**Structure of the microbial plankton community in the NW
Iberian margin at the end of the upwelling season**

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ABSTRACT

At the end of summer, seasonal upwelling at the NW Iberian Peninsula relaxes and nutrient depleted oceanic waters conveyed by the Iberian Poleward Current (IPC) converge on the western shelf with outwelled waters from the four bays known as Rías Baixas. At that time, with virtual absence of diatoms and dominance of nanoflagellates in the autotrophic microbial plankton community, *Synechococcus* accounted for ~ 20% of the total autotrophic biomass (AB). This situation contrasts with that previously reported for spring upwelling (Crespo et al., 2011), when diatoms were present in upwelled waters on the shelf and *Synechococcus* accounted for a minor fraction ($0.16 \pm 0.13\%$) of AB. Primary production ($624 \pm 585 \text{ mg C m}^{-2} \text{ d}^{-1}$), 5-fold lower than in the spring upwelling, was particularly low ($246 \pm 104 \text{ mg C m}^{-2} \text{ d}^{-1}$) at the IPC, though higher values ($1616 \pm 389 \text{ mg C m}^{-2} \text{ d}^{-1}$) were measured on the western shelf in front of the Rías Baixas. Phytoplankton gross growth rates (μ) showed the same pattern of extremely low values in the IPC ($\mu = 0.06 \text{ d}^{-1}$) and the highest ($\mu = 0.27 \text{ d}^{-1}$) in the waters leaving the Rías Baixas. Whereas AB exceeded heterotrophic biomass (HB) in this upwelling system in spring, at the end of summer HB was higher than AB. However, both biomasses were linearly related with a HB:AB ratio of 1 and the excess of HB (intercept = $2.8 \pm 0.5 \text{ g C m}^{-2}$) being similar to the rather constant bacterial biomass ($2.7 \pm 0.5 \text{ g C m}^{-2}$). This fact points to the key role played by bacteria during the oligotrophic conditions at the end of summer. Bacteria can maintain the functioning of the microbial community through processing dissolved organic matter and supplying food to heterotrophs and mixotrophs.

Keywords: Microbial community, nanoflagellates, *Synechococcus*, bacteria, heterotrophy, coastal upwelling, NW Iberian Peninsula

1. Introduction

It is well known that the size-structure and composition of the planktonic community influence the dynamics of the microbial food web and consequently biogeochemical cycles in the ocean (Azam et al., 1983). Although multiple stages of the microbial plankton community can be found in the sea, those commonly found in oligotrophic and eutrophic areas set the limits for all possible situations (Legendre and Rassoulzadegan, 1995). Picoplankton cells dominate in the microbial community of the oligotrophic ocean (Buck et al., 1996), where the microbial loop drives recycling (Azam et al., 1983). On the contrary, large plankton and export (Malone, 1980) characterise the structure and functioning of the microbial community in eutrophic areas.

According to this scheme, coastal upwelling systems should be classified as eutrophic areas because export and dominance of large cell-sized plankton are among the most obvious characteristics of these zones. However, coastal upwelling systems are essentially dynamic, with temporal and spatial variability modulating the structure and composition of the microbial community (Lorenzo et al., 2005; Sherr et al., 2005). Thus, picoplankton from the nearby oligotrophic ocean can be advected shoreward during downwelling, while intense upwelling can expand to the adjacent oligotrophic ocean the presence or dominance of large plankton. Coastline (bays and capes) and hydrographic singularities (eddies and fronts) can also modify the structure and composition of the microbial community through the changes they cause in upwelling and downwelling intensity (Figueiras et al., 2006; Crespo et al., 2011). The microbial community in eastern boundary upwelling ecosystems also experiences variability over the growth season, which is basically characterised by the continuous increase in the abundance of heterotrophs (Figueiras et al., 2002). Furthermore, in a changing world, with climate change altering the upwelling intensity and the extension of the upwelling

season (Álvarez-Salgado et al., 2008), it is reasonable to expect future changes in the structure and composition of the microbial community following changes in oceanographic conditions (Cermeño et al., 2008; Li et al., 2009; Morán et al., 2010). Therefore, knowledge of the structure and composition of the microbial community in these highly productive zones is not only important to understand the functioning of the microbial food web during the several upwelling-downwelling phases that these systems undergo, it is also relevant to track future modifications induced by environmental changes.

The NW Iberian Peninsula (Fig. 1a) constitutes the northern limit of the eastern boundary Large Marine Upwelling Ecosystem that extends from the African coast (10°N) to the European coast (44°N) (Wooster et al., 1976). Upwelling is seasonal at these latitudes (42-44 °N), with northerly winds causing upwelling from spring to autumn and southerly winds inducing downwelling in winter (Wooster et al., 1976; Figueiras et al., 2002). Upwelling and downwelling force contrasting circulation patterns in the four bays located on the west of the NW Iberian Peninsula, collectively known as Rías Baixas (Fig. 1a). Upwelling induces positive estuary-like circulation in the Rías causing the export of material to the nearby shelf, whereas the reverse circulation occurs during downwelling, with shelf water entering the Rías (Figueiras et al., 2002).

