

PISCES

(Pelagic Interaction Scheme for Carbon and ecosystem studies)



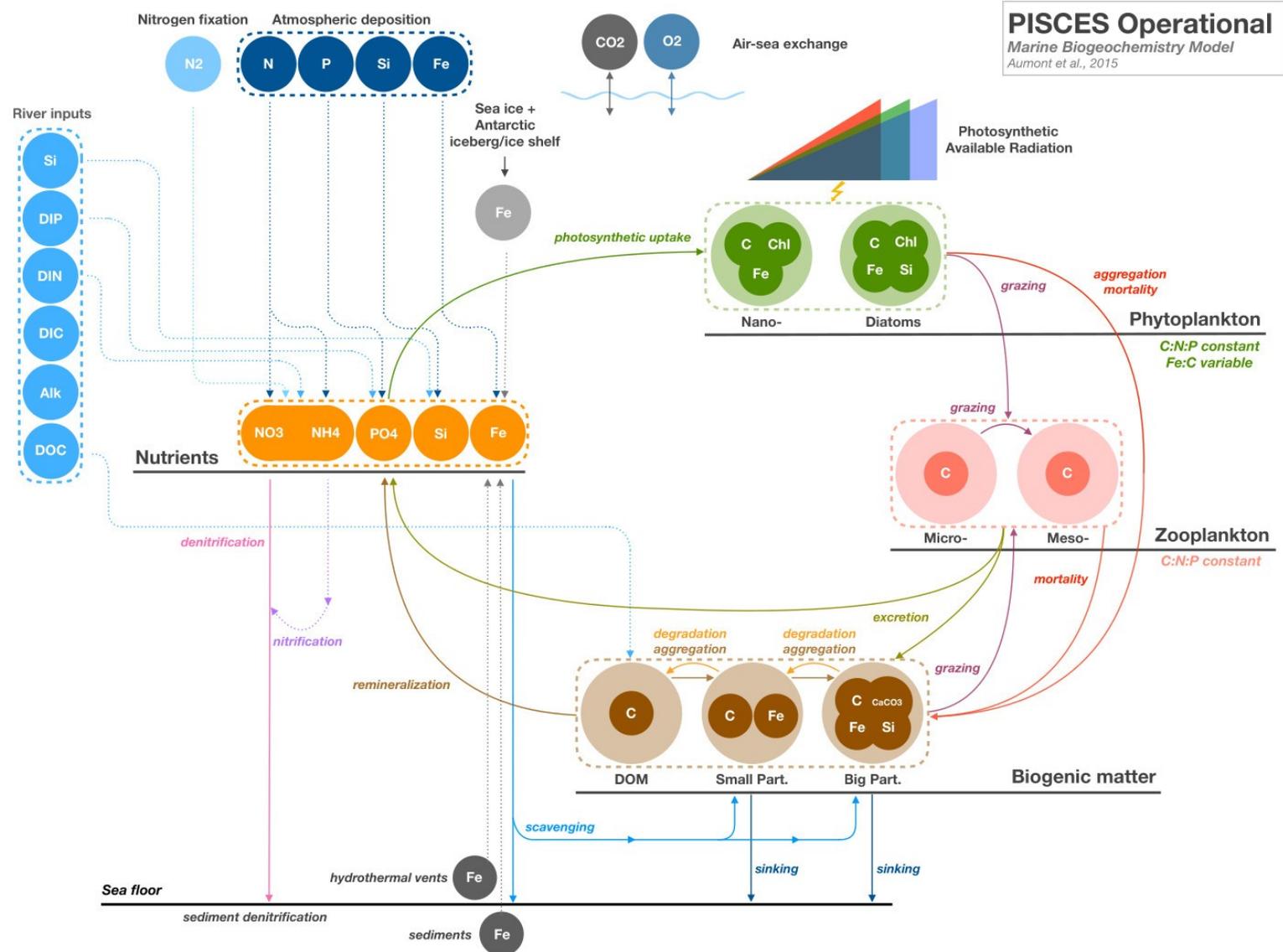
PISCES

Basic information on the model options

PISCES-std vs. PISCES-QUOTA

`ln_p4z = .true.`

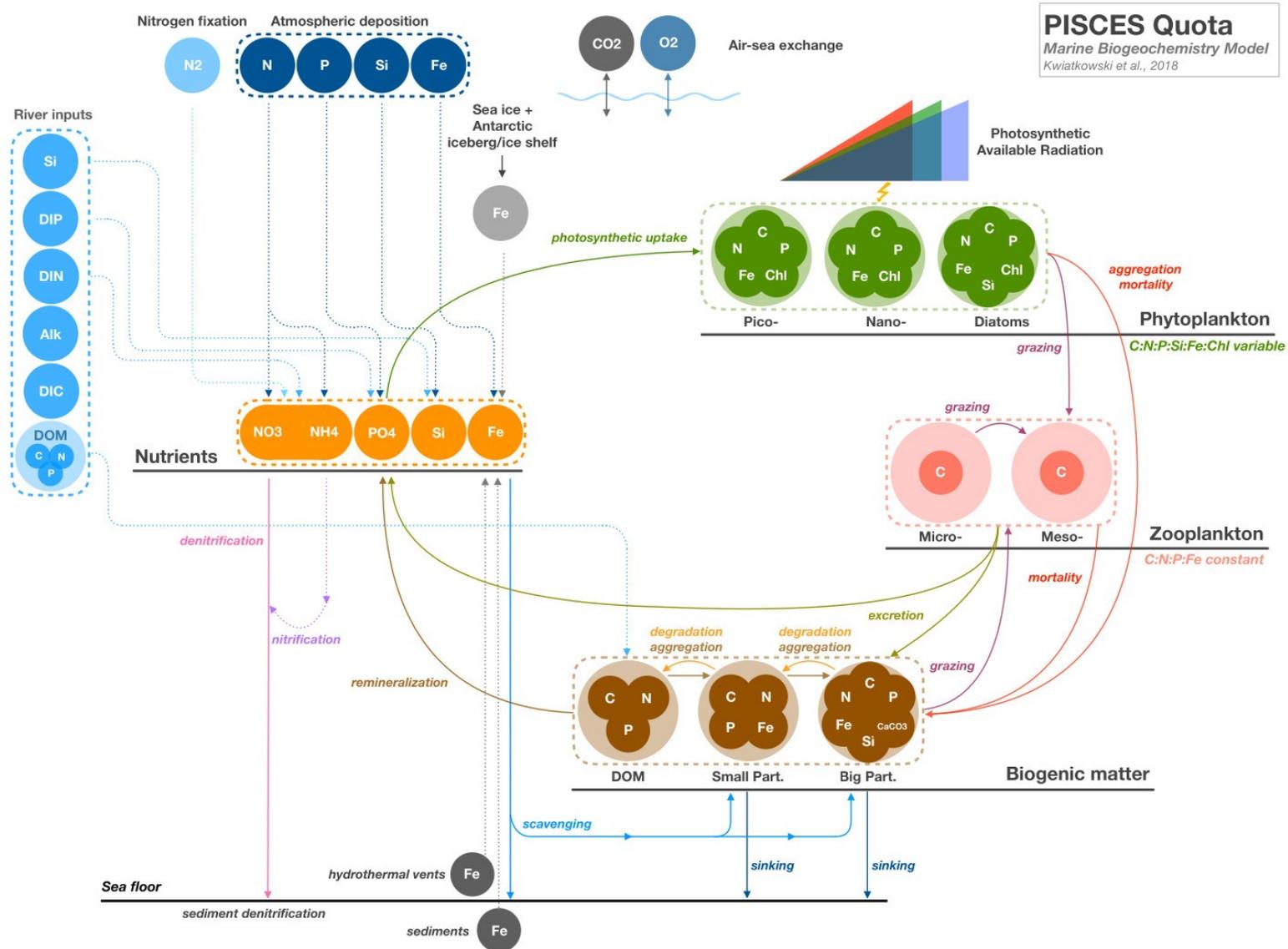
PISCES-std (24/25 tracers)



PISCES-std vs. PISCES-QUOTA

`ln_p5z = .true.`

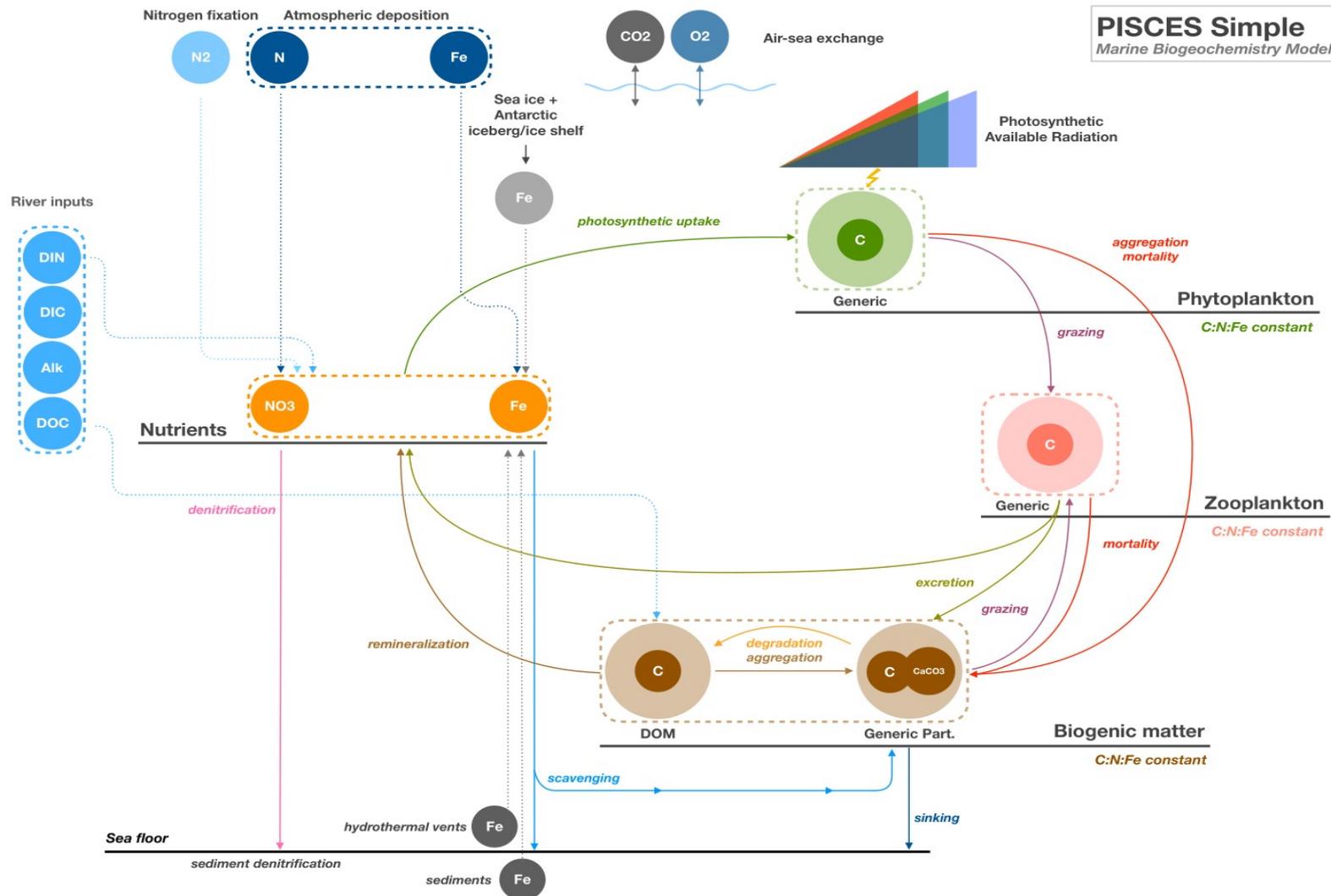
PISCES-QUOTA (39/40 tracers)



