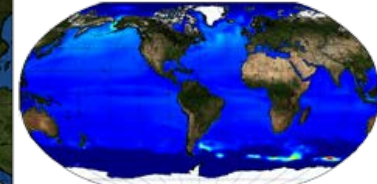
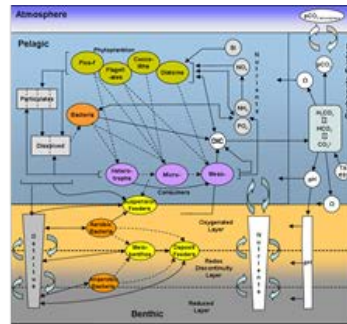
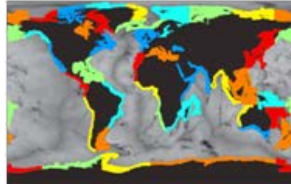


## Marine Matters



# Towards next generation of Marine ecosystem models

Icarus Allen,

Y Artioli, J Blackford, J Bruggeman, M Butenschon, J Clark, L de Mora, L Polimene, S Salliey.

## Why Model Marine Ecosystems

- Improve understanding of the regulation of key ecosystem services
- Integrate improved knowledge in models
- Apply models to potential management solutions

**1. Described**  
by ecosystem model

### Marine Ecosystem

#### *Components*

Habitats

Functional diversity

#### *Processes*

Production

Decomposition

Foodwebs

Ecological interactions

**2. Emergent properties**  
of ecosystem model

### Intermediate Services

#### *Supporting*

Primary production

Nutrient cycling

Oxygen

#### *Regulating*

Biological control

Carbon sequestration

**3. Informed by**  
ecosystem model

### Final Services

#### *Provisioning*

Fish & Shellfish

Seaweed

#### *Regulating*

Climate regulation

#### *Cultural*

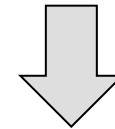
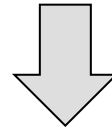
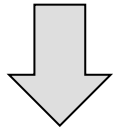
Healthy environment

# Marine ecosystem model

Parameters

Equations

Emergence

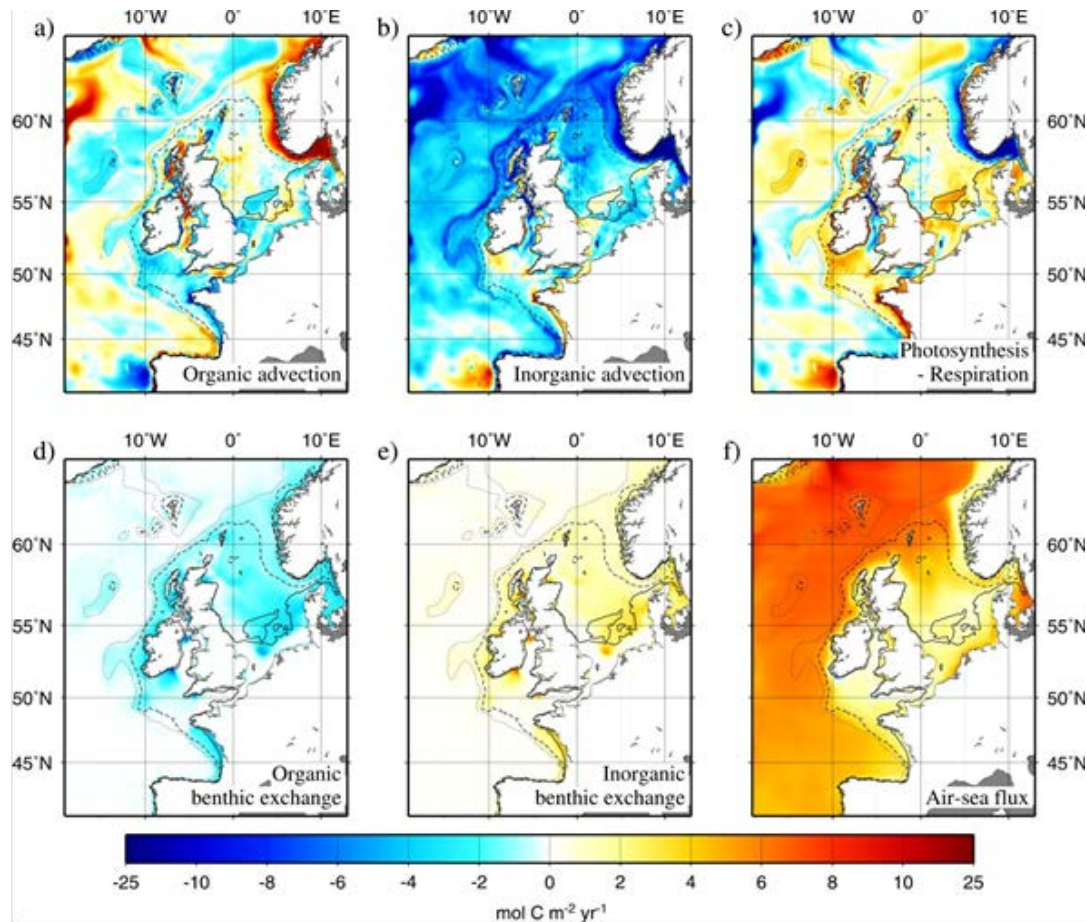


Molecular biol.  
Physiology  
genomics

Biogeography  
Physiology

Biogeochemistry  
Ecology  
?

