

1 Role of zooplankton dynamics for Southern Ocean phytoplankton biomass and global
2 biogeochemical cycles

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1 Abstract

2 Global ocean biogeochemistry models currently employed in climate change projections use highly
3 simplified representations of pelagic food webs. These food webs do not necessarily include critical
4 pathways by which ecosystems interact with ocean biogeochemistry and climate. Here we present a
5 global biogeochemical model which incorporates ecosystem dynamics based on the representation of
6 ten plankton functional types (PFTs); six types of phytoplankton, three types of zooplankton, and
7 heterotrophic procaryotes. We improved the representation of zooplankton dynamics in our model
8 through (a) the explicit inclusion of large, slow-growing macrozooplankton, and (b) the introduction
9 of trophic cascades among the three zooplankton types. We use the model to quantitatively assess the
10 relative roles of iron versus grazing in determining phytoplankton biomass in the Southern Ocean
11 High Nutrient Low Chlorophyll (HNLC) region during summer. When model simulations do not
12 include macrozooplankton grazing explicitly, they systematically overestimate Southern Ocean
13 chlorophyll biomass during the summer, even when there is no iron deposition from dust. When
14 model simulations include a slow-growing macrozooplankton and trophic cascades among three
15 zooplankton types, the high chlorophyll summer bias in the Southern Ocean HNLC region largely
16 disappears. Our model results suggest that the observed low phytoplankton biomass in the Southern
17 Ocean during summer is primarily explained by the dynamics of the Southern Ocean zooplankton
18 community, despite iron-limitation of phytoplankton community growth rates. This result has
19 implications for the representation of global biogeochemical cycles in models as zooplankton faecal
20 pellets sink rapidly and partly control the carbon export to the intermediate and deep ocean.

22 1 Introduction

23 Phytoplankton, zooplankton and heterotrophic bacteria (including both *Bacteria* and *Archaea*, herein
24 called 'bacteria') in the oceans control important ecosystem processes and services (Ducklow, 2008),
25 including primary, secondary and export production. Primary production, i.e. the production of
26 organic matter by photoautotrophs using inorganic nutrients, can be either particulate and serve as
27 food for heterotrophs, from protists to fish larvae, or dissolved and used by bacteria. Secondary
28 production, the fraction produced by zooplankton grazing on phytoplankton, other zooplankton, or
29 organic detritus, serves as food for larger organisms in the ocean, including fish and mammals.
30 Export production, the fraction of primary production that sinks below the surface mixed layer, exerts
31 an influence on marine biogeochemistry and climate as sinking organic matter remineralised to
32 inorganic matter at depths becomes isolated from the atmosphere for decades to centuries. Export
33 production responds primarily to the activity of large plankton, particularly the production and
34 sinking of faecal pellets of zooplankton (e.g. copepods and euphausiids) as well as the aggregation of
35 diatoms, for example, during intense blooms. Export production reduces the surface concentration of
36 inorganic carbon and maintains atmospheric CO₂ about 200 ppm lower than it would be in the
37 absence of biological activity (Maier-Reimer et al., 1996). In contrast, bacteria and small zooplankton
38 (e.g. heterotrophic flagellates and ciliates) remineralise and recycle organic matter in the upper
39 ocean, thus reducing the quantity of organic matter that is exported. These ecosystem processes are
40 controlled by the state of the environment (e.g. temperature, light, available nutrients, vertical
41 mixing), and are modulated by the ecosystem structure of the planktonic community.

42 Dynamic Green Ocean Models have been developed and used in global biogeochemical studies to
43 understand and quantify the interactions between marine ecosystems and the environment. In these
44 models, phytoplankton and zooplankton are grouped by taxa into plankton functional types (PFTs)
45 according to their specific and unique roles in marine biogeochemical cycles (Hood et al., 2006; Le
46 Quéré et al., 2005). Although generally only a small number of PFTs are treated explicitly, their
47 inclusion has been shown to improve the realism of model simulations. For example, the explicit
48 inclusion of diatoms in marine ecosystem models is required to reproduce the observed response to

1 natural or purposeful iron fertilisation in the ocean (Aumont and Bopp, 2006), and observed changes
2 in export production during glacial cycles (Bopp et al., 2002). The representation of diazotrophs (i.e.
3 N₂-fixing organisms) is necessary to simulate the feedbacks between iron and the nitrogen
4 inventories of the ocean (Moore et al., 2006; Moore and Doney, 2007) and to reproduce observed
5 N:P ratios (Weber and Deutsch 2010; 2012), of coccolithophores to simulate large blooms of
6 phytoplankton (i.e. chlorophyll) biomass (Gregg and Casey, 2007) and phytoplankton succession
7 (Gregg et al., 2003), and of *Phaeocystis* to reproduce the ecosystem structure in the Southern Ocean
8 (Wang and Moore, 2011).

9 Fewer studies have examined the role of different zooplankton PFTs in global ocean
10 biogeochemistry, even though there are zooplankton physiological datasets (e.g. Hirst and Bunker,
11 2003; Straile, 1997). The simulation of phytoplankton biomass was improved in published studies
12 when more mechanistic parameterisations of zooplankton dynamics constrained by observations
13 were included in a global model (Buitenhuis et al., 2006; Buitenhuis et al., 2010). Similarly, the
14 seasonal cycle of phytoplankton (Aita et al. 2003) and the open-ocean oxygen depletion (Bianchi et
15 al. 2013) were improved when the influence of zooplankton vertical migration was included in global
16 biogeochemical models. The choice of the grazing formulation in particular was found to influence
17 phytoplankton diversity (Prowse et al., 2012; Vallina et al., 2014b) and the resulting food web
18 dynamics (Sailley et al., 2013; Vallina et al., 2014a), and to have implications for energy flow to
19 higher trophic levels (Stock et al., 2014).

20 Zooplankton can influence the fate of exported materials through several processes, including
21 grazing, repackaging of organic matter in faecal pellets, and the vertical migrations and transport of
22 carbon and nutrients into the mesopelagic zone (e.g. Stemmann et al., 2000; Steinberg et al. 2008).
23 Furthermore, there are important interactions among grazing, nutrient cycles, and environmental
24 conditions as was shown in studies based on regional models and observations in the equatorial
25 Pacific (Landry et al., 1997; Price et al., 1994), North Pacific (Frost, 1991), the Atlantic (Daewel et
26 al., 2014; Steinberg et al., 2012) and the Southern Ocean (Banse, 1995; Bishop and Wood, 2009).
27 The importance of grazing was also highlighted during iron enrichment experiments (Henjes et al.,
28 2007; Latasa et al., 2014), in part explaining why some experiments led to increased carbon export
29 and others did not (Martin et al., 2013). Thus, a more explicit representation of different zooplankton
30 PFTs in global models could provide important clues for the functioning of marine biogeochemistry.

31 Here, we present a new global ocean biogeochemistry model with ten PFTs. The parameterisation of
32 vital rates associated with these PFTs is based on an extensive synthesis of published information on
33 growth rates and other relevant parameters. We use the model to examine a long-standing paradox in
34 biological oceanography: the low phytoplankton biomass in the Southern Ocean despite the high
35 concentrations of macronutrients. This has been attributed to lack of iron (Fe) because of the distance
36 to continental dust sources (Geider and La Roche, 1994; Martin, 1990). Increases in phytoplankton
37 biomass have been produced in more than a dozen open ocean iron fertilisation experiments (Boyd
38 and al., 2007; Smetacek et al., 2012). The influx of Fe has been proposed as a driver for the
39 drawdown of atmospheric CO₂ during glaciations (Kohfeld et al., 2005; Watson et al., 2000), and
40 intentional Fe-fertilisation has been considered as a means to both geo-engineer climate (Rickels et
41 al., 2012) and to sell carbon credits (Tollefson, 2012). However, ocean biogeochemistry models that
42 explicitly include the effect of Fe-limitation on phytoplankton growth fail to reproduce the low Chl
43 biomass observed during summer in the Southern Ocean (Aumont and Bopp, 2006; Dutkiewicz et al.,
44 2005; Le Quéré et al., 2005; Moore et al., 2004). This raises the question of the relative control
45 exerted by Fe-limitation on biomass versus that exerted by the grazing pressure of zooplankton
46 (Banse, 1996; Price et al., 1994) and more generally on the suitability of the current generation of
47 models to explore ecosystem – climate interactions. Our study addresses this question directly.

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2 Methods

2.1 Model description and development

The PlankTOM10 Dynamic Green Ocean Model is a global ocean biogeochemistry model that includes plankton ecosystem processes based on the representation of ten PFTs and their interactions with the environment. PlankTOM10 incorporates six autotrophic and four heterotrophic PFTs: picophytoplankton (pico-eukaryotes and non N₂-fixing cyanobacteria such as *Synechococcus* and *Prochlorococcus*), N₂-fixers (*Trichodesmium* and N₂-fixing unicellular cyanobacteria), coccolithophores, mixed-phytoplankton (e.g. autotrophic dinoflagellates and chrysophytes), diatoms, colonial *Phaeocystis*, bacteria (here used to subsume both heterotrophic *Bacteria* and *Archaea*), protozooplankton (e.g. heterotrophic flagellates and ciliates), mesozooplankton (predominantly copepods), and crustacean macrozooplankton (euphausiids, amphipods, and others, called ‘macrozooplankton’ for simplicity; Fig. 1). Gelatinous macrozooplankton are not included in the model. Diversity within groups are not considered, and the physiological parameters for each PFT are the same everywhere in the ocean, although some are dependent on environmental conditions (i.e. nutrients, light, food, temperature).

The current version of the PlankTOM10 model was developed from the model of Buitenhuis et al. (2013a), using the strategy for regrouping PFTs described by Le Quéré et al. (2005). It does not include new equations for growth and loss terms compared with previous versions of the PlankTOM model, but it includes an additional trophic level in the zooplankton PFTs (i.e. macrozooplankton). Parameterisations are based on more data related to the vital rates of individual PFTs, where new information was available. Previous studies have shown that model results are highly sensitive to PFT growth rates (Buitenhuis et al. 2006; 2010), and considerable effort was made to constrain these rates using observations from LaRoche and Breitbarth (2005), Bissinger et al. (2008), Buitenhuis et al. (2008), Sarthou et al. (2005), Schoemann et al. (2005), Rivkin and Legendre (2001), Buitenhuis et al. (2010), Hirst and Bunker (2003), and Hirst et al. (2003).

The complete set of model equations and parameter values are provided in the Supplementary Information. Here, we describe the elements that are most important for the analysis of the Southern Ocean and the strategy used to determine parameter values for PFT growth and loss processes.

PlankTOM10 simulates the growth of ten PFTs in response to environmental conditions. The PFT biomasses are produced by the model for each grid box based on the growth and loss term equations presented in Supplementary Material. The model includes three detrital pools: large and small particulate organic matter, and semi-labile dissolved organic matter. The sinking rate of large particles is based on the mineral (ballast) content of particles following Buitenhuis et al. (2001), while the sinking rate of small particles is constant at 3 m d⁻¹. The model includes full cycles of carbon (C), oxygen (O₂), and phosphorus (P), which are assimilated and released by biological processes at a constant ratio of 122:172:1 (Anderson and Sarmiento, 1994). Phytoplankton and particulate organic matter have a variable Fe/C ratio, while zooplankton and bacteria have a fixed ratio of 2e⁻⁶, which is lower than the minimum phytoplankton Fe/C ratio (Schmidt et al. 1999). Zooplankton and bacteria release excess iron. The model also includes a full cycle of silica (Si) and calcite (CaCO₃) as in Maier-Reimer (1993), and simplified cycles for Fe and nitrogen (N). CO₂ and O₂ are exchanged with the atmosphere using the gas exchange formulation of Wanninkhof (1992). The Fe cycle is represented as in Aumont and Bopp (2006). Iron is deposited with dust particles using the monthly fields of Jickells et al. (2005), the Fe content of dust is assumed to be 3.5% everywhere. We assume an Fe solubility from dust of 1% (Jickells et al. 2005). Iron is also delivered to the ocean via river fluxes following the outflow scheme of da Cunha et al. (2007) with 95% sedimentation in estuaries. Dissolved inorganic nitrogen (DIN) is the sum of nitrate and ammonium. The N:P ratio of organic processes is set to the Redfield ratio of 16:1. N₂-fixers can use N₂ and thus have access to unlimited N from the atmosphere.

1 The growth rate parameters for the ten PFTs in PlankTOM10 are based on a compilation of growth
2 rates as a function of temperature (Section 2.2). Phytoplankton PFT growth rates are also limited by
3 light and inorganic nutrients (P, N, Si, and Fe) using a dynamic photosynthesis model that represents
4 the two-way interaction between photosynthetic performance and Fe/C and Chl/C ratios (Buitenhuis
5 et al. 2013a). Light limitation is constrained by the slope of the photosynthesis-irradiance curve (α)
6 and the maximum Chl/C ratio (θ_{\max}). We could not distinguish PFT-specific values for α (Geider et
7 al., 1997) and used a mean value of $1.0 \text{ mol C m}^2 (\text{g Chl mol photons})^{-1}$ for all PFTs. Observed θ_{\max}
8 for diatoms are systematically higher than those of other PFTs (Geider et al., 1997). There are too
9 few direct observations to parameterize θ_{\max} for other PFTs, so we fitted the observations (Geider et
10 al., 1997) for θ_{\max} to the maximum growth rate (μ^{\max}) presented in that paper. The fit showed θ_{\max}
11 increasing with growth rate ($n=19$, $p=0.02$). We thus used a θ_{\max} higher than average for *Phaeocystis*
12 and diatoms, and a lower than average θ_{\max} for N_2 -fixers.

