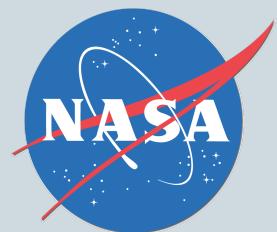
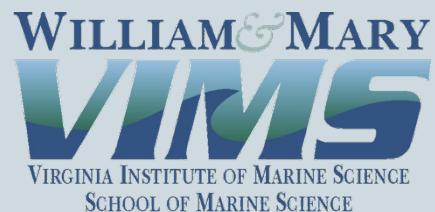


Model Study of Phytoplankton Phenology on the Northeast U.S. Continental Shelf



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Marjorie Friedrichs

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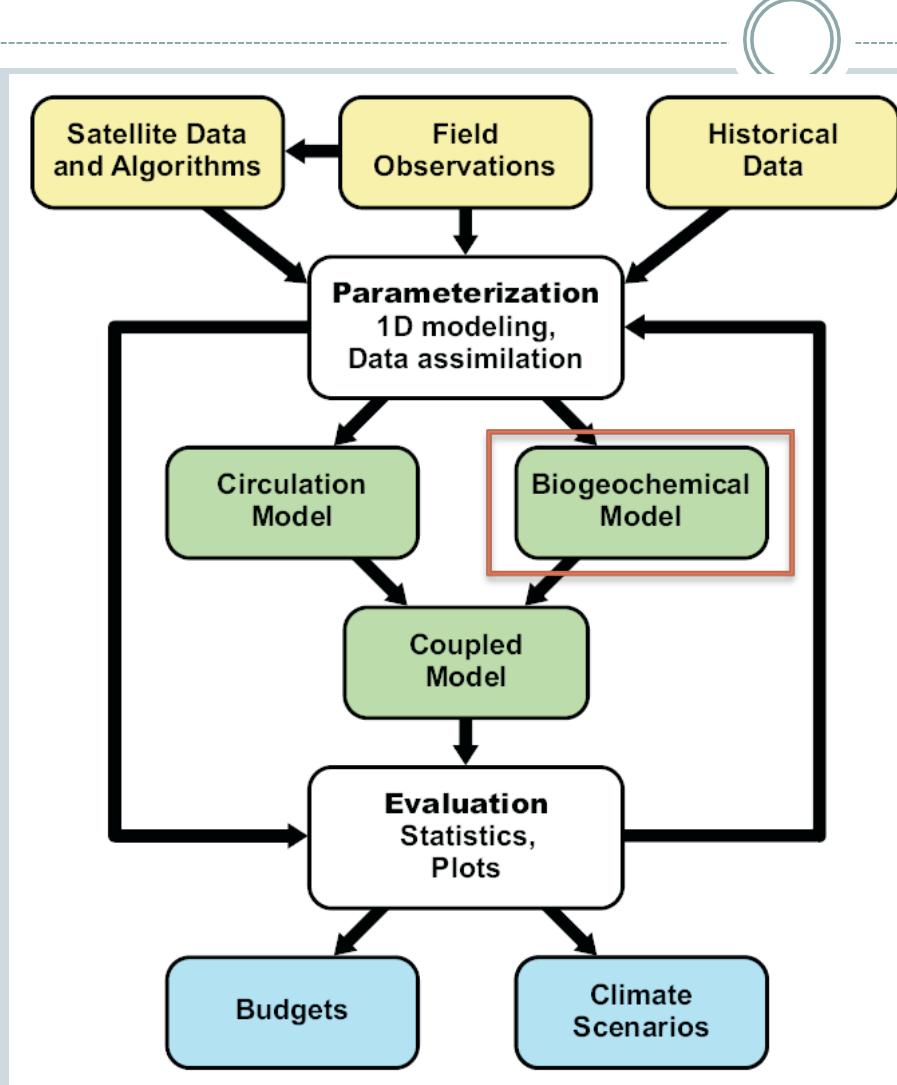


Outline



- Introduction
- Model refinement
- Results
 - Sensitivity analysis
 - Effects of model refinement on bloom timing
 - More realism/complexity
- Future work
- Summary

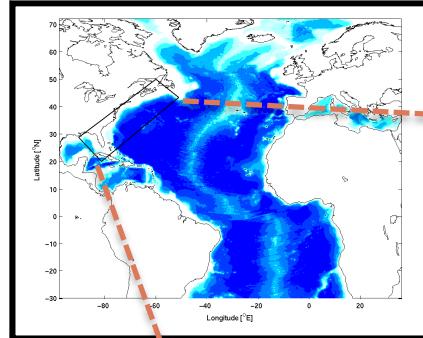
NASA U.S.ECoS Project



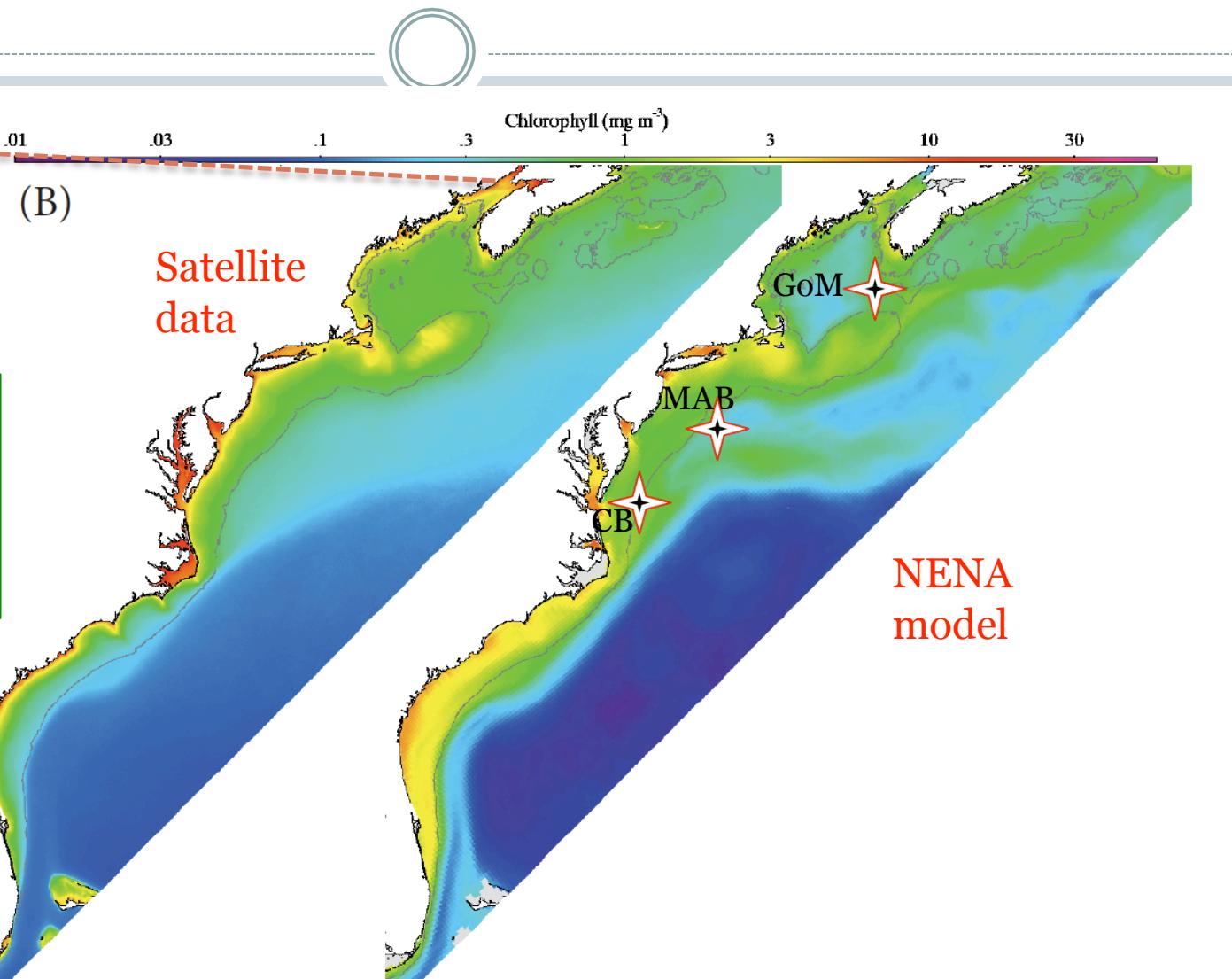
**U.S. Eastern Continental Shelf Carbon Budget:
Modeling, Data Assimilation, and Analysis**