The seasonal upwelling-downwelling transition at the end of summer coincides with the advection toward the coast of warm and saline oceanic water, which then flows northward on the slope forming the Iberian Poleward Current (IPC) (Haynes and Barton, 1990; Álvarez-Salgado et al., 2003; Torres and Barton, 2006). During this transition, with downwelling not totally established yet and upwelling relics still present in the region, a subsurface chlorophyll maximum (SCM) is typically observed in a well

1 stratified water column (Castro et al., 1997; Torres and Barton, 2006; Crespo et al.,
2 2008). The microplankton community at the NW Iberian region during this transition
3 time has been relatively well studied and now it is known that it is basically composed
4 of small ($\sim 30\ \mu\text{m}$) dinoflagellates (Castro et al., 1997; Crespo et al., 2008), because
5 downwelling precludes diatom growth and confines large dinoflagellates ($>60\ \mu\text{m}$)
6 inside the Rías (Fermín et al., 1996; Crespo et al., 2007). There is also some evidence
7 pointing to the dominance of pico- and nanoplankton within the autotrophic community
8 (Lorenzo et al., 2005; Rodríguez et al., 2006). In contrast, the trophic structure of the
9 whole microbial community is relatively unknown.

10 Here we present a detailed analysis of the structure and composition of the
11 autotrophic and heterotrophic microbial communities at the end of the upwelling season
12 in this coastal upwelling region. The results, derived from data collected nearly 20 years
13 ago (September 1991), are compared to those obtained from a contrasting upwelling
14 situation in the spring of the same year (Crespo et al., 2011). This information is
15 essential to understand the functioning of the pelagic system and develop an accurate
16 plankton functional type (PTF) model for the region (Le Quéré et al., 2005) as a
17 baseline to which future environmental changes can be referred.

18 19 **2. Materials and methods**

20 *2.1. Sampling*

21 The northwestern Iberian margin (Fig. 1a) was sampled at 37 stations located along 8
22 sections perpendicular to the shoreline at the end of the summer in 1991 (September 10
23 to 18). Starting at the northernmost section (Stns 11 to 14), sampling was carried out
24 with a conductivity-temperature-depth (CTD) probe (SBE 9/11) equipped with a Sea
25 Tech fluorometer and a rosette with 12 PVC Niskin bottles. Seawater samples were

collected from the CTD upcasts at several depths, between surface and bottom on the shelf and from surface to ~100 m at the other stations. To ensure that the SCM was sampled, the depths were selected after inspecting the fluorescence profiles. Nutrients and chlorophyll *a* (chl *a*) were determined at all sections, while the structure and composition of the microbial community were analysed at 18 stations situated along alternate sections (black symbols in Fig. 1a). Primary production was estimated from determinations of photosynthesis-irradiance relationships and light transmittance in the water column at 10 stations (triangles in Fig. 1a) randomly distributed on the sampling area. This distribution was due to the constraint imposed by sampling during day light hours. Seawater for photosynthesis-irradiance determinations was collected at 3-4 depths within the photic layer.

2.2. Ekman transport, nutrients and chl *a*

The cross-shelf Ekman transport (Fig. 1b) was estimated according to Bakun (1973) using geostrophic winds off Cape Finisterre (43°N, 11° W).

Nitrate and ammonium concentrations ($\mu\text{mol kg}^{-1}$) were determined onboard by segmented flow analysis (Technicon AAII system) according to Hansen and Grasshoff (1983).

For chl *a* determinations 100 ml of seawater were filtered through 25 mm GF/F filters under low vacuum pressure. The filters were immediately frozen at -20°C until pigment extraction in 90% acetone over 24 h in dark at 4°C. Chl *a* concentrations (mg m^{-3}) were determined using a Turner Designs fluorometer calibrated with pure chl *a* (Sigma).

2.3. Microbial plankton biomass

Sub-samples of 10 ml were processed according to Porter and Feig (1980) to enumerate autotrophic and heterotrophic pico- ($<2\ \mu\text{m}$) and nanoplankton ($2\text{--}20\ \mu\text{m}$) by epifluorescence microscopy (see details in Figueiras et al., 2006). *Prochlorococcus*, which can be found in shelf waters of the region at this time of the year (Rodríguez et al., 2006), were not well discriminated as autotrophs with this technique due to their low fluorescence. Thus, the abundance of picoautotrophs can be underestimated and the abundance of heterotrophic bacteria overestimated (Buck et al., 1996). Bacterial abundance was converted to carbon biomass according to Lee and Furhmann (1987). Dimensions were taken for the other groups and their biovolumes were calculated assuming a spherical shape. Cell carbon conversions were made according to Verity et al. (1992) for pico- and nanoflagellates and Bratbak and Dundas (1984) for *Synechococcus*.

Sub-samples of 100 ml preserved in Lugol's iodine were settled in composite sedimentation chambers and examined with an inverted microscope to identify and count microplankton ($20\text{--}200\ \mu\text{m}$) species of diatoms, dinoflagellates and ciliates. The small species were enumerated from two transects scanned at $\times 250$ and $\times 400$, whereas the less abundant larger species were counted from the whole slide scanned at $\times 100$. Epifluorescence microscopy and the list of species given by Lessard and Swift (1986) were used to differentiate between phototrophic and heterotrophic species. Dimensions were taken to estimate cell and plasma volumes by approximation to the nearest geometrical shape. The plasma volume of diatoms and the biovolume of dinoflagellates were converted to cell carbon following Strathmann (1967). Carbon biomass of ciliates was estimated according to Putt and Stoecker (1989).

