# iMarNet: An ocean biogeochemistry model

# 2 inter-comparison project within a common

# 3 physical ocean modelling framework.

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### Abstract

Ocean biogeochemistry (OBGC) models span a wide range of complexities from highly 24 simplified, nutrient-restoring schemes, through nutrient-phytoplankton-zooplankton-25 detritus (NPZD) models that crudely represent the marine biota, through to models that 26 represent a broader trophic structure by grouping organisms as plankton functional types 27 (PFT) based on their biogeochemical role (Dynamic Green Ocean Models) and ecosystem 28 models which group organisms by ecological function and trait. OBGC models are now 29 integral components of Earth System Models (ESMs), but they compete for computing 30 resources with higher resolution dynamical setups and with other components such as 31 atmospheric chemistry and terrestrial vegetation schemes. As such, the choice of OBGC in 32 ESMs needs to balance model complexity and realism alongside relative computing cost. 33 34 Here, we present an inter-comparison of six OBGC models that were candidates for implementation within the next UK Earth System Model (UKESM1). The models cover a 35 36 large range of biological complexity (from 7 to 57 tracers) but all include representations of at least the nitrogen, carbon, alkalinity and oxygen cycles. Each OBGC model was coupled to 37 38 the Nucleus for the European Modelling of the Ocean (NEMO) ocean general circulation model (GCM), and results from physically identical hindcast simulations were compared. 39 40 Model skill was evaluated for biogeochemical metrics of global-scale bulk properties using conventional statistical techniques. The computing cost of each model was also measured in 41 42 standardised tests run at two resource levels. No model is shown to consistently outperform all other models across all metrics. Nonetheless, the simpler models are broadly closer to 43 44 observations across a number of fields, and thus offer a high-efficiency option for ESMs that prioritise high resolution climate dynamics. However, simpler models provide limited insight 45 46 into more complex marine biogeochemical processes and ecosystem pathways, and a parallel approach of low resolution climate dynamics and high complexity biogeochemistry 47 48 is desirable in order to provide additional insights into biogeochemistry – climate interactions. 49

#### 1 Introduction

- Ocean biogeochemistry is a key part of the Earth System: it regulates the cycles of major
- 52 biogeochemical elements and controls the associated feedback processes between the land,
- ocean and atmosphere. As a result, changes in ocean biogeochemistry can have important
- 54 implications for climate (Reid et al., 2009). Marine ecosystems are indirectly affected by
- anthropogenic environmental change (Jackson et al., 2001), particularly through climate-
- induced changes in physical properties and CO<sub>2</sub>-induced ocean acidification. Understanding
- 57 and quantifying the response of ocean biogeochemistry to global changes and their
- 58 feedbacks with the Earth System is essential to improve our capacity to maintain ecosystem
- 59 services this century and beyond.
- 60 With the recent publication of the Intergovernmental Panel on Climate Change (IPCC) 5<sup>th</sup>
- Assessment Report (AR5), global efforts are already underway to develop the next
- 62 generation of Earth System Models (ESMs) to support climate policy development and any
- 63 further IPCC Assessment Report. OBGC coupled to ESMs can help address a series of
- overarching scientific questions: How will the ocean contribute to atmospheric trace gas
- composition (e.g. CO<sub>2</sub>, CH<sub>4</sub>, N<sub>2</sub>O, DMS) in a changing climate? Are there tipping points in
- 66 marine biogeochemistry (e.g. oceanic anoxic events, methane hydrate release) that could be
- 67 triggered by a changing climate? Are there interactions between ESM processes and
- society's management of resources (e.g. fisheries, land use, agriculture) in the marine
- 69 environment? Furthermore, as ESMs are increasingly being evaluated based on their
- 70 capacity to understand past variability (Braconnot et al., 2012), further questions might
- 71 include: What controlled variations in atmospheric trace gas concentrations and isotopic
- 72 composition over the geological past?
- 73 For an anticipated 6<sup>th</sup> IPCC assessment report it is generally considered that these global-
- scale questions, with direct implications for climate policies, will again be the main focus of
- 75 ocean biogeochemical models within ESMs. In addition, the ESM model archive is
- increasingly being used by activities within the Inter-Sectoral Impact Model Intercomparison
- 77 Project (http://www.pik-potsdam.de/research/climate-impacts-and-
- vulnerabilities/research/rd2-cross-cutting-activities/isi-mip/scientific-publications) to
- 79 address socioeconomically-directed questions such as: How will climate change affect ocean
- primary production (e.g. Bopp et al., 2013), fisheries (Barange et al., 2014; Cheung et al.,
- 2012), and harmful algal and jellyfish blooms (e.g. Codon et al., 2013, Gilbert et al., 2014)?
- What is the potential for geoengineering schemes such as ocean fertilisation (Buesseler &
- 83 Boyd, 2003) and alkalinity addition (Kheshgi, 1995; Harvey, 2008) to affect the climate
- system, and how do they affect the rest of the Earth System?

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Within the UK, the Integrated Global Biogeochemical Modelling Network (iMarNet) project aims to advance the development of ocean biogeochemical models through collaboration between existing modelling groups at Plymouth Marine Laboratory (PML), National Oceanography Centre (NOC), University of East Anglia (UEA) and the Met Office-Hadley Centre (UKMO). As part of iMarNet we conducted an intercomparison of 6 current UK models, to help inform the selection of a baseline OBGC model for the next UK Earth System Model (UKESM1). This intercomparison focused on model skill at reproducing global-scale bulk properties - such as nutrient and carbon distributions - that broadly characterise the activity of marine biota (and, thus, the carbon cycle) in the ocean. To limit the role of errors originating with modelled physics, all of the examined model simulations were performed within the same physical ocean GCM, under the same external forcing and following the same experimental protocol. As all of the models examined have been previously published, our analysis does not include an assessment of their underlying biological fidelity (i.e. the extent to which structures, parameterisations and parameter sets of candidate models are apriori realistic). However, while primarily focused on model skill, the intercomparison also considers the computational cost of the models in relation to the realism that they offer.

