# Supplemental Material: A diatom extension of the cGEnIE Earth system model – EcoGEnIE 2.0

EcoGEnIE 2.0 is a trait-based functional type ecosystem model developed from EcoGEnIE 1.0 (Ward et al., 2018) to include diatom dynamics and silicon cycle in the Earth System model, cGEnIE. EcoGEnIE 1.0 captured the dynamic of size classes of phytoplankton and zooplankton based on allometric relationships for, e.g., uptake rate, grazing rate and nutrient affinity. EcoGEnIE 2.0 depicts three new functional phytoplankton groups: picoplankton, eukaryote and diatom, each with two to three size classes relevant to their type (Table S1). To distinguish the different groups of phytoplankton, we used the allometric relationships as defined by Dutkiewicz et al. (2020) for the maximum photosynthetic rate for picoplankton and diatoms (Table S1).

	EcoGEnIE 1.0	EcoGEnIE 2.0	
	(Ward et al., 2018)	This study	
<u>ECOSYSTEM</u>			
	8 size classes of phytoplankton	3 size classes of diatoms (2.0, 20.0, 200.0)	
Ecosystem	and zooplankton (0.6, 1.9, 6.0,	2 size classes of picoplankton (0.6, 2.0)	
structure	19.0, 60.0, 190.0, 600.0, 1900.0)	2 size classes of eukaryote (20.0, 200.0)	
		4 size classes of zooplankton (6.0, 20.0, 200.0, 2000.0)	
Maximum photosynthetic rate	Unimodal relationship for all	Diatoms: 3.9V <sup>-0.08</sup> d-1 (Dutkiewicz et al., 2020)	
	plankton	Picoplankton: 0.9V <sup>0.08</sup> d <sup>-1</sup> (Dutkiewicz et al., 2020)	
		Eukaryotes: 2.2V <sup>-0.08</sup> d <sup>-1</sup> (assuming 20% lower than	
		diatoms)	
Diatom trade-offs	N/A	Benefits: Higher Pmax and grazing protection	
		Cost: Si requirement	
Light limitation		Modified EcoGEnIE 1.0 with diagnosed mixed-layer	
		light penetration and no light under sea-ice	
BIOGEOCHEMISTRY			
Iron cycle	Dust source: (Mahowald et al.,	Dust source: Albani et al. (2016)	
	1999)	TDFe and TL	
	TDFe and TL	$DOM\tau = 0.5$	
	$DOM\tau = 1$	Fesol = $0.00244$	
	Fesol = 1	Kscav = 0.225	
	Kscav = 1.338		
Silicon cycle	N/A	bg_par_bio_red_POC_opal=0.65	
		bg_ctrl_bio_remin_opal_fixed=.false.	
		bg_par_bio_remin_sinkingrate=125.0	

Extend the	N/A	gm_par_geochem_Tmin = -2.0	
temperature range		gm_par_geochem_Tmax = 45.0	
for solubility and		gm_par_carbchem_Tmin = -2.0	
geochemical		gm_par_carbchem_Tmax = 45.0	
constants			
Instantaneous	N/A	bg_par_bio_remin_sinkingrate_physical=9.9E9;	
remineralisation		bg_par_bio_remin_sinkingrate_reaction=125.0	
Geochemical	N/A	bg_par_bio_geochem_tau=90.0	
reaction timescale			
Relative	N/A	eg_par_beta_POCtoDOC=0.75	
partitioning of C			
into DOM			
<u>PHYSICS</u>			
Configuration	Worlg4 (Worjh2 with modified	Worjh2 (Cao et al., 2009)	
	wind)		
Mixed-layer	On	Off	
scheme			

Table S1: EcoGEnIE 2.0 model setup in comparison to EcoGEnIE 1.0

# Plankton physiology

State variables and their rates of change are defined by a variety of ecological processes. Further state variables are fully described in Ward et al. (2018).

## **Temperature limitation**

Metabolic processes within cells are temperature limited. We account for temperature limitation applying the following exponential function of temperature to all plankton.

$$\gamma^T = e^{A(T - T_{ref})} \tag{1}$$

Temperature sensitivity is described by the constant A, T is ambient water temperature in  ${}^{\circ}$ C and  $T_{ref}$  is a reference temperature where  $\gamma^T = 1$ .

