

1 **Role of microzooplankton during a *Phaeocystis* sp. bloom in the**
2 **Oosterschelde (SW Netherlands)**

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4 I. G. Teixeira^{a,*}, B. G. Crespo^{a,1}, T. G. Nielsen^{b,2}, F. G. Figueiras^a

5
6 ^a Instituto de Investigaciones Mariñas (CSIC), Eduardo Cabello 6, 36208 Vigo, Spain

7
8 ^b Department of Marine Ecology, The National Environmental Research Institute, University
9 of Aarhus, Frederiksborgvej 399, PO Box 358, 4000 Roskilde, Denmark.

10
11 ¹ Present address: Departament de Biologia Marina i Oceanografia, Institut de Ciències del
12 Mar (CSIC), Pg Marítim de la Barceloneta 37-49, 08003 Barcelona, Spain

13
14 ² Present address: National Institute of Aquatic Resources, DTU Aqua Section for
15 Oceanecology and Climate, Technical University of Denmark, Kavalergården 6, 2920
16 Charlottenlund, Denmark

17
18
19 *Corresponding author:

20 Email: isabeltx@iim.csic.es

21 Telephone: +34 986231930

22 Fax: +34 986 292767

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24 **RUNNING HEAD: Role of microzooplankton during a *Phaeocystis* sp. bloom**

1 ABSTRACT

2 The impact of microzooplankton on the plankton community was assessed during a bloom of
3 *Phaeocystis* sp. in the Oosterschelde basin (SW Netherlands). Chlorophyll *a* concentration
4 and *Phaeocystis* sp. abundance reached maxima values in the middle of the sampling period
5 ($\sim 27 \text{ mg chl } a \text{ m}^{-3}$ and $3.4 \times 10^{10} \text{ cells m}^{-3}$, respectively). The bloom collapse coincided with
6 the last sampling day, when accentuated decreases in chlorophyll *a* concentrations ($\sim 11 \text{ mg}$
7 $\text{chl } a \text{ m}^{-3}$) and *Phaeocystis* sp. cells numbers ($\sim 1.3 \times 10^{10} \text{ cells m}^{-3}$) were recorded.

8 Microzooplankton organisms were significant consumers of both phytoplankton and
9 heterotrophic plankton. Although *Phaeocystis* sp. was the most consumed organism (336 ± 71
10 $\text{mg C m}^{-3} \text{ d}^{-1}$), microzooplankton impact on its standing stock was lower than on the stocks of
11 other less abundant organisms. This impact was also lower during the peak of the bloom,
12 when colonial forms of *Phaeocystis* sp. presumably predominated, than during the bloom
13 collapse, when free-living cells were supposedly more abundant. The impact of
14 microzooplankton on heterotrophic organisms was higher than on phytoplankton, and
15 increased when the bloom collapsed. Picoheterotrophs experienced the highest impact on their
16 standing stock ($\sim 75\% \text{ d}^{-1}$) and production ($\sim 90\% \text{ d}^{-1}$). These results demonstrate that during a
17 *Phaeocystis* sp. bloom, the microbial food web was responsible for channelling a significant
18 fraction of plankton biomass, either from direct consumption of *Phaeocystis* sp. cells or
19 through consumption of heterotrophs, which would have been favoured by the high quantities
20 of organic matter released during the bloom collapse.

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22 KEY WORDS: *Phaeocystis* sp., microzooplankton, microbial food web, Oosterschelde, SW
23 Netherlands

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1 **1. Introduction**

2 *Phaeocystis* (Prymnesiophyceae, Haptophyta) is a worldwide distributed phytoplankton
3 genus with a significant role in biogeochemical processes, primarily due to the production of
4 high amounts of organic carbon and the production of sulphur compounds involved in climate
5 regulation (e.g. Schoemann et al., 2005). The major particularity of this organism is related to
6 its polymorphic life cycle, in which the solitary stage characterised by flagellate or non-motile
7 free-living small cells (3–9 µm) can alternate with the colonial stage composed of non-motile
8 cells embedded in a mucilaginous matrix (e.g. Rousseau et al., 1994, 2007). When in the form
9 of these mucilaginous colonies, which can reach several millimetres in diameter, *Phaeocystis*
10 spp. frequently forms nearly mono-specific blooms of huge biomass in polar and temperate
11 regions during the spring (e.g. Schoemann et al., 2005). The magnitude of these blooms is
12 such that locally they can account for up to 70% of the annual primary production (Joiris et
13 al., 1982).

14 Despite numerous investigations, several details on the life cycle of *Phaeocystis* spp. are
15 still not totally understood. This must be due to its morphological and physiological
16 variability and the contrasting environmental conditions of the different systems where this
17 alga can temporally dominate. High nutrient concentrations and high irradiance have been
18 assigned as responsible for promoting the transformation of solitary cells into colonial forms
19 (Veldhuis et al., 1986; Verity et al., 1988a; Peperzak, 1993). Conversely, the bloom demise,
20 habitually occurring within a short time period, has been attributed to nutrient limitation
21 and/or changes in temperature, both causing the colony disruption and the release of the
22 colonial cells into the water column (Veldhuis et al., 1986; Verity et al., 1988b). Pelagic
23 consumption, cell lysis and sedimentation are other processes that can be responsible for the
24 sudden disappearance of biomass from the water column (Wassmann, 1994; Weisse et al.,
25 1994; Brussaard et al., 1996). Though sedimentation has been observed to occur on several

1 occasions, cell lysis mediated by viral infection and pelagic consumption are frequently
2 identified as the main responsible factors for the termination of *Phaeocystis* spp. blooms
3 (Rousseau et al., 2000; Brussaard et al., 2005a; Baudoux et al., 2006).

4 The magnitude of the pelagic consumption of *Phaeocystis* spp. cells and derived material,
5 which determines the fraction of biomass that is transferred through the food web, has major
6 implications for the fluxes of matter and energy in the systems where this organism
7 dominates. Colonial forms, which are protected by a tough skin (Hamm et al., 1999) that aids
8 them to avoid viral infection and grazing on individual cells, apparently are suitable for
9 consumption by copepods and larger animals. Thus, colonial forms could contribute to sustain
10 the classical short food chain. In fact, it has been recently reported that colony formation and
11 enlargement in *Phaeocystis* spp. is enhanced by the presence of micrograzers, which could
12 indicate that this strategy corresponds to a mechanical defence against predation by
13 microzooplankton (Jakobsen and Tang, 2002; Tang, 2003). However, the trophic significance
14 of colonial biomass for metazoans is uncertain, and divergent reports regarding ingestion
15 rates, the nutritive value of preys and adverse effects on consumers are widespread through
16 literature (Weisse et al., 1994). These contrasting results seem to be related to the variability
17 in the size ratio between predators and colonies, the physiological state of preys and whether
18 experiments were performed in laboratory with *Phaeocystis* spp. as single preys or *in situ*
19 (Nejstgaard et al., 2007). In general, crustacean grazers depict a much lower grazing impact in
20 the field than in the laboratory, which could be due to the presence of alternative preys *in situ*,
21 and changes in the chemical properties of *Phaeocystis* spp. in the laboratory conditions which
22 could favour their consumption. Rather than consuming *Phaeocystis* spp. biomass during
23 bloom periods, copepods have been observed to feed on alternative preys, namely
24 microzooplankton (Hansen et al., 1993; Gasparini et al., 2000). In contrast, microzooplankton
25 apparently shows preference for solitary cells of only a few micrometers (Admiraal and

1 Venekamp, 1986; Weisse and Scheffel-Möser, 1990; Tang et al., 2001; Brussaard et al.,
2 2005a), although large naked dinoflagellates, such as *Noctiluca scintillans* and *Gyrodinium*
3 *spirale*, have been observed to feed on colonies (Jakobsen and Tang, 2002; Stelfox-
4 Widdicombe et al., 2004). In addition, both colony matrix disintegration and cell lysis at the
5 end of the bloom period, which increase the quantities of dissolved organic matter in the
6 water column (Alderkamp et al., 2007), should enter the microbial food web through
7 enhanced bacterial production and microzooplankton bacterivory (Fernández et al., 1992; van
8 Boeckel et al., 1992; Brussaard et al., 1995; Brussaard et al., 1996; Rousseau et al., 2000).