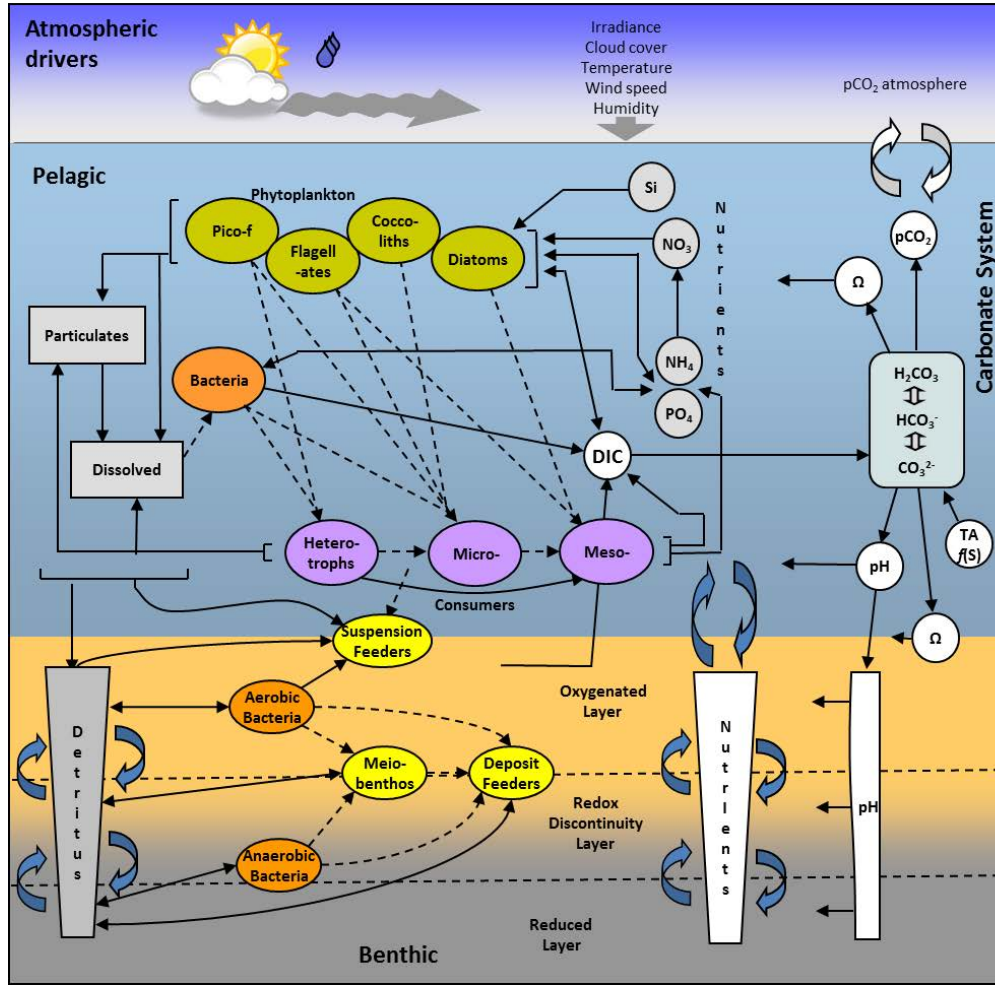
“traditional models are static structures which are not able to evolve under changing environmental conditions”  
Capacity to evolve emergent properties  
(Hood et al., *Oceanography*, 2007)



*Shelf Carbon Budget  
Wakelin et al 2012*

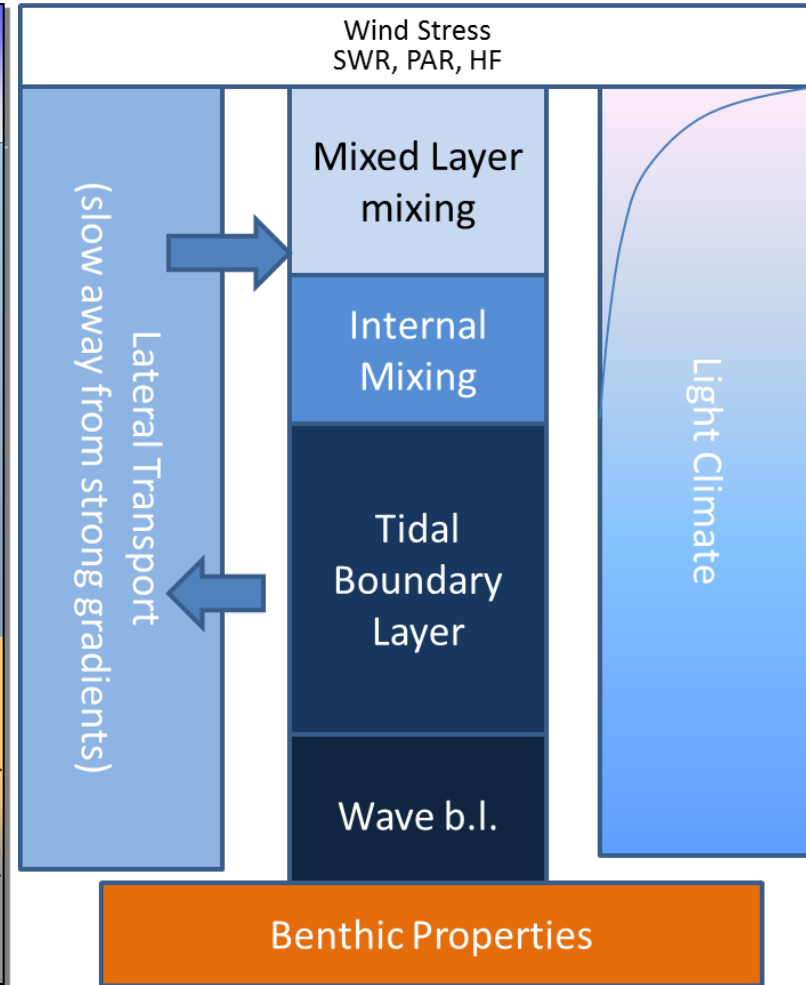
The overarching scientific goal is to enhance our capacity to assess the controls on biogeochemical cycling and hence to quantify with uncertainties the budgets of carbon, nitrogen, phosphorous and silicon including their response to climate, natural variability and anthropogenic stress.