Previous authors have performed biogeochemical model intercomparisons with parallels to this study (e.g. Friedrichs et al., 2007; Kriest et al., 2010; Steinacher et al., 2010; Popova et al., 2012). These have differed from this study, and each other, in a number of ways. For instance, this study is 3D rather than 1D (cf. Friedrichs et al., 2007); global rather than regional (cf. Popova et al., 2012); uses identical rather than diverse physics (cf. Steinacher et al., 2010); and spans a more functionally diverse range of biogeochemical models (cf. Kriest et al., 2010). The latter two factors, in particular, distinguish this study, permitting us to both formally separate the impact of physics from that of biogeochemical dynamics, and to do so across a broad range of model complexity from NPZD through to state-of-the-art PFT models with considerable ecological sophistication. This study is still constrained by the use of a single ocean circulation, and by a bespoke gradation of model complexity (PlankTOM6 and PlankTOM10 partially inform this). Nonetheless, this study represents an intercomparison along separate lines to those previously conducted.

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# 2 Method

## 2.1 Experimental Design

All participating models made use of a common version (v3.2) of the NEMO physical ocean general circulation model (Madec, 2008) coupled to the Los Alamos sea-ice model (CICE) (Hunke and Lipscomb, 2008). This physical framework is configured at approximately 1×1 degree horizontal resolution (ORCA100; 292×362 grid points), with a focusing of resolution

around the equator to improve the representation of equatorial upwelling. Vertical space is divided into 75 fixed levels, which increase in thickness with depth, from approximately 1m at the surface to more than 200m at 6000m. Partial level thicknesses are used in the specification of seafloor topography to improve the representation of deep water circulation. Vertical mixing is parameterized using the turbulent kinetic energy scheme of Gaspar et al., (1990), with modifications made by Madec (2008). To ensure that the simulations were performed by the different modelling groups using an identical physical run, a Flexible Configuration Management (FCM) branch of this version of NEMO was created, and all biogeochemical models were implemented in parallel within this branch and run separately. 

Simulations were initialised at year 1890 from an extant physics-only spin-up (ocean and sea-ice), to minimise undesirable transient behaviour in ocean circulation. In terms of ocean biogeochemistry, all model runs made use of a common dataset of three-dimensional fields for the initialisation of major tracers. Nutrients (nitrogen, silicon and phosphorus) and dissolved oxygen in this dataset were drawn from the World Ocean Atlas 2009 (Garcia et al., 2010a; Garcia et al., 2010b), while dissolved inorganic carbon (DIC) and alkalinity were drawn from the Global Ocean Data Analysis Project (GLODAP) (Key et al., 2004). GLODAP does not include a DIC field that is directly valid for 1890, so a temporally-interpolated field was produced based on GLODAP's "pre-industrial" (i.e. ~1800) and "1990s" fields of DIC. As there is currently no comprehensive spatial dataset of the micronutrient iron, participating models were permitted to make use of different initial distributions of iron (typically those routinely used by the models in other settings). All other biogeochemical fields (e.g. plankton, particulate or dissolved organic material) were initialised to arbitrary small initial conditions.

After initialisation at 1890, the models were run for 60 years (1890-1949 inclusive) under the so-called "normal year" of version 2 forcing for common ocean-ice reference experiments (CORE2-NYF; Large and Yeager, 2009). Subsequently, the models were run under transient, interannual forcing from the same dataset (CORE2-IAF) for a further 58 years (1950-2007 inclusive). CORE2 provides observationally derived geographical fields of downwelling radiation (separate long- and short-wave), precipitation (separate rain and snow), and surface atmospheric properties (temperature, specific humidity and winds), and is used in conjunction with bulk formulae to calculate net heat, freshwater and momentum exchange between the atmosphere and the ocean.

For all models, some degree of tuning occurred prior to this study, albeit in physical frameworks different (to varying degrees) to that used here. Tuning during this study was limited or absent between models, but some models, such as HadOCC and MEDUSA, may have benefitted from being previously tuned within the NEMO framework (although in a

different version and grid configuration). 158 Supplementary Figure S7 shows an intercomparison of the common NEMO physics with 159 observations for several key physical fields. In terms of SST, NEMO represents observed 160 patterns well, although simulates a warmer Gulf Stream and noticeably cooler temperatures 161 in the vicinity of the Labrador Sea. In conjunction with fresher salinities in the North Atlantic 162 (results not shown), these differences result in shallower depths of the mixed layer and 163 pycnocline in this region. By contrast, in the Southern Ocean both mixed layer depths and 164 the modelled pycnocline are markedly deeper than in observations. This latter regional bias 165 has biogeochemical consequences across all of the models examined here (see later). 166 2.2 Candidate model structures 167 168 The models evaluated within this study vary significantly in biological complexity. The key 169 features of the participating models are summarized below: HadOCC (Palmer & Totterdell, 2001): the Hadley Centre Ocean Carbon Cycle model 170 (HadOCC) model is a simple NPZD (Nutrient, Phytoplankton, Zooplankton, Detritus) 171 representation that uses N nutrient as its base currency but with coupled flows of C, 172 alkalinity and O2. The model was the ocean biogeochemistry component of the UK Met 173 174 Office's HadCM3 climate model, and was used for the first ever fully coupled carbon-climate study (Cox et al., 2000). 175 Diat-HadOCC (Halloran et al., 2010): is a development of the HadOCC model which includes 176 two phytoplankton classes (diatoms and "other phytoplankton") and representations of the 177 Si and Fe cycles, as well as a dimethyl sulphide (DMS) sub-model. The model is the ocean 178 biogeochemistry component of HadGEM2-ES (Collins et al., 2011), the UK Met Office's Earth 179 System model used to run simulations for CMIP5 and the Intergovernmental Panel on 180 Climate Change (IPCC) 5<sup>th</sup> Assessment Report (AR5). 181 182 MEDUSA-2 (Yool et al., 2011; Yool et al., 2013): Model of Ecosystem Dynamics, nutrient Utilisation, Sequestration and Acidification (MEDUSA) is an "intermediate complexity" 183 184 plankton ecosystem model designed to incorporate sufficient complexity to address key 185 feedbacks between anthropogenically-driven changes (climate, acidification) and oceanic biogeochemistry. MEDUSA-2 resolves a size-structured ecosystem of small 186 187 (nanophytoplankton and microzooplankton) and large (microphytoplankton and mesozooplankton) components that explicitly includes the biogeochemical cycles of N, Si 188 189 and Fe nutrients as well as the cycles of C, alkalinity and O2. As such, MEDUSA-2 is broadly

similar in structure to Diat-HadOCC, but includes several more recent parameterisations.

