

# Plankton community structure and carbon cycling in a coastal upwelling system. I. Bacteria, microprotozoans and phytoplankton in the diet of copepods and appendicularians

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**ABSTRACT:** Copepod and appendicularian grazing experiments using naturally occurring planktonic assemblages from a coastal embayment (Mejillones Bay, northern Chile upwelling system at 23° S) were conducted between October 2000 and October 2001. Total carbon ingestion rates based on size-fractionated chlorophyll data showed that dominant copepods (*Acartia tonsa*, *Centropages brachiatus*, *Oithona similis* and *Paracalanus parvus*) ingested between 2 and 8  $\mu\text{gC ind.}^{-1} \text{d}^{-1}$ , while appendicularians (*Oikopleura dioica* and *O. longicauda*) ingested ~3 to 4  $\mu\text{gC ind.}^{-1} \text{d}^{-1}$ . Even when most copepods were feeding on larger cells (>23  $\mu\text{m}$ ) at high rates, the smaller copepods also grazed at similar rates on nanoplankton (5 to 23  $\mu\text{m}$ ) and picoplankton (<5  $\mu\text{m}$ ). In contrast, chain-forming diatoms were cleared at very low rates by copepods. Bacteria were cleared only by appendicularians (~170 to ~400  $\text{ml ind.}^{-1} \text{d}^{-1}$ ) but not by any copepod, while heterotrophic protists constituted a substantial proportion in the diet of both copepods and appendicularians (~10 to 100% body carbon  $\text{d}^{-1}$ ), particularly during austral spring. Occasionally, copepod C-specific ingestion on heterotrophs was similar to that on autotrophic cells. Large ciliates and dinoflagellates were cleared but not ingested by the appendicularian *O. dioica*, suggesting a mechanism of trapping large cells in their houses and implying a rapid export of fresh material. Since heterotrophs are a common component in the diet of these 2 groups (omnivory by copepods and bacterivory by appendicularians), they can potentially affect microbial food webs in this upwelling system and thus carbon export.

**KEY WORDS:** Carbon flux · Omnivory · Microzooplankton · Clearance rate · Copepods · Appendicularians

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## INTRODUCTION

Coastal upwelling systems play an important role in biological productivity and carbon cycling within oceans. While comprising a small portion of the surface area, continental margins account for more than 50% of primary productivity in the oceans (Jahnke & Shimmield 1995). Even though the quantitative contribution of upwelling systems to carbon fluxes is still uncertain, they have a widely recognized role in global carbon export (Liu et al. 2000). As in coastal and oceanic areas, microbially dominated food webs of continental margins en-

hance carbon retention in surface waters, while zooplankton activities enhance carbon export by algal grazing and production of sinking faeces (Peinert et al. 1989).

The coastal upwelling area off northern Chile is one of the major coastal upwelling regions in the Humboldt Current System (HCS). The area is a region of high production as a result of fertilization of surface waters by nutrients transported from depth during upwelling (Thomas et al. 1994, Morales et al. 1996). Most research on the fate of primary production in the Chilean upwelling system has only studied herbivory by the abundant copepod genera *Calanus*, *Paracalanus* and

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*Acartia* (e.g. Gonzalez et al. 2000, Grünewald et al. 2002). It is often assumed that copepods feed mostly on phytoplankton, mainly diatoms; therefore, trophic coupling between copepods and protists such as ciliates, dinoflagellates and heterotrophic nanoflagellates has been less thoroughly examined in this region. Predation by omnivorous copepods on other components of the phytoplankton and microzooplankton, especially small heterotrophic nanoflagellates and ciliates, may be important. For other coastal areas it has been suggested that protists occasionally constitute the main food source for calanoid and cyclopoid copepods (Kleppel et al. 1991, Pierce & Turner 1992, Levinsen et al. 2000), which would have important implications for food web dynamics.

On the other hand, some attention has been given to the role of large-sized microphages, e.g. appendicularian tunicates. Appendicularians may play an important role in channeling carbon from bacteria and small-sized phytoplankton into downward export (Fortier et al. 1994). In contrast with copepods, which have long been considered the major component of secondary production in the sea, appendicularians use their mucus network to graze on bacteria, small nanoflagellates (Alldredge & Madin 1982, Deibel & Powell 1987), fine-grained lithogenic material (Dagg et al. 1996) or even colloidal material (Flood et al. 1992). While the biomass of appendicularians is lower than that of copepods, the production of the former is >30 to 100% of the latter in coastal areas (Nakamura et al. 1997, Hopcroft & Roff 1998).

The sparse information on this group in the coastal upwelling area off northern Chile makes it difficult to assess the quantitative and ecological importance of appendicularians in this ecosystem and to determine their position in and influence on food-web dynamics. In addition, no information is available on the feeding rates of either dominant copepods or appendicularians on microprotozoans and bacteria, or on the factors regulating their feeding in Chilean upwelling ecosystems; this precludes any speculations about their potential role in the pelagic food web.

If small-sized calanoid and cyclopoid copepods and appendicularian tunicates are able to determine or modify the fate of the production of small-sized phytoplankton and microprotozoans, they could have a significant impact on the pathway of carbon export and cycling in coastal upwelling ecosystems. In order to provide insight into this, the present study aimed to study the feeding of small copepods and appendicularians on the full food size-spectrum (phytoplankton, microprotozoan and bacteria) in a coastal upwelling area off northern Chile by measuring their clearance and ingestion rates with natural assemblages of protozoan (dinoflagellates ciliates and heterotrophic nanoflagellates) and 5 different fractions of phytoplankton

(chain-forming diatoms, pennate and centric diatoms, autotrophic nanoflagellates and dinoflagellates) and bacteria.

## MATERIALS AND METHODS

**Feeding experiments.** Experiments were conducted on 4 occasions: (1) austral spring, 17 to 22 October 2000; (2) austral summer, 9 to 10 February 2001; (3) austral winter, 1 to 2 August 2001; (4) austral spring, 21 to 22 October 2001. Copepods and appendicularians were sampled at a coastal station (Stn Exp) in Mejillones Bay (23° S, 73° 20' W; Fig. 1) by slow vertical hauls in the upper 25 m using a WP-2 net (mesh size 200 µm) with a large non-filtering cod-end (~40 to 60 l). Immediately after collection, the contents of the cod-end were transferred to a thermobox and brought to the coastal laboratory in Mejillones.

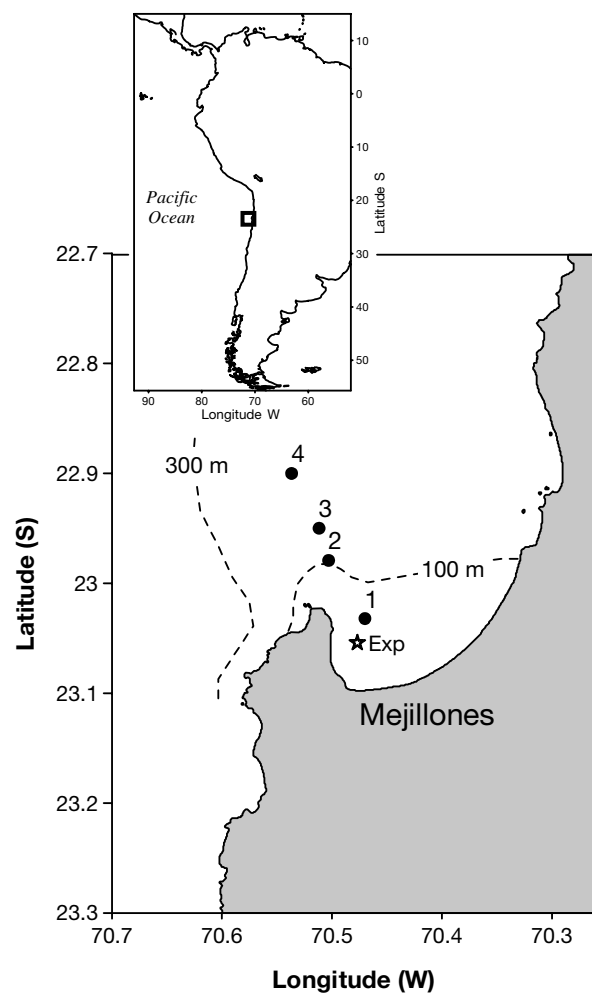


Fig. 1. Study area in Mejillones Bay showing position of the sampling station (Stn Exp)

