

# Influence of zooplankton grazing on free dissolved enzymes in the sea

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**ABSTRACT:** In the Northern Adriatic Sea, extracellular enzymatic activity was measured during a Lagrangian study following a drifting buoy for 40 h. Dissolved free enzymatic activity represented 20 to 70 % of total activity depending on the type of enzyme.  $\alpha$ - and  $\beta$ -glucosidases exhibited a significantly higher free activity than proteolytic enzymes. In subsequent laboratory experiments we investigated the effect of zooplankton on the free enzyme pool. The 4-step approach included: (1) determination of the enzymatic activities in copepods (mainly *Acartia clausi*); (2) enzymatic activity in fecal pellets; (3) short- and long-term grazing experiments; and (4) degradability of free glucosidase in seawater.  $\alpha$ - and  $\beta$ -glucosidases, leu-aminopeptidase, lipase and chitinase were examined. Experiments in which zooplankton were selectively enriched revealed a significant increase in both particle-bound (due to the increase of bacterial density) and dissolved free enzymatic activity. Incubating water enriched in free enzymes released by zooplankton with natural bacterial consortia, we found that 70 % of the original  $\alpha$ - and  $\beta$ -glucosidase activity remained after 22 h. The presence of microorganisms did not enhance the degradation of these enzymes as compared to autoclaved controls. We found that a considerable amount of free dissolved enzymes is lost by 0.2  $\mu\text{m}$  filtration using Nuclepore filters, thereby leading to an underestimation of dissolved enzymes by ~30 % in our experiments. Based on our results we conclude that mesozooplankton contribute to the free enzymatic activity in natural waters especially during periods of high grazing activity.

**KEY WORDS:** Extracellular enzymes · Zooplankton · Dissolved enzymes · Fecal pellet · Northern Adriatic Sea

## INTRODUCTION

Extracellular enzymatic activity represents one of the most important parameters determining microbial activity in the sea (Hoppe 1983). Bacterial extracellular enzymes located on the outer bacterial membrane or in the periplasm (e.g. Chróst 1990) have received considerable attention since a major part of degradation of polymers in the sea seems to be particle-associated (Hollibaugh & Azam 1983, Hoppe 1983, Okami 1986, Rosso & Azam 1987, Billen 1991, Hoppe et al. 1993). A smaller fraction of enzymes, however, is freely dissolved in water (Reichardt et al. 1967, Priest 1984,

Vives Rego et al. 1985). Free dissolved enzymes are secreted by intact viable cells (e.g. from bacterial surfaces) or they can be liberated after cell damage as result of grazing activity such as 'sloppy feeding' by zooplankton (Meyer-Reil 1981, Azam & Ammerman 1984, Chróst 1990). Reports of free dissolved enzymatic activity are very diverse. Somville & Billen (1983) reported that most of the exoprotease activity in the eutrophic Belgian coastal zone is freely dissolved. Hashimoto et al. (1985) found that 10 to 50 % of the carboxypeptidase activity was cell-free activity. Authors of other studies such as Christison & Martin (1971) for the English channel, Vives Rego et al. (1985) for the Southern Bight of the North Sea and Rosso & Azam (1987) for the Santa Monica Basin (California), have shown that most of the enzymatic activity is related to particles in a size range of 0.2 to 0.8  $\mu\text{m}$ . Even Rosso &

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Azam (1987), while stating that most of the enzymatic activity is bacterial-bound, nevertheless found high rates of free dissolved enzymatic activity in the surface waters (upper 100 m) ranging from ~30 to ~70% of the total enzymatic activity (their Fig. 1a–c). Recently, Karner & Rassoulzadegan (unpubl.) obtained high free dissolved enzymatic activity ranging from 20 to >90% of the total enzymatic activity in 0.1  $\mu\text{m}$  filtered seawater in the Western Mediterranean Sea. In the surface waters of the Northern Adriatic Sea free dissolved enzymatic activity did not exceed 10% for  $\alpha$ - and  $\beta$ -glucosidases and for leu-aminopeptidase during summer 1991 (Karner & Herndl 1992) and ranged between 20 and 70% of total enzymatic activity in June 1992 (Rath & Herndl 1994).

Digestive enzymes have been extensively studied in copepods mainly to investigate how enzyme levels reflect their feeding activities (Hirche 1981, Mayzaud & Mayzaud 1981, Baars & Oosterhuis 1984, Head et al. 1984, Mayzaud et al. 1984). For some enzymes, correlations of their activity with the concentration and quality of food were found in calanoid copepods. Amylase activity in copepodite stage V of *Calanus finmarchicus* (Tande & Slagstad 1982) and carbohydrase activities in total zooplankton population (Mayzaud et al. 1984) showed a diel cycle, whereas it took several days to induce laminarinase in *Euphausia pacifica* and *Calanus pacificus*, when the proper substrate was available (Cox 1981). Hassett & Landry (1988) found no short-term changes in digestive enzymes, assuming a long-term acclimation of the enzyme level.

Recently the release of dissolved organic matter including enzymes by heterotrophic flagellates (Nagata & Kirchman 1992a, b, Karner et al. 1994) has been investigated. These enzymes are either digestive enzymes or remains of bacterial membranes (Nagata & Kirchman 1992b).  $\alpha$ - and  $\beta$ -glucosidase activity was correlated with cladoceran abundance in a eutrophic reservoir (Vrba et al. 1992). However, this provides no direct proof of enzyme release by cladocerans since chlorophyll *a* (chl *a*) and bacterial concentrations were positively correlated with enzymatic activity as well (Vrba et al. 1992). Moreover, most of the enzymes were found attached to particles and less than 10% were dissolved. So far, there is no information available on the release of active enzymes during grazing by metazoans. A certain but unknown amount of digestive enzymes may be released from the gut during defecation together with the undigested fraction of the food (Jumars et al. 1989). It is questionable whether all the enzymes remain in the fecal pellets, or whether they leak through the surface membrane and diffuse into the ambient water similar to the release of amino acids reported by Fuhrman (1987), Roy & Poulet (1990) and Poulet et al. (1991). Whether or not they accumulate in

the ambient water during intense grazing activity depends on the residence times of those enzymes in natural seawater. Since different methods were used for estimating enzymatic activity in zooplankton and bacteria, direct comparison between enzymatic activity of the copepods and microbiota has not been made so far.