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PlankTOM6 & PlankTOM10 (Le Quéré et al., 2005): PlankTOM is a Dynamic Green Ocean 191 192 Model that represents lower-trophic level marine ecosystems based on Plankton Functional 193 Types (PFTs). A hierarchy of PlankTOM models exists that vary in the number of PFTs 194 resolved. Two members drawn from this stable were used in iMarNet. PlankTOM6 includes 195 six PFTs - diatoms, coccolithophores, mixed-phytoplankton, bacteria, protozooplankton and 196 mesozooplankton - while PlankTOM10 includes an additional four PFTs - Nitrogen-fixers, 197 Phaeocystis, picophytoplankton and macrozooplankton (Le Quéré et al. 2005; Buitenhuis et 198 al., 2013). The models include the marine cycles of C, N, O<sub>2</sub>, P, Si, a simplified Fe cycle, and 199 three types of detrital organic pools including their ballasting properties and estimates the 200 air-sea fluxes of CO<sub>2</sub>, O<sub>2</sub>, DMS, and N<sub>2</sub>O. PlankTOM6 and PlankTOM10 were developed by an 201 international community of ecologists and modellers to quantify the interactions between 202 climate and marine biogeochemistry, particularly those mediated through CO<sub>2</sub>. They make 203 use of extensive synthesis of data for the parameterisation of growth rates of PFTs (e.g. 204 Buitenhuis et al., 2006; 2010) and for the model evaluation (Buitenhuis et al., 2013).

ERSEM (Baretta et al., 1995; Blackford et al., 2004): European Regional Seas Ecosystem 205 206 Model (ERSEM) is a generic lower-trophic level/ model designed to represent the 207 biogeochemical cycling of C and nutrients as an emergent property of ecosystem 208 interaction. The ecosystem is subdivided into three functional types: producers 209 (phytoplankton), decomposers (bacteria) and consumers (zooplankton), and then further 210 subdivided by trait (size, silica uptake) to create a foodweb. Physiological (ingestion, respiration, excretion and egestion) and population (growth, migration and mortality) 211 212 processes are included in the descriptions of functional group dynamics. Four phytoplankton 213 (picophytoplankton, nanophytoplankton, diatoms and non-siliceous macrophytoplankton), 214 three zooplankton (microzooplankton, heterotrophic nanoflagellates and mesozooplankton) 215 and one bacteria are represented, along with the cycling of C, N, P, Si and O<sub>2</sub> through pelagic 216 (Blackford et al., 2004) and benthic (Blackford, 1997) ecosystems. ERSEM is used for shelf 217 seas water quality monitoring and climate impact assessment, has been coupled to fisheries 218 models (e.g. Barange et al., 2014), and is run operationally by the UK Met Office (e.g. 219 Siddorn et al., 2007).

The intercomparison process required limited changes to model organisation and code, and models retained disparate parameterisations for several overlapping processes, including ocean carbonate chemistry and air-sea exchange (HadOCC, Diat-HadOCC – Dickson & Goyet 1994, Nightingale et al., 2000; MEDUSA - Blackford et al., 2007; PlankTOM-6, PlankTOM-10 - Orr et al., 1999; ERSEM - Artoli et al., 2012). In the case of calcium carbonate (CaCO<sub>3</sub>) production, the models utilised a range of different parameterisations. HadOCC and Diat-HadOCC use a simple empirical relationship that ties CaCO<sub>3</sub> production to primary production. MEDUSA relates CaCO<sub>3</sub> production to export production, with a PIC:POC ratio (particulate inorganic carbon:particulate organic carbon ratio) dependent on calcite

229 230 231 232 233 234 235 236	saturation state. In PlankTOM-6 and PlankTOM-10, coccolithophore algae are explicitly modelled, with a fixed PIC:POC ratio. ERSEM relates CaCO <sub>3</sub> production to export production driven by nanophytoplankton losses, with a variable PIC:POC ratio dependent on temperature, nutrient limitation and calcite saturation state. Meanwhile, CaCO <sub>3</sub> dissolution was a simple exponential function of depth in the HadOCC models, with the other models modifying similar vertical dissolution with reference to the ambient saturation state of CaCO <sub>3</sub> .
237 238	The representation of biogeochemical cycles and biota in each model are summarized in Tables 1 and 2 respectively.
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240	2.3 Model evaluation
241	Assessment against observational datasets was made for a set of bulk ocean biogeochemical
242	properties that were common across all models: pCO <sub>2</sub> , alkalinity, dissolved inorganic carbon
243	(DIC), dissolved inorganic nitrogen (DIN), chlorophyll and primary production In all cases,
244	model results were regridded to the same geographical grid (World Ocean Atlas) and guided
245	by literature on appropriate skill metrics (e.g. Doney et al., 2009; Stow et al., 2009) model
246	skill was assessed through statistical techniques such as global surface field standard
247	deviation and spatial pattern correlation coefficients. In the biogeochemical regions of the
248	North Atlantic, Equatorial Pacific and Southern Ocean, depth profiles of model outputs were
249	also assessed against observations within the top 1000m of the water column.
250	Observational fields used within the model intercomparison are comprised of World Ocean
251	Atlas 2009 DIN (Garcia et al., 2010a), chlorophyll (O'Reilly et al., 1998) and pCO $_2$ (Takahashi
252	et al., 2009). Because of its biogeochemical importance, and the diversity in published
253	estimates, observational primary production is an average of three empirical models:
254	(Behrenfeld and Falkowski, 1997); (Carr et al., 2006) and (Westberry et al., 2008)- which are
255	all estimates derived from satellite ocean colour and SST. The observational fields of
256	chlorophyll and primary production used here represent averages over the 2000-2004 time
257	period. This same period is used throughout the following analysis as a standard interval
258	except in the case of DIC and alkalinity, which are analysed over the mean 1990-1999 period
259	corresponding to the GLODAP data product.
260	These fields were selected for several reasons. Firstly, they are ocean or biogeochemical

bulk properties for which there are global-scale observations. Secondly, these fields broadly

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represent foundational aspects of marine biogeochemical cycles. For instance, nutrients play a critical role in regulating the distribution and occurrence of marine plankton, while phytoplankton photosynthesis represents the vast majority of the primary energy source to marine ecosystems. Thirdly, the measurement of these fields is relatively well-defined with long-established standard methodologies. Properties that are directly related to biological entities, for instance biomass abundances, can be less precisely defined, difficult to match up with modelled quantities, or even absent from some models examined here. That said, the observational field of global scale primary production used here has a relatively high uncertainty because it is drawn from three methodologies which exhibit a large range (cf. Yool et al., 2013). Finally the examined properties are those which, if modelled poorly, legitimately cast doubt over the wider utility of a biogeochemical model in an earth systems context. Model results always depart from observations, but systematic disagreement with these basic observations is strongly suggestive of problems with process representation within a model. The model comparison focuses on the mean and seasonal cycle. It does not include evaluation of variability over interannual or longer timescales, in part because of limited data availability.