2.4. Irradiance at the sea surface and in the water column

The photosynthetic active radiation (PAR , $\lambda = 400-700$ nm, $\mu\text{mol quanta m}^{-2} \text{ s}^{-1}$) during the cruise was measured on deck with a Li-190SA cosine-sensor at 1 min intervals. These readings were then hourly integrated to obtain the daily distribution of incident irradiance at sea surface.

The PAR irradiance in the water column was measured at the 10 stations where primary production was determined. Readings were taken at 2 m intervals with a spherical quantum sensor LI-193SA. Incident irradiance at the sea surface (PAR_{0+}) was recorded at the same time with a cosine-sensor LI-190SA. Then, the PAR profile in the water column (PAR_z) was fitted to:

$$PAR_z = PAR_{0-} e^{-kz} \quad (1)$$

where k (m^{-1}) is the light attenuation coefficient and PAR_{0-} is the irradiance just below the sea surface. The ratio PAR_{0-}/PAR_{0+} provides the light transmittance at the air-sea interface.

2.5. Photosynthesis-irradiance relationships, primary production and phytoplankton growth rates

The photosynthesis-irradiance experiments were conducted in linear incubators (Arbones et al., 2000) illuminated on the front side with tungsten-halogen lamps (50W, 12 V) and refrigerated with circulating seawater. Each incubator housed 14 subsamples collected in 75 ml Corning tissue flasks, which were inoculated with 1.85×10^5 Bq (5 μCi) of ^{14}C -bicarbonate. One of these flasks was covered with aluminium foil and placed at the end of the incubator to check dark carbon fixation. Following incubation

(2 h), samples were filtered under low vacuum pressure through 25 mm GF/F filters that were then exposed to HCl fumes for 12 h to remove unincorporated inorganic ^{14}C . Radioactivity was measured with a liquid scintillation counter using the external standard and the channel ratio methods to correct for quenching.

In absence of photo-inhibition the data were fitted to the model of Webb et al. (1974):

$$P_z^B = P_m^B [1 - \exp(-\alpha^B PAR / P_m^B)] \quad (2)$$

or to the model of Platt et al. (1980) when photo-inhibition was evident:

$$P_z^B = P_s^B [1 - \exp(-\alpha^B PAR / P_s^B)] \exp(-\beta^B PAR / P_s^B) \quad (3)$$

to derive the chlorophyll-specific light saturated rate of photosynthesis (P_m^B , mg C mg chl $a^{-1} h^{-1}$), light saturated rate of photosynthesis in the absence of photo-inhibition (P_s^B , mg C mg chl $a^{-1} h^{-1}$), light-limited slopes (α^B , mg C mg chl $a^{-1} h^{-1}$ ($\mu\text{mol quanta m}^{-2} \text{ s}^{-1}$) $^{-1}$) and coefficients of photo-inhibition (β^B , mg C mg chl $a^{-1} h^{-1}$ ($\mu\text{mol quanta m}^{-2} \text{ s}^{-1}$) $^{-1}$). For the cases with photo-inhibition P_m^B was estimated when $\partial P^B / \partial PAR = 0$ (Platt et al., 1980). P_z^B is the chlorophyll-specific rate of carbon fixation (mg C mg chl $a^{-1} h^{-1}$) at each sampled depth.

The daily incident irradiance at sea surface and the vertical profiles of PAR and chl a concentration were combined in equations (2) and (3) to estimate the daily water-column primary production (PP, mg C $m^{-2} d^{-1}$) down to 1% of incident irradiance at sea surface. Integration was carried out at hourly steps and 1 m intervals.

Phytoplankton gross growth rate (μ , d^{-1}) was estimated at each sampled depth where carbon fixation was determined:

$$\mu = \ln \left(1 + \frac{dC/dt}{C_{ph}} \right) \quad (4)$$

where dC/dt is the volumetric carbon fixation ($\text{mg C m}^{-3} \text{ d}^{-1}$) and C_{ph} is phytoplankton carbon (mg C m^{-3}).

3. Results and discussion

3.1. Meteorological and oceanographic conditions: the onset of the IPC

Northerly and southerly winds, coastal upwelling and coastal downwelling favourable respectively, alternated in the region in September 1991 (Fig. 1b). Thus, although the cruise began during strong downwelling ($Q_x = -595 \text{ m}^3 \text{ s}^{-1} \text{ km}^{-1}$) on September 10th, the situation gradually reverted until the sampling ended under moderate upwelling on September 18th ($Q_x = 270 \text{ m}^3 \text{ s}^{-1} \text{ km}^{-1}$). This is a wind regime commonly registered at the NW Iberian Peninsula during the seasonal upwelling-downwelling transition that leads to the onset of the IPC, while upwelling vestiges are still observed in the region (Castro et al., 1997; Álvarez-Salgado et al., 2003; Torres and Barton, 2006). The onset of the IPC regularly occurs with the advection of warm and saline oceanic surface water toward the west coast (Crespo and Figueiras, 2007; Relvas et al., 2007) as found during this study. Thus, warm ($>20^\circ\text{C}$; Fig. 2a) and saline (>35.7 ; Fig. 2b) waters observed at the most oceanic stations correspond to thermohaline properties characteristic of IPC advected waters. However, we observed no signal of these salty waters at the most northern sections, indicating that the IPC was not

1 completely developed. Therefore, we were capturing the onset of the IPC, as observed
2 in detail by Haynes and Barton (1990) at the same time of the year. In fact, traces of
3 upwelled waters with temperature $<19^{\circ}\text{C}$ were seen on the shelf at the northern
4 locations (Fig. 2a). On the western shelf, the less saline waters (Fig. 2b) with high
5 chlorophyll concentrations (Fig 2c) were exported from the Rías Baixas through their
6 positive estuary-like circulation forced by upwelling (Figueiras et al. 2002; Crespo et
7 al., 2007).

