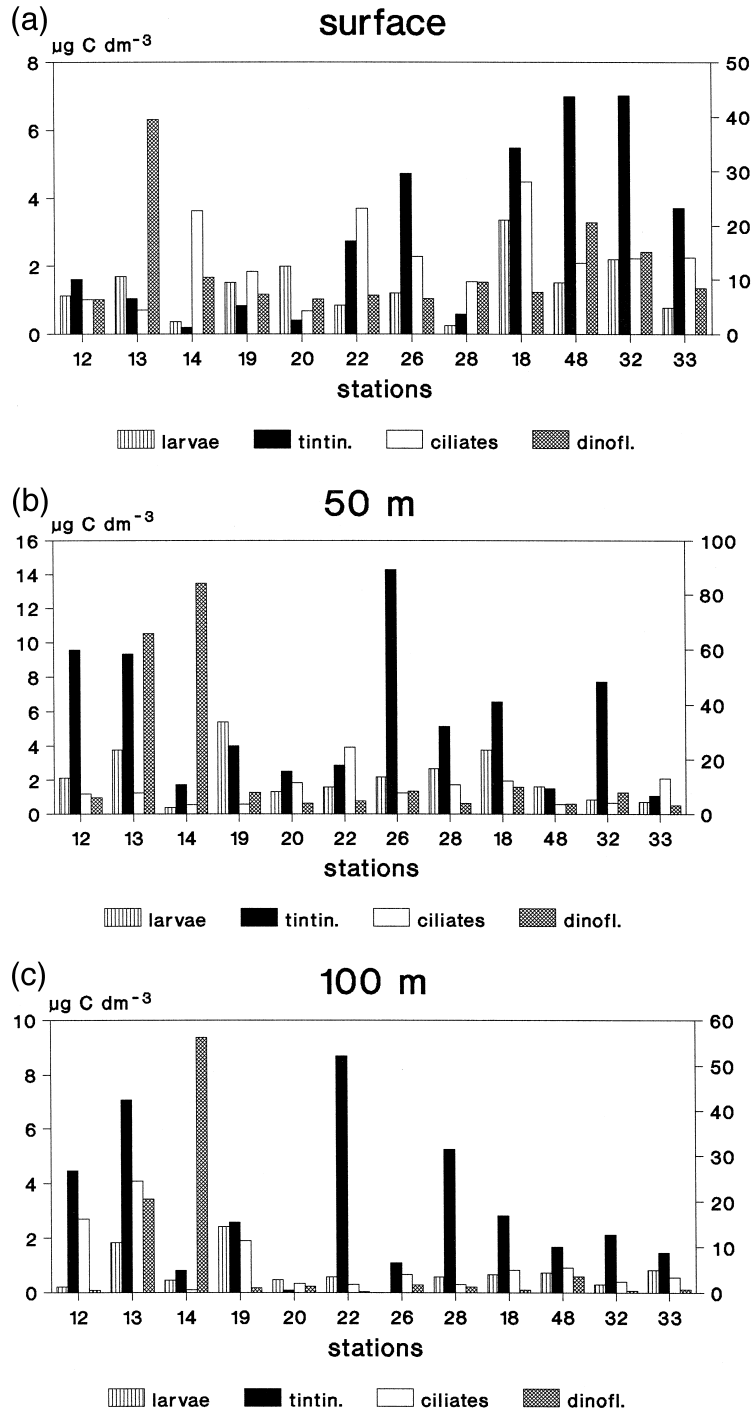


Fig. 9. Heterotrophic biomass ( $\mu\text{g C dm}^{-3}$ ) of dinoflagellates, tintinnids, ciliates, larvae and eggs, at surface (a), 50 m (b) and 100 m (c). A second scale is used to show dinoflagellate biomass.