9 As the fate of the high biomass produced during *Phaeocystis* spp. blooms has a major
10 implication for biogeochemical cycles in marine systems, further assessment of the role of
11 microzooplankton during these blooms is fundamental to reach a better understanding of the
12 functioning of the marine systems dominated by these organisms. In this context, an intensive
13 and multidisciplinary research was conducted in the Oosterschelde (SW Netherlands) (Fig. 1)
14 between 4 and 13 May 2004, coinciding with the development and decay of a *Phaeocystis* sp.
15 bloom when phytoplankton biomass, primary production and the microzooplankton feeding
16 impact were studied.

17

18 **2. Materials and methods**

19 *2.1. Experimental procedure*

20 Sampling took place on board of R/V *Luctor* between 4 and 13 May 2004. Seawater samples
21 to determine chlorophyll *a* (chl *a*) concentration, phytoplankton biomass and primary
22 production were taken at three depths (surface, middle and bottom) during high tide in a
23 location close to a mussel bed (Fig. 1) using Niskin bottles. The sampling depths (1-6 m)
24 depended on the tidal range, which varied according to the sampling days. On 9 and 12 May
25 samples were not taken. Temperature, practical salinity and light penetration in the water

1 column were measured with a CTD to which a spherical PAR quantum sensor was attached.
2 Practical salinity was converted to absolute salinity (g kg^{-1}) according to TEOS-10 Primer,
3 where $\delta S_A = 0$ in this shallow water system.

4

5 2.2. Analyses

6 For chl *a*, three replicates of 50 ml were filtered onto GF/F filters. Pigments were extracted
7 overnight in 5 ml 96% ethanol in the dark and chl *a* concentration was determined against a
8 chl *a* standard after reading fluorescence before and after acidification on a Turner Designs
9 Model 700 Fluorometer.

10 Samples for microplankton counts were preserved in Lugol's iodine solution and
11 sedimented in composite sedimentation chambers. Diatoms, dinoflagellates, other flagellates
12 $>20 \mu\text{m}$ and ciliates were identified and counted at species level, when possible, using an
13 inverted microscope. Differentiation of phototrophic species of dinoflagellates, other
14 flagellates (hereafter, flagellates) and ciliates was basically done following Lessard and Swift
15 (1986) and Larsen and Sournia (1991); however, epifluorescence microscopy (see below) was
16 also used to assign phototrophy within these plankton groups. The biovolumes of each taxon
17 were calculated from the dimensions and shapes according to Hillebrand et al. (1999). The
18 plasmatic volumes of diatoms and total cell volume of dinoflagellates, flagellates $>20 \mu\text{m}$ and
19 ciliates were converted to cell carbon following Strathmann (1967) for diatoms and
20 dinoflagellates, Verity et al. (1992) for flagellates and Putt and Stoecker (1989) for ciliates.
21 Fixation procedure did not allowed us to quantify the number of colonies of *Phaeocystis* sp.,
22 and consequently the carbon biomass calculated for this species does not include the carbon
23 content of the colonial matrix (Rousseau et al., 1990; van Rijssel et al., 1997).

24 Primary production was measured as carbon assimilation after 24h *in situ* incubations.

25 Samples were placed in tissue culture flasks (3 light flasks and 2 dark flasks) and immediately

1 inoculated with ca. 3.7×10^5 Bq of $\text{NaH}^{14}\text{CO}_3$. Then, samples were placed in a rope with
2 specially constructed holders. A weight kept the line vertical, whereas the surface of the line
3 was maintained afloat by a large buoy. After 24h of incubation samples were recovered and
4 filtered through 25 mm GF/F filters under low vacuum pressure (<20 kPa). Filters were
5 placed in scintillation vials and left in HCl fumes atmosphere for 12h to remove unassimilated
6 ^{14}C . Radioisotope incorporation was determined with a liquid scintillation counter using the
7 external standard and the channel ratio methods to correct for quenching.

8

9 *2.3. Microzooplankton feeding impact*

10 The feeding impact of microzooplankton on the several organisms of the plankton
11 community was determined using the dilution technique (Landry and Hassett, 1982) on 4, 6,
12 8, 10 and 13 May 2004. Water was collected from the surface with a Niskin bottle as soon as
13 possible in the morning. All experimental containers, bottles and tubing were soaked in
14 10% HCl and rinsed with Milli Q water before each experiment. Filtered seawater was
15 obtained by gravity filtration through 0.8/0.2 μm Suporcap filter into a 25 l container. Sample
16 water was transferred to another 25 l container, very gently through a tube. In the laboratory,
17 dilutions were prepared directly in 2 l polycarbonate bottles (Nalgene). A total of 6 dilutions,
18 in the proportions of 100, 80, 60, 40, 20 and 10% of sample relatively to filtered seawater
19 were performed, with two replicates each. These bottles were incubated for 24h in running
20 seawater and at *in-situ* temperature and surface light conditions. During incubation period,
21 these bottles were regularly gently hand-rotated to avoid massive sedimentation.

22 Samples to determine the biomasses of plankton organisms ≤ 200 μm were taken from the
23 natural seawater at the beginning of the incubation and from each incubation bottle at the end
24 of the incubation. Samples for microplankton biomass determinations were processed as
25 described above. Metazooplankton <200 μm and mesozooplankton were found occasionally

1 and at very low concentrations in the dilution bottles. So, their importance as grazers in these
2 experiments was not considered. Pico- and nanoplankton were determined in subsamples of
3 10 ml fixed with buffered 0.2 μm filtered formaldehyde (2% final concentration) and stained
4 with DAPI at 0.1 $\mu\text{g ml}^{-1}$ final concentration (Porter and Feig, 1980). After 10 minutes in the
5 dark, samples were filtered through 0.2 μm black Millipore-Isopore filters. The filters were
6 then immersed in low fluorescence immersion oil and examined at x1000 magnification using
7 an epifluorescence microscope. Autotrophic organisms were enumerated under blue light
8 excitation and heterotrophic organisms were counted under excitation with UV light.
9 Bacterial biomass was estimated according to Lee and Furhmann (1987). Dimensions of
10 several individuals of the other groups were taken and cell volumes were calculated assuming
11 spherical shape. Cell carbon was estimated following Verity et al. (1992) for pico- and
12 nanoflagellates and Bratbak and Dundas (1984) for *Synechococcus*-type cyanobacteria.

13 Changes in total plankton ($\leq 200 \mu\text{m}$) biomass and in the several plankton groups
14 occurring between the beginning (C_0) and the end (C_t) of the incubation time ($t = 1$ day) were
15 used to calculate the net growth rates (k , d^{-1}):

$$16 \quad k = \frac{1}{t} \cdot \ln\left(\frac{C_t}{C_0}\right) \quad (1)$$

17 Mortality rates due to microzooplankton (m , d^{-1}) were calculated as the slope and the
18 specific growth rates (μ , d^{-1}) as the y-axis intercept of the linear regression between the net
19 growth rates and the fraction of unfiltered seawater (X) (Landry and Hassett, 1982):

$$20 \quad k = \mu - m \cdot X \quad (2)$$

21 In the cases of non-linear responses (saturated and saturated-increased, see Teixeira and
22 Figueiras, 2009), the specific growth rate (μ) was obtained by the y-axis intercept of the
23 regression of the linear part of the response. The mortality rate (m) was then obtained by

1 equation 2, replacing the specific growth rate calculated and using the net growth rate
2 obtained in the undiluted sample (Gallegos, 1989; Teixeira and Figueiras, 2009).

3 The daily mortality impact on the standing stock (%SS, %d⁻¹) and on production (%P, %d⁻¹)
4 were calculated as:

$$5 \quad \%SS = (1 - e^{-m}) \times 100 \quad (3)$$

$$6 \quad \%P = \frac{m}{\mu} \times 100 \quad (4)$$

7 The quantity of carbon consumed (G , mg m⁻³ d⁻¹) and produced (P , mg m⁻³ d⁻¹) were
8 estimated as:

$$9 \quad G = m \times C_m \quad (5)$$

$$10 \quad P = \mu \times C_m \quad (6)$$

11 where C_m (mg m⁻³) is:

$$12 \quad C_m = C_0 \left[\frac{e^{(\mu-m)t} - 1}{(\mu - m)t} \right] \quad (7)$$

13

14

15 **3. Results**

16 *3.1. Hydrographic conditions*

17 The water column (maximum depth = 6.5m) was well mixed from surface to bottom
18 during all days (Figs. 2a & b). There was a slight increase in the salinity of the water column,
19 which varied between 30.8 g kg⁻¹ on the first 3 days and values higher than 31 g kg⁻¹ on the
20 following days (Fig. 2a). Temperature did not show any tendency, varying between 12.6 and
21 13.3 °C (Fig. 2b).

