

Plankton community structure and carbon cycling in a coastal upwelling system.

II. Microheterotrophic pathway

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ABSTRACT: Planktonic food-web structure and carbon dynamics were studied in Mejillones Bay (23° S, off northern Chile) on 3 occasions: February, August and October 2001. Mejillones Bay was influenced by moderate upwelling events during February and October and presented a shallow (30 to 40 m depth) oxygen minimum layer (OML). On all sampling occasions, chain-forming diatoms that were grazed by small zooplankton and heterotrophic (h-)dinoflagellates comprised most of the autotrophic biomass. Heterotrophic (h-)nanoflagellates were largely bacterivorous, and responsible for a substantial removal of bacterial biomass, mostly associated with productive surface waters and the OML. Ciliate biomass was relatively low during all periods, but ciliates removed a large fraction of h-nanoflagellate production (12 to 22 % d⁻¹). Thus, in the microbial food web bacterial carbon can be transferred to ciliates and then subsequently to zooplankton. The impact of small zooplankton on primary production (PP) was relatively low on all sampling occasions. Small copepods and appendicularians removed from 0.6 to 5 %, and from 0.08 to 0.6 % PP d⁻¹ respectively. Appendicularians were the only zooplankton able to graze on bacteria, but grazing impact was not sufficient to regulate bacterial biomass. The microheterotrophic pathway could be an important link in this highly productive upwelling embayment. Our results showed that a large part of the photosynthetically fixed carbon was channeled through the microbial food web, with only a small part allocated to copepods and appendicularians. The food-web structure of Mejillones Bay can be classified as multivorous, with herbivorous and microbial grazing playing an important role in carbon export.

KEY WORDS: Microbial food web · Protozoans · Upwelling system · Copepods · Appendicularians · Carbon flow · Humboldt Current System

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INTRODUCTION

In the past, our understanding of coastal upwelling systems was strongly influenced by classical studies (e.g. Ryther 1969) which, when estimating fish production, assumed that there were few trophic levels and a high trophic efficiency in upwelling regions. Even now, the importance of the microbial food web to the food webs of fishes is commonly disregarded by fisheries scientists (e.g. Boudreau & Dickie 1992, Cury et al. 2000). Carbon fixed by primary producers is transferred along upwelling food webs through both 'microbial' and 'classical' pathways, and the proportion di-

rected along each route probably depends largely on the size of the primary producers (Walker & Peterson 1991, Mann 1992). There is increasing evidence that the microbial food web is a fundamental and almost permanent feature of not only oligotrophic but also eutrophic marine systems (Neuer & Cowles 1994, Tamigneaux et al. 1997). The significance of bacteria and microprotozoans as components of the biomass, and their role in biogeochemical cycles in upwelling systems have been widely recognized (Newel & Turley 1987, Painting et al. 1992, Ducklow et al. 2001). Previous studies in upwelling regions along the west coast of North America (Heinbockel 1978, Landry & Hassett

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1982, Landry & Lorenzen 1989); and off Peru (Beers et al. 1971, Sorokin 1978) have shown that protists can be an important component of the plankton community. However, trophic relationships within the microbial-loop food web in upwelling systems are poorly understood and greatly simplified (Painting et al. 1992). Because of the high production of upwelling regions and their importance for carbon fluxes from continental shelves to slopes (Walsh et al. 1981), more knowledge on the magnitude and variability of protist activity in upwelling regions is needed.

Like the rest of the northern part of the Humboldt Current System (HCS), the nearshore environment off northern Chile is influenced by intermittent upwelling events and it is also one of the most biologically productive areas within the HCS. Although new data on primary production in this region has been reported (Daneri et al. 2000), there is little data on the fate of this production in either the microheterotrophic or zooplankton pathways. Carbon fluxes through the pelagic food web off northern Chile are poorly known. In a multidisciplinary study in northern Chile off Antofagasta before and during the development of an El Niño event, González et al. (1998), suggested that zooplankton could remove from 4 to 6% of primary production (PP). However, most research in the HCS has only studied zooplankton herbivory as the main pathway by which carbon is exported from the food web. In addition, carbon flow dynamics, especially those referring to the microbial loop, is still lacking, and it is therefore not known how much of the PP channeled through the microbial and classical food webs is available to higher trophic levels, including large fish populations. While large copepods such as *Calanus* spp. and *Eucalanus* spp. represent a major fraction of the biomass in offshore waters (e.g. Escribano & Hidalgo 2000), the dominant small copepods (e.g. *Paracalanus parvus*, *Acartia tonsa* and *Oithona similis*) and appendicularians (*Oikopleura* spp.) are important components in nearshore waters and embayments along the HCS (Grünwald et al. 1998, González et al. 2000, Aravena & Palma 2002). Small calanoid and cyclopoid copepods and appendicularians are known to be largely omnivorous and bacterivorous respectively (e.g. Tiselius 1989, Dagg et al. 1996, Vargas et al. 2002, Zeldis et al. 2002, Vargas & González 2003, this issue). Therefore, when small zooplankton species feed on protozoans (which in turn feed effi-

ciently on pico- and nanoplankton), the energy transfer from the microbial loop (including bacteria and protists), and also from the microbial food web (including autotrophic pico- and nanoplankton) to higher trophic levels may be greater than previously assumed.

The main objectives of the present study were to quantify the total grazing impact on PP by measuring and describing the trophic activity of various components of the planktonic food web in surface waters, and to determine the percentage of PP that could be channeled by the microprotozoan and small zooplankton pathways in an embayment off northern Chile. We estimated potential carbon fluxes and analyzed the interaction between the microheterotrophic food web and higher trophic levels on 3 occasions representing periods of different productivity and zooplankton community structure in order to assess the trophic interactions of different plankton groups.

MATERIALS AND METHODS

The investigation was conducted along a transect of 4 stations representing bay (Stn 1), shelf (Stns 2 and 3) and slope (Stn 4) conditions in Mejillones Bay, northern Chile (23° S, 73° 20' W) (Fig. 1). Sampling was performed on board the RV 'Purihaalar' (Universidad de Antofagasta) on 3 occasions: 10 to 14 February, 1 to 5 August and 20 to 25 October 2001; on each occasion, the stations were sampled twice (i.e. 2 legs).

Hydrography. Temperature, salinity and oxygen profiles were recorded from near the bottom to the sur-

Table 1. Spatial-temporal oceanographic and biological data for 3 cruises: 10 to 14 February, 1 to 5 August and 20 to 25 October 2001. Each cruise sampled 4 stations, with 2 legs per cruise. N: no of measurements at each station

Measurement	Depths	N	Instruments
Temperature	Profiles	3 profiles	SeaBird SBE-19
Salinity			CTD with Beckman
Oxygen			oxygen sensor
Fluorescence			
Light extinction		~ 3	Secchi Disk
Chlorophyll <i>a</i>	1, 5, 10, 25, 50 and 100 m	2 profiles	30 l Niskin bottles
Bacterial biomass	1, 25, 50 m	3 samples per depth	30 l Niskin bottles
Protozoan biomass	1, 25, 50 m	3 samples per depth	30 l Niskin bottles
Phytoplankton biomass	1, 25, 50 m	3 samples per depth	30 l Niskin bottles
Zooplankton biomass	4 strata: 0–25; 25–50; 50–100 and 100–150 m	3 WP-2 plankton tows	WP-2 net
	Integrated	1 Bongo net tow	Bongo net

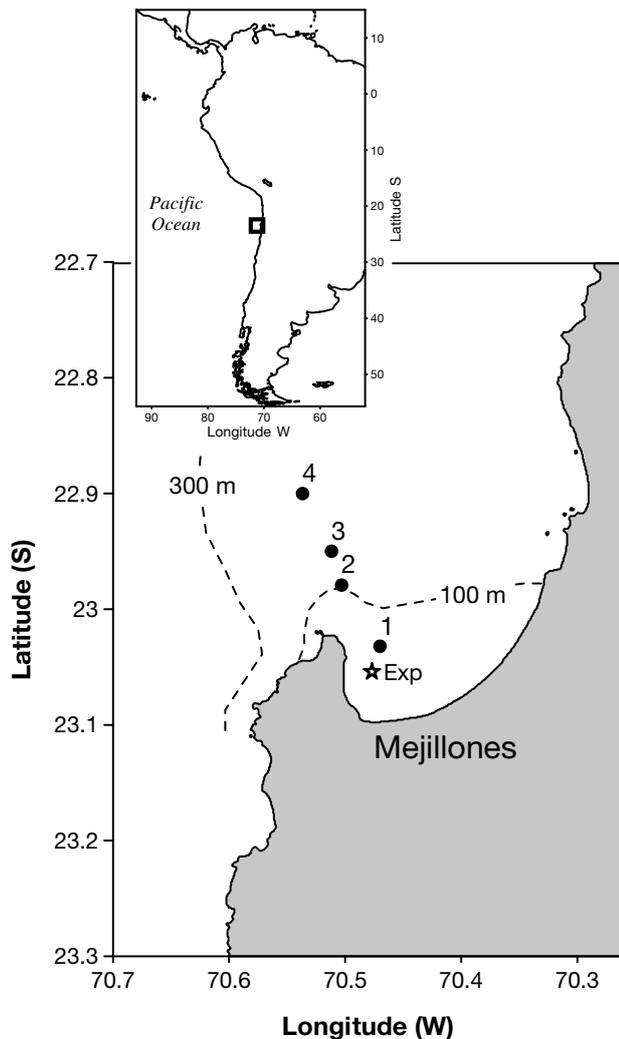


Fig. 1. Location of 4 sampling stations in Mejillones Bay. Iso-baths are included. Stn Exp: station used in study of Vargas & González (2003, this issue)

face using a SeaBird SBE-19 CTD equipped with an YSI-calibrated Beckman oxygen-sensor and a Westar fluorometer. Water samples for chlorophyll *a* (chl *a*), phytoplankton biomass, bacterial biomass, heterotrophic (h-) nanoflagellates, dinoflagellates and ciliates were collected at 1, 5, 10, 25, 50 and 100 m depths with 30 l Niskin bottles (Table 1).

