







Modeling BGC in the Ice Interior: the CICE release and beyond

Nicole Jeffery, Scott Elliott, and Elizabeth Hunke

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Boulder, CO.





Antarctic







Does sea ice physics largely determine which type of algal bloom occurs? And when? And how?

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This Talk

- How recent CICE developments help us model biogeochemistry throughout sea ice:
 - a) Temperature, Salinity, bottom biogeochemistry
 - b) Vertical tracer transport modeling
- Compare and contrast bottom bgc and zbgc
 - a) Flux and transport assumptions
 - b) Does it matter?
- Conclusions

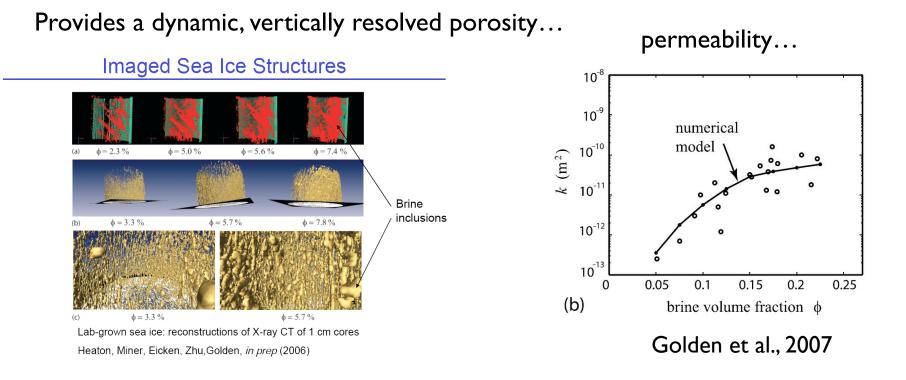




Recent CICE developments (I)

Thermodynamics: mushy layer in addition to B&L99

Salinity: mushy layer and scheme for use with B&L99 2)



and critical information for parameterizing brine motion.

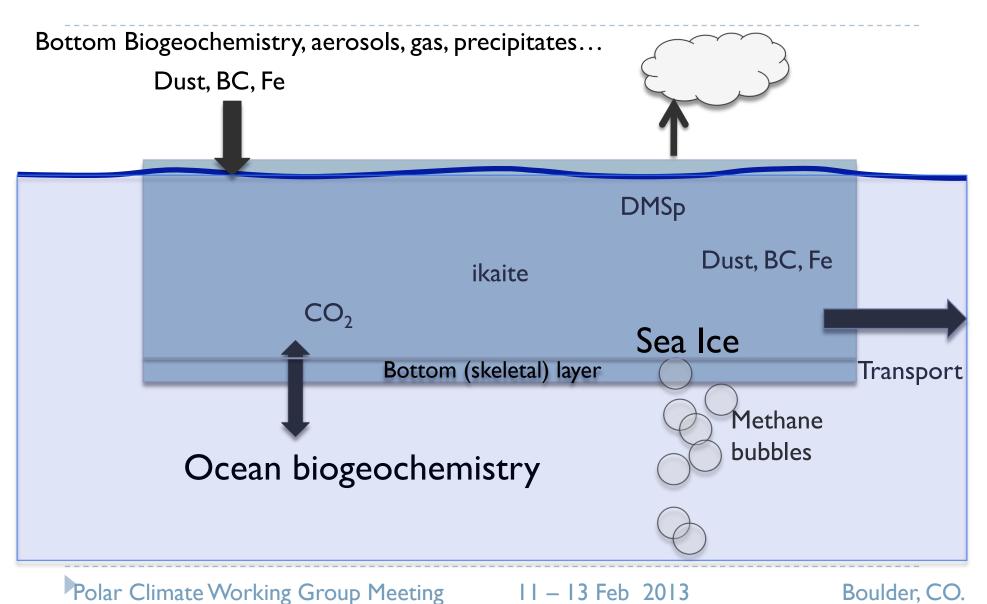
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CICE developments and current work (II)