Within 1 h of collection, undamaged copepods were sorted under a stereomicroscope and appendicularians with intact houses were removed with a wide-mouth pipette and transferred to 200 ml beakers and stored at *in situ* temperature until the experiment began (Table 1). Water for incubation was collected from 10 m with clean Teflon-coated Niskin bottles (Go-Flo, 5 l) and subsequently screened through a 200  $\mu\text{m}$  net to remove most grazers. The animals were pipetted into 500 ml acid-washed polycarbonate bottles with ambient water and filled to avoid air bubbles; 3 control bottles without animals and 3 bottles with 3 to 5 individuals each were placed on a plankton rotating wheel (0.2 rpm) in darkness at *in situ* temperature for approximately 15 to 20 h. Copepod and appendicularian concentrations in the experimental bottles ranged between 6 and 10 ind.  $\text{l}^{-1}$  (Table 1), since during our study their abundance ranged between ~300 and 750 ind.  $\text{m}^{-3}$  in the upper 0 to 25 m layer. Initial control bottles were immediately preserved with 2% acid Lugol's solution and a subsample was preserved in glutaraldehyde (6.0% w/v in 0.2  $\mu\text{m}$  prefiltered seawater). At the end of incubation, subsamples were taken from all bottles: 5 ml were preserved in glutaraldehyde for determination of bacterial biomass, 20 ml for nanoflagellate counts (when not stained immediately with proflavine) and 60 ml were preserved with acid Lugol's solution for determination of cell concentration. In addition, for determination of size-fractionated chlorophyll *a* (chl *a*) (<5  $\mu\text{m}$ ; 5–20  $\mu\text{m}$  and >23  $\mu\text{m}$ ) three 100 ml subsamples were also filtered and dark-extracted in acetone 95% before measurement on a TD 700 Turner fluorometer (Strickland & Parsons 1972). The rest of each sample was gently poured through a 20  $\mu\text{m}$  sieve to collect copepods or appendicularians, their eggs and fecal pellets and appendicularian houses. The cephalothoracic length of copepods and the trunk length of appendicularians were measured under a dissecting microscope and fecal pellets and houses were counted. The carbon content of the animals was calculated using the length–weight regressions of Klein Breteler et al. (1982: for *Acartia* sp. and *Centropages* sp.) Hirche & Mumm (1992: for *Oithona similis*), Uye (1982: for other copepods), and Paffenhöfer (1976) and Gorsky et al. (1988: for *Oikopleura* spp.). The chl *a* in the appendicularian houses was measured to correct ingestion rates due to particles potentially cleared from the experimental bottles but trapped in the houses.

Table 1. Grazing experiments with various copepod species and appendicularian *Oikopleura dioica* in Mejillones Bay during austral spring, summer and winter situations. n: no. of replicate grazing bottles

Expt, Date	Density (no. $\text{l}^{-1}$ )	Size range (mm)	n	Duration (h)	Temp ( $^{\circ}\text{C}$ )
1 21–22 Oct 2000					
<i>Acartia tonsa</i>	8	0.7 – 0.9	3	18.5	12
<i>Centropages brachiatus</i>	6	1.0 – 1.2	3	19	12
<i>Oithona similis</i>	10	0.6 – 0.8	3	19.5	12
2 10–11 Feb 2001					
<i>Oikopleura dioica</i>	6	0.5 – 0.7	3	15	19
<i>Acartia tonsa</i>	8	0.7 – 0.9	3	14	19
<i>Paracalanus parvus</i>	8	0.6 – 0.7	3	14	19
3 1–2 Aug 2001					
<i>Oikopleura longicauda</i>	6	0.5 – 0.55	3	20	14
<i>Acartia tonsa</i>	8	0.8 – 1.0	3	19.5	14
<i>Paracalanus parvus</i>	10	0.7 – 0.8	3	20	14
4 21–22 Oct 2001					
<i>Oithona similis</i>	10	0.5 – 0.65	3	19.8	16.5
<i>Centropages brachiatus</i>	6	1.2 – 1.6	3	21.0	16.5
<i>Paracalanus parvus</i>	8	0.65 – 0.9	3	20.3	16.5

**Cell counts and calculation of clearance and ingestion rates.** Nanoflagellate and bacteria samples were analysed using a color-image analysis system similar to that described by Verity & Sieracki (1993). Samples were counted with the aid of a video camera connected to a Zeiss epifluorescence microscope and a Sony® monitor. A Windows-based imaging software package (Optimas® V. 6.0) controlled image-capture, measurement, analysis and output. Bacteria were quantified by the acridine orange technique (Hobbie et al. 1977). Bacterial volume was calculated from length and width measurements of at least 50 cells per sample. Biovolume was converted to carbon using the equation  $C$  (fg) =  $90.06 \times V$  ( $\mu\text{m}^3$ )<sup>0.59</sup>, where  $V$  = bacterial volume (Simon & Azam 1989, Riemann & Bell 1990).

For the enumeration of nanoflagellates, after incubation, subsamples were immediately filtered at 0.8  $\mu\text{m}$  and stained with proflavine (0.033% w/v in distilled water) according to Haas (1982) and fixed with glutaraldehyde (as described above) until subsequent analysis. Nanoflagellates were counted and divided into 2 groups: <5  $\mu\text{m}$  and 5–20  $\mu\text{m}$ , and autotrophic cells were identified by autofluorescence. In October 2000 and October 2001, enumeration of nanoflagellates was done using acridine orange as fluorochrome (Davis & Sieburth 1982).

Large cells were counted under an inverted microscope (Leica Leitz DMIL). Subsamples of 50 ml were allowed to settle for 24 h in sedimentation chambers (Utermöhl 1958) before diatoms, dinoflagellates and ciliates were identified, counted and measured under the microscope. Plasma volumes were calculated (Edler 1977) and averaged from a minimum of 50 cells

species<sup>-1</sup>. Ciliate biovolume was calculated assuming conical shapes, with a length to diameter ratio of 1.25 for ciliates <50 µm and 2 for ciliates >50 µm (Tiselius 1989). We used carbon to plasma volume ratios of 0.11 pgC µm<sup>-3</sup> for diatoms (Edler 1979), 0.3 and 0.19 gC µm<sup>-3</sup> for heavily thecate and athecate dinoflagellates forms, respectively (E. J. Lessard unpubl. data, in: Gifford & Caron 2000), and 0.148 pgC µm<sup>-3</sup> for ciliates (Ohman & Snyder 1991).

Clearance and ingestion rates were calculated according to Frost (1972) for the following groups: bacteria, autotrophic (a-) and heterotrophic (h-) nanoflagellates (<5 µm and 5–20 µm), a- and h-dinoflagellates, silicoflagellates, ciliates, pennate and centric diatoms and chain-forming diatoms. Chl *a* ingestion was converted to carbon ingestion by multiplying by the mean carbon concentration of cells derived from 10–20 m Niskin-bottle sampling. Clearance was calculated only when the difference in prey concentration between control and experimental bottles proved to be significant (Student's *t*-test: *p* < 0.05, Table 2). A *Q*<sub>10</sub> = 2.8 was used to eliminate temperature differences among experiments (Hansen et al. 1997).

One of the biases arising from the incubation method is that the prey suspension contains several trophic levels. In order to correct for this bias, we applied a 3-component equation template that considers interactions among 3 grazers in differently structured food chains (Tang et al. 2001). The values for protozoan grazing used in this template were estimated using a model proposed by Peters (1994), which includes temperature, cell volumes, and concentrations of both prey and predator as explanatory variables and predicts ingestion rates over a wide range of biological and

environmental conditions. Corrections were done for the interactions bacteria–flagellates–zooplankton, flagellates–ciliates–zooplankton and diatoms–h-dinoflagellates–zooplankton.

Field cell concentration and biomass for bacteria, nanoflagellates, ciliates and phytoplankton was estimated using water samples collected from 10 to 20 m with clean Teflon-coated Niskin bottles (Go-Flo; 5 l). Biomass was determined using the methods described in the preceding subsection.

## RESULTS

The natural phytoplankton and protozoans provided as food varied in abundance among the experiments. During the study period, the major contribution to total biomass was by chain-forming diatoms (Fig. 2), mainly species of *Chaetoceros*, *Guinardia*, *Eucampia*, and *Pseudonitzschia*. Highest diatom biomass occurred during the August experiment (8.6 ± 3.6 gC m<sup>-2</sup>). In October 2000, autotrophic nanoflagellates constituted a large fraction of the autotrophic biomass (2.6 ± 0.9 gC m<sup>-2</sup>, 32% of total autotrophs). Autotrophic and heterotrophic cells contributed equally to the nanoplankton pool (*t*-test: *p* > 0.05), while most dinoflagellates were autotrophs (*t*-test: *p* < 0.05), mainly represented by *Ceratium* and *Prorocentrum* species. Autotrophic (a-) dinoflagellates were more abundant in the summer and winter samples (2 to 4 gC m<sup>-2</sup>), when chain-forming diatoms dominated. Dinoflagellates, ciliates and bacteria were also less important in terms of biomass during our study, with highest biomass during the October 2001 sampling.