Phytoplankton biomass and production. Chl *a* and phaeopigment concentration was measured after filtration of 500 ml seawater, followed by extraction for 24 h in 90 % acetone and fluorometric analysis (TD 700 Turner fluorometer) before and after acidification (Strickland & Parsons 1972). Water samples for phytoplankton and protozoan cell counts were also collected with 30 l Niskin bottles at 3 depths: 1, 25 and 50 m. Cells were counted and measured 24 h after settling by inverted microscopy at 200× magnification (Uter-

möhl 1958). Phytoplankton composition and biomass was determined by the methodology reported in Vargas & González (2003).

Size-fractionated primary production was measured in parallel at a coastal station (Stn Exp) using the method of Steemann Nielsen (1952). In the present study, PP data has been used only to construct a carbon budget. For details of the methodology see Iriarte et al. (2000).

Bacteria, microprotozoan biomass, production and grazing. Samples for the determination of bacteria, microprotozoan and phytoplankton biomass were collected from the same Niskin bottles, and 3 depths were also chosen: 1, 25 and 50 m. Bacteria, and autotrophic (a-) and heterotrophic (h-)nanoflagellates were quantified in 3 replicates from each sample by the acridine orange technique (Davis & Sieburth 1982, Hobbie et al. 1977) and analyzed using a color image-analysis system similar to that described by Verity & Sieracki (1993). Autofluorescence of a-nanoflagellates was revealed by blue light excitation. Biovolume was converted to carbon using the equation of Simon & Azam (1989) and Riemann & Bell (1990). Samples of ~100 ml unfiltered seawater were preserved in 2% acid Lugol solution and refrigerated until counts of microprotozoans (ciliates, a- and h-dinoflagellates) could be made in the laboratory. Microprotozoans were counted and measured, and their carbon content was estimated according to the methods of Vargas & González (2003). All naked dinoflagellates >20 μm were considered as heterotrophic (Hansen 1991, Levinsen et al. 1999).

We estimated protozoan (h-nanoflagellates, ciliates and h-dinoflagellates) ingestion with a model proposed by Peters (1994), since this model estimated of potential microprotozoan ingestion better than other models we tested previously (e.g. the temperature-dependent model of Vaqué et al. 1994, and the size-scaling model of Hansen et al. 1997), which did not consider important sources of variability such as prey and predator size. Peters' model predicts ingestion rates for a wide range of freshwater and marine ecosystems, and considers temperature, cell volumes, and concentrations of both prey and predator as explanatory variables:

$$\log GR = 2.701 - 0.344[\log Vol_{prey}] + 0.477[\log Vol_{predator}] + 0.489[\log AB_{prey}] - 0.270[\log AB_{predator}] + 0.033T$$

where GR = grazing rate (prey h^{-1}); Vol_{prey} = biovolume of preys (μm^3); $Vol_{predator}$ = biovolume of predators (μm^3); AB_{prey} = abundance of preys (ind. l^{-1}); $AB_{predator}$ = abundance of predators (ind. l^{-1}), and T = temperature ($^{\circ}C$).

Carbon demand ($mgC\ m^{-2}\ d^{-1}$) was estimated for microprotozoan ingestion on a- and h-nanoflagellate

and diatom prey, and integrated for the 0 to 25 m depth stratum. We attempted to predict predator–prey interactions on the basis that h-nanoflagellates feed mainly on bacteria and a-nanoflagellates, ciliates feed mainly on a- and h-nanoflagellates, and h-dinoflagellates feed mainly on diatoms. Since not all small choreotrich ciliates may feed efficiently on bacteria (Sherr et al. 1989, Kivi & Setälä 1995), we took only bacterivory into account for small aloricate ciliates. We could not estimate some feeding interactions, such as those of mixotrophic nanoflagellates, or of the abundant mixotrophic dinoflagellate *Ceratium furca* feeding on choreotrich ciliates (i.e. *Strobilidium* spp. and tintinnids; Smalley et al. 1999). A growth efficiency of 33% was used to calculate the production of protozooplankton (Hansen et al. 1997).

Zooplankton biomass and grazing. Vertical net hauls to estimate zooplankton abundance and biomass were done twice in 4 depth strata using a WP-2 net with 200 μm mesh size, closing devices, and a calibrated General Oceanic flowmeter. The depth strata were 0–25, 25–50, 50–100 and 100–150 m (Table 1). In order to correct estimation of euphausiid abundance for evasion, additional oblique net hauls were done with a Bongo net (0.6 m mouth size) equipped with 200 and 500 μm meshes and a calibrated flowmeter. All plankton samples were preserved immediately after collection with 5% formalin solution in seawater buffered with sodium borate. In the laboratory, subsamples (1/5 fraction) from the net-tows were counted, and the organisms identified and measured with a dissecting microscope. Biomass was calculated using length–weight regressions of Klein Breteler et al. (1982) (*Acartia* sp. and *Centropages* sp.), Hirche & Mumm (1992) (*Oithona similis*, *Metridia* spp.), Uye (1982) (other copepods), and Paffenhöfer (1976) and Gorsky et al. (1988) (*Oikopleura* spp.).

The grazing impact of zooplankton feeding on natural assemblages of protozoans (h-nano, dinoflagellates and ciliates), phytoplankton (diatoms, a-nanoflagellates and dinoflagellates) and bacteria were estimated using the ingestion rate data reported by Vargas & González (2003) for dominant small copepods and appendicularians at Stn Exp (Fig. 1). Total ingestion by small copepods and appendicularians was calculated as the sum of the integrated abundance (0–25 m depth) of each species multiplied by the species-specific ingestion rate (estimated from incubation experiments). Since ingestion rates were estimated for the most abundant small copepod species and appendicularians (typically around 80 to 95% of the total biomass of each group), this was assumed to represent the total grazing impact by the small copepod and appendicularian communities.

RESULTS

Hydrography

Vertical cross-sections showed that in February and October the water column in the study area was thermally stratified down to 20 to 25 m depth, with a gradient of 14 to 18°C (Fig. 2a, g), while during August the water column was more mixed (Fig. 2d). On all sampling occasions, salinity was relatively uniform throughout the whole water column (Fig. 2b, e, h). In February and October, the 14°C isotherm rose from ~40 to ~15 m depth in an offshore–inshore direction suggesting that despite the stratification, a moderate wind-driven upwelling might have developed in the area around Mejillones peninsula. In fact, the depth of the oxygen minimum layer (OML; defined here as $\leq 0.5 \text{ ml O}_2 \text{ l}^{-1}$) also rose nearshore, and during February and August was located at ~30 m depth (Fig. 2c, f). Conversely, in October the OML depth remained in deep waters at ~60 to 70 m depth (Fig. 2i).

Phytoplankton biomass

A dominant feature during the whole sampling period was the occurrence of the highest chl *a* concentration near the surface (upper 20 to 30 m) and always associated with Stn 1, inside Mejillones Bay (Fig. 2c, f, i). The highest chl *a* concentrations were found during the August sampling, with concentrations up to 35 mg m^{-3} in surface waters (Fig. 2f). At this time, high chl *a* concentrations (15 to 20 mg m^{-3}) were also observed along the transect. In October, surface chl *a* concentrations were the lowest of the entire study period (~2 mg m^{-3}) (Fig. 2i). Most of the chl *a* biomass was restricted to the well-oxygenated upper 25 m layer.