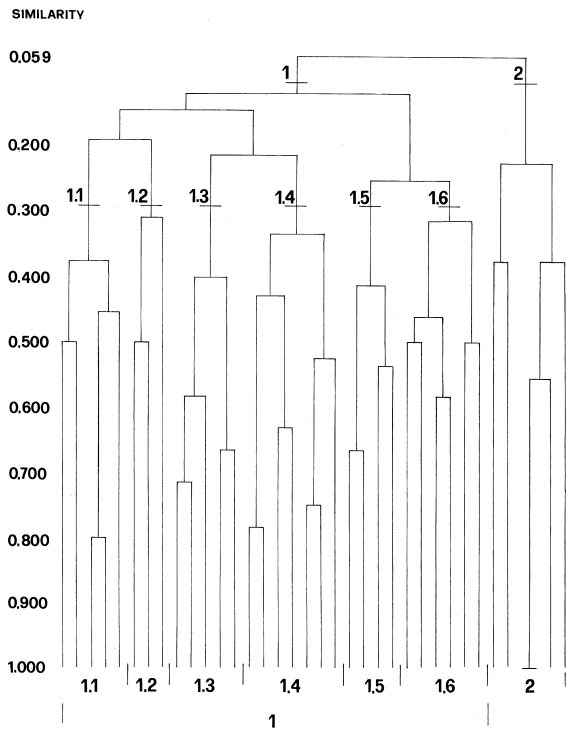


Fig. 10. Grouping of stations using the dendrogram classification obtained by processing the species/samples matrix with the similarity ratio and complete linkage method.

sumers which generally feed on particles  $< 20 \mu\text{m}$  (Rassoulzadegan et al., 1988; Pierce and Turner, 1992). Below the 25% of irradiance depth (4.5–11 m) there was an increase in the picoplankton contribution to total primary production (Magazzù and Decembrini, 1992). The relative abundance of this fraction as well as that of nanoplankton at the base of the photic zone could be considered the most important factor influencing the vertical distribution of heterotrophic microplankton biomass even if other factors (i.e., vertical stability of the water column, lower inter specific competition, etc.) could have contributed to determining this distribution pattern.

The role of microzooplankton consumers in the Southern Ocean carbon appears less important at the surface than in aphotic layers where they utilize primarily pico- and nanoplankton producers and slow down the sinking of organic material through the production of small faecal pellets with lower settlement rates than large pellets produced by macrograzers (Longhurst, 1991).

Our values of total heterotrophic biomass appear consistently higher than those found in the Weddell-Scotia Confluence Area in 1988/1989 where maximum average values were less than  $4 \mu\text{g C dm}^{-3}$  at 20 m depth (Alder and Boltovskoy, 1993) or less than  $8 \mu\text{g C dm}^{-3}$  in the Weddell Sea in 1989 (Boltovskoy and Alder, 1992). This is probably due to the great contribution of heterotrophic dinoflagellates which were present in our samples, averaging  $12.23 \pm 19.1 \mu\text{g C dm}^{-3}$ . Dinoflagellates generally show lower values: e.g., from 0.35 to  $1.95 \mu\text{g C dm}^{-3}$  in the Weddell Sea in 1987 (Boltovskoy et al., 1989) and from 0.02 to  $13.2 \mu\text{g C dm}^{-3}$  in the Confluence Area (Alder and Boltovskoy, 1993). Their values of tintinnid carbon ranged respectively from 1.64 to  $9.05 \mu\text{g C dm}^{-3}$  and from 0.02 to  $9.28 \mu\text{g C dm}^{-3}$ . Again in the Weddell Sea, Buck et al. (1992) found tintinnid carbon ranging from 0.02 to  $1.3 \mu\text{g C dm}^{-3}$ . Our mean value of  $3.88 \pm 9.3 \mu\text{g C dm}^{-3}$  fits well in the range found in the previous cruises.