22 The light attenuation coefficient in the water column (Fig. 2c) varied between 0.66 and
23 1.03 m⁻¹, with the highest value occurring on 8 May and the lowest on 13 May, at the end of

1 the sampling. Relatively low values (0.71 m^{-1}) were also recorded at the beginning of the
2 sampling, on 4 May (Fig. 2c).

3

4

5 3.2. *Chl a* and *Phaeocystis sp.*

6 *Chl a* concentration increased from $20.4 (\pm 1.0)$ to $26.8 (\pm 1.3)$ $\text{mg chl } a \text{ m}^{-3}$ during the
7 first 5 days, falling to $11.1 (\pm 0.2)$ $\text{mg chl } a \text{ m}^{-3}$ on the last day (Fig. 3a). Differences in *chl a*
8 concentrations between the 3 sampling depths were not significant ($0.40 < p < 0.88$; t-test for
9 paired samples).

10 *Chl a* concentration was positively correlated with phytoplankton abundance ($r = 0.91$, $p <$
11 0.01) and carbon biomass ($r = 0.89$, $p < 0.01$), two variables that showed a similar evolution
12 during the sampling period (Figs. 3b & c). The phytoplankton community was dominated by
13 *Phaeocystis sp.*, which accounted for $\sim 98\%$ of the total phytoplankton abundance and $\sim 90\%$
14 of the total phytoplankton biomass (Figs. 3b & c). This species reached an abundance of $3.4 \times$
15 $10^{10} (\pm 1.3 \times 10^{10})$ cells m^{-3} and a biomass of $871 (\pm 343)$ mg C m^{-3} during the peak of the
16 bloom on 8 May, and then dropped to $1.3 \times 10^{10} (\pm 2.5 \times 10^9)$ cells m^{-3} and $345 (\pm 63)$ mg C
17 m^{-3} on the last day. Differences between depths in abundance and biomass of *Phaeocystis sp.*
18 and total phytoplankton were not significant ($0.40 < p < 0.95$; t-test for paired samples).
19 Despite *Phaeocystis sp.* was mainly forming colonies (as it was observed in fresh samples and
20 deduced from the high accumulation of gelatinous structures on filters), the lugol's iodine that
21 was added to preserve the phytoplankton samples destroyed these structures. This resulted in
22 a much higher abundance of *Phaeocystis sp.* as solitary cells (99.4%) than as colonial forms
23 (0.6%) in the counted samples.

24 As a result of the correlation between phytoplankton carbon biomass and *chl a*
25 concentration, the autotrophic carbon to *chl a* (AC:*chl a*) ratio (Fig. 3d) remained relatively

1 constant around a mean value of 34 ± 3 during the sampling period. Nevertheless, the AC:chl
2 *a* ratio was slightly but significantly lower ($p < 0.05$, t-test for two samples) when chl *a* and
3 phytoplankton biomass were increasing (31 ± 2 , 4 to 7 May) than when these two variables
4 were decreasing (37 ± 2 , 8 to 13 May).

5

6 3.3. Primary production

7 Primary production decreased with depth (Fig. 4) in accordance with the light attenuation
8 coefficient in the water column (Fig. 2c). Thus, primary production varied between 102 and
9 $608 \text{ mg C m}^{-3} \text{ d}^{-1}$ at the surface, 76 and $444 \text{ mg C m}^{-3} \text{ d}^{-1}$ at the middle depth and between 0
10 and $89 \text{ mg C m}^{-3} \text{ d}^{-1}$ at the bottom layer (Fig. 4). Highest differences between depths were
11 observed on 8 May. The zero value recorded at the bottom layer that day coincided with the
12 strongest light attenuation in the water column (Fig. 2c), which led to a photic layer shallower
13 than the depth at the station. Depth integrated primary production varied between 353 and
14 $1626 \text{ mg C m}^{-2} \text{ d}^{-1}$, with lowest values at the beginning and the end of the sampling period
15 (Fig. 4) and highest values in between, on 8 and 10 May. Integrated primary production was
16 positively correlated with integrated phytoplankton carbon biomass ($r = 0.74$; $p < 0.05$).

17

18 3.4. Microzooplankton feeding impact

19 In addition to *Phaeocystis* sp., which represented ~90% of total phytoplankton biomass
20 (Fig. 5a), diatoms were also present in the initial plankton populations of dilution
21 experiments. Nevertheless, diatoms accounted for a minor fraction (3-7%) of the total
22 autotrophic carbon (AC). The biomass of diatoms increased from ~ 25 mg C m^{-3} during the
23 first days to ~ 40 mg C m^{-3} on the last days. Among this group, *Rhizosolenia imbricata*,
24 *Guinardia delicatula*, *Cerataulina pelagica* and small and large *Pseudo-nitzschia* spp. were
25 the most abundant species. The phototrophic ciliate *Mesodinium rubrum*, autotrophic

1 dinoflagellates and picoautotrophs (*Synechococcus*-like cyanobacteria and picoflagellates)
2 were also found, but their contribution to phytoplankton biomass was very low (Fig. 5a). The
3 evolution of heterotrophs was different from that of autotrophs (Fig. 5b), showing a peak on 6
4 May (155 mg C m⁻³) due to the high biomass of heterotrophic nanoflagellates (HNF) on that
5 day (121 mg C m⁻³). HNF, which dominated within microzooplankton community at the
6 beginning of the sampling period, accounting almost for 90% of the microzooplankton
7 biomass, decreased from 6 May onwards. On the last day of sampling, HNF with a biomass of
8 12 mg C m⁻³ only represented 20% of microzooplankton carbon. Heterotrophic picoflagellates
9 (HPF) showed also a decreasing tendency during the sampling period, but their biomass was
10 much smaller, varying between 3.8 and 1.3 mg C m⁻³. On the contrary, the importance of
11 heterotrophic dinoflagellates (mainly naked forms >20 µm), ciliates (mainly aloricate
12 choreotrichs >50 µm) and heterotrophic bacteria (HB) increased through the sampling period
13 (Fig. 5b); heterotrophic dinoflagellates from 3 to 26 mg C m⁻³, ciliates from 7 to 23 mg C m⁻³
14 and HB from 19 to 37 mg C m⁻³.

15 According to the outcome of the dilution experiments (Table 1), heterotrophic pico- and
16 nanoplankton (HB, HPF and HNF), *Synechococcus* and *Phaeocystis* sp. were consumed
17 during all the sampling period. Diatoms were also consumed, but only on May 13. Growth
18 and mortality rates were highly variable amongst the several organisms, with growth rates
19 varying between 0.15 and 3.08 d⁻¹ and mortality rates between 0 and 2.64 d⁻¹. The rates were
20 generally higher for total heterotrophic carbon (HC) (1.03 ≤ µ ≤ 1.48 d⁻¹ and 0.69 ≤ m ≤ 1.48
21 d⁻¹) than for total AC (0.16 ≤ µ ≤ 0.91 d⁻¹ and 0.23 ≤ m ≤ 0.58 d⁻¹). Growth and grazing rates
22 of AC were very similar to those obtained for *Phaeocystis* sp. (0.15 ≤ µ ≤ 0.96 d⁻¹ and 0.26 ≤
23 m ≤ 0.62 d⁻¹).

24 On average, 40 ± 5% d⁻¹ of the whole plankton standing stock ≤200 µm was consumed by
25 microzooplankton (Fig. 6a). HC was more impacted (Fig. 6a, 67 ± 11% d⁻¹) than AC (Fig. 6a,

1 $34 \pm 9\% \text{ d}^{-1}$). Moreover, there was an increase in the consumption of the standing stock of HC
2 from $50\% \text{ d}^{-1}$ being ingested at the beginning of the sampling to almost $80\% \text{ d}^{-1}$ at the end
3 (Fig. 6a). Among autotrophs, the lowest grazing impact ($21\% \text{ d}^{-1}$) was found on 8 May (Fig.
4 6a), coinciding with the maximum peak in the abundance of *Phaeocystis* sp. (Fig. 3b & c).
5 The impact on the standing stock of *Phaeocystis* sp. ($37 \pm 9\% \text{ d}^{-1}$) was relatively low, always
6 below $50\% \text{ d}^{-1}$ (Fig. 6b). Regarding the several plankton groups, the highest impact occurred
7 on the standing stock of picoheterotrophs ($75 \pm 6\% \text{ d}^{-1}$ for HB and $74 \pm 19\% \text{ d}^{-1}$ for HPF),
8 followed by HNF ($62 \pm 36\% \text{ d}^{-1}$), though HNF were not consumed on 4 May (Fig. 6c).