8 Although other effects may contribute, the wind-forced upwelling-downwelling
9 response describes to first order the predominant dynamics over the NW Iberian shelf.
10 Interactions with spatial inhomogeneity of the wind field and coastal orography (Torres
11 et al., 2003) are detectable in the variability of the shelf regime, but are of secondary
12 importance. Remote forcing of upwelling via coastally-trapped waves, although not
13 observed in the region (e.g. Relvas et al., 2007) presumably must play a minor role.
14 However, the direct wind response explains most of the variability observed in the NW
15 Iberian upwelling system.

16 The whole region showed a strongly stratified water column (Figs. 3a-c) with a well-
17 developed nitracline at $\sim 50\text{ m}$ (Figs. 3d-f) to which a SCM was associated (Figs. 3g-i).
18 The SCM, however, surfaced on the shelf (Figs. 3g-i) and this outcropping was
19 reflected in the distribution of chl *a* concentration at sea surface (Fig. 2c), with higher
20 values near the coast than in the IPC. Nevertheless, the chl *a* distribution recorded along
21 the section situated in front of the Rías Baixas (Fig. 3i) points to the discontinuity
22 between the SCM and the surface maximum nearest the coast. This suggests that
23 phytoplankton exported from the Rías Baixas significantly contributed to the chl *a*
24 maximum registered at stn 71 (Fig. 3i).

While nitrate was depleted in the upper water column (Figs. 3d-f), ammonium was relatively important at sea surface (Fig. 2d), with concentrations being particularly high ($>1 \mu\text{mol kg}^{-1}$) at the front separating IPC from shelf waters.

3.2. Autotrophic plankton biomass: the importance of picophytoplankton

The autotrophic community was largely dominated by autotrophic nanoflagellates (ANF), with significant contributions of *Synechococcus*, autotrophic picoflagellates (APF) and autotrophic dinoflagellates (AD) (Table 1). Diatoms and autotrophic ciliates (AC) accounted for a minor fraction ($< 2\%$) of the total autotrophic biomass (AB). As the biomasses of these four dominant phytoplankton groups were linearly correlated ($r \geq 0.53$; $P < 0.001$; $n = 135$), spatial variations in AB (Figs. 4a-c) matched variations in the biomasses of these groups, mainly variations in ANF biomass (Figs. 4d-f). However, there were small differences along the sections. Thus, the biomass of *Synechococcus* was higher near the coast at all sections sampled (Figs. 4g-i; Table 1), whereas APF biomass was slightly higher in coastal waters at the northernmost section (Figs. 4j-l; Table 1). ANF (Figs. 4d-f) and AD (Figs. 4m-o) showed highest biomass values in surface waters off the Rías Baixas (Stn 71; Figs. 4f and o). Small species (*Heterocapsa niei*, *Prorocentrum minimum*) dominated in the AD community at all sites (Crespo et al., 2008), while the presence of large pigmented dinoflagellates (e.g. *Ceratium fusus*, *Prorocentrum micans*, *Dinophysis acuminata*) was restricted to surface waters of low salinity off the Rías Baixas (Crespo et al., 2008). These dinoflagellates were exported from the Rías to contribute to the maximum of biomass recorded at the surface of station 71 (Fig. 4o). It is also remarkable the low AB ($< 80 \text{ mg C m}^{-3}$) carried out by surface oceanic waters $> 20^\circ \text{C}$ (Fig. 4). However, the presence of *Prochlorococcus* at the NW Iberian margin, which were not satisfactorily enumerated on this occasion, is

1 tightly coupled to the onset of the IPC (Rodríguez et al., 2006), and so it could be
2 present in the warm and saline surface oceanic water advected to the region.
3 Nevertheless, the NW Iberian Peninsula is near the northern limit of the distribution of
4 *Prochlorococcus* in the Atlantic Ocean (Buck et al., 1996) and their abundance in IPC
5 waters is always typically low ($<10^3$ cells ml^{-1} ; Calvo-Díaz et al., 2004), around 2-3 fold
6 lower than the abundance of *Synechococcus* recorded during this cruise in the surface
7 waters of the IPC.

8 Despite this apparent coastal-ocean variability in the biomass of autotrophic
9 plankton, the water-column integrated biomass values (Table 1) at coastal (the two most
10 inshore stations) and oceanic stations were not significantly different ($P > 0.05$, t-test for
11 two samples). This situation markedly contrasts with that found in the region during
12 spring upwelling (Fig. 5), when a coastal environment with relatively low AB (8.7 ± 4.1
13 g C m^{-2}) and high contribution of diatoms ($27 \pm 17\%$) was clearly discernible from an
14 oceanic environment with higher AB ($24.5 \pm 9.6 \text{ g C m}^{-2}$) and almost complete
15 dominance of ANF ($89 \pm 6\%$). Another major difference between these two seasons
16 refers to the importance of autotrophic picoplankton (*Synechococcus* and APF).