In general, diatoms dominated phytoplankton biomass at all stations and on all sampling occasions. However, there were marked differences in species composition between the different sampling occasions. In February, the dominant genera and species (>25% of the identified cells) were *Chaetoceros* spp., *Cylindrotheca* spp. and *Guinardia delicatula*. In the August sampling, the phytoplankton was dominated by a bloom of the chain-forming diatoms *G. delicatula* and *Detonula pumila*, with individual cells <30 μm . In October, diatoms were represented mostly by *Eucampia cornuta* and *Chaetoceros socialis*. Despite the small area covered by our study, there was a high spatial patchiness along the transect, with the contribution of diatoms to carbon biomass differing among close stations. The highest diatom biomass was observed outside of the bay, at Stn 2 in February and at Stn 3 in August (Fig. 3a, b). Varying C:chl ratios were evident

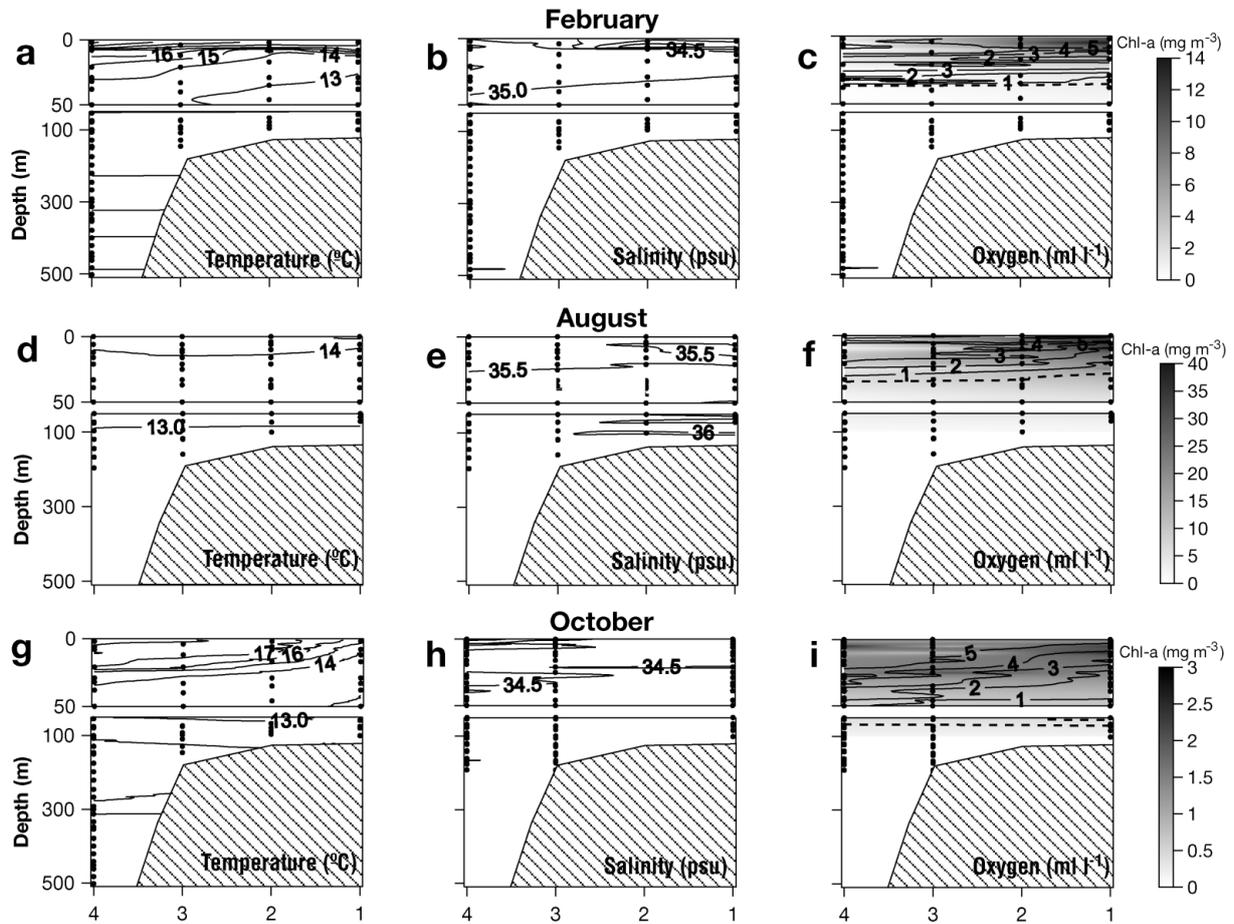


Fig. 2. Isopleths of (a, d, g) temperature, (b, e, h) salinity, (c, f, i) oxygen and chlorophyll *a* on (a, b, c) February, (d, e, f) August and (g, h, i) October 2001 sampling occasions. Dotted line in plots (c), (f) and (i) represents depth of oxygen minimum layer, defined here as $0.5 \text{ ml O}_2 \text{ l}^{-1}$

along the transect in August, as maximum chl *a* concentration did not match with highest cell biomass (Fig. 2f, 3b). A-nanoflagellates were more abundant inside Mejillones Bay (Stn 1). The contribution of a-nanoflagellates to the total biomass at this station was particularly significant during the February sampling (Fig. 3a), with a biomass even higher than that of diatoms, with a dominance of cells $>5 \mu\text{m}$. The biomass of a-dinoflagellates was represented mostly by large armored *Ceratium* species, and highest values were observed during the August sampling (Fig. 3b).

Bacteria and protozoan biomass, production and grazing

The depth-integrated bacterial biomass (i.e. 0 to 50 m depth) was on the order of 0.05 to 0.7 gC m^{-2} , and the highest biomass was, on average, observed during the August sampling (Figs. 3b & 4). Most bacterial bio-

mass was associated with productive surface waters (i.e. the upper 5 m layer) and the OML at 50 m depth (Fig. 4). H-nanoflagellate biomass ranged between 0.1 and 2.5 gC m^{-2} , with maximum values inside the bay (Stn 1) during February and August (Fig. 3a). Ciliates and h-dinoflagellate biomass was relatively low on all sampling occasions. The ciliates present consisted primarily of the large tintinnids *Helicostomella* spp., *Eutintinnus* spp. and *Tintinnopsis* spp. and some aloricate choreotrichs *Strombidium* spp. Although the ciliate biomass was low, it was significantly higher at the slope stations (Stns 2 and 3) than inside the bay (Stn 1) (*t*-test; 1-tailed; $p = 0.007$) (Fig. 3, see also Tables 3 & 4). Thecate forms dominated h-dinoflagellates $>20 \mu\text{m}$, with *Protoperdinium* spp. and the myxotroph *Prorocentrum micans* constituting the dominant genera. Maximum biomass of ciliates and h-dino flagellates was observed in February at Stn 2 ($\sim 1600 \text{ mgC m}^{-2}$). Most of the protozoan biomass inside Mejillones Bay was restricted to surface waters, with values between

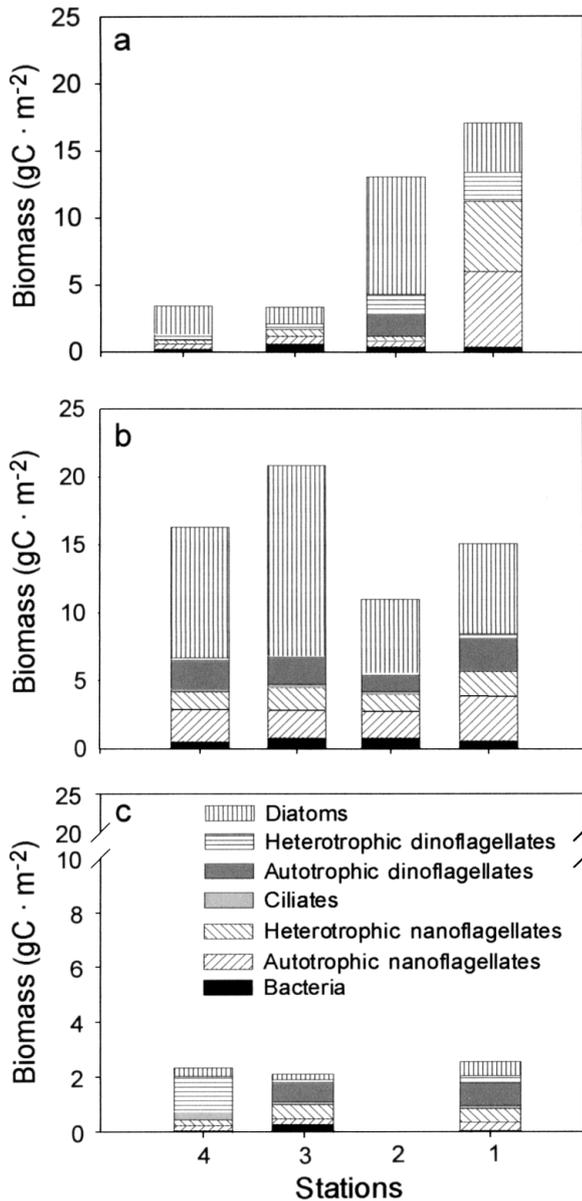


Fig. 3. Mean contribution of major taxonomic groups to integrated (0 to 50 m depth) biomass (gC m^{-2}) of autotrophic and heterotrophic community along transect on (a) February, (b) August and (c) October 2001 sampling occasions (average of Legs 1 and 2). Note different y-axis in October plot

10 and 150 mgC m^{-3} (t -test; 1-tailed; $p = 0.0008$). The highest biomass was observed during the February sampling (Fig. 4).