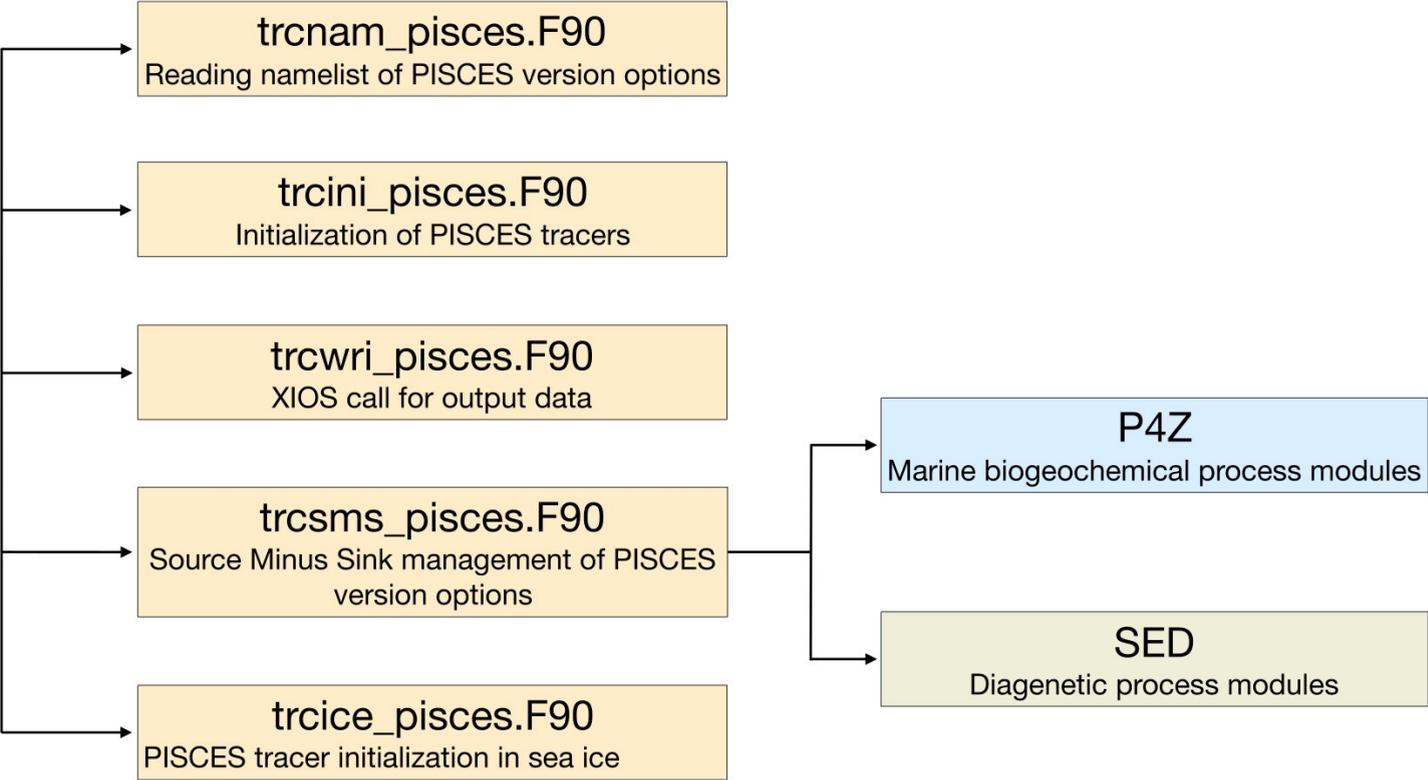
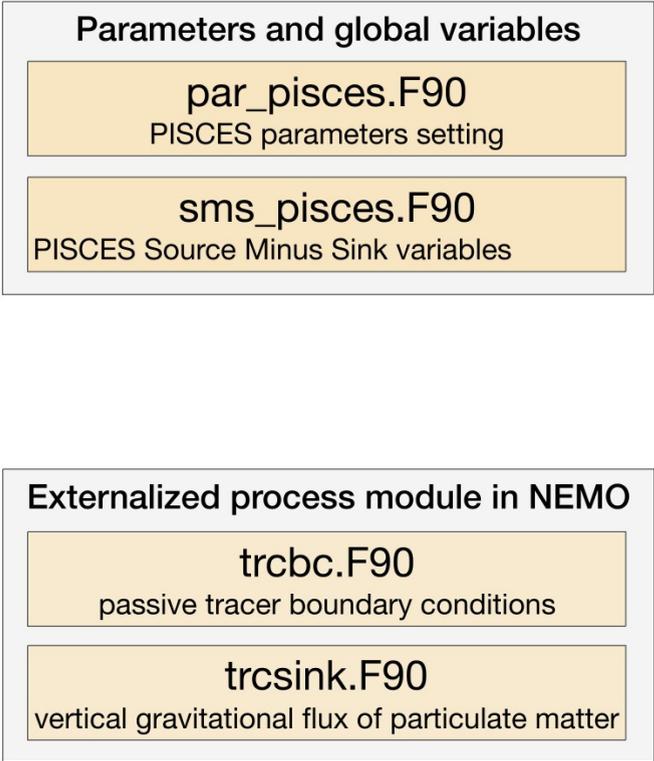
PISCES-simple vs. PISCES-STD

`ln_p2z = .true.`

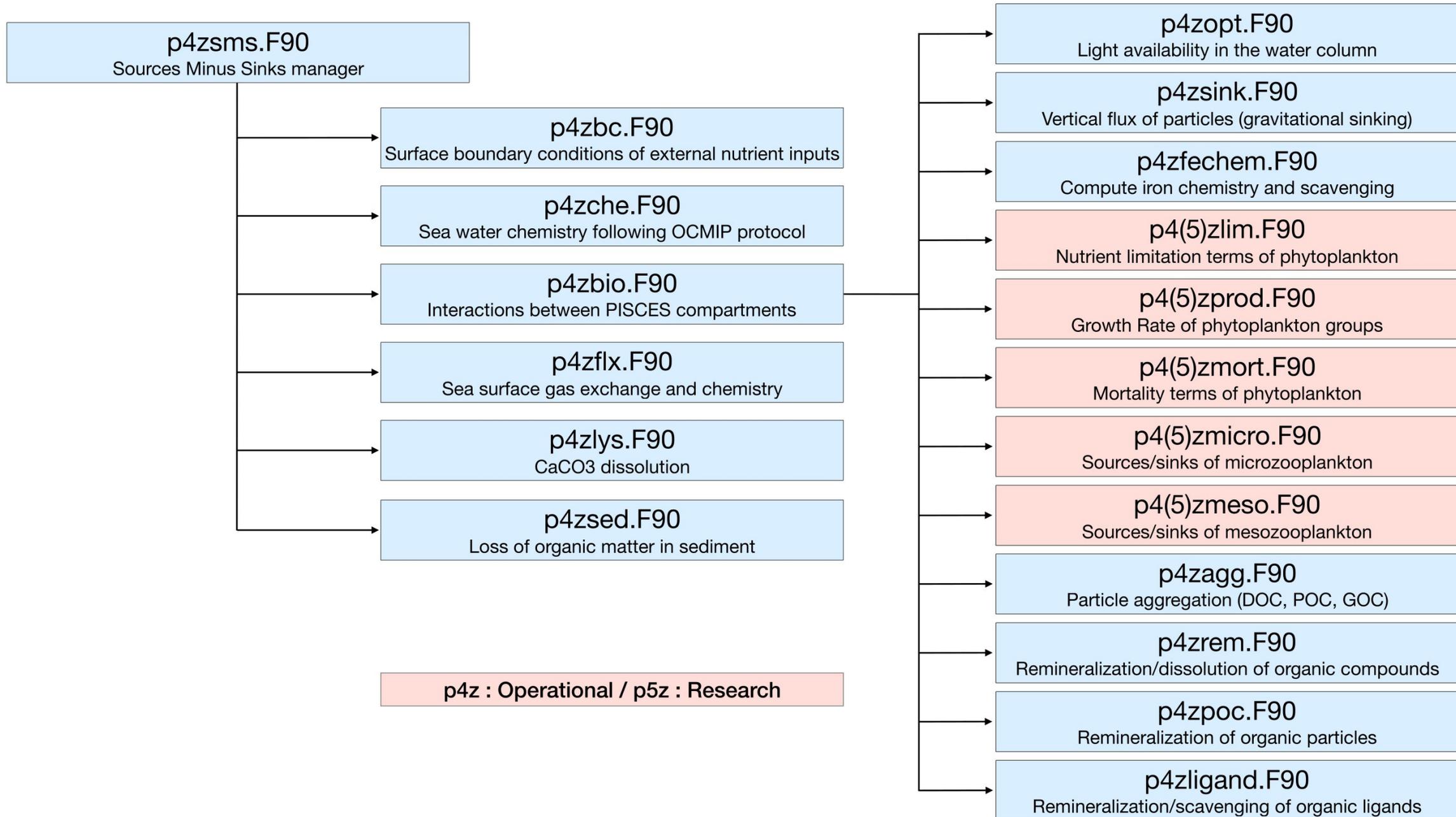
PISCES-Simple (9 tracers)