Zooplankton and the microbial community have many carbohydrases and proteases in common, like  $\alpha$ - and  $\beta$ -glucosidases, laminarinase and aminopeptidase. In order to investigate the effect of enzymes in ecological studies, fluorescent artificial substrates have been introduced to aquatic microbial ecology (Hoppe 1983). Although there are still considerable uncertainties as to what extent these model substrates reflect cleavage of natural substrates (Chróst 1990), this method provides a powerful tool for detecting changes of microbial metabolic activity. This study is the first attempt to directly compare both microbial and metazoan-zooplankton enzymatic activity using identical methods and a minimum of experimental manipulation in order to evaluate the role of metazoans on the distribution of free dissolved enzymatic activity in the euphotic zone.

## MATERIAL AND METHODS

**Diel fluctuation of total and free dissolved enzymatic activity in the Northern Adriatic Sea.** During a Lagrangian study in the Northern Adriatic Sea in spring 1992 water samples were taken with Niskin bottles at 5 m (i.e. the position of the drifting bags) over 40 h at intervals of 4 h. In order to determine free enzymatic activity, half of the sample was filtered immediately through 0.2  $\mu\text{m}$  Nuclepore filters and the enzymatic activity in the filtrate measured as described below. The activity of the filtrate is considered to be the free dissolved enzymatic activity while the activity in the raw seawater represents the total enzymatic activity. In order to minimize cell disruption the filter was changed after allowing 20 ml to pass through and the vacuum was kept at <200 mb (see also Fuhrman & Bell 1985). All subsamples were tested for  $\alpha$ - and  $\beta$ -glucosidases, lipase, chitinase and leu-aminopeptidase (representative for most of the proteolytic activity; Delange & Smith 1971).

**Enzymatic activity in copepods.** In order to estimate enzyme activities in *Acartia clausi*, 100 freshly caught specimens (from the Northern Adriatic Sea) were collected with fine forceps and homogenized with a glass tissue grinder in 0.2  $\mu\text{m}$  filtered, autoclaved seawater in May and June 1991. In November 1992, 100 copepods of about 1 mm in length were randomly selected and either homogenized in 0.2  $\mu\text{m}$  filtered, autoclaved seawater or in Tris buffered, double-distilled water (pH = 8.2). The results obtained were identical using the 2 methods, therefore we pooled the data.

**Enzymatic activity in fecal pellets.** Between 100 and 500 fecal pellets were collected with the fecal pellet collecting device as described by Bochdansky & Herndl (1992a) and washed 3 times with 0.2  $\mu\text{m}$  filtered, autoclaved seawater prior to homogenizing them with a precombusted glass tissue grinder in a known volume of autoclaved seawater. Enzymatic activity was estimated for  $\alpha$ - and  $\beta$ -glucosidase, leu-aminopeptidase, lipase and chitinase using fluorescent substrates as described below. Prior to homogenizing the fecal pellets, a subsample of the water from the last rinse was taken and its enzymatic activity subtracted from the activity measured in the fecal pellet suspension.

**Enzyme release by zooplankton.** In order to determine the release of enzymes by zooplankton, seawater from the Northern Adriatic Sea was amended with a mixed zooplankton community (>200  $\mu\text{m}$ ) mainly consisting of copepods and incubated in rolling tanks under dim light for 3 to 8 h. The zooplankton sample was rinsed 3 times in 0.2  $\mu\text{m}$  filtered seawater and subsequently transferred into a plankton-splitter. Half of the sample was immediately poured into a rolling tank; the second half was filtered through a net with a mesh size of 200  $\mu\text{m}$  (to remove the zooplankters) and then added to the 0.2  $\mu\text{m}$  filtered water in the control tanks. This procedure ensured identical conditions of the water in both treatments. Together with the zooplankton we removed or added attached bacteria but assumed that their contribution to overall enzymatic activity was negligible. After incubation, the zooplankton were removed and a subsample of the water was filtered through 0.2  $\mu\text{m}$  Nuclepore polycarbonate filters again in order to distinguish free from total enzymatic activity.

To determine the influence of zooplankton on the enzymatic activity in the water, 200 individuals of *Acartia clausi* were placed into a 1 l rolling tank filled with about 800 ml 65  $\mu\text{m}$  filtered seawater from the Northern Adriatic Sea. Tanks filled with water from the same site but without copepods served as controls. All incubations were performed in triplicates over a period of 3 d. The experiments were performed under continuous illumination of 60  $\mu\text{E m}^{-2} \text{s}^{-1}$  at 18°C. During the course of incubation, the oxygen supply was sufficient due to the large air space in the rolling tanks. The mortality of *A. clausi* was less than 8% in all of the 3 tanks after 3 d although the copepods probably suffered from shortage of food. Subsamples for chl *a*, pheopigment, primary production, bacterial numbers, bacterial secondary production and enzymatic activity ( $\alpha$ - and  $\beta$ -glucosidase, leu-aminopeptidase) were collected daily. Primary production and bacterial production were estimated by  $\text{NaH}^{14}\text{CO}_3$  and [methyl- $^3\text{H}$ ]-thymidine incorporation (Parsons et al. 1984), respectively, and bacteria were enumerated by acridine orange direct counting (Hobbie et al. 1977).

At the beginning and at the end of the experiment, subsamples were withdrawn to determine the dry weight of the particulate material and the concentration of monomeric dissolved carbohydrates using the 3-methyl-2-benzothiazolinone hydrazone hydrochloride (MBTH) assay. For all analytical procedures (except the enzyme assay) we followed the instructions given in Parsons et al. (1984).

In order to evaluate the role of defecation on enzyme release by net-zooplankton, a mixed zooplankton sample (starved for ~1 h) was added to 0.2  $\mu\text{m}$  sterile-filtered seawater together with glass beads (1 to 20  $\mu\text{m}$  in diam.). The glass beads served as an inert model to simulate gut passage of food; control vessels to which no glass beads were added served as control. After a 3.2 and a 4 h incubation period, zooplankton were removed and the total and dissolved free enzymatic activity determined as described below.