Estimates of grazing by protozoans showed that h-nanoflagellates may ingest from 12 to 76 bacteria $\text{ind.}^{-1} \text{ h}^{-1}$, similarly to many reports for different marine ecosystems (Table 2). We presumed that only small aloricate ciliates were feeding on bacteria, and values ranged between ~ 270 and 660 bacteria $\text{ciliate}^{-1} \text{ h}^{-1}$. Ciliate ingestion of a- and h-nanoflagellates was rela-

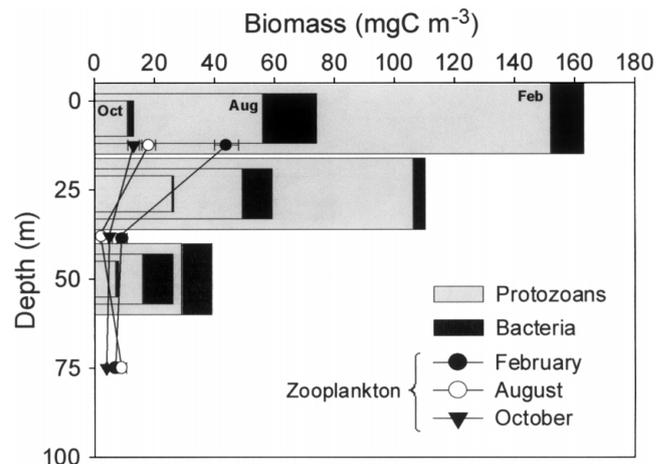


Fig. 4. Vertical distribution of bacteria, protozoan and zooplankton biomass (mgC m^{-3}) at Stn 1, inside Mejillones Bay, on the 3 sampling occasions. Differences in bar width distinguish sampling period. Errors bars = SD

tively low (~ 1 to 20 cells $\text{ciliate}^{-1} \text{ h}^{-1}$, Table 2). Ingestion of diatoms by h-dinoflagellates was ~ 0.5 cells $\text{ind.}^{-1} \text{ h}^{-1}$, which also fits well within the range of values reported in the literature (Table 2). Production of protozooplankton was closely associated with biomass level, with a high spatial heterogeneity for each protozoan group. However, maximum total protozooplankton production mainly occurred nearshore at Stns 1 and 2 (Tables 3, 4 & 5). H-dinoflagellates played a major role in phytoplankton consumption, with the highest carbon demand by protozooplankton (ANOVA, $p < 0.01$). The highest total grazing impact by protozooplankton was observed inside the bay, except in October, when grazing was slightly higher at Stn 4 than at Stn 1 (t -test: $p < 0.05$) (Tables 3, 4, & 5). Inside Mejillones Bay, protozooplankton ingested between 350 and $2500 \text{ mgC m}^{-2} \text{ d}^{-1}$.

Zooplankton biomass and grazing

Copepods were the dominant components of the zooplankton biomass in the study area. A total of 13 genera or species were identified in the samples. Small calanoid and cyclopoid copepods were the dominant group comprising 64, 48 and 72% of the total biomass inside Mejillones Bay in February, August and October respectively. *Paracalanus parvus*, *Acartia tonsa*, *Centropages brachiatus* and *Oithona similis* were the dominant species in this group, especially inside Mejillones Bay (Stn 1) (Table 6). The contribution of large copepods (e.g. *Calanus chilensis* and *Eucalanus* sp.) was slightly higher than that of small copepods only during

February and October at Stn 4 (ANOVA; $p < 0.05$) (Table 6). The distribution of copepods reflected the increase in phytoplankton biomass rather than chl *a* concentration, with the highest biomass at Stns 1 and 3 in February, Stns 2 and 3 in August and Stn 1 in October. Appendicularians, mostly dominated by the species

Oikopleura dioica, constituted around 5% of the total biomass inside the bay during February (64.4 mgC m^{-2}). During August, another coastal appendicularian species, *O. longicauda*, constituted around 7% of the total biomass inside the bay (Table 6: 'Appendicularians'). This species was also very abundant on the shelf

Table 2. Feeding data for flagellates, ciliates and dinoflagellates from this study, 2001 (Peters' 1944 model) and from the literature. HNF: heterotrophic nanoflagellates; ANF: autotrophic nanoflagellates

Predators		Prey		
Species	Ingestion (cell $pred^{-1} h^{-1}$)	Species	Nos. ($\times 10^6 ml^{-1}$)	Source
Flagellates				
Mixed HNF > 5 μm	24–64	Mixed bacteria	0.1–0.4	This study, February (16°C)
	28–63	Mixed bacteria	0.5–1	This study, August (14°C)
	12–76	Mixed bacteria	0.3–1	This study, October (15°C)
Mixed HNF	54	Mixed bacteria	0.2–1.4	Cuevas (1999) (9.5–13°C)
<i>Monosiga</i> sp.	27	<i>Pseudomonas</i> spp.	5–35	Fenchel (1982): batch cultures at 20°C
<i>Pseudobodo tremulans</i>	84	<i>Pseudomonas</i> spp.	5–50	
<i>Monas</i> spp.	10–75	Mixed bacteria	25–8000	Sherr et al. (1983)
<i>Pseudobodo tremularis</i>		Mixed bacteria	2.4–4.8*	Andersen & Fenchel (1985) ^a
<i>Pseudobodo</i> sp.	45–73	Mixed bacteria		
Mixed HNF	5–28	Fluorescently labeled bacteria	2.5–9.2	Sherr et al. (1988): 12–20°C
<i>Spurnella</i> sp.	12–48	<i>Synechococcus</i> spp.	10–12	Boenigk et al. (2001): 18–20°C
<i>Cafeteria</i> sp.	2–14	<i>Synechococcus</i> spp.	10–12	
($\times 10^3 ml^{-1}$)				
Ciliates				
Mixed ciliates	305–592	Mixed bacteria	0.1–0.4	This study, February
	479–647	Mixed bacteria	0.5–1	This study, August
	270–663	Mixed bacteria	0.3–1	This study, October
	6–15	Mixed ANF	0.4–4	This study, February
	6–21	Mixed ANF	0.7–5	This study, August
	1–11	Mixed ANF	0.01–0.4	This study, October
	13	Mixed HNF	0.9–10	This study, February
	11	Mixed HNF	0.7–2	This study, August
	6	Mixed HNF	0.2–0.6	This study, October
	<i>Strobilidium hexakinetum</i>	110–650	Mixed bacteria	
<i>Halteria grandinella</i>	470–2690	Mixed bacteria		
<i>Strombidium sulcatum</i>	900–926	<i>Prochlorococcus</i> + <i>Synechococcus</i> spp.	500	Christaki et al. (1999)
<i>Tintinnopsis acuminata</i>	29–80	<i>Isochrysis</i> spp.		Verity (1985)
<i>Tintinnopsis vasculum</i>	34–83	<i>Dicrateria</i> spp.		
<i>Uronema marina</i>	710	Fluorescently labeled bacteria	14400	Sherr et al. (1988): 19.5–22°C
Mixed spirotrichs	380		1500	
<i>Strombidium vestitum</i>	~17–25	<i>Isochrysis galbana</i>		Tang et al. (2001)
<i>Rimostrombidium conicum</i>	~20–75	or <i>Phaeocystis globosa</i>		
(ml^{-1})				
Dinoflagellates				
Mixed H-dinoflagellates (mostly <i>Protoperidinium</i>)	0.4	Mixed diatoms	5–25	This study, February
	0.4	Mixed diatoms	1–5	This study, August
	0.5	Mixed diatoms	0.9–3	This study, October
<i>Protoperidinium</i>	0.08–0.2	Red tide dinoflagellates		Jeong & Latz (1994)
<i>Gyrodinium galatheanum</i>	0–0.01	Mixed cryptophytes		Li et al. (2001)

^aVery low bacterial biomass at 15°C

Table 3. Integrated distribution of biomass, production and ingestion of auto- and heterotrophs (H-) across transect in Mejillones Bay area in February 2001. Data integrated for oxygenated upper 25 m of water column. PP: primary production