Code structure - Main



Code structure - SMS

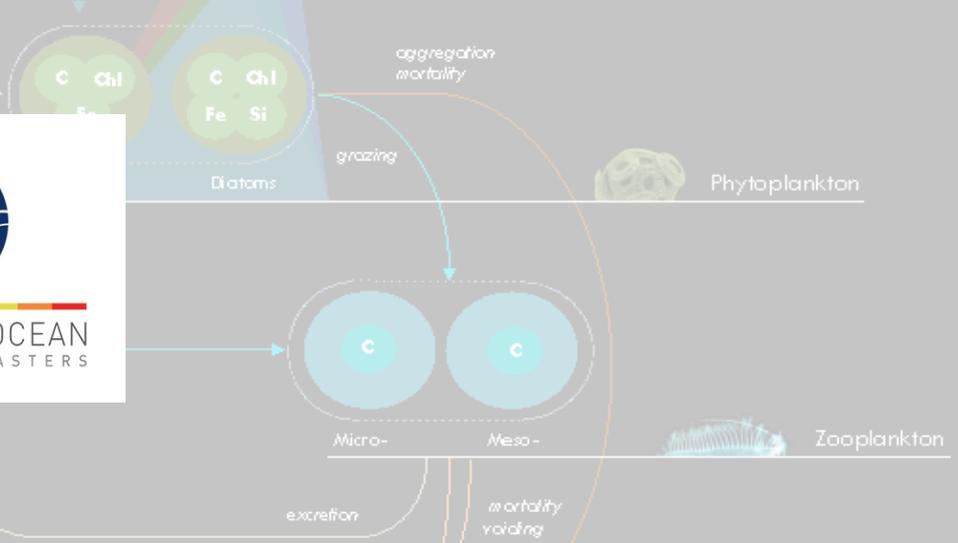
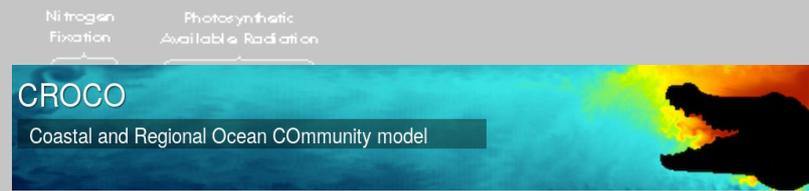
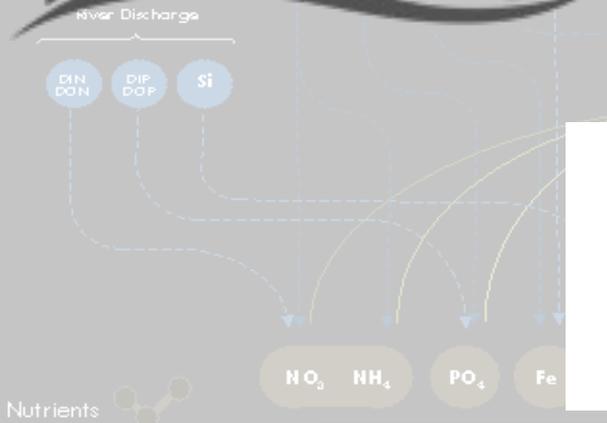


Modeling platforms

PISCES

Ocean Biogeochemical Model
Aumont and Bopp, 2006

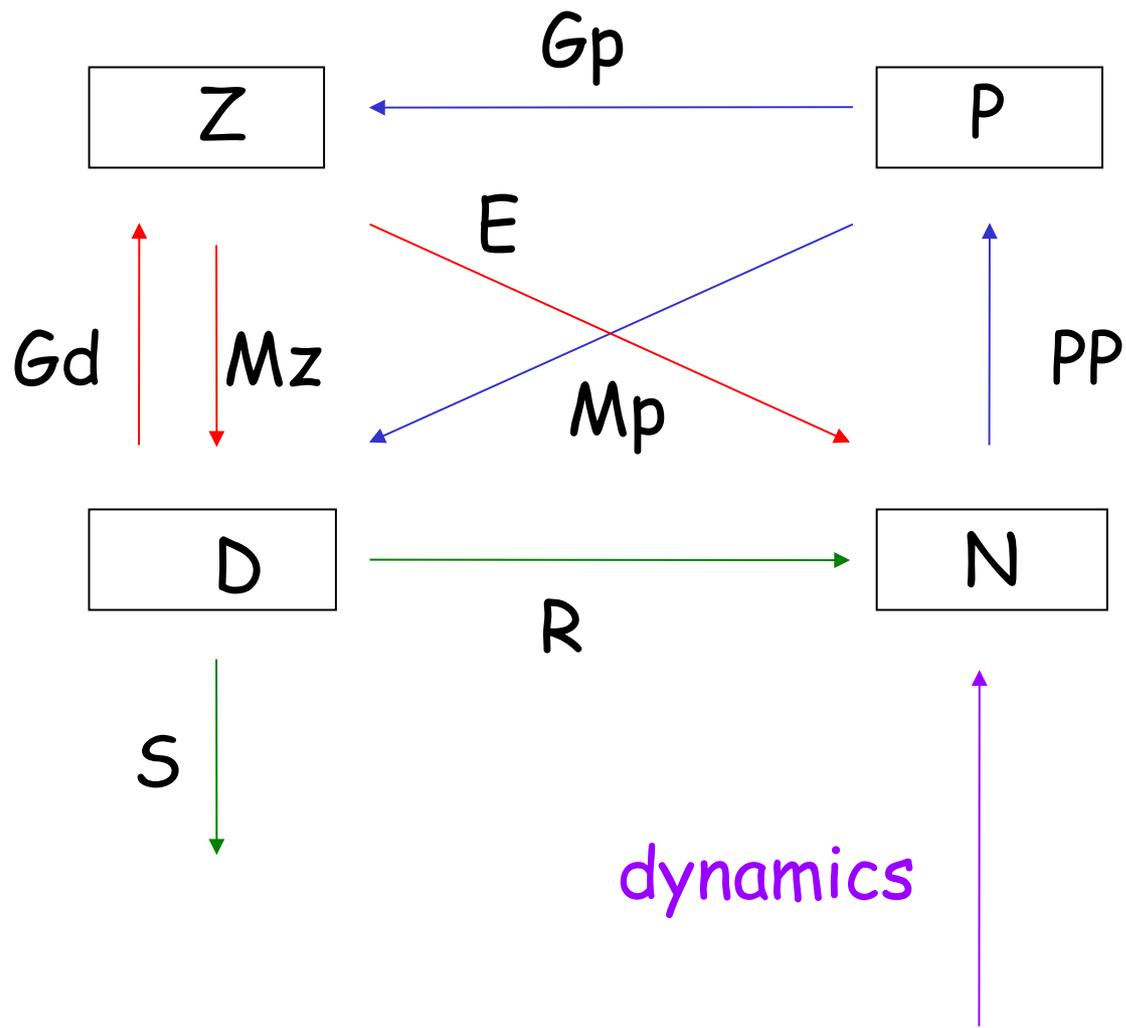
C/N/P
Redfieldian



Parameterizing marine biogeochemistry

Olivier Aumont

LOCEAN



(1) Phyto

$$S(P) = PP$$

$$P(P) = Gp + Mp$$

(2) Zoo

$$S(Z) = Gp + Gd$$

$$P(Z) = Mz + E$$

(3) Detritus

$$S(D) = Mz + Mp$$

$$P(D) = Gd + S + R$$

(4) Nutrients

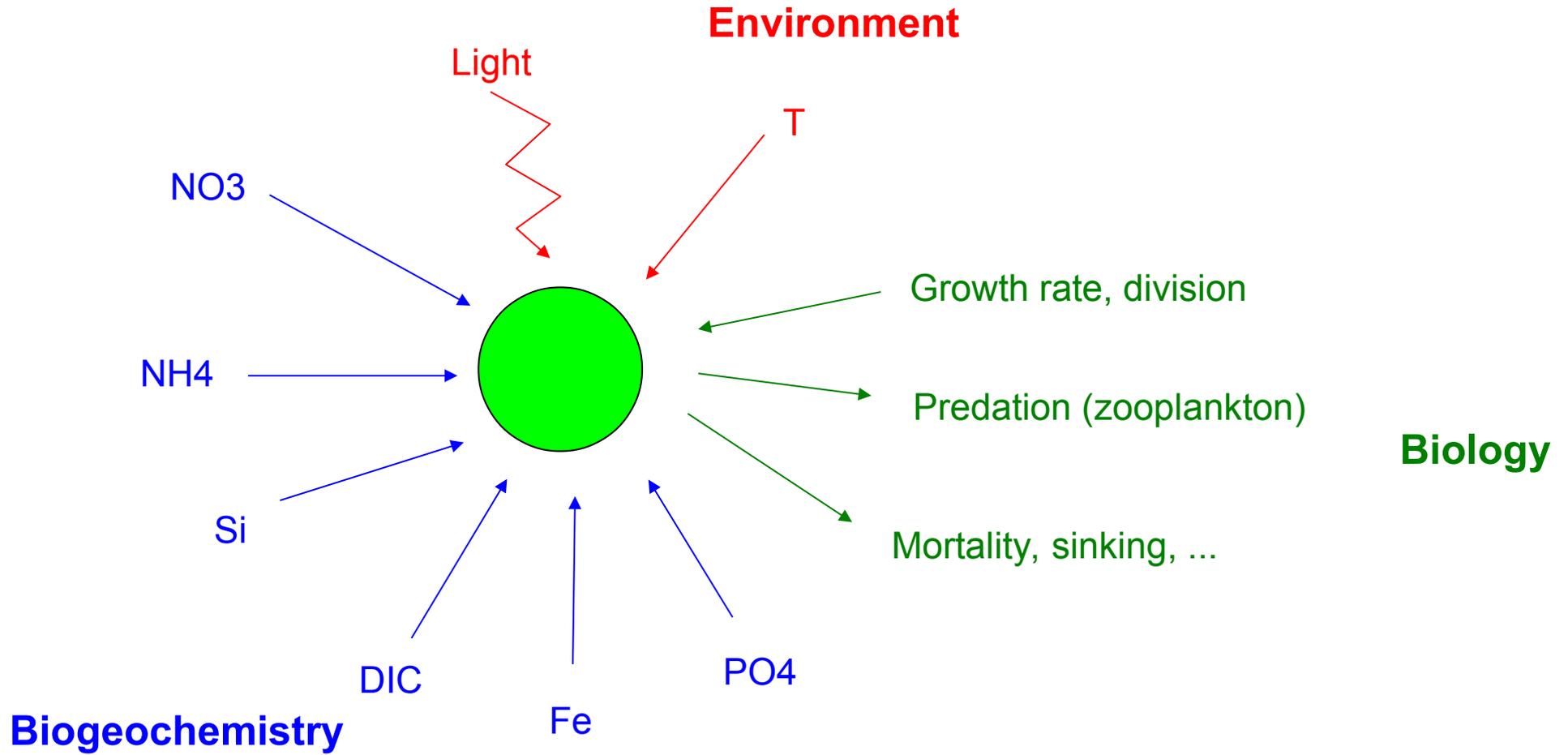
$$S(N) = R + E$$

$$P(N) = PP$$

Conservation:

$$S(P) + S(Z) + S(D) + S(N) - P(P) - P(Z) - P(D) - P(N) = 0$$

Colimitations: Background



Photosynthesis, growth rate

Photosynthesis

- Photosynthesis: Process by which autotrophic organisms use solar energy to produce organic matter