Table 2. Significance levels of difference between prey densities in control and experimental bottles (Student's *t*-test): \*\**p* < 0.01, \**p* < 0.05, ns: no significant difference. ChainD: chain-forming-diatoms; PenD: pennate diatoms; CenDand: centric diatoms; AuDino: autotrophic nanoflagellates; HeDino: heterotrophic dinoflagellates; AN: autotrophic nanoflagellates (<5 µm); HN: heterotrophic nanoflagellates (>5 µm); Bact: bacteria. Experiment dates in Table 1

Expt, Species	Diatoms			Dinoflagellates		Ciliates	Silicoflagellates	Nanoflagellates				Bacteria
	ChainD	PenD	CenD	AuDino	HeDino			AN>5	HN>5	AN<5	HN<5	
1 <i>Acartia tonsa</i>	*	*	ns	ns	ns	ns	*	**	**	*	*	ns
<i>Centropages brachiatus</i>	*	*	ns	*	ns	*	ns	*	*	*	*	ns
<i>Oithona similis</i>	*	*	ns	*	*	*	ns	*	*	*	*	ns
2 <i>Oikopleura dioica</i>	*	*	*	*	ns	*	ns	**	*	**	*	*
<i>Acartia tonsa</i>	**	*	*	**	**	*	*	**	**	**	*	ns
<i>Paracalanus parvus</i>	**	*	*	**	*	*	**	*	**	ns	ns	ns
3 <i>Oikopleura longicauda</i>	*	*	*	*	*	*	*	*	*	*	*	**
<i>Paracalanus parvus</i>	*	ns	*	*	**	*	*	*	*	ns	ns	ns
<i>Acartia tonsa</i>	ns	*	*	*	*	*	*	*	*	*	ns	ns
4 <i>Oithona similis</i>	*	*	*	*	*	*	ns	*	**	*	*	ns
<i>Centropages brachiatus</i>	*	*	*	**	*	*	ns	**	*	**	*	ns
<i>Paracalanus parvus</i>	*	*	*	**	ns	**	ns	**	*	*	*	ns

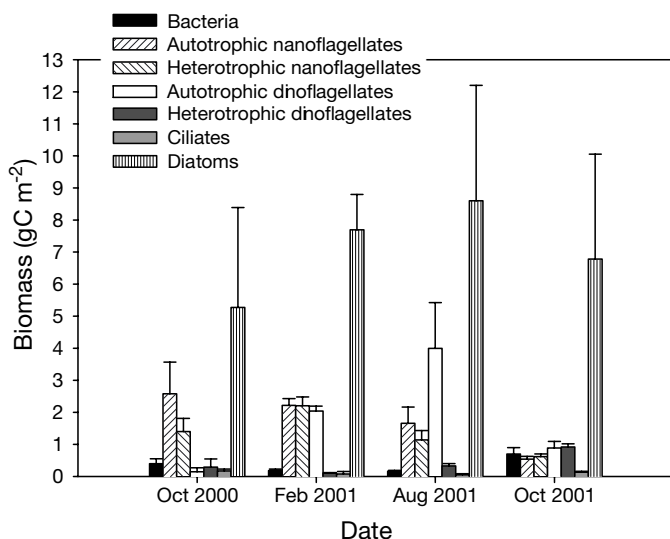


Fig. 2. Contribution of major taxonomic groups to integrated (0 to 25 m depth) biomass of autotrophic and heterotrophic community at sampling station during feeding experiments. Error bars = SD of replicate samples ( $n = 3$ )

### Copepod and appendicularian clearance and ingestion

Grazing on natural food assemblages was relatively similar among the small-copepod species. Small copepods such as *Acartia tonsa* and *Oithona similis* did not ingest cells in proportion to their abundance. For instance, although during our study diatoms made the most important contribution to biomass (Fig. 2), size-fractionated chlorophyll ingestion by *A. tonsa* and *O. similis* showed that they were feeding mostly on the 5–23 and <5  $\mu\text{m}$  fractions respectively (Table 3). Cell counts showed that *A. tonsa* mostly ingested a- and h-nanoflagellates >5  $\mu\text{m}$  (Figs. 3 & 4). However, during August, when a high diatom biomass was observed, *A. tonsa* was grazing mostly on centric diatoms (>23  $\mu\text{m}$  fraction) (Table 3 & Fig. 5). At that time, carbon ingestion was the highest of all experiments. Similarly, during the October 2001 experiment, a low nanoflagellate biomass was observed and *O. similis* was also grazing on large cells >23  $\mu\text{m}$ , mostly centric diatoms (Fig. 6).

Since dinoflagellates and ciliates were very scarce during all experiments, clearance on h-dinoflagellates and ciliates was occasionally higher than on phytoplankton (e.g. February) and predation was so intense that ciliates were reduced to ca. 10% of their initial concentrations (data not shown). For instance, in the October 2001 experiment, high

carbon ingestion on h-dinoflagellates by *Oithona similis* was observed (Fig. 6). Predation by *Acartia tonsa* and *O. similis* on h-nanoflagellates and other microprotozoans (e.g. h-dinoflagellates and ciliates), resulted in heterotrophic prey contributing, on average, one-third of the total ingested carbon, the other two-thirds arising from autotrophic prey (Table 4).

Size-fractionated chlorophyll ingestion showed that in all experiments *Centropages brachiatus* ingested large cells (>23  $\mu\text{m}$ ), with carbon ingestion rates relatively similar between experiments (Table 3). However, heterotrophic prey such as dinoflagellates and ciliates were always cleared faster than diatoms (Figs. 3 & 6). A-nanoflagellates were also ingested, but only in October 2000, when they were most abundant (Fig. 2). Since h-dinoflagellates were abundant in October 2001, high clearance and ingestion of this prey was observed. In this experiment, the percentage of body carbon (BC) ingested from heterotrophs almost equalled that from autotrophs (Table 4). Similarly, *Paracalanus parvus* fed mostly on

Table 3. Carbon ingestion rates ( $\mu\text{gC ind.}^{-1} \text{d}^{-1}$ ; mean  $\pm$  SD) based on size-fractionated chlorophyll data for various zooplankton species studied seasonally in Mejillones Bay. During February 2001, only fractions <23 and >23  $\mu\text{m}$  were analyzed. Chl:C ratio ranged between 90 and 150

Species	Oct 2000	Feb 2001	Aug 2001	Oct 2001
Fraction ( $\mu\text{m}$ )				
<i>Acartia tonsa</i>				
<5	0.7 $\pm$ 0.9		0.1 $\pm$ 0.4	
5–23	1.3 $\pm$ 0.5	2.8 $\pm$ 0.4 (< 23)	0.1 $\pm$ 0	
>23	1.6 $\pm$ 0.3	1.2 $\pm$ 0.2	7.0 $\pm$ 4.6	
Total	3.6 $\pm$ 1.8	4.0 $\pm$ 0.7	7.0 $\pm$ 5.0	
<i>Centropages brachiatus</i>				
<5	0.5 $\pm$ 0.2			0.1 $\pm$ 0.01
5–23	2.5 $\pm$ 1.7			0.6 $\pm$ 0.1
>23	0.4 $\pm$ 0.3			2.9 $\pm$ 0.3
Total	3.4 $\pm$ 2.3			3.6 $\pm$ 1.5
<i>Oithona similis</i>				
<5	1.3 $\pm$ 0.7			0.1 $\pm$ 0.1
5–23	0.5 $\pm$ 0.4			0 $\pm$ 0
>23	0.8 $\pm$ 0.8			2.8 $\pm$ 1.4
Total	2.6 $\pm$ 1.9			2.9 $\pm$ 1.5
<i>Oikopleura dioica</i>				
<23		3.3 $\pm$ 0.4		
>23		0.8 $\pm$ 1.5		
Total		4.1 $\pm$ 1.9		
Total corrected		3.4 $\pm$ 1.8		
<i>Paracalanus parvus</i>				
<5			0 $\pm$ 0	0.2 $\pm$ 0.1
5–23		0.6 $\pm$ 0.2 (< 23)	0 $\pm$ 0	0 $\pm$ 0
>23		2.5 $\pm$ 0.7	8.1 $\pm$ 1.6	3.1 $\pm$ 0.03
Total		3.1 $\pm$ 0.9	8.1 $\pm$ 1.6	3.3 $\pm$ 0.1
<i>Oikopleura longicauda</i>				
<5			0.1 $\pm$ 0.5	
5–23			1.6 $\pm$ 0.5	
>23			3.6 $\pm$ 1.5	
Total			5.3 $\pm$ 2.5	
Total corrected			4.2 $\pm$ 2.2	