Species	Station			
	1	2	3	4
Biomass (mgC m⁻²)				
Phytoplankton	6376	5671	846	351
Bacteria	202	146	230	54
H-nanoflagellates	2540	246	242	118
Ciliates	15	1660	5	5
H-dinoflagellates	23	1658	0	0
Small copepods	768	427	704	64
Medium-large copepods	217	171	675	198
Appendicularians	58	6	32	3
Euphausiids	33	0	0	0
Production (mgC m⁻² d⁻¹)				
Phytoplankton	5566 ^a			
H-nanoflagellates	6764	654	645	314
Ciliates	21	2366	7	7
H-dinoflagellates	24	1703	0	0
Ingestion (mgC m⁻² d⁻¹)				
H-nanoflagellates	401	172	146	76
Ciliates	753	281	223	229
H-dinoflagellates	1351	900	0	0
<i>Total</i>	<i>2505</i>	<i>1353</i>	<i>369</i>	<i>305</i>
% PP ⇒ protozooplankton	45.0			
Small copepods	108	80	151	21
Appendicularians	34	3	14	1
Euphausiids	2	0	0	0
<i>Total</i>	<i>144</i>	<i>83</i>	<i>165</i>	<i>22</i>
% PP ⇒ zooplankton	2.6			

^aUnpubl. data of J. Iriarte & H. González

(Stn 3), with an integrated biomass of 87 mgC m⁻² in the upper 25 m depth (12% of the total biomass) (Table 4). Euphausiids were infrequently collected in samples taken with both the WP-2 and Bongo nets, and very few specimens were caught in February. Therefore we did not consider this group as significant grazers during our study (Table 4). The zooplankton community was mainly present in the well-oxygenated upper 25 m, where biomass ranged between 10 and 45 mgC m⁻³. Maximum biomass was observed during February, and decreased with depth below the surface (Fig. 4). Small copepods accounted for the highest carbon consumption on autotrophs (*t*-test: *p* = 0.005). Inside Mejillones Bay, appendicularians ingested only between 25 and 34 mgC m⁻² d⁻¹ (Tables 3 & 4) and the maximum carbon ingestion by appendicularians (~72 mgC m⁻² d⁻¹) was observed in August at Shelf Stn 3. Total zooplankton carbon consumption was higher at Stn 3 during February and August, and at Stn 1 during October. Inside Mejillones Bay, the total zooplankton ingested between 63 and 144 mgC m⁻² d⁻¹ in the upper 25 m depth, with maximum ingestion in February.

Table 4. Integrated distribution of biomass, production and ingestion of auto- and heterotrophs (H-) across transect in Mejillones Bay area in August 2001. Data integrated for oxygenated upper 25 m of water column. PP: primary production

Species	Station			
	1	2	3	4
Biomass (mgC m⁻²)				
Phytoplankton	9250	5637	17235	8680
Bacteria	359	423	615	316
H-nanoflagellates	1113	537	945	567
Ciliates	77	89	118	132
H-dinoflagellates	144	88	63	121
Small copepods	202	544	440	180
Medium-large copepods	120	88	131	20
Appendicularians	30	62	87	0
Euphausiids	0	0	65	17
Production (mgC m⁻² d⁻¹)				
Phytoplankton	9415 ^a			
H-nanoflagellates	2964	1429	2517	1509
Ciliates	109	127	169	188
H-dinoflagellates	147	99	65	124
Ingestion (mgC m⁻² d⁻¹)				
H-nanoflagellates	39	18	28	11
Ciliates	1013	853	655	741
H-dinoflagellates	1360	1335	0	559
<i>Total</i>	<i>2412</i>	<i>2206</i>	<i>683</i>	<i>1311</i>
% PP ⇒ protozooplankton	26			
Small copepods	61	103	120	42
Appendicularians	25	51	72	0
Euphausiids	0	0	4	1
<i>Total</i>	<i>86</i>	<i>154</i>	<i>196</i>	<i>43</i>
% PP ⇒ zooplankton	1			

^aUnpubl. data of J. Iriarte & H. González

DISCUSSION

Hydrography and productivity

The physical environment of the Mejillones Bay area has been characterized in previous studies (Navea & Miranda 1980, Rodríguez et al. 1991, Marín et al. 2001, Sobarzo & Figueroa 2001), in which the development of quasi-permanent upwelling events along the Mejillones Peninsula was described. During the period of our study, the temperature and oxygen distribution suggested the occurrence of upwelling events in February and October. In these periods the OML penetrated the upper 30 m depth layer inside the bay. Previous studies (Morales et al. 1996, Escribano 1998) indicated that low-oxygen waters may reach 40 to 60 m depth in this area. Furthermore, October corresponds to the time of maximum upwelling (Marín & Olivares 1999). Such periods of low oxygen are typically characterized by high nitrate concentrations (Rodríguez et al. 1991), and thus may have enhanced primary production in the area. In

Table 5. Integrated distribution of biomass, production and ingestion of auto- and heterotrophs (H-) across transect in Mejillones Bay area in October 2001. Data integrated for oxygenated upper 25 m of water column. PP: primary production

Species	Station		
	1	3	4
Biomass (mg C m⁻²)			
Phytoplankton	752	522	1514
Bacteria	33	98	31
H-nanoflagellates	266	294	102
Ciliates	35	43	198
H-dinoflagellates	132	74	5
Small copepods	237	165	118
Medium-large copepods	23	43	23
Appendicularians	2	2	1
Euphausiids	63	0	0
Production (mg C m⁻² d⁻¹)			
Phytoplankton	1124*		
H-nanoflagellates	708	784	272
Ciliates	50	62	283
H-dinoflagellates	136	76	5
Ingestion (mg C m⁻² d⁻¹)			
H-nanoflagellates	116	187	84
Ciliates	76	92	317
H-dinoflagellates	157	77	76
Total	349	356	477
% PP ⇒ protozooplankton	31		
Small copepods	57	35	26
Appendicularians	1	1	0.5
Euphausiids	5	0	0
Total	63	36	26.5
% PP ⇒ zooplankton	6		

*Unpubl. data of J. Iriarte & H. González

fact, phytoplankton biomass was very high during our study, in agreement with data from other studies (e.g. Marín & Olivares 1999). The biomass was dominated by chain-forming diatoms, with maximum values inside Mejillones Bay and the shelf area (Stn 3).

Primary production and microphytoplankton biomass are usually high in the euphotic zone of the study area (Marín & Olivares 1999). The depth of the euphotic zone inside Mejillones Bay is around 10 to 40 m (Rodríguez et al. 1991). During our study, the mean euphotic layer (measured by Secchi disk, Table 1) occurred at around 20 to 25 m depth. Integrated PP in the euphotic zone (J. Iriarte & H. González unpubl.), was 5.5, 9.4, and 1.1 g C m⁻² d⁻¹, during February, August and October respectively, values that are among the highest reported for this area, and microphytoplankton >23 µm accounted for >50% of the PP over the whole study period.

Table 6. Integrated biomass (mg C m⁻²) of zooplankton in upper 50 m depth layer in coastal and oceanic stations during different sampling periods

Taxon	February	August	October	Mean (SE)
Coastal stations				
Small-medium copepods				
<i>Paracalanus parvus</i>	625.7	134.9	288.1	349.6 (251.1)
<i>Acrocalanus</i> sp.	0.9	1.5	0	08 (0.7)
<i>Acartia tonsa</i>	131.4	52.5	1.2	61.7 (65.6)
<i>Centropages brachiatus</i>	48.3	12.1	17	25.8 (19.6)
<i>Temora longicornis</i>	0.5	0	0	0.2 (0.3)
<i>Oithona similis</i>	2.1	4.3	4.8	3.7 (1.4)
<i>Oncaea</i> sp.	29.8	7.4	9.3	15.5 (12.4)
<i>Corycaeus anglicus</i>	2.4	0.3	9.9	4.2 (5.1)
<i>Clytemnestra</i> sp.	0.3	0	0	0.1 (0.1)
Copepodids (>500 µm)	0.6	0.7	0	0.4 (0.4)
Total	841.9	213.7	330.3	462 (356.8)
Large copepods				
<i>Calanus chilensis</i>	56.4	94.1	28.2	72.9 (54.8)
<i>Eucalanus</i> sp.	274.6	45.4	6.1	112.0 (143.0)
<i>Candacia</i> sp.	0	20.4	0	6.8 (11.8)
<i>Metridia longa</i>	8.7	1.4	0.3	3.5 (4.5)
Total	339.8	161.3	34.6	195.2 (214.1)
Appendicularians				
Total	64.4	32.6	7	34.7 (28.8)
Euphausiids				
Adults	43.4	0	0	14.5 (25.1)
Juvenile	0	0	63.1	21.0 (36.4)
Calyptopis	13.6	32.7	1.5	16.0 (15.7)
Furcilia	6.8	4.5	21.7	11.0 (9.3)
Total	63.9	37.2	86.3	62.5 (86.6)
Oceanic stations				
Small-medium copepods				
<i>Paracalanus parvus</i>	80.1	125.3	99.5	101.0 (22.7)
<i>Acrocalanus</i> sp.	1.1	0	0.4	0.5 (0.6)
<i>Acartia tonsa</i>	3.5	19.8	1.4	8.2 (10.1)
<i>Centropages brachiatus</i>	0	7.3	2.0	3.1 (3.8)
<i>Temora longicornis</i>	0.8	0	0.7	0.5 (0.4)
<i>Oithona similis</i>	2.7	1.4	1.9	2.0 (0.7)
<i>Oncaea</i> sp.	9.7	1.0	11.6	7.4 (5.6)
<i>Corycaeus anglicus</i>	3.3	0.5	4.6	2.8 (2.1)
<i>Clytemnestra</i> sp.	0	0	0	0
Copepodids (>500 µm)	0.3	0.1	0	0.2 (0.2)
Total	101.5	155.4	122.1	126.4 (46.1)
Large copepods				
<i>Calanus chilensis</i>	11.3	49.2	165.7	75.4 (80.5)
<i>Eucalanus</i> sp.	401.4	51.4	69.9	174.2 (197)
<i>Candacia</i> sp.	10	1.6	12.6	8.1 (5.7)
<i>Metridia longa</i>	2.8	0	0	0.9 (1.6)
Total	425.6	102.2	248.2	258.6 (284.8)
Appendicularians				
Total	6.4	0.3	2.4	3.1 (3.1)
Euphausiids				
Adults	0	0	0	0 (0)
Juvenile	0	0	0	0 (0)
Calyptopis	19.2	0	1.7	7 (10.6)
Furcilia	0	0	56.5	18.8 (32.6)
Total	19.2	0	58.1	25.8 (43.2)