Since larval zooplankton has been shown to occur in high densities in marine snow in the Northern Adriatic Sea, we tried to evaluate the influence of this larval zooplankton on the overall extracellular enzymatic activity in marine snow (Bochdansky & Herndl 1992b). One major component of the zooplankton found in the aggregates were polychaete larvae of *Prionospio* sp. in the Nectochaeta stage which occurred in densities of 200 ind.  $l^{-1}$  marine snow (Bochdansky & Herndl 1992b). All zooplankton larger than ~60  $\mu\text{m}$  were removed from the marine snow samples by means of a mouth-pipette under a dissecting microscope. The marine snow was subsequently transferred into 8 sterile petri dishes each containing a final volume of 10 ml. Ten *Prionospio* sp. larvae were added to 4 petri dishes. The 4 remaining petri dishes served as controls. After 4 h in darkness at 24°C, which is roughly the surface water temperature of the Northern Adriatic Sea in summer, the polychaete larvae were removed and the total enzymatic activity was determined as described below.

#### **Degradability of free dissolved enzymes in seawater.**

In order to test the degradability of free dissolved enzymes, water from a tank was taken in which a mixture of zooplankton (ca 100 ind.  $l^{-1}$ ) was kept overnight. After removing the zooplankton with a 200  $\mu\text{m}$  net, the water was filtered first through a Whatman GF/F glass-fiber filter and then through a 0.2  $\mu\text{m}$  Nuclepore polycarbonate filter in order to obtain a sterile, particle-free water. One part of this water was added to 3 parts of: (1) freshly collected raw seawater; (2) autoclaved seawater; and (3) 0.2  $\mu\text{m}$  (Nuclepore, polycarbonate) filtered, autoclaved seawater. Free dissolved (again filtered through 0.2  $\mu\text{m}$  Nuclepore after the incubation period) and total enzymatic activities of  $\alpha$ - and  $\beta$ -glucosidase together with bacterial numbers were determined after 1, 2, 4.5, 7 and 22 h of incubation.

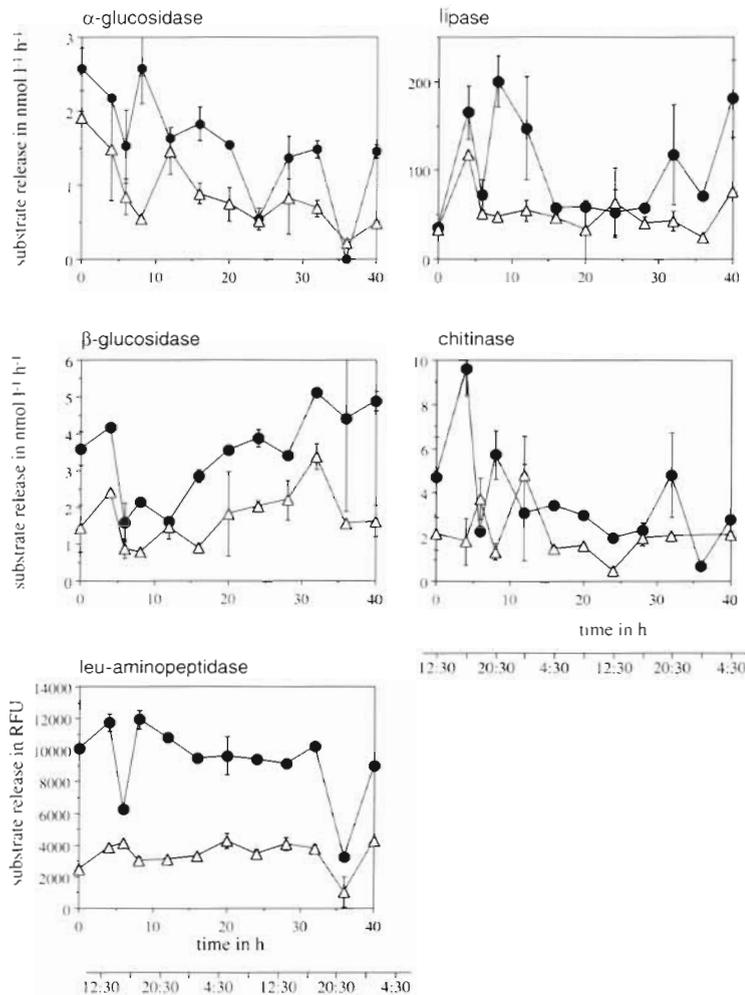


Fig. 1. Free dissolved ( $\Delta$ ) and total enzymatic activity ( $\bullet$ ) during a Lagrangian study in the Northern Adriatic Sea in June 1992. A drifting buoy deployed at 5 m depth about 18 nautical miles east of the Po river delta was followed over a period of 40 h. Leu-aminopeptidase activity given as relative fluorescence units (RFU). Abscissae show time elapsed (in h) and time of day of the measurement. Bars indicate range of the 2 subsamples

**Determination of enzymatic activity.** Extracellular enzymatic activity was measured using the following fluorescent artificial substrates: 4-methyl umbelliferyl  $\alpha$ -D-glucopyranoside and 4-methyl umbelliferyl  $\beta$ -D-glucopyranoside for estimating  $\alpha$ - and  $\beta$ -glucosidase activity respectively, 4-methyl umbelliferyl butyrate for lipase activity, 4-methyl umbelliferyl *n*-acetyl-D-glucosaminide for chitinase and L-leucine 7-amino-4-methyl coumarin for proteolytic activity. For details of this method see Karner & Herndl (1992). Briefly, the artificial substrates were added to the respective water or suspension at 125  $\mu$ M final concentration (2.5  $\mu$ M for experiments with polychaete larvae) and incubated at *in situ* temperature for 1 h. This concentration was found to ensure maximum fluorescent yield. In the

samples the increase in fluorescence over time was determined and calibrated with known concentrations of 4-methyl umbelliferylone (MUF). Since we had no suitable standard for leu-aminopeptidase, its activity is expressed as relative fluorescence unit (RFU).

