



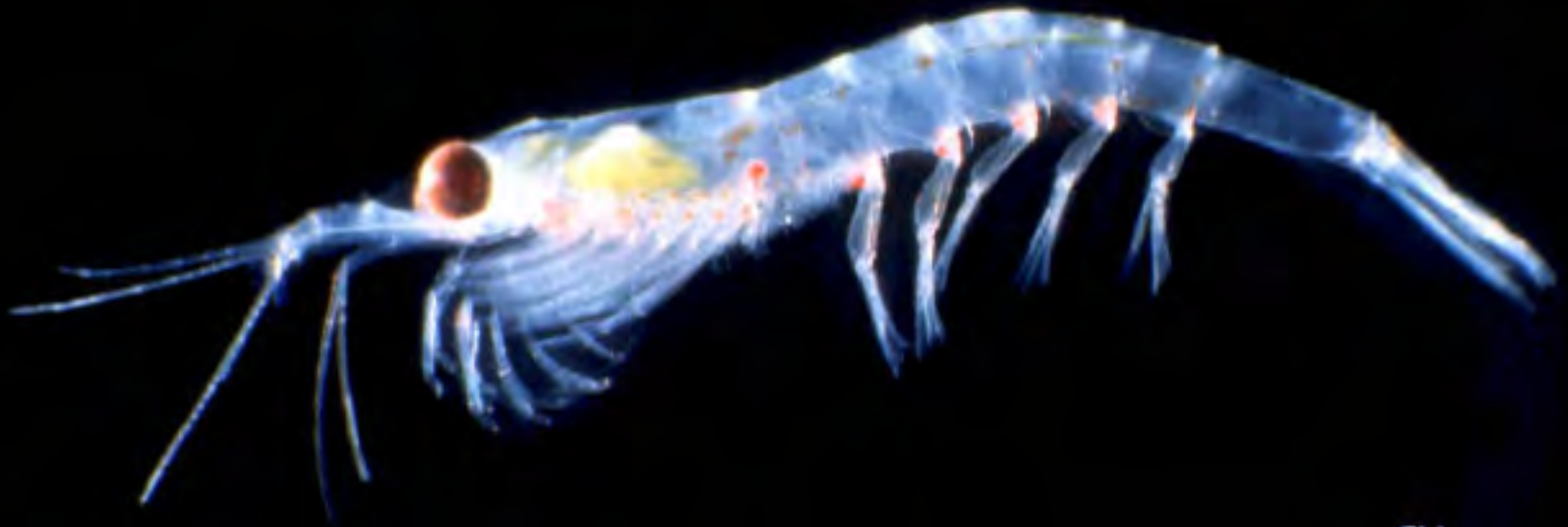
# On Adding a Stage-Structured Model of Krill to NEMURO



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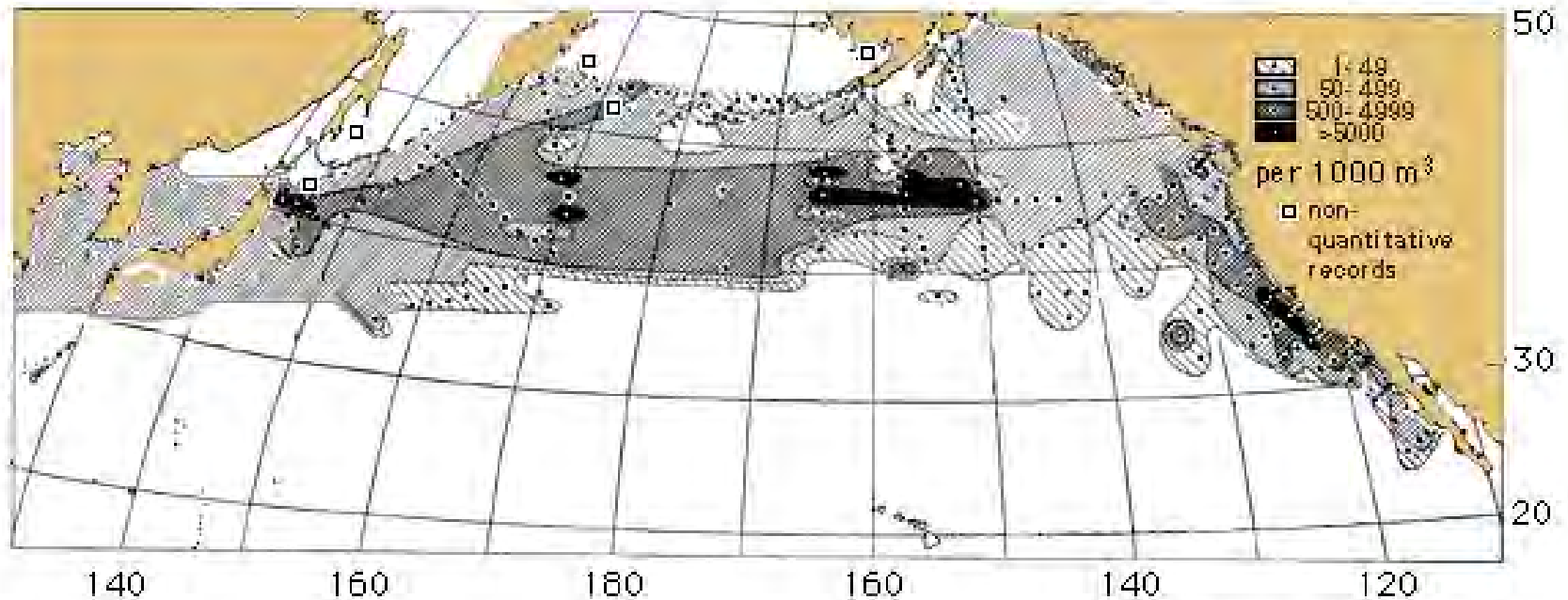
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## *Euphausia pacifica*



- Distributed around Pacific Rim and across open ocean
- Occupies a broad variety of habitats from cold subarctic waters to warm subtropical waters.
- A key species in food chains as grazer and prey for commercially-important fishes, as well as many birds and mammal species.
- How might climate change affect distribution, population dynamics and production of this species in different regions of the Pacific?

PICES FUTURE is interested in forecasts.