9 Concerning the microzooplankton impact on production, a sudden increase was observed
10 for AC and TC on the last day of sampling (Fig. 6d), when consumption shifted from $63 \pm$
11 $18\% \text{ d}^{-1}$ of AC and $74 \pm 23\% \text{ d}^{-1}$ of TC produced during the previous days to $373\% \text{ d}^{-1}$ of the
12 AC and $265\% \text{ d}^{-1}$ of the TC produced on 13 May. This evolution closely resembled that
13 observed for the impact on the production of *Phaeocystis* sp. (Fig. 6e), which derived from a
14 consumption of $67 \pm 20\% \text{ d}^{-1}$ of the production between 4 and 10 May to a value of $422\% \text{ d}^{-1}$
15 on 13 May. The impact on the production of the other two phytoplankton groups with
16 significant responses in the dilution experiments was highly variable, ranging from 17 to
17 $100\% \text{ d}^{-1}$ for *Synechococcus* and from 52 to $98\% \text{ d}^{-1}$ for autotrophic picoflagellates (APF)
18 (Fig. 6e). In contrast, the microzooplankton impact on the production of HC gradually
19 increased over the sampling period, varying from $56\% \text{ d}^{-1}$ on the first day to $113\% \text{ d}^{-1}$ on the
20 last day (Fig. 6d), showing a relatively constant and significant impact ($96 \pm 16\% \text{ d}^{-1}$) on the
21 production of HB (Fig. 6f). The other two heterotrophic groups, HPF and HNF, also
22 experienced a high impact ($87 \pm 33\% \text{ d}^{-1}$ and $73 \pm 45\% \text{ d}^{-1}$, respectively) on their production,
23 although it was highly variable (Fig. 6f).

24 In spite of these percentages, consumption of autotrophic biomass ($255\text{--}420 \text{ mg C m}^{-3} \text{ d}^{-1}$)
25 1) was always higher than consumption of heterotrophic biomass ($90\text{--}161 \text{ mg C m}^{-3} \text{ d}^{-1}$) (Fig.

1 7a). *Phaeocystis* sp. was virtually the only phytoplankton species consumed from 4 to 10 May
2 and represented 95% of the total AC removed on 13 May, when diatoms were also consumed
3 (Fig. 7b). Among heterotrophs (Fig. 7c), HNF, which were not consumed on the first day of
4 sampling, were highly consumed on 6 May ($131 \text{ mg C m}^{-3} \text{ d}^{-1}$), showing after that a
5 decreasing tendency. In contrast, the HB biomass daily removed by microzooplankton
6 increased over the sampling period, from $28 \text{ mg C m}^{-3} \text{ d}^{-1}$ on the first day to $45 \text{ mg C m}^{-3} \text{ d}^{-1}$
7 in the last dilution experiment. The autotrophic biomass produced ($68 - 980 \text{ mg C m}^{-3} \text{ d}^{-1}$)
8 during the sampling period showed higher variations than the biomass consumed (Figs. 7a &
9 d). *Phaeocystis* sp. was the phytoplankton species with the highest production (Figs. 7e),
10 representing >96% of the total autotrophic carbon produced. In contrast, consumption ($90-161$
11 $\text{mg C m}^{-3} \text{ d}^{-1}$) and production ($120-164 \text{ mg C m}^{-3} \text{ d}^{-1}$) of heterotrophic biomass were more
12 tightly coupled (Figs. 7a,d,c,f), with HNF showing higher production during the first 3 days
13 of sampling and HB acquiring more importance at the end.

14 Growth and microzooplankton grazing rates derived from dilution experiments (Table 1)
15 were used to predict the expected *Phaeocystis* sp. biomass in the water column on the
16 following day (Fig. 8). The observed and predicted biomasses were in reasonable agreement
17 at the beginning of the bloom, with predicted biomass exceeding the observed biomass on 6
18 and 49 mg C m^{-3} on May 5 and 7, respectively. However, this difference was much higher
19 when the bloom started to decay (May 11), with observed biomass being 403 mg C m^{-3} lower
20 than expected biomass estimated considering only growth and microzooplankton grazing rates
21 obtained in dilution experiments.

22

23 **4. Discussion**

24 *4.1. The Phaeocystis sp. bloom*

1 Blooms of *Phaeocystis* spp. in the Oosterschelde (Laanbroek et al., 1985; Bakker et al.,
2 1990) and in the adjacent North Sea (e.g. Veldhuis et al., 1986; Weisse & Scheffel-Möser,
3 1990; Rousseau et al., 2000) are frequent in spring. Here, we followed an almost
4 monospecific *Phaeocystis* sp. bloom in the Oosterschelde well reflected in the evolution of
5 chl *a* concentration and cell abundance (Figs. 3a & b). The bloom was also discernible in the
6 evolution of primary production (Fig. 4), which was correlated with *Phaeocystis* sp. biomass
7 (Fig. 3c), and in the evolution of the light attenuation coefficient (Fig. 2c). Even though the
8 progress of this bloom was observed inside the Oosterschelde, its advection from the North
9 Sea cannot be disregarded, because the increase in salinity recorded during the sampling
10 period (Fig. 2a) points to the possible input of water from the North Sea, saltier than the water
11 in the Oosterschelde (Laanbroek et al., 1985).

12 The high chl *a* concentration (20 mg chl *a* m⁻³) and the high cell abundance (~23 x 10⁹
13 cells m⁻³) of *Phaeocystis* sp. (Figs. 3a & b) recorded on the first day of sampling were above
14 the threshold reported for the formation of colonies (Schoemann et al., 2005). This indicates
15 that our sampling started after the onset of the bloom, when *Phaeocystis* sp. was already in the
16 colonial form. The bloom rapidly evolved to reach its maximum 4 days later, and then
17 declined. Changes in temperature and/or nutrient limitation have been assigned as triggering
18 factors for the beginning of bloom collapse (Veldhuis et al., 1986; Verity et al., 1988b). In
19 this case temperature did not show any significant change (Fig. 2b), but nutrient limitation
20 could have been responsible for finishing this event. Although not having a nutrient record
21 during the sampling period, we do have nutrient data before (on April 29) and after (May 17)
22 the survey. Thus, on April 29 nitrate concentration was 25.44 µmol l⁻¹ while on May 17 was
23 8.53 µmol l⁻¹ (T. Ysebaert, personal communication), indicating a high nutrient decrease
24 during that period. By its turn, phosphate concentrations were very low on both days (0.1 and
25 0.23 µmol l⁻¹, respectively) and this nutrient is usually limiting *Phaeocystis* blooms in the

1 North Sea (Veldhuis et al., 1986; Peperzak et al., 1998). Nonetheless, the decrease in nitrate
2 concentrations could lead to a reduction in *Phaeocystis* biomass that can be viewed as the
3 bloom collapse. Also the changes observed in the AC:chl *a* ratios (Fig. 3d), with slightly but
4 significant higher ratios during the demise phase, could point to nutrient limitation as the
5 possible factor triggering the bloom decline. However, this could also be a consequence of a
6 decrease in the light attenuation coefficient in the water column with the progress of the
7 bloom (Fig. 2c). Other loss factors, as sedimentation, have also been reported (Wassmann,
8 1994), but sinking seems unusual in this region where tidal currents are strong enough to
9 prevent it. In fact, in this work we did not observe accumulation of phytoplankton in bottom
10 layers at the end of sampling (Fig. 4).