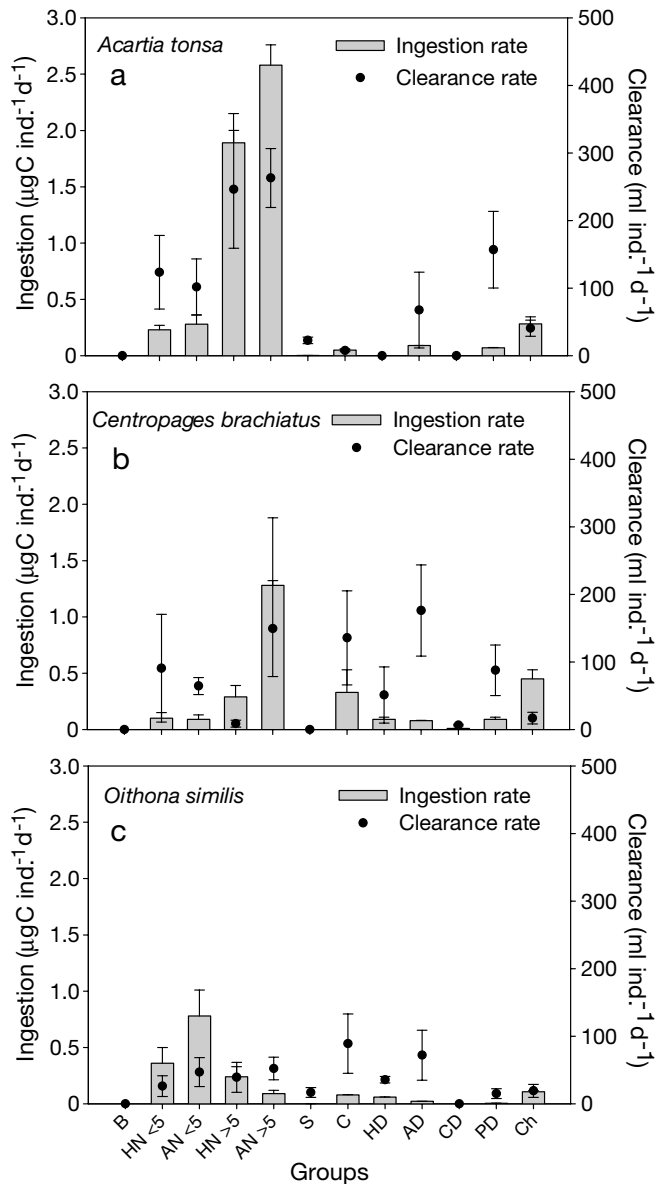


Fig. 3. Clearance and ingestion rate of major autotroph (a) and heterotroph (h-) groups by small copepods in austral spring, October 2000. B: bacteria; HN: h-nanoflagellates; AN: a-nanoflagellates; S: silicoflagellates; C: ciliates; HD: h-dinoflagellates; AD: a-dinoflagellates; CD: solitary centric diatoms; PD: pennate diatoms; Ch: chain forming diatoms. <5 or >5 = <5 or >5  $\mu\text{m}$  cell size. Error bars = SD of replicate samples ( $n = 3$ )

large cells during all experiments. Nanoflagellates, ciliates and dinoflagellates were not selected by this copepod, and most of its ingested carbon originated from centric diatoms and, occasionally in October 2001, on abundant h-dinoflagellates (Figs. 4 to 6). Therefore, the high % BC  $\text{d}^{-1}$  ingested by *P. parvus* was mostly based on autotrophic cells (Table 4).

Clearance of appendicularians was always higher than that of copepods (Figs. 5 & 6). Despite their abundance,

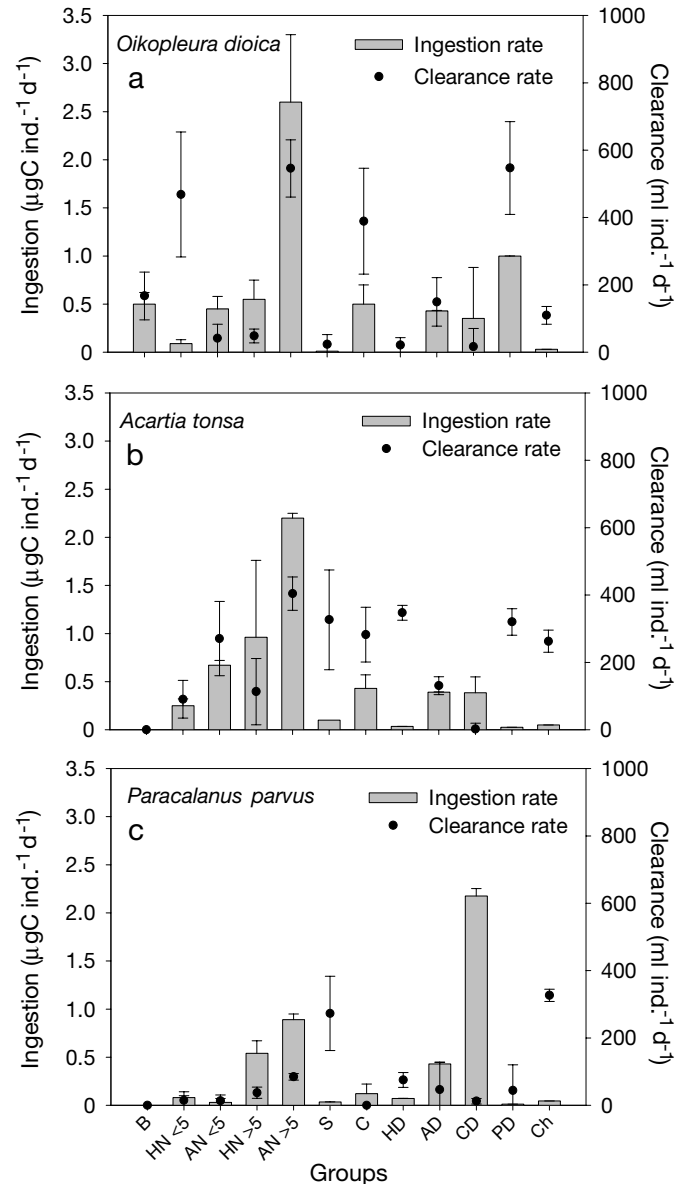


Fig. 4. Clearance and ingestion rates of major autotroph and heterotroph groups by small copepods and appendicularians in austral summer, February 2001. Further details as in Fig. 3

long chain-forming diatoms were not eaten by the appendicularian *Oikopleura dioica*, whose highest clearance and ingestion rates were on a-nanoflagellates >5  $\mu\text{m}$  (~545  $\text{ml ind.}^{-1} \text{d}^{-1}$  and 2.6  $\mu\text{gC ind.}^{-1} \text{d}^{-1}$  respectively). Higher ingestion on autotrophs than heterotrophs was observed (Table 4). Another appendicularian, *O. longicauda*, ingested 4.2  $\mu\text{gC ind.}^{-1} \text{d}^{-1}$  during an experiment in August. Due to the low abundance of ciliates in this experiment, high clearance but low carbon ingestion was observed (500  $\text{ml ind.}^{-1} \text{d}^{-1}$ ) (Fig. 5). As with copepods, larger cells were also selected by *O. longicauda* and most carbon inges-

ted also came from centric diatoms ( $\sim 3.5 \mu\text{gC ind.}^{-1} \text{d}^{-1}$ ). *O. longicauda* showed the highest carbon ingestion of all species studied (auto + heterotrophs cells =  $\sim 9 \mu\text{gC ind.}^{-1} \text{d}^{-1}$ ) (Table 4). A large amount of carbon was not ingested, but was trapped and potentially exported on or into appendicularian houses ( $0.7$  and  $1.1 \mu\text{gC ind.}^{-1} \text{d}^{-1}$  for *O. dioica* and *O. longicauda* respectively). Bacteria and small nanoflagellates ( $< 5 \mu\text{m}$ ) were cleared at high rates by both appendicularian species ( $\sim 200$  to  $500 \text{ ml ind.}^{-1} \text{d}^{-1}$ ), but not by any copepod (*t*-test:  $p > 0.05$ , Table 2).