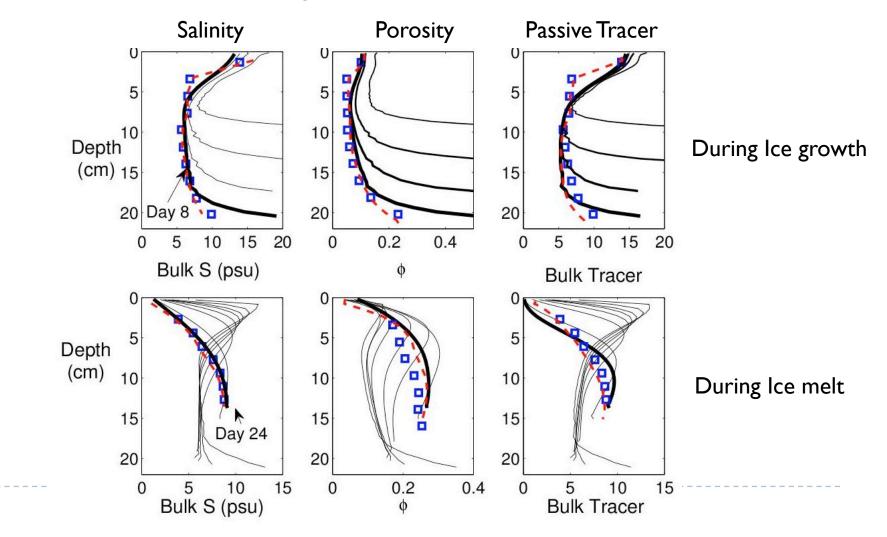






Recent CICE development (III)

Vertical tracer transport







Modeling ice biochemistry

If we assume that physical properties and processes determine the when/where of an algal bloom...

Some important processes:

- 1) Flushing (surface and snow melt, meltponds)
- 2) Gravity drainage (modulate ocean-ice fluxes during ice growth)
- 3) Snow loading (high snow thickness/ice thickness ratio)

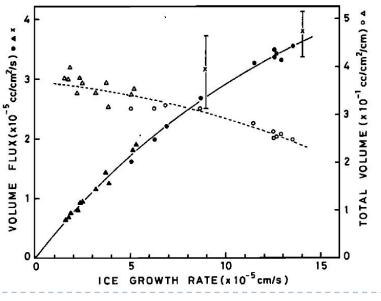
For the Arctic, *any* model of ice bgc needs a representation of the first two processes.