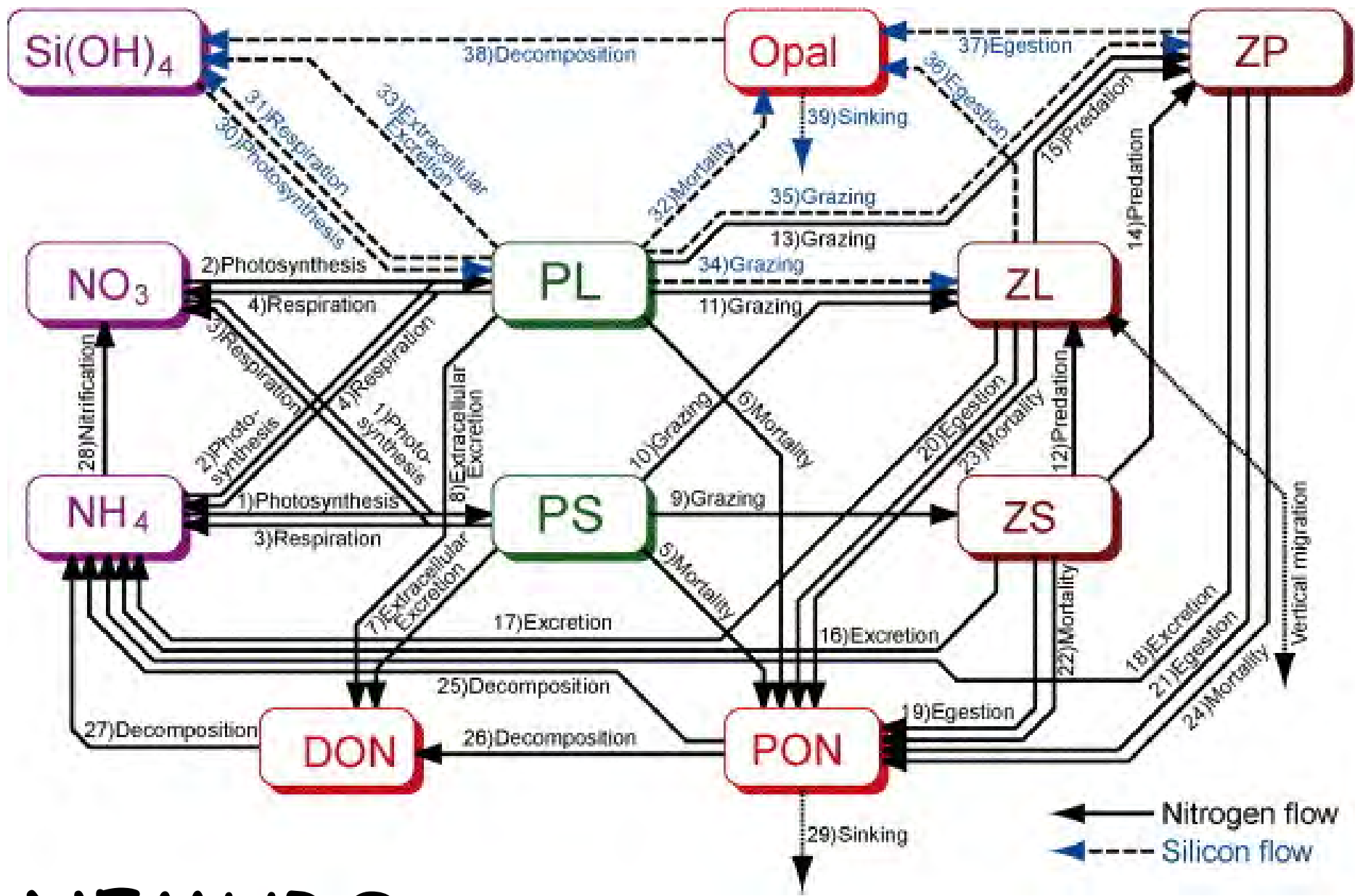
Wolfgang Fennel summarized models in the HAB topic session yesterday.

- 1) Forecasts means models.
- 2) "Start with simple toy models. Then move to full 3D models."

The model shown here is a toy model (0-D).

**Continuous Distributions or IBMs?**

Depends on the question.



# NEMURO



## SOME BIOMASS

**Vitals:**  
**320 lbs, 7'6";**





**Vitals:**  
**320 lbs, 7'6";**

**MORE BIOMASS**



**Vitals:**  
**~320 lbs,**



**Vitals:**  
**320 lbs, 7'6";**

**?**  
**==**



**Vitals:**  
**~320 lbs,**





**Vitals:**  
**320 lbs, 7'6";**



Smaller Organisms have

- 1) Higher weight specific rates (feeding, respiration, growth)
- 2) Different allocation to growth, maintenance and reproduction.
- 3) Consume different types of prey
- 4) Different active behaviors (migrations and movements)

than **Larger Organisms**

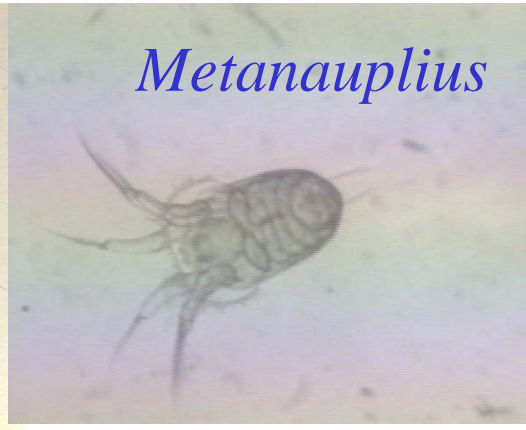
**Individual Size Matters!**



# *Euphausia pacifica* life stages



*N2*



*Metanauplius*



*Calyptopi*

*Adult*



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# In Zooplankton, Individual Size

- Impacts preferred prey type (abundance/size)
- Impacts growth rate
- Impacts size-dependent mortality
- Impacts behavior
- Impacts internal pools (lipid reserves)

Stage-specific CW

*Euphausia pacifica* life stages

~7  $\mu\text{g ind}^{-1}$

N2

*Metanauplius*

571 indiv.

$\Sigma R=529.2 \text{ ug C d}^{-1}$

$\Sigma G=519.6 \text{ ug C d}^{-1}$

R/G=1

Assim/Wt=0.26

~3.2  $\mu\text{g ind}^{-1}$

1250 indiv.

$\Sigma R=633.6 \text{ ug C d}^{-1}$

$\Sigma G=425 \text{ ug C d}^{-1}$  R/G=1.5

Assim/Wt=0.26

$\Sigma R=122.9 \text{ ug C d}^{-1}$

$\Sigma G=26 \text{ ug C d}^{-1}$  R/G=4.7

Assim/Wt=0.04

~4000  $\mu\text{g ind}^{-1}$

1 indiv.

9b

## Advantages of IBMs

- 1) Biology is often **mechanistically explicit**. (not hidden in differential equations).
- 2) Biological-Physical-Chemical Interactions are clearly detailed.
- 3) Individual is the fundamental biological unit, thus it is **natural and intuitive to model at that level**, rather than at the population level.
- 4) Allows explicit inclusion of an **individual's history and behavior**.
- 5) History-Spatial Heterogeneity interactions '*easily*' handled.



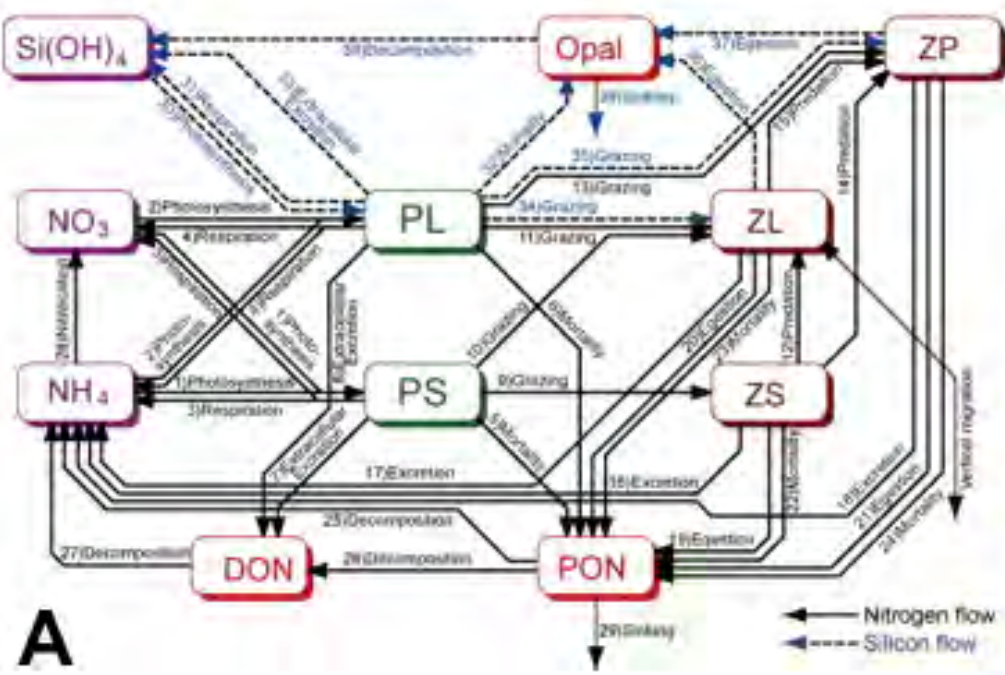
## Costs Involved in IBM Approach

- 1) Difficult to implement **feedback** from IBM (Lagrangian) to underlying Eulerian model, esp. across multiple trophic levels
- 2) Requirement for Large Numbers of Particles
  - Difficult to simulate **realistic abundances**
- 3) Difficult (Impossible?) to simulate **density dependence**
- 4) Extensive **Computation Penalty**