- The ratio between the different chemical elements is called the Redfield ratio
- The amount of organic matter produced by the photosynthesis is called Gross Primary Production

Growth rate

- Growth refers to the increase in biomass (in C units generally)

$$\mu = P^B \frac{\text{Chl}}{C} - r \quad \text{Where } r \text{ is the respiration rate}$$

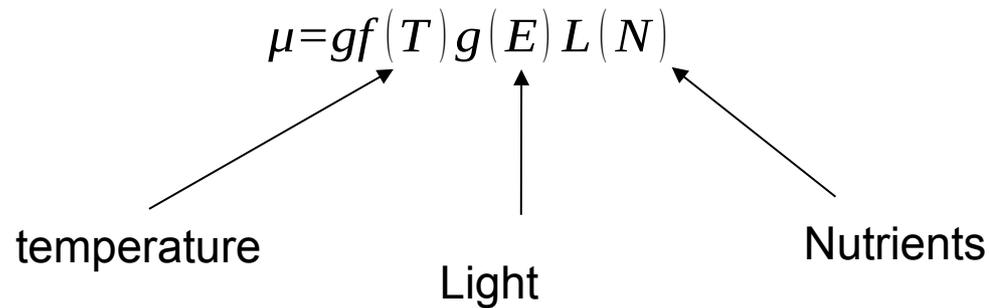
- From the growth rate, net primary production can be defined as the accumulation of organic matter

Growth rate: General background

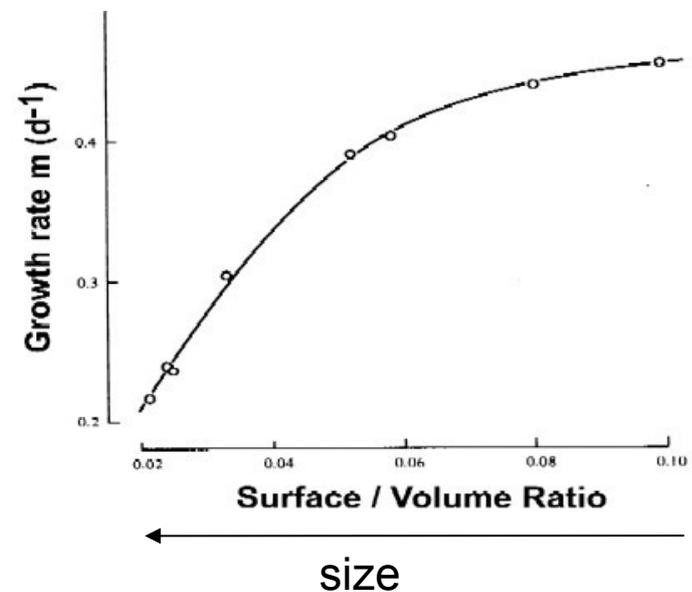
- Growth rate is a function of the environmental and biogeochemical conditions and of the species
- It can be expressed as follows:

$$\mu = gf(T)g(E)L(N)$$

temperature Light Nutrients

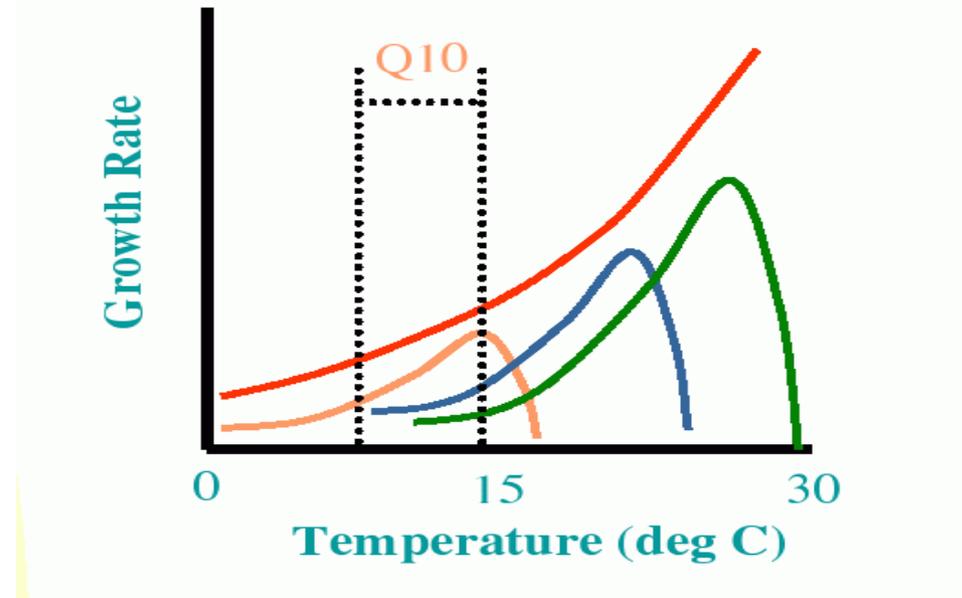


- The specific growth rate varies with species. In general, it tends to decrease with size.



Growth rate: Temperature

- Growth rate increases with temperature until a critical level



- A relationship for the envelope has been proposed for the first time by Eppley (1972) :

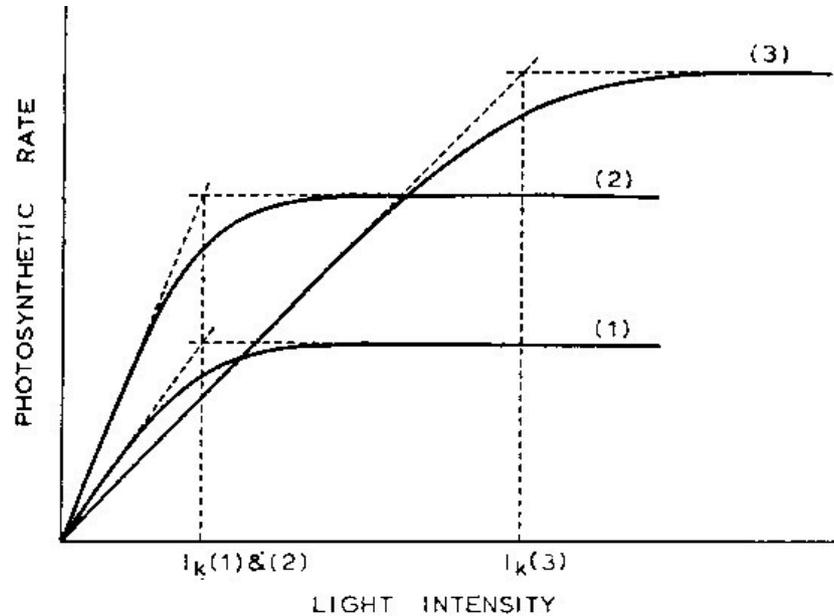
$$f(T) = 1.066^T$$

Growth rate increases by 1.9 times every 10°C (Q10).

- Since, several alternative expressions have been proposed, but Eppley's relationship remains the most commonly used one.

Growth rate: Light

- Growth rate increases with light until a maximum value at which it saturates or even decreases



α initial slope

μ_{max} maximum growth rate

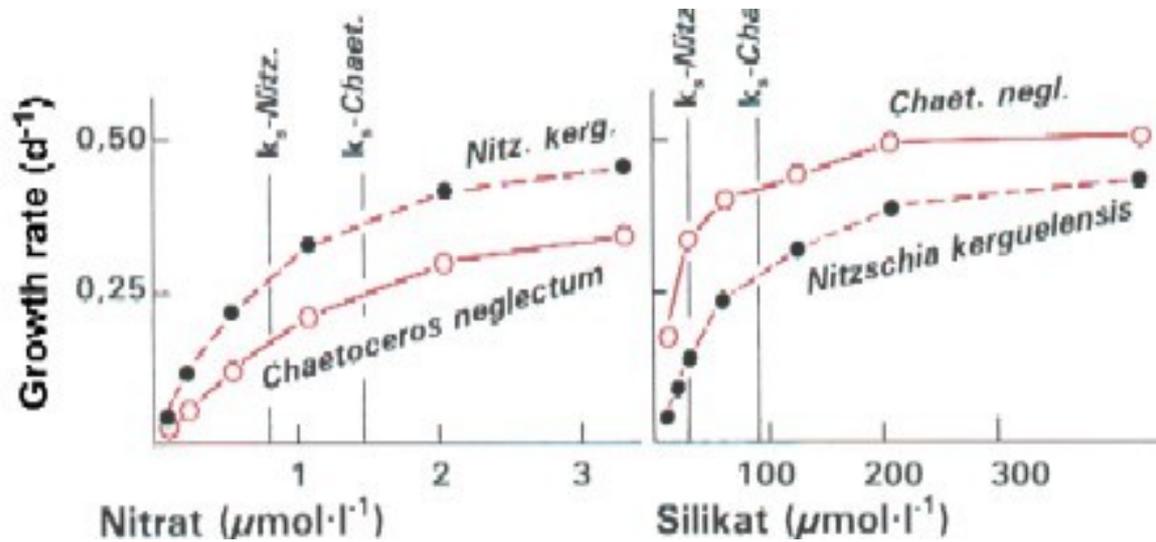
$$I_k = \mu_{max} / \alpha$$

$$\mu = \mu_{max} \left(1 - \exp\left(\frac{-I}{I_k}\right) \right)$$

- I_k is extremely variable between species. For instance, in cyanobacteria, *synechococcus* spp have a high I_k whereas some *prochlorococcus* spp have very low I_k .
- I_k strongly varies with the average received light (photoacclimation)

Growth rate: Monod model (1942)