### Effects of cell-size and concentration on clearance

Mean weight-specific, temperature-corrected clearance rates showed a functional response for *Acartia tonsa* grazing on natural food assemblages. When taking into account all particles available, clearance decreased at high carbon concentrations. However, a dome-shape curve emerged when food sources were considered separately (e.g. a- and h-nanoflagellates). Maximum clearance was  $\sim 250 \text{ ml } \mu\text{gC d}^{-1}$  at  $19^\circ\text{C}$  (Fig. 7a) at a concentration of  $\sim 11 \mu\text{gC l}^{-1}$  for pennate diatoms. At high carbon concentrations, clearance decreased considerably. Maximum clearance on a- and h-dinoflagellates was  $\sim 60 \text{ ml } \mu\text{gC d}^{-1}$  at concentrations between  $6$  and  $12 \mu\text{gC l}^{-1}$ . Centric diatoms did not show any pattern and clearance of them was relatively low. Similarly, the clearance rate of *Paracalanus parvus* also decreased with increasing carbon concentration. Clearance of both a- and h-nanoflagellates by *P. parvus* varied almost independently of prey concentration (Fig. 7b). A maximum weight-specific clearance on dinoflagellates of  $\sim 200 \text{ ml } \mu\text{gC d}^{-1}$  at h-dinoflagellate concentrations around  $13 \mu\text{gC l}^{-1}$  was observed. At higher concentrations, clearance decreased considerably. Feeding on pennate and centric diatoms decreased at concentrations higher than  $\sim 100 \mu\text{gC l}^{-1}$ , with a maximum clearance of  $119$  and  $183 \text{ ml } \mu\text{gC d}^{-1}$  respectively.

The size of prey cells also had an effect on the clearance rate of copepods and appendicularians (Fig. 8). For the entire size spectrum, the volume cleared increased in a generally hyperbolic fashion for both copepods and appendicularians. *Acartia tonsa* grazed on all cells available, but clearance increased from about  $0$  to  $350 \text{ ml } \mu\text{gC d}^{-1}$  within a prey size range of  $\sim 0.5 \mu\text{m ESD}$  (bacteria) to  $\sim 85 \mu\text{m ESD}$  (mainly ciliates *Eutimnopsis* spp. and *Tintinopsis* spp.). However, clearance decreased when *A. tonsa*

Table 4. Daily ingestion rate (IR) ( $\mu\text{gC ind.}^{-1} \text{d}^{-1}$ ; mean  $\pm$  SD) on autotroph (A) and heterotroph (H) cells based on cells counts. Mean zooplankton carbon content ( $\mu\text{gC ind.}^{-1}$ ) and IR as percentage of body carbon are also shown

Species	Trophic status	Mean IR ( $\mu\text{gC ind.}^{-1} \text{d}^{-1}$ )	Mean carbon content ( $\mu\text{gC ind.}^{-1}$ )	IR (% body carbon)
<b>Expt 1</b>				
<i>Acartia tonsa</i>	A	$3.2 \pm 0.3$	$4.1 \pm 0.8$	78.0
	H	$2.2 \pm 0.6$	$4.1 \pm 0.8$	53.6
<i>Centropages brachiatus</i>	A	$2.0 \pm 0.7$	$8.7 \pm 2.3$	22.9
	H	$0.8 \pm 0.6$	$8.7 \pm 2.3$	9.2
<i>Oithona similis</i>	A	$1.0 \pm 0.3$	$2.0 \pm 0.1$	50
	H	$0.7 \pm 0.6$	$2.0 \pm 0.1$	35
<b>Expt 2</b>				
<i>Oikopleura dioica</i>	A	$4.9 \pm 1.3$	$4.1 \pm 0.8$	119.5
	H	$1.6 \pm 0.5$	$4.1 \pm 0.8$	39.0
<i>Acartia tonsa</i>	A	$3.8 \pm 0.3$	$6.7 \pm 2.3$	56.7
	H	$1.7 \pm 1.0$	$6.7 \pm 2.3$	25.3
<i>Paracalanus parvus</i>	A	$3.6 \pm 0.2$	$2.5 \pm 0.1$	144
	H	$0.8 \pm 0.2$	$2.5 \pm 0.1$	32
<b>Expt 3</b>				
<i>Oikopleura longicauda</i>	A	$6.9 \pm 0.9$	$1.6 \pm 0.1$	431.2
	H	$1.9 \pm 0.4$	$1.6 \pm 0.1$	118.7
<i>Acartia tonsa</i>	A	$4.9 \pm 0.3$	$6.4 \pm 0.4$	76.5
	H	$0.8 \pm 0.3$	$6.4 \pm 0.4$	12.5
<i>Paracalanus parvus</i>	A	$5.7 \pm 0.3$	$3.7 \pm 0.1$	154.0
	H	$1.1 \pm 0.1$	$3.7 \pm 0.1$	29.7
<b>Expt 4</b>				
<i>Oithona similis</i>	A	$2.8 \pm 0.1$	$1.2 \pm 0.4$	233.3
	H	$1.4 \pm 0.1$	$1.2 \pm 0.4$	116.6
<i>Centropages brachiatus</i>	A	$2.8 \pm 0.3$	$10.4 \pm 2.3$	26.9
	H	$2.3 \pm 0.2$	$10.4 \pm 2.3$	22.1
<i>Paracalanus parvus</i>	A	$3.7 \pm 0.3$	$3.7 \pm 0.4$	100
	H	$1.0 \pm 0.06$	$3.7 \pm 0.4$	27

was feeding on the large dinoflagellate *Ceratium* spp. ( $175 \mu\text{m ESD}$ ). *Centropages brachiatus* also showed a similar pattern, but high clearance was maintained when this copepod was feeding on large cells. Small cells of  $< 30 \mu\text{m ESD}$  (nanoflagellates and diatoms) and larger than  $100 \mu\text{m ESD}$  (dinoflagellates) were not efficiently cleared by *Paracalanus parvus*. The small cyclopid *Oithona similis* was efficient at feeding on small particles. However, maximum clearance was also observed when *O. similis* were feeding on ciliates and dinoflagellates. Even though maximum clearance rates of *Oikopleura dioica* were also observed on large-size particles such as pennate and centric diatoms ( $20$  to  $50 \mu\text{m ESD}$ ) and ciliates and dinoflagellates ( $> 70 \mu\text{m ESD}$ ), these cells could not be ingested by this appendicularian, mainly because the mesh of its inlet filter is smaller ( $\sim 30 \mu\text{m}$ ) than these cells (Fenaux 1986).

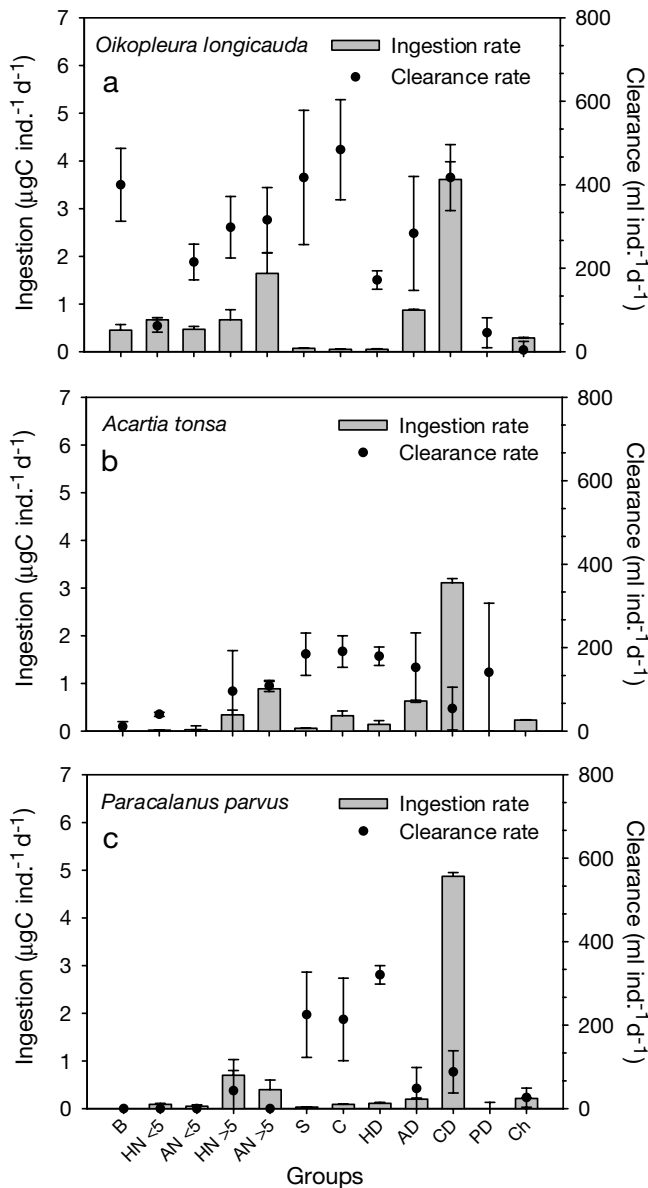


Fig. 5. Clearance and ingestion rate of major autotroph and heterotroph groups by small copepods and appendicularians in austral winter, August 2001. Further details as in Fig. 3