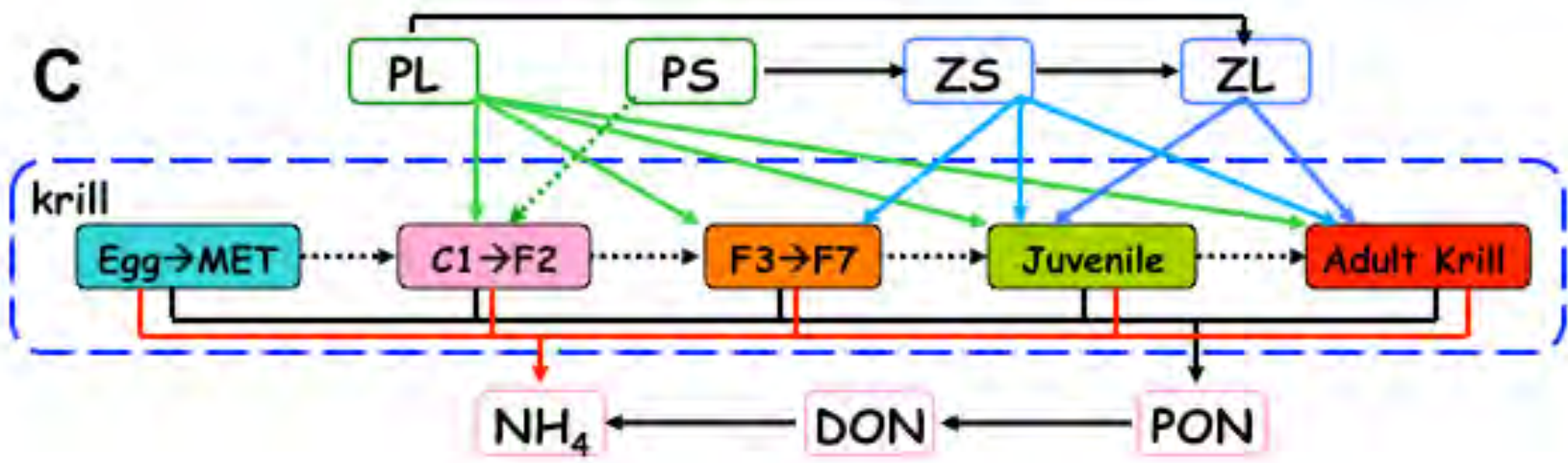
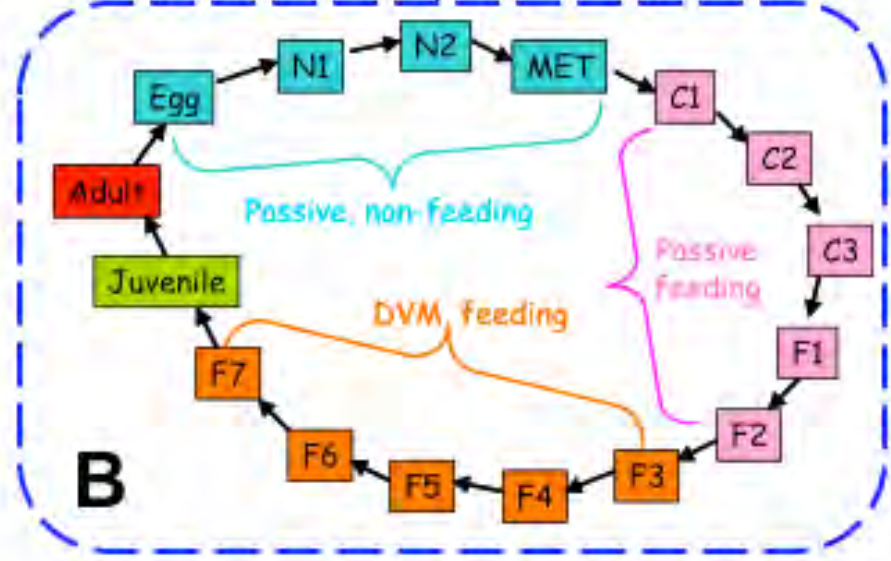
For many research questions, we can live with these problems (or work around them).

But, when considering future climate change scenarios and responses of ecosystem structure and function, it is difficult to ignore biomass density (concentration).

Consequently, the dominance of Eulerian concentration based models in climate scenario assessment.



More detailed life-stage groupings for krill (dashed blue outline), where lifestages are aggregated into 5 groups determined by DVM, feeding behaviors and reproduction.



# A Stage Progression Model

*E. pacifica* Belehradec function for time to stage as function of temperature

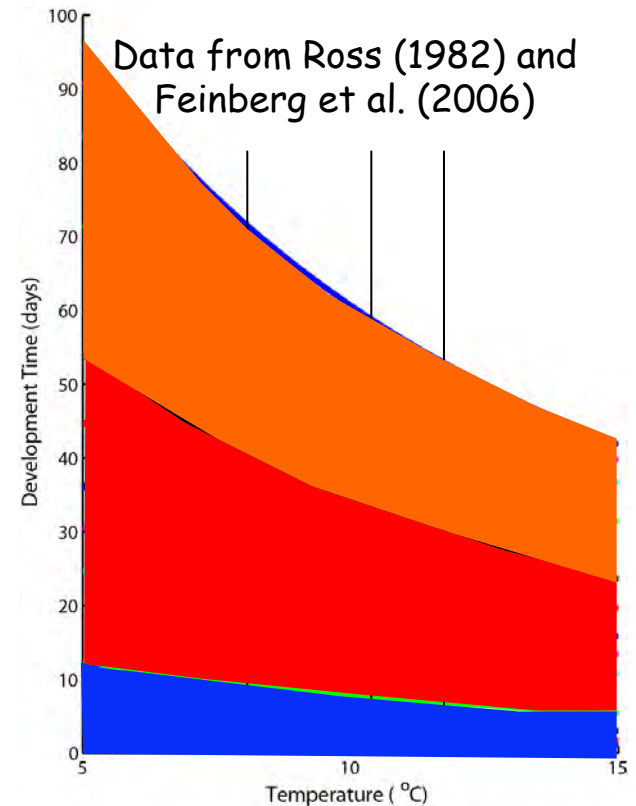
Basic Form is:  $D_i = a_i (T + b)^c$

$D_i$  is the time (days) from egg to stage  $i$

$a_i$  is a stage specific constant

$b$  is a stage-independent shift in temperature

$c$  is assumed to be -2.05 (commonly observed from experiments; determines the curvature)



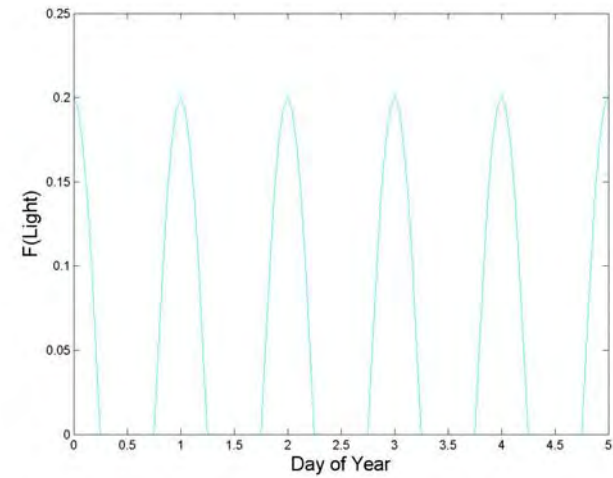
$MR_i = 1/[a_i (T + b)^c] \rightarrow$  Leads to numerical diffusion

$MR\text{-corr}_i$  based on Hu et al. (2008; MEPS) algorithm that uses mean age of state $_i$  and a probability density function of transfer.