Quantitative significance of microheterotrophs in a coastal upwelling area

The abundance and biomass of different protozoan groups have not been previously described for upwelling areas along the Humboldt Current System. Our estimates of protozooplankton carbon biomass are consistent with values in the literature. In the upper 25 m depth layer, the biomass of h-dinoflagellates ranged between 1 and 6 mgC m⁻³ (i.e. 23 to 144 mgC m⁻²) inside Mejillones Bay, with a peak of 66 mgC m⁻³ at Stn 2 during February. The biomasses of h-dinoflagellates generally range between 0.1 and 50 mgC m⁻³ (Lessard 1991). For comparison, in the Benguela upwelling system, dinoflagellate biomass ranged between 7 and 16 mgC m⁻³ in Elands Bay (Pitcher 1986) and from 5 to 49 mgC m⁻³ in St Helena Bay (Pitcher 1988). Conversely, the contribution of ciliates to carbon biomass was low during our study, ranging between 0.6 and 3 mgC m⁻³ (15 to 77 mgC m⁻²) inside Mejillones Bay (upper 25 m). In general, ciliate biomass is relatively low in upwelling areas. In the Benguela system, Painting et al. (1992) reported a mean biomass of around 7 mgC m⁻³. Ciliates and h-dinoflagellates biomass also compares well with that in the Oregon upwelling system, which ranged from 0.2 to 9 mgC m⁻³ for choreotrich ciliates and from 0.1 to 16 mgC m⁻³ for thecate dinoflagellates (Neuer & Cowles 1994). Sorokin (1978) found protozoan biomass (ciliates and flagellates) to be between 20 and 50 mgC m⁻³ in surface waters of the Peruvian upwelling region. These values are also well within the range found for total protozoan biomass in our study (10 to 150 mgC m⁻³).

Earlier studies by Goldman & Caron (1985), Parslow et al. (1986) and Sherr et al. (1986) indicated that many microprotozoan organisms cannot be allocated into exclusively autotrophic, heterotrophic or bacterivorous compartments. We attempted to estimate predator-prey interactions, taking into account the main trophic interactions reported in the literature. Potential ingestion rates estimated using Peters' (1994) model were in the range of those reported in different studies in coastal and oceanic areas. Fenchel (1982) demonstrated that flagellates could be major consumers of bacteria, and described them as the main grazers in the pelagic ecosystem. Since ingestion rates reported in the literature for marine and freshwater systems typically range between 10 and 75 bacteria h-nanoflagellate⁻¹ h⁻¹ (e.g. Fenchel 1982, Andersen & Fenchel 1985, Sherr et al. 1988, Boenigk et al. 2001), we believe that our values are a realistic approximation for grazing on bacteria. In fact, in the upwelling area off Central Chile (36° S), Cuevas (1999) observed a mean ingestion rate for h-nanoflagellates (h-nano) of ~54 bacteria h-nano⁻¹ h⁻¹.

Trophic relationships between ciliates and phytoplankton are more unusual, since previous studies have shown that naked oligotrich and choerotrich ciliates feed almost exclusively on nanoplankton (Rassoulzadegan et al. 1988, Bernard & Rassoulzadegan 1990). The estimated ingestion rates in our study were similar to those reported in the literature (Verity 1985, Sherr et al. 1988, Šimek et al. 1996, Christaki et al. 1999, Tang et al. 2001). Occasionally, small aloricate ciliates may consume bacteria at rates similar to flagellates, and display a greater preference for larger bacteria than flagellates (Epstein & Shiaris 1992). In our study, estimated ciliate ingestion rates were ~270 to 660 bacteria ciliate⁻¹ h⁻¹, which fit very well with results reported in the literature for small aloricate ciliates (Table 2). In experiments with fluorescent labeled bacteria, Sherr et al. (1988) reported high ingestion rates of ciliates on bacteria, whereby *Uronema marina* ingested ~710 bacteria ciliate⁻¹ h⁻¹ and *Strombidium sulcatum* ingested 1095 bacteria ciliate⁻¹ h⁻¹ at 22 and 19.5°C respectively.

Marine heterotrophic thecate dinoflagellates feed with a pseudopod or 'pallium'. This structure is highly plastic, and easily stretches to accommodate spines and most large diatoms chains (Gaines & Taylor 1984, Jacobson & Anderson 1986). In our study, we considered only trophic interactions between h-dinoflagellates and diatoms, a decision based on the fact that *Protoperidinium* was the most abundant h-dinoflagellates genus during our study: with few exceptions, *Protoperidinium* species have been observed to feed only on diatoms (Jacobson & Anderson 1986), and small flagellates are generally not able to support *Protoperidinium* spp. growth (Naustvoll 2000). Although h-dinoflagellates were not a dominant constituent of biomass, they were important herbivores with important impacts on carbon fluxes.

Planktonic production and carbon flow

Carbon flux in the ocean depends mainly on the magnitude of primary production and the biogeochemical processes within the euphotic zone, as well as on the complexity of the pelagic food web. Since the photic layer during our study ranged between 20 and 25 m depth, biomass, production and ingestion rate measurements were integrated between 0 and 25 m to construct a carbon budget for each sampling period (Figs. 5, 6 & 7). Although the hydrography of the sampling area is highly dynamic, and advection could greatly influence our results, the southern boundary of Mejillones Bay (Stn 1: Fig. 1) has been characterized as a region of reduced upwelling (i.e. of 'upwelling shadow' sensu Graham & Largier 1997), where oceanographic conditions generate an internal eddy and more stable conditions than outside