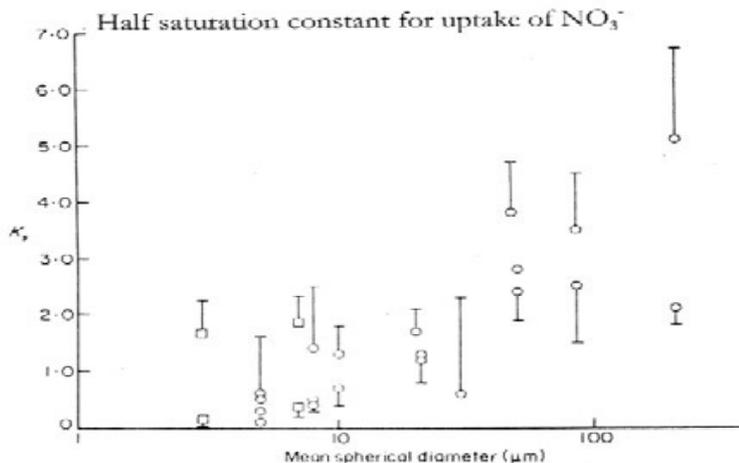
Monod model = growth rate is a function of the external concentration of nutrients



$$\mu = \mu_{max} \frac{N}{K_N + N}$$

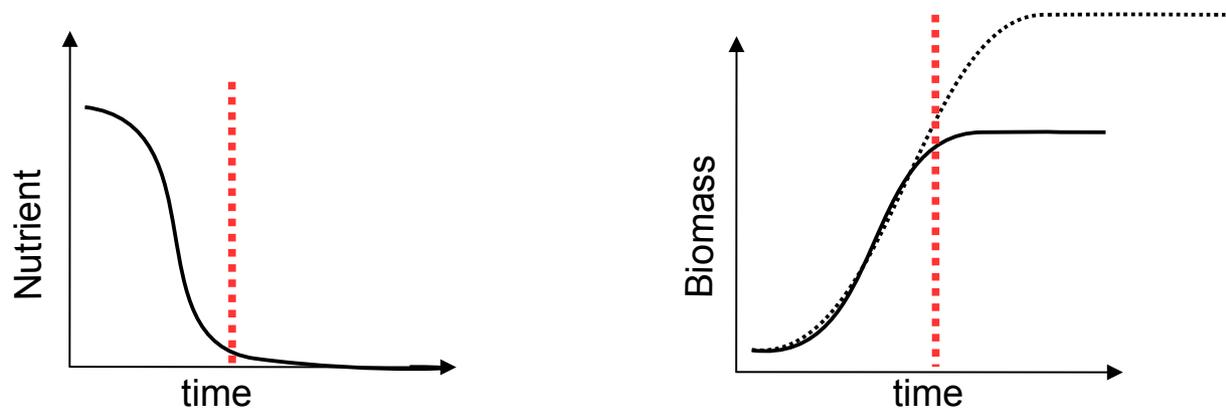
K_N : half-saturation constant

K_N increases with size because the S/V ratio decreases. Furthermore, as a result of acclimation processes, K_N decreases with the nutrient concentration



Limitation of the Monod model

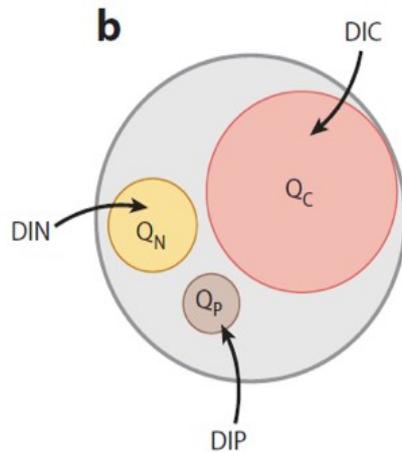
- Measured & works best under relatively steady nutrients (or slow change)
- Growth stops when nutrients fall to 0



- Assumes constant stoichiometry
- No luxury uptake of transiently elevated nutrients
- Can be difficult to estimate K_N

Growth rate: Droop model (1968)

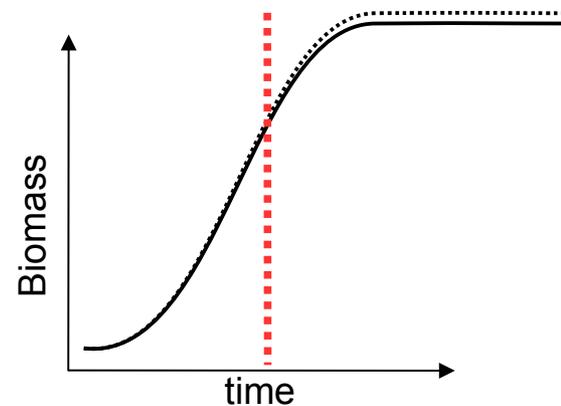
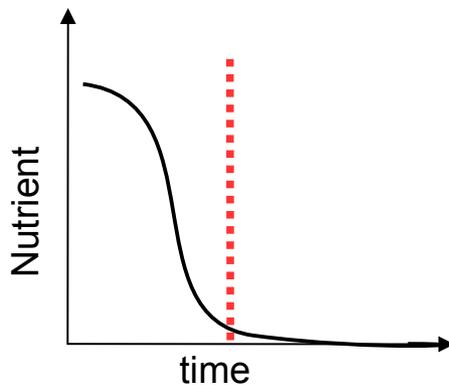
- Droop model = growth rate is a function of the internal pool of nutrients (quota)
- The internal pool (quota) is a function of the external concentrations of N



$$\text{growth} = \mu_{\infty} \left(1 - \frac{Q_{\min}}{Q} \right)$$

$$V = V_{\max} \frac{N}{K_N + N}$$

- Generally works better, more physiologically grounded, more general



Growth rate: multi-nutrients

- Currently, there is no clear consensus on the law which drives growth rate with multiple nutrients. 2 different laws are generally used:

- The multiplicative law:

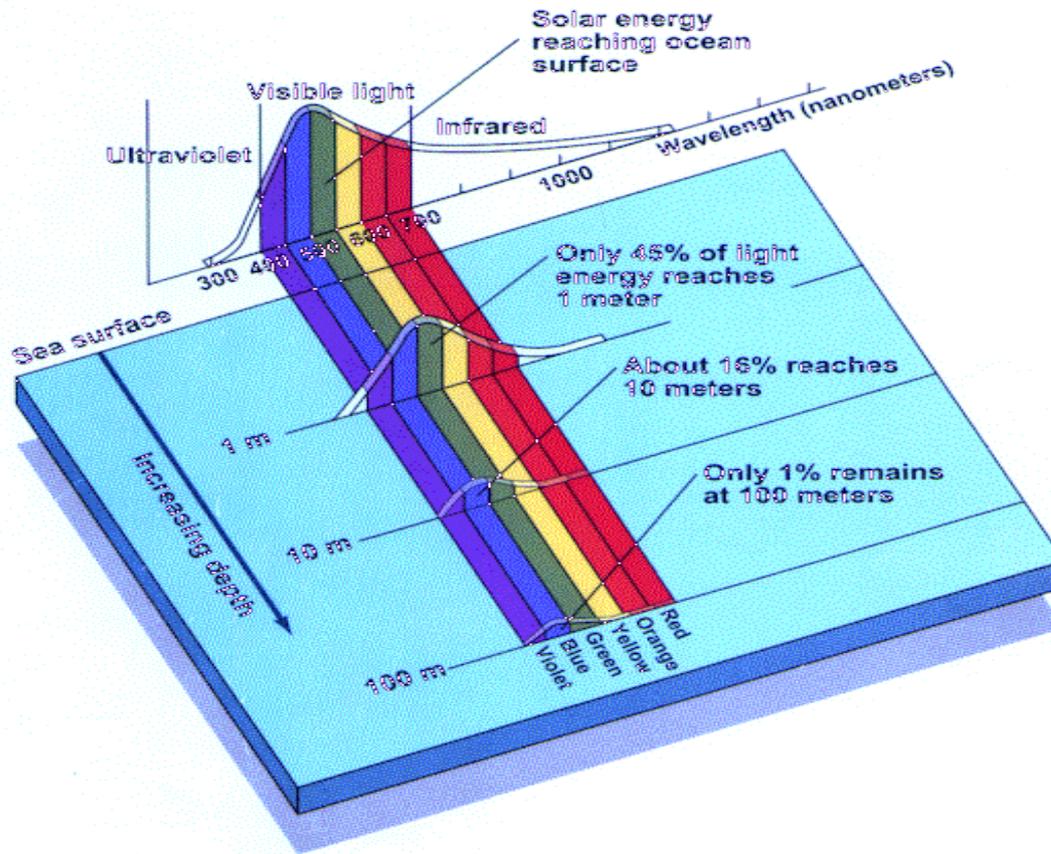
$$\mu = \mu_{max} \frac{N_1}{K_{N_1} + N_1} \frac{N_2}{K_{N_2} + N_2} * \dots$$

- The law of the minimum or Liebig's law (1840) :

$$\mu = \mu_{max} \min \left(\frac{N_1}{K_{N_1} + N_1}, \frac{N_2}{K_{N_2} + N_2}, \dots \right)$$

- Many other laws do exist but they are not commonly used.

Light in the ocean



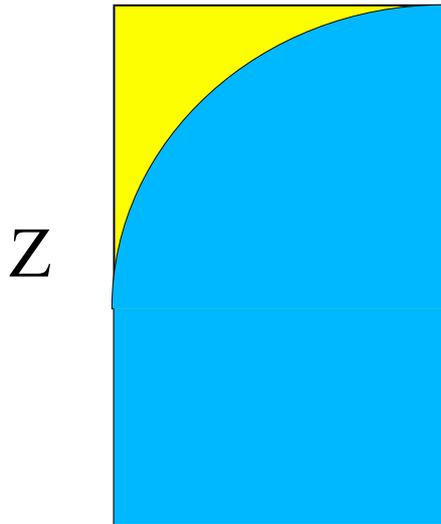
- Light decreases with depth as (Loi de Beer): $I(z) = I(0)e^{-kz}$
- In pure water, the attenuation length is 37m. for a chlorophyll concentration of 0.2 mg Chla/m³, its value is about 20m.
- Blue light penetrates much deeper than red light which remains trapped in the top 10 to 20m of the ocean.

Light in the ocean

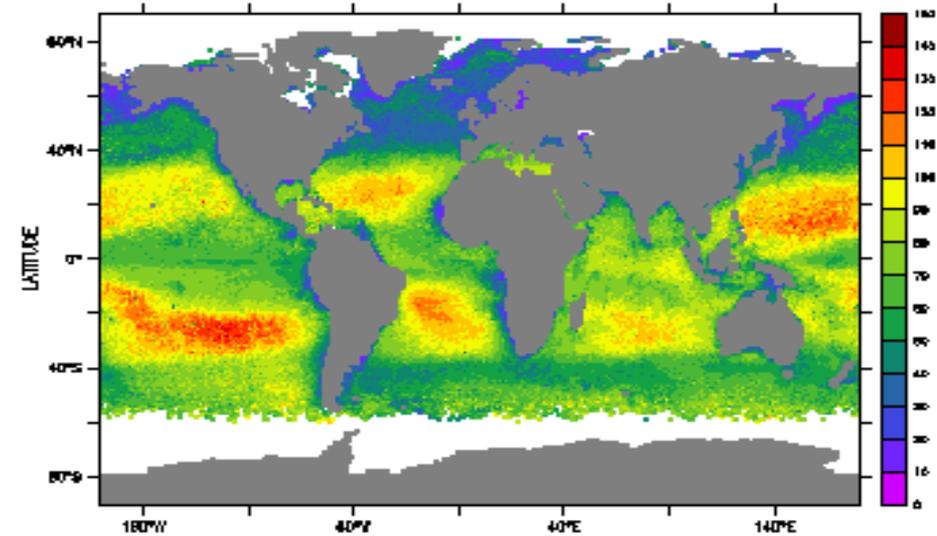
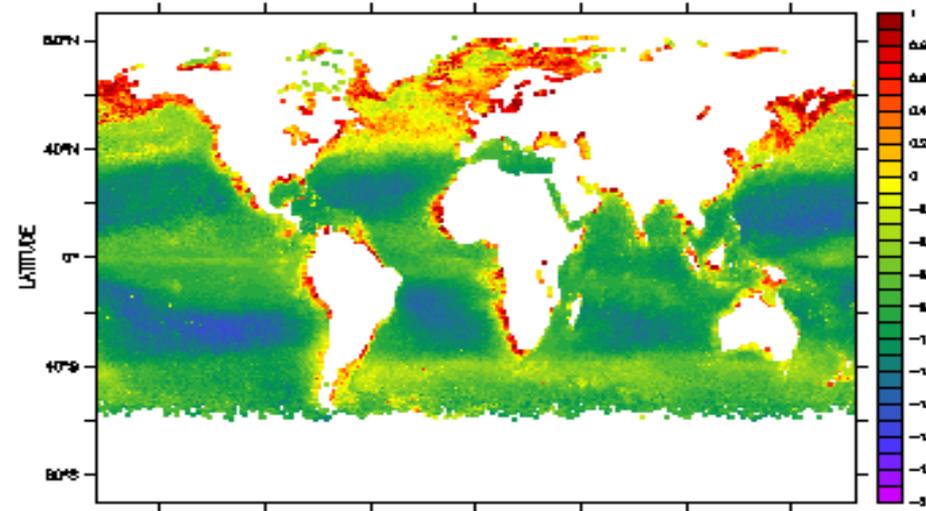
Euphotic zone