## Biogeochemical Processes



European Regional Seas Ecosystem Model  
ERSEM

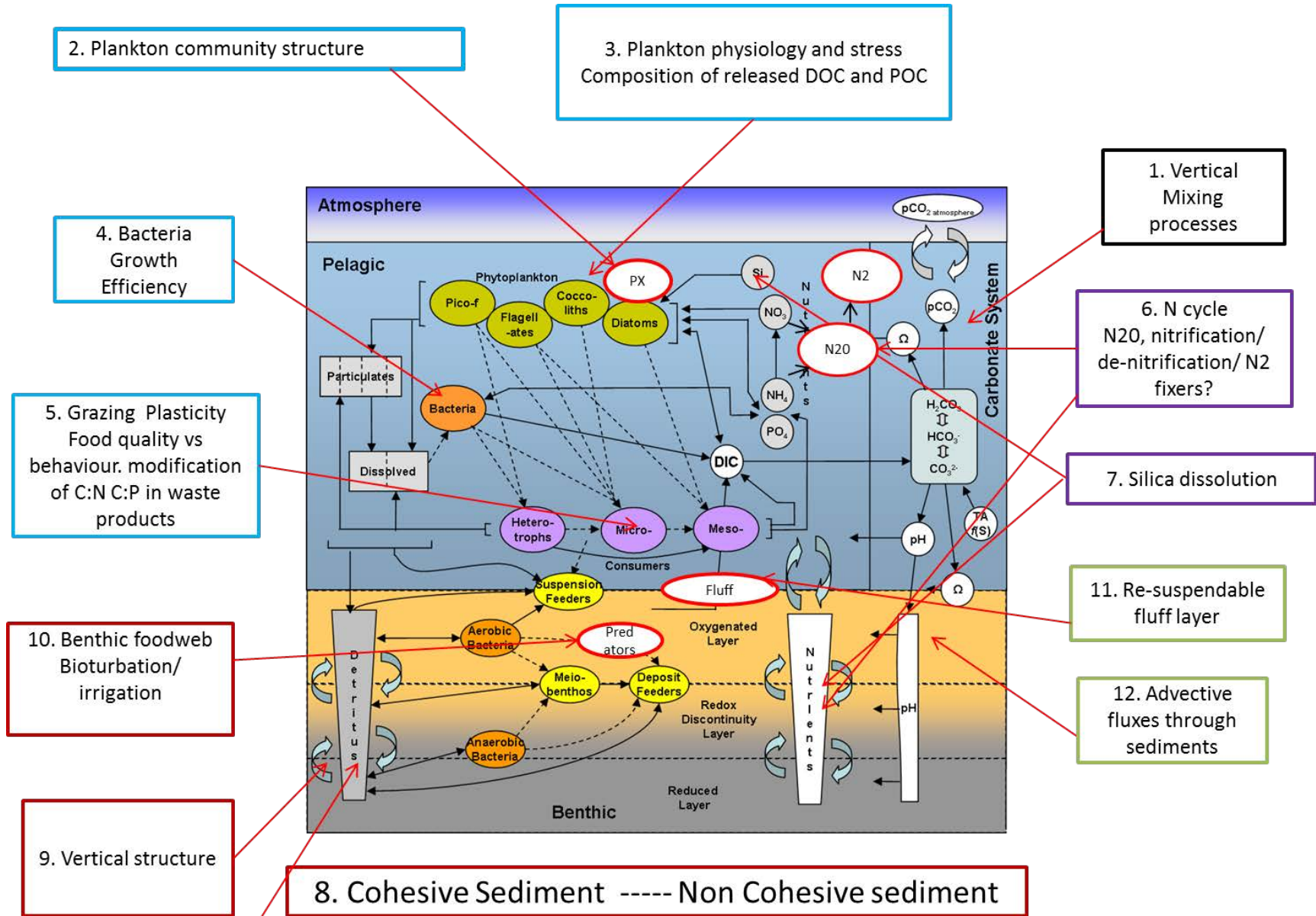
## Physical Processes



3D: NEMO- Shelf  
1D: GOTM

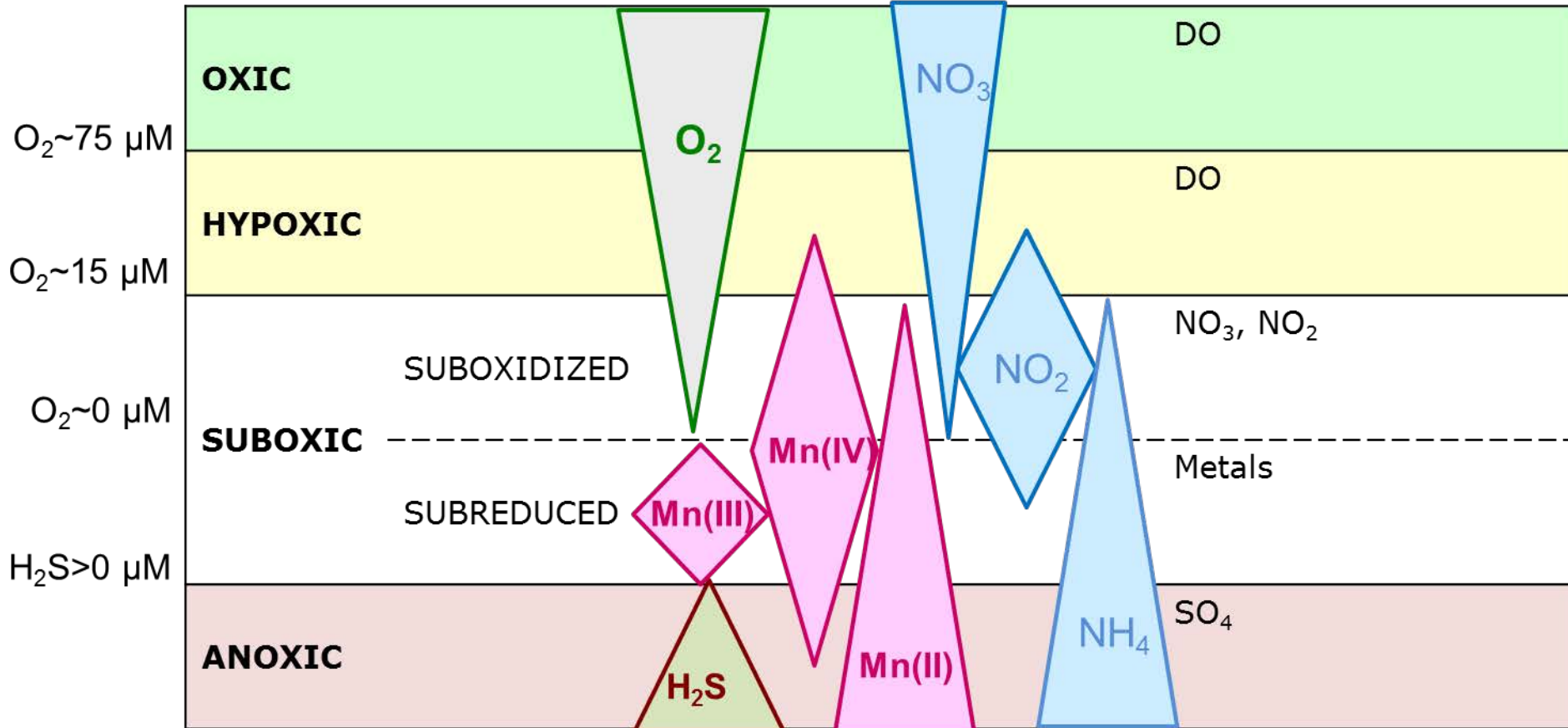


# Summary of SSB Model Developments

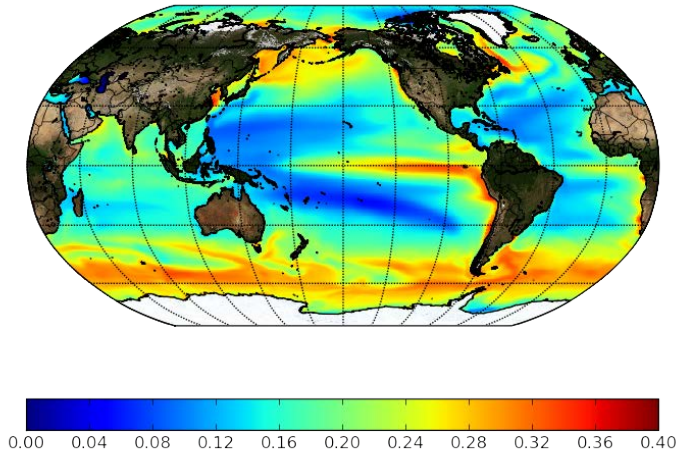