13 We used a two-step approach to define the nutrient limitation parameters, which are not well
14 constrained by observations. Firstly, we assigned initial PFT-specific half-saturation values to each
15 phytoplankton PFT based on literature-derived values, using the value for a similar-sized PFT when
16 PFT-specific information was not available. We then examined the covariations of surface Chl
17 concentrations with the limiting nutrient concentrations as shown in Figure 3, and adjusted the
18 magnitude of the half-saturation parameters of phytoplankton PFT to approximately fit the
19 observations, keeping the ratios of k-half values between phytoplankton PFTs approximately the
20 same as the initial ratios. With this approach, we use the observed k-half values as an initial starting
21 point but tune the model to match the emerging properties highlighted in Figure 3.

22 Initial values for the half-saturation concentrations of P (k_P) and N (k_N) for phytoplankton growth
23 rates were based on observations. For N_2 -fixers, coccolithophores and diatoms, the half-saturation
24 values for growth were computed using the half-saturation values of uptake reported in Riegman et
25 al. (1998), LaRoche et al. (2005), and Sarthou et al. (2005) multiplied by the minimum/maximum
26 N:C ratio (0.33) to account for the acclimation of nutrient saturated vs. nutrient limited growth
27 (Morel, 1987). For picophytoplankton, reported values for the half-saturation extend over three
28 orders of magnitude. We assigned low half-saturation values as these organisms grow even under
29 very low nutrient conditions (Timmermans et al., 2005). For mixed phytoplankton, we assigned a
30 value intermediate between picophytoplankton and diatoms. For *Phaeocystis*, we used half-saturation
31 values that characterise colonies (Schoemann et al., 2005). The selected set of parameter values
32 shown in Figure 3 are reported in Table 2.

33 Iron uptake was computed using a cell quota model (Buitenhuis and Geider, 2010; Geider et al.,
34 1997), where the Fe uptake by phytoplankton PFTs is explicitly regulated by the light conditions. The
35 three parameters needed are the minimum, the maximum and the optimal Fe quotas. The minimum
36 and maximum quotas were set at the same value of 2.5 and 20 $\mu\text{mol Fe/mol C}$ for all PFTs based on
37 the analysis of Buitenhuis and Geider (2010). The optimal quota was set to the minimum quota plus
38 $2 * \mu_{20}^{\max}$ based on (Sunda and Huntsman, 1995) for all PFTs. In addition, phytoplankton PFTs also
39 respond to the concentration of Fe in water which is parameterised with a half saturation constant.
40 The half saturation of Fe uptake (k_{Fe}) is lower for picophytoplankton (Timmermans et al., 2005) than
41 other phytoplankton, and higher for N_2 -fixers (LaRoche and Breitbarth, 2005) and diatoms (Sarthou
42 et al., 2005). Intermediate values for k_{Fe} have been reported for the other phytoplankton PFTs (Le
43 Vu, 2005; Schoemann et al., 2005). The selected set of parameter values after adjustments produces
44 no systematic covariation between Chl and Fe, as observed (Fig. 3, Table 2).

45 The half-saturation parameters of zooplankton grazing rate were initially based on the relationship
46 between metabolic rates and body volume of Hansen et al. (1997). We used the same approach as for
47 nutrient limitation of the phytoplankton PFTs, and adjusted the half-saturation parameters for grazing
48 based on the observed covariations between surface Chl concentrations and zooplankton biomass

1 (Fig 3). The selected set of parameter values that approximately fit the observed covariations in
2 Figure 3 is reported in Table 2.

3 Zooplankton food preferences were assigned based on predator-prey size ratio (Table 3), as there
4 were insufficient data to determine these parameters directly. This approach assumes that
5 protozooplankton generally have a high preference for bacteria and a low preference for diatoms, that
6 mesozooplankton have a higher preference for protozooplankton and a low preference for N₂-fixers
7 and bacteria, and macrozooplankton have a lower preference for N₂-fixers, picophytoplankton and
8 bacteria than other groups. We assume that all zooplankton graze on organic particles (Table 3) but
9 prefer to graze on other PFTs. The weighing factors influenced primarily the biomass of the prey and
10 predators, but had little influence on their geographic distribution. We thus used the model results on
11 biomass (Table 4) to guide the size of the relative preferences among PFTs for each grazer.

12 The gross growth efficiency (the part of grazing that is incorporated into biomass) was defined based
13 on the mean across available observations: 0.21 for bacteria (data from Rivkin and Legendre, 2001),
14 and 0.29, 0.25, and 0.30 for protozooplankton, mesozooplankton and macrozooplankton, respectively
15 (data from Straile, 1997). Respiration and mortality parameters were based on observations from
16 Buitenhuis et al. (2010) for protozooplankton, Buitenhuis et al. (2006) for mesozooplankton, and
17 Moriarty (2013) for macrozooplankton. The temperature-dependence of respiration and mortality was
18 fitted to all data as for the growth rate (Section 2.2), except for the mortality of macrozooplankton
19 and mesozooplankton. There are nine observations on macrozooplankton mortality and we tuned this
20 term based on the resulting biomass. The fitted relationship for the mortality of mesozooplankton was
21 reduced by a factor of ~2 to account for the explicit mortality from macrozooplankton represented in
22 the model. This correction preserves the temperature-dependence of mortality, but it recognises that
23 explicit grazing by macrozooplankton already takes place in the model, which does not represent the
24 grazing by other organisms (e.g. salps, fish larvae). In total, grazing accounts for 2/3 to 3/4 of the
25 mortality of mesozooplankton (Hirst and Kiorboe, 2002).

26 **2.2 Growth rates as a function of temperature**

27 The most important trait that distinguishes the various PFTs is the rate at which they grow under
28 different conditions (Buitenhuis et al., 2006; Buitenhuis et al., 2010). We compiled maximum growth
29 rates as a function of temperature (Table 1). We fit an exponential growth relationship to the
30 observations by optimising the relation $\mu^T = \mu_0 * Q_{10}^{T/10}$ where T and μ^T are the observed temperature
31 and associated growth rate, μ_0 is the growth at 0°C, and Q_{10} is the derived temperature-dependence of
32 growth (Table 1). The parameter values for μ_0 and Q_{10} were estimated by minimising the error,
33 quantified as the least squares cost function $\Sigma((\mu^T - \mu_{obs}^T) / \mu_{obs}^T)^2$. Normalising to observations helps
34 ensure a good fit of μ^T in cold waters where growth rates are low. We used exponential growth,
35 rather than a temperature-optimal growth, to avoid biases caused by the lack of observations for some
36 PFTs at low or high temperatures. The p-value of a linear regression between observations and the
37 exponential fit (Table 1) provides a measure of how well the relationship is constrained by the
38 observations. The fit assigns equal weight for all the data, rather than following the 99% quantile
39 (e.g. Eppley, (1972); Bissinger et al., (2008)) to provide a better representation of the mean
40 community for each PFT.

41 Growth rate parameters estimated with this method are well constrained (p-values < 0.05) for seven
42 of the ten PFTs, including all of the heterotrophic PFTs (Table 1). There are insufficient data to
43 provide significant constraints on the growth rates of N₂-fixers (p = 0.76), and some uncertainty in
44 the growth data for coccolithophores (p = 0.06) and *Phaeocystis* (p = 0.23; Table 1). However, the
45 growth of N₂-fixers is less than that of other phytoplankton PFTs (Fig. 2), and the fitted relationship
46 produces μ^T less than that of other PFTs despite these uncertainties. An exponential function may not
47 be appropriate for growth rates of coccolithophores and *Phaeocystis* (Schoemann et al. 2005). The
48 growth rate of coccolithophores was overestimated at low temperatures due to high growth rates at

1 20°C and the absence of observations for temperatures below 5°C. We reduced the fitted growth rate
2 of coccolithophores linearly to 0 below 10°C to match the observed reduced coccolithophore biomass
3 in cold regions (O'Brien et al., 2013).

4 **2.3 Covariation between Chl and nutrients or zooplankton**

5 We used relationships between observed concentrations of Chl and both inorganic nutrients (e.g.
6 NO₃, PO₄ and Fe), and zooplankton biomasses (protozooplankton, mesozooplankton and
7 macrozooplankton; Fig. 3) to provide additional constraints on model parameters. Specifically, we
8 used observations for *in situ* NO₃ and PO₄ concentrations from the World Ocean Atlas 2009; *in situ*
9 Fe concentration data from Tagliabue et al. (2012); protozooplankton biomass data from Buitenhuis
10 et al. (2010); mesozooplankton biomass data from Buitenhuis et al. (2006); macrozooplankton
11 biomass data from Atkinson et al. (2004) and Moriarty et al. (2013). All the data were binned into
12 1x1° grid boxes. Most observations are for the surface ocean. Mesozooplankton and
13 macrozooplankton data are from depth-integrated tows of typically 200 m depth and may
14 underestimate surface concentrations (by a factor 1.5-2 based on our model simulations). All data are
15 monthly except for mesozooplankton, which are seasonal. Chl concentration is from SeaWiFS
16 satellite averaged over 1998-2009 and interpolated to the same grid. The model output was averaged
17 over the same time period, and sampled for the same month and on the same grid box as the
18 observations. The data intervals were chosen to include approximately the same number of grid
19 boxes, except for macrozooplankton where the lowest interval was set to 0 – 0.05 μmol C L⁻¹ because
20 of the large number of grid boxes with very low macrozooplankton concentration. Ten concentration
21 intervals were used for the nutrients (Fig. 3).

22 Chlorophyll concentrations covary with NO₃ concentrations at <3 μmol L⁻¹, and with PO₄ in the
23 range 0.3-0.5 μmol L⁻¹ (Fig. 3; Spearman ranked correlations for data in the 25-75% interquartile
24 range gives r = 0.72 for NO₃ and r = 0.73 for PO₄). These relationships are consistent with our
25 understanding of the growth limitation of phytoplankton in the subtropics, where NO₃ and PO₄
26 concentrations are low. There is no observed covariation between Chl and Fe concentration (r = -
27 0.16). The strongest covariations are between Chl and protozooplankton at concentrations <0.6 μmol
28 C L⁻¹ (r = 0.83) and between Chl and mesozooplankton at concentrations <0.3 μmol C L⁻¹ (r = 0.77).
29 There is no covariation between Chl concentration and macrozooplankton biomass (r = -0.19; Fig. 3).
30 We use these relationships to tune the growth limitations parameters in the model, so that the
31 functional relationships between Chl and nutrients or zooplankton are close to the observed
32 relationships overall.

33 **2.4 Simulations**

34 PlankTOM10 is coupled to the Ocean General Circulation Model (OGCM) NEMO version 3.1
35 (NEMOv3.1). We used the global configuration (Madec and Imbard, 1996), which has a resolution of
36 2° of longitude and a mean resolution of 1.5° of latitude, with enhanced resolution up to 0.3° in the
37 tropics and at high latitudes. The model resolves 30 vertical levels, with 10 m depth resolution in the
38 upper 100 m. NEMOv3.1 calculates vertical diffusion explicitly and represents eddy mixing using the
39 parameterisation of Gent and McWilliams (1990). The model thus generates its own mixed-layer
40 dynamics and associated mixing based on local buoyancy fluxes and winds. NEMOv3.1 is coupled to
41 a dynamic-thermodynamic sea-ice model (Timmermann et al., 2005).

42 PlankTOM10 is initialised from observations of dissolved inorganic carbon (DIC) and alkalinity
43 from Key et al. (2004), O₂ and nutrients from Garcia et al. (2006a) and Garcia et al. (2006b), and
44 temperature and salinity from the World Ocean Atlas 2005 (Antonov et al., 2006; Locarnini et al.,
45 2006). Fe is initialised with a constant concentration of 0.6 nmol Fe L⁻¹ north of 30°S and 0.2 nmol
46 Fe L⁻¹ in the Southern Ocean, consistent with observations (Parekh et al., 2005; Tagliabue et al.,
47 2012). The PFTs equilibrated within three years and were not influenced by initialisation. The model

1 is forced by daily winds and precipitation from the ECMWF interim reanalysis (Simmons et al.,
2 2006) from 1989 to 2009. Results for standard simulations are averaged over 1998-2009. A series of
3 sensitivity tests are presented for the model parameters that influence the key results the most.

4 To understand the interaction pathways among ecosystems, biogeochemistry and climate, we
5 developed a simplified version of the model that included only six PFTs (PlankTOM6) (Fig. 1).
6 PlankTOM6 is identical to PlankTOM10 except that the growth rates of N₂-fixers, mixed-
7 phytoplankton, *Phaeocystis*, and macrozooplankton are zero, and the mortality of the
8 mesozooplankton is increased to account for the lack of macrozooplankton predation until the point
9 when primary production is at its maximum. Given the otherwise similar model structure,
10 parameters, initialisation and simulation protocol, comparison of results from PlankTOM6 and
11 PlankTOM10 provide information on the specific roles of zooplankton dynamics in the model.