# **Nutrient uptake**

Environmental availability ( $[R_{i_r}]$ ) of a nutrient ( $i_r = i_b$ ) governs phytoplankton uptake rate, along with maximum uptake rate ( $V^{max}$ ), quota saturation term, temperature limitation and nutrient affinity ( $\alpha_{i,i_r}$ ).

$$V_{j,i_r} = \frac{v_{j,i_r}^{max} \alpha_{j,i_r}[R_{i_r}]}{v_{j,i_r}^{max} + \alpha_{j,i_r}[R_{i_r}]} Q_{j,i_b}^{stat} \cdot \gamma^T$$

$$\tag{2}$$

This equation modifies the Michaelis-Menten-type response by including nutrient affinity rather than the half saturation constant.

# Plankton "quota" saturation

Saturation of nutrient biomass relative to carbon is prevented by setting the uptake capacity to zero when the cellular nutrient quota, Q, is satisfied (Ward et al., 2012). This quota is determined by the ratio of nutrients assimilated to carbon biomass.

$$Q_{j,i_b}^{stat} = \left(\frac{q_{j,i_b}^{max} - q_{j,i_b}}{q_{j,i_b}^{max} - q_{j,i_b}^{min}}\right)^h \tag{3}$$

The general uptake regulation term,  $i_b$ , for a given element j is a linear function of the nutrient status, which is altered by the shape parameter h = 0.1 (Geider et al., 1998).

#### **Photosynthesis**

We use a photosynthesis model for phytoplankton adapted from Geider et al. (1998) and Moore et al. (2001) where light limitation  $(\gamma_{j,I})$  relates to the Poisson function. This function depends on local irradiance (I) depends on the chlorophyall a: carbon ratio  $(Q_{j,Chl})$  and iron-dependent initial slope of the P-I curve  $(\alpha \cdot \gamma_{j,Fe})$ .

$$\gamma_{j,l} = 1 - \exp\left(\frac{-\alpha \cdot \gamma_{j,Fe} \cdot Q_{j,Chl} \cdot l}{P_{j,C}^{sat}}\right) \tag{4}$$

 $P^{sat}$  refers to the maximum light-saturated growth rate, which depends on a maximum rate of  $P^{max}$  with respect to the temperature and nutrient limitations.

$$P_{j,C}^{sat} = P_{j,C}^{max} \cdot \gamma^{T} \cdot \min \left[ \gamma_{j,P}, \gamma_{J,Fe} \right]$$
 (5)

The resulting gross photosynthetic growth rate  $(P_{j,C})$  is thus calculated as:

$$P_{j,C} = \gamma_{j,I} P_{j,C}^{sat} \tag{6}$$

## Grazing

We determine a predator's  $(j_{pred})$  predator-biomass specific grazing rate on its prey  $(j_{prey})$  by the overall grazing rate, the prey switching term  $(\phi)$  and prey refuge  $(1 - e^{\Lambda \cdot F_{j_{pred},C}})$  where grazing will reduce if availability is low.

$$G_{j_{pred},j_{prey},C} = \gamma^T \cdot G_{j_{pred},C}^{max} \cdot \frac{F_{j_{pred},C}}{k_{j_{pred},C} + F_{j_{pred},C}} \cdot \phi_{j_{pred},j_{prey}} \cdot (1 - e^{\Lambda \cdot F_{j_{pred},C}})$$
(7)

 $G^{max}$  represents the maximum grazing rate, with a half-saturation concentration for all prey denoted  $k_{jprey,C}$  and total available food as  $F_{jpred,C}$ . The "prey-switching" term is optional with the details outlined in Ward et al. (2018). We assume that predators do not feed on detrital organic matter.