17 Whereas in spring upwelling (Crespo et al., 2011) picophytoplankton represented a
18 minor fraction of AB at both coastal and oceanic domains ($2.3 \pm 0.8\%$), especially
19 evident for *Synechococcus* ($0.2 \pm 0.1\%$), their contribution at the end of summer ($32 \pm$
20 9%) was appreciably higher (Table 1; Fig. 5). In addition, the biomass of autotrophic
21 picoplankton at the end of summer ($2.24 \pm 0.88 \text{ g C m}^{-2}$) was substantially higher than
22 in spring upwelling ($0.42 \pm 0.40 \text{ g C m}^{-2}$; Fig. 5), and this difference between the two
23 seasons was particularly noticeable for *Synechococcus*, with an average biomass 35
24 times higher at the end of summer ($1.41 \pm 0.61 \text{ g C m}^{-2}$, Table 1) than in spring ($0.04 \pm$
25 0.05 g C m^{-2}). In contrast, the biomass of ANF was lower and less variable at the end of

summer ($3.69 \pm 1.16 \text{ g C m}^{-2}$; Table 1) than in spring upwelling ($12.99 \pm 10.34 \text{ g C m}^{-2}$), when differences in ANF biomass between coastal and ocean domains were considerable (Fig. 5).

Higher biomass of autotrophic picoplankton in the rather oligotrophic waters at the end of summer than in the eutrophic conditions of spring, agrees well with similar situations described for other coastal upwelling systems (Sherr et al., 2005) and other coastal waters in general (Echevarría et al., 2009). It is also consistent with the macroecological pattern (Li, 2002) that describes an increase in autotrophic picoplankton abundance and a decrease in the abundance of larger cells following the degree of oligotrophy. In such a scenario, with dominance of nano- and picophytoplankton (Table 1), nitrate depletion (Figs. 3d-f) and relatively high concentrations of ammonium in the surface layer (Fig. 2d), considerable recycling within the microbial community and hence low f-ratios (Probyn, 1992) would be expected.

3.3. Carbon to chl *a* ratio

The distribution of AB (Figs. 4a-c) roughly followed that of chl *a* (Figs. 3g-i), according to a phytoplankton carbon to chl *a* (Cph:chl *a*) ratio of 170 ± 6 (Fig. 6). This relationship slightly improves to give a somewhat higher Cph:chl *a* ratio of 178 ± 5 when 6 points (3 from the northernmost section and other 3 points from the southernmost section) were removed from the regression. One of these points corresponds to a surface sample at the northernmost section with a high phytoplankton carbon value (305 mg C m^{-3}) and a relatively low chl *a* concentration (0.52 mg m^{-3}). The other 5 points belong to samples from the SCM (2 at the northernmost section and 3 at the southernmost section) with low Cph:chl *a* ratios (Fig. 6).

1 Ratios like this have been reported for oligotrophic areas in the open ocean (e.g.
2 Buck et al., 1996) and they are commonly interpreted as a signal of nutrient limitation
3 for phytoplankton growth (Marañón, 2005). However, high Cph:chl *a* ratios can also
4 result from mixotrophic nutrition (Skovgaard et al., 2000), which is widespread within
5 pigmented pico- (Zubkov and Tarran, 2008) and nanoplankton (Unrein et al., 2007).
6 Therefore, mixotrophic nutrition at the NW Iberian margin under the oligotrophic
7 conditions at the end of summer should not be disregarded. This is a nutrition mode that
8 has also been suggested for pigmented nanoplankton in the region during spring
9 (Crespo et al., 2011) to explain the high Cph:chl *a* found at the oceanic environment.

10 The existence of a unique Cph:chl *a* ratio for the whole area contrasts with the
11 situation found during spring upwelling (Fig. 5), when near-shore waters with direct
12 influence of upwelling showed a Cph:chl *a* ratio (38 ± 3) considerably lower than
13 oceanic waters (157 ± 8). The occurrence of a unique ratio, without differences between
14 surface and SCM, also suggests that photo-acclimation in the SCM was not important,
15 which in turn points to the recent formation of the SCM. Then, the SCM formation
16 probably occurred through the subduction of coastal surface waters forced by the
17 advection of surface oceanic water with low chl *a* concentration (Fig. 2c) and low
18 phytoplankton biomass (Fig. 4). If this is the process responsible for the formation of
19 the SCM, the corollary is that the phytoplankton populations, including
20 picophytoplankton, found in the area were not transported from the open ocean. On the
21 contrary, picophytoplankton must be viewed as a typical component of the
22 phytoplankton community in coastal upwelling systems that get more importance under
23 non upwelling conditions (Lorenzo et al., 2005; Sherr et al., 2005).