12 13 **3 Results**

14 **3.1 Temperature and size – dependence of PFT growth rates**

15 The data show systematic patterns in growth rates that differ among PFTs. The growth rates of all
16 PFTs increase with increasing temperature, but not to the same extent (Fig. 2). The growth rate of
17 phytoplankton PFTs increases with PFT size, from 0.15 d⁻¹ for N₂-fixers to 1.87 d⁻¹ for *Phaeocystis*,
18 and the growth rate of heterotrophic PFTs decreases with size, from 1.22 d⁻¹ for bacteria to 0.19 d⁻¹
19 for macrozooplankton (Table 1). The sign of the relationship between growth rate and size between
20 phytoplankton PFTs is the opposite of the sign of this relationship within specific PFTs, including
21 diatoms (Sarthou et al., 2005), picophytoplankton (Chen and Liu, 2010) and coccolithophores
22 (Buitenhuis et al., 2008).

23 **3.2 Ecosystem properties in the PlankTOM10 model**

24 PlankTOM10 reproduces the main characteristics of observed surface Chl, with high concentrations
25 in the high latitudes and low concentrations in the subtropics, higher Chl concentration in the
26 Northern compared to the Southern hemisphere, and in the South Atlantic compared to the South
27 Pacific Ocean (Fig. 4). The global biogeochemical fluxes simulated by PlankTOM10 are generally
28 below or at the low end of the range of observed values (in Table 4, ‘model’ and ‘data’, respectively),
29 with global primary production of 42.4 PgC yr⁻¹, export production of 7.6 PgC yr⁻¹, export of CaCO₃
30 and SiO₂ of 0.4 PgC yr⁻¹ and 2.9 PgSi yr⁻¹, respectively, and N₂ fixation of 165 TgN yr⁻¹.

31 PlankTOM10 produces distinctive geographical distributions of carbon biomasses among PFTs (Fig.
32 5). About a third of the phytoplankton biomass occurs as picophytoplankton, followed in descending
33 abundance by diatoms and *Phaeocystis*, mixed-phytoplankton, coccolithophores and N₂-fixers (Table
34 4). This distribution is broadly consistent with observations (Buitenhuis et al., 2013b) but the
35 simulated phytoplankton biomass is generally on the low side of the observational range, which is
36 consistent with the results of the global biogeochemical fluxes. The simulated biomass of
37 coccolithophores is overestimated (i.e. 0.077 PgC compared with 0.001-0.032 PgC) although CaCO₃
38 export is underestimated, suggesting either that the model calcification or aggregation rates are too
39 low or that zooplankton calcifiers contribute significantly to CaCO₃ export.

40 The model underestimates bacterial biomass by a factor of 10 compared with observations. This
41 possibly reflects the fact that the model only represents highly active bacteria and a substantial
42 fraction of observed biomass is from low activity and ghost cells. The model underestimates
43 protozooplankton by a factor of 1.5-5 (in absolute value) or 2-3 (as a fraction of total biomass value)
44 compared to observations (Table 4). This discrepancy could be caused by the underestimation of
45 bacterial biomass, as bacteria are an important source of food for protozooplankton. The simplified
46 representation of the range of protozooplankton grazers in a single PFT representing both

1 heterotrophic nanoflagellates and microzooplankton could also play a role. Simulated
2 mesozooplankton biomass is only slightly below the observed range, while simulated
3 macrozooplankton biomass is within the observed range, although the uncertainty here is large (0.010
4 – 0.64 PgC). Overall the balance is slightly skewed towards relatively more biomass than observed in
5 the larger zooplankton (53% compared to 3-47%) compared to the smaller zooplankton groups (13%
6 compared to 27-31%; Table 4).

7 The geographic distribution of each simulated PFT is also distinctive (Figs. 6-7). Satellite data
8 products indicate that small phytoplankton (picophytoplankton and N₂-fixers) are generally dominant
9 in the tropics, haptophytes (coccolithophores and *Phaeocystis*) in mid to high latitudes, and diatoms
10 in high latitudes (Alvain et al., 2005; Brewin et al., 2010). The simulated phytoplankton distribution
11 generally matches the distribution inferred from satellite normalised radiance (Fig. 6), except in the
12 temperate zones where observations suggest a balance between picophytoplankton and haptophytes
13 and the model shows a dominance of haptophytes. PlankTOM10 also reproduces the locations of
14 blooms of colonial *Phaeocystis* and coccolithophores (Fig. 7). The simulated geographic distributions
15 of zooplankton PFTs are particularly distinctive, with protozooplankton abundant in the tropics and
16 subtropics, mesozooplankton at high latitudes of both hemisphere, and macrozooplankton with high
17 biomass in the North Pacific and South Atlantic and along the coasts (Fig. 5).

18 The marine ecosystem as a whole appears to function realistically: Mesozooplankton grazing on
19 phytoplankton is somewhat overestimated relative to the 5.5 Pg/y estimated by Calbet 2001, so they
20 have taken over the role of principal herbivores. Possibly the faster turnover rates of small copepods
21 are overrepresented in the observational data on mesozooplankton, leading to a trophic position of
22 mesozooplankton somewhat too low in the foodchain. Export production, phytoplankton biomass and
23 metazoan zooplankton biomass are realistic in the model, leading to realistic seasonal cycles, but the
24 regenerated part of primary production is underestimated, concomitant with low protozooplankton
25 biomass, which impacts the model on shorter timescales of days.

26 **3.3 Comparison of PlankTOM6 and PlankTOM10**

27 PlankTOM10 and PlankTOM6 generally produce similar results in surface Chl concentration,
28 nutrient distribution, primary and export production (Fig. 8), except that PlankTOM6 fails to
29 reproduce the observed low Chl concentration in summer in the Southern Ocean (Fig. 4; Section 3.4).
30 The overall difference between the two models, quantified statistically using a Taylor distribution
31 (Taylor, 2001), are less than 0.1 in either correlation or normalised standard deviation (Fig. 8).
32 PlankTOM10 does slightly better than PlankTOM6 for the distribution of Chl, primary and export
33 production, but slightly worse for the distribution of silica and nitrate, with similar performance for
34 phosphate (Fig. 8). These differences are small in part because of the short duration of the
35 simulations presented here (20 years), which allow equilibration of the ocean surface only. The
36 models are generally similar also in their representations of the distribution of biomass among
37 phytoplankton PFTs, with most of biomass being in picophytoplankton in both models (Fig. 9 and
38 Table 4). However PlankTOM6 allocates more biomass to protozooplankton compared to
39 PlankTOM10, though PlankTOM6 is still at the low end of observed concentrations (Table 4).

40 The failure of PlankTOM6 to reproduce the observed low Chl concentration in the Southern Ocean
41 during summer is further highlighted in Fig. 10, which shows the seasonal cycle of mean Chl for the
42 Northern Hemisphere and the Southern Ocean, where it is most pronounced. In PlankTOM6, the
43 seasonal cycle in the North and South are very similar, with the slightly lower concentrations in
44 Southern Ocean during summer caused by a slightly deeper summer time mixed-layer depth (29m
45 compared to 19m). In contrast in PlankTOM10, the seasonal cycle of Chl in the South is smaller and
46 concentrations are always below those in the North, as is the case for observations. As PlankTOM6
47 and PlankTOM10 have identical physical environments (including mixed-layer depth), the North-

1 South differences are due to ecosystem structure. In the following sections, we focus our analysis to
2 the model parameters that influence the low Chl concentration in the Southern Ocean the most.

3 **3.4 Role of zooplankton dynamics for HNLC regions**

4 The observed phytoplankton biomass, including the low Chl concentrations in HNLC regions,
5 reflects the balance between phytoplankton growth and loss. Phytoplankton growth rates vary with
6 temperature, light, and nutrient supply, whereas losses result mainly from grazing by zooplankton,
7 respiration, cell death, sinking to depth, and dilution by vertical mixing. Any process that reduces the
8 net rate of increase of phytoplankton biomass (i.e. differences between growth and loss) may lead to
9 low residual Chl concentration. For example, Platt et al. (2003a) showed that deep mixing by wind
10 dilutes Chl in the surface layer and reduces the average irradiance experienced by the phytoplankton.
11 This results in low growth rate and demand for nitrate; the conditions generally observed in HNLC
12 regions. Here we further examine the consequences of high zooplankton-mediated grazing losses.

13 We use the North/South ratio in surface Chl concentration as a metric to quantify model
14 performance, focusing on the Pacific Ocean where the contrast between the Northern Hemisphere
15 and the Southern Ocean is most pronounced. This metric is simple and easy to quantify with data
16 (geographic locations: boxes in Fig. 4).). Satellite observations indicate a North/South Chl ratio of
17 2.16 ± 0.35 (1998-2009 mean \pm 2SD of annual values). To ensure that the ratio is not affected by
18 potential biases in the SeaWiFS Southern Ocean data (Johnson et al., 2013), we also used *in situ* data
19 from the World Ocean Atlas which indicates a similar North/South Chl ratio of 2.0. This ratio is 1.72
20 ± 0.051 in the PlankTOM10, and 1.21 ± 0.074 in the PlankTOM6 simulations (Fig. 11). Controlling
21 factors on this ratio are examined here through a set of sensitivity tests.

22 **3.4.1 Role of trophic level and top zooplankton**

23 We tested the specific effect of macrozooplankton on Chl by running four additional model
24 experiments (Fig. 11): in the Z1 simulation, we added macrozooplankton to PlankTOM6, in Z2 we
25 parameterised the top grazer in PlankTOM6 using the same growth and loss rate parameters as
26 macrozooplankton, in Z3 we removed macrozooplankton from PlankTOM10, and in Z4 we
27 parameterised the top grazer in PlankTOM10 using the same growth and loss rate parameters as
28 mesozooplankton. These sensitivity studies were identical to the PlankTOM10 (or PlankTOM6)
29 simulation in all other respects. Experiments Z1 and Z2 both include macrozooplankton, but in
30 different food-web positions. These experiments maintain a high North/South Chl ratio of 1.64 and
31 1.46, respectively (Fig. 11). Experiments Z3 and Z4 did not include macrozooplankton but had
32 grazing structures as in the standard PlankTOM6 and PlankTOM10 models, the North/South Chl
33 ratio was 1.26 and 1.11 respectively. These four experiments show that the presence in the model of
34 slow-growing zooplankton, such as macrozooplankton, plays a pivotal role in determining the
35 relative average concentrations of Chl in the Northern versus Southern hemisphere (difference
36 between PlankTOM6 and both Z1 and Z2). More realistic patterns are achieved by including a third
37 zooplankton food-web compartment (higher ratio in Z1 than in Z2) and three additional
38 phytoplankton compartments (higher ratio in PlankTOM10 than in Z1).

39 **3.4.2 Role of macrozooplankton growth rate**

40 We examined the impact of macrozooplankton grazing in sensitivity tests in which the grazing rate of
41 macrozooplankton was varied within the range of the observed growth rates (Fig. 2; Table 1). These
42 simulations show that macrozooplankton grazing rate has a strong influence on the Chl North/South
43 ratio (Fig. 12). The PlankTOM10 simulation that uses the mean growth rate from observations
44 (Section 2.2) produces results that are closest to the observed North/South Chl ratio. When the
45 grazing rate is decreased (by up to 2σ), the macrozooplankton biomass decreases by over 50% and
46 the North/South Chl ratio decreases from 1.72 to 1.05. When the grazing rate is increased, the
47 macrozooplankton biomass decreases because of pressure on the food sources (Fig. 12) and the Chl

1 North/South ratio also decreases. These simulations suggest that the observed Chl North/South
2 distributions are a consequence of trophic balances among PFTs.

3 **3.4.3 Role of atmospheric iron deposition**

4 We tested the relative role of atmospheric iron deposition compared with grazing for the North/South
5 Chl distribution by applying five different dust deposition scenarios, all (except one) with realistic
6 but different regional distributions, to the PlankTOM10 and PlankTOM6 models: D0 is an extreme
7 case with no atmospheric dust deposition (where phytoplankton use iron sources from deep waters),
8 D1 dust deposition including the effect of dust particle size on iron solubility (Mahowald et al.,
9 2009), and D2-D4 iron deposition using the three distinct dust fields (Ginoux et al., 2001; and Luo,
10 2003; Tegen et al., 2004) averaged by Jickells et al. (Jickells et al., 2005). The simulated North/South
11 Chl ratios vary from 1.62 and 1.85 in these experiments (Fig. 11). These differences are smaller than
12 the differences between the PlankTOM10-like (1.46-1.85) and the PlankTOM6-like simulations
13 (1.08-1.26) for all experiments. In PlankTOM6, even the simulation with no iron deposition from
14 dust (D0) produces Southern Ocean Chl concentrations that are too high during summer. This result
15 is consistent with the observation that although Fe is lower in the Southern Ocean than elsewhere,
16 concentrations average around $0.3 \text{ nmol Fe L}^{-1}$ (range of $0.15\text{--}0.6 \text{ nmol Fe L}^{-1}$) in the summer
17 (January and February, $n=79$) in the Subantarctic region (Tagliabue et al., 2012), which is near the
18 half-saturation for growth of most phytoplankton as well as those used in the model (Le Quéré et al.,
19 2005; Sarthou et al., 2005). Thus Fe concentrations may be limiting for phytoplankton growth, but
20 nevertheless the observed very low Chl concentration during summer months seem to reflect losses
21 due to other processes, such as grazing mortality rather than reduced growth rates from low Fe
22 supply.