Goal:
To develop carbon budgets for the U.S. East coast continental shelf

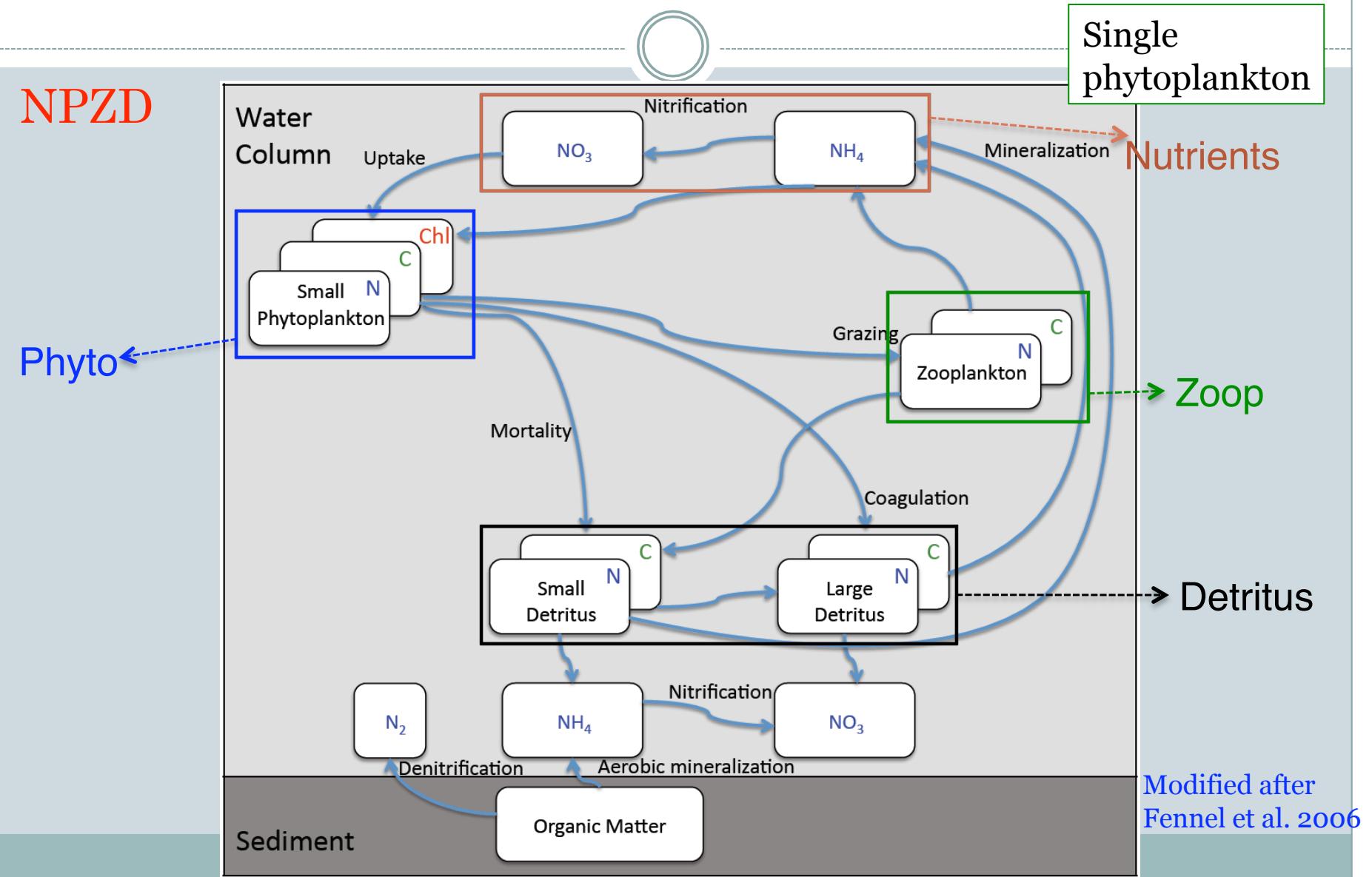
3D NENA simulation - Chlorophyll



North Eastern
North American
(NENA)
continental shelf

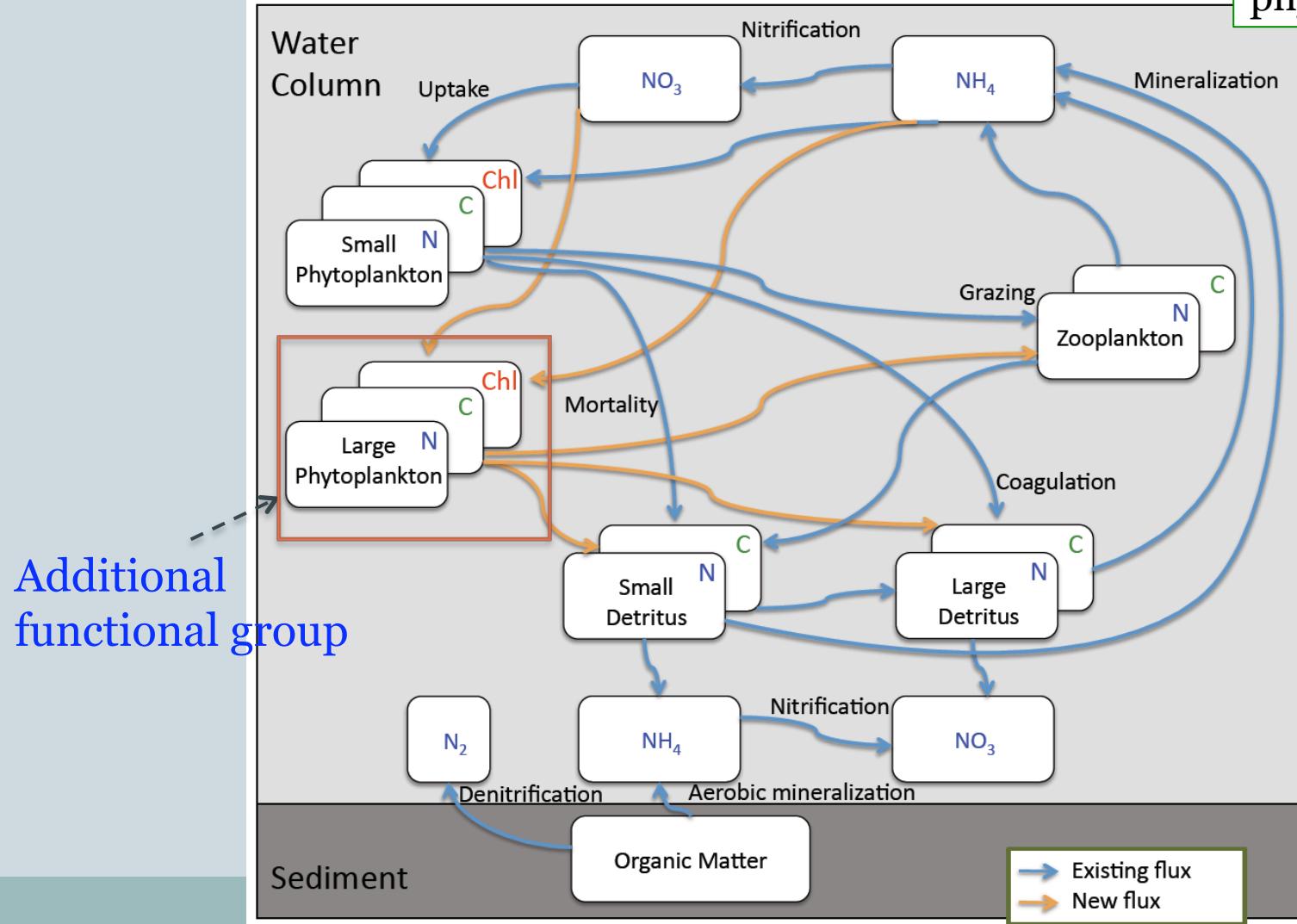


Ecosystem model



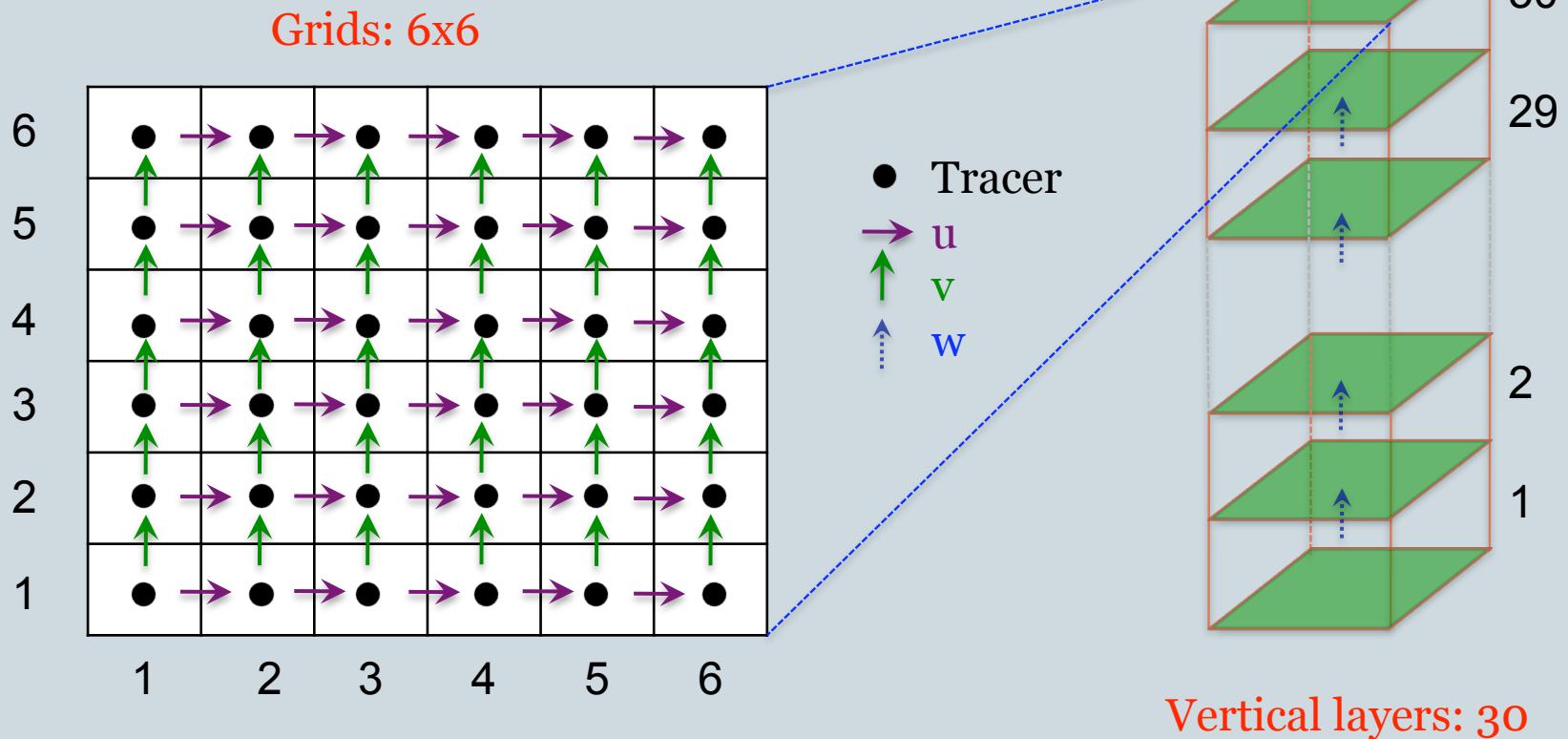
Model refinement

Two phytoplankton



ROMS bio-toy grids

- Circulation model: 1D ROMS Bio-toy



1D model set up



- Physics: vertical advection + diffusion
- Forcing: NCEP Daily North American Regional Reanalysis 2004
 - Wind
 - Air temperature
 - Radiation
 - ...
- Initial condition: 3D NENA Year 2003 simulation



Parameter sensitivity analysis



Normalized sensitivity $S_c(k)$:

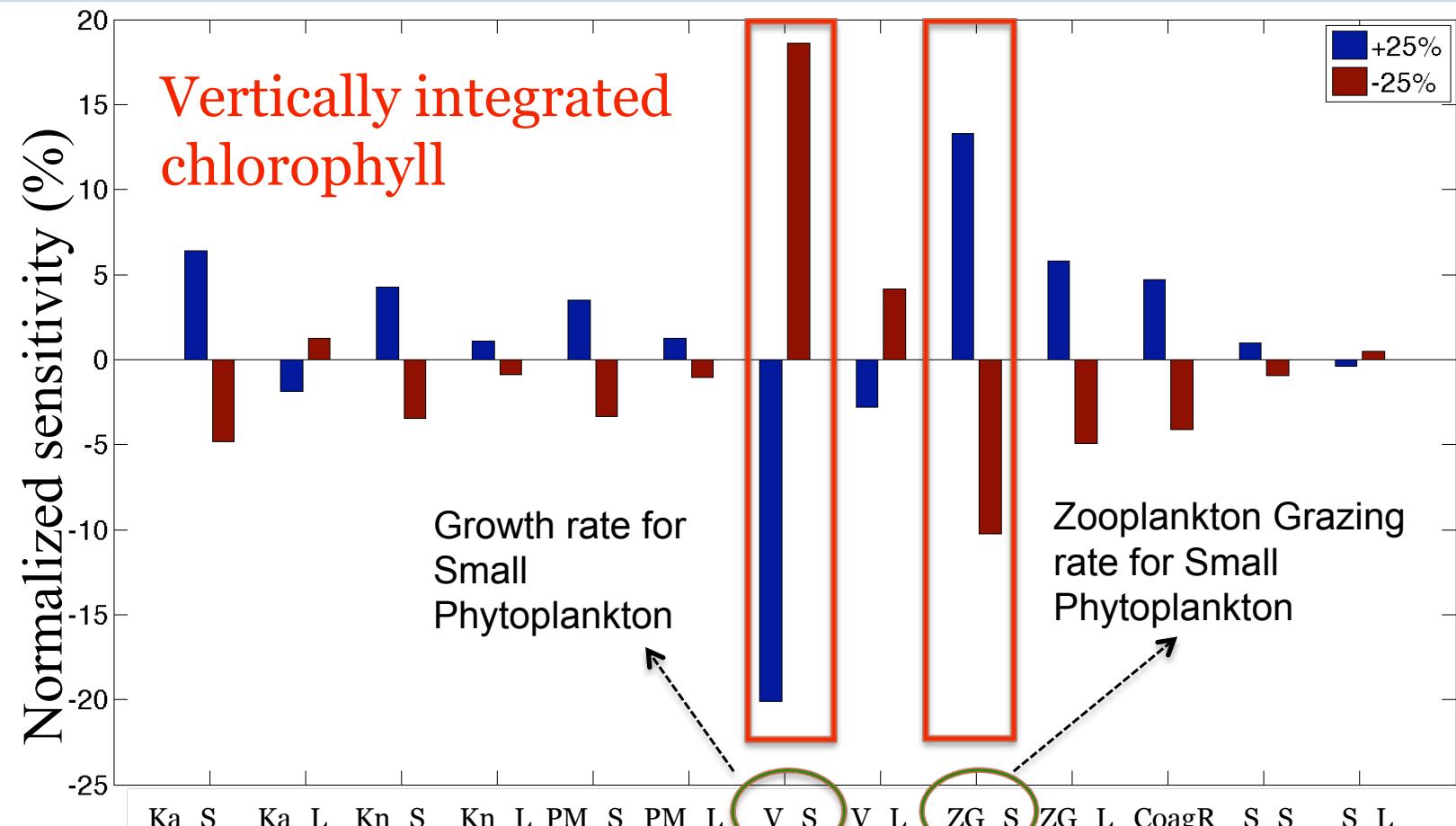
$$S_c(k) = \frac{\frac{C(k) - C_s}{C_s}}{\frac{k - k_s}{k_s}}$$

Change in Chl
(vertically integrated)

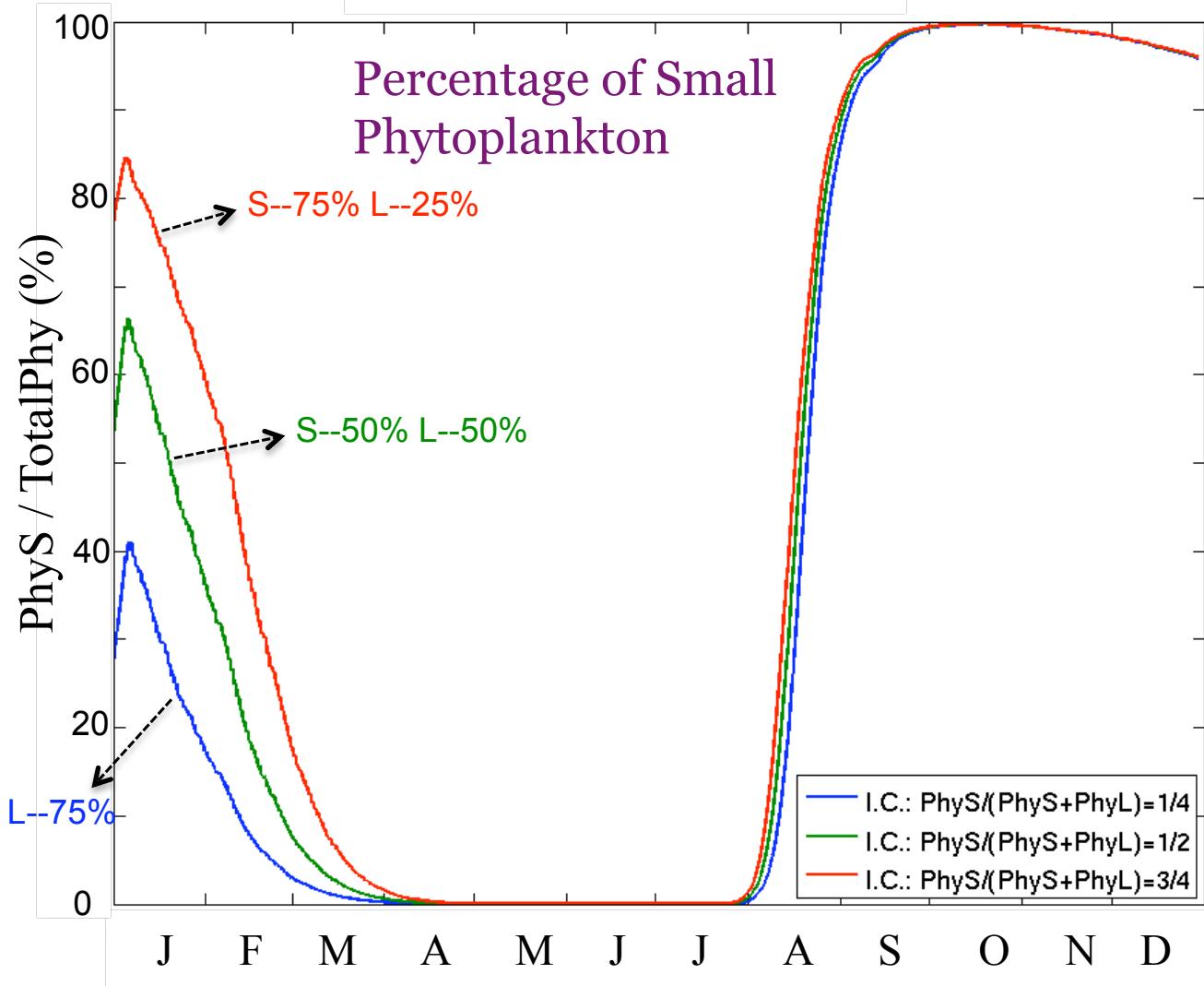
Change in parameter by 25%

- k : parameter value
- k_s : reference parameter value
- C_s : Chl concentration for the reference case with k_s
- $C(k)$: Chl concentration when using k