$$I > 1\% I_0$$

Solar radiation

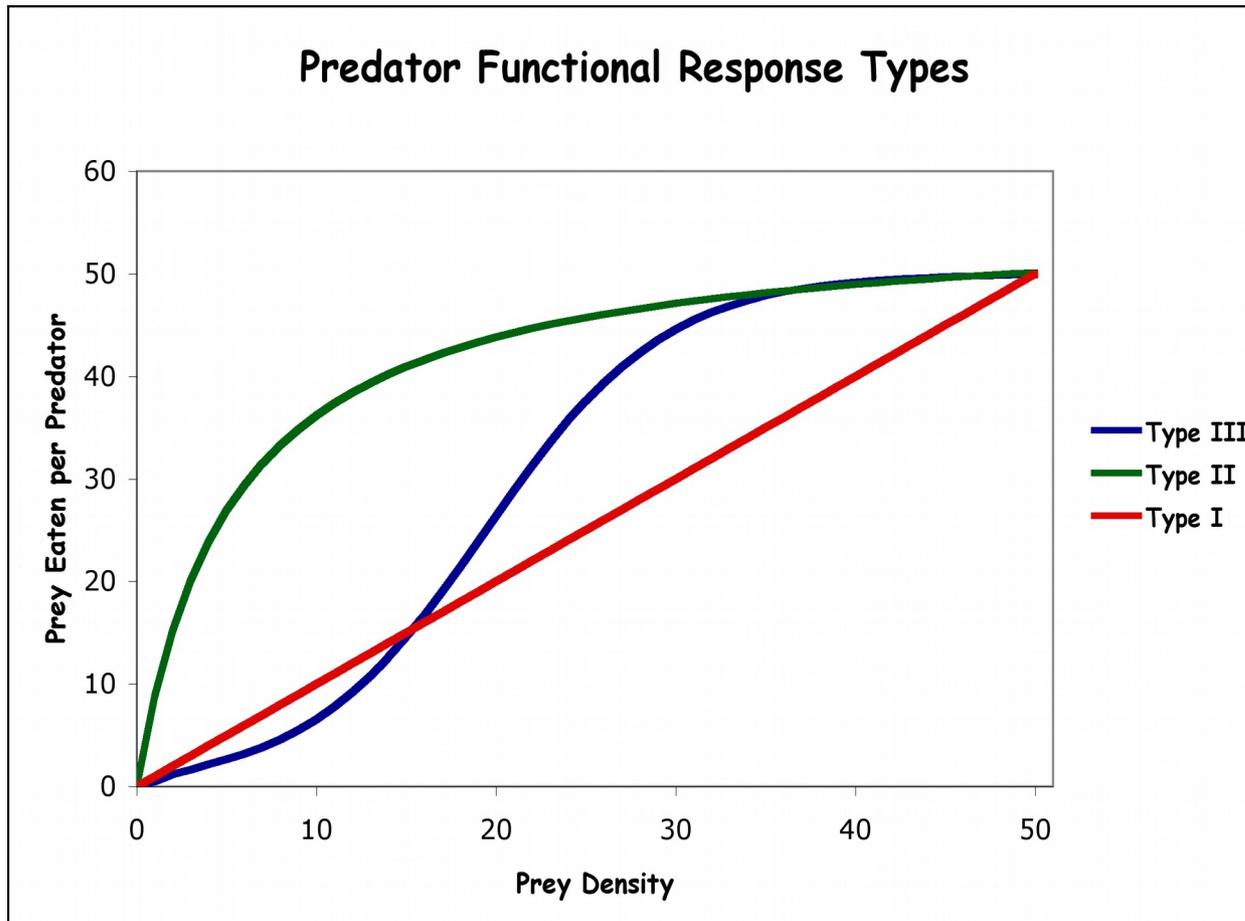


Chlorophyll



Euphotic depth

Predation by zooplankton



- Holling Type I

$$G_p = gP$$

- Holling Type II

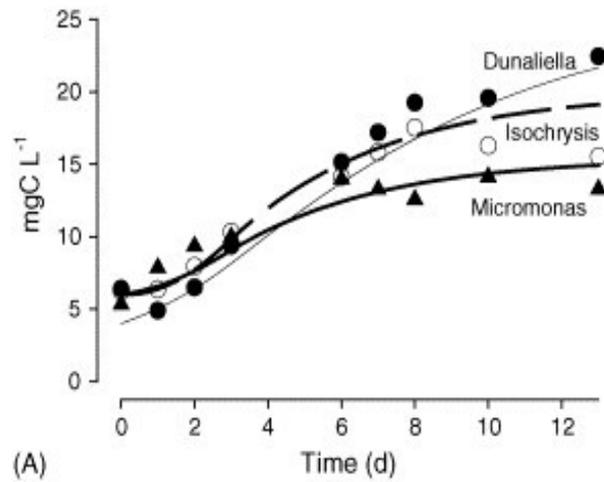
$$G_p = g \frac{P}{K_z + P} Z$$

- Holling Type III

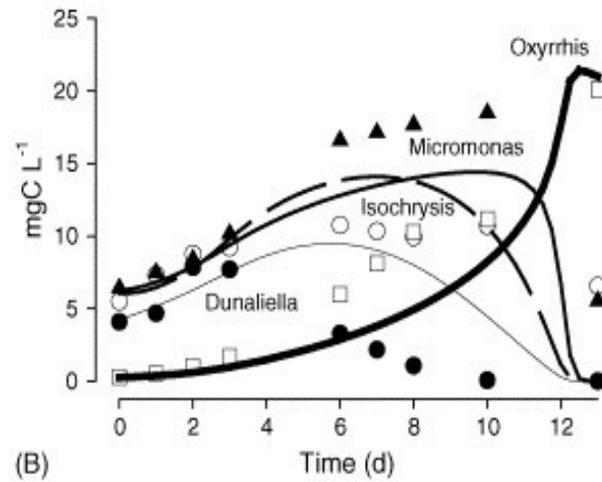
$$G_p = g \frac{P^2}{K_z^2 + P^2} Z$$

Predation on several preys

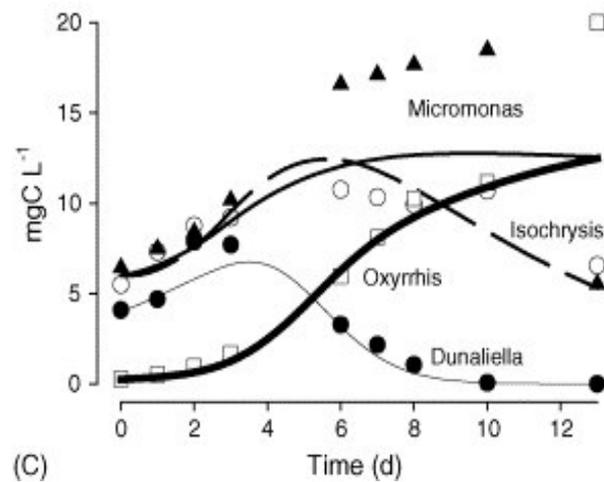
Very complex to properly model!



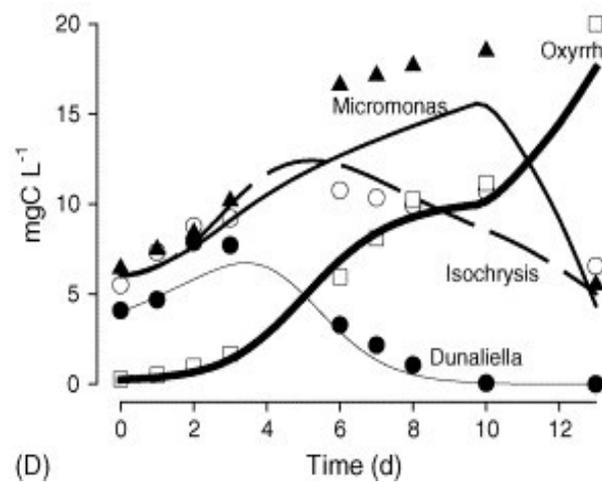
(A)



(B)



(C)



(D)