# Redox conditions

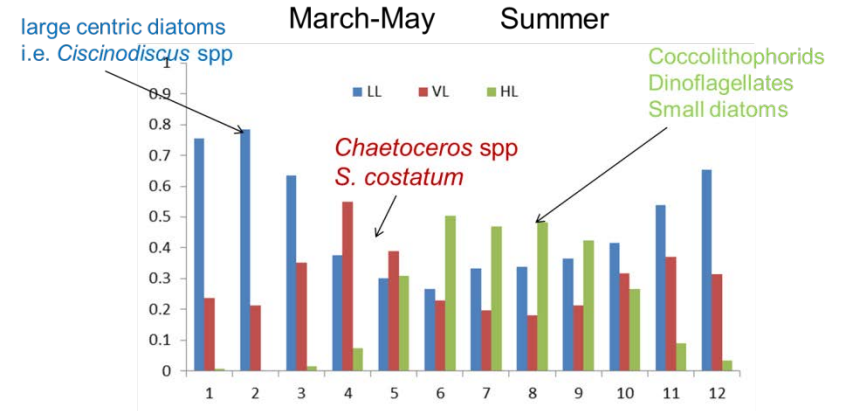
Dominating oxidizer:



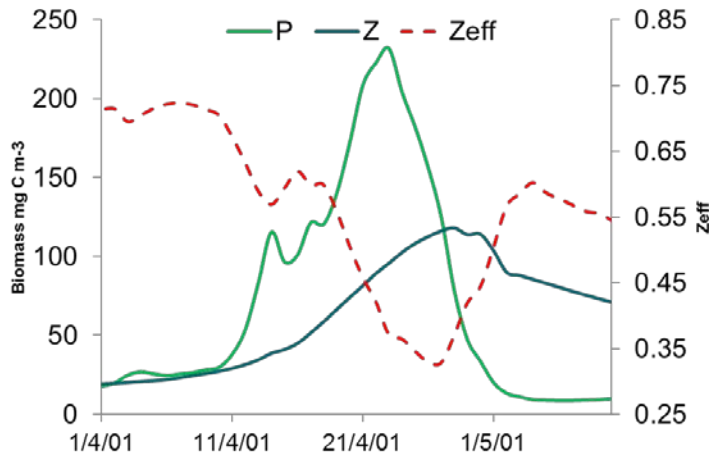
## Bacterial Growth Efficiency



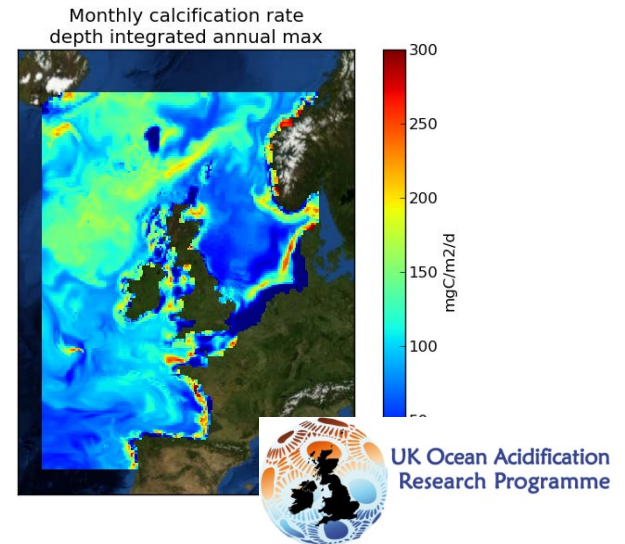
## Phytoplankton Succession



## Stoichiometry Modulation of Predation



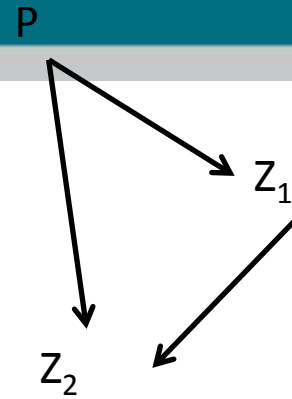