## RESULTS

### Diel fluctuation of free dissolved enzymatic activity in the Northern Adriatic Sea

The fluctuation of total and free enzymatic activity of the 5 enzymes tested during the Lagrangian study is depicted in Fig. 1. The percentage of free dissolved activity ranged from 20 to more than 100% of the total activity depending on the type of enzymes. The contribution of free enzymatic activity to overall activity was lowest in leu-aminopeptidase and highest in chitinase (Fig. 1).

### Enzymatic activity in copepods

The enzymatic activities of freshly collected *Acartia clausi* (June and July) and similar-sized copepods (November) are shown in Table 1. Leu-aminopeptidase activity was 2 orders of magnitude higher in the November samples than in either June or July; all the other enzymes were within the same order of magnitude despite the difference in temperature among the different sampling dates.

### Enzymatic activity in fecal pellets

Although there was enzymatic activity detectable in the fecal pellet suspension, it never exceeded the activity in the water with which the fecal pellets were rinsed in any of the 7 measurements. Fecal pellet-bound enzymatic activity was therefore not detectable with our method.

### Enzyme release by zooplankton

In the long-term experiment (over 3 d), bacterial growth as well as enzymatic activity increased (Fig. 2). Bacterial density approximately doubled in the cope-

Table 1. *Acartia clausi* and other copepods. Activity of different enzymes in 100 freshly caught and homogenized *A. clausi* collected in June and July and similar-sized (ca 1 mm) copepods in November. Values expressed in nmol 4-methyl umbelliferone (MUF) released ind.<sup>-1</sup> h<sup>-1</sup> (mean  $\pm$  SD); leu-aminopeptidase in relative fluorescence units (RFU) ind.<sup>-1</sup> h<sup>-1</sup>

Enzyme	1–6 June 1991 (n = 6, 15.6°C, <i>in situ</i> temp.)	2–6 July 1991 (n = 5, 24.1°C, <i>in situ</i> temp.)	13 November 1992 (n = 4, 21.0°C, room temp.)
$\alpha$ -Glucosidase	0.001 ( $\pm$ 0.00058)	0.002 ( $\pm$ 0.0025)	0.006 ( $\pm$ 0.0035)
$\beta$ -Glucosidase	0.001 ( $\pm$ 0.00059)	0.009 ( $\pm$ 0.0092)	0.004 ( $\pm$ 0.0021)
Leu-aminopeptidase	0.024 ( $\pm$ 0.0044)	0.053 ( $\pm$ 0.031)	5.6 ( $\pm$ 4.0)
Lipase	–	0.152 ( $\pm$ 0.15)	0.32 ( $\pm$ 0.040)
Chitinase	0.005 ( $\pm$ 0.0016)	0.015 ( $\pm$ 0.013)	0.009 ( $\pm$ 0.0032)

pod tanks, whereas it was rather stable in the control tanks. Bacterial growth rate did not vary significantly between copepod and control tanks and remained relatively constant ranging from 1 to 2.4 d<sup>-1</sup>. As indicated by the declining pigment concentrations, algal biomass was grazed efficiently in the copepod tanks resulting in higher pheopigment than chl *a* concentrations (Fig. 2). In the control tanks, algal biomass increased within the first day and declined thereafter. Specific primary production (production per unit chl *a*) was highest at the beginning of the experiments and declined during the course of the incubation. Mean specific primary production was significantly higher in the copepod tanks than in the controls (Mann-Whitney,  $p < 0.01$ ,  $n = 12$ ; Fig. 2). The concentration of particulate dry weight increased in the control tanks during the 3 d incubation period, whereas it remained more or less constant in the copepod tanks. Dissolved monomeric carbohydrates remained approximately at the same level in both treatments indicating that released monomers are rapidly consumed by the microbiota. Particle-bound enzymatic activity was significantly higher in the copepod tanks (Mann-Whitney,  $p < 0.01$ ,  $n = 12$ ), whereas it remained rather constant in the controls. Also, free dissolved enzymatic activity was significantly higher in the copepod tanks than in the controls (Mann-Whitney,  $p < 0.01$ ,  $n = 12$ ). Free dissolved  $\alpha$ -glucosidase activity ranged from 4 to 11% of the total activity in both controls and copepod tanks. Free  $\beta$ -glucosidase amounted to 0.6–5.1% of the total activity in the control tanks and to 6–10% in the copepod tanks. Dissolved free leu-aminopeptidase activity comprised up to 86% of the total activity in the copepod tanks (Fig. 2), whereas in the controls the highest value was 60% of the total activity at the beginning of the experiment.