#### **278 3 Results**

#### 3.1 Model skill assessment

#### 3.1.1 Surface fields

Figures 1-3 (and Supplementary Figures S1-S3) show annual average fields from each of the models for a series of ocean properties, together with comparable observational fields. The figures also include a panel that shows the corresponding model-observation Taylor diagram (Taylor, 2001). These illustrate both the correlation between (azimuthal position) and relative variability (radial axis) of model and observations, such that models more congruent with observations generally appear closer to the reference marker on the x-axis of the diagram. As Taylor diagrams do not account for mean field biases (Joliff et al., 2009) these are provided separately in figure legends.

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Figure 1 shows annual average surface  $pCO_2$  fields for both models and observations, with correlation coefficients ranging from r=0.01 to r=0.68 (Takahashi et al., 2009). In general, the simpler models (HadOCC, Diat-HadOCC and MEDUSA-2) better capture the global spatial pattern of  $pCO_2$  (r=0.54 to r=0.68), but they overestimate the standard deviation in global surface  $pCO_2$  by up to a factor of 2. This overestimation of the variance in global surface  $pCO_2$  is a result of high modelled  $pCO_2$  values in the equatorial Pacific and in particular the

Eastern equatorial Pacific. In contrast, the more complex models (PlankTOM6, PlankTOM10 296 and ERSEM) perform considerably worse in terms of capturing global spatial patterns of 297 surface ocean pCO<sub>2</sub>. In particular, all three models underestimate the observed high pCO<sub>2</sub> 298 values along the equatorial Pacific ocean as well as the high coastal pCO2 values in that 299 300 region, opposite to the bias found in simpler models. However, the PlankTOM models 301 overall show comparable standard deviations in mean global surface pCO<sub>2</sub> to that seen in 302 observations. The negative pCO<sub>2</sub> biases in the equatorial Pacific exhibited by the PlankTOM6, PlankTOM10 303 and ERSEM models may be explained, at least in part, by the positive biases that these 304 models show for surface alkalinity in this region (Figure S3). The models with positive pCO<sub>2</sub> 305 biases in the equatorial Pacific (HadOCC, Diat-HadOCC and MEDUSA-2), do not have 306 307 negative surface alkalinity biases in this region but values are much closer to observations 308 (Figure S3). The root of these alkalinity biases lies in variation in PIC production by the 309 models in this region as discussed in greater detail below. Figure 2 illustrates model performance for annual average surface dissolved inorganic 310 nitrogen (DIN) concentrations. Here, all models capture global patterns relatively well, with 311 correlation coefficients >0.8, in part because of the initialisation from observations in 1890. 312 The model with the highest spatial pattern correlation coefficient is ERSEM, although it 313 314 slightly underestimates the global variability of DIN. The other models have lower spatial pattern correlation coefficients and generally overestimate the global variability of DIN. 315 PlankTOM6 performs below other models, while PlankTOM10 has similar performance as 316 the simpler models. In general, aside from ERSEM and PlankTOM10, most models show 317 elevated Pacific DIN, with the simpler models, MEDUSA-2 in particular, exhibiting high 318 319 equatorial anomalies. Finally, while ERSEM shows good agreement throughout most of the world ocean, both the North Atlantic and North Pacific show anomalously low annual 320 321 average DIN concentrations. 322 "Surface DIN concentrations are influenced by both the efficiency of primary production and 323 the efficiency of remineralisation both of which differ between models. Although we don't 324 explore the differences in remineralisation, the models which show positive DIN biases in the equatorial Pacific (HadOCC, Diat-HadOCC and MEDUSA-2), are generally shown to also 325 326 have positive integrated primary production biases in this region (Figure S1). To a lesser 327 extent the reverse is true of the models with negative DIN biases in the equatorial Pacific 328 (PlankTOM10 and ERSEM)." 329

331 332	models. The models with the highest correlation coefficients are PlankTOM10 (0.49) followed by MEDUSA-2 (0.36). All other models have correlation coefficients <0.2.
333	Anomalously high chlorophyll values in the equatorial Pacific and, especially, the Southern
334	Ocean significantly elevate the spatial variability of Diat-HadOCC above that of observations
335	(and all other models). More generally, with the exception of PlankTOM10, all of the models
336	show some degree of excess chlorophyll in the Southern Ocean, with Diat-HadOCC
337	exhibiting very high concentrations in this relatively unproductive region.
338	In addition to the ocean properties shown in Figures 1-3, complementary figures for
339	alkalinity, DIC and primary production can be found in the supplementary material (Figures
340	S1-S3). In each case, global annual average fields are shown together with the
341	corresponding Taylor diagram.
342	Table 3 shows the correlation coefficients and standard deviations normalised relative to
343	observations of the models for all six of the ocean properties (five surface fields plus depth-
344	integrated primary production). These are additionally colour-coordinated according to the
345	rank order of model performance, and the range of correlation coefficients over all of the
346	models is shown for each field. As already suggested above, model performance varies both
347	between fields and between models. All models perform consistently and relatively well for
348	DIN and DIC in part because of the "memory" of initial distributions. Model performance
349	varies more widely for pCO <sub>2</sub> and primary production and varies most widely for chlorophyll,
350	although it is consistently poor across all models.
254	Fig. 1. A second describe data to Table 2 be about the distribution of a of a consequence
351	Figure 4 summarises the data in Table 3 by showing the distribution of performance
352	rankings (both correlation coefficients and normalised standard deviations) across the
353 354	selected fields for each model, i.e. the number of first, second, etc., rankings for each
355	model. No model is shown to consistently outperform all other models across all metrics.  Indeed all models perform best in at least one metric, and similarly all models perform
356	worst in at least one metric. There is little discernable relationship between model
357	complexity and model performance. Indeed Table 3 shows that for 4 out of 6 fields the best
358	performing model in terms of correlation coefficients is a simpler model (i.e. HadOCC, Diat-
359	HadOCC or MEDUSA-2) and for 5 out of 6 fields the best performing model in terms of
360	normalised standard deviations is a more complex model (i.e. PlankTOM6, PlankTOM10 or
361	ERSEM).
362	These findings in annual average model performance are found to be consistent when
363	examined at monthly timescales (Figure 5).