## Calcification



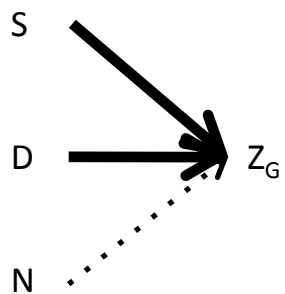
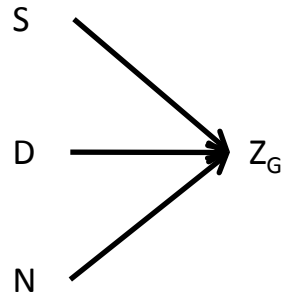


# Analysis of Grazing interactions

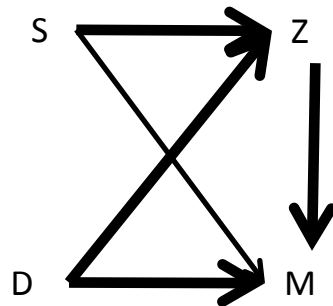
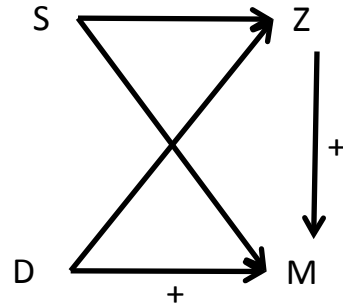
Sailley et al 2013