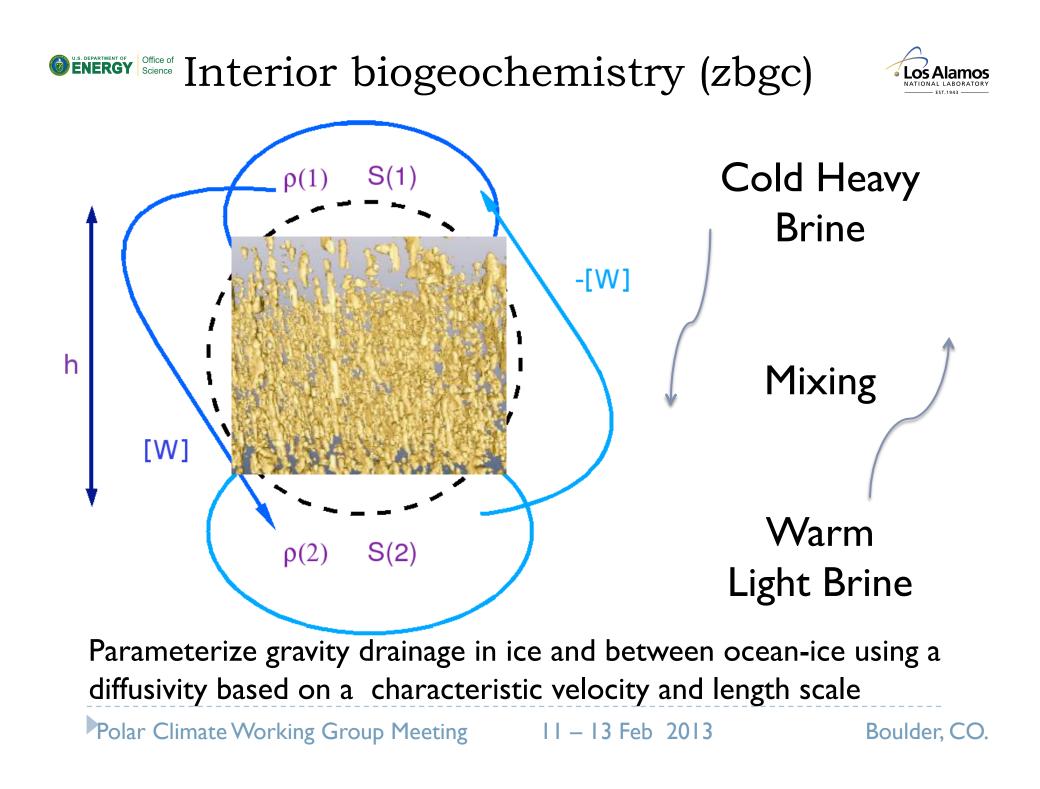


Bottom (skeletal) Ice Algae - Assumptions

- Skeletal layer thickness and porosity
- Compounds (nitrate, silicate...) are advected between the ice/ocean via a piston velocity (Vp)
- > When ice growth dominates, Vp is a function of the growth rate after brine flux measurements of Wakatsuchi&Ono
 - multiplied by a tuning
 - parameter.
- > When ice melt dominates, Vp is a function of melt.



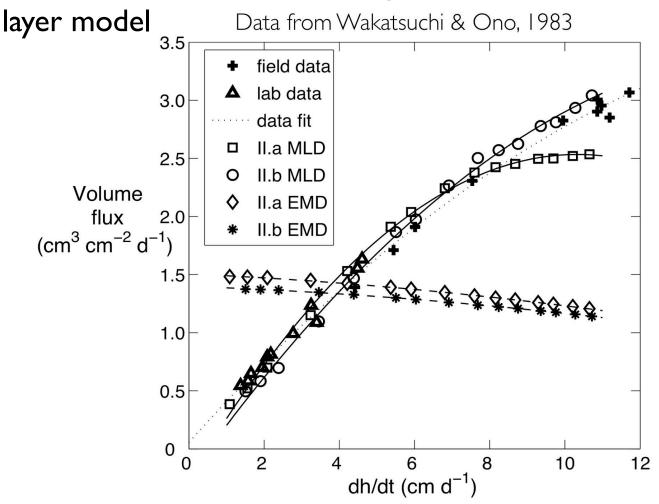
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Consistent with Piston velocity formulation of the skeletal



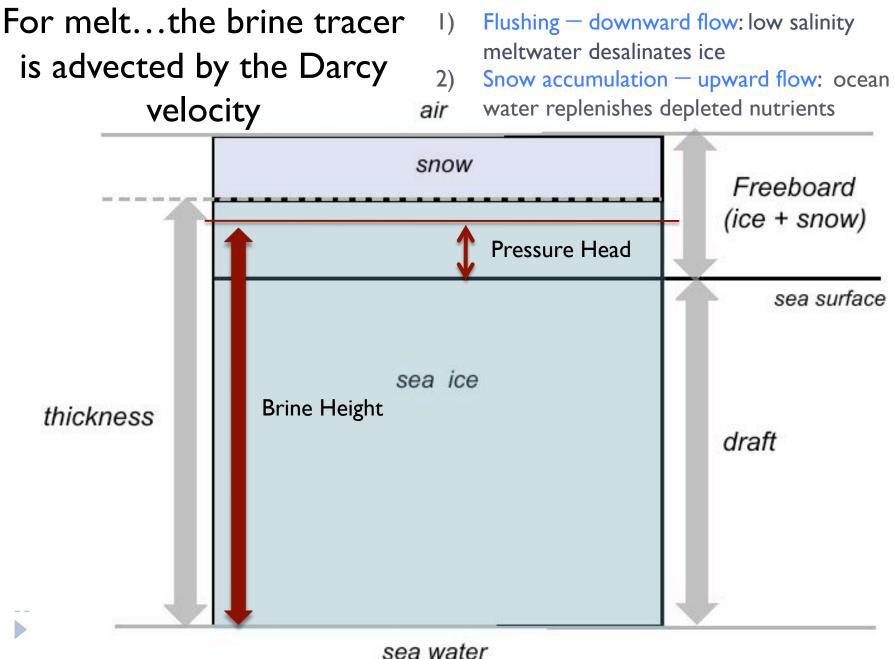
A tuning parameter (unknown length scale) is still needed, but it can be determined independently of the biochemistry using the salinity contours.