11

12 4.2. *Microzooplankton feeding impact*

13 Our results showed that microzooplankton organisms were significant consumers in the
14 pelagic system, but their impact varied during the bloom period (Fig. 6). The highest quantity
15 of carbon biomass consumed by microzooplankton corresponded to *Phaeocystis* sp. cells
16 (Figs. 7a & b). However, this pattern was forced by the dominance of this species during the
17 sampling period resulting in higher quantities consumed even though mortality rates were
18 lower than those of the other plankton groups (Table 1). Despite these high quantities
19 consumed, microzooplankton was unable to control the bloom development, consuming only
20 a fraction of the *Phaeocystis* sp. standing stock and production (Figs. 6b & e). However, on
21 the last sampling day, when the bloom decayed, microzooplankton consumption exceeded 4
22 times the production of this haptophyte (Fig. 6e). These variations in consumption must be
23 related to the structural changes occurring in *Phaeocystis* sp. through the sampling period.
24 With exception of few dinoflagellate species, such as *Noctiluca scintillans* (Jakobsen and
25 Tang, 2002) or *Gyrodinium spirale* (Stelfox-Widdicombe et al., 2004), microzooplankton is

1 not an efficient consumer either of the whole colonial structure or the colonial single cells
2 protected by the colony skin (Hamm et al., 1999). However, when *Phaeocystis* sp. is in the
3 form of single cells in the water column, microzooplankton organisms are able to ingest them
4 (Admiraal and Venekamp, 1986; Weisse and Scheffel-Möser, 1990; Tang et al., 2001). Here
5 we found a higher microzooplankton grazing on *Phaeocystis* sp. on the last day (Figs. 6b &
6 e), when colonies presumably had collapsed and cells were free-living. As on the previous
7 days *Phaeocystis* sp. was probably forming colonies, the impact of microzooplankton would
8 have occurred on the less abundant solitary cells that usually are present in the water column
9 coexisting with colonial structures (Rousseau et al. 1994). Consumption of some colonial
10 forms could also have occurred, because large naked forms predominated among
11 heterotrophic dinoflagellates and these organisms could feed on *Phaeocystis* sp. colonies as
12 *Gyrodinium spirale* do it (Stelfox-Widdicombe et al., 2004). It must be noted however that
13 the accentuated impact on *Phaeocystis* sp. production on May 13 was not due to a higher
14 grazing mortality, but to the low growth rate of *Phaeocystis* sp. on this day, several times
15 lower than the mortality rate (Table 1). This low growth rate can indicate nutrient limitation,
16 as stated before, but also that microzooplankton was not the only mortality factor for
17 *Phaeocystis* sp. on this day. When viral lysis is significant, the conventional dilution protocol
18 can under-estimate the specific growth rates (Evans et al., 2003; Baudoux et al., 2006). Cell
19 lysis mediated by viral attack has been shown to be very important in the termination of the
20 *Phaeocystis* spp. blooms (Rousseau et al., 2000; Brussaard et al., 2005a; Baudoux et al.,
21 2006). Thus, the high microzooplankton grazing on *Phaeocystis* sp. production at the end of
22 the bloom must be taken carefully, as probably both viral lysis and grazing by micro-
23 heterotrophs were acting on this alga simultaneously. This can also explain the results shown
24 in Fig. 8, where other loss factors than microzooplankton grazing were apparently responsible
25 for the disappearance of *Phaeocystis* sp. biomass from the water column on May 11, when the

1 bloom started to decay. Besides viral lysis and grazing by micro-heterotrophs, other loss
2 processes can also include pelagic consumption by mesozooplankton (Weisse et al., 1994) or
3 even benthic consumption by mussels (Smaal and Twisk, 1997), which are abundant in this
4 system. Colony destruction occurring when the bloom started to collapse probably made
5 *Phaeocystis* sp. more susceptible to all consumers.

6 Consumption by microzooplankton during the *Phaeocystis* sp. bloom was not restricted to
7 this alga. Picoautotrophs (APF and *Synechococcus*) were also grazed (Table 1), although the
8 quantities of carbon consumed of these organisms were insignificant due to their low
9 biomasses (Fig. 7b). Grazing on diatoms, which only occurred on the last day (Table 1), could
10 be related to the presence of *Pseudo-nitzschia* spp. Pennate diatoms, including these species,
11 are known to be attached to *Phaeocystis* sp. colonies (Peperzak et al., 1998; Sazhin et al.,
12 2007). Thus, *Pseudo-nitzschia* spp. could have been attached to the colonial forms during the
13 bloom and hence be protected against grazing. On the last day, with the disintegration of
14 colonies, *Pseudo-nitzschia* spp. could become free-living, and as *Phaeocystis* sp. free cells, be
15 more susceptible to pelagic consumption.

16 In addition to phytoplankton, heterotrophs were also an important nutritional source for
17 microzooplankton (Figs. 7a & c). In relative terms of biomass and abundance,
18 microzooplankton impact on heterotrophs was actually higher than on phytoplankton (Figs.
19 6a). Among this group, picoheterotrophs (HPF and HB) were highly impacted by
20 microzooplankton, with ~75% of their standing stock and ~90% of their production being
21 channelled to the microbial food web (Figs. 6c & f). The microzooplankton consumption was
22 not enough to fully control HB, which showed slight increases in biomass (Fig. 5b) and
23 production (Fig. 7f) over the sampling period, reaching maximum values during the bloom
24 decay. This increase can be associated with the response of HB to the increase in dissolved
25 organic matter concentrations due to colony disintegration (Laanbroek et al., 1985; Veldhuis

1 et al., 1986; Verity et al., 1988b; Noordkamp et al., 2000; Rousseau et al., 2000; Brussaard et
2 al., 2005b). HNF were also consumed (Fig. 7c), in agreement with other studies indicating
3 consumption of these organisms by microzooplankton during *Phaeocystis* sp. blooms
4 (Brussaard et al., 1995, 1996; Rousseau et al., 2000).

5 Noteworthy, microzooplankton impact increased towards the end of the sampling period
6 (Fig. 6). This was not only observed for grazing on phytoplankton, but also for predation on
7 heterotrophic components. This increasing importance of the microbial food web in the
8 transfer of biomass with the progress of the bloom adds new evidences to previous reports
9 (Admiraal and Venekamp, 1986; Weisse and Scheffel-Möser, 1990; Brussaard et al., 1995;
10 Peperzak et al., 1998; Rousseau et al., 2000) supporting the view that *Phaeocystis* sp. blooms
11 stimulate the dominance of the microbial food web.

12

13 4.3. Concluding remarks

14 These results indicate that a significant fraction of the *Phaeocystis* sp. bloom was
15 channelled through the microbial food web, with two pathways in which microzooplankton is
16 a direct intervenient. Firstly, direct consumption of *Phaeocystis* sp. cells by microzooplankton
17 occurred in all experiments and increased during the bloom decay. Secondly,
18 microzooplankton consumption of small heterotrophs allowed the return to the food web of
19 the high amounts of organic matter released during the collapse of the bloom (Alderkamp et
20 al., 2007): HB taking advantage of the released organic matter, and HNF being favoured by
21 the increase in bacterial production. Thus, microzooplankton, which is consumed by copepods
22 during *Phaeocystis* blooms (Hansen et al., 1993; Gasparini et al., 2000), transfers the
23 photosynthesised organic matter to higher trophic levels, linking the microbial and classical
24 food webs and therefore enhancing the efficiency of the pelagic food web.

25

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9

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1 **Figure legends**

2

3 Fig. 1. Map of the Oosterschelde with the location of the sampling site (black dot) and its
4 position in the Dutch coast (inset).

5

6 Fig. 2. Vertical profiles of (a) absolute salinity (g kg^{-1}), (b) temperature ($^{\circ}\text{C}$), and (c) light
7 attenuation coefficient (m^{-1}) in the water column.

8

9 Fig. 3. Mean values in the water column of (a) chl *a* concentration ($\text{mg chl } a \text{ m}^{-3}$), (b) cell
10 abundance of *Phaeocystis* sp. and total phytoplankton (cells m^{-3}), (c) carbon biomass of
11 *Phaeocystis* sp. and total phytoplankton (mg C m^{-3}) and (d) autotrophic carbon to chl *a* ratio.
12 Vertical bars in a, b, and c are standard deviations.

13

14 Fig. 4. Primary production rates at the surface, middle and bottom depths ($\text{mg C m}^{-3} \text{ d}^{-1}$).

15 Values of integrated primary production ($\text{mg C m}^{-2} \text{ d}^{-1}$) are also given.

16

17 Fig. 5. Initial carbon biomass (mg C m^{-3}) for each dilution experiment of (a) autotrophic and
18 (b) heterotrophic plankton community. APF: autotrophic picoflagellates, AC: autotrophic
19 carbon, HPF: heterotrophic picoflagellates, HB: heterotrophic bacteria, HNF: heterotrophic
20 nanoflagellates, HC: heterotrophic carbon. Left y-axis corresponds to bars and right y-axis
21 corresponds to lines.