As regards areal distribution, dinoflagellates strongly influence the general pattern which exhibits

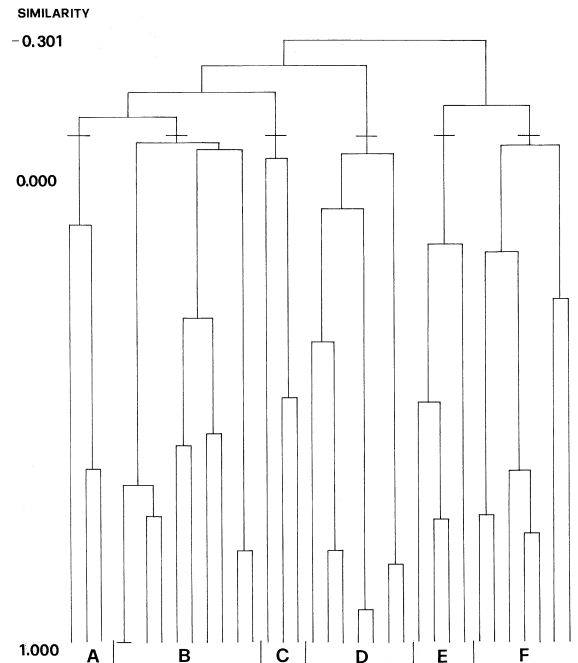


Fig. 11. Grouping of species using the dendrogram obtained by processing the species/samples matrix with the coefficient correlation and complete linkage method.



Table 2 (continued)

1.4						1.5				1.6					2								
18 50	28 50	26 50	32 50	22 50	33 50	12 50	13 50	13 100	14 50	14 100	18 100	48 0	48 50	48 100	20 0	20 50	19 100	20 100	22 100	12 100	33 100	26 100	
								204	1134	816													
									2079	1020		2068		288									
		324																					
		2																					
154	156	535	266	10	20		304	178		36		83	28	16	30	8	3						
		2																					
2337	1296	1620	2096	474	694	474	996		189	816		3619			779	1056							
												6											
						10	62	43	4	4	33	50	5										
		12	3				2	5			5	17	14	5	4	8							
							1660																
							106	18															
										4			2	16								2	
																							16
		5		8		8	28	3	4	2	5		9	23	19	29		3	5	5			2
				8	40	8					2			3		5	17	8	3	3	2		
3		14	3	18	29	18	2	3	4											5	5	7	5
9	3	7	2	6	20	6	2										6	5					
				3		3																	
	168	7	10	699	148	699	14	13	86	80	141	182	104	101	22	66	171	24	127	127	81	108	
6	15	17		34	3	34	2	20	28		31	6			24		31		154	154			
				3	3	3			2					8					16	16	7		
		648	262															156					
				237		237										352							
14		53		5	31	5																	
91	64	82	46	36	20	36	71	33	12	4	24	28	28	21	67	48	37		59	59	51	39	
34	8	17	35	3	3	3	32	15		2	2	6		5					3	3			
43	8	7	8				2																
29	10					6																	
145	6	22	127	3	6						5	704	5	3									
		2	12			3																	
3	3														4								
29	3		20			3					14					3						2	

their contribution both in terms of abundance and biomass is still high also at 50 m depth. Again this might be explained by a higher availability of appropriate food items at this depth.

On the basis of cluster analysis, each layer was separated from the others, indicating depth to be the most influential parameter in structuring community composition.

Stations 20 and 48 appeared different from all the others, as well as stations 13 and 14 at lower depths, due to the peculiarity of their populations.

Summing up, it appears that water mass characteristics and structures, as well as timing of sampling, strongly influenced the distribution of phytoplankton biomass and composition which mostly influence the

distribution of microzooplankton biomass. The microzooplankton community composition appears more related to patchiness of dominant species, which changed over the sampling time.

### Acknowledgements

This work was supported by the Italian National Antarctic Research Program. Thanks are due to crew members of R/V *Polar Stern* and especially to Prof. Letterio Guglielmo who collected the samples during the cruise. The authors wish to thank Dr. Mike Lucas for reviewing the manuscript and for his constructive comments, and the three anonymous reviewers for suggestions and critical remarks on the manuscript.

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