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While the majority of biological activity in the ocean is concentrated in its surface layers,
biogeochemical fields in the deep ocean have a complex structure created through the
interaction of ocean physics with biologically-mediated processes such as export and
remineralisation. As such, model performance cannot be solely assessed from surface fields
of ocean BGC properties. To examine this, Figures 6 and 7 show the annual average depth
profiles of DIC and alkalinity for three important regions: the North Atlantic (Atlantic 060°N), Southern Ocean (≥60°S) and Equatorial Pacific (Pacific Ocean 15°S-15°N).

In Figure 6, all models are shown to capture the DIC profile in the Equatorial Pacific though HadOCC, Diat-HadOCC and MEDUSA-2 are somewhat closer to observations than ERSEM and the PlankTOM models. A similar situation is seen in the North Atlantic where the depth profiles of MEDUSA-2, HadOCC and Diat-HadOCC are closest to observations, although surface agreement is greater than that at depth. All models are shown to perform relatively poorly in the Southern Ocean, with much weaker gradients with depth than observations. HadOCC, Diat-HadOCC and ERSEM show gradients that are marginally closer to that observed, but all of the models consistently fail to reproduce the observed >100 mmol m<sup>-3</sup> surface-1000m increase. As Figure S7 shows, this common problem of vertical homogeneity between the models is driven by systematic biases in vertical mixing in this region, as well as known errors in ocean circulation (e.g. Yool et al., 2013).

The annual average depth profiles of alkalinity are shown in Figure 7. In the North Atlantic, HadOCC and Diat-HadOCC are closer to observations while ERSEM and, particularly, MEDUSA-2 are further away from observations (but in opposite directions). Again, and for the same reasons as outlined above, no model performs well at capturing the depth profile observed in the Southern Ocean. In the Equatorial Pacific all of the models have similar alkalinity at depth but diverge from observations towards the surface. The near-surface depth profiles in HadOCC, Diat-HadOCC and MEDUSA-2 are closest to observations in that region. Alkalinity shows very little variability with depth in the PlankTOM6, PlankTOM10 and ERSEM models and is higher than observations in near-surface waters (>100 meg m<sup>-3</sup>). This excess alkalinity may explain the broadly lower pCO<sub>2</sub> values visible in this region in Figure 1. The source of this bias in surface alkalinity is, at least in part, due to disparity in modelled CaCO<sub>3</sub> production in this region. As Supplementary Figures S8-S10 show, PlankTOM6, PlankTOM10 and ERSEM export negligible particulate inorganic carbon (PIC; Figure S9) relative to particulate organic carbon (POC; Figure S8) in this region. This results in low rain ratios (Figure S10) and the divergence of DIC and alkalinity performance of these models in this region. The lack of PIC export in these models runs contrary to observations (e.g. Dunne et al., 2007), but reflects the current difficulty in modelling CaCO<sub>3</sub> production – which HadOCC, Diat-HadOCC and MEDUSA-2 circumvent by simplistic empirical parameterisations.

The depth profiles of DIN and  $O_2$  are given in the supplementary material (Figures S4-5).

## 3.2 Computational benchmarking

Computational timing tests (CPU time) were carried out relative to the ocean component of the HadGEM3 (Hewitt et al., 2011) model (ORCA1.0L75), on standard configurations of 128 and 256 processors on an IBM Power7 machine. As would be intuitively expected, the cost of candidate ocean biogeochemical models is found to be higher for models with more tracers regardless of the number of processors used. While there are deviations in both directions between the models, broadly there is a linear relationship between number of model tracers and compute cost (Figure S6) reflecting the significant cost of applying advection and mixing terms to each tracer.

Using ERSEM (the computationally most expensive model) increases computational cost approximately 6-fold relative to HadOCC when 128 processors are used. This relative increase in computational cost is reduced to approximately 4.5-fold when 256 processors are used. PlankTOM10 has the greatest relative reduction (36.6%) in computational cost when run on 256 processors as opposed to 128, although this model would still increase the total cost of the ocean component by a factor of 5 relative to a physics-only ocean, compared to a factor of 1.5 for HadOCC (Table 4).

## 4 Discussion

Our model comparison suggests that for global annual average surface fields, global monthly average surface fields and annual average depth profiles in three oceanographic regions there is little evidence that increasing the complexity of OBGC models leads to improvements in the representation of large scale ocean patterns of bulk properties. In some cases, the comparison suggests that simpler OBGC are closer to observations than intermediate or complex models for the standard assessment metrics used here.

The biologically simpler models HadOCC, Diat-HadOCC and MEDUSA-2 are shown to have generally higher global spatial pattern correlation coefficients of pCO<sub>2</sub>, DIC and alkalinity at both annual and monthly temporal resolution (Figures 1, 5 and Table 3). The more complex models PlankTOM6, PlankTOM10 and, in the case of DIC, ERSEM, have annual and monthly standard deviations that are generally closer to observations than the simplest two models (HadOCC and Diat-HadOCC). As such, we find no robust relationship between model complexity and model skill at capturing global scale distributions of surface pCO<sub>2</sub>, DIC and alkalinity. The biologically simpler models are shown to generally best capture the depth