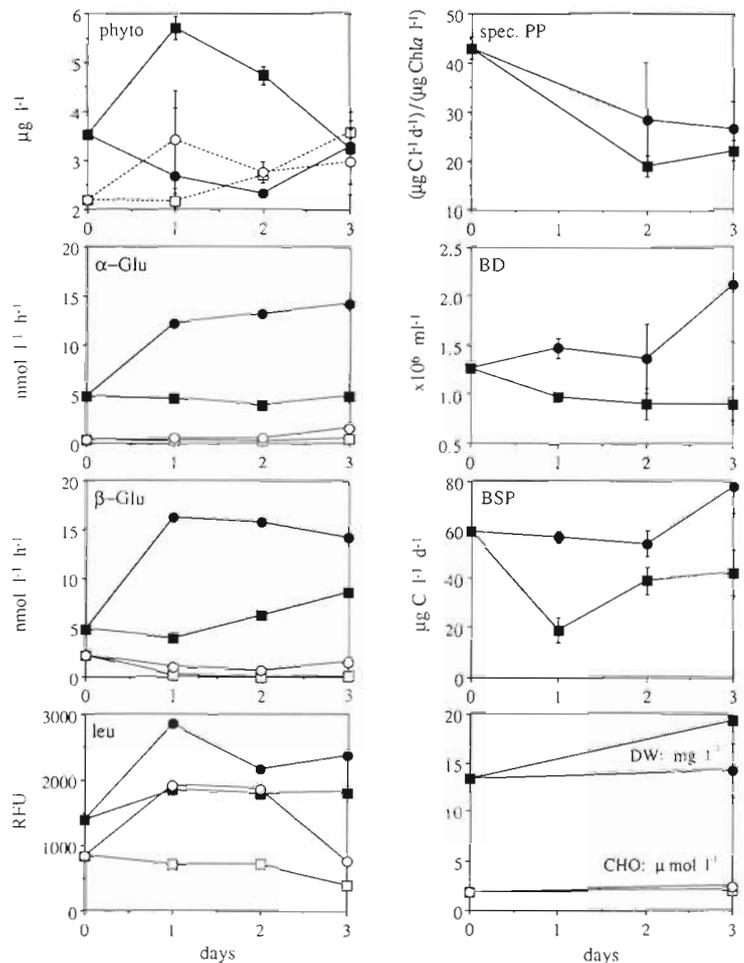


Fig. 2. *Acartia clausi*. Long-term experiments in which *A. clausi* was incubated in natural seawater (○, ●) as compared to natural seawater without copepods added (□, ■). Phyto: phytoplankton biomass as indicated by pigment levels, i.e. chl *a* (open symbols) and pheopigments (filled symbols).  $\alpha$ -Glu:  $\alpha$ -glucosidase;  $\beta$ -Glu:  $\beta$ -glucosidase; leu: leucine-aminopeptidase activity. Open symbols indicate free dissolved and filled symbols total enzymatic activity. Spec. PP: specific primary production (primary production per unit of chl *a*); BD: bacterial density; BSP: bacterial secondary production; DW and CHO: dry weight and dissolved monomeric carbohydrates, respectively, measured only at start and end of experiment. Bars represent SD of 3 replicate experiments with 2 subsamples each

Table 2. Development of bacterial biomass, free dissolved and total enzymatic activity in zooplankton (>200  $\mu\text{m}$ ) enriched batch cultures and in raw seawater controls without zooplankton >65  $\mu\text{m}$  added. n: number of zooplankton organisms (>200  $\mu\text{m}$ )  $\text{l}^{-1}$  seawater; D: duration of the experiment in h;  $t_0$ ,  $t_1$ : values at beginning and end of experiment, respectively, for (C) controls and (Z) zooplankton enriched tanks; Bac: bacterial numbers  $\times 10^3 \text{ ml}^{-1}$ ; Fd EA: free dissolved enzymatic activity; total EA: total enzymatic activity in nmol substrate released  $\text{l}^{-1} \text{ h}^{-1}$  (leu-aminopeptidase in RFU  $\text{ind}^{-1} \text{ h}^{-1}$ ); + gb and - gb: with and without addition of glass beads, respectively. All values are means of duplicate measurements; 0: not detectable

Zooplankton added	Type of water	n	D	$t_0$			$t_1(\text{C})$			$t_1(\text{Z})$		
				Bac	Fd EA	Total EA	Bac	Fd EA	Total EA	Bac	Fd EA	Total EA
<b><math>\alpha</math>-Glucosidase</b>												
Mixed	Natural seawater	417	2.5	217	–	0.274	263	0	0.250	288	0.576	0.756
Mixed	Natural seawater	481	4	–	–	–	245	0.153	0.548	296	1.173	1.614
Mixed	Natural seawater	850	6	67	0.249	0.956	64	0.665	2.119	78	3.905	6.107
Mixed	Natural seawater	602	7	327	0.226	0.422	395	0.869	0.172	443	0.706 <sup>c</sup>	7.134
Mixed	Filtered – gb	650	4	–	0	–	–	–	–	18	1.031	1.307
Mixed	Filtered + gb	650	4	–	0	–	–	–	–	24	0.831	0.831
Mixed	Filtered – gb	1236	3.2	–	–	–	5.4	0.625	0.546 <sup>b</sup>	5.7	2.186	1.952 <sup>b</sup>
Mixed	Filtered + gb	1236	3.2	–	–	–	5.4	0.625	0.546 <sup>b</sup>	5.6	2.030	2.108 <sup>b</sup>
Polychaete larvae	Marine snow <sup>a</sup>	500	4	–	–	4.25 ( $\pm 0.43$ )	–	–	15.45 ( $\pm 2.64$ )	–	–	17.41 ( $\pm 1.78$ )
<b><math>\beta</math>-Glucosidase</b>												
Mixed	Natural seawater	417	2	217	0.025	0.189	263	–	0.030	288	0.176	0.527
Mixed	Natural seawater	481	4	–	–	–	245	0.491	0.951	296	0.764	1.356
Mixed	Natural seawater	850	6	67	0	0.955	64	0.241	1.528	78	1.487	2.694
Mixed	Natural seawater	602	7	327	0.115	0.562	395	0.462	1.605	443	1.363	2.618
Mixed	Filtered – gb	650	4	–	0.159	–	–	–	–	18	1.065	1.201
Mixed	Filtered + gb	650	4	–	0.159	–	–	–	–	24	1.193	0.800
Mixed	Filtered – gb	1236	3.2	–	–	–	5.4	0.461	1.469 <sup>b</sup>	5.7	2.073	2.227 <sup>b</sup>
Mixed	Filtered + gb	1236	3.2	–	–	–	5.4	0.461	1.469 <sup>b</sup>	5.6	1.920	2.227 <sup>b</sup>
Polychaete larvae	Marine snow <sup>d</sup>	500	4	–	–	5.22 ( $\pm 0.47$ )	–	–	13.68 ( $\pm 3.44$ )	–	–	21.35 ( $\pm 3.11$ )
<b>I-Amino-peptidase</b>												
Mixed	Natural seawater	417	2	217	163.08	727.70	263	143.42	722.42	288	257.40	685.93
Mixed	Natural seawater	481	4	–	–	–	245	319.71	1105.07	296	482.55	1067.10
Mixed	Natural seawater	850	6	67	0	1830	64	230	2684	78	1051	4325
Mixed	Natural seawater	602	7	327	130.76	462.46	395	305.10	575.93	443	446.19	1890.87
Mixed	Filtered – gb	650	4	–	1815.09	–	–	–	–	18	3550.11	3489.55
Mixed	Filtered + gb	650	4	–	1815.09	–	–	–	–	24	2682.26	3306.80
Mixed	Filtered – gb	1236	3.2	–	–	–	5.4	642	358 <sup>b</sup>	5.7	1177	1196 <sup>b</sup>
Mixed	Filtered + gb	1236	3.2	–	–	–	5.4	642	358 <sup>b</sup>	5.6	819	716 <sup>b</sup>
Polychaete larvae	Marine snow <sup>d</sup>	500	4	–	–	1736 ( $\pm 269$ )	–	–	1934 ( $\pm 308$ )	–	–	5133 ( $\pm 660$ )
<b>Lipase</b>												
Mixed	Natural seawater	417	2	217	14.07	13.96	263	17.34	29.91	288	31.47	56.50
Mixed	Natural seawater	481	4	–	–	–	245	22.20	16.91	296	32.02	77.68
Mixed	Natural seawater	850	6	67	12.29	39.15	64	18.88	33.46	78	97.21	121.16
Mixed	Natural seawater	602	7	327	–	–	395	27.87	28.01	443	109.37	126.61
Mixed	Filtered – gb	650	4	–	24.82	–	–	–	–	18	30.64	57.91
Mixed	Filtered + gb	650	4	–	24.82	–	–	–	–	24	43.37	51.11
Mixed	Filtered – gb	1236	3.2	–	–	–	5.4	17.13	7.39 <sup>b</sup>	5.7	27.88	11.86 <sup>b</sup>
Mixed	Filtered + gb	1236	3.2	–	–	–	5.4	17.13	7.39 <sup>b</sup>	5.6	32.86	9.51 <sup>b</sup>
<b>Chitinase</b>												
Mixed	Natural seawater	417	2	217	0.051	0.048	263	0	0.348	288	0.281	0
Mixed	Natural seawater	481	4	–	–	–	245	0	0.138	296	0.327	0.465
Mixed	Natural seawater	850	6	67	0	0.125	64	0	0.645	78	0.817	1.461
Mixed	Natural seawater	602	7	327	0.051	0.048	395	0	0.348	443	0.281	0
Mixed	Filtered – gb	650	4	–	2.430	–	–	–	–	18	1.925	2.726
Mixed	Filtered + gb	650	4	–	2.430	–	–	–	–	24	2.703	2.243
Mixed	Filtered – gb	1236	3.2	–	–	–	5.4	0.439	0.659 <sup>b</sup>	5.7	1.317	2.927 <sup>b</sup>
Mixed	Filtered + gb	1236	3.2	–	–	–	5.4	0.439	0.659 <sup>b</sup>	5.6	0.805	1.171 <sup>b</sup>