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Does it matter?

Tracers : Algal nitrogen and Nitrate

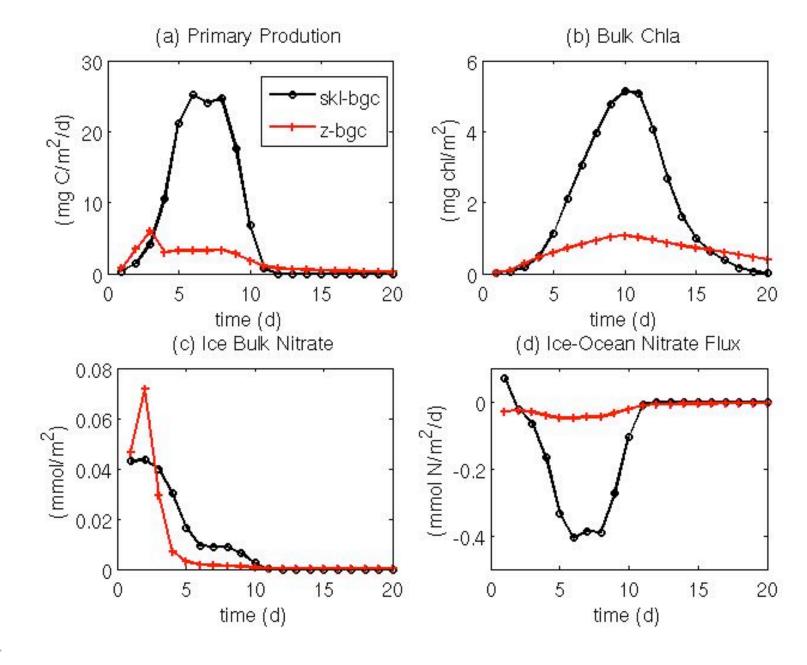
Column Test Case: 25 days of ice growth and melt with light

- Light and Nitrate can limit growth
- Skeletal biochemistry and Z biochemistry are identical
- Dynamic salinity in both cases
- Ice growth and melt are identical

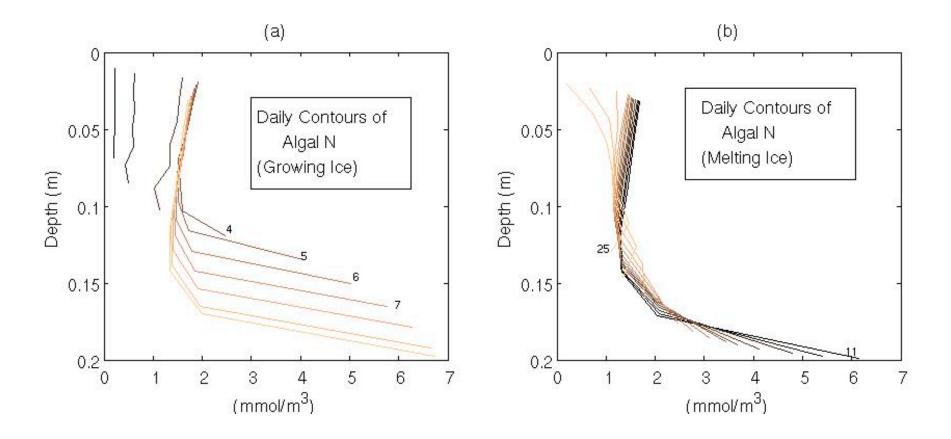
Differences: Ice-Ocean and Ice bottom-ice interior fluxes

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Z-bgc Model: Peak in Algal Nitrogen in the ice bottom