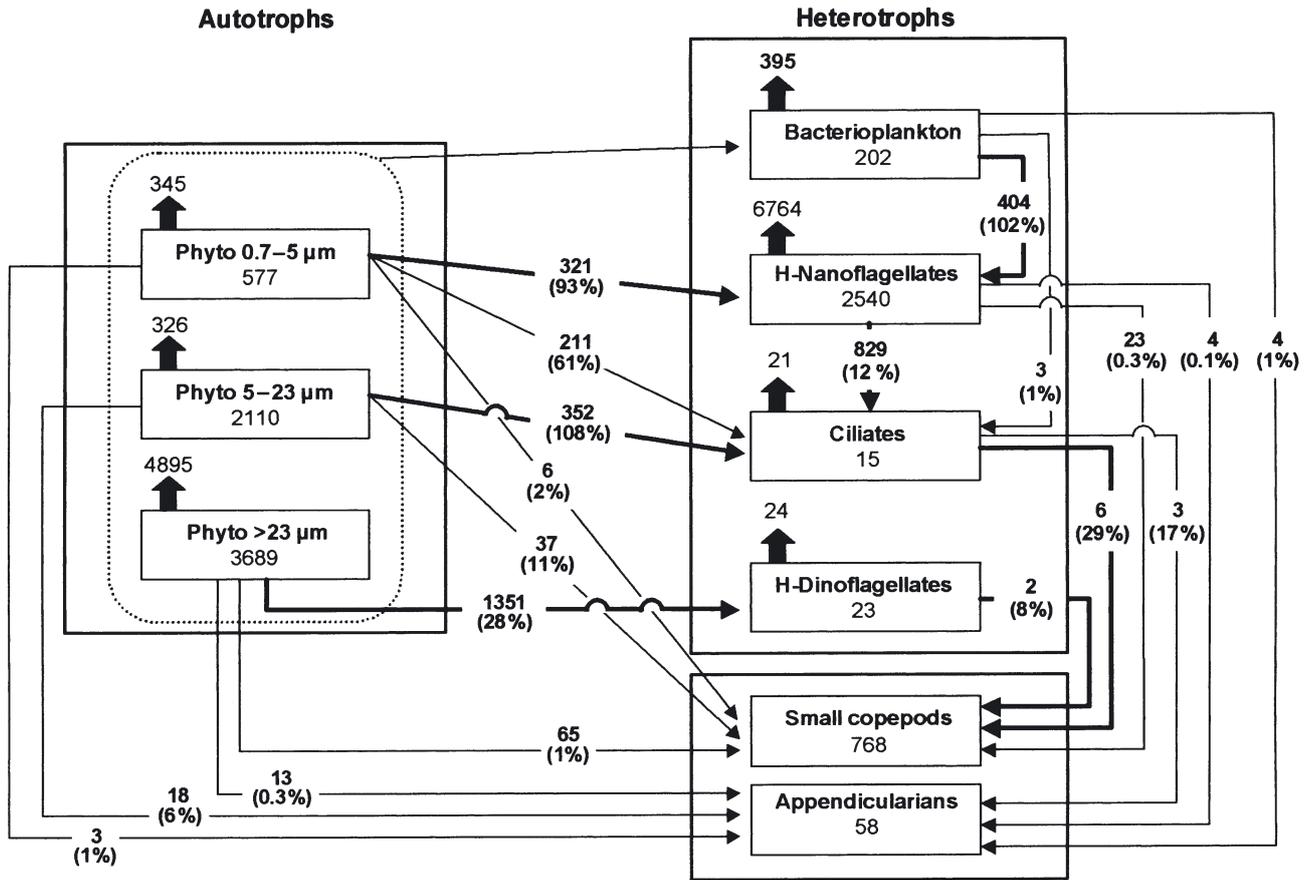


Fig. 5. Carbon budget in February 2001. Nos. inside boxes: biomass (mg C m^{-2}); nos. above black arrows: production ($\text{mg C m}^{-2} \text{d}^{-1}$); nos. on thin continuous arrows: consumption ($\text{mg C m}^{-2} \text{d}^{-1}$). Values in parentheses: % of production ingested. Biomass, production and consumption values integrated between 0 and 25 m depths. Continuous lines represent main fluxes between boxes

the bay (Marín et al. 2001). Therefore, a carbon budget was constructed for Stn 1, inside Mejillones Bay, where J. Iriarte & H. González (unpubl.) also measured size-fractionated PP. During our study, we did not measure bacterial production (BP). However, in a coastal station close to Mejillones Bay, Iriarte et al. (2000) measured BP during the summer, recording values of $\sim 15.8 \mu\text{g C l}^{-1} \text{d}^{-1}$ which, integrated over the upper 25 m depth layer, would be $\sim 395 \text{ mg C m}^{-2} \text{d}^{-1}$ (except during October when extremely low bacterial biomass was recorded). We considered that our results could have been substantially influenced by PP variability, since BP is closely related to PP (Kirchman et al. 1995). BP was assumed to derive primarily from photosynthetically produced dissolved organic carbon (DOC) and 'sloppy feeding' of zooplankton, and to represent only $\sim 8\%$ of total PP (Figs. 5 & 6). This is supported by the fact that Mejillones Bay lacks a freshwater contribution from rivers, and is a highly productive system (Rodríguez et al. 1991, Marín & Olivares 1999). In addition, the diatom species found during our study typically showed the highest percent-

age of extracellular release of DOC (e.g. *Chaetoceros* spp.: Nagata 2000).

Calculation of grazing pressure showed that h-nanoflagellates were capable of removing a high percentage of BP during February and August (102 and 55% BP d^{-1} , respectively; Figs. 5 & 6). H-nanoflagellates were also able to daily remove between 69 and 93% of PP from small cells (0.7 to 5 μm), and had the highest impact on this fraction. The grazing impact of ciliates on small autotrophs was important, particularly in February, when ciliates removed 108% d^{-1} of a-nanoflagellates PP (5 to 23 μm ; Fig. 5). Ciliates were also the most important grazers of h-nanoflagellates, and were possibly able to remove between 12 and 22% of the daily h-nanoflagellate production. Since only small aloricate ciliates (ca. 25 to 30% of total ciliates in this study) can exploit bacteria (Sherr et al. 1989), an insignificant percentage of the BP (0.2 to 2% d^{-1}) was ingested by total ciliates. These results are in agreement with those of other studies, which have shown that ciliates strongly influenced the nanoflagel-

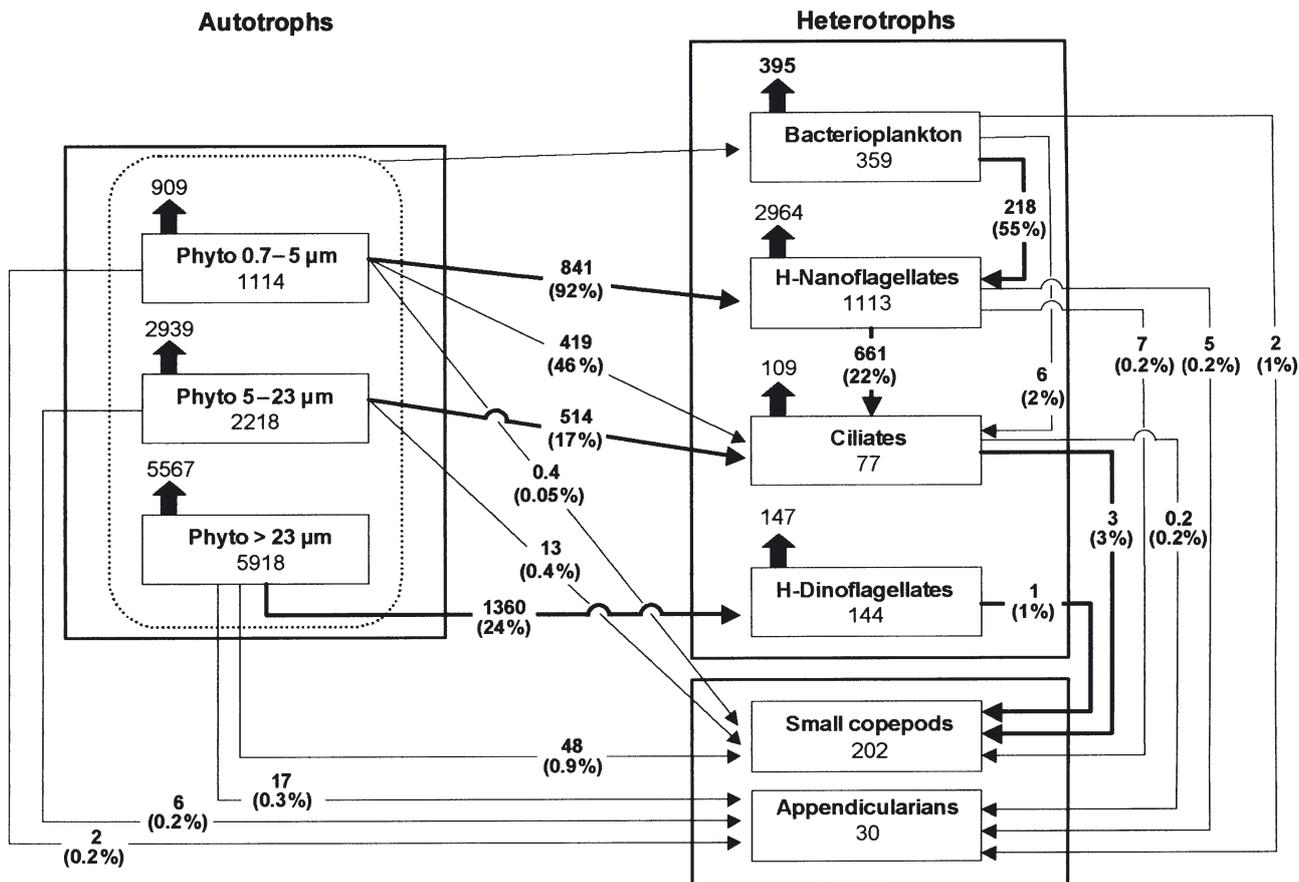


Fig. 6. Carbon budget in August 2001. Further details as in legend to Fig. 5

late community, even when present in low abundance (Epstein & Shiaris 1992).