23 As a means of validating the model results, we also tested the response of PlankTOM10 to Fe-
24 fertilisation to verify that the model reproduced the observed Chl blooms under Fe enrichment
25 conditions (Boyd and al., 2007). This was done by saturating the surface layer of the ocean with Fe
26 for one month (February). In this experiment, surface Chl south of 40°S increased by $2.1\pm 2.2 \text{ mg}$
27 Chl/m^3 (mean \pm 1SD) with a maximum concentration of 14.2 mg Chl/m^3 . This is similar to the
28 responses observed at sea during Fe-fertilisation experiments (Boyd and al., 2007). Thus Planktom10
29 predicts that net phytoplankton growth can escape the constraint imposed by zooplankton grazing and
30 bloom when superabundant Fe is provided as is the case in during the meso-scale Fe-fertilization
31 experiments. The response of the model to Fe enrichment provides further support of our hypothesis
32 that grazing is responsible for the low Chl concentration in the Southern Ocean during summer under
33 realistic Fe inputs.

34 **3.4.4 Role of combined effects**

35 Model simulations could be influenced by the model structure and parameters, the physical transport,
36 meteorological data, or the choice of dust deposition fields. We assessed the combined effects of
37 model choices by comparing our results with outputs from seven other models: a version of the
38 PISCES model (Aumont and Bopp, (2006), the CCSM-BECs model (Doney et al., (2009), and the
39 NEMURO model (Kishi et al., (2007), IPSL-CM5A-LR (Dufresne et al., 2013), GRDL-ESM2M
40 (Jones et al., 2011), HadGEM2-ES (Giorgetta et al., submitted), and CanESM2 (Arora et al., 2011).
41 All of these other models focus on the representation of phytoplankton groups and parameterise
42 grazing pathways in a simpler fashion than PlankTOM10. They produce a North/South Chl ratio in
43 the range from 0.60 to 1.36, lower than the value (1.72) obtained using PlankTOM10. Previous
44 studies have suggested that the overestimation of Chl may result from a generalised model bias
45 towards too shallow mixing depth in the Southern Ocean in summer, but S  f  rian et al. (2013) have
46 shown that while better representation of sub-grid scale processes and mixed layer depth improves
47 the simulation of Chl overall it does not lead to a more realistic North/South Chl ratio (Fig. 11).

1 Thus, the comparison between PlankTOM10 and other ocean biogeochemistry models supports our
2 contention that it is important to simulate grazing pathways explicitly.

4 4 Discussion

5 The development of PlankTOM10 has benefited from the existence of the very extensive range of
6 observations to develop realistic parameterisations of key processes, particularly PFT growth rates.
7 Although the simulated global biogeochemical fluxes are generally below or at the low end of the
8 range of observed values and several regional discrepancies exist between observed and modelled
9 biomass and fluxes, the model reproduces both the relative importance of different PFTs and the
10 geographic patterns in their abundance. Thus, while not perfect, the model is sufficient to explore the
11 role of ecosystem dynamics in determining ocean biogeochemistry.

12 Our analyses suggest that Southern Ocean Chl during summer is primarily controlled by zooplankton
13 grazing and the structure of the pelagic food web, rather than the low supply rate of iron. Trophic
14 cascading appears to account for the differences between the results from PlankTOM10 and
15 PlankTOM6 (Fig. 13; Zollner et al. 2009). For example, protozooplankton graze on phytoplankton
16 (and bacteria), which reduces their prey's biomass. However, mesozooplankton graze on
17 phytoplankton and protozooplankton, and macrozooplankton graze on phytoplankton and both
18 protozooplankton and mesozooplankton. Thus the grazing pressure of larger zooplankton on smaller
19 zooplankton can indirectly reduce the overall grazing pressure on phytoplankton. In PlankTOM10,
20 macrozooplankton concentration is higher in winter in the Northern Hemisphere Pacific sector where
21 the surface layer is more stratified and food is abundant, compared with the Southern Ocean Pacific
22 sector where the surface layer is more mixed and food is scarce. Thus when the spring bloom starts in
23 the North, the biomass and grazing pressure exerted by macrozooplankton are high enough to reduce
24 the biomass of smaller zooplankton consequently reducing the grazing pressure on Chl and leading to
25 an increase in Chl. However, in the South macrozooplankton biomass is too low to cause significant
26 losses of smaller zooplankton. Hence, the high proto- and mesozooplankton biomasses prevent a
27 phytoplankton bloom from developing in that region. Although PlankTOM6 simulates some degree
28 of trophic cascade with the presence of two zooplankton PFTs, our sensitivity tests presented in Fig.
29 11 show that the difference in growth rates between the two zooplankton PFTs is too small to impact
30 the phytoplankton significantly.

31 The higher concentration of macrozooplankton biomass in the North compared to the South is
32 consistent with the observations, where the mean biomasses of macrozooplankton was reported as
33 three times higher in the Northern Hemisphere compared to the Southern Hemisphere (Moriarty et
34 al., 2013). A similar contrast is found between the Atlantic and Pacific sectors of the Southern Ocean,
35 where the high macrozooplankton biomass observed in the Atlantic (Atkinson et al., 2004) would
36 reduce the abundance of smaller zooplankton resulting in higher Chl concentrations in the Atlantic
37 sector, as simulated in PlankTOM10 (Fig. 4). Such trophic cascades have been observed in diverse
38 ecosystems on land and in the ocean (Casini et al., 2009). Furthermore, many observational-based
39 studies have highlighted the important role of zooplankton grazing for controlling phytoplankton
40 biomass (Atkinson et al., 2001; Banse, 1996; Dubischar and Bathmann, 1997; Granli et al., 1993).
41 Although some processes are missing from the model (e.g. vertical migration of zooplankton, which
42 mostly contributes to downward export), the model suggests that the primary cascading effect of
43 grazing is sufficient to account for a large part of the North/South Chl differences.

44 Our results indicate that zooplankton grazing exerts an important control on Southern Ocean Chl.
45 This propagates through to influence phytoplankton biomass. Indeed, the North/South ratio of
46 phytoplankton biomass at surface is greater in PlankTOM10 (1.62) compared to PlankTOM6 (1.18),
47 very close to the modelled North/South ratio of Chl. The difference between the PlankTOM10 and

1 PlankTOM6 also persists through depth until about 300 m. Because of these marked differences, it is
2 clear that the representation of global biogeochemical cycles in ocean models is influenced by the
3 ecosystem structure. In both PlankTOM6 and PlankTOM10, the mesozooplankton and
4 macrozooplankton faecal pellets aggregate into the same large, fast-sinking particle pool, thus
5 limiting the effect of different size classes of zooplankton on carbon export. To distinguish the effects
6 of different food web structures on export production, a wider spectrum of particle size classes
7 sinking at different speeds are needed (e.g. Kriest; 2002). In addition, an improved vertical dynamics
8 of the mesopelagic zone, together with the enhanced representation of zooplankton dynamics in the
9 present study would allow further exploration of the interactions between iron fertilisation, grazing,
10 and mixed-layer dynamics, which have led to large differences among ocean iron fertilization
11 experiments (Smetacek and Naqvi 2008; Boyd et al. 2008).

12 There are a number of limitations to the current version of PlankTOM10, including simplified
13 overwintering strategies for zooplankton, the use of a coarse Fe model, and the lack of representation
14 of semi-refractory organic matter. In addition, the model does not include some ecosystem pathways,
15 such as viral lysis (Evans et al., 2009), and the zooplankton representation does not include salps,
16 pteropods, and auto- and mixotrophic dinoflagellates. The nano- and microzooplankton are also
17 combined into a single compartment. The realism of the simulations may also be affected by the
18 relatively coarse resolution of the physical ocean model. However these biases affect both
19 PlankTOM6 and PlankTOM10, and thus the experiments still provide information on the processes
20 that differ between the two models. Our work suggests that improved representation of the
21 zooplankton components could help further constrain the processes that regulate Chl distribution in
22 models. The effect of further ecosystem model developments will be explored in follow-up studies.

23 5 Conclusions

24 The development of global marine ecosystem models is hampered in particular because of our poor
25 understanding of several critical ecosystem processes and food-web interactions (Smetacek et al.
26 2004), and the paucity of global-scale observation of physiological rates and biomass for
27 parameterisation and validation (Le Quéré and Pesant, 2008; Barton et al. 2013). For example, the
28 wide range in observed growth rates for the same temperature is an indication of the challenges met
29 by marine ecosystem modellers, particularly in representing the within-PFT diversity, which is
30 unaccounted for in our model. Much more work is needed to understand the specific pathways by
31 which matter circulates within ecosystems, taking into account the regional distributions of
32 zooplankton groups and interactions with the environment including seasonal mixed layer dynamics.

33 The role of macrozooplankton highlighted here has implications for carbon export to depth because
34 faecal pellets of some macrozooplankton have very fast sinking rates (Fortier et al., 1994; Turner
35 2002). Hence, a more explicit representation of the pelagic food web in global models is needed to
36 capture the full range of interactions between marine ecosystems, marine biogeochemistry and
37 climate. The synthesis and analysis of observations and model results by the MAREDAT and
38 MAREMIP projects provide valuable insights into the processes that control marine ecosystems,
39 including the contributions that different PFTs make to ocean biomass (Buitenhuis et al., 2013a;
40 Hashioka et al., 2012; Sailley et al., 2013).

41 Our simulations examining the effects of grazing on phytoplankton biomass raise questions about the
42 biological and biogeochemical bases for the current projections of the feedbacks between climate
43 (and other environmental changes) and marine ecosystems. It also highlights potential complications
44 for the large-scale proposed use of purposeful Fe-fertilisation to enhance the deep ocean storage of
45 CO₂ (Ciais et al., 2013). Assessments of the impact of such geo-engineering techniques will be
46 unreliable, at least until the full ecosystem response including the grazing pathways (Landry et al.,
47 1997) and the relationship between ecosystem dynamics and deep water carbon export (Smetacek et

1 al., 2012) can be reproduced with models, which could be used to make quantitative predictions of
2 deliberate Fe-fertilisation over large areas.

3 Our results on the important role of grazing do not contradict the results on the importance of Fe-
4 fertilisation as highlighted in Fe enrichment experiments (Boyd and al., 2007), because additional Fe
5 would trigger further growth provided that Fe were initially below an optimal concentration (Blain et
6 al., 2007). However, our results suggest that low Fe concentrations by themselves are insufficient to
7 account for the very low Chl levels observed in the Southern Ocean HNLC region in summer, and
8 that differences in zooplankton trophic and community structure, and concomitant grazing dynamics
9 play an important role in controlling phytoplankton blooms and maintaining very low Chl levels in
10 that region. Although previous studies emphasised the role of phytoplankton community structure
11 (Arrigo et al., 1999) and mixed layer dynamics for nutrient supply and demand (Platt et al., 2003a;
12 Platt et al., 2003b) in ocean biogeochemical cycles, our analysis makes it clear that it is important to
13 consider the whole pelagic ecosystem, including the zooplankton, when studying and predicting
14 ecosystem responses to Fe (or any essential nutrient) fertilisation. This complex interplay has
15 received less attention than either the drivers of primary production or the representation of Fe
16 cycling in global biogeochemical modelling. Our results suggest that representing zooplankton
17 interactions more explicitly in models would improve the representation of biogeochemistry –
18 climate interactions, and could bring new insights to understand changing global biogeochemical
19 cycles.

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33 Performance Computing Cluster of the University of East Anglia.

1 Table 1. Growth rates of PFTs at 0 and 20°C (μ_0 and μ_{20}), and rate increase for a 10°C increase in
 2 temperature (Q_{10}). The uncertainty in μ_0 and Q_{10} represents ± 1 standard deviation from an optimal
 3 parameter value in the parameter space. Full references for the phytoplankton growth rate data are
 4 provided in the Supplementary Information. The zooplankton growth rate data are from published
 5 data synthesis cited here.