## DISCUSSION

### Comparative total carbon ingestion of copepods and appendicularians

During our study, total carbon ingestion rates based on size-fractionated chlorophyll data showed that small copepods ingested between 2 and 8  $\mu\text{gC ind.}^{-1} \text{d.}^{-1}$ . (Table 3), values similar to those reported for other upwelling regions. Boyd et al. (1980) found a carbon-based ingestion rate of between 0 and 7  $\mu\text{gC ind.}^{-1} \text{d.}^{-1}$  for *Centropages brachiatus* in the Peru upwelling

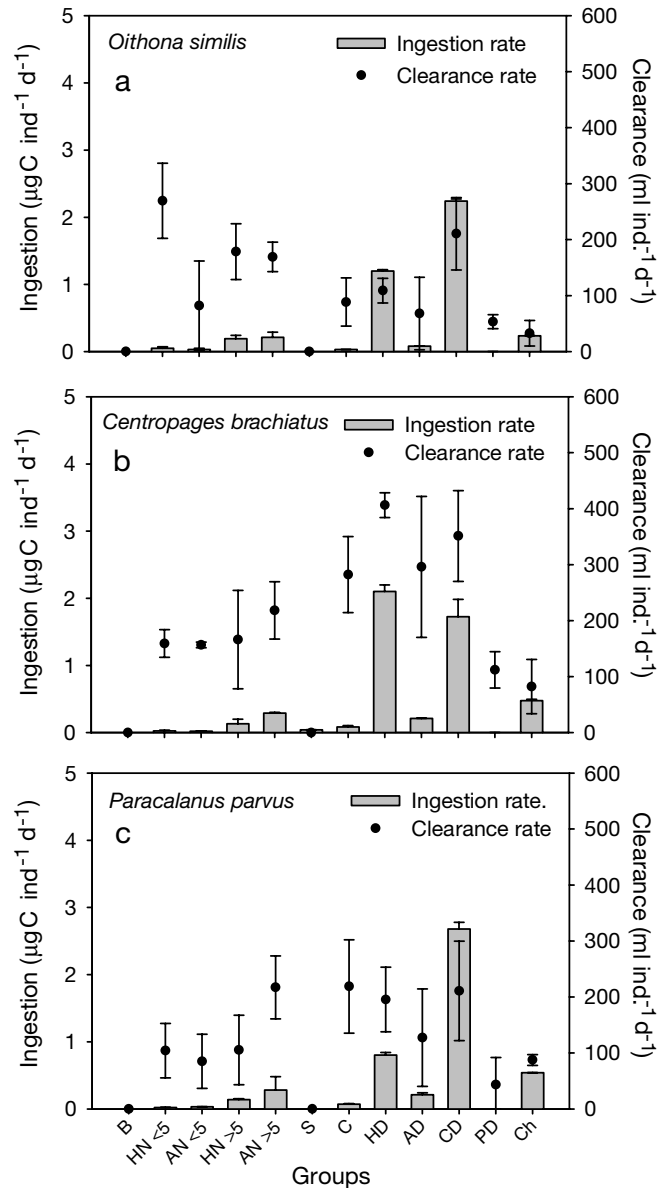


Fig. 6. Clearance and ingestion rate of major autotroph and heterotroph groups by small copepods in austral spring, October 2001. Further details as in Fig. 3

system and *Paracalanus* spp. ingested between 2 to 3.9  $\mu\text{gC ind.}^{-1} \text{d.}^{-1}$  in the Benguela upwelling system (Peterson et al. 1988, in Verheye et al. 1992). In northern Chile, off Antofagasta, González et al. (2000) estimated an ingestion rate of 5 to 22  $\mu\text{gC ind.}^{-1} \text{d.}^{-1}$  for a pool of small calanoid copepods (e.g. *Temora* spp., *Acartia* spp. and *Paracalanus* spp.). However, all these values were for gut-pigment content, and considered only carbon ingestion of autotrophic cells. Further, such ingestion estimates are controversial due to pigment loss to colorless, non-fluorescent compounds (Tirelli & Mayzaud 1998).



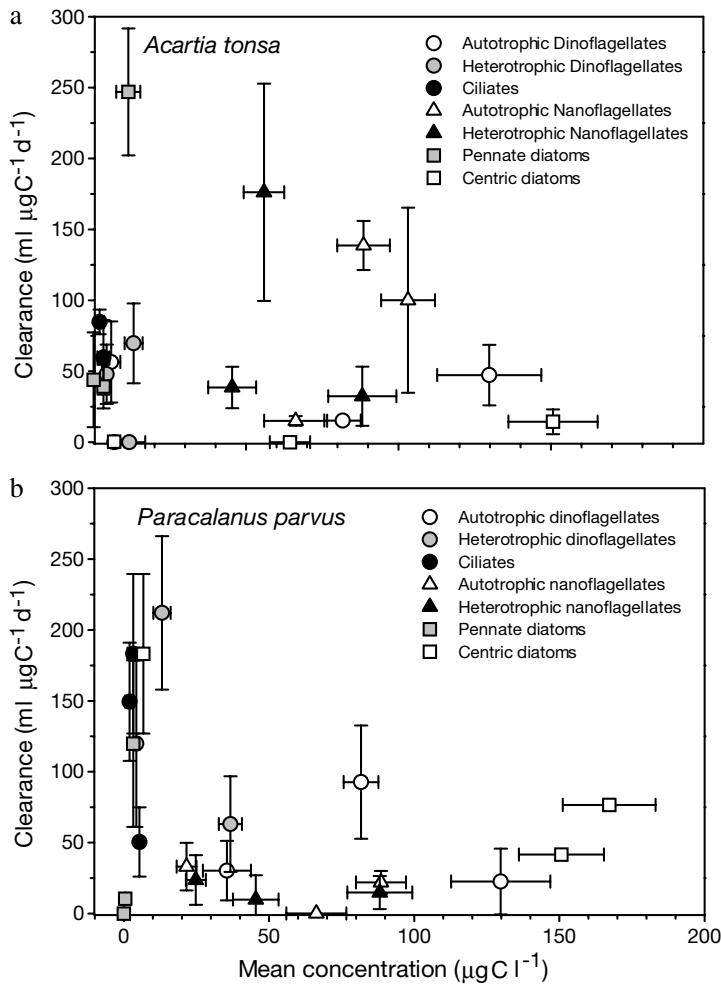


Fig. 7. *Acartia tonsa* and *Paracalanus parvus*. Weight-specific clearance of a- and h-dinoflagellates, ciliates, a- and h-nanoflagellates, pennate diatoms and centric diatoms at different natural carbon concentrations. Data were corrected to 19°C using a  $Q_{10}$  of 2.8. Errors are  $\pm$  SE of mean

There is no prior information on appendicularian feeding in upwelling areas. In the present study, appendicularians had a chlorophyll-based ingestion of  $\sim 4 \mu\text{gC ind.}^{-1} \text{d}^{-1}$ . Ingestion rates of appendicularians are difficult to estimate, because some of the cells removed from suspension adhere to the house and are not ingested (Gorsky 1980). In fact, our results from corrected ingestion, CI (CI = total chl *a* cleared—chl *a* trapped in houses) revealed that between 0.7 and  $1.1 \mu\text{gC}$  were trapped daily in appendicularian houses (house production =  $\sim 5.3$  houses  $\text{ind.}^{-1} \text{d}^{-1}$ , data not shown). This value is similar to potential ingestion by *Oikopleura dioica* on cells  $> 23 \mu\text{m}$  ( $0.8 \mu\text{gC ind.}^{-1} \text{d}^{-1}$ ). If the inlet filter of *O. dioica* retains particles larger than  $\sim 30 \mu\text{m}$  (Fenaux 1986), according to the individual body size (M. Kiefer pers. comm.), then the fraction  $> 23 \mu\text{m}$  may not be ingested, but trapped in the inlet

filter of the house and therefore exported to the benthos. Apart from the exceptions mentioned previously, there are few reports on appendicularian ingestion rates in the literature. Paffenhöfer (1976) found that *O. dioica* ingests around 100 to 200 % BC  $\text{d}^{-1}$ . This is in agreement with the daily ration of  $\sim 150\%$  for *O. dioica* in our study, but much higher than the 60 % BC  $\text{d}^{-1}$  for *O. dioica* estimated from the ingestion of radiolabeled bacterioplankton by Sorokin (1973) (around  $60 \mu\text{gC l}^{-1}$ ). However, the *in situ* concentration of particles in the previous studies were much lower than the average concentration of ca. 200 to  $300 \mu\text{gC l}^{-1}$  reported for northern Chile (González et al. 2000, H. González unpubl. data), and appendicularians may not reach saturation level even at the highest natural particle concentrations (Bochdansky & Deibel 1999). The small *O. longicauda* (1 mm trunk length) ingested  $\sim 550\%$  BC  $\text{d}^{-1}$  (i.e. auto- + heterotrophic prey). High values for daily ration in appendicularians have been also reported for other appendicularians. Deibel (1988) found that *O. vanhoeffeni* can ingest 980 % BC  $\text{d}^{-1}$  in cold Newfoundland waters ( $-2$  to  $4^\circ\text{C}$ ), and estimated that its total daily ration in the Northeast Polynya would be about 250 % BC  $\text{d}^{-1}$  ( $-1^\circ\text{C}$ ) (Deibel 1988). This indicates that clearance by appendicularians is not only high in upwelling temperate regions (i.e.  $14^\circ\text{C}$  in this study) but is also not significantly lower at very low temperatures.