Despite their low abundance in this coastal upelling system, h-dinoflagellates had a major predation impact on large cells (>23 μm), since they may remove between 18 and 28% of PP of this fraction. The estimates of grazing impact on diatom biomass and production illustrate that h-dinoflagellates play an important role in controlling the autotrophic production of cells >23 μm. Thus, ciliates may compete with h-nanoflagellates for bacteria and small flagellates <5 μm, while the larger h-dinoflagellates may compete with copepods for large diatoms (Nielsen & Hansen 1995). The impact of small zooplankton on PP was relatively low during the whole sampling period. Small copepods removed between 0.6 and 5% PP d⁻¹, with the highest impact in October when PP was relatively low. Appendicularians (mostly *Oikopleura dioica*) removed 0.08 to 0.6% PP d⁻¹, and both copepods and appendicularians had a major grazing impact on the nanoplankton fraction. However, the impact of appendicularians could have been more significant offshore (Stn 3) in August, when they constituted 12% of zooplankton biomass (Table 4).

González et al. (1998) constructed a conceptual model of carbon flow from estimated PP. In the coastal area off Peninsula Mejillones, approximately 12% of the total PP was consumed by mesozooplankton. However, it should be noted that in the study by González et al. (1998), total average PP ranged between 2 and 3 g C m⁻² d⁻¹, a much lower range than the 1 to 9 g C m⁻² d⁻¹ PP recorded during the present study in February and August by J. Iriarte & H. González (unpubl.). In fact, population grazing-impact in the present study was similar to that recorded by González et al. (1998, 2000). The population grazing impact measured by González et al. (1998, 2000) was around 250 to 300 mg C m⁻² d⁻¹, while during our study it was between 60 and 150 mg C m⁻² d⁻¹. However, we did not consider grazing by large copepods (*Calanus chilensis* and *Eucalanus* sp.). Our biomass data showed that large copepods were not abundant, and salps were not found in Mejillones Bay; therefore, most grazing in our study was by small copepods and appendicularians, and large copepods could have removed only a small percentage of the PP in the >23 μm fraction. Abundance of both small cycloids and appendicularians may also have been somewhat

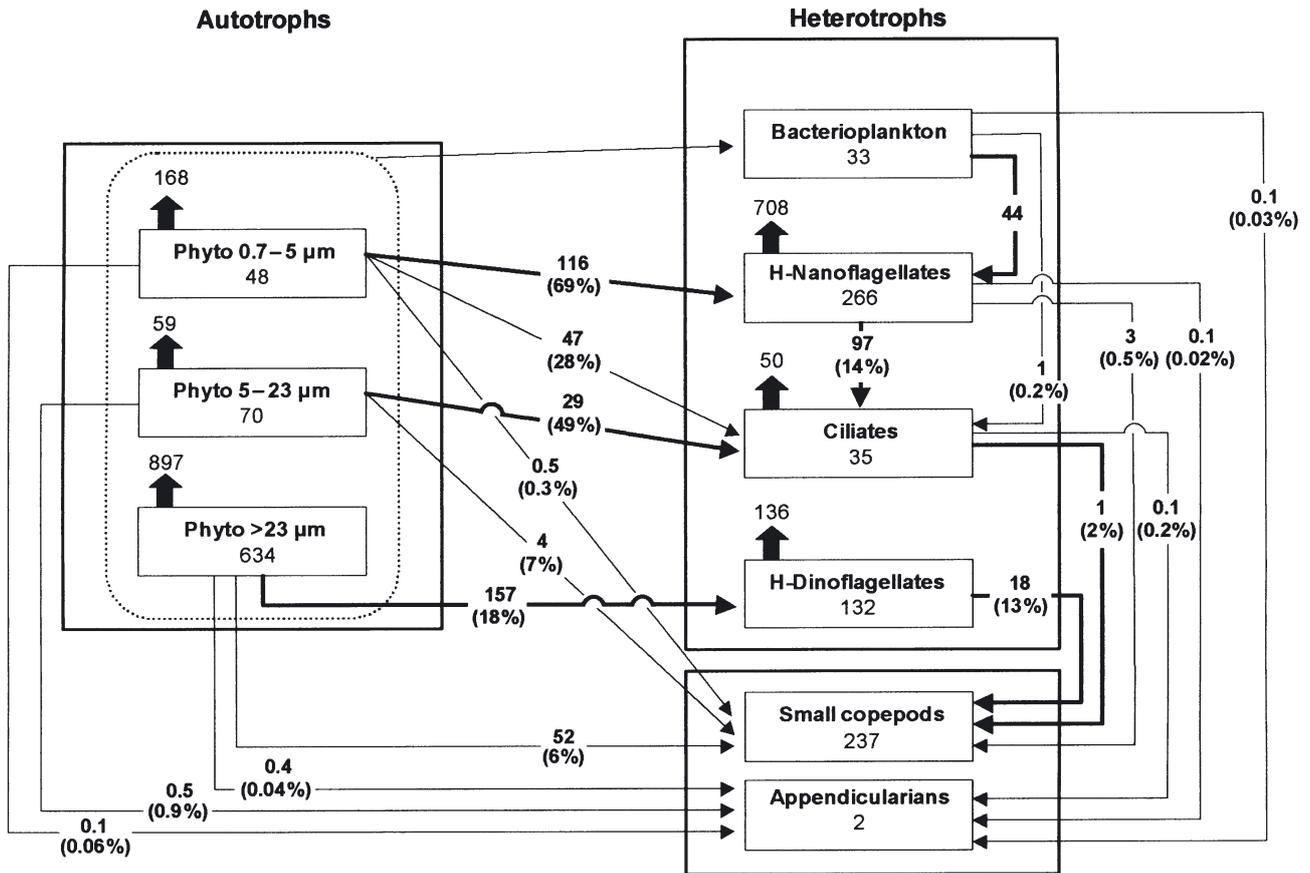


Fig. 7. Carbon budget in October 2001. Further details as in legend to Fig. 5

underestimated during our study, which used a 200 μm net (Krsinic & Lucic 1994, Gallienne & Robins 2001). However, since *Oikopleura dioica* individuals in our samples ranged between 0.8 to 1.2 mm in size, we assume that only the early developmental stages of *O. dioica* could have been undersampled. Appendicularians were the only zooplankton group able to graze on bacteria at ~1% BP d⁻¹, a rate which is not sufficient to have any effect on bacterial biomass but important enough to transfer a significant amount of bacterial carbon directly to higher trophic levels (e.g. fish larvae) (King et al. 1980, Gorsky & Fenaux 1998). Small copepods showed a strong top-down control of ciliates and dinoflagellates, which transferred nanoplanktonic carbon towards higher trophic levels. In February, small copepods removed 29% of the ciliate production, and during October, when PP was low, small copepods fed on h-dinoflagellates, consuming 13% of their production (Figs. 5 & 7). Thus, high clearance rates of copepods on h-dinoflagellates may also result in a significant impact on these protozoans, which are important competitors for diatoms. Because of the extremely high PP in this coastal embayment, a potential high produc-

tion of DOC could support bacterial production, which is efficiently utilized by h-nanoflagellates and ciliates and subsequently by copepods and appendicularians, fuelling the microbial carbon pathways (Uitto et al. 1997). Even though appendicularians may extend the size range of plankton grazed upon, the grazing exerted by both small copepods and appendicularians was not high enough to control the phytoplankton blooms generated during upwelling events. Accordingly, a high percentage of PP would be available for other metabolic processes in the water column (e.g. respiration), for offshore exportation or for sedimentation. Measurements of pCO₂ in the ocean-atmosphere interphase and vertical carbon fluxes during El Niño 1997/1998 by González et al. (1998) showed that carbon sequestration in Mejillones Bay accounted for only a small fraction of PP (3 to 8% PP d⁻¹), suggesting that recycling is important in this bay. Comparison with other biological carbon fluxes measured in a parallel study by Eissler & Quiñones (1999) suggests that microplankton respiration comprises the highest non-photosynthetic carbon flux, and consequently is a crucial factor for the flux of carbon to the deep ocean.

Our results show that microprotozoans can remove a significant percentage of PP (26 to 45 % d⁻¹), and have a much higher impact than small zooplankton (1 to 6 % d⁻¹). Our estimates of microprotozoan impacts are within the range found in coastal waters off Washington, USA (17 to 52%: Landry & Hassett 1982), the Baltic Sea (23%: Uitto et al. 1997); and Gulf of St Lawrence (40 to 100%: Tamigneaux et al. 1997). Therefore, the microheterotrophic pathway may also be an important link in highly productive upwelling areas. The present study is one of the first to provide data that clearly show that the microbial-loop pathway is an important link in carbon fluxes of coastal upwelling ecosystems. In this type of ecosystem, some of the carbon from bacterial and protozoan production may pass to larger zooplankton, and subsequently to large metazoans (e.g. fish larvae). According to the classification of trophic pathways by Legendre & Rassoulzadegan (1995), the food-web structure found during our study would be classified as multivorous, whereby herbivorous and microbial grazing modes both have significant roles in carbon export. Nevertheless, the concept of carbon export in a highly dynamic upwelling area is very complex; in addition to enter trophic pathways, autotrophic production may have several other pathways such as exportation to adjacent areas or burial in local sediments.

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