profiles of DIC and alkalinity in the North Atlantic and Equatorial Pacific (Figures 6-7), 435 436 possibly because their biological export production can more easily be tuned to maintain 437 the observed vertical gradients. There are however ocean biogeochemical fields where models of greater biological 438 complexity tend to equate to improved model skill. The annual and monthly global 439 correlation coefficients of the PlankTOM models are shown to be closest to observations for 440 chlorophyll and primary production fields (Figures 3 and Table 4). These PlankTOM models 441 do not consistently produce the annual chlorophyll and primary production field standard 442 deviations closest to observations (Table 4), however at monthly resolution their field 443 standard deviations are the most consistent across models (Figure 5). 444 445 The comparison of depth profiles shows that despite all models being initialised from the same observational fields, there is quite a lot of divergence even at depths of less than 446 447 1000m. In some cases, such as alkalinity in the Southern Ocean (Figure 7), all models have a similar systematic bias compared to observations. This is suggestive of the influence of 448 449 errors within the physical ocean model. That is, the ocean biogeochemistry may be 450 influenced to a greater extent by the physical ocean model and hence there is a common 451 response across models. For other fields such as DIN in the Southern Ocean and Equatorial 452 Pacific (Figure S5), models have both positive and negative biases compared to observations 453 suggestive of a greater relative role of the OBGC model than the physical model. 454 It is clear that more biologically complex models are required to more completely assess the impacts of environmental change on marine ecosystems. By representing processes that are 455 456 not present in simpler models, the more complex models are also able to represent 457 additional factors such as climatically-active gases (e.g. DMS, N2O). Assessment of such 458 representations however fell outside the scope of this paper. Models of intermediate 459 complexity (e.g. Diat-HadOCC and MEDUSA-2) are shown in this inter-comparison to 460 reproduce large scale ocean biogeochemistry features relatively well, yet minimise 461 computational cost and have sufficient biological complexity to allow important ESM 462 questions to be explored, including those that require an explicit iron cycle (e.g. ocean iron 463 fertilisation). It should be noted that models implemented within the NEMO physical ocean framework 464 prior to this inter-comparison project had an advantage over those new to this framework. 465 This is a somewhat unavoidable consequence of what is also one of this inter-comparison 466 467 study's main strengths, namely that the models were adapted to use the same ocean physics framework. Specifically, the HadOCC and MEDUSA-2 model developers were familiar 468 469 with NEMO v3.2 and had some previous opportunity to tune models. Linked to this is the

470 question of how dependent the results were on parameter values. Although model 471 developers were afforded a limited opportunity to tune parameters, given further time to 472 tune one would expect improved performance, especially for those models that had not 473 been previously implemented within NEMO v3.2. The rationale for the chosen fields of intercomparison was, as stated previously, that they 474 are common across all models and are key facets of global marine biogeochemistry. It could 475 however be argued that these bulk fields were insufficient to adequately assess all models 476 and in particular the most complex models. Further analysis, beyond the scope of this paper 477 will undertake as thorough an analysis of the biological components as each model will 478 479 support. 480 Finally although computational cost is discussed as a pragmatic driver of OBGC model selection, it should be noted that computer power is continuously increasing and the 481 482 intercomparison results presented here may differ for an alternative spatial resolution 483 ocean grid requiring greater computational resources. In addition, ongoing efforts to 484 transport passive ocean tracers on degraded spatial scales (e.g. Levy et al., 2012) have the 485 potential to result in computational savings that would realistically permit the 486 implementation of higher complexity OBGC models within ESMs. **5 Conclusions** 487 488 The 6 ocean biogeochemical models analysed within this inter-comparison cover a large range of ecosystem complexity (from 7 tracers in HadOCC to 57 in ERSEM), and therefore 489 490 result in a range of approximately 5 in computational costs (from increasing the cost of the physical ocean model by a factor of 2 to a factor of 10). Results suggest little evidence that 491 492 higher biological complexity implies better model performance in reproducing observed global-scale bulk properties of ocean biogeochemistry. 493 As no model is found to have the highest skill across all metrics and all are most or least 494 495 skilful for at least one metric, our results suggest that it is in the interest of the international 496 climate modelling community to maintain a diverse suite of ocean biogeochemical models. 497 One priority for the next generation of Earth System Models (CMIP6) is to enhance model resolution in the hope that it will resolve some of the existing biases in climate 498 499 models. This puts pressure on the computing time available for representing biological complexity. Our results suggest that intermediate complexity models (such as MEDUSA-2 500 501 and Diat-HadOCC) offer a good compromise between the representation of biological complexity (through their inclusion of an iron cycle) and computer time, given their 502 503 relatively good performance in reproducing bulk properties. However, intermediate complexity models are limited in the detail to which they can address climate feedbacks and 504

505 506	it may be that more complex models can in future provide additional insight, based on ongoing measurements and data syntheses.
507 508	The quest for increasing resolution in ESMs is unlikely to end soon, as the resolution needed to resolve eddies in the ocean (1/8 degree or less) needs to be achieved before
509	important improvements in representing climate dynamics are achieved. Most ESMs being
510	developed for the next CMIP phase will have a grid of 1/2 to 1/4 degree. Even with
511	increasing computational power and schemes for accelerating transport of passive tracers
512	(Levy et al., 2012) available, other priorities (e.g. ensemble simulations for risk assessments)
513	may still make it difficult to prioritise the representation of biogeochemical complexity in
514	ESMs. In order to achieve scientific progress on important questions of the
515	interactions between marine biogeochemistry and climate, it is thus important that
516	lower resolution ESMs that prioritise biogeochemical complexity are maintained and used in
517	CMIP exercises in parallel to higher resolution models.
	ever ever ever ever the important resolution models.
518	
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# Table 1. Biogeochemical cycles represented in each candidate model.

	HadOCC	Diat-HadOCC	MEDUSA-2	PlankTOM6	PlankTOM10	ERSEM
N	<b>✓</b>	1	1	1	<b>✓</b>	1
P					1	1
Si		1	✓	1	1	1
Fe		1	1	1	1	1
С	1	1	1	1	<b>✓</b>	1
Alkalinity	1	1	1	1	<b>✓</b>	1
O <sub>2</sub>	1	1	1	1	1	1

Table 2. Composition of the marine ecosystems represented in each candidate model, along with the total number of biogeochemical tracers (including those detailed in Table 1).

	HadOCC	Diat-HadOCC	MEDUSA- 2	PlankTOM6	PlankTOM10	ERSEM
Generic Phytoplankton	1	1		1	1	
Diatoms		1	1	1	1	1
Large Phytoplankton						1
Picophytoplankton			1		1	1
Coccolithophores				1	1	1
N <sub>2</sub> fixers					1	
Flagellates						1
Phaeocystis					1	
Generic Zooplankton	1	1				
Microzooplankton			1	1	1	1
Mesozooplankton			1	1	1	1
Macrozooplankton					1	
Heterotrophic						1
Nanoflagellates						
Picoheterotrophs				1	1	1
Tracers	7	13	15	25	39	57

Table 3. Model-observation correlation coefficients (R) and standard deviations normalised by the standard deviation of observations ( $\sigma$ ) for all examined annual surface fields and depth integrated primary productivity. Colours indicate model ranking and are organised through the worst performing model in red to the best performing model in dark blue (through orange, yellow, green and light and dark blue).