22

23 Fig. 6. Percentages of the standing stock (a-c) and production (d-f) daily consumed by
24 microzooplankton of the bulk plankton groups (a & d), autotrophic plankton groups (b & e)
25 and heterotrophic plankton groups (c & f). AC, autotrophic carbon; HC, heterotrophic carbon;

1 TC, total carbon; APF, autotrophic picoflagellates, HPF, heterotrophic picoflagellates; HNF,
2 heterotrophic nanoflagellates; HB, heterotrophic bacteria.

3

4 Fig. 7. Plankton carbon biomass (mg C m^{-3}) daily ingested by microzooplankton (left panels)
5 and daily produced (right panels). (a & d) total autotrophic and heterotrophic plankton, (b &
6 e), autotrophic plankton groups, (c & f) heterotrophic plankton groups. AC, autotrophic
7 carbon; HC, heterotrophic carbon; APF, autotrophic picoflagellates, HPF, heterotrophic
8 picoflagellates; HNF, heterotrophic nanoflagellates; HB, heterotrophic bacteria. Right y-axis
9 in b & e corresponds to lines.

10

11 Fig. 8. Comparison between *Phaeocystis* sp. biomass (mg C m^{-3}) observed *in situ* and
12 estimated using the growth and microzooplankton grazing rates obtained from the dilution
13 experiments started the previous day.

14

15

Table 1[Click here to download Table\(s\): Table 1.doc](#)

Table 1. Mortality (m , d^{-1}) and growth (μ , d^{-1}) rates obtained from the dilution experiments. AC: total autotrophic carbon, APF: autotrophic picoflagellates, HC: total heterotrophic carbon, HPF: heterotrophic picoflagellates, HNF: heterotrophic nanoflagellates, HB: heterotrophic bacteria, TC: total plankton carbon biomass ($\leq 200\mu m$). r^2 : coefficient of determination for the linear regressions ($p < 0.05$) between the net growth rate and the fraction of unfiltered seawater. In cases of saturated or saturated-increased responses, r^2 was obtained for the linear part of the response (Teixeira and Figueiras, 2009). When r^2 is not given, the growth rate was obtained with only the two highest diluted bottles (Gallegos, 1989). a: non-linear responses. ns: not significant.

| | 04 May | | | 06 May | | | 08 May | | | 10 May | | | 13 May | | |
|------------------------|--------|-------|-------------------|--------|-------|-------------------|--------|-------|-------|--------|-------|-------------------|--------|-------|-------------------|
| | m | μ | r^2 | m | μ | r^2 | m | μ | r^2 | m | μ | r^2 | m | μ | r^2 |
| AC | 0.39 | 0.66 | 0.84 ^a | 0.51 | 0.60 | 0.73 ^a | 0.23 | 0.36 | a | 0.39 | 0.91 | 0.76 ^a | 0.58 | 0.16 | 0.91 |
| Diatoms | | ns | | | ns | | | ns | | | ns | | 0.54 | 0.64 | 0.90 ^a |
| <i>Synechococcus</i> | 0.43 | 0.70 | 0.99 ^a | 0.21 | 1.26 | 0.93 ^a | 0.35 | 0.30 | 0.84 | 1.02 | 1.10 | 0.90 | 1.33 | 1.33 | 0.95 ^a |
| APF | 0.36 | 0.49 | 0.93 | | ns | | 1.02 | 1.04 | 0.71 | 0.87 | 1.67 | 0.70 | | ns | |
| <i>Phaeocystis</i> sp. | 0.43 | 0.70 | 0.86 ^a | 0.57 | 0.61 | 0.74 ^a | 0.26 | 0.38 | a | 0.43 | 0.96 | 0.76 ^a | 0.62 | 0.15 | 0.89 |
| HC | 0.69 | 1.24 | 0.89 | 0.97 | 1.03 | 0.80 | 1.36 | 1.37 | 0.97 | 1.25 | 1.48 | 0.87 ^a | 1.48 | 1.31 | 0.95 |
| HPF | 0.78 | 1.48 | 0.84 ^a | 0.80 | 1.49 | 0.80 ^a | 1.42 | 1.09 | 0.85 | 2.42 | 2.45 | 0.88 ^a | 2.64 | 2.74 | 0.91 |
| HNF | 0.00 | 1.09 | 0.95 ^a | 1.20 | 1.00 | 0.93 | 1.15 | 1.59 | 0.91 | 1.60 | 1.70 | 0.72 | 2.36 | 3.08 | 0.85 |
| HB | 1.73 | 1.42 | 0.93 | 1.45 | 1.78 | 0.86 ^a | 1.15 | 1.20 | 0.81 | 1.52 | 1.86 | 0.96 ^a | 1.20 | 1.23 | 0.91 ^a |
| TC | 0.44 | 0.77 | 0.96 ^a | 0.52 | 0.53 | 0.67 | 0.48 | 0.53 | a | 0.51 | 1.00 | 0.75 ^a | 0.65 | 0.25 | 0.90 |



Fig. 1
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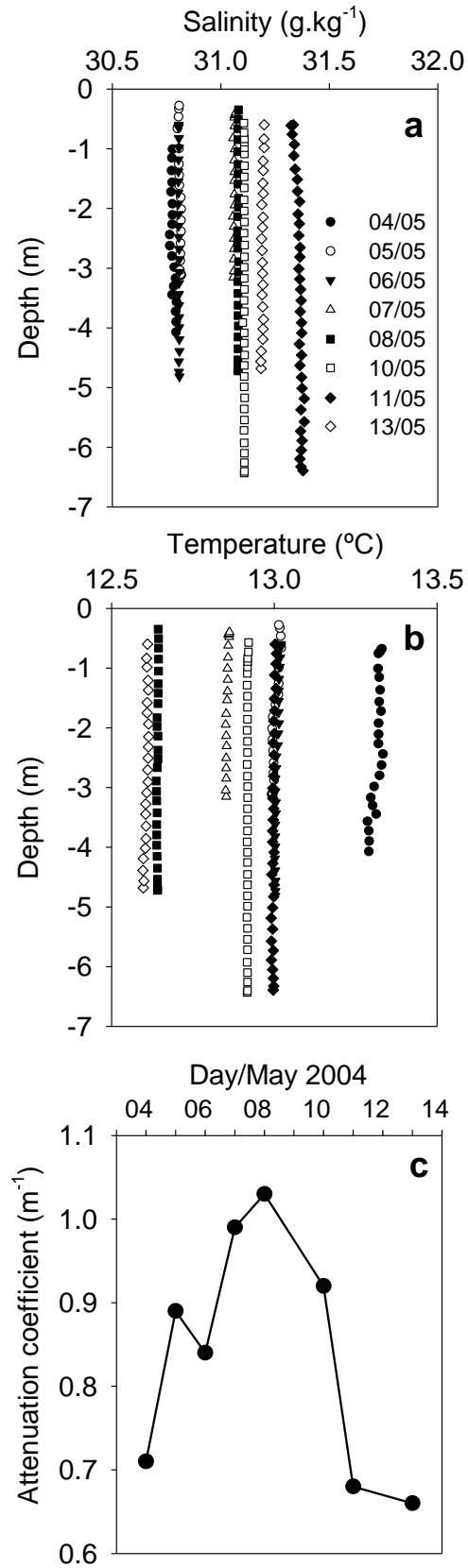


Fig. 2
Teixeira et al.

