

## Simulating ecosystem dynamics and marine

 biogeochemical cycles with multiple plankton functional typesJun Yu, Kristen M. Krumhardt, J. Keith Moore, Robert T. Letscher, Shanlin Wang, Nicola Wiseman, Matthew C. Long, Keith Lindsay, Michael Levy, Colleen M. Petrik, Adam C. Martiny
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## The old ecosystem model (CESM2-4P1Z)

(a) MARBL-4P1Z model


## The new ecosystem model (CESM2-8P4Z)

(b) MARBL-8P4Z model


## The grazing terms

$$
\begin{aligned}
G & =\frac{g^{\max } \cdot T_{f} \cdot \sum_{i=1}^{n} \boldsymbol{d}_{\boldsymbol{i}} p_{i}}{K+\sum_{i=1}^{n} \boldsymbol{d}_{\boldsymbol{i}} p_{i}} \\
\boldsymbol{d}_{\boldsymbol{i}} & =\frac{p_{i}}{\sqrt{p_{1}^{2}+p_{2}^{2}+\cdots+p_{n}^{2}}}
\end{aligned}
$$

- The feeding strategy follows a modified Holling Type II relationship;
- We initially set decreased maximum specific grazing rates ( $\mathrm{g}^{\text {max }}$ ) with predator size (Hansen et al., 1997);
- Density-dependent grazing to stabilize model dynamics by creating a refuge for the less abundant prey species.



## Model performance: nutrients field



## Phytoplankton biomass


(b) model prochlorococcus ( $0-50 \mathrm{~m}$ ) mmol m-3


(d) model synechococcus ( $0-50 \mathrm{~m}$ ) $\mathrm{mmol} \mathrm{m}-3$

${ }^{\text {sos }}$ (e) in situ pico-eukaryotes ( $0-50 \mathrm{~m}$ ) mmol m-3
(f) model pico-eukaryotes ( $0-50 \mathrm{~m}$ ) mmol m-3


* MAREDAT dataset faces considerable uncertainties related to cell counts, unidentified species, and conversions of cell abundance to biomass, thus carbon conversion estimates are susceptible to large errors if cell size is not accurately assessed.



## Other constraints

(a) Modeled Total Alkalinity ( $r=0.85$ )

(b) Glodap Alkialinity



Other constraints for modeled phytoplankton performance:

- Modeled Si performance for diatoms;
- Modeled Alk performance for coccolithophores;
- Modeled N -fixation performance for diazotrophs.


## Zooplankton biomass


 (c) macrozoo consume $5.36 \%$ of NPP

(b) mesozoo consume 19.72\% of NPP, Ref: 10\%-15\% (Calbet 2001)


Table 1. Regional comparisons of system characteristics from the full data set of dilution experiments. Data are distinguished among oceanic, coastal (overlying the continental shelf), and estuarine habitats in the upper table and among tropical/subtropical, temperate/subpolar, and polar habitats in the lower table. Mean values ( $\pm$ standard errors) are given for initial Chl $a$, phytoplankton growth rate ( $\mu \mathrm{g}$ ), grazing mortality (m), \% Chl $a$ grazed day ${ }^{-1}$, and \% primary production (PP) grazed day ${ }^{-1}$. Growth and mortality rate averages are significantly different among zones and climates ( $p<0.05$, Tukey-Kramer test), except for oceanic and coastal ( $\mu \mathrm{m}$ and m ), and for tropical and temperature $(\mu)$.

|  | Chl $a\left(\mu \mathrm{~g} \mathrm{~L}^{-1}\right)$ | $\mu\left(\right.$ day $\left.^{-1}\right)$ | $\mathrm{m}\left(\right.$ day $\left.^{-1}\right)$ | \% Chl $a$ grazed | \% PP grazed |
| :--- | :---: | :---: | :---: | :---: | :---: |
| Oceanic | $0.58 \pm 0.03$ | $0.59 \pm 0.02$ | $0.39 \pm 0.01$ | $41.5 \pm 1.4$ | $69.6 \pm 1.5$ |
| Coastal | $3.06 \pm 0.53$ | $0.67 \pm 0.05$ | $0.40 \pm 0.04$ | $47.3 \pm 4.4$ | $59.9 \pm 3.3$ |
| Estuarine | $13.0 \pm 1.8$ | $0.97 \pm 0.07$ | $0.53 \pm 0.04$ | $78.7 \mp 7.3$ | $59.7 \pm 2.7$ |
| Tropical | $1.01 \pm 0.21$ | $0.72 \pm 0.02$ | $0.50 \pm 0.02$ | $55.1 \pm 2.3$ | $74.5 \pm 2.0$ |
| Temperate | $5.18 \pm 0.66$ | $0.69 \pm 0.03$ | $0.41 \pm 0.02$ | $51.4 \pm 2.9$ | $60.8 \pm 1.8$ |
| Polar | $0.62 \pm 0.06$ | $0.44 \pm 0.05$ | $0.16 \pm 0.01$ | $19.5 \pm 2.1$ | $59.2 \pm 3.3$ |

## Biogeochemical fluxes









## Improved seasonal variations

## 8P4Z



## 4P1Z


(a) Chlorophyll Northern box

(b) Chlorophyll Southern box



## Improved phytoplankton community composition

Satellite
(a) SeaWiFS picoC \% (0.5-2 um in diameter)


8P4Z
4P1Z

sown (b) 8 P 4 Z picoC \%



## Conclusions

- We developed a next-generation marine biogeochemical model with an expanded ecosystem in CESM2-MARBL, with the ability to model key species observed in the ocean;
- The 8P4Z model generally reproduces the observed phytoplankton community composition and stoichiometry, facilitating reasonable estimations of integrated biogeochemical elemental fluxes;
- The 8P4Z model improves the modeled chlorophyll spring bloom compared with the simplified ecosystem CESM, benefiting from a better representation of bottom-up (resource competition) and top down (target grazers) controls.