6

PFT	μ_0	Q_{10}	μ_{20}	number of obs.	p values	Size range (μm)	Main references
<i>Autotrophs</i>							
N ₂ -fixers	0.05 ± 0.05	1.83 ± 0.71	0.1 5	34	0.76	0.5-2.0	LaRoche and Breitbarth (2005) ^g
Picophytoplankton	0.26 ± 0.06	1.81 ± 0.18	0.8 9	150	<0.01	0.7-2.0	Agawin et al. (1998); Johnson et al (2006); Moore et al. (1995)
Coccolithophores	0.70 ± 0.17	1.14 ± 0.17	0.9 0	322	0.06	5-10	Buitenhuis et al. (2008); S. Larsen (t paper)
Mixed-phytoplankton	0.35 ± 0.05	1.57 ± 0.12	0.8 7	95	<0.01	2-200	Bissinger et al. (2008) ^g
Diatoms	0.44 ± 0.02	1.93 ± 0.07	1.6 3	439	<0.01	20-200	Sarthou et al. (2005) ^g
<i>Phaeocystis</i>	0.68 ± 0.07	1.66 ± 0.16	1.8 7	67	0.23	120-360	Schoemann et al. (2005) ^g
<i>Heterotrophs</i>							
Bacteria	0.66 ± 0.04	1.45 ± 0.06	1.2 2	1429	<0.01	0.3-1.0	Rivkin and Legendre (2001) ^g ; Cho and Giovannoni (2004)
Protozooplankton	0.46 ± 0.07	1.48 ± 0.13	1.0 3	1057	0.01	5-200	Buitenhuis et al. (2010) ^g
Mesozooplankton	0.31 ± 0.02	1.27 ± 0.05	0.4 9	2745	<0.01	200-2000	Hirst and Bunker (2003) ^g
Macrozooplankton	0.03 ± 0.01	3.01 ± 0.52	0.1 9	253	<0.01	>2000	Hirst et al. (2003) ^g

7 ^gThese references include syntheses of data from other papers

8

9

1 Table 2. Model parameters constraining the resource limitations of growth rates. See model equations
 2 in Supplementary Information for definitions of parameters.

PFT

Autotrophs

	Light		Fe ^{opt} μmolFe molC ⁻¹	Nutrients half saturation ^b		
	α ^a	θ _{max} gChl gC ⁻¹		k _{Fe} nmol L ⁻¹	k _P μmol L ⁻¹	k _N μmol L ⁻¹
N ₂ -fixers	1	0.025	8.6	40	0.2	13
Picophytoplankton	1	0.033	8.6	10	0.13	2
Coccolithophores	1	0.033	8.6	25	0.13	2
Mixed-phytoplankton	1	0.033	8.6	25	0.1	2
Diatoms	1	0.058	8.6	40	0.06	2
<i>Phaeocystis</i>	1	0.042	8.6	25	0.8	3

Heterotrophs

	Food half saturation K _{Food} μmolC L ⁻¹
Bacteria	10
Protozooplankton	10
Mesozooplankton	10
Macrozooplankton	9

3 ^aunits of molC gChl⁻¹ m² (mol photons)⁻¹

4 ^bThe reported values are half saturation for uptake for Fe, and half saturation for growth for P and N.

5

1 Table 3. Relative preference of zooplankton for food. The preferences are weighted with the biomass
 2 to obtain the model parameter value as in Buitenhuis et al. (2010).

3

Plankton Functional Type	Protozooplankton	Mesozooplankton	Macrozooplankton
<i>Autotrophs</i>			
N ₂ -fixers	2	0.1	0.1
Picophytoplankton	2	0.75	0.5
Coccolithophores	2	0.75	1
Mixed-phytoplankton	2	0.75	1
Diatoms	1	1	1
<i>Phaeocystis</i>	2	0.75	1
<i>Heterotrophs</i>			
Bacteria	4	0.1	0.1
Protozooplankton	0	2	1
Mesozooplankton	0	0	1
Macrozooplankton	0	0	0
<i>Particulate matter</i>			
Small organic particles	0.1	0.1	0.1
Large organic particles	0.1	0.1	0.1

4

1 Table 4. Global mean values for rates and biomass from observations (data) and PlankTOM10
 2 (model) averaged over 1998-2009. The reported confidence level are from the author's assessment of
 3 confidence with high (H): most likely within $\pm 25\%$ of reported value; medium (M): most likely
 4 within $\pm 50\%$ of reported value; low (L): could be more than $\pm 50\%$ of reported value. For the
 5 biomass of phytoplankton and zooplankton, the percentage of the total biomass is also indicated in
 6 parentheses (excluding mixed-phytoplankton for which no observations are available).
 7

	PlankTOM10	PlankTOM6	Data	Confidence	Reference for the data
Rates					
Primary production (PgC y ⁻¹)	42.6	35.4	51-65	H	Buitenhuis et al. (2013b)
Export production at 100 m (PgC y ⁻¹)	7.6	7.7	9-10	M	Schlitzer (2004); Lee (2001)
CaCO ₃ export at 100 m (PgC y ⁻¹)	0.40	0.80	0.6-1.1	M	Lee (2001); Sarmiento et al. (2002)
SiO ₂ export at 100 m (Pg Si)	2.9	4.5	3.4	H	Tréguer et al. (1995)
N ₂ fixation (TgN y ⁻¹)	165	—	60-200	H	Gruber (2008)
Phytoplankton biomass 0-200 m (PgC)^a					
N ₂ -fixers	0.062 (9.8%)	—	0.008-0.12 (2-8%)	M	Luo et al. (2012)
Picophytoplankton	0.21 (34%)	0.23 (50%)	0.28-0.52 (35-68%)	M	Buitenhuis et al. (2012b)
Coccolithophores	0.077 (12%)	—	0.001-0.032 (0.2-2%)	M	O'Brien et al. (2013)
Mixed-phytoplankton	0.079 (12%)	0.023 (5.0%)			
<i>Phaeocystis</i>	0.080 (13%)	—	0.11-0.69 (27-46%)	M	Vogt et al. (2012)
Diatoms	0.12 (19%)	0.20 (45%)	0.013-0.75 (3-50%)	M	Leblanc et al. (2012)
Heterotrophs biomass 0-200 m (PgC)^a					
Bacteria	0.031	0.030	0.25-0.26	H	Buitenhuis et al. (2012a)
Protozooplankton	0.067 (12%)	0.12 (44%)	0.10-0.37 (27-31%)	M	Buitenhuis et al. (2010)
Mesozooplankton	0.18 (34%)	0.15 (56%)	0.21-0.34 (25-66%)	M	Moriarty and O'Brien (2013)
Macrozooplankton	0.28 (53%)	—	0.010-0.64 (3-47%)	L	Moriarty et al. (2013)

8 ^aThe biomass ranges have been computed using the method described in Buitenhuis *et al.* (2013b).
 9

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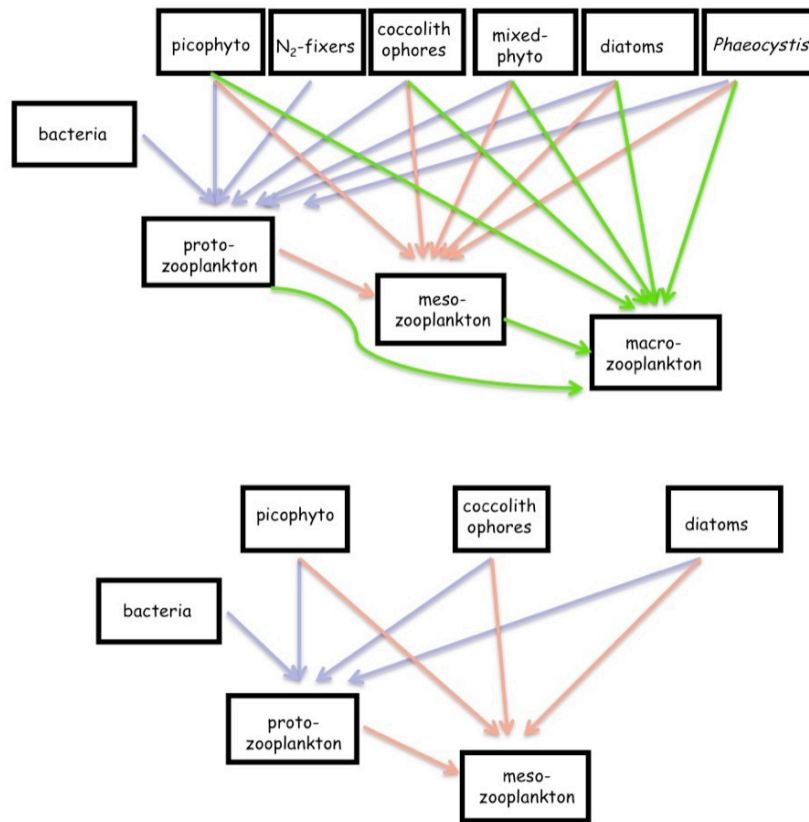
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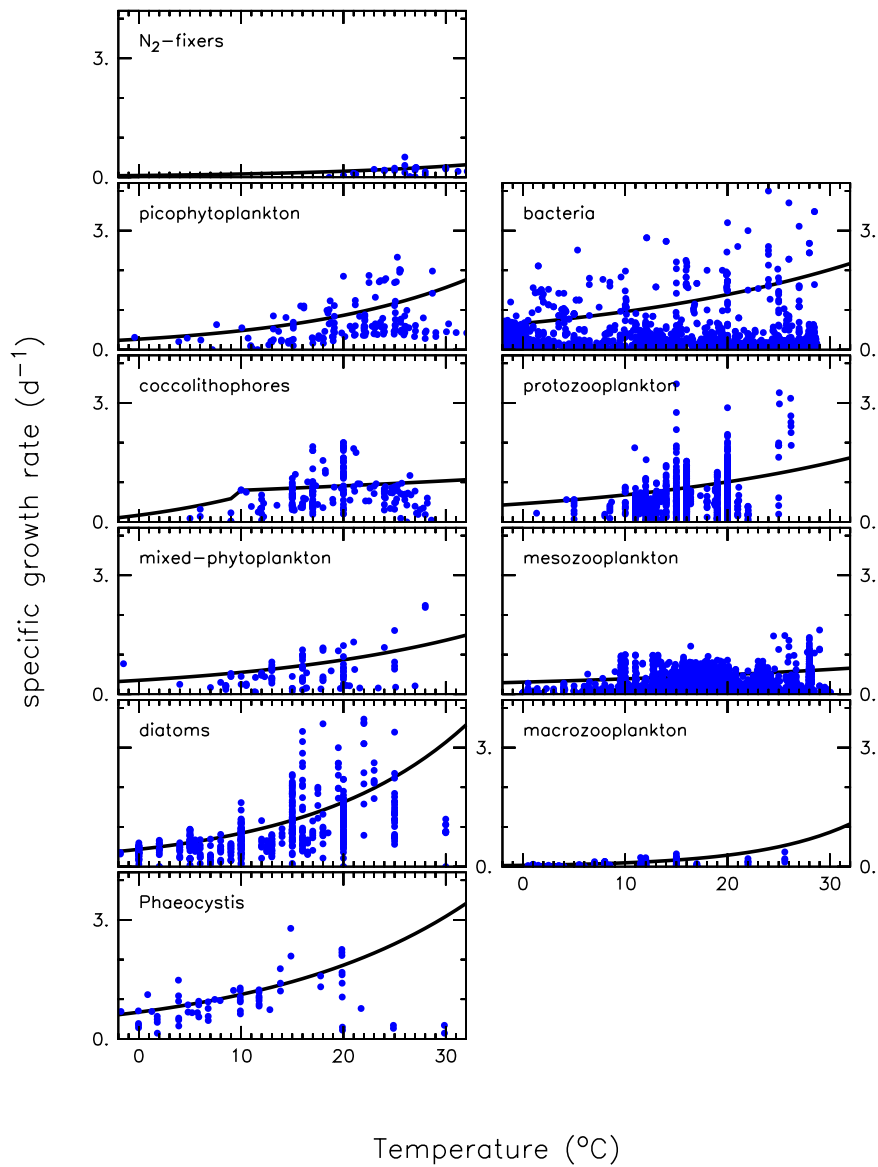
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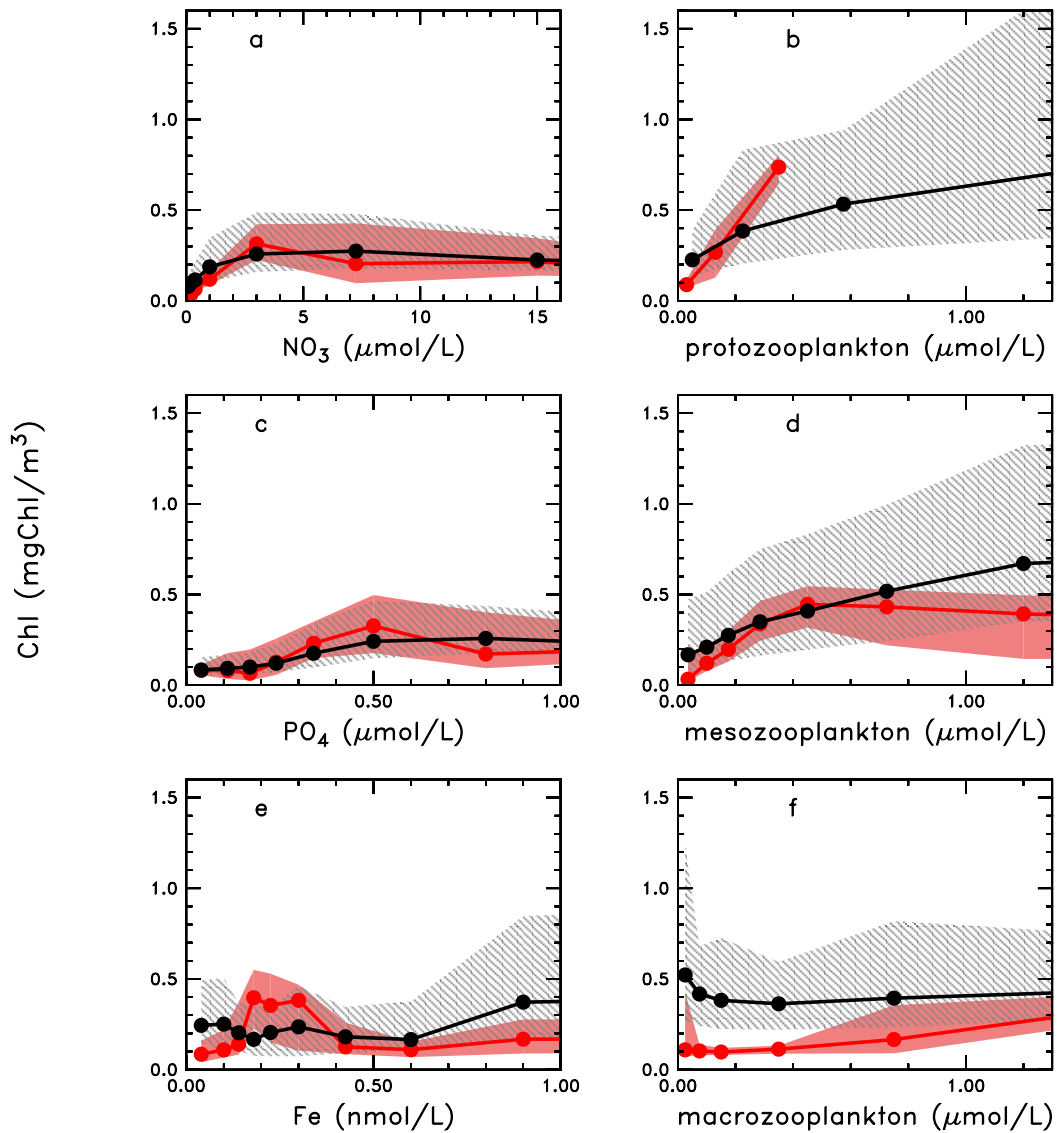
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3 **Figure 1.** Schematic representation of the PlankTOM10 (top) and PlankTOM6 (bottom) marine
4 ecosystem models. The arrows show grazing fluxes by protozooplankton (purple), mesozooplankton
5 (red), and macrozooplankton (green). Only fluxes with weighing factors above 0.1 are shown (Table
6 3).