Figure 3
Click here to download Figure(s): Fig3.doc

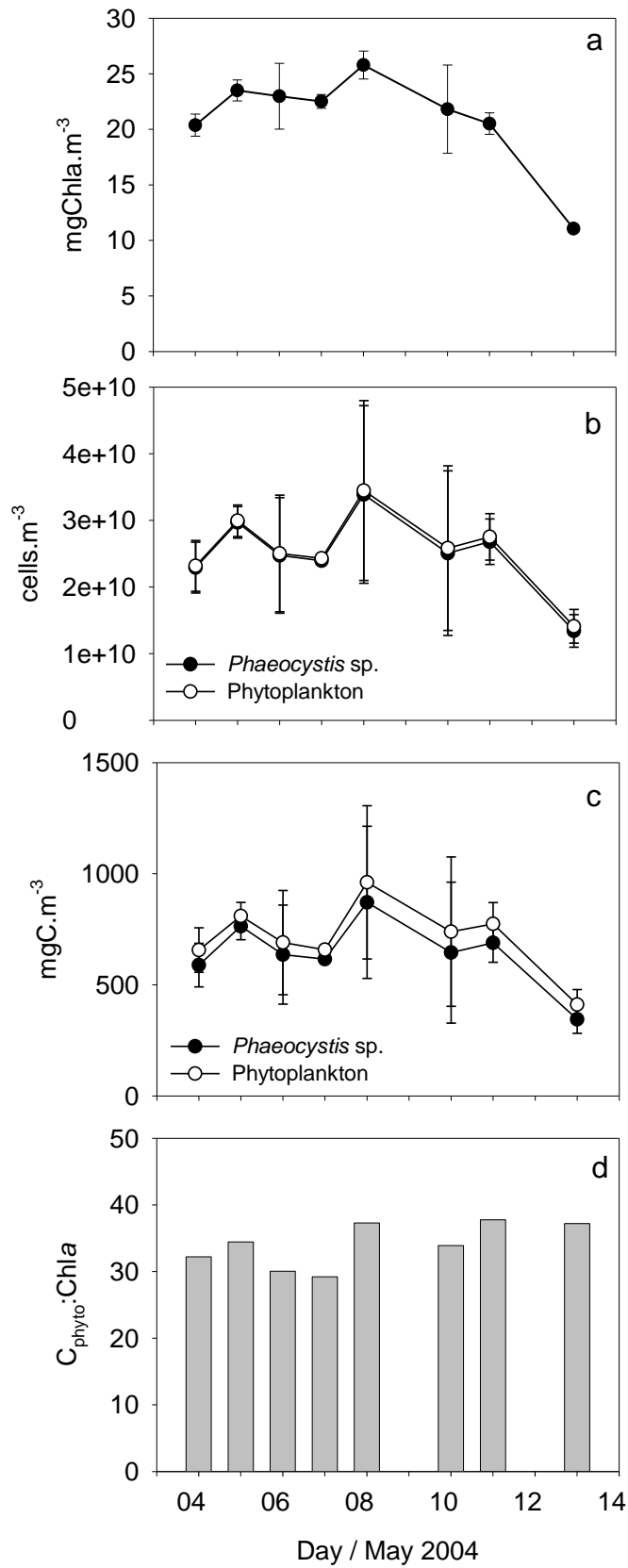


Fig. 3
Teixeira et al.

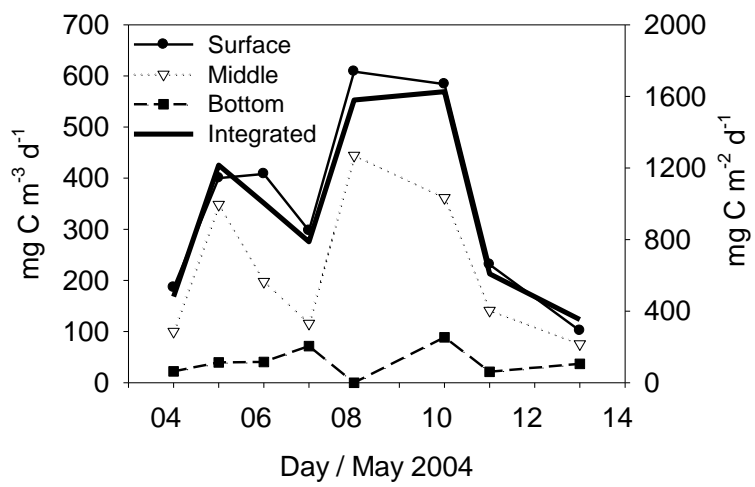


Fig. 4
Teixeira et al.

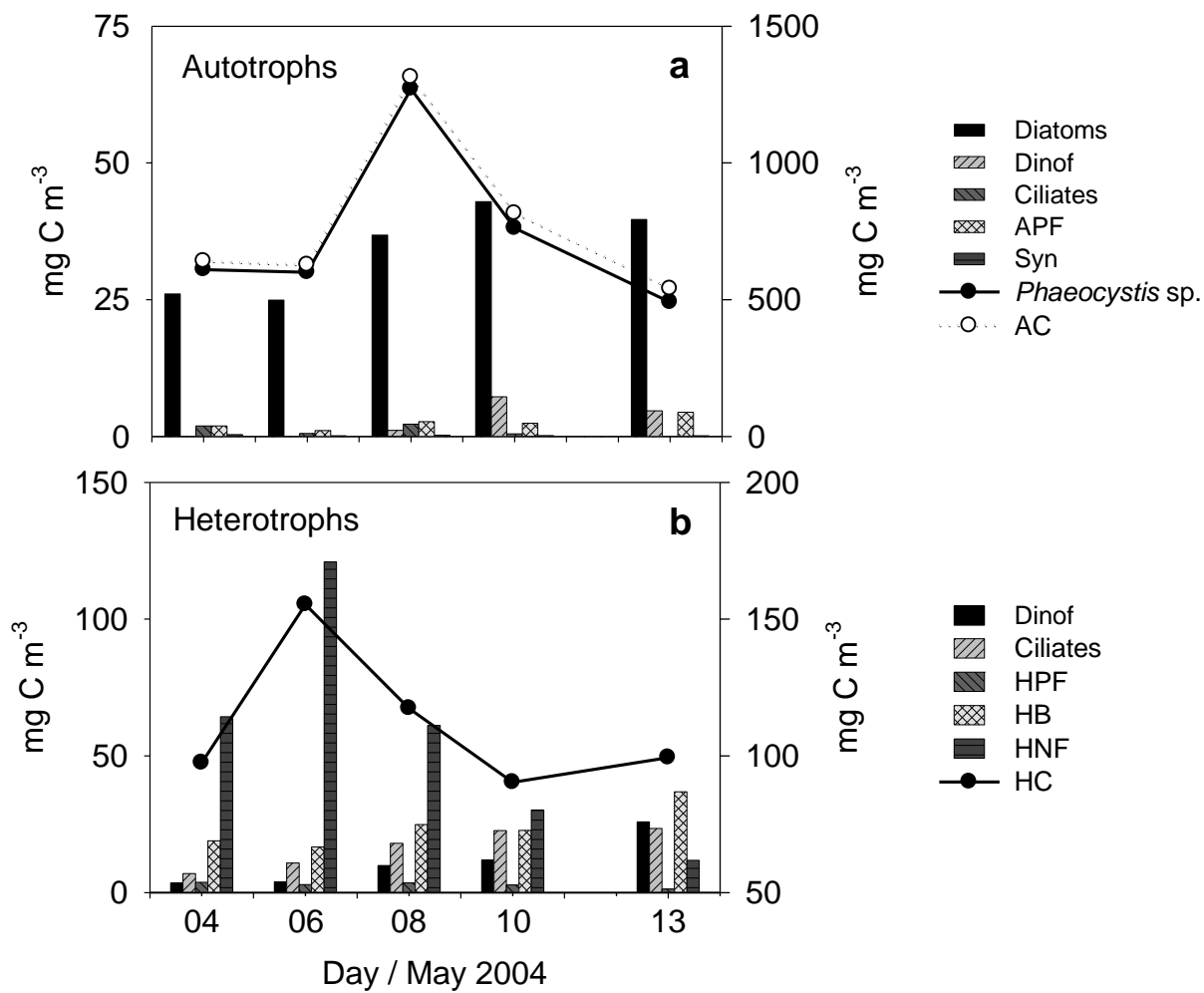


Fig. 5
Teixeira et al.