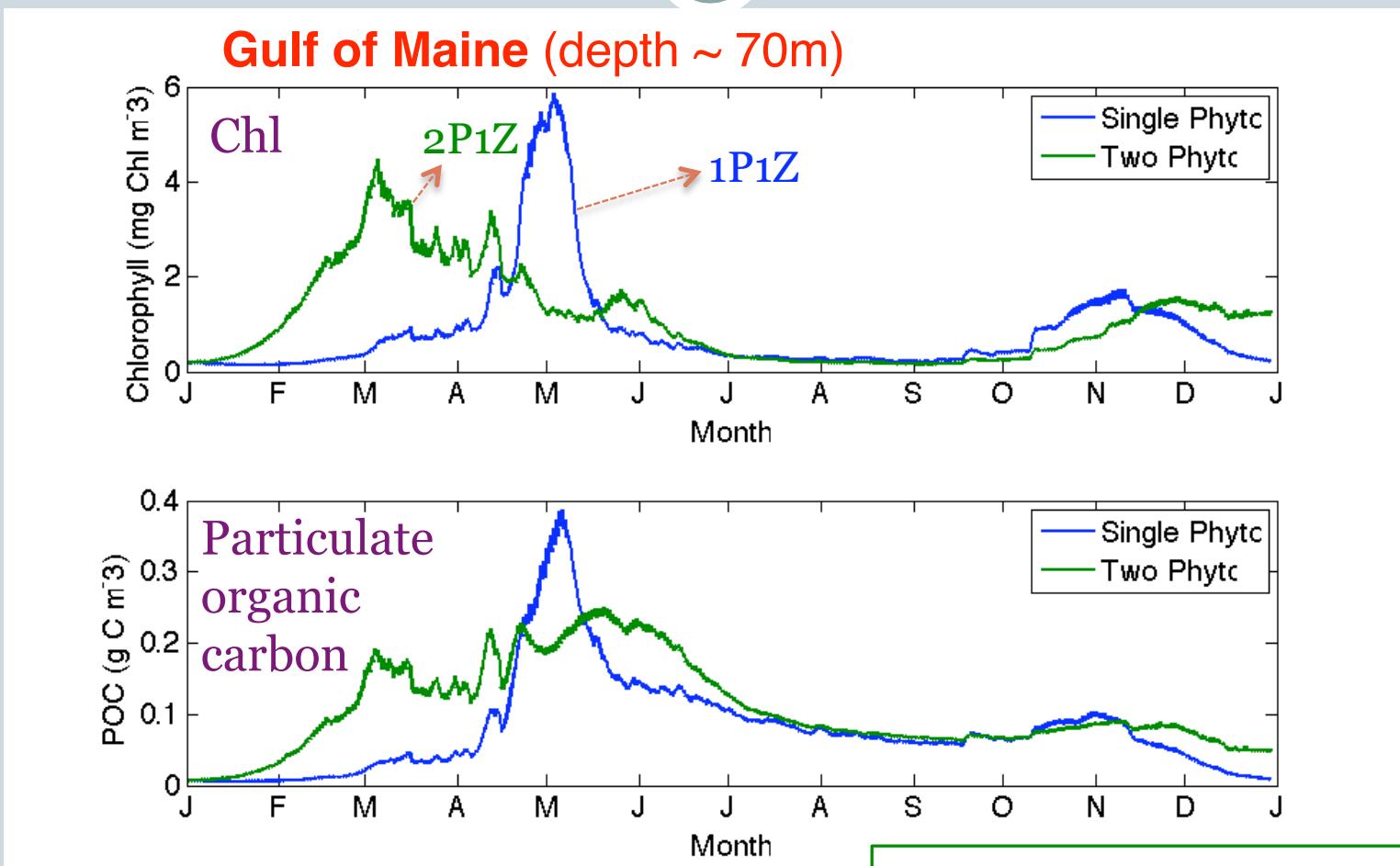
Parameter sensitivity analysis



Sensitivity to initial phytoplankton size fraction



Single vs. two phytoplankton – site 1

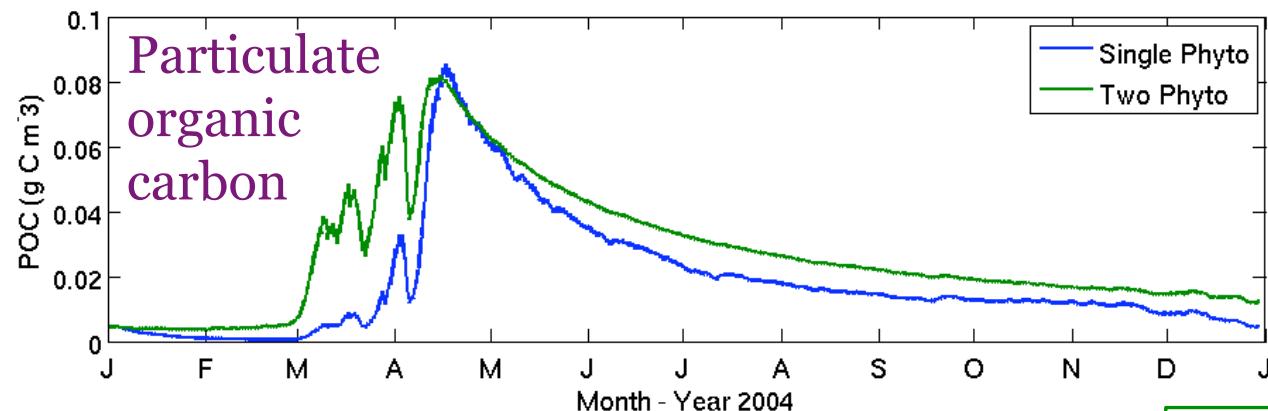
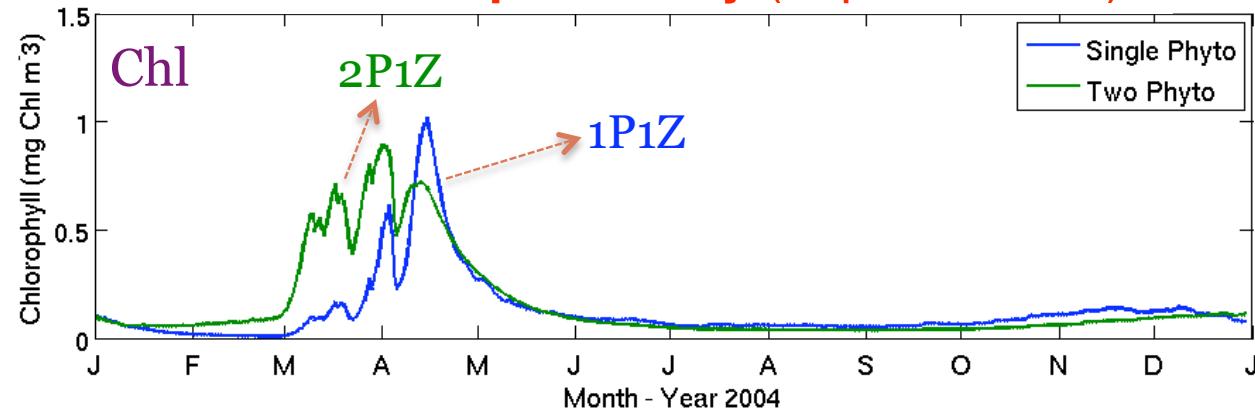


Change in timing + magnitude

Single vs. two phytoplankton – site 2



Outside Chesapeake Bay (depth ~ 460m)