<sup>a</sup> 4 replicates  $\pm$  SD

<sup>b</sup> Triton X-100 was used to determine total enzymatic activity. Triton X-100 is not applicable for estimating total lipase activity. Micelles were presumed formed which prevented the enzyme from complexing and therefore strongly reduced lipase-artificial substrate affinities

<sup>c</sup> The only occurrence in which free dissolved enzymatic activity was higher in the control tanks than in the copepod tanks

With one exception (indicated by 'c' in Table 2,  $\alpha$ -glucosidase) free dissolved enzymatic activity was higher in the batch cultures with copepods added than in the controls (Table 2). During short incubation periods ( $\leq 7$  h) bacterial biomass increased slightly, but was most likely not sufficient to explain the strong increase of enzymatic activity in the copepod tanks assuming that the 'per cell activity' of the bacteria did not change dramatically. In the experiments where glass beads were offered to copepods (Table 2), release of enzymes was not enhanced although fecal pellets containing glass beads were abundant. In filtered seawater, copepods also produced fecal pellets best described as 'ghost pellets' (Lampitt et al. 1990) without any visible content. Since fecal pellets might have been disrupted by the copepods during the experiment in the rolling tanks, no attempt was made to quantify the amount of fecal pellets produced. Enzymatic activity levels were higher in the copepod tanks than in the controls (Table 2).

As shown in Table 2 for the experiments with polychaete larvae,  $\alpha$ - and  $\beta$ -glucosidase and leu-amino-peptidase activity increased in both the controls and in marine snow containing the larvae. Total enzymatic activity, however, was significantly higher in the presence of the larvae than in the controls (Mann-Whitney,  $p < 0.1$  for all 3 enzymes tested,  $n = 4$ ). No distinction was made between free and particle-bound activity. Parallel experiments run separately but with the same experimental setup showed no changes in bacterial numbers or production rates during the incubation period (unpubl. data).

#### Degradability of free dissolved enzymes in seawater

As shown in Fig. 3, total as well as free dissolved  $\alpha$ - and  $\beta$ -glucosidase activity declined rapidly within 1 h after starting the experiment but remained at this level for 22 h in autoclaved and in 0.2  $\mu\text{m}$  filtered seawater. After 22 h, total enzymatic activity increased in natural seawater. Filtration through 0.2  $\mu\text{m}$  filters led to a significant reduction of enzymatic activity in all treatments (Student  $t$ -test,  $p < 0.05$ ,  $n = 36$ ). With the exception of  $t_5$  (22 h), there was no significant difference in enzymatic activity between natural seawater, the 0.2  $\mu\text{m}$  and autoclaved controls (Fig. 3). Bacterial growth provoked an increase in total  $\alpha$ - and  $\beta$ -glucosidase activity in the treatment with natural seawater (Fig. 4) since the enzyme activity in neither the 0.2  $\mu\text{m}$  filtered and autoclaved seawater, nor in the autoclaved seawater, followed this trend. Bacterial densities at the end of the incubation period