#### **Mortality**

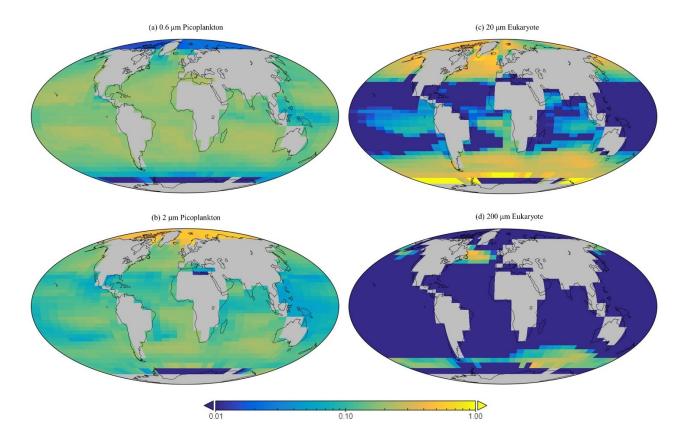
Biomass loss also occurs via mortality via a linear biomass mortality rate  $(m_i)$ .

$$m_j = m_p (1 - e^{-10^{10} \cdot B_{j,C}})$$
 (8)

 $m_j$  reduces when population carbon biomass (represented by the vector  $B_C$ ) is smaller than  $10^{-10}$  mmol C m<sup>-3</sup>, to help support low biomass population to ensure that every surface grid cell houses a viable population.

## Other plankton distribution

The carbon biomass distributions of the base functional types of our plankton community and their grazers (zooplankton) are shown in Figure S1 and S2. Global distributions of 0.6 µm picoplankton are generally consistent across high and low latitudes, with subtropical gyres housing far less of the 2 µm class. Eukaryotes, being of the micro size, exhibit distinct regions of habitancy relative to picoplankton, their larger 200 µm class is restricted to high latitudes whilst the 20 µm class also emanates in equatorial zones. Respective zooplankton size classes are intuitively determined by the presence of their prey (i.e. groups at least 10 times smaller than them), thus we observe larger zooplankton occupying increasingly smaller ecological niches.



**Figure S1.** Surface concentrations of carbon biomass for picoplankton ((a) and (b)), and eukaryotes ((c) and (d)) size classes (mmol C m<sup>-3</sup>).

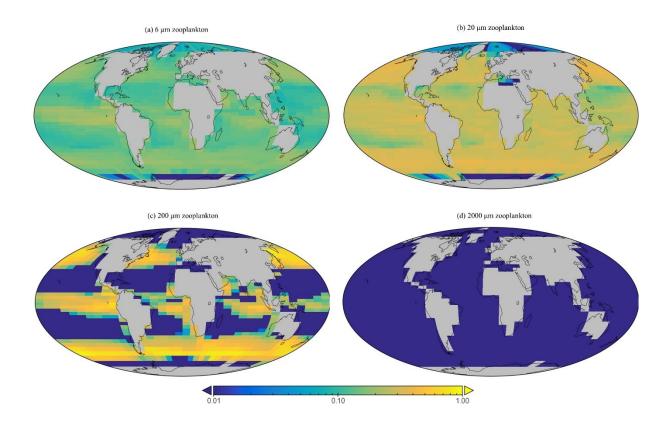


Figure S2. Surface concentrations of carbon biomass for each zooplankton size class (mmol C m<sup>-3</sup>).

# M-score trade-off

Figure S3 shows a trade-off in oxygen and phosphate M-scores, a factor we took into consideration when selecting our best run.

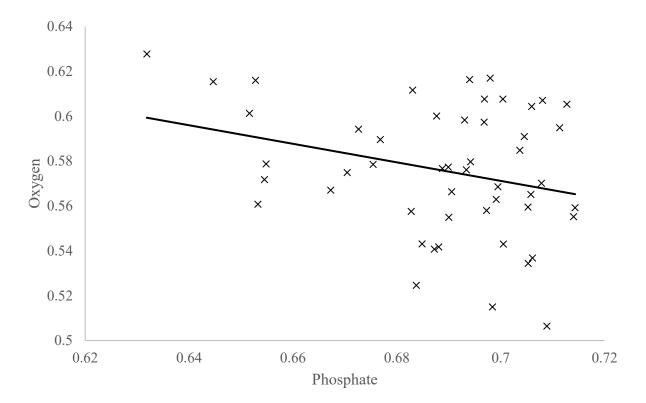


Figure S3. Top 50 average M-score runs with oxygen and phosphate plotted against each other.

## **Supplemental References**

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