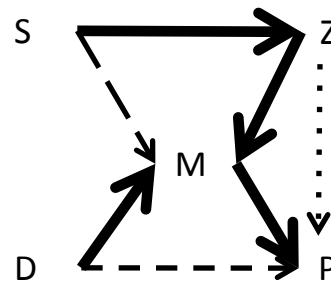
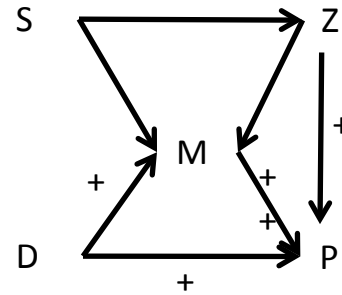
CCSM-BEC



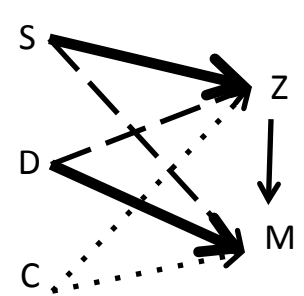
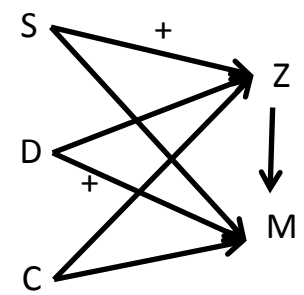
PISCES



NEMURO

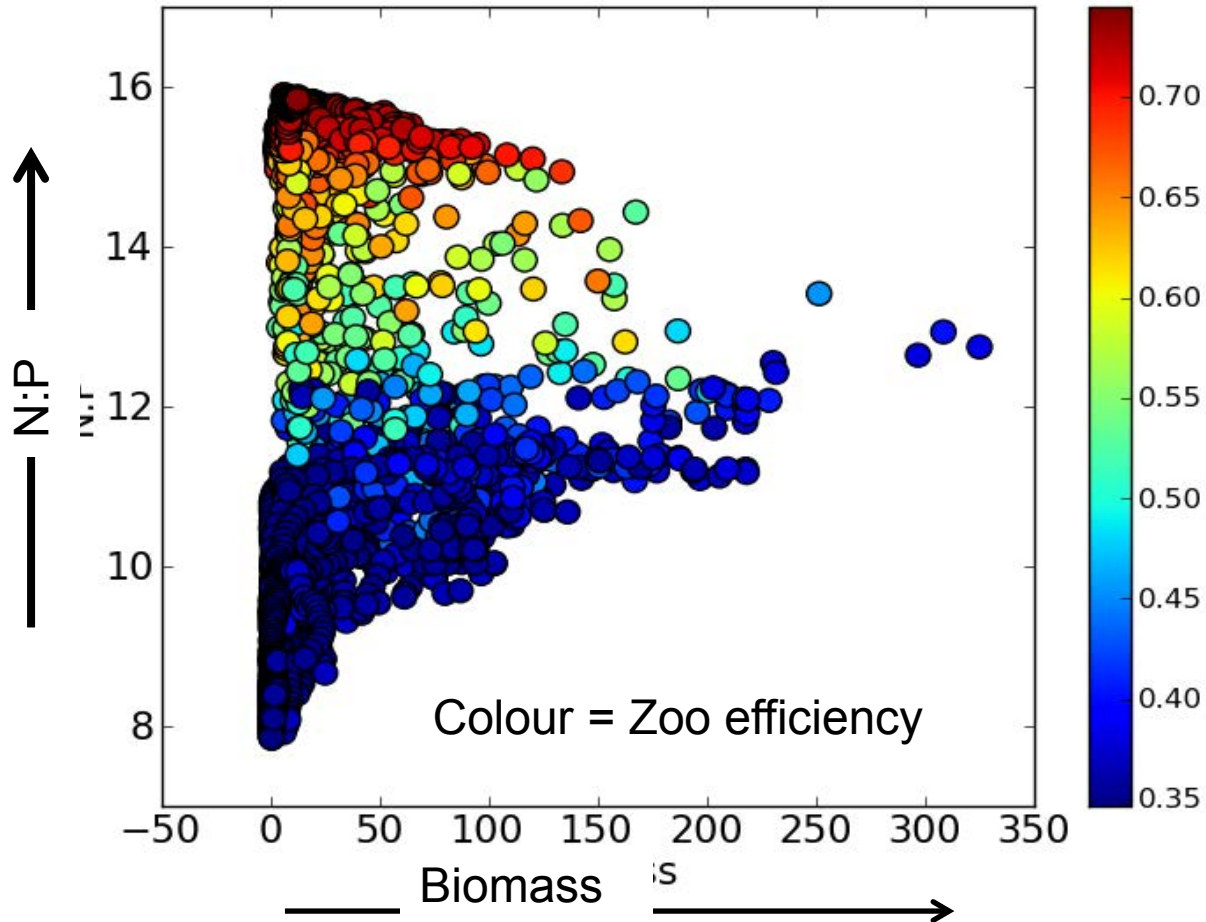


PlankTOM5





Zooplankton efficiency is dependent on the nutritional quality (nutrient content) of the prey



Preliminary results  
(GOTM-ERSEM at L4)

(Mitra et al, 2008;  
Polimene et al., in prep)