#### Contribution of bacteria, microprotozoans and phytoplankton to carbon ingestion

Most feeding studies on copepods have shown that phytoplankton is consumed primarily in proportion to its abundance (e.g. Turner & Tester 1989). However, the present study revealed that the less-abundant heterotrophic protists constituted a substantial proportion of the diet of both copepods (*Acartia tonsa*, *Centropages brachiatus* and *Oithona similis*) and appendicularians (*Oikopleura dioica* and *O. longicauda*). This microprotozoan–zooplankton coupling was particularly evident in some experiments. Even though chain-forming diatoms made a major contribution to biomass, a- and h-nanoflagellates  $> 5 \mu\text{m}$  were an important component in the diet of *A. tonsa* and *C. brachiatus*. Turner & Granéli (1992) found similar results, with *A. clausi* and *C. hamatus* selectively grazing flagellates. In the present study, in October 2001, when a high biomass of h-dinoflagellates was observed, the trophic link between h-dinoflagellates and copepods was important to the carbon flux in the study area, as a high carbon ingestion was observed for the copepods *C. brachiatus* and *Oithona similis*. The C-specific ingestion of heterotrophs during this experiment was close

to the ingestion of autotrophic cells by both *C. brachiatus* (2.3 and 2.8  $\mu\text{gC ind.}^{-1} \text{d}^{-1}$  respectively) and *O. similis* (1.8 and 2.8  $\mu\text{gC ind.}^{-1} \text{d}^{-1}$  respectively), and almost half the body carbon ingested originated from heterotrophs (~20 and 100% BC  $\text{d}^{-1}$ ). Conversely, *Paracalanus parvus* showed a more herbivorous strategy, as most of its ingested carbon originated from autotrophs during all study periods (Table 4). However, large prey such as dinoflagellates and ciliates were also ingested. The specific carbon clearance rates of *P. parvus* for h-dinoflagellates and ciliates were on

average  $\sim 100 \text{ ml } \mu\text{gC d}^{-1}$ . The value for clearance of h-dinoflagellates is comparable to that reported for *P. parvus* by Suzuki et al. (1999) (33  $\text{ml } \mu\text{gC d}^{-1}$ ), but about 1 order of magnitude higher than for clearance of ciliates (17  $\text{ml } \mu\text{gC d}^{-1}$ ).

The small cyclopid *Oithona similis* cleared small a- and h-nanoflagellates  $< 5 \mu\text{m}$ , which were inefficiently grazed by the other copepods. These results disagree those of Nakamura & Turner (1997), who found that *O. similis* did not significantly ingest small a- and h-nanoflagellates, preying instead on particles  $> 10 \mu\text{m}$ .

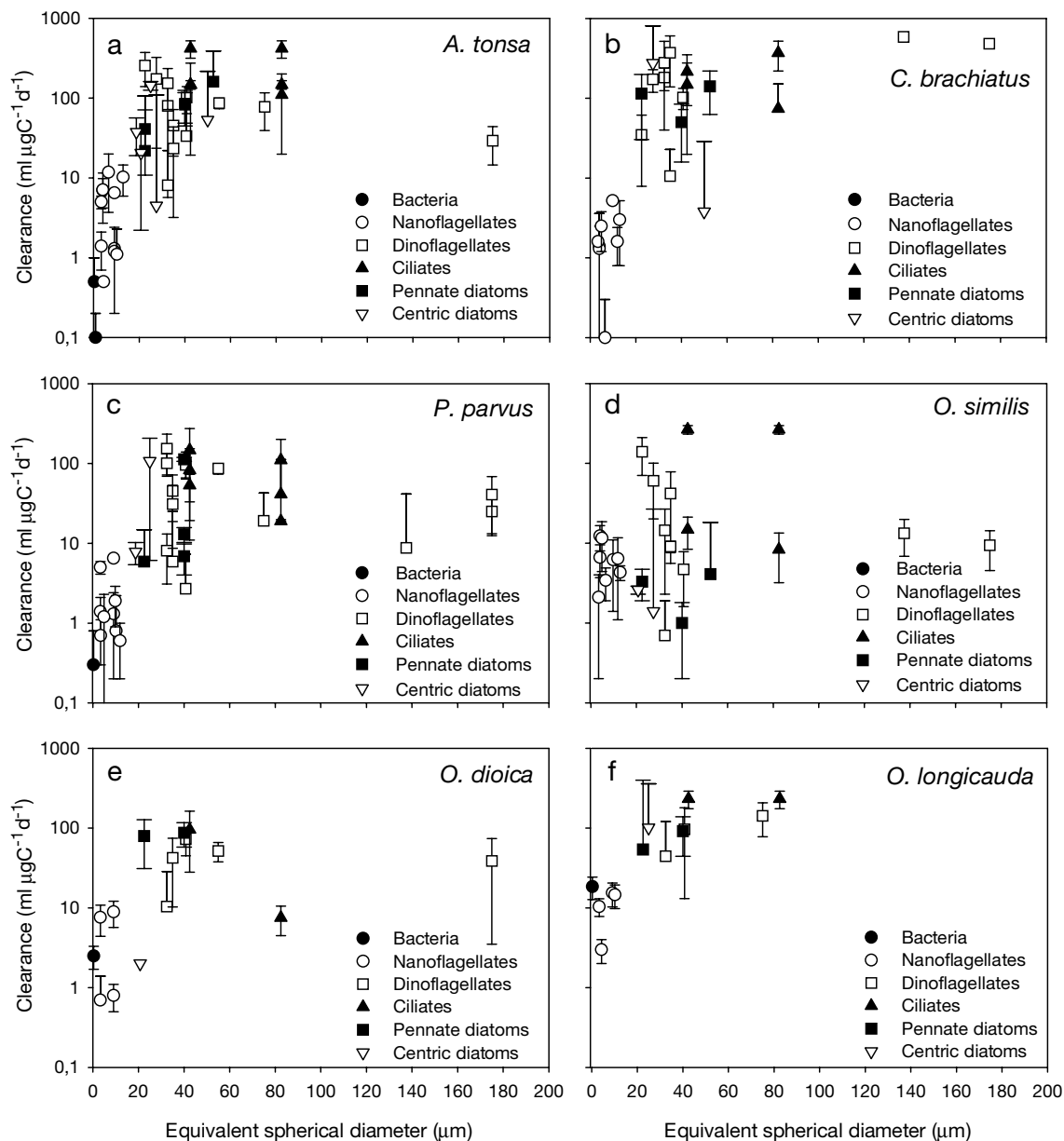


Fig. 8. Clearance by small copepods and appendicularians with different taxon-specific cell sizes of bacteria, nanoflagellates, dinoflagellates, ciliates, pennate and centric diatoms corrected to 19°C. Errors bars are  $\pm$  SE of mean. Note log scale for clearance (y-axis)

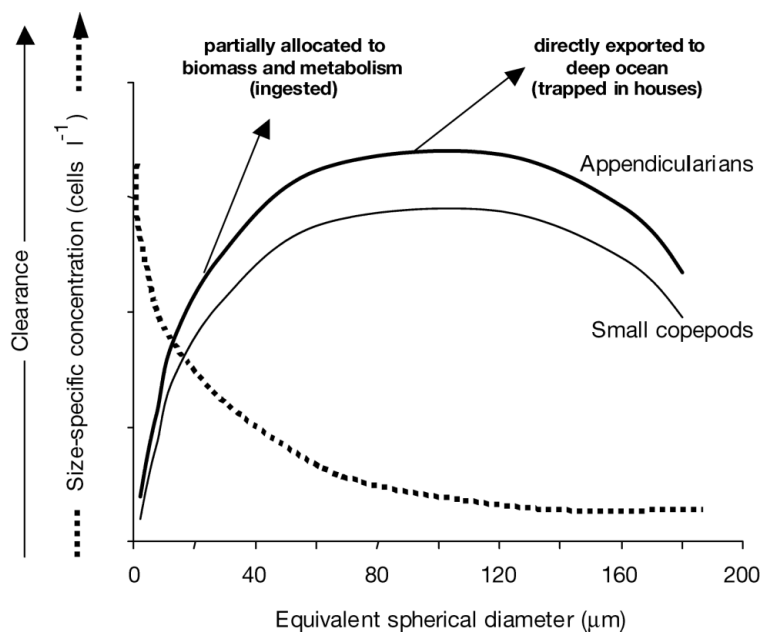


Fig. 9. Conceptual model of relationship between cell concentration and clearance rate of copepods and appendicularians in an upwelling system as a function of cell size

Our results extend the size spectrum that can be efficiently grazed by this species ( $>10 \mu\text{m}$ , Nakamura & Turner 1997), and support Nielsen & Sabatini's (1996) hypothesis that *Oithona* spp. may act as a link between small protozooplankton and fish larvae.