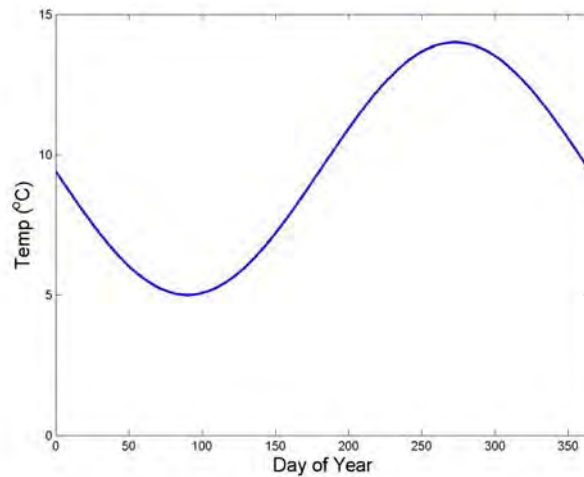


# External Forcing

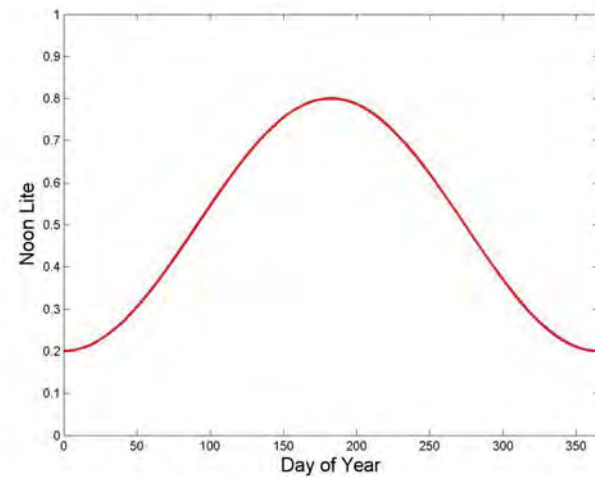
## Diurnal Light Cycle



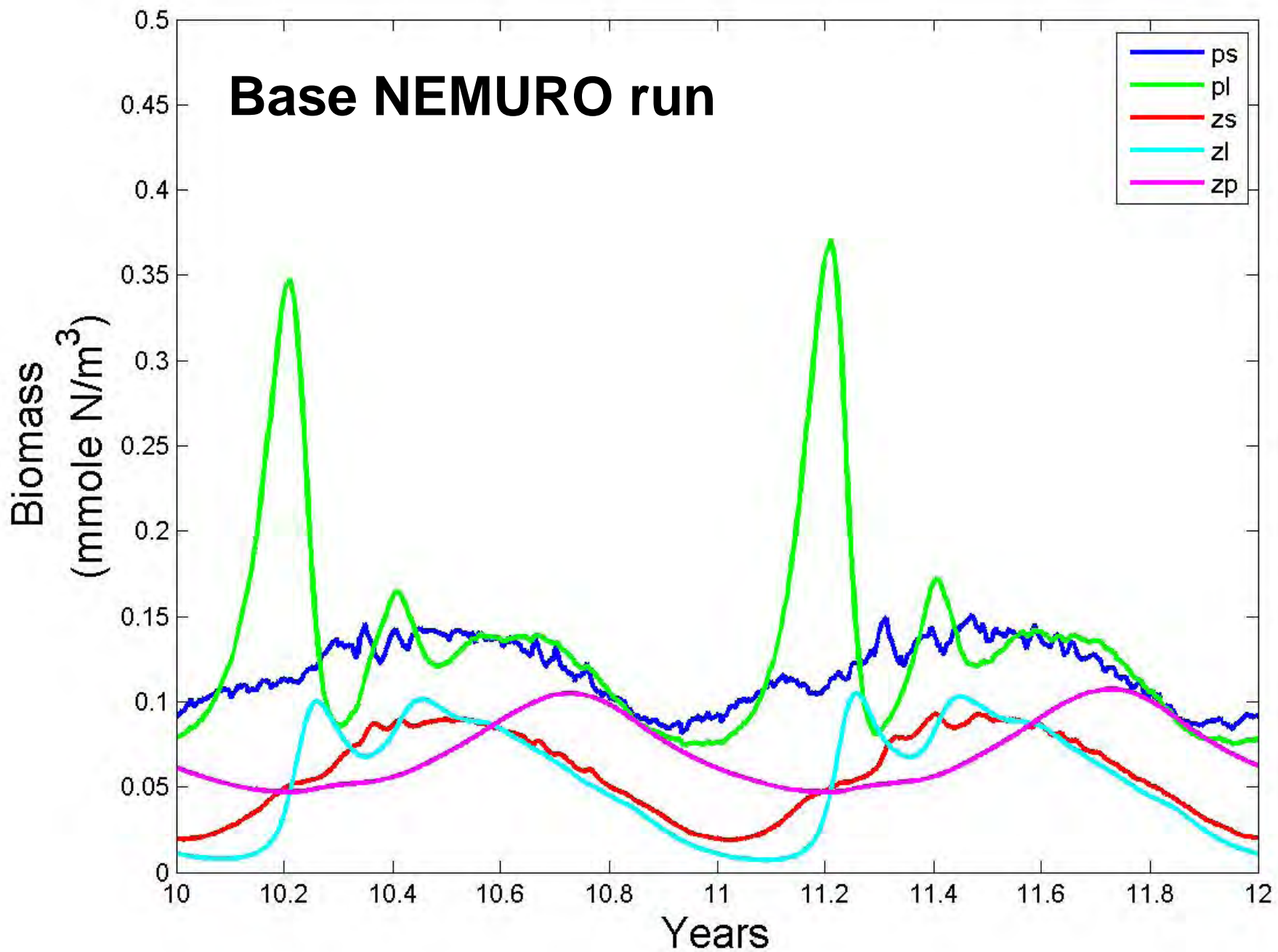
