Can pCO₂ observations help constrain predatorprey interactions in a biological model?

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Biogeochemical ROMS implementation for Atlantic Canada



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- 10 km horizontal resolution
- 30 vertical layers
- Physical B.C. from Urrego-Blanco & Sheng (2012)
- Biochemical B.C. from observations
- 3-hourly ECMWF ERA-Interim atmospheric forcing
- 12 major rivers
- Tides
- No ice
- HSIMT advection scheme









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Why does our model not capture the rapid drawdown of pCO₂?

(1) Magnitude of P growth(2) Timing & duration of P growth

---- Growth rates ---- Grazing rates



switching to a 1-D environment



Spring Bloom Diagnostics:
1. P-Z dynamics
2. pCO₂ seasonality
3. Cumulative sinking



CASE 1 low P growth high Z grazing

CASE 2 low P growth low Z grazing **CASE 3 high P growth** low Z grazing





CASE 1 low P growth high Z grazing

CASE 2 low P growth low Z grazing **CASE 3 high P growth** low Z grazing





CASE 1 low P growth high Z grazing

CASE 2 low P growth low Z grazing **CASE 3 high P growth** low Z grazing







50 to 150 µatm over 30 days







Observed drawdown of 50 to 150 µatm over 30 days





- By just optimizing our biological model to Chl & NO₃ observations, we were not able to capture the bloom related pCO₂ drawdown
- TAKE-HOME POINTS

- By **including pCO**₂ observations into our model analysis, we believe we can better **constrain the dynamics driving the spring bloom** on the Scotian Shelf
 - **Predator-prey dynamics** play a key role in both the **magnitude**, and **timing and duration** of the spring bloom
- More work needs to be done towards implementing these predator-prey interactions in our models.

THANK YOU!



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