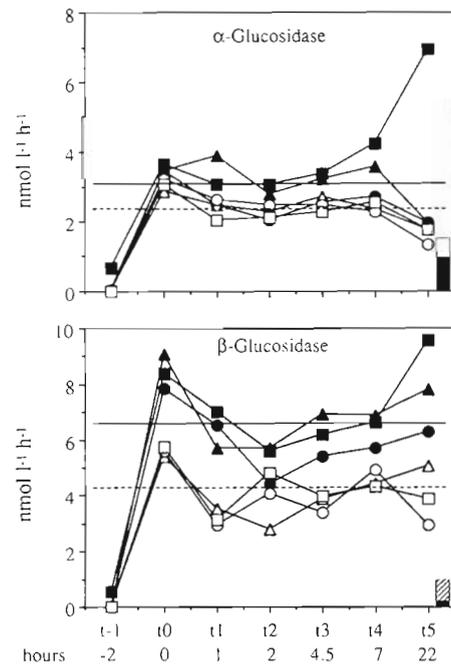


Fig. 3. Time course of  $\alpha$ - and  $\beta$ -glucosidase activity. Sterile water containing high concentrations of copepod-derived enzymes was added at  $t_0$  to autoclaved seawater ( $\Delta$ ,  $\blacktriangle$ ), 0.2  $\mu\text{m}$  filtered, autoclaved seawater ( $\circ$ ,  $\bullet$ ) and raw seawater ( $\square$ ,  $\blacksquare$ ). Total enzymatic activity indicated by filled symbols; open symbols stand for free dissolved enzymatic activity. Continuous line indicates mean total enzymatic activity of all 3 treatments from  $t_0$  to  $t_4$ . Dashed line represents mean enzymatic activity after additional 0.2  $\mu\text{m}$  filtration. On average, 0.2  $\mu\text{m}$  filtration reduced enzymatic activity by 27% ( $\alpha$ -glucosidase) and 35% ( $\beta$ -glucosidase). For comparison, bar on right shows free dissolved enzymatic activity (light section) and particle-bound enzymatic activity (dark section) in the original seawater at  $t_6$  without addition of the enzyme-enriched water

( $t_5$ ) were  $1.65 \times 10^3$  cells  $\text{ml}^{-1}$  in the 0.2  $\mu\text{m}$  filtrate and  $0.85 \times 10^3$  cells  $\text{ml}^{-1}$  in the autoclaved seawater, corresponding to 0.2% of the bacteria present in the natural seawater.

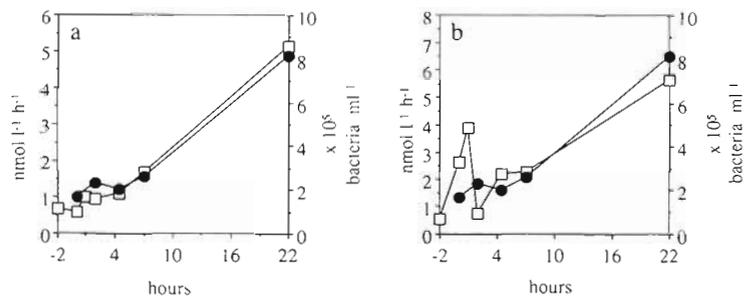


Fig. 4. Development of bacterial density ( $\bullet$ ) and particle-bound ( $\square$ ) activity of (a)  $\alpha$ -glucosidase and (b)  $\beta$ -glucosidase in natural seawater. Particle-bound activity calculated by subtracting free dissolved enzymatic activity in the 0.2  $\mu\text{m}$  filtrate from the total enzymatic activity

## DISCUSSION

During the Lagrangian study variations in enzymatic activity were high although we were following the same water body using a drifting buoy (Fig. 1). The course of temperature and salinity during this study, however, indicated mixing with more oligotrophic waters from the eastern part of the Northern Adriatic Sea. Free dissolved enzymatic activity always comprised a substantial portion of the total enzymatic activity. The percentage of free activity was lower for leu-aminopeptidase than for the other enzymes. Occasionally, free enzyme levels exceeded total chitinase and lipase activities. Similar observations were made by Rosso & Azam (1987) in sediment-pore water, who attributed this to 'blank problems'.

The fluorogenic substrate analogs used in this and other studies seem to be suitable for estimating enzymatic activity in zooplankton and, in addition, for allowing direct comparison of enzymatic activity of microorganisms and metazoans. All 5 enzymes investigated were found to be associated with copepods. Nevertheless, the pattern of activity and the relative importance of the enzymes is quite different: while  $\alpha$ - and  $\beta$ -glucosidase and proteolytic activity have previously been found in copepods, lipase activity has not received as much attention and was assumed to play only a minor role in energy gathering processes (Mayzaud & Mayzaud 1981). In fact, we found a higher level of lipase activity in copepods than in seawater (Fig. 1). Lipase might be involved in the degradation of lipid storage products and does not necessarily reflect digestive processes. We found relatively high chitinase activity in copepod homogenates; we could not distinguish, however, between chitinase produced by the copepod glands or possibly by 'enteric' bacteria. Since no chitinase activity was detectable in fecal pellets, we assume that the measured activity is derived from the copepod itself and not from the contents of the gut. This is supported by the notion that free chitinase activity is associated with molting processes of many arthropods (Smucker & Kim 1991). Whether chitinase is involved in digestive processes still remains questionable (Lampitt et al. 1990). Even for  $\alpha$ - and  $\beta$ -glucosidases and for leu-aminopeptidase, which are definitely digestive enzymes, measurements of zooplankton homogenates potentially include both enzymes from the gland tissues and from the digestive tract (Baars & Oosterhuis 1984).

In copepods, B-cells of the vacuolar region of the mid-gut burst and hydrolytic enzymes are released into the lumen (Nott et al. 1985). This mechanism mediates the extracellular digestion of food in the gut which continues within the fecal pellet produced in the

distal region. Therefore, the products of hydrolysis will have to diffuse rapidly through the surface membrane of the fecal pellet (Nott et al. 1985). As demonstrated in this study enzymes are not retained by this membrane either, since the enzymatic activity inside the fecal pellet was not higher than in the water used for rinsing the pellets; Cox (1981) arrived at a similar conclusion. Since bacteria are scarce in freshly egested fecal pellets (Gowing & Silver 1983, Jacobsen & Azam 1984) their contribution to overall activity can only be small, if not negligible. Obviously, enzymes are rapidly dispersed into the ambient water similar to the release of undigested dissolved organic material and the non-adsorbed end products of hydrolysis (Jumars et al. 1989).