## Annual Temp Cycle



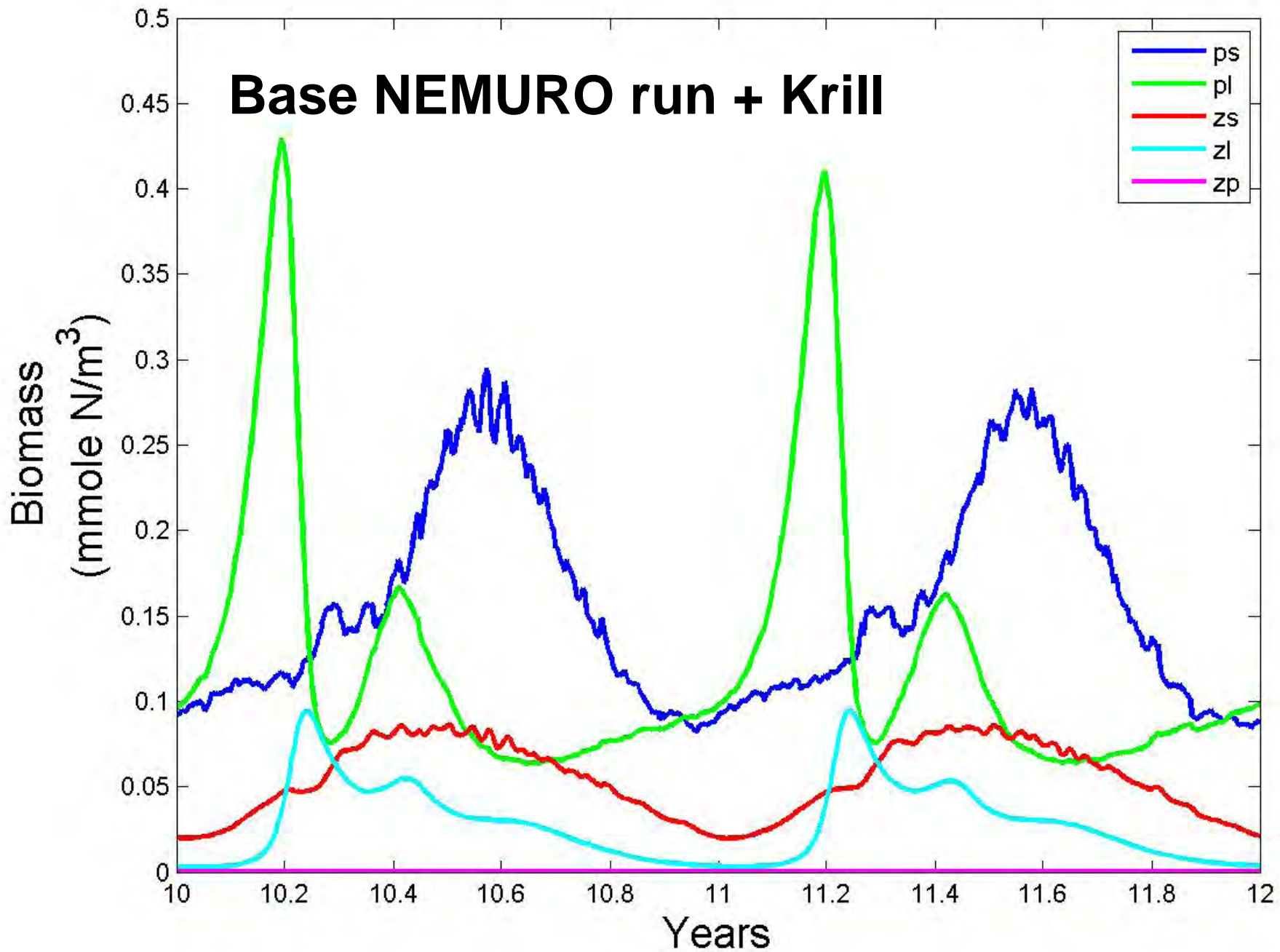
## Annual Light Cycle



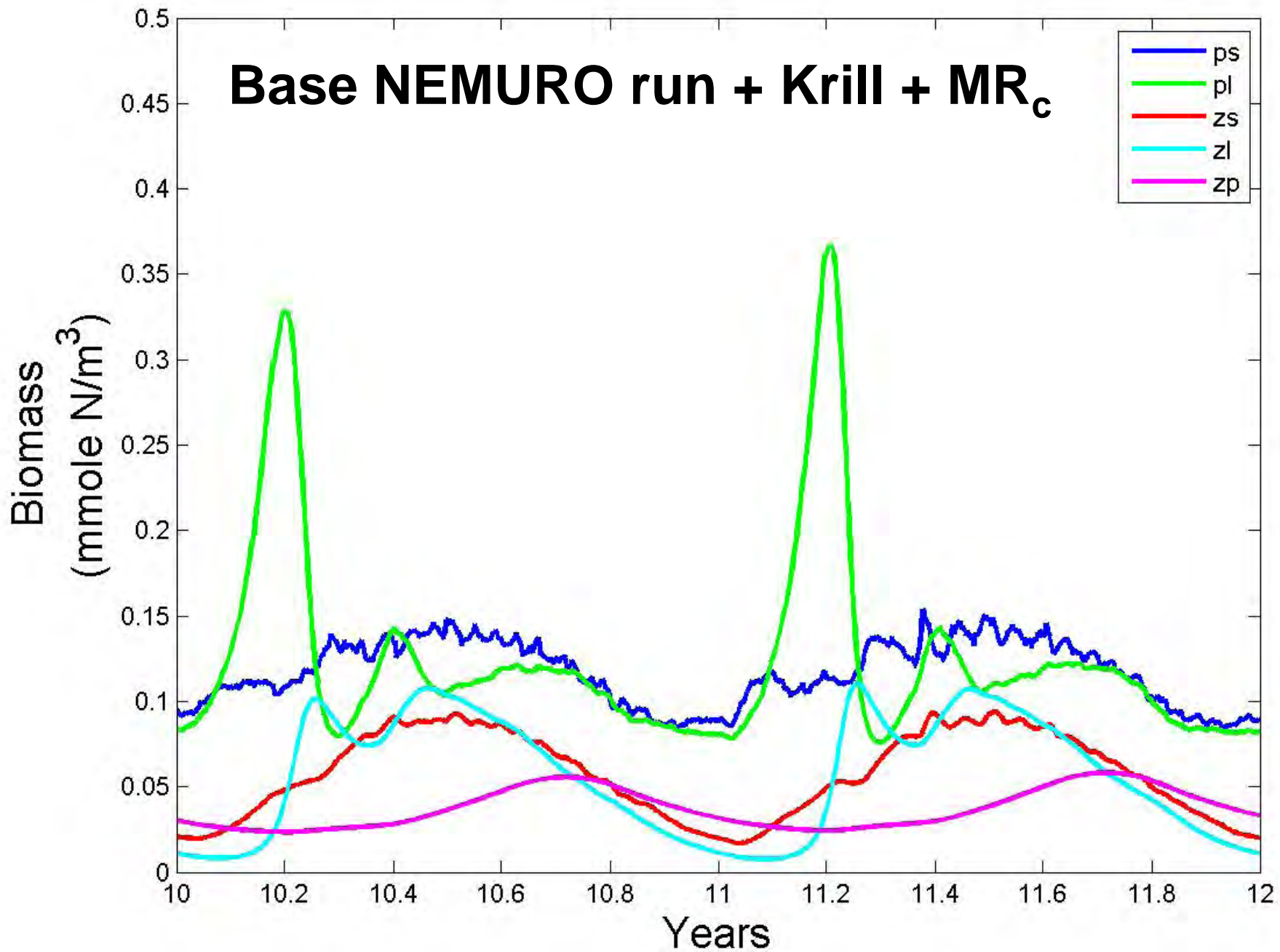