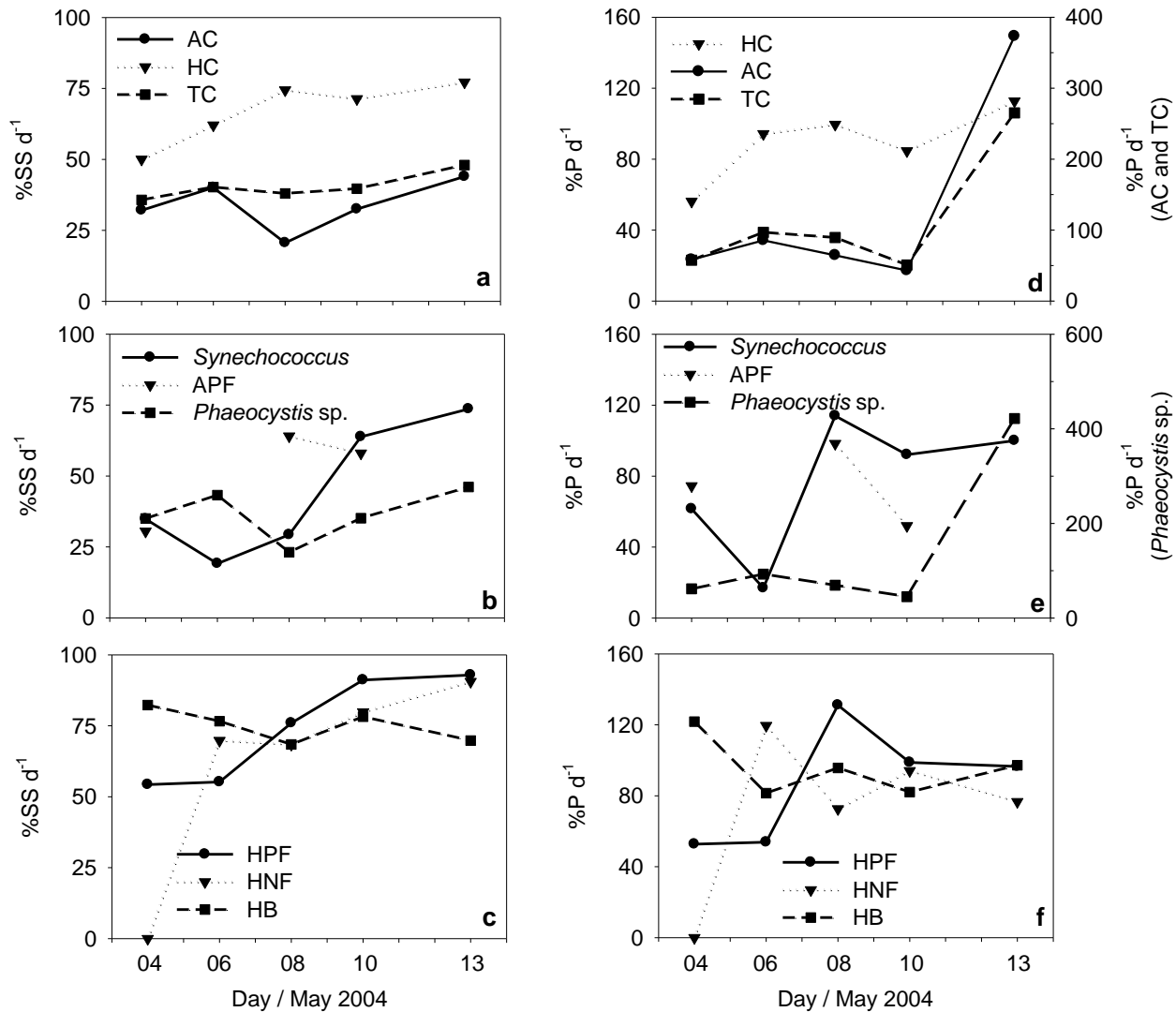


Fig. 6
 Teixeira et al.

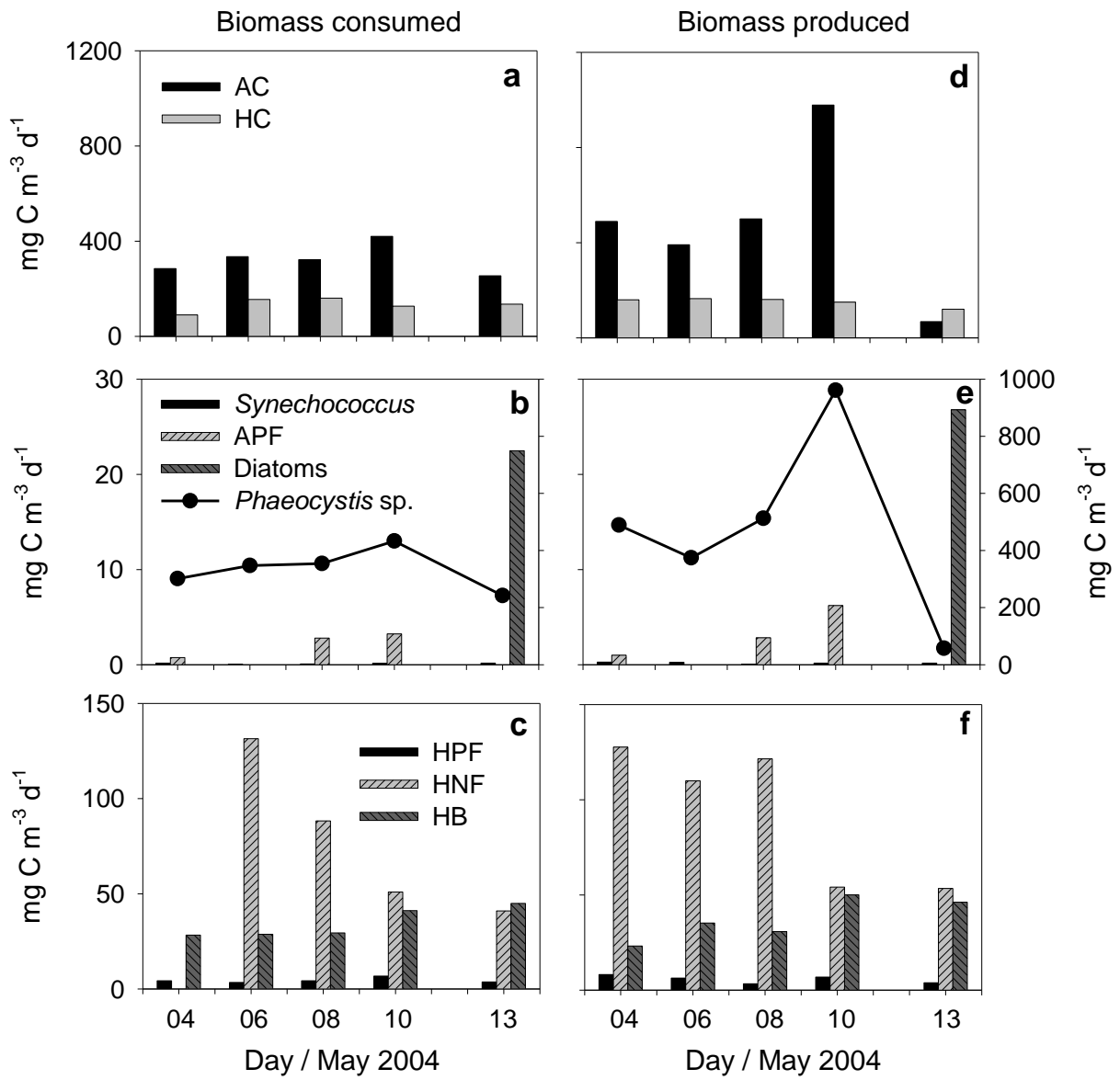


Fig.7
 Teixeira et al.

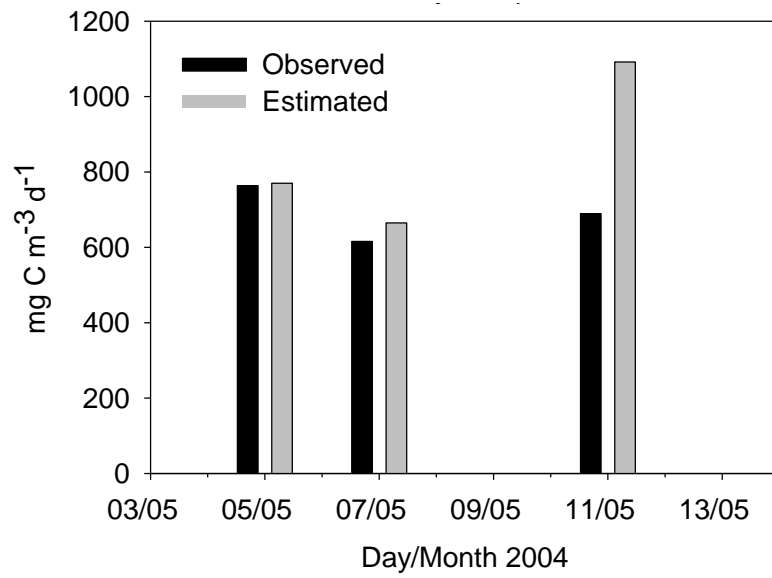


Fig. 8
Teixeira et al.

*Highlights

Microzooplankton consumed a significant fraction of *Phaeocystis* sp. biomass.

Microzooplankton impact was higher during the bloom collapse, when cells were free-living.

Heterotrophs were also consumed by microzooplankton simultaneously.

We observed two main pathways for channelling *Phaeocystis* sp bloom biomass.