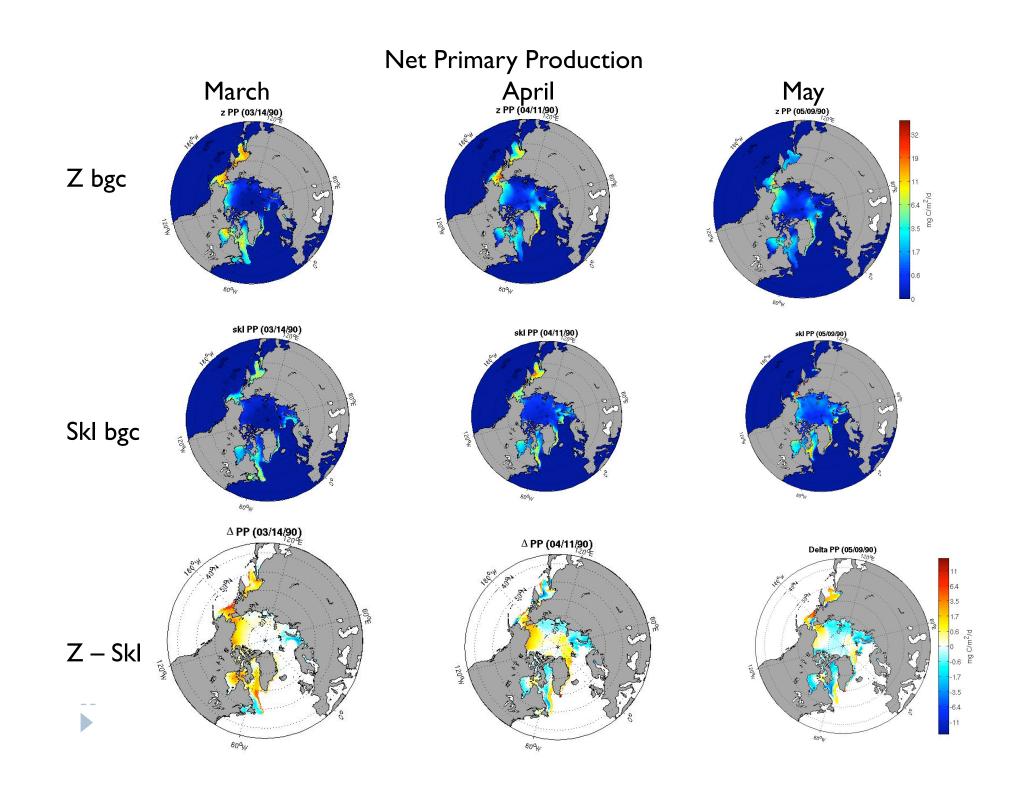


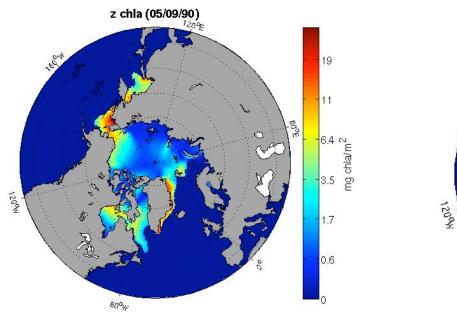


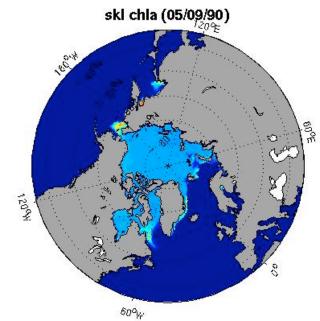
Does it matter? Arctic Scale

Tracers : Algal nitrogen and Nitrate

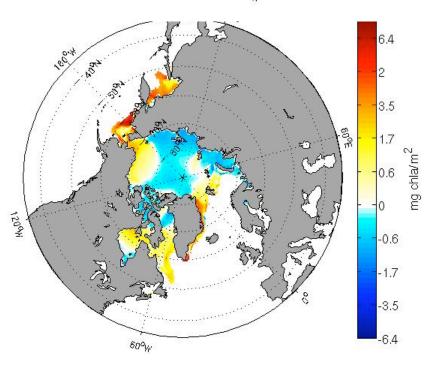
- Simulations from a 9 year Arctic spinup with salinity
- Biochemistry added Dec 1, 1990
- Nitrate uses Arctic ocean climatology
- Light and Nitrate can limit growth
- Skeletal biochemistry and Z biochemistry are identical
- No additional salinity, light, iron or temperature Inhibition.







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Summary

- Tracer transport portion of zbgc scheduled for release but not with biochemistry
- Works with any salinity model or prescription, but works better with more realistic salinity profiles.
- Representation of ice-ocean fluxes can radically change algal production estimates.
- In zbgc, the gravity drainage strength is tuned independently from the biochemistry. Could help constrain the fluxes in bottom bgc.