# Base NEMURO run



# Base NEMURO run + Krill

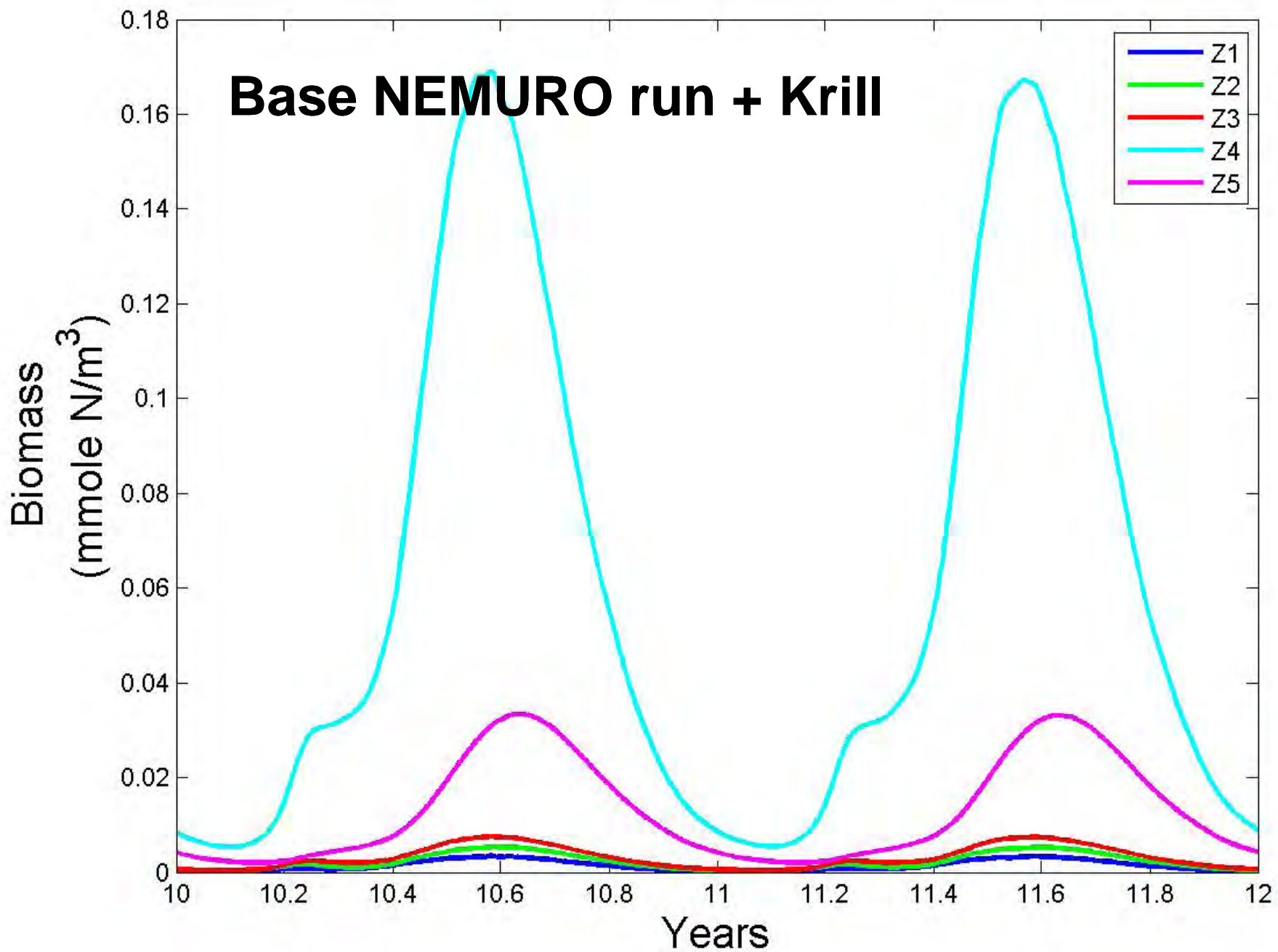


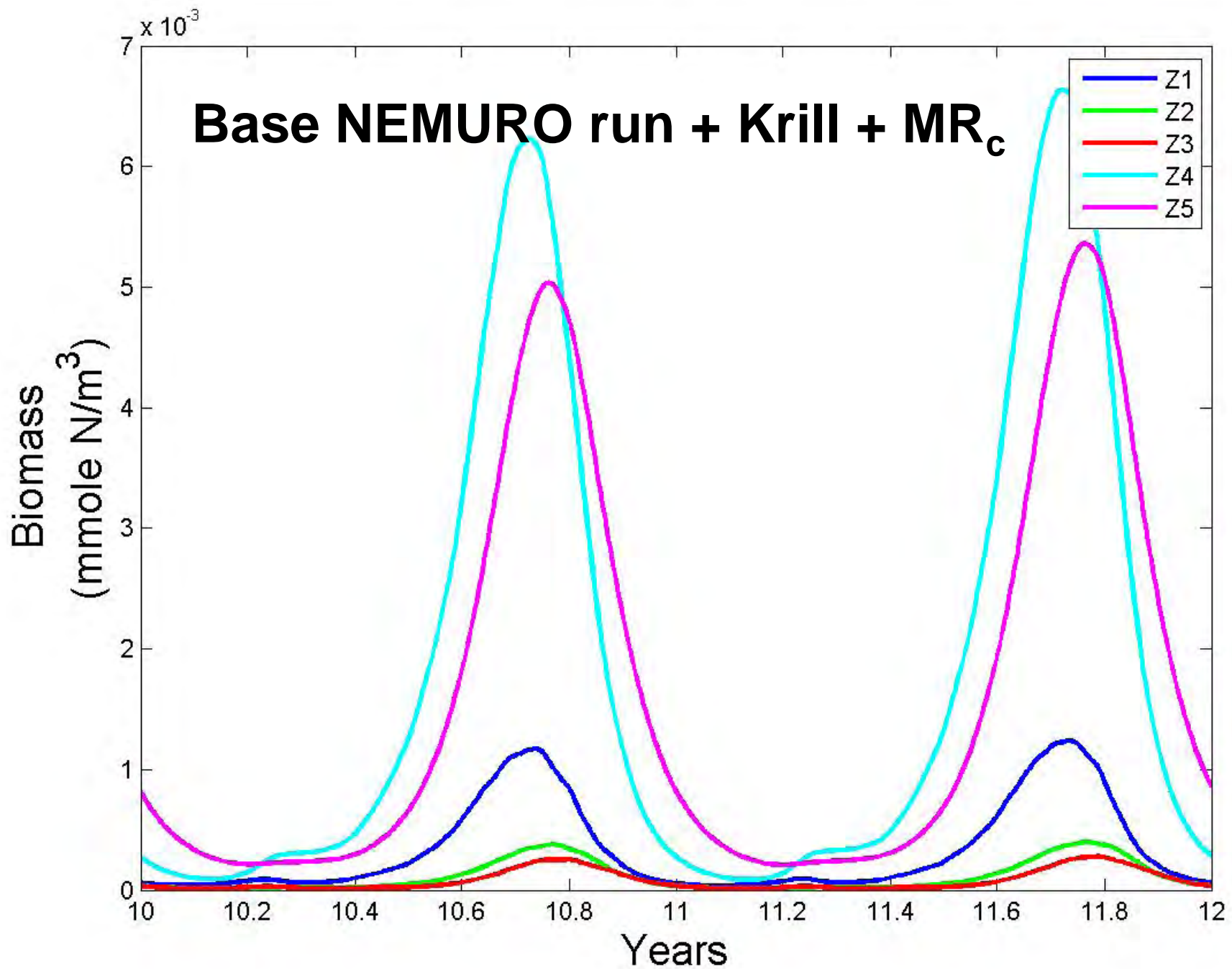
# Base NEMURO run + Krill + MR<sub>c</sub>