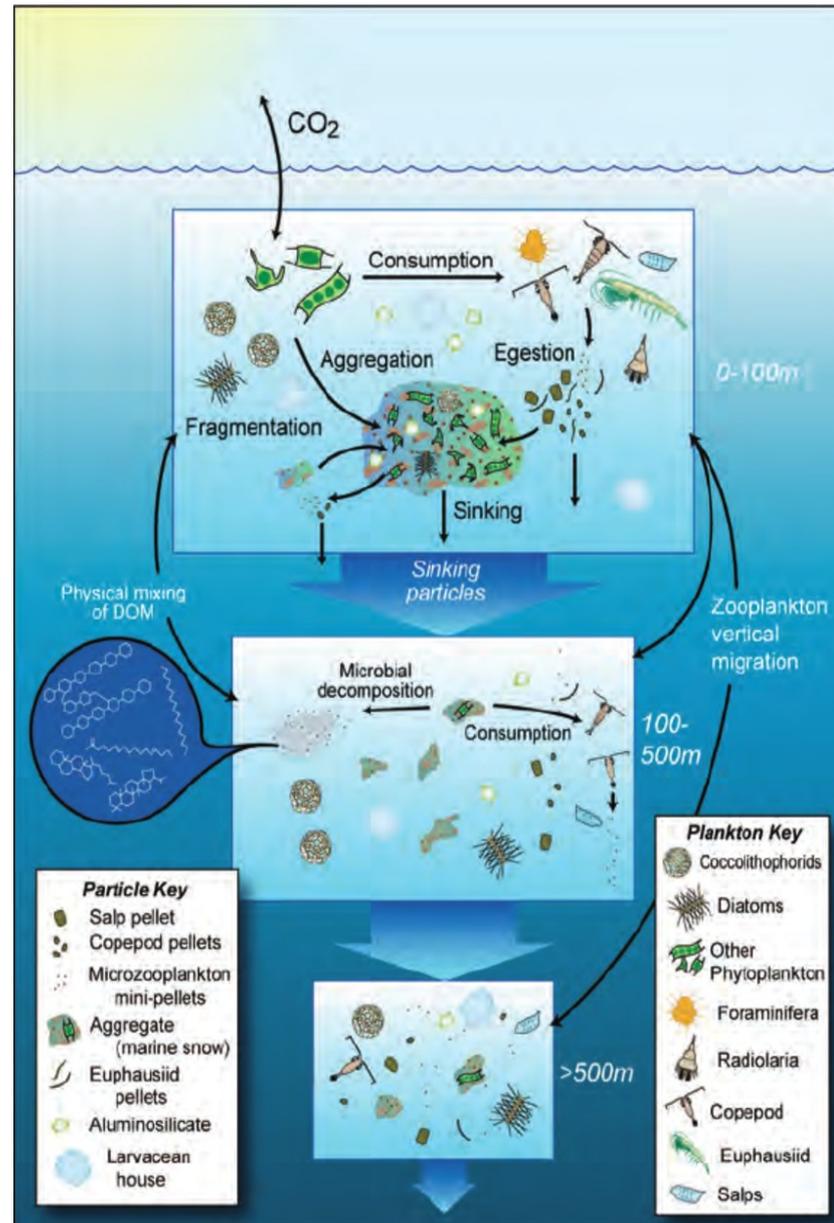
Mortality in models

- Mortality in models does not necessarily represent senescence. It may model :
 - senescence
 - viral attacks
 - aggregation/sinking
 - predation by unresolved higher trophic levels
- Numerous formulations exist but the two most common expressions are:

$$M = m_p P$$

$$M = m_p P^2$$

Representing particles in models



Representing particles in models: POC

- If bacterial degradation and sinking are the only active processes

$$\frac{\partial D}{\partial t} = \underbrace{-\lambda D}_{\text{Remineralization}} - \underbrace{\frac{\partial vD}{\partial z}}_{\text{Sinking}}$$

- Specific cases :

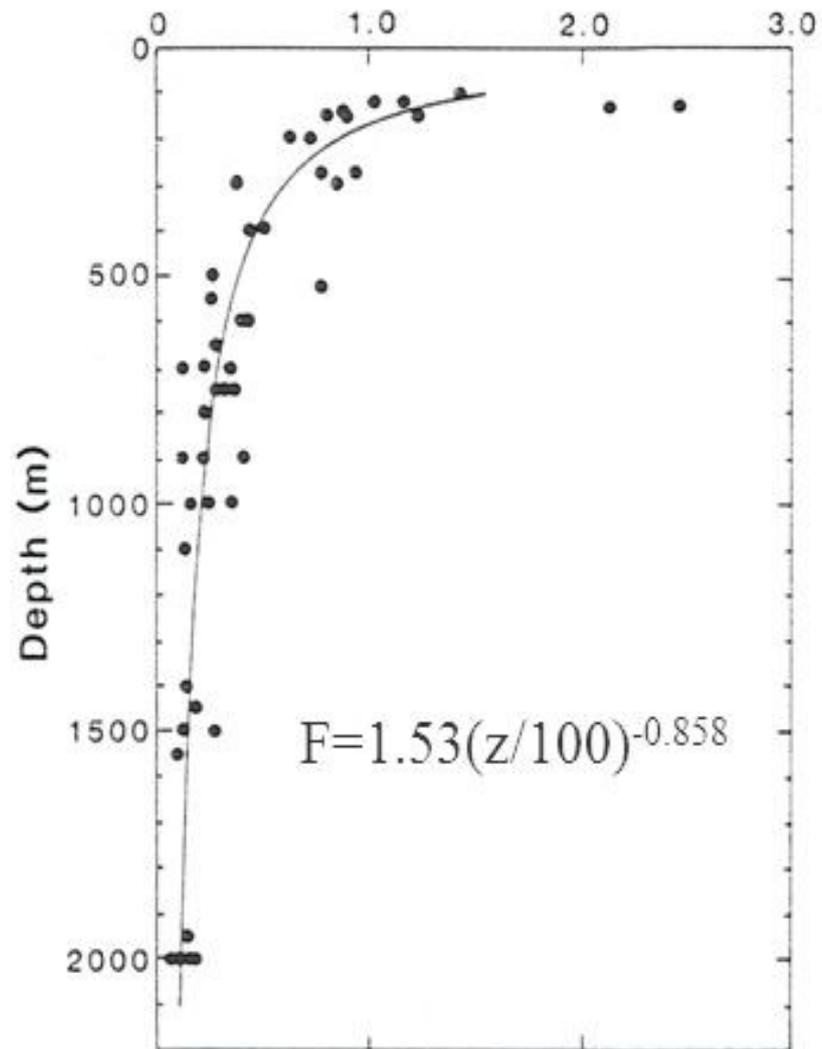
- If v and λ are constant :

$$F = vD = F(\tilde{z}) e^{-\frac{\lambda}{v}(z - \tilde{z})}$$

- If λ is constant and $v = Az$

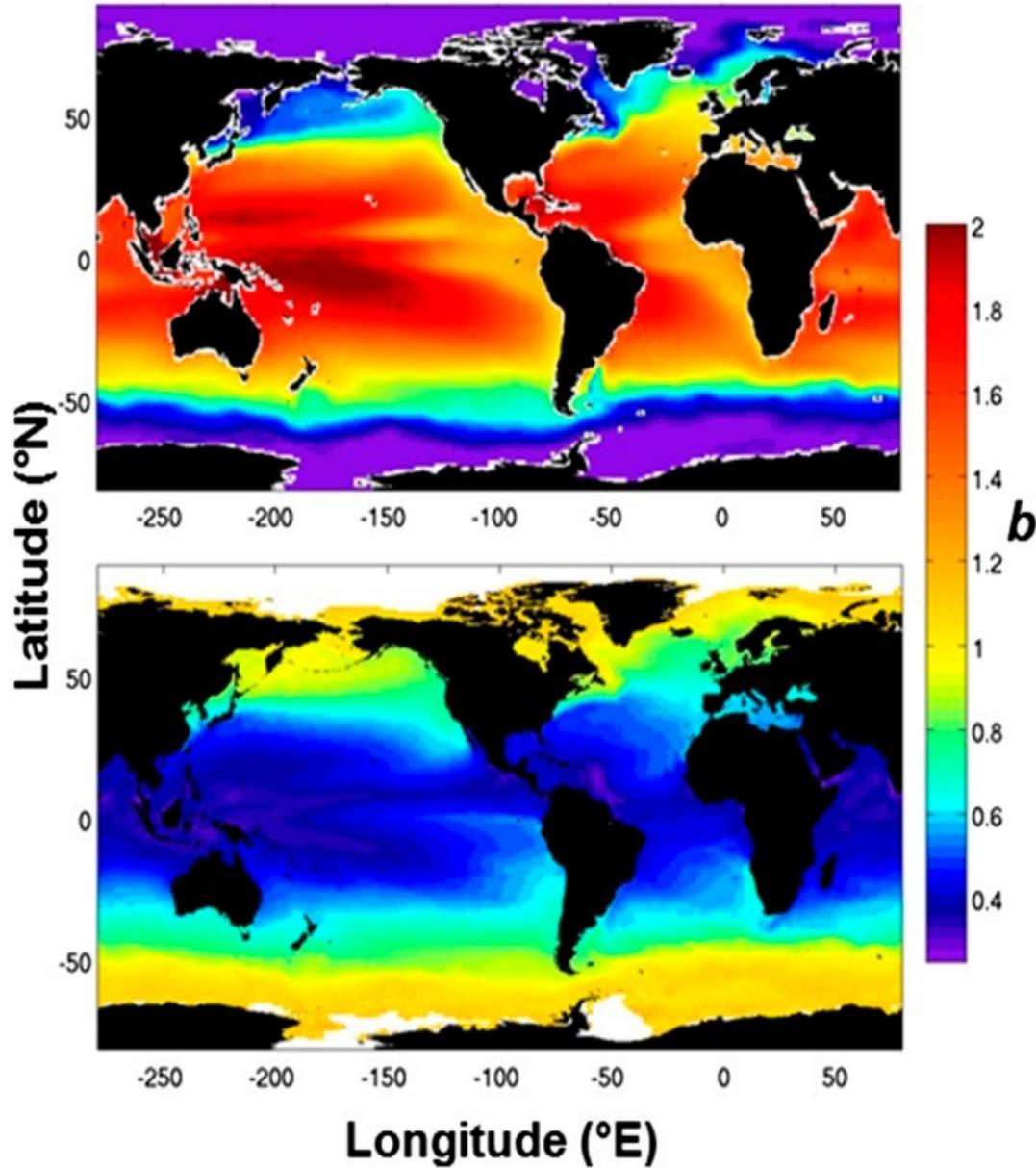
$$F = vD = F(\tilde{z}) \left(\frac{z}{\tilde{z}} \right)^{-b} \quad \text{where} \quad b = \frac{\lambda + A}{A}$$

Observed fluxes vs. modeled fluxes



Fluxes seem to be well approximated by a power law function

But far from being that simple



From Marsay et al. (2015)

A simple set of 4 equations

$$(S - P)(P) = \mu_{max} L_{NUT} L_I P - g \frac{pP}{K_z + pP + (1-p)D} Z - m_p P$$

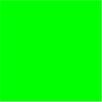
$$(S - P)(Z) = ga \frac{pP + (1-p)D}{K_z + pP + (1-p)D} Z - m_z Z^2$$

$$(S - P)(D) = g(1-a) \frac{pP + (1-p)D}{K_z + pP + (1-p)D} Z + m_z Z^2 + m_p P$$

$$-g \frac{(1-p)D}{K_z + pP + (1-p)D} Z - v \frac{\partial}{\partial Z} D - t_d D$$

$$(S - P)(N) = -m_{max} L_{NUT} L_I P + t_d D$$

Non-linear equations



Parameters needed to constrain the model

$\mu_{\max}, K_N, K_I, m_p$ phytoplankton

g, K_z, p, a, m_z zooplankton

v, t_d détritus

Difficult estimation

- lab experiments, species dependant, equilibrium state
- large variability: non constant
- agregate many processses
- inverse methods (data assimilation)
- empirical estimation