Model	pCO <sub>2</sub>		DIN		Chl.		Alkal	inity	DIC		Primai	γ
											Production	
	R	σ	R	σ	R	σ	R	σ	R	σ	R	σ
HadOCC	0.68	1.92	0.88	1.20	0.30	0.68	0.91	1.19	0.93	1.18	0.19	0.92
Diat-	0.54	1.77	0.90	1.20	0.15	2.65	0.91	1.19	0.93	1.13	0.13	1.51
HadOCC												
MEDUSA-2	0.64	1.56	0.85	1.21	0.36	0.40	0.88	1.14	0.92	1.17	0.64	1.10
PlankTOM6	0.34	1.03	0.79	1.20	0.32	1.08	0.70	0.88	0.75	0.96	0.47	0.61
PlankTOM10	0.29	0.94	0.88	1.19	0.50	0.43	0.58	1.16	0.65	1.08	0.53	0.74
ERSEM	0.01	2.04	0.94	0.95	0.04	0.91	0.84	1.18	0.86	1.07	-0.08	1.12
Range	0.64	1.09	0.15	0.26	0.46	2.64	0.33	0.31	0.28	0.23	0.72	0.90

Table 4. Computational cost of each candidate model when coupled to the ocean component of HadGEM3, relative to a physics-only simulation with the same ocean model (ORCA1.0L75). A cost of 2.0 indicates that adding the biogeochemistry model doubles total simulation cost. Timings are shown for simulations carried-out on 128 and 256 processors of an IBM Power7 machine.

	Cost	Cost
Model	(128 processors)	(256 processors)
HadOCC	1.75	1.48
Diat-HadOCC	2.36	1.88
MEDUSA-2	2.73	2.10
PlankTOM6	5.11	3.52
PlankTOM10	7.74	4.90
ERSEM	10.36	6.87

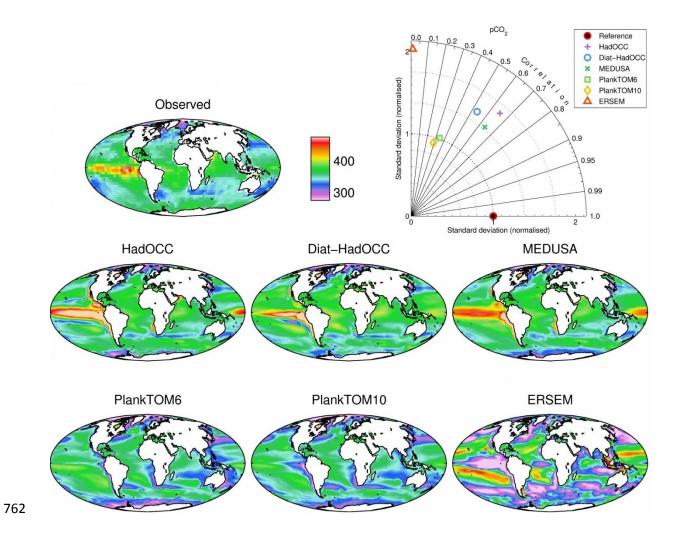


Figure 1. Observational (Takahashi et al., 2009; top left) and modelled annual average surface ocean pCO $_2$  (µatm) for year 2000. Mean field values: observations 357.7; HadOCC 368.8; Diat-HadOCC 369.2; MEDUSA 368.5; PlankTOM6 349.8; PlankTOM10 349.5; ERSEM 343.0.

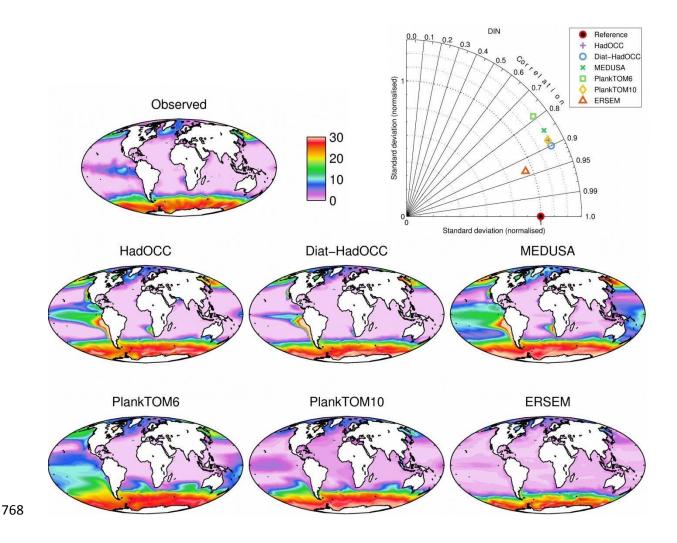


Figure 2. Observational (World Ocean Atlas, 2009; top left) and modelled annual average surface ocean Dissolved Inorganic Nitrogen (mmol m<sup>-3</sup>) for the period 2000-2004. Mean field values: observations 5.24; HadOCC 7.88; Diat-HadOCC 6.33; MEDUSA 10.18; PlankTOM6 9.45; PlankTOM10 7.25; ERSEM 4.58.

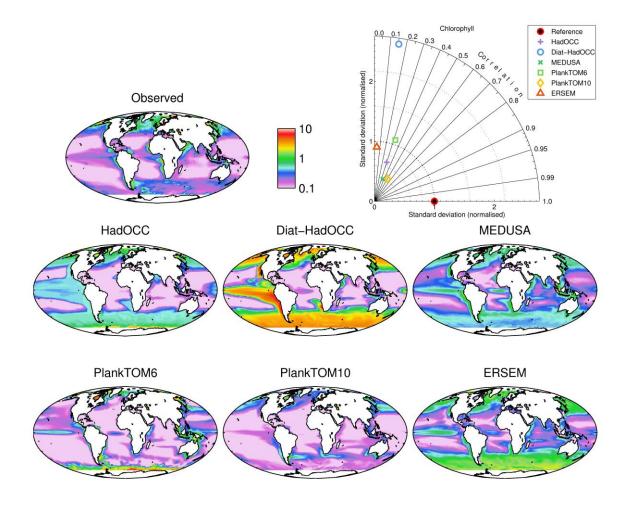


Figure 3. Observational (SeaWiFS; top left) and modelled annual average surface ocean chlorophyll (mg m<sup>-3</sup>) for the period 2000-2004. To avoid biasing the plots, observational data and model output are only shown for regions in which all months were represented at least once across all of the sampled years. Mean field values: observations 0.215; HadOCC 0.347; Diat-HadOCC 1.170; MEDUSA 0.346; PlankTOM6 0.312; PlankTOM10 0.160; ERSEM 0.501.