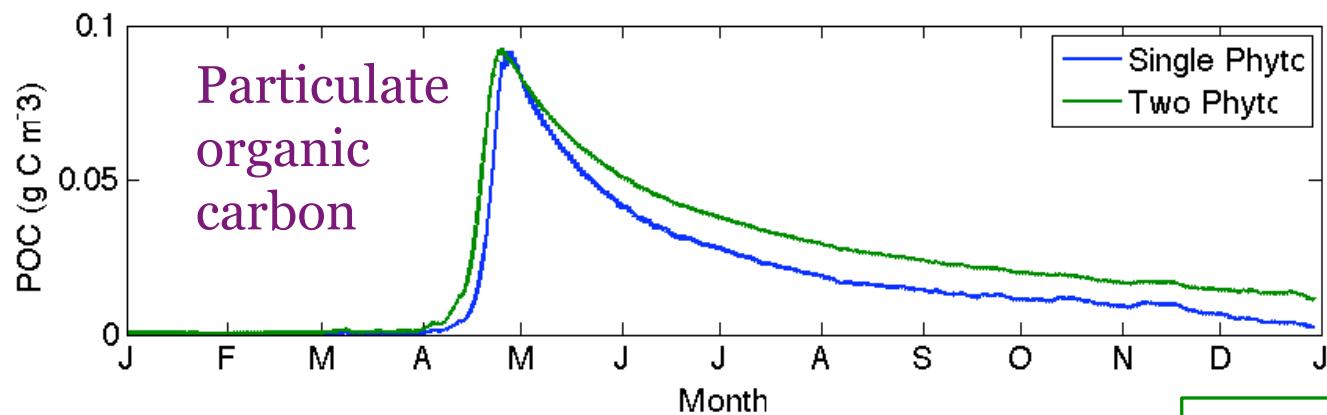
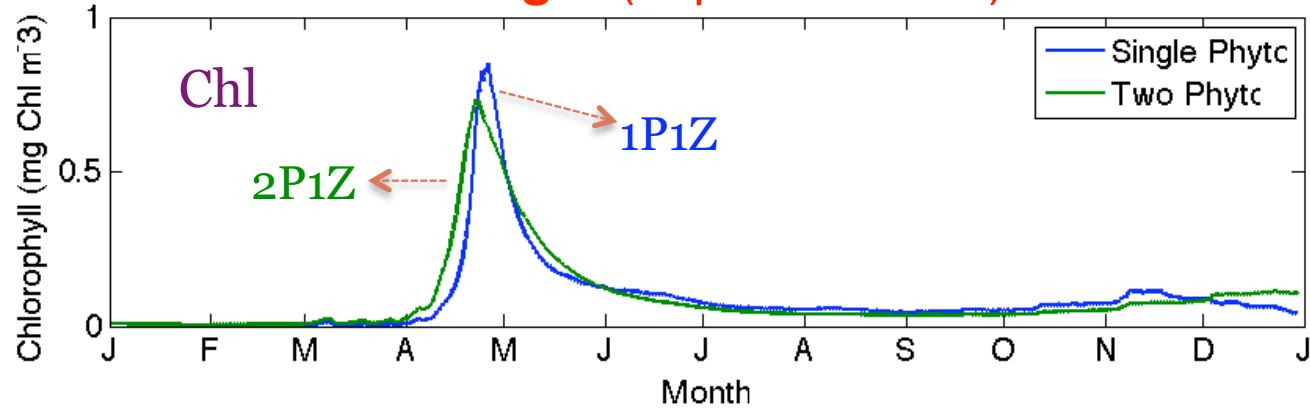


Change in timing

Single vs. two phytoplankton –site 3

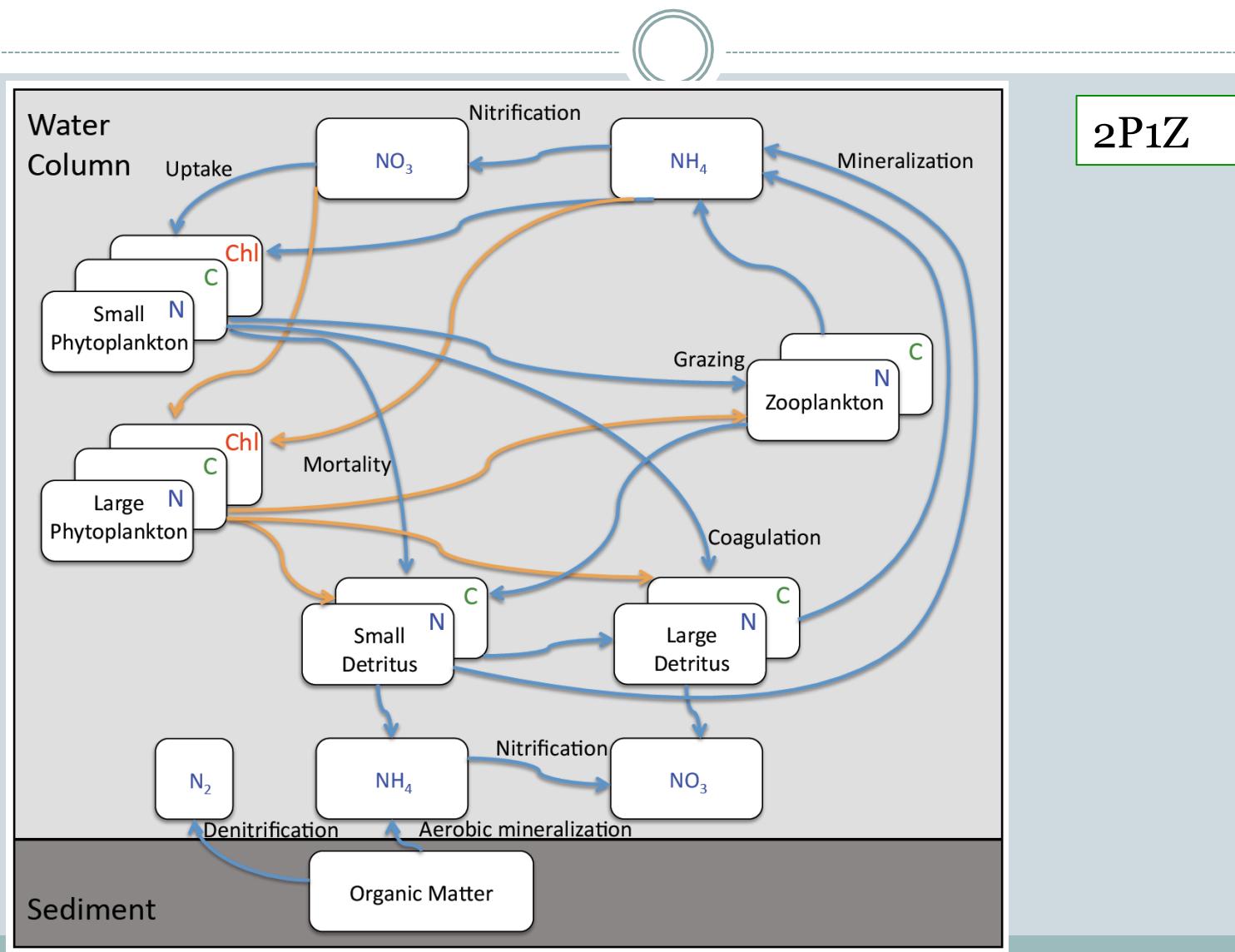


Mid-Atlantic Bight (depth ~ 2400m)