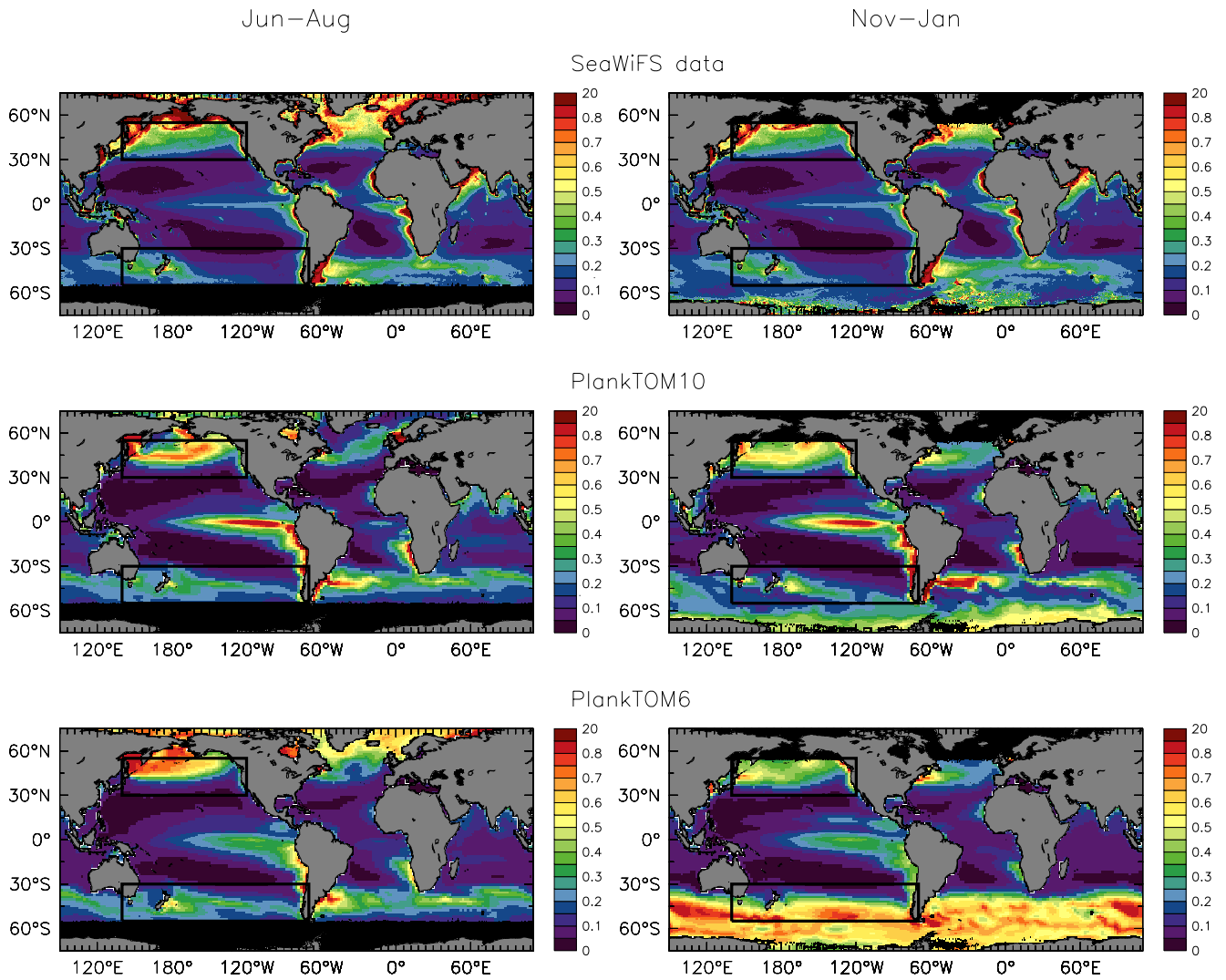
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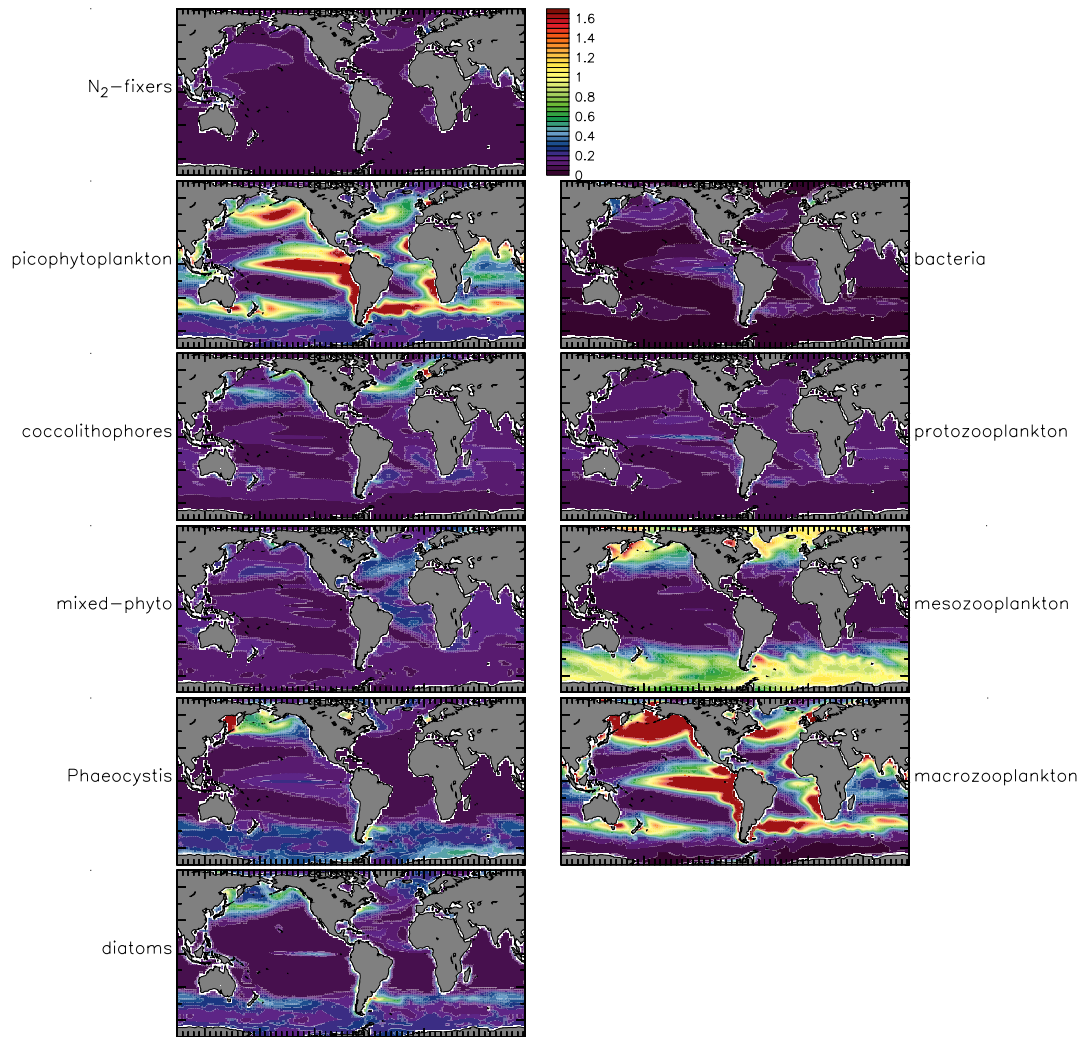
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 2 **Figure 2.** Maximum growth rates for 10 Plankton Functional Types as a function of temperature for
 3 the phytoplankton PFTs (left) and for the heterotrophic PFTs (right). The PFTs are presented from
 4 the smallest (top) to the largest (bottom) in size. The fit to the data used in the model is shown in
 5 black, using the parameter values from Table 1. See Table 1 for references.
 6



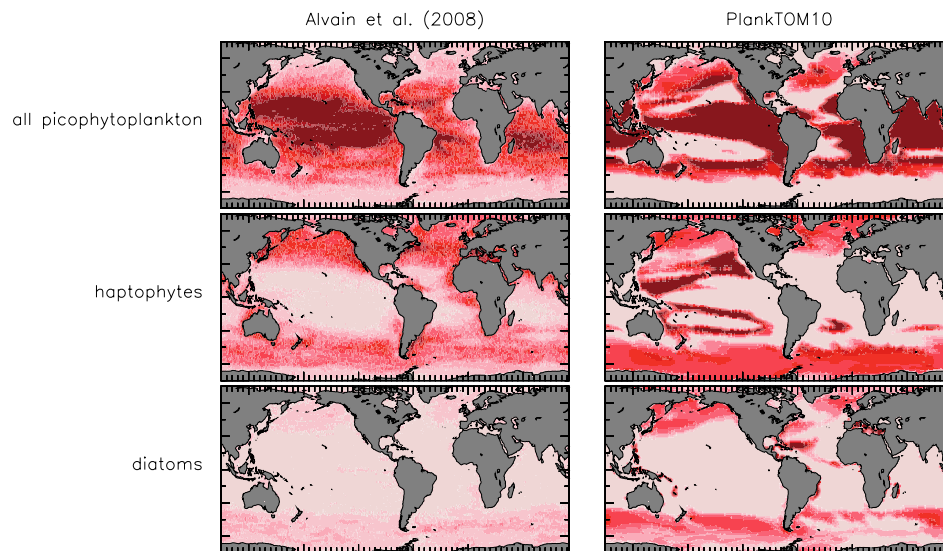
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2 **Figure 3.** Covariation between Chl concentration and (left) potentially limiting nutrients and (right)
3 biomass of zooplankton groups for the World Ocean. Chlorophyll data from SeaWiFS satellite are
4 the same in each panel, and are averaged over 1998-2009. The NO_3 and PO_4 data are from the World
5 Ocean Atlas 2009, updated from (Garcia, 2006b). Fe data are from (Tagliabue *et al.*, 2012). The
6 protozooplankton biomass data are updated from Buitenhuis *et al.* (2010), the mesozooplankton
7 biomass data from Buitenhuis *et al.* (2006), and the macrozooplankton biomass data include all krill
8 data from Atkinson *et al.* (2004) and other crustacean data from (Moriarty *et al.*, 2013). All data are
9 monthly averages except for the mesozooplankton, which are seasonal. All data are for the surface,
10 generally corresponding to the mixed layer, except for observed Chl, which is seen by satellite over
11 one optical depth, and observed mesozooplankton and macrozooplankton, which are from depth-
12 integrated tows and may underestimate surface concentrations (by a factor 1.5-2; see text). The black
13 lines are medians, and grey shadings the 25-75% interquartile range for Chl concentration. The
14 median from the PlankTOM10 model is shown in red.
15