## Benthic

Benthic model component is the biggest challenge

- Historically a poor relation to pelagic / ocean modelling in terms of effort
- Current ERSEM benthic models – enabled ~200 pubs
- Computationally efficient at expense of accessibility – black box
- Need to open (Pandora's?) box, no longer fit for purpose
- Important?: e.g. 90% shelf calcification on sea floor

Opportunity to restructure, re-conceptualise

Flexible basic structure that enables SSB, OA, CCS and Ecosystems work

Break down conceptual barriers between pelagic and benthic

Include multiple sediment types

Sands: Advective physics

Physical burial

Detritus resuspension

Revise functional groups

Biogeochemical functionality

Improve bioturbation

Trophic transfer

Improve redox chemistry

Improve inorganic carbon / carbonate system, inc alkalinity

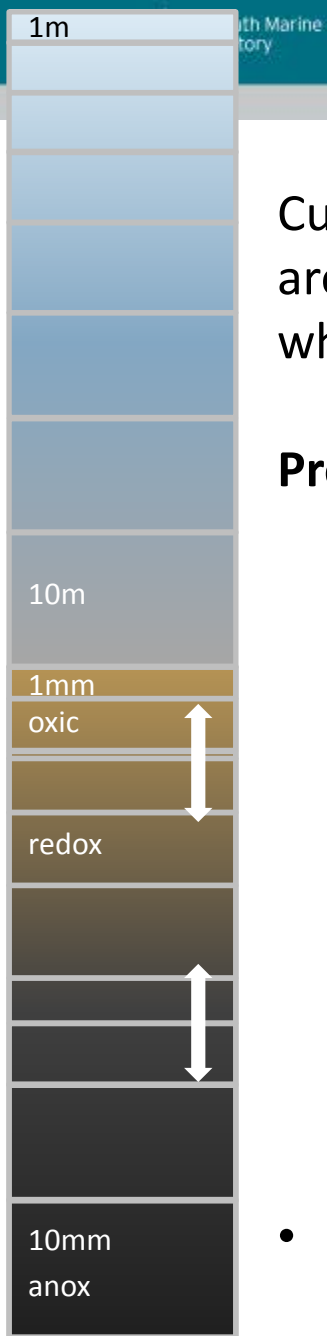
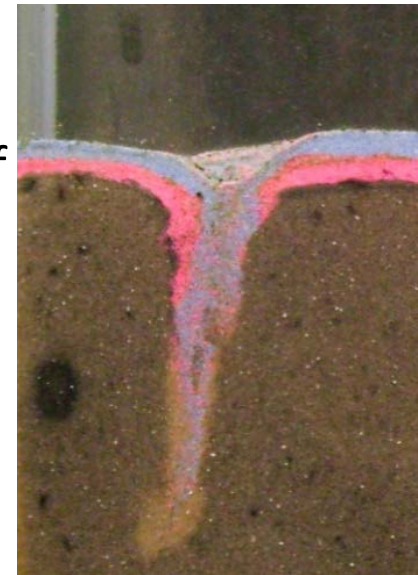
# Potential Physical Structure

Currently have a three layer implicit model: Standard chemical profiles are fitted to the model chemical concentrations ( $O_2$ ,  $NO_3$ ,  $NH_4$ ), from which the depths of 3 layers oxic – redox – anoxic are derived.

## Proposal: convert to Z level configuration.

- Perfect for physical advection
- Good for characterisation of sediments (variable porosity)
- Similar conceptually to pelagic
- Need care in choosing z coordinate (mm scale structures at surface)
- Problematic in dealing with biota that live across several layers or create intrusions of surface chemical environment into deeper layers

- Sub grid scale spatial variability?



10mm

# Which Functional types?

- Need to be right for chemistry / bioturbation / trophic transfer
- Limited in numbers (competitive exclusion)