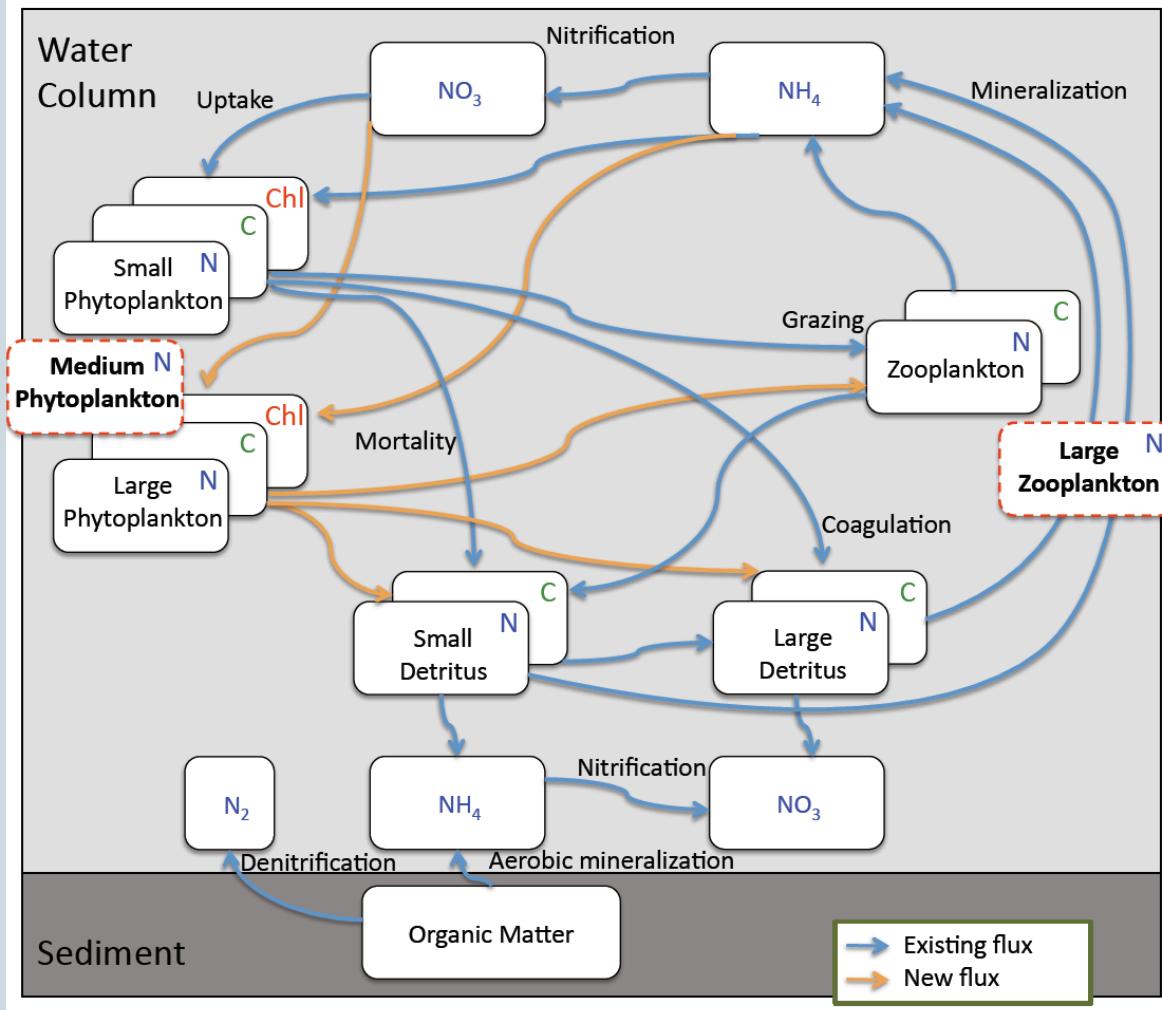


Minor change

Two phytoplankton model: complex enough?

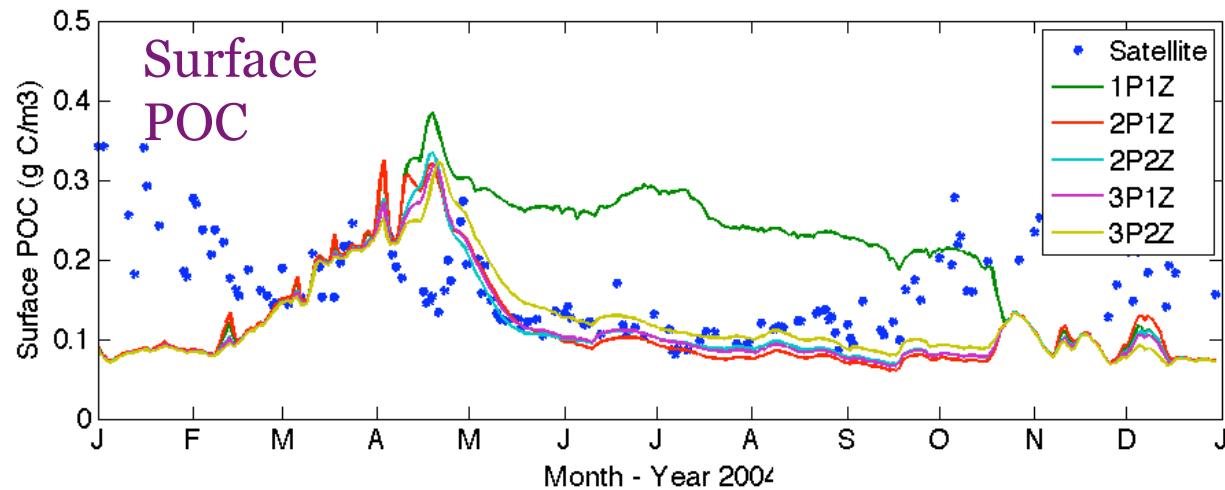
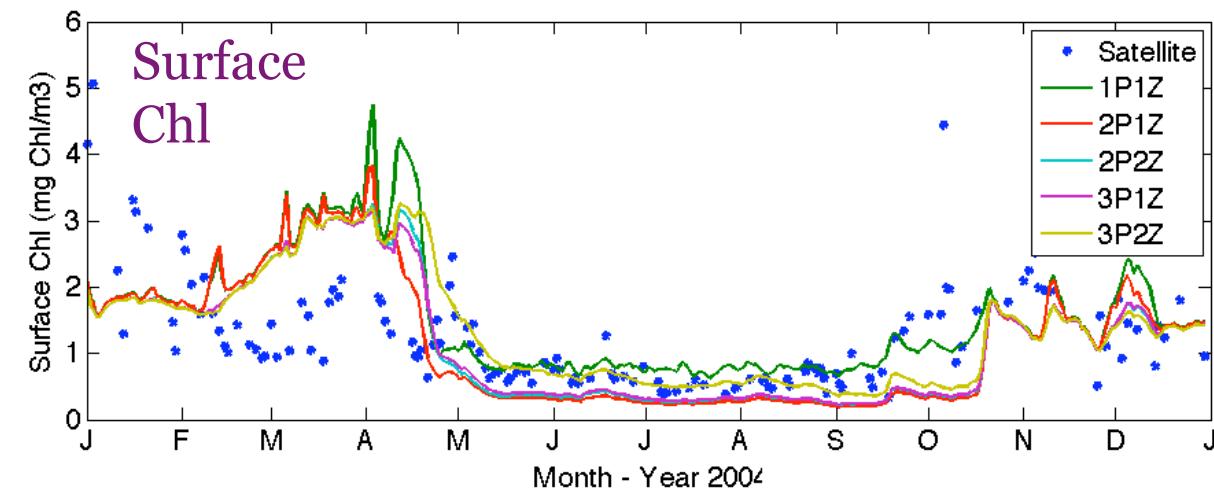


More plankton?