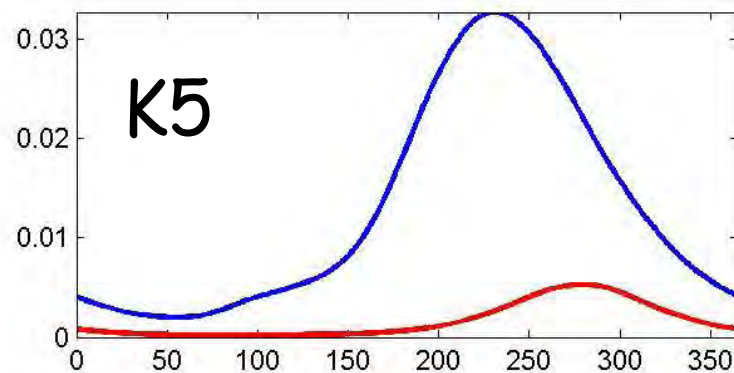
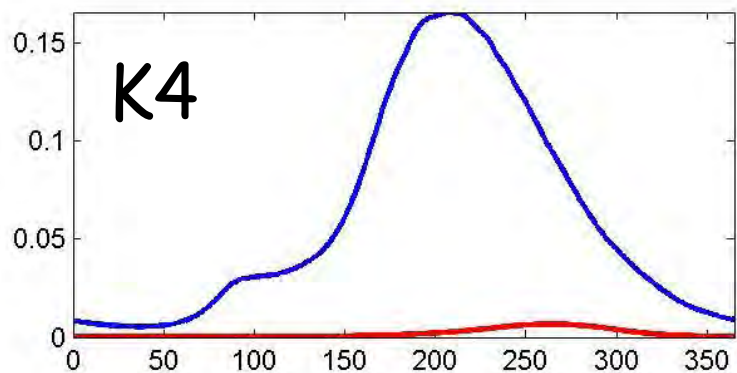
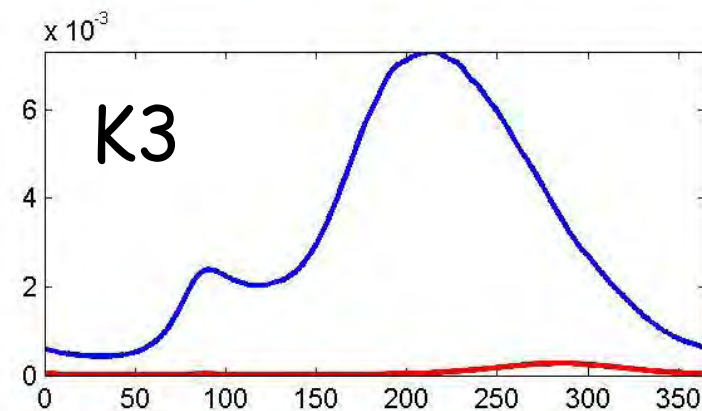
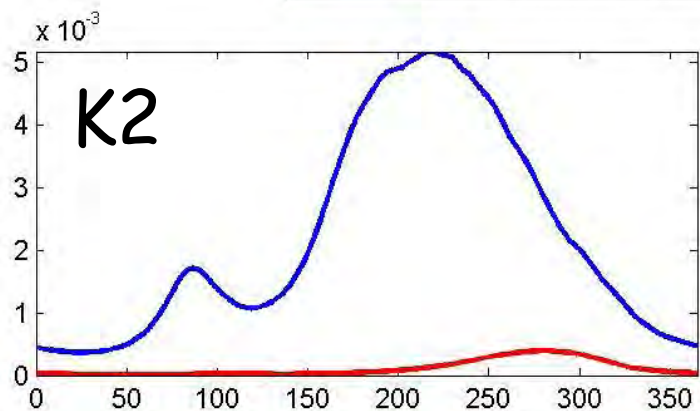
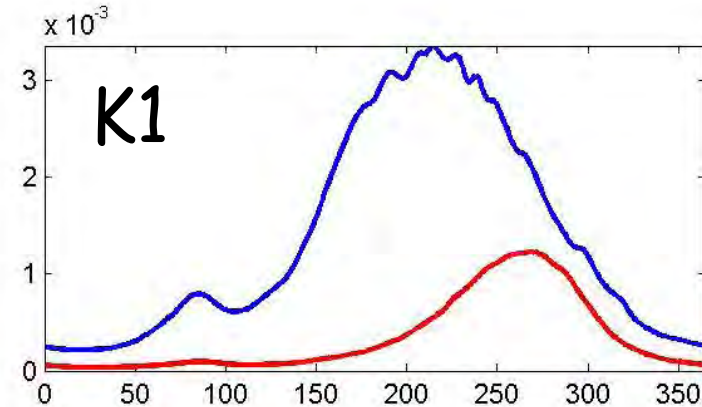
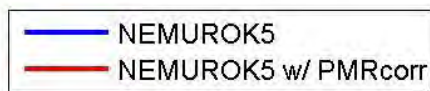


# Base NEMURO run + Krill

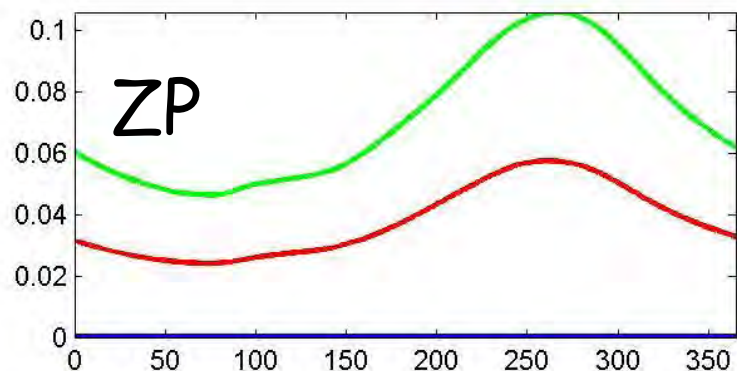
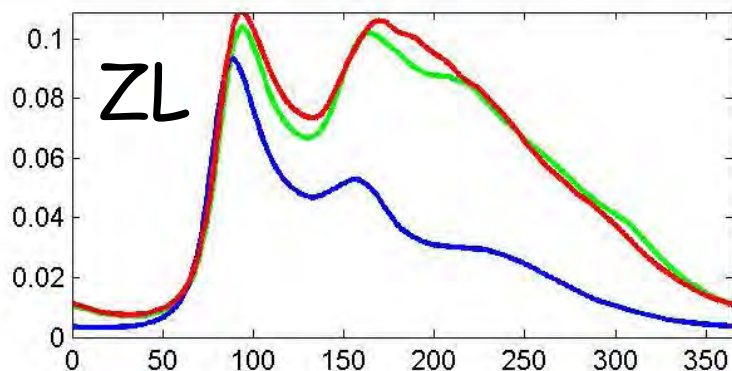
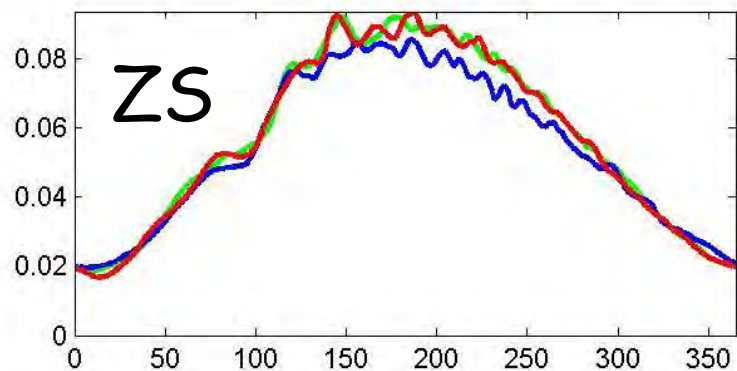
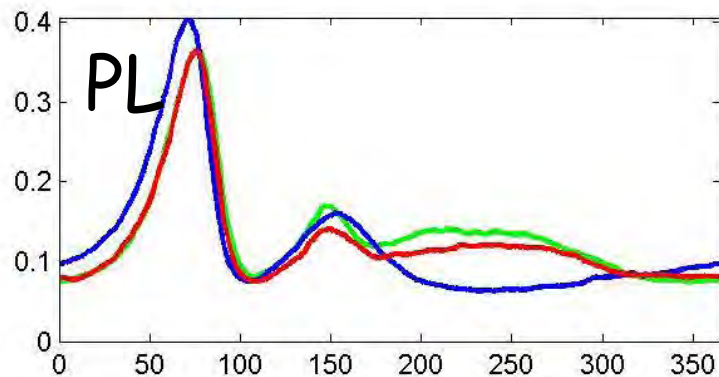
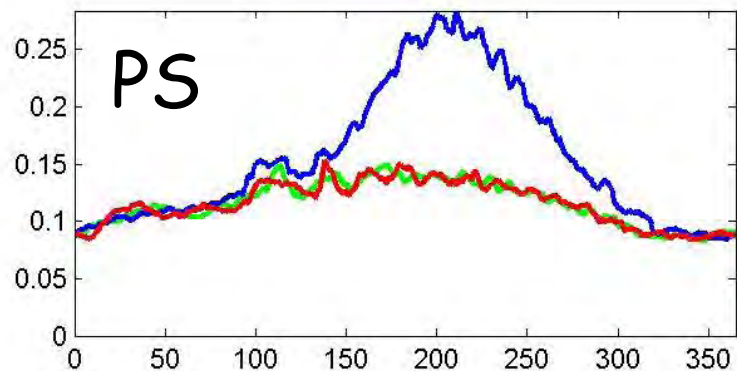
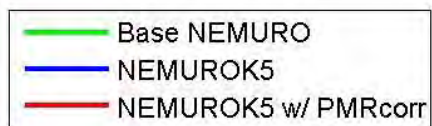