3.4. Primary production and phytoplankton growth

Primary production (Table 2), was highly variable ($624 \pm 585 \text{ mg C m}^{-2} \text{ d}^{-1}$), with the lowest value ($128 \text{ mg C m}^{-2} \text{ d}^{-1}$) occurring in the IPC waters at the southwest corner (stn 85; Fig 1a) and the highest ($1891 \text{ mg C m}^{-2} \text{ d}^{-1}$) in waters leaving the shelf at 42.5°N (stn 65, Fig. 1a). The low PP ($246 \pm 104 \text{ mg C m}^{-2} \text{ d}^{-1}$) measured in the IPC (Table 2) is typical of oligotrophic waters (Frazel and Berberian, 1990; Lorenzo et al. 2004) and was similar to values previously reported for the IPC at the NW Iberian Peninsula (Tilstone et al., 2003), when microbial community respiration exceeds carbon fixation (Teira et al., 2001). Primary production in western shelf waters ($1616 \pm 389 \text{ mg C m}^{-2} \text{ d}^{-1}$) lies within the range of values reported for stratified or not strong upwelling conditions on the shelf (Tilstone et al., 2003; Lorenzo et al., 2005). However, it was appreciably lower than the values commonly measured during upwelling (Teira et al., 2001; Tilstone et al., 2003). The PP determined at the northern shelf not affected by the IPC ($593 \pm 319 \text{ mg C m}^{-2} \text{ d}^{-1}$) can be considered as representative of summer stratified conditions (Teira et al., 2001; Tilstone et al., 2003), when the microbial community tends to be in metabolic balance (Teira et al., 2001). Overall, PP during this cruise at the end of summer was 5-fold lower than PP measured during spring upwelling in the same area (Fig. 5).

Although phytoplankton gross growth rates were generally higher at sea surface ($0.17 \pm 0.10 \text{ d}^{-1}$) than at SCM ($0.06 \pm 0.04 \text{ d}^{-1}$), they were extremely low in the whole photic layer. As the rates estimated using C_{ph} actually determined (equation 4) were not significantly different ($P = 0.61$; t-test for paired samples) to the rates estimated with C_{ph} derived from the C_{ph}:chl *a* ratio of 170 (Fig. 6), the mean gross growth rates ($\bar{\mu}$, d^{-1}) in the photic layer ($Z_{1\%}$) were estimated as:

$$\bar{\mu} = \frac{1}{Z_{1\%}} \int_{Z_{1\%}} \ln \left(1 + \frac{dC/dt}{C_{ph}} \right) \quad (5)$$

where C_{ph} was derived from the profiles of chl a concentration at each station where PP was measured.

The resulting gross growth rates varied from a minimum value of 0.06 d^{-1} at stn 72 in the IPC to a maximum value of 0.27 d^{-1} in the shelf waters in front of the Rías (stn 63), and they are similar to the gross growth rates recorded in the oceanic domain in spring (Fig. 5), where mixotrophic nutrition was presumably high (Crespo et al., 2011). These low rates, which correspond to doubling times of the phytoplankton community varying between 11.25 and 2.54 days (Table 2), are within the range of values found in the oligotrophic waters of subtropical gyres (Marañón, 2005) where nutrient limitation of phytoplankton growth is assumed. However, it must be noted that these gross growth rates were estimated for photosynthetic carbon fixation, and consequently they could be lower than the achieved rates if mixotrophic nutrition of ANF and APF occurs.

3.5. Microbial heterotrophic biomass and heterotrophic to autotrophic biomass ratio

The heterotrophic community was dominated by heterotrophic nanoflagellates (HNF) and bacteria (Table 3), with the two groups accounting for $77 \pm 10\%$ of the total integrated heterotrophic biomass (HB). Heterotrophic picoflagellates (HPF) and heterotrophic dinoflagellates (HD) also showed an appreciable contribution ($\sim 10\%$) to HB (Table 3). Unlike the strong reduction (57%) experienced by AB when compared to the AB registered in the region during spring upwelling, the drop in HB (22%) was substantially lower (Fig. 5). This leads to a system in which HB exceeded AB, in agreement with the expected situation for low nutrient conditions (Duarte et al., 2000),

with heterotrophic metabolism predominating in the microbial community (Teira et al., 2001). The reduction in HB occurred through the strong decrease (56%) in the HNF biomass (Table 3; Fig. 5), because bacterial biomass remained invariant (Table 3; Fig. 5) and the biomass of the other heterotrophic groups increased.

Although the distribution of HD (Figs. 7m-o) and heterotrophic ciliates (data not shown) show a preference of these two groups for coastal waters, the distributions of the other groups (Figs. 7d-l) as well as that of HB (Figs. 7a-c) were basically similar to the distributions of autotrophic plankton (Fig. 4). In fact, both HB and AB covaried (Fig. 8) according to a linear relationship with slope of 1, which suggests a rapid response of heterotrophic plankton to variations in the autotrophic components of the microbial community. On other hand, the intercept of this relationship ($\sim 30 \text{ mg C m}^{-3}$) that accounts for the excess of HB recorded in the system (Tables 1 and 3), roughly coincides with the average biomass of bacteria in a water column 100 m depth (Table 3). This constant biomass indicates that there is top-down control of bacteria and they can be viewed as playing a pivotal role at the NW Iberian margin under the oligotrophic conditions at the end of summer. Bacteria, through processing the dissolved organic matter, mainly exported from the Rías, and accumulated in the upper water column over the growth season (Álvarez-Salgado et al., 1999), can supply food to other heterotrophs and mixotrophs.