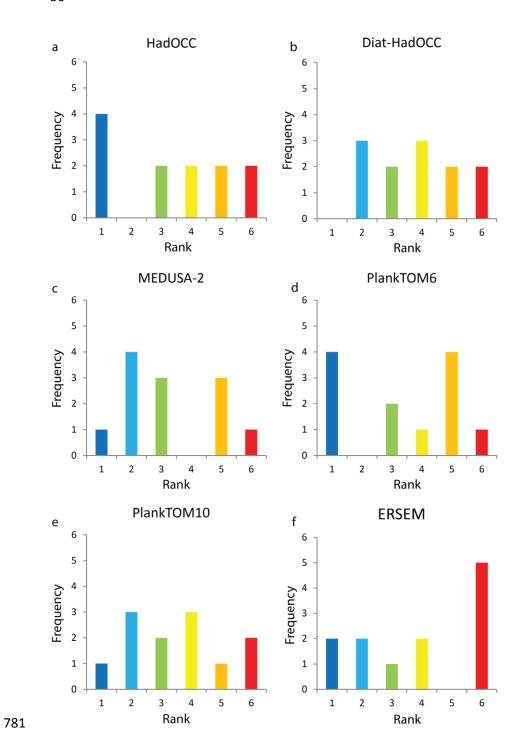


Figure 4. Frequency distributions of best- to worst-performances for each model, in terms of correlation coefficients and normalised standard deviations of annual surface fields and depth integrated primary productivity. Colours follow those of Table 1.

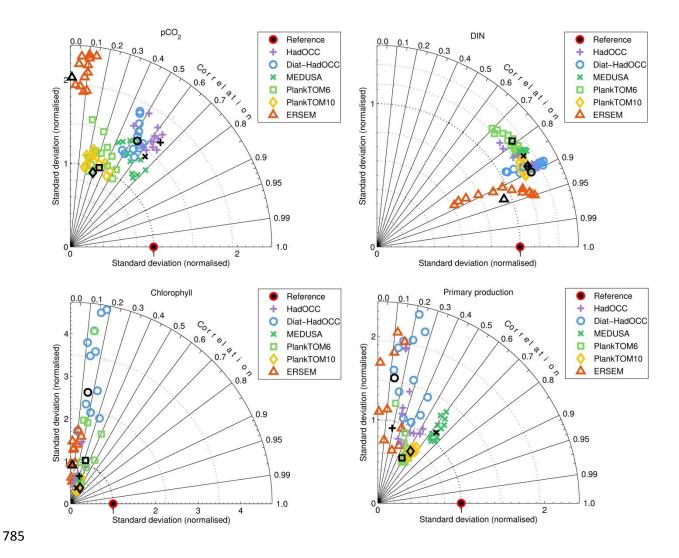


Figure 5. Monthly Taylor plots for pCO<sub>2</sub>, Dissolved Inorganic Nitrogen (DIN), chlorophyll and primary production for all models relative to observations. Annual averages are shown in black. Note that negative correlation coefficients are not shown in the Taylor plot.

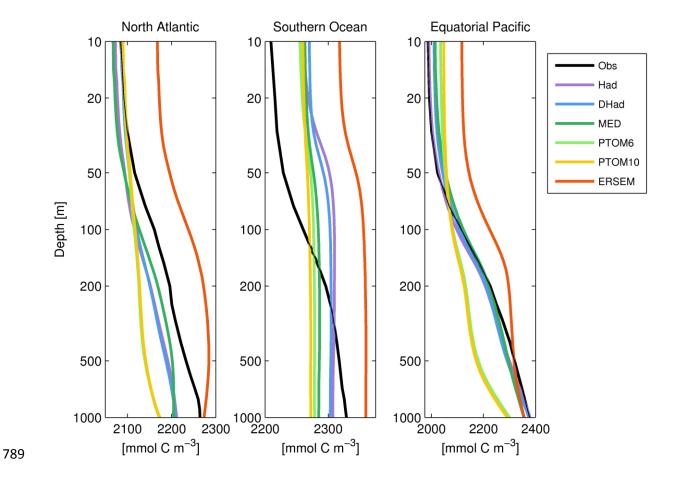


Figure 6. Observed (black; GLODAP) and modelled profiles of Dissolved Inorganic Carbon (mmol C  $\rm m^{-3}$ ) in the North Atlantic (0°N to 60°N), Southern Ocean (90°S to 60°S) and Equatorial Pacific (15°S to 15°N). Vertical scaling is logarithmic (log<sub>10</sub>).

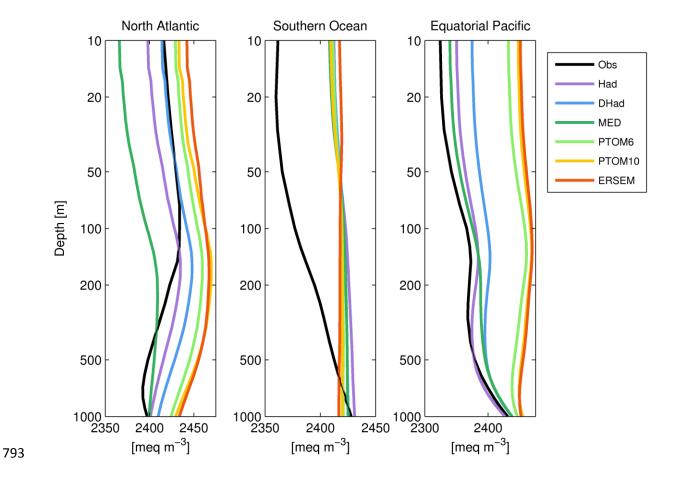


Figure 7. Observed (black; GLODAP) and modelled profiles of alkalinity (meq m $^{-3}$ ) in the North Atlantic (0°N to 60°N), Southern Ocean (90°S to 60°S) and Equatorial Pacific (15°S to 15°N). Vertical scaling is logarithmic (log<sub>10</sub>).