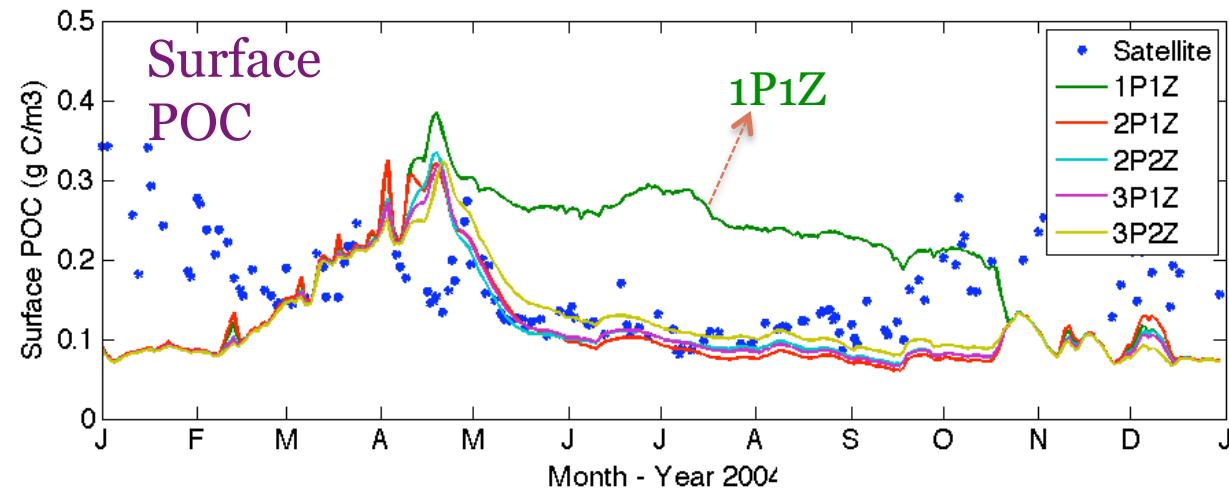
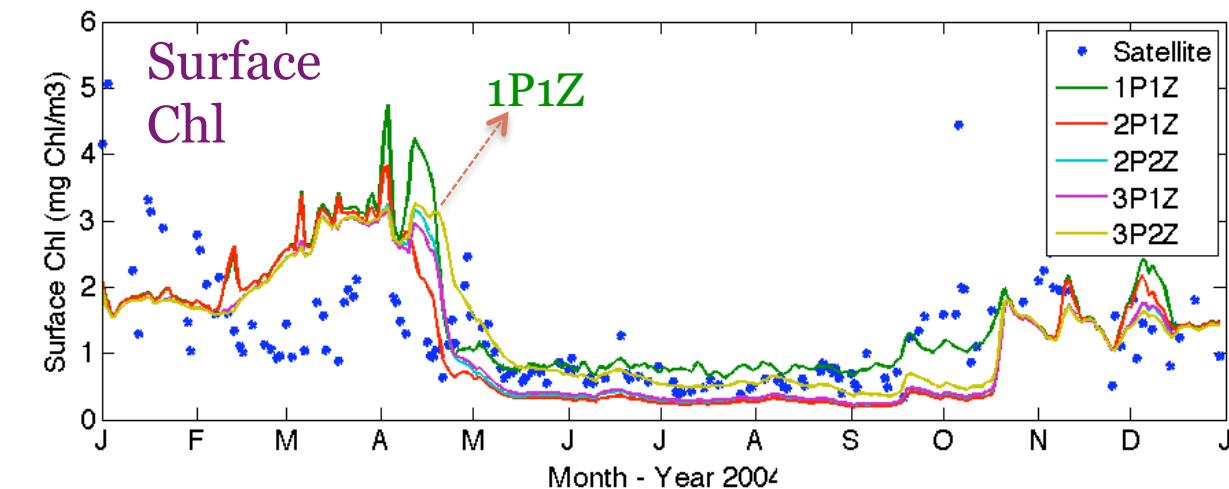


- 1P1Z
- 2P1Z
- 2P2Z
- 3P1Z
- 3P2Z

More comparisons



More plankton? Not necessarily better

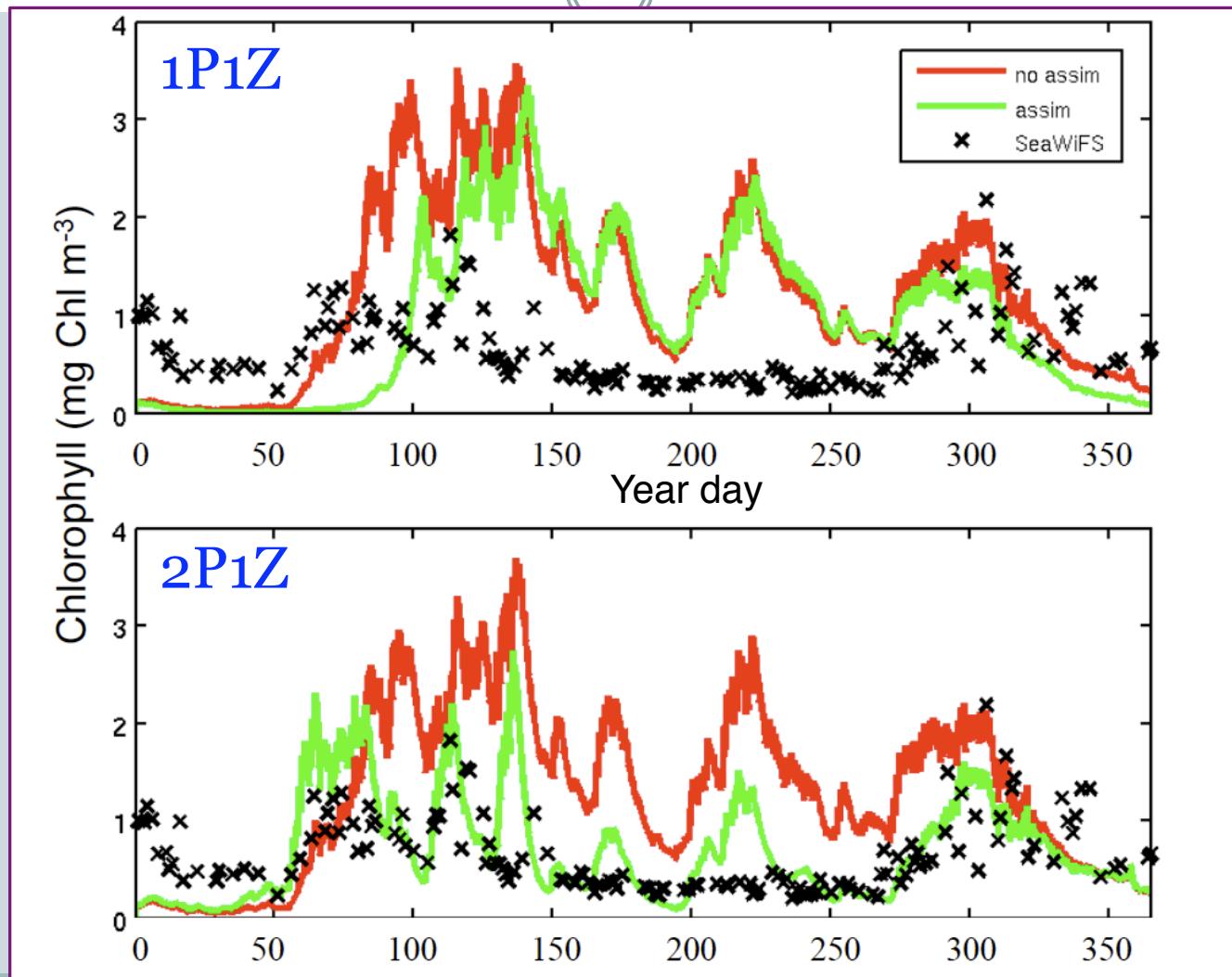


Future work: data assimilation



- Variational adjoint method
- Method: nonlinear weighted least square
- Data:
 - Satellite Chl+POC
 - in situ data
- Optimized parameters:
 - Growth rates
 - Grazing rates

Preliminary result - data assimilation



Summary



1. Phytoplankton community structure can significantly impact model estimate of bloom.
2. The impact is larger in regions where chlorophyll is high.
3. Preliminary results indicate that adding more plankton beyond 2P1Z does not improve the model-data comparison for surface chlorophyll or surface POC.



Thanks for your attention!