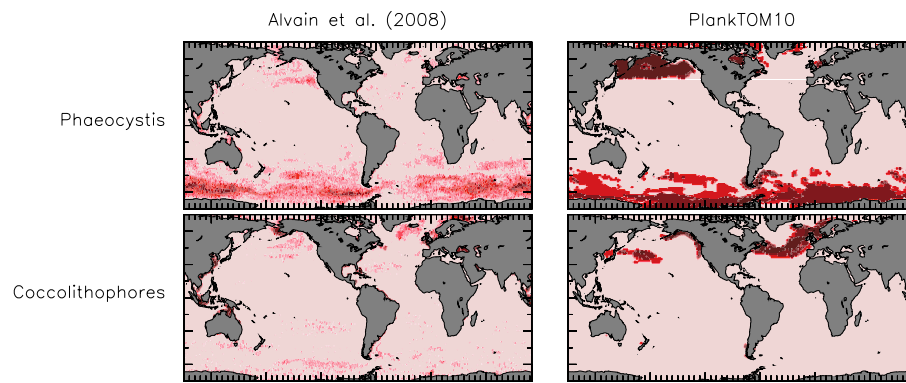
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 2 **Figure 4.** Surface Chl (mg m^{-3}) for (left) Southern Ocean winter (Jun-Aug) and (right) Southern
 3 Ocean summer (Nov-Jan). Data are from (top) SeaWiFS satellite, (middle) PlankTOM10, and
 4 (bottom) PlankTOM6. All datasets are averages for 1998-2009. Model results are shown for the top
 5 10-m deep surface box. The boxes highlight the regions used in Fig. 11
 6



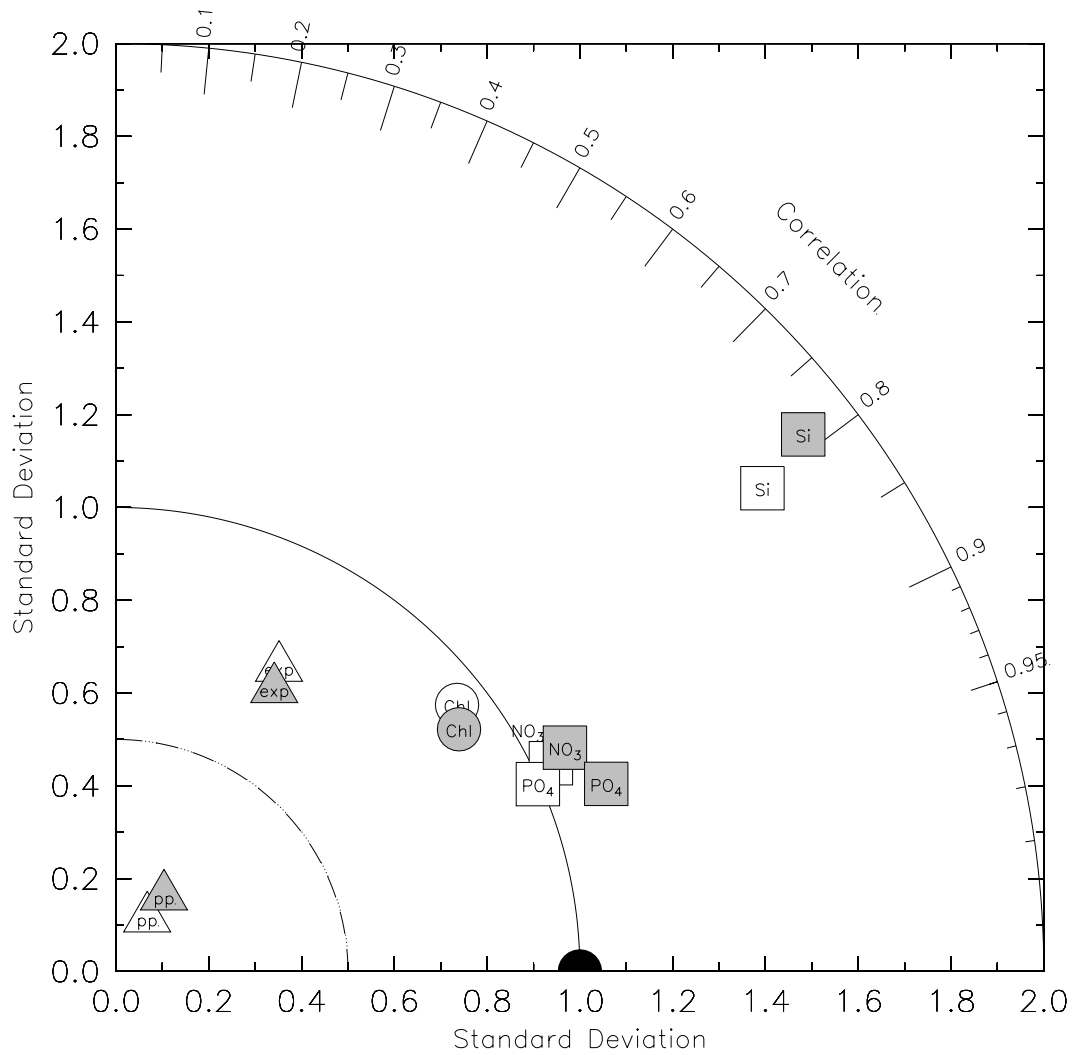
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 2 **Figure 5.** Annual mean surface carbon biomasses for individual Plankton Functional Types as
 3 simulated by the PlankTOM10 model ($\mu\text{mol C L}^{-1}$). Model results are averaged for 1998-2009 and
 4 shown for the top 10-m deep surface box.
 5



1
2 **Figure 6.** Dominance of picophytoplankton (top), haptophytes (middle) and diatoms (bottom) in the
3 ocean surface (fraction of time). Left panels show the frequency of the dominance of each PFT
4 detected from satellite data by Alvain et al. (2005) for each pixel during 1998-2006. Right panels
5 show model results, as the surface Chl for each PFT divided by the total Chl. For the model results,
6 picophytoplankton include both the picophytoplankton and N₂-fixers groups; haptophytes include
7 coccolithophores, DMSP-producers and mixed-phytoplankton. The data provides information on the
8 spatial patterns, but not on the absolute amplitude of the dominance. To best highlight the spatial
9 patterns in the model, a PFT is assumed to be dominant if it accounts for at least 45% of the biomass
10 for picophytoplankton and haptophytes, and 30% of the biomass for diatoms. The dark red represents
11 area with highest dominance of a PFT, while in the lightest red the PFT is absent.
12

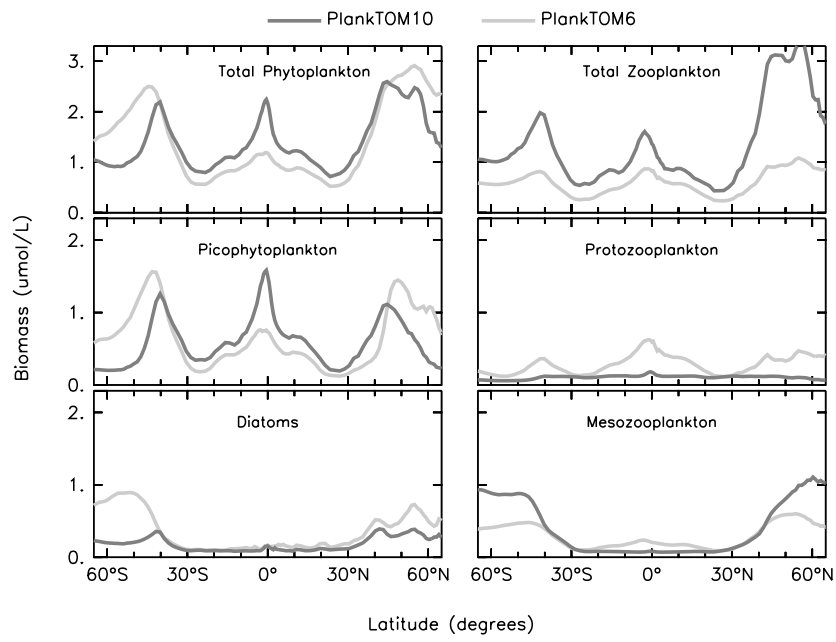


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2 **Figure 7.** Frequency of blooms of *Phaeocystis* (top) and coccolithophores (bottom) in the surface
3 ocean. *Phaeocystis* data are from Alvain et al. (2005); coccolithophore blooms are updated from
4 Brown and Yoder (1994). A bloom is defined in the model when the PFT accounts for at least 30%
5 of the biomass and when Chl exceeds 0.3 mgChl/m^3 . The dark red represents area with highest
6 dominance of a PFT, while in the lightest red the PFT is absent.
7

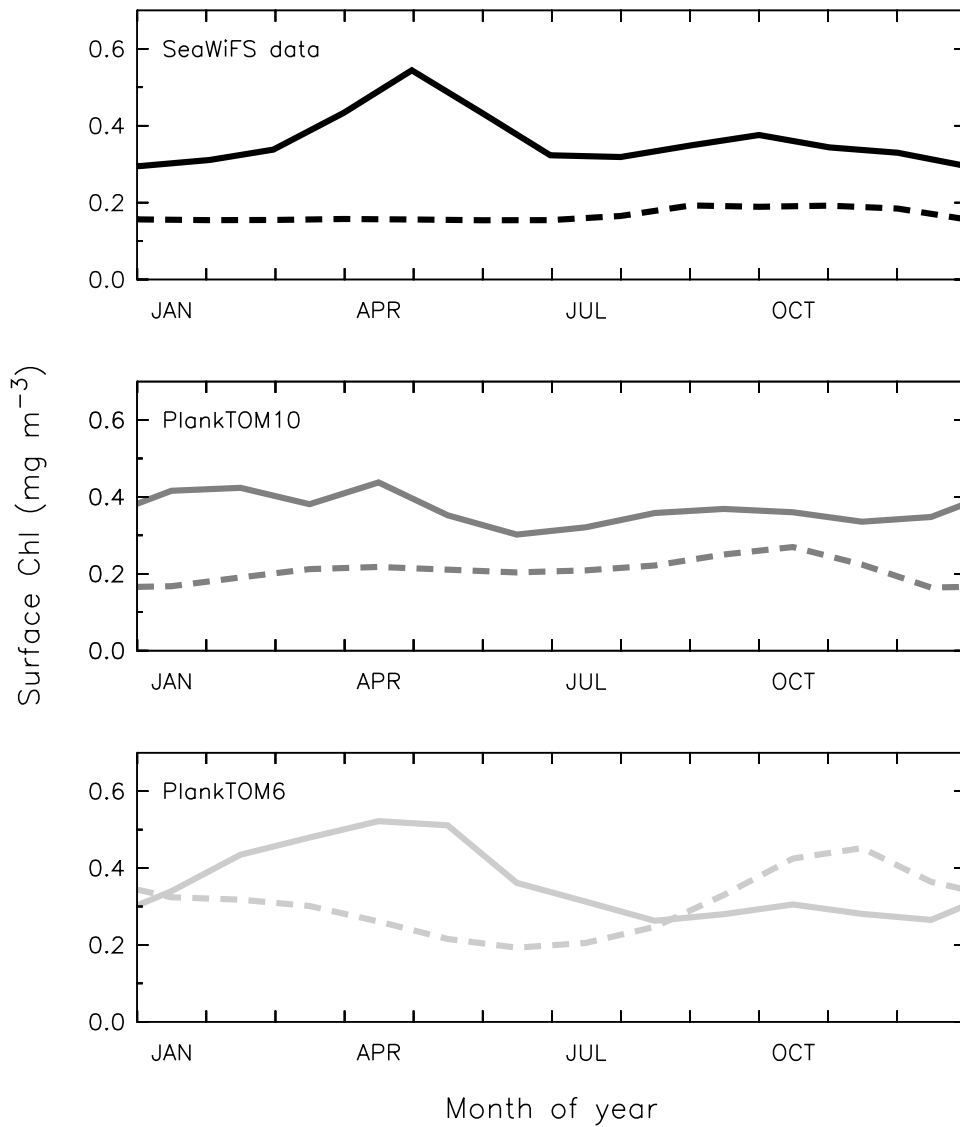


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 2 **Figure 8.** Taylor diagram comparing the distributions of surface concentration in annual and monthly
 3 mean Chl (Chl and Chl_s), NO₃, PO₃, Si, primary production (pp) and export production (exp) for
 4 PlankTOM10 (in grey) and PlankTOM6 (in white) with observations. Chl, biomass and nutrient
 5 observations are as in Figure 3. Export production is from (Schlitzer, 2004) and represents annual
 6 mean flux at 100 m. Primary production is from Buitenhuis et al. (2013) and includes monthly mean
 7 values for the surface 300 m. The black dot shows the location where the model results should be if it
 8 was perfect and there were no errors in the observations. The distance from the black dot quantifies
 9 the performance of the model (Taylor, 2001).