With one exception, the water containing zooplankton exhibited consistently higher free dissolved enzymatic activity (Table 2, Fig. 2). Therefore, the question arises whether the dissolved enzymes originate from cell damage due to sloppy feeding and digestion or whether they are derived from the copepods. Experiments with zooplankton kept without food showed that enzymatic activity in the ambient water is elevated in the presence of zooplankton even when no food was offered indicating substantial release of digestive enzymes (Table 2).

We did not attempt to quantify the enzyme release per zooplankton since we were using natural seawater under varying conditions. Complex interactions on the microbial level, especially between phytoplankton, bacteria and their microzooplankton grazers (e.g. heterotrophic flagellates; Karner et al. 1994) and depletion of food prevent serious calculations on the actual enzyme release per copepod. We also did not know the nutritional status of the freshly-collected zooplankton, and enzyme levels have been shown to be dependent on the feeding history of the marine animals (Oosterhuis & Baars 1985). Nevertheless, the separate treatment of every experiment and the comparison of the enzymatic activity of the water in which zooplankton were added with controls lacking zooplankton indicate that zooplankton were responsible for a significant increase in free dissolved enzymatic activity even during short-term incubations. Bacterial growth had little effect during short term experiments ( $\leq 7$  h) and certainly did not contribute to the enzymatic activity detected in the 0.2  $\mu\text{m}$  filtered seawater (Table 2).

Although zooplankton grazing caused an increase in enzymatic activity, no direct correlation was found between enzymatic activity and the number of zooplankters added or the incubation period; the only exception was free  $\beta$ -glucosidase activity which was directly proportional to the number of zooplankton in the experiments (Table 2,  $r^2 = 0.837$ ,  $n = 8$ ). Vrba et al.

(1992) found close correlations of the abundance of copepods and  $\beta$ -N-acetylglucosaminidase in a reservoir, and recently Karner et al. (1994) showed increased free dissolved enzymatic activity in relation to the production of heterotrophic flagellates. In our experiments, most likely the physiological conditions of the copepods (including feeding history) and/or the composition of the food had a major influence on the amount of enzymes released.

During the 3 d experiment (Fig. 2), zooplankton grazing reduced phytoplankton biomass within 1 d, releasing probably significant amounts of readily utilizable material, which promoted bacterial growth. At the beginning of the experiment the percentage of free enzymatic activity was low but increased within 1 d to levels significantly higher than in the controls. Especially the amount of free protease activity increased drastically during the first day and reached a level equal to total enzymatic activity in the controls. Whether this increase was related to high release of protease by copepods or was caused by bacteria remains unclear. However, since bacterial numbers and growth rates increased while free dissolved leu-aminopeptidase decreased again at Day 3, it is more likely that the grazing activity of copepods was responsible for the high free proteolytic activity. On the other hand,  $\alpha$ - and  $\beta$ -glucosidase activity were mainly particle-bound and therefore probably of bacterial origin (Fig. 4). Similar increase of bacterial biomass and production in the presence of zooplankton has been observed by others (Eppley et al. 1981, Roman et al. 1988). In contrast to Roman et al. (1988), however, the bacterial growth rates in our experiments remained rather constant.

The enzyme degradation experiment with  $\alpha$ - and  $\beta$ -glucosidases indicates that: (1) 0.2  $\mu$ m filtration through Nuclepore filters reduces the amount of free dissolved activity recovered; (2) about 70% of the initial enzymatic activity remained after 22 h; and (3) the presence of microorganisms does not enhance degradation of dissolved enzymes significantly compared to the sterile controls. Filtration through 0.2  $\mu$ m polycarbonate filters reduced the amount of dissolved enzymes recovered by 27% (for  $\alpha$ -glucosidase) and by 35% (for  $\beta$ -glucosidase). This strongly indicates adsorption of dissolved enzymes on Nuclepore filters. Adsorption of enzymes on filter surfaces can introduce substantial errors especially at low enzymatic activity. While this mechanism leads to an underestimation of free dissolved enzymes, possible cell damage and enzyme release from the cell surface during the filtration process can potentially cause overestimation of free enzymatic activity. Besides direct adsorption on the filter surface, attachment to any kind of small particles and removal of those by means of 0.2  $\mu$ m filtra-

tion could be another explanation for the loss of enzymes. Adsorption of enzymes to these particle surfaces has not been studied in detail (Hoppe 1991), but there is evidence that bacterial enzymes are embedded in an exopolymeric matrix (Decho 1990). Free dissolved enzymes may become attached to this matrix forming complexes similar to enzyme-humic complexes observed in soils (Chróst 1990, Lähdesmäki & Piispanen 1992). Another potential mechanism of incorporation of enzymes into particle-like complexes was mentioned by Nagata & Kirchman (1992b): digestive enzymes could be trapped within partially degraded bacterial membranes which act as micelles (liposomes). Such bound enzymes may be better protected from proteolysis than those freely dissolved. Therefore, not all enzymes retained by ultrafiltration are necessarily remains of enzymes from cell fragments as Vives Rego et al. (1985) suggested. A large fraction of the initially dissolved enzymes (also those of zooplankton origin) may be trapped by particles including colloids or liposomes which pass through 0.2  $\mu$ m filters but not ultrafiltration. This would also explain why initially dissolved digestive enzymes of flagellate origin can be sedimented by ultracentrifugation (Nagata & Kirchman 1992b). The unbound, dissolved enzymes would consequently be degraded faster. The rapid degradation of a certain amount of  $\alpha$ - and  $\beta$ -glucosidase activity during the first hour (Fig. 3) seems to support the hypothesis of 2 fractions of dissolved enzymes: a more labile one which is degraded rapidly and a more stable fraction which remains active for at least 1 d.

In summary, we have demonstrated that artificial fluorogenic substrate analogs are suitable for direct comparison of the enzyme levels in zooplankton and bacteria. Our experiments showed elevated enzymatic activities in seawater in the presence of zooplankton, due to the increase in the carrying capacity of the microbial community by releasing readily utilizable material and by direct release of enzymes via defecation or 'sloppy feeding'. We have shown that free  $\alpha$ - and  $\beta$ -glucosidase are degraded slowly and could therefore accumulate during periods of high zooplankton grazing activity. Besides bacteria, enzyme release by zooplankton should therefore be considered as an additional source of free dissolved enzymatic activity in the sea.

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