There is not previous information on feeding of appendicularians on natural assemblages of bacteria, microprotozoans and ciliates. The feeding of appendicularians on bacteria extends the size range of grazed particles, that were not available to copepods; the weight-specific clearance of appendicularians on bacteria and small flagellates was much higher than that of copepods. Although appendicularians feed on small cells, they may also affect larger cells through their unique feeding mode: Centric diatoms, as well as ciliates around  $80 \mu\text{m}$  ESD ( $\sim 500 \text{ ml ind. d}^{-1}$ ) were cleared at high rates by *Oikopleura longicauda*, but as *O. longicauda* has no incurrent filter, most large diatoms, dinoflagellates and ciliates would probably have been caught in the filter mesh and collected on the pharyngeal filter ( $610 \times 150 \text{ nm}$ , Deibel & Powell 1987). Although *O. dioica* cleared pennate and centric diatoms ( $20$  to  $50 \mu\text{m}$  ESD), ciliates and dinoflagellates ( $>70 \mu\text{m}$  ESD), these cells were probably trapped by the mesh of its inlet filter ( $\sim 30 \mu\text{m}$ ).

Chain-forming diatoms were not strongly selected by either copepods or appendicularians. Copepods were probably unable to feed on the entire large chains of the most abundant species found during our study: *Eucampia cornuta*, *Chaetoceros* spp., and *Guin-*

*ardia delicatula*. However, they could feed on solitary forms such as *G. delicatula* ( $25$  to  $30 \mu\text{m}$ ) during the February experiment. Schnack (1983) observed similar pattern in the northwest African upwelling area. Schnack (1983) observed that copepods were unable to feed on entire colonies of *Thalassiosira partheneia* (which has a cell size of about  $9 \mu\text{m}$  but forms colonies of up to  $5 \text{ cm}$  in length), but that they consumed cells once the colonies had disintegrated. One of the potential biases in our study could have been a 'food-chain effect' during incubations: as incubation proceeded, diatom growth in the experimental bottles was higher than that in controls, since diatoms in the experimental bottles were released from ciliate and h-dinoflagellate grazing pressure because these were consumed by copepods or appendicularians. This could have created an apparently low grazing rate on diatoms, even if some zooplankton grazing did actually occur. However, correction of our estimations by the 3 component equation template of Tang et al. (2001) resulted in higher grazing values on phytoplankton than uncorrected values (data not shown). A further bias in our experiments could

be related to the trophic interactions between nanoflagellates being eaten by ciliates, which in turn were eaten by appendicularians. However, ciliates were very scarce during our study ( $1.8 \text{ ciliates ml}^{-1}$ ). Ciliates in control bottles could potentially remove  $\sim 0.5 \mu\text{gC}$  from nanoflagellates during 15 h incubations, while the 3 appendicularians in the experimental bottles removed around  $6.3 \mu\text{gC}$  during the same time. In addition, appendicularians removed around 96% of ciliate biomass in the bottles; therefore, grazing rates on nanoflagellates were underestimated by approximately 8%. In consequence, we believe that trophic artifacts have been partially corrected in our clearance-rate estimates on autotrophic and heterotrophic prey.

#### Implications of microprotozoan–zooplankton coupling in an upwelling ecosystem

Clearance rates by *Acartia tonsa* generally decreased with increasing abundance of nanoflagellates, dinoflagellates and diatoms. Similar results were found for *Paracalanus parvus* feeding on nanoflagellates and centric diatoms. These results are in agreement with the model of the relationship between filtering rate and food concentration suggested by Marin et al. (1986), whereby clearance rate will decrease above a certain critical concentration of food. A similar pattern was found by Dagg & Walser (1987) for *Neocalanus*

*plumchrus* feeding on *Thalassiosira weissflogii*, Ohman (1987) for *N. tonsus* and Paffenhöfer (1988) for *Paracalanus* sp. Paffenhöfer (1988) suggested that the different slopes and shapes of these curves were the result of the adaptation of copepods to different trophic environments. Maximal clearance rates at different concentrations of prey suggest that copepods adapt to continuous low or high supplies of different autotrophic and heterotrophic prey of different sizes. Copepod and appendicularian ingestion of autotrophic plus heterotrophic prey typically represented more than 80% of copepod body carbon. This means that small-sized copepods and appendicularians were not food-limited, since they are able to obtain sufficient daily rations even at low *in situ* chl *a* concentrations by feeding on many particles other than phytoplankton (González et al. 2000).

Size-dependency is well documented for copepods feeding on phytoplankton (e.g. Cowles 1979). Size-differentiated clearance rates showed that clearance rates increase with increasing prey size in a hyperbolic fashion. Since large cells such as ciliates and large dinoflagellates (i.e. from 60 to 180 µm ESD) were very scarce during all our experiments, these were cleared at the highest rates, which indeed revealed a size-selectivity by copepods (Fig. 9). Similarly, Levinsen et al. (2000) observed that *Calanus* spp. generally grazed ciliates more efficiently than phytoplankton, and Frost (1972) also found an increase in clearance by *C. pacificus* with increasing size of diatoms up to the largest species examined (~67 µm ESD).

Although the appendicularian *Oikopleura dioica* feeds on small particles, it may also affect larger algae through its unique feeding mode. Therefore, there should be differences in the fate of small and large cells cleared from the bottles. For instance, while small cells such as bacteria and nanoflagellates are efficiently ingested and allocated to biomass, metabolism and fecal pellet production, cells larger than ~30 µm cannot be ingested by this appendicularian. Thus, even though *O. dioica* had high clearance rate of large cells, these cells probably accumulate and adhere to the house before this is abandoned and exported to the deep ocean depths (Fig. 9). Accumulation of cells on the houses of *O. dioica* could have important implications. Not only could house production dominate vertical carbon fluxes (Kjørboe et al. 1996, Vargas et al. 2002), but also sinking rates could be accelerated by the weight of attached particles, such as dinoflagellates, cells or chains that typically dominate the biomass in this upwelling area and that are too large for appendicularians to feed on.

Since clearance decreased with increasing food concentrations, but increased with increasing cell size, it seems that the clearance rate of copepods is a function

of food concentration and food size, and that a copepod can vary the size range of cells ingested as a function of concentration and available cell size. This selectivity model correlates well with the size-structure of heterotrophs in this environment, where small prey (nanoflagellates) are very abundant, but large prey (dinoflagellates and ciliates) are scarce. This implies that large dinoflagellates and ciliates may be under a stronger top-down regulation than autotrophs (diatoms and nanoflagellates) in this coastal upwelling ecosystem.

Our results on feeding by small calanoid and cyclopooid copepods and appendicularians suggest that heterotrophs are a significant component of their diets. Since most studies in the Humboldt Current System have considered neither the contribution of heterotrophic microprotozoans to zooplankton grazing nor their implications for carbon fluxes (e.g. Peterson et al. 1988, Castro et al. 1991, González et al. 2000, Grunewald et al. 2002), grazing experiments on heterotrophs are needed for comparison with data based on the gut-fluorescence method (Mackas & Bohrer 1976). Given the high abundance of small copepods and appendicularians in these coastal areas (González et al. 2000, Grunewald et al. 1998), omnivory would also suppress microbial food webs, and enable zooplankton to utilize small picoplankton and convert it to exportable biogenic carbon (Zeldis et al. 2002). This important link needs to be considered in future plankton studies and food-web models of upwelling ecosystems along this coastal margin.

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