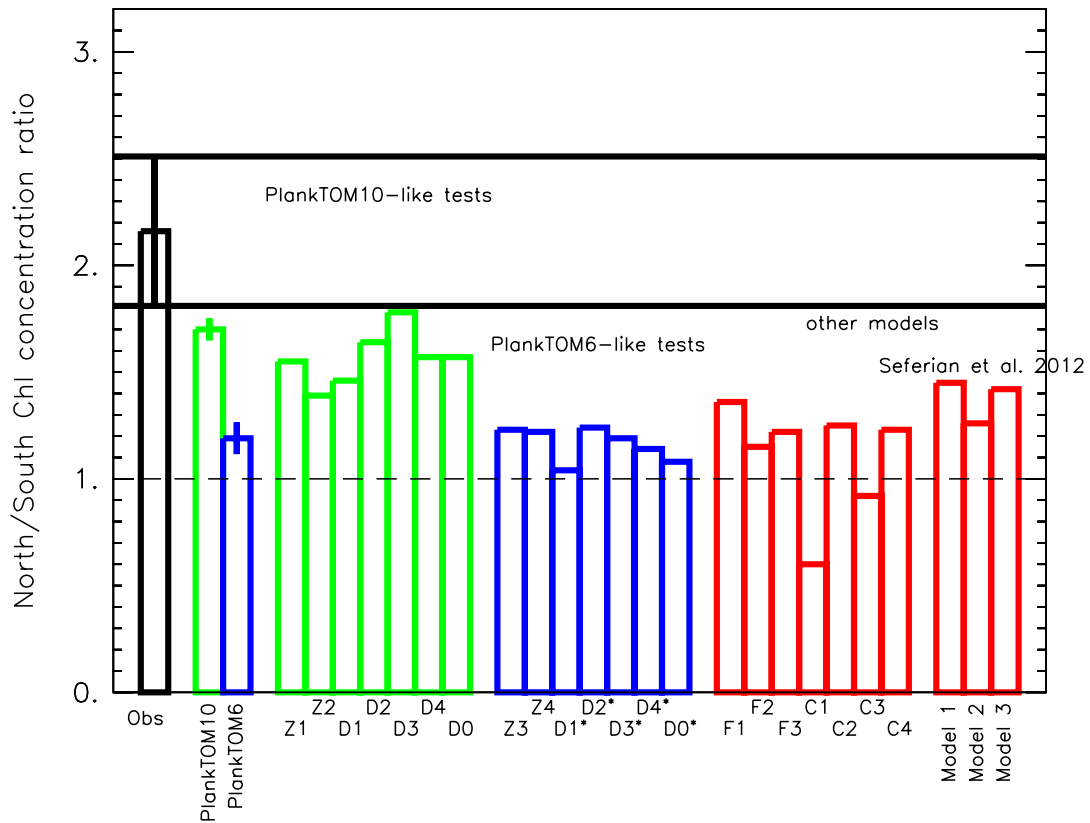
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1
 2 **Figure 9.** Zonal mean distribution of phytoplankton (left) and zooplankton (right) PFTs for the
 3 PlankTOM10 (dark grey) and PlankTOM6 (light grey) models ($\mu\text{mol C L}^{-1}$).
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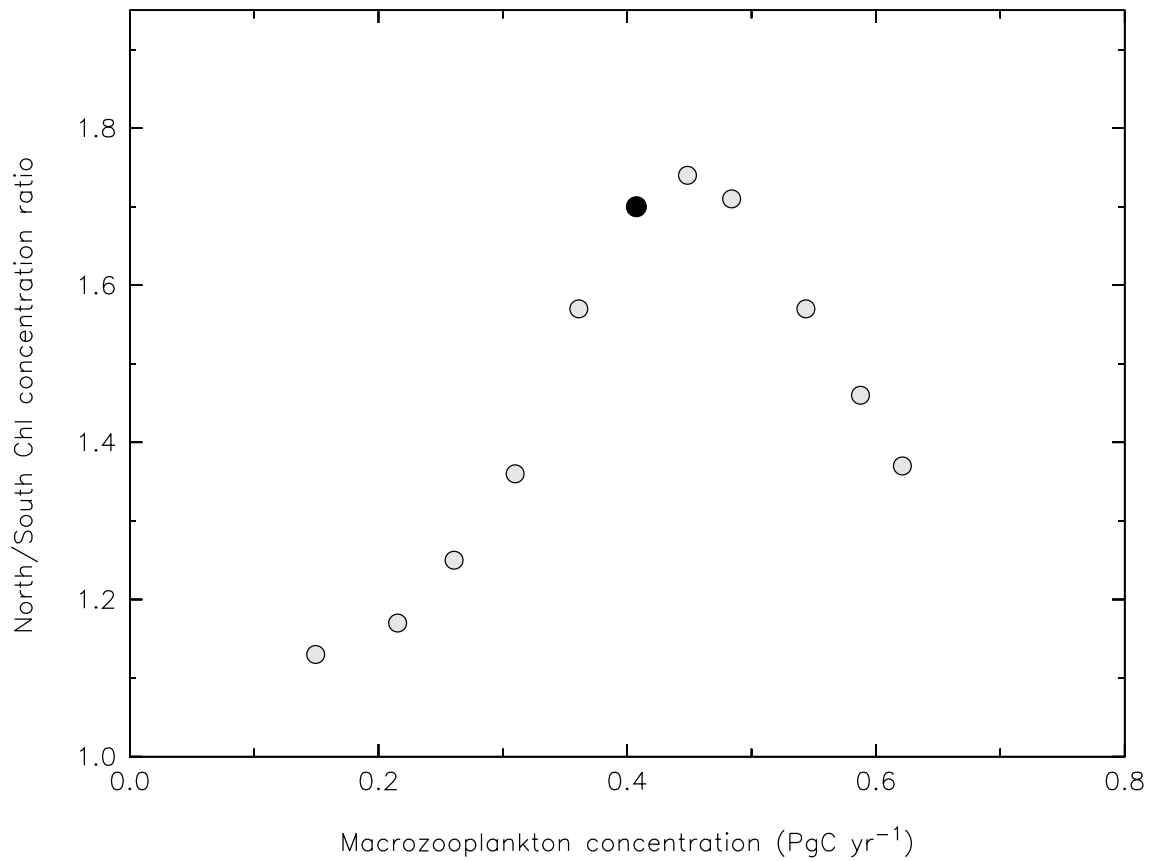


1
2 **Figure 10.** Monthly variations of surface Chl concentration in the North (full solid lines) and South
3 (dashed lines; mgChl m⁻³) Pacific Ocean. Data are from (top) SeaWiFS satellite, (middle)
4 PlankTOM10, and (bottom) PlankTOM6. All datasets are averages for 1998-2009. Model results are
5 shown for the top 10-m deep surface box. All data are averaged between 30 and 55 degrees latitude
6 in both hemispheres; 140°E-240°E in the North and 140°E-290°E in the South as highlighted in Fig.
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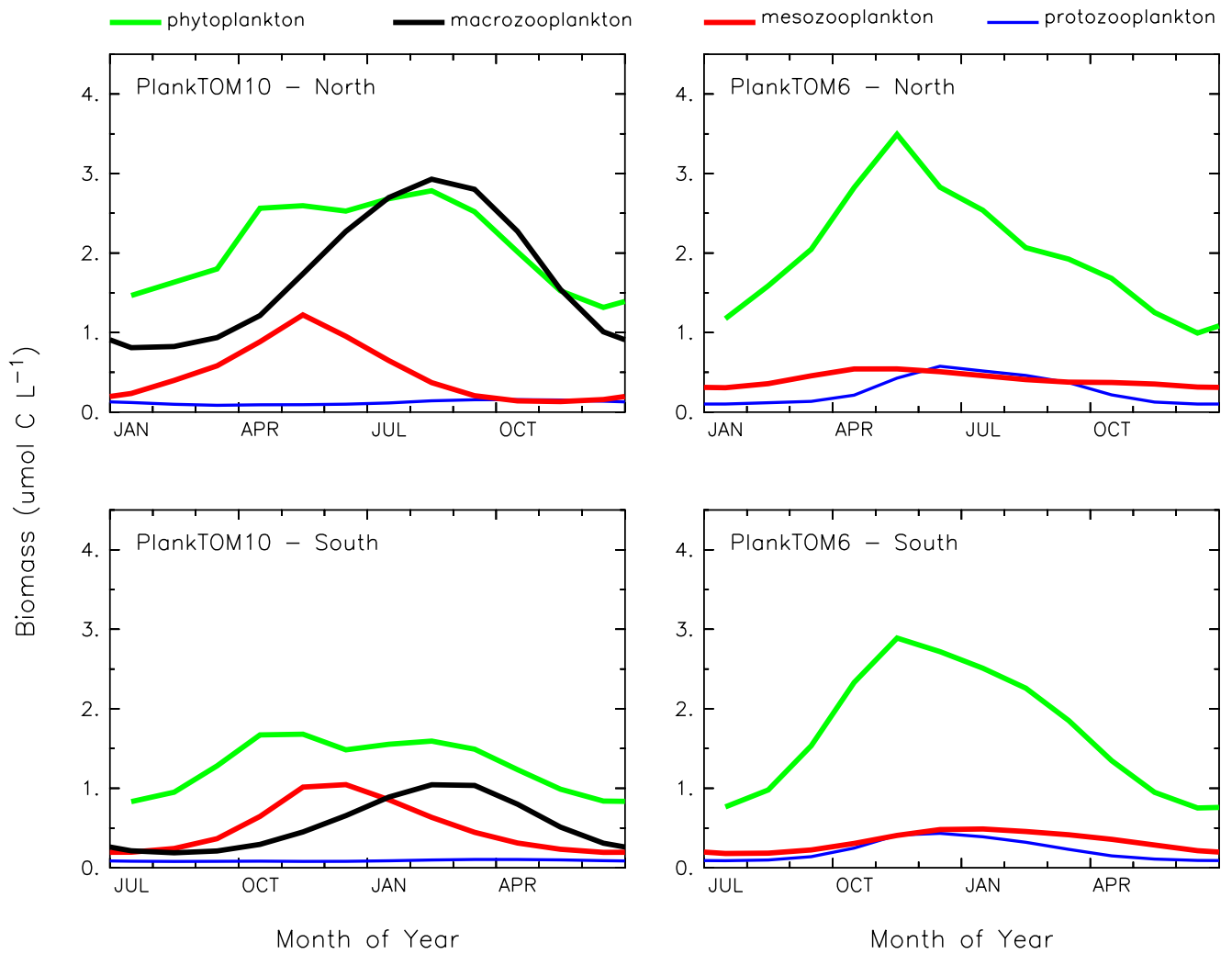


1 Model simulations

2 **Figure 11.** North/South ratio of surface Chl concentration in the Pacific Ocean. Observations are
3 from SeaWiFS. Model results in green correspond to model runs with slow-growing zooplankton:
4 PlankTOM10 (includes macrozooplankton), (Z1) PlankTOM6 plus macrozooplankton, (Z2)
5 PlankTOM6 with mesozooplankton parameterised like macrozooplankton, (D0-D4) PlankTOM10
6 with no dust deposition or with dust fields from (Mahowald *et al.*, 2009), (Tegen *et al.*, 2004),
7 (Ginoux *et al.*, 2001) and (Mahowald *et al.*, 2003), respectively. Model results in blue correspond to
8 model runs without slow growing zooplankton: PlankTOM6, (Z3) PlankTOM10 minus
9 macrozooplankton, (Z4) PlankTOM10 with macrozooplankton parameterised like mesozooplankton,
10 and (D0*-D4*) as above with PlankTOM6. Results from (F1-F3) are model simulations available
11 through the MARine Ecosystem Model Intercomparison Project and (C1-C4) the Climate Model
12 Intercomparison Project 5 (Arora *et al.*, 2011, Dufresne *et al.*, in revision, Giorgetta *et al.*, submitted,
13 Jones *et al.*, 2011). Results from (S  ferian *et al.*, 2012) mainly differ through their representation of
14 sub-grid scale processes, with improvements in the representation of summer mixed layer depth from
15 Model 1 to Model 3. All data are averaged between 30 and 55 degrees latitude in both hemispheres;
16 140  E-240  E in the North and 140  E-290  E in the South as highlighted in Fig. 4
17



1
 2 **Figure 12.** North/South ratio of surface Chl concentration in the Pacific Ocean as in Fig. 9 versus the
 3 surface biomass of macrozooplankton (PgC yr⁻¹). The standard PlankTOM10 results are shown by
 4 the filled circle. Results from ten sensitivity tests are shown by the empty circles, where the
 5 maximum growth rate of macrozooplankton is varied within $\pm 2\sigma$ within the range of the data (Fig. 2).
 6



1
 2 **Figure 13.** Mean surface concentrations of the biomass of phytoplankton (green), macrozooplankton
 3 (black), mesozooplankton (red), and protozooplankton (blue). Results are show for (left) the
 4 PlankTOM10 model and (right) the PlankTOM6 model, and for (top) the North, and (bottom) the
 5 South. All data are averaged for 1998-2009, and between 30 and 55 degrees latitude in both
 6 hemispheres; 140°E-240°E in the North and 140°E-290°E in the South as highlighted in Fig. 4

7