4. Summary

The microbial plankton community in the NW Iberian margin at the end of the upwelling season appeared dominated by heterotrophs, with nano- and picoplankton cells prevailing in both the autotrophic and heterotrophic community. This composition of the microbial community seems to be characteristic of the system, rather than being

advection from the adjacent oligotrophic ocean by downwelling. It markedly contrasts with the structure and composition of the microbial community recorded during spring upwelling, which was characterised by the dominance of autotrophs with presence of diatoms in a near-shore band directly affected by upwelling (Crespo et al., 2011). The high C_{ph}: chl *a* ratios, the low primary production and the extremely low phytoplankton growth rates recorded at the end of the upwelling season points to a microbial community with predominance of heterotrophic processes, as the excess of heterotrophic biomass and the tight coupling of variations in autotrophic and heterotrophic biomasses suggest. The relatively constant bacterial biomass, which roughly matches the excess of heterotrophic biomass recorded in the system, points to the key role played by bacteria at the NW Iberian upwelling during the regularly observed oligotrophic conditions at the end of summer. In this context, bacteria can maintain the microbial community functioning through processing the dissolved organic matter accumulated over the upwelling season and so supplying food to heterotrophs and mixotrophs. Under a possible future scenario with lower upwelling intensity and shorter upwelling season in the region (Álvarez-Salgado et al., 2008), the structure and composition of the microbial community reported here, with heterotrophy dominating in the upper layer, will prevail in the zone. In the actual regime, the export from the Rías could supply the organic matter needed to maintain the functioning of the microbial community. However, with possible decreases in upwelling intensity and shortness of the upwelling season, the microbial community of the Rías Baixas, now predominantly autotrophic (Arbones et al., 2008), could eventually shift to heterotrophy leading to oligotrophy in the entire NW Iberian margin.

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Figure legends

Fig. 1. (a) The NW Iberian margin showing the location of the four Rías Baixas on the west coast and the stations sampled. The sections selected for vertical representations in Figs. 3, 4 and 7 are marked with black lines. Black symbols denote the stations where the structure and composition of the microbial plankton community were determined. The ten stations where primary production and phytoplankton growth rates were estimated are marked with black and open triangles. (b) Cross-shore Ekman transport component ($-Q_x$) from September 1 to 20, 1991, estimated at 43° N, 11° W. Positive/negative values of $-Q_x$ indicate offshore/onshore transport (upwelling/downwelling) of surface waters.

Fig. 2. Sea surface distributions of (a) temperature, (b) salinity, (c) chl *a* concentration and (d) ammonium concentration.

Fig. 3. Vertical distributions at 3 selected sections (see Fig. 1a) of (a to c) temperature, (d to f) nitrate concentration and (g to i) chl *a* concentration. Section 11 to 14 was sampled on 10th September, section 51 to 55 on 13th September and section 71 to 75 on 16th September.

Fig. 4. Vertical distributions at 3 selected sections (see Fig. 1a) of (a to c) total autotrophic biomass (AB), (d to f) biomass of autotrophic nanoflagellates (ANF), (g to i) biomass of *Synechococcus* (SYN), (j to l) biomass of autotrophic picoflagellates (APF) and (m to o) biomass of autotrophic dinoflagellates (AD). Dashed line corresponds to the isotherm of 20° C at the oceanic stations (see Fig. 2a). Section 11 to 14 was sampled on 10th September, section 51 to 55 on 13th September and section 71 to 75 on 16th September.

Fig. 5. Cartoon summarising the contrasting structures of the microbial plankton community found in the NW Iberian upwelling at the end of summer (this work) and in spring upwelling (Crespo et al., 2011), when two domains (coastal and oceanic) were evident. The total area of the circles, as well as the areas of the portions representing the different plankton groups in each circle, is proportional to the respective biomass values. In the upper panel, representing the autotrophic community, the average value for phytoplankton carbon to chlorophyll *a* ratios (Cph:chl *a*), integrated primary production (PP) and phytoplankton growth rates estimated from carbon fixation (μ) are also given. Syn: *Synechococcus*; APF: autotrophic picoflagellates; ANF: autotrophic nanoflagellates; Diat: diatoms; AD: autotrophic dinoflagellates; AC : autotrophic ciliates. The heterotrophic community, with lesser variability than the autotrophic community, is represented in the lower panel. HPF: heterotrophic picoflagellates; HNF: heterotrophic nanoflagellates; HD: heterotrophic dinoflagellates; HC: heterotrophic ciliates.

Fig. 6. Phytoplankton carbon (Cph) to chl *a* relationships including all samples (solid line) $Cph = (170 \pm 6) chl\ a, r^2 = 0.68, n = 135$; and removing 6 samples (open circles) from the regression (dashed line) $Cph = (178 \pm 5) chl\ a, r^2 = 0.79$. See text for more details.

Fig. 7. Vertical distributions at 3 selected sections (see Fig. 1a) of (a to c) total heterotrophic biomass (HB), (d to f) biomass of heterotrophic nanoflagellates (HNF), (g to i) biomass of heterotrophic bacteria, (j to l) biomass of heterotrophic picoflagellates (HPF) and (m to o) biomass of heterotrophic dinoflagellates (HD). Dashed line corresponds to the isotherm of 20° C at the oceanic stations (see Fig. 2a). Section 11 to 14 was sampled on 10th September, section 51 to 55 on 13th September and section 71 to 75 on 16th September.

Fig. 8. Autotrophic biomass (AB) *versus* heterotrophic biomass (HB). Solid line, $HB = (28.17 \pm 5.40) + (1.02 \pm 0.05)AB, r^2 = 0.73, n = 135$, defines the relationship considering all points, which is not markedly different from the relationships obtained after excluding 3 samples (open circles, dashed line), $HB = (30.15 \pm 3.06) + (0.97 \pm 0.03)AB, r^2 = 0.90$.