Evaluation of a model



Experiments in controlled environments



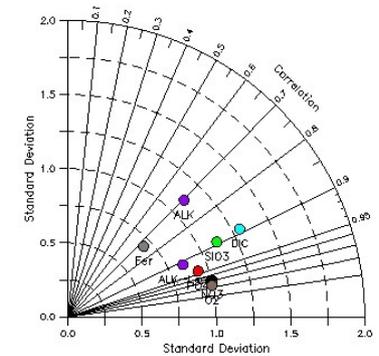
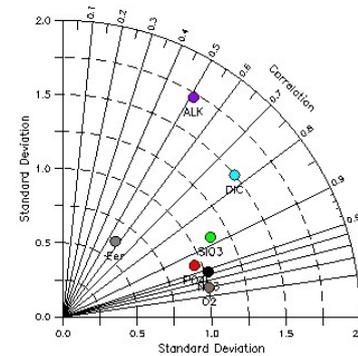
Observations

Tuning the models



```
Model Editor: AdMod - Model: Global - Mode:
shampisbio | biological parameters
-----
nuttrtc = 1 | time step frequency for biology
wbl0 = 2. | POC sinking speed
wbl0t = 2.E-7 | half saturation constant for mortality
ferat3 = 10.E-6 | Fe/C in zooplankton
wbl02 = 30. | Big particles sinking speed
niter1max = 1 | Maximum number of iterations for POC
niter2max = 1 | Maximum number of iterations for GOC
-----
shampislim | parameters for nutrient limitations
-----
concn03 = 1.E-6 | Nitrate half saturation of nanophytoplankton
concn03 = 3.E-6 | Phosphate half saturation for diatoms
concn04 = 1.E-7 | NH4 half saturation for phyto
concn04 = 3.E-7 | NH4 half saturation for diatoms
concn04 = 1.E-9 | Iron half saturation for phyto
concn04 = 3.E-9 | Iron half saturation for diatoms
concn04 = 1.E-11 | Half-saturation for Fe limitation of Bacteria
concn04 = 2.E-8 | NH4 half saturation for phyto
concn03 = 2.E-7 | Phosphate half saturation for diatoms
xsize04 = 1.E-6 | Minimum size criteria for diatoms
xsizephy = 1.E-6 | Minimum size criteria for phyto
xsize04 = 3.0 | Size ratio for nanophytoplankton
xsize04 = 3.0 | Size ratio for diatoms
xsl1 = 2.E-6 | half saturation constant for Si/C uptake
xsl2 = 20E-6 | half saturation constant for Si/C
xldoc = 437.E-6 | half-saturation constant of DOC remineralization
qfella = 7.E-6 | Optimal quota of phyto
qfella = 7.E-6 | Optimal quota of diatoms
ccabr = 0.3 | mean rain ratio
-----
36,1 10%
```

10, 20, 100x, ...

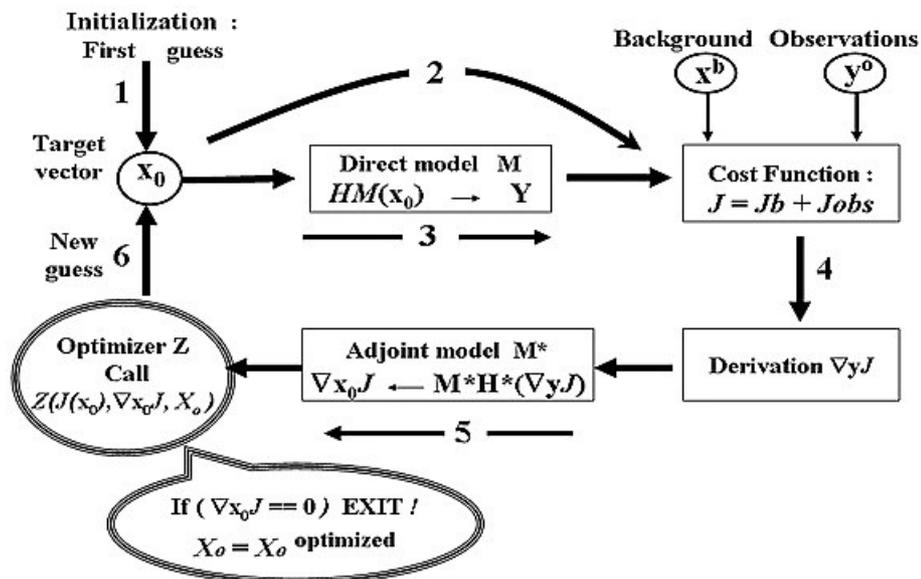


- Can be very painful
- The more complex the model is, the more difficult it is

We learn a lot on the model dynamics/behavior

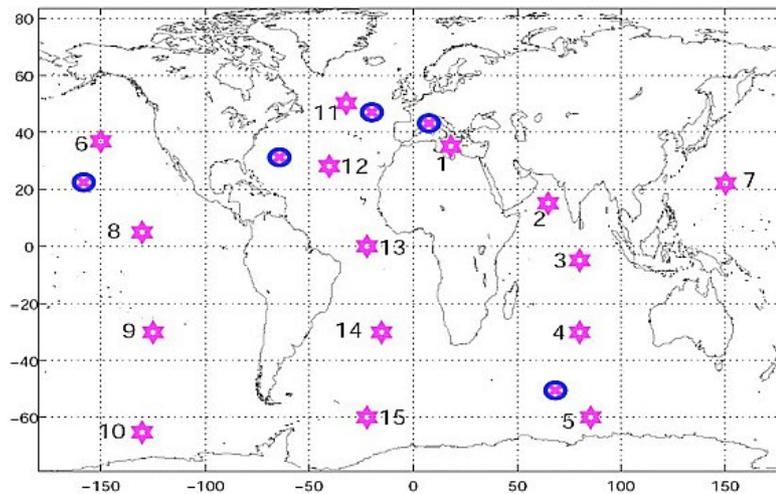
Optimizing the parameters : Assimilation

Variational assimilation

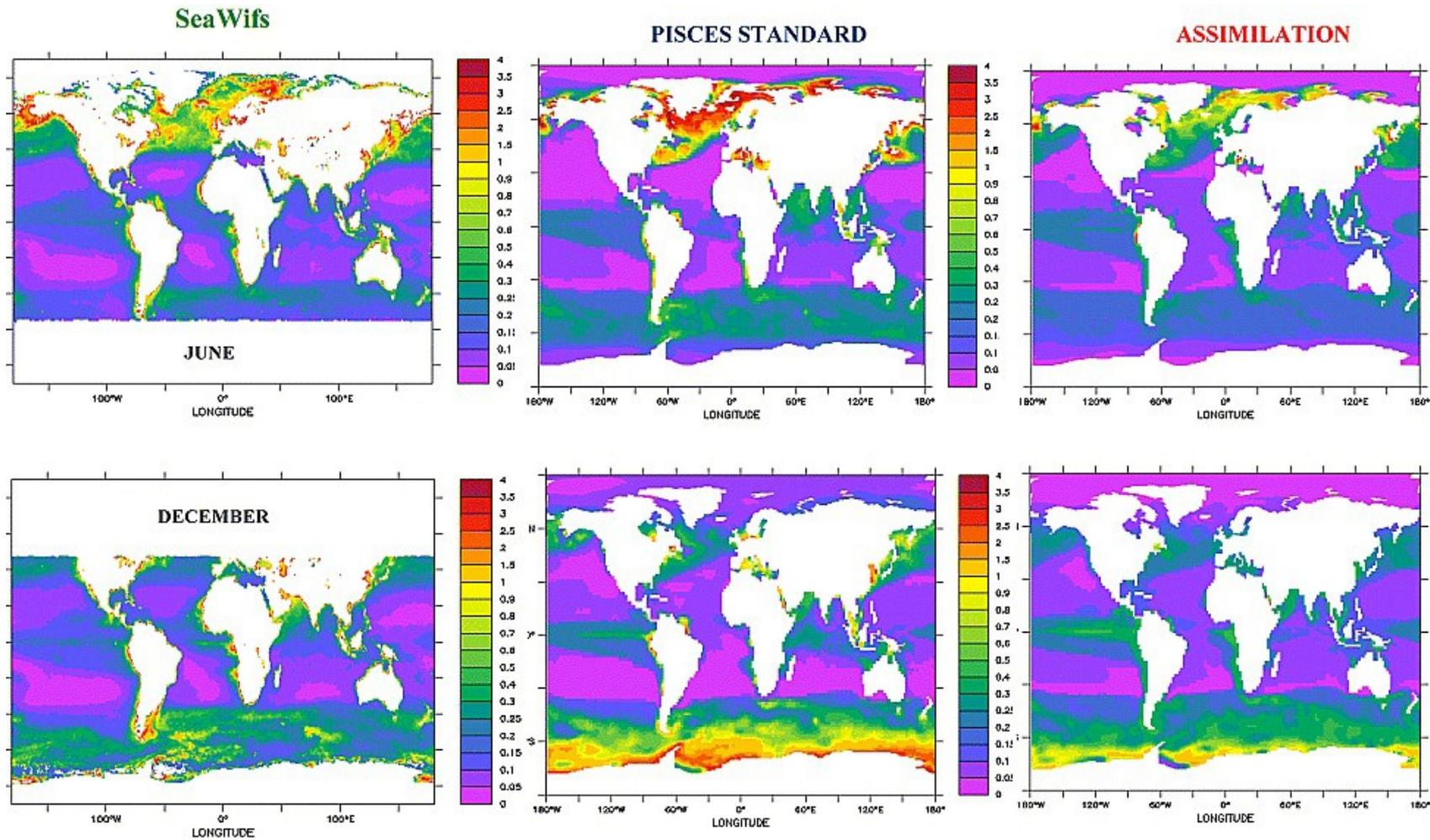


Example: Simultaneous assimilation at 5 1-D stations: Chla, NO3, POC et Si

45 optimized parameters



Optimizing parameters



Kane et al., 2011

► Can be difficult (not the universal cure) !

- Are stations representative of the system?
- What should be done with non assimilated variables?
- Some parameters may not be well constrained by available data

Models Intercomparison Projects (MIPs)

Motivation of these exercises

- Evaluate and intercompare the participating models
- Identify the converging and diverging behaviors and stimulate the model developments
- Estimate the uncertainties, for instance in projections

Ocean biogeochemistry: 1st exercise started in 1995

- 4 participating groups : IPSL, GFDL, MPIM, Hadley

Since, many projects have been organized:

- Carbon cycle (ocean models only) : OCMIP-2, OCMIP-3, OMIP
- Iron cycle : FeMIP
- Carbon cycle (Earth System Models) : CMIP, C4MIP
- Marine ecosystems : FISHMIP