# Krill conc. (mmoles m<sup>-3</sup>)

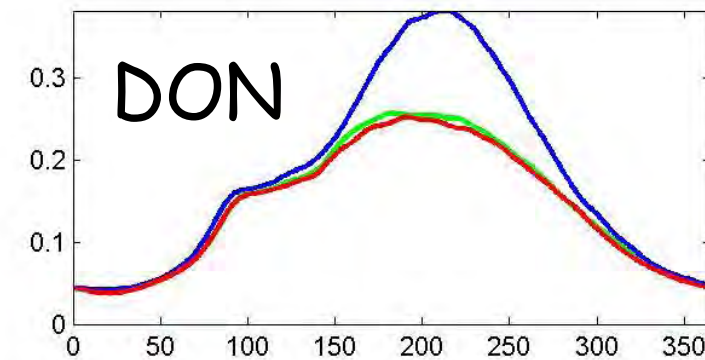
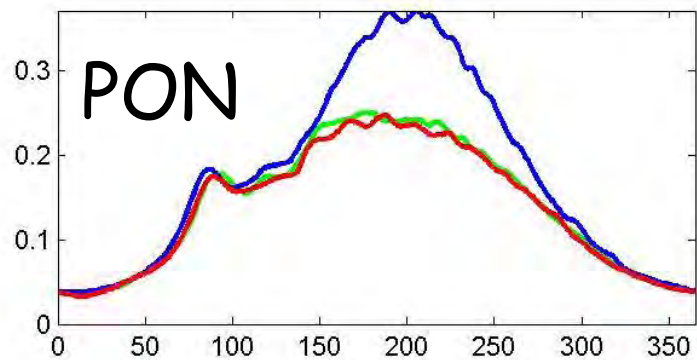
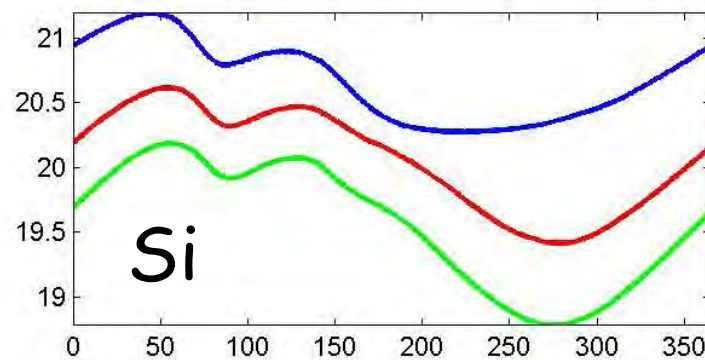
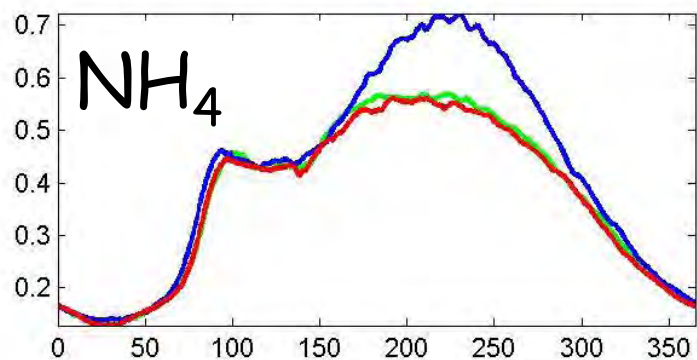
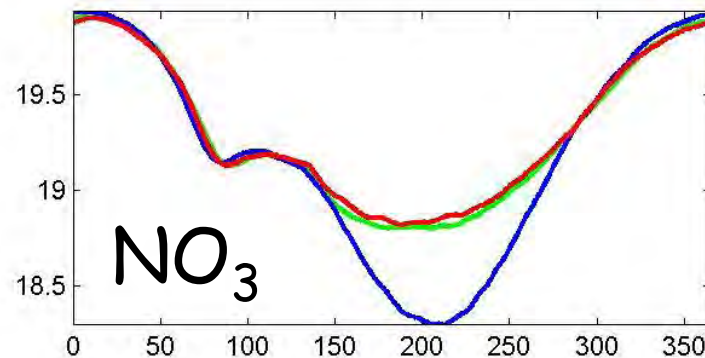
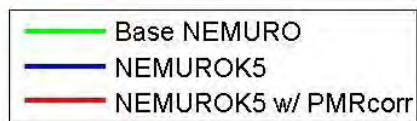


# NEMURO conc. (mmoles m<sup>-3</sup>)





# NEMURO conc. (mmoles m<sup>-3</sup>)





## Summary & Preliminary Findings

- 1) Adding krill to NEMURO **decreases ZL and ZP**, the latter to extinction, and **increases PS** (through a trophic cascade).
- 2) Adding krill+MRc reduces ZP by about 50% from the base NEMURO, but has little effect on other Z's and P's in NEMURO.
- 3) The **ZP reduction** (krill+MRc) or extinction (krill) must be **occurring through competition**, since no life stage of krill feeds directly upon ZP.
- 4) Krill concentrations with MRc are much lower (~5%) of concentrations in the krill model w/o MRc. **Delayed transfer** from young stages of krill to older stages **exposes krill to the higher mortality rates** of the younger stages for a long time.
- 5) **Timing of peak krill biomass is delayed** by ~60 days in simulations with the MRc.
- 6) The **mean age & concentration approach** of Hu et al. (2008) can be applied to molting of krill in Eulerian models, and will have value also for **implementing DVM behaviors (swimming speeds)** in Eulerian models.

**THE  
END**