Table 1. Integrated (surface-bottom on the shelf and 0-100 m at the ocean) values of chl *a* concentration (mg m⁻²) and autotrophic plankton biomass (g C m⁻²) at each station sampled. AB: total autotrophic biomass; *Syn*: *Synechococcus*; APF: autotrophic picoflagellates; ANF: autotrophic nanoflagellates; Diat: diatoms; AD: autotrophic dinoflagellates; AC: autotrophic ciliates. Average values and contribution of each group to AB are also given.

Stn	chl <i>a</i>	AB	<i>Syn</i>	APF	ANF	Diat	AD	AC
11	45	6.79	1.60	1.78	2.71	0.03	0.52	0.15
12	24	4.31	0.77	0.39	1.79	0.03	1.32	0.02
13	22	6.81	0.75	0.28	4.94	0.02	0.81	0.01
14	29	5.00	0.70	0.30	3.19	0.05	0.75	0.00
31	34	6.86	1.12	0.76	4.25	0.09	0.58	0.06
32	52	10.38	2.12	1.35	6.17	0.05	0.64	0.06
33	41	6.97	2.18	0.95	3.07	0.02	0.74	0.01
34	32	5.61	1.14	0.59	3.24	0.07	0.55	0.02
51	42	6.36	1.46	0.83	2.83	0.26	0.42	0.56
52	48	8.25	2.51	1.02	3.21	0.02	1.05	0.43
53	39	7.89	2.00	0.74	3.91	0.02	1.02	0.19
54	39	6.89	1.12	0.62	4.31	0.02	0.75	0.06
55	41	6.37	0.91	0.67	4.16	0.03	0.59	0.01
71	67	10.89	1.52	0.86	5.68	0.84	1.89	0.09
72	32	7.87	2.20	0.96	3.64	0.02	1.00	0.05
73	36	8.98	1.83	1.18	4.49	0.04	1.42	0.02
74	43	4.79	0.74	0.96	2.34	0.02	0.71	0.02
75	35	4.96	0.63	0.85	2.53	0.02	0.91	0.01
Average (SD)	39 (11)	7.00 (1.83)	1.41 (0.61)	0.84 (0.37)	3.69 (1.16)	0.09 (0.19)	0.87 (0.37)	0.10 (0.16)
Percentage of AB (SD)			19.7 (6.1)	12.1 (5.1)	52.8 (9.8)	1.1 (1.9)	12.9 (5.7)	1.4 (2.3)

Table 2. Primary production (PP, mg C m⁻² d⁻¹), mean phytoplankton growth rate in the photic layer ($\bar{\mu}$, d⁻¹) and doubling time (1/k, days) in the 3 hydrographic zones identified at the end of summer in the NW Iberian margin. $k = \ln(2)/\bar{\mu}$.

Zone	Stn	PP	$\bar{\mu}$	1/k
North	12	225	0.10	7.25
	14	770	0.15	4.63
	24	783	0.20	3.45
IPC	34	365	0.11	6.13
	44	345	0.23	3.04
	55	215	0.10	7.08
	72	177	0.06	11.25
	85	128	0.13	5.22
West-Rías	63	1341	0.27	2.54
	65	1891	0.22	3.21

Table 3. Integrated (surface-bottom on the shelf and 0-100 m at the ocean) values of microbial heterotrophic plankton biomass (g C m⁻²) at each station sampled. HB: total heterotrophic biomass; HPF: heterotrophic picoflagellates; HNF: heterotrophic nanoflagellates; HD: heterotrophic dinoflagellates; HC: heterotrophic ciliates. Average values and contribution of each group to HB are also given.

Stn	HB	Bacteria	HPF	HNF	HD	HC
11	13.03	3.10	1.90	3.36	4.07	0.60
12	10.91	2.41	0.95	3.50	3.75	0.30
13	9.21	2.20	0.89	5.19	0.84	0.09
14	7.58	2.02	0.60	4.28	0.48	0.20
31	9.30	2.05	0.81	5.18	0.66	0.61
32	12.40	3.01	1.28	7.11	0.30	0.70
33	8.56	2.76	1.09	4.31	0.22	0.19
34	7.79	2.65	0.82	3.74	0.39	0.19
51	8.35	2.26	0.79	3.47	1.53	0.31
52	9.48	3.48	0.99	4.12	0.50	0.40
53	10.37	3.02	0.74	5.07	1.05	0.49
54	9.60	2.66	0.85	5.31	0.47	0.30
55	9.21	2.30	0.96	5.53	0.35	0.06
71	12.80	3.45	1.07	7.00	1.01	0.27
72	10.90	3.21	1.25	5.03	1.15	0.27
73	10.92	2.95	1.26	6.25	0.13	0.33
74	7.98	2.46	1.21	3.92	0.12	0.27
75	7.06	2.09	1.03	3.56	0.10	0.28
Average	9.75	2.67	1.03	4.77	0.95	0.32
(SD)	(1.79)	(0.48)	(0.29)	(1.17)	(1.15)	(0.17)
Percentage of HB (SD)		27.7 (4.0)	10.6 (2.4)	49.4 (9.1)	9.1 (9.6)	3.3 (1.5)

Fig. 1
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Fig. 2
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Fig. 3
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