## Current minimum implementation

Suspension feeders

Deposit feeders

Meiobenthos

Aerobic bacteria

Anaerobic bacteria

Surficial Bioturbators  
Surficial Bulldozers  
Surficial Suspension Feeders

Intermediate Bioturbators

Head-up Feeders  
Head-down Feeders

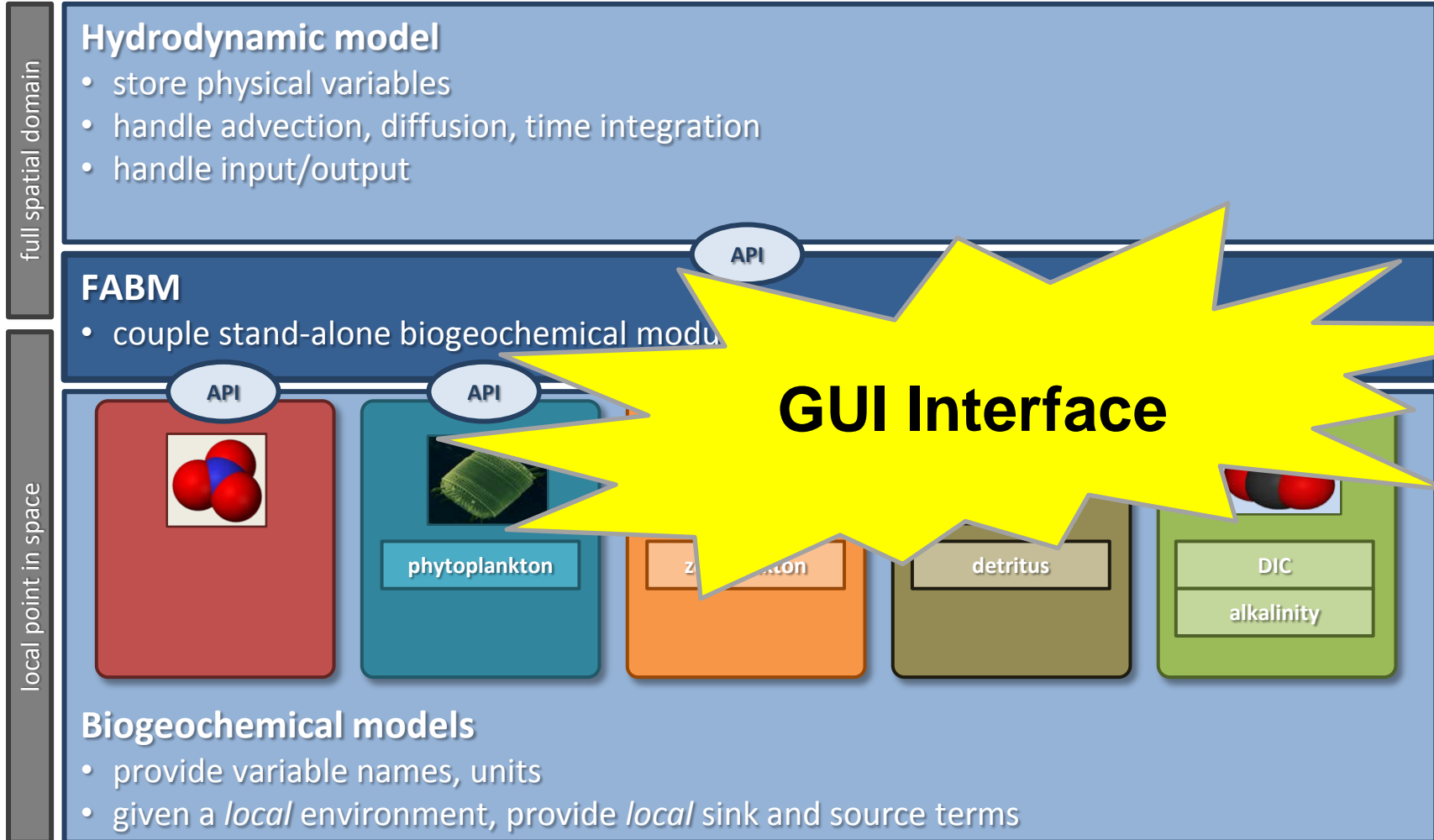
Biodiffusers  
Regenerators

Meiobenthic predators  
Microbial feeders

Deposit feeders  
Cyanobacter

Diatoms  
Other microphytobenthos  
Aerobes  
Sulphur oxidisers





<http://www.shelfseasmodelling.org/>

• **Currently configured:**

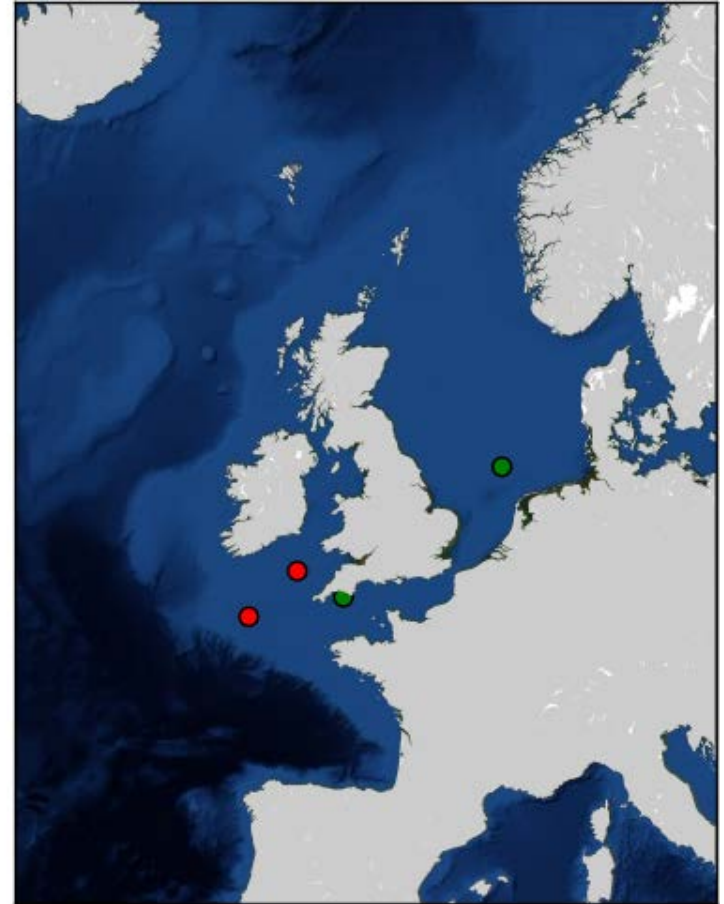
- Station L4 (4°13 W, 50°15 N)
- Oyster Grounds (4°02 E, 54°25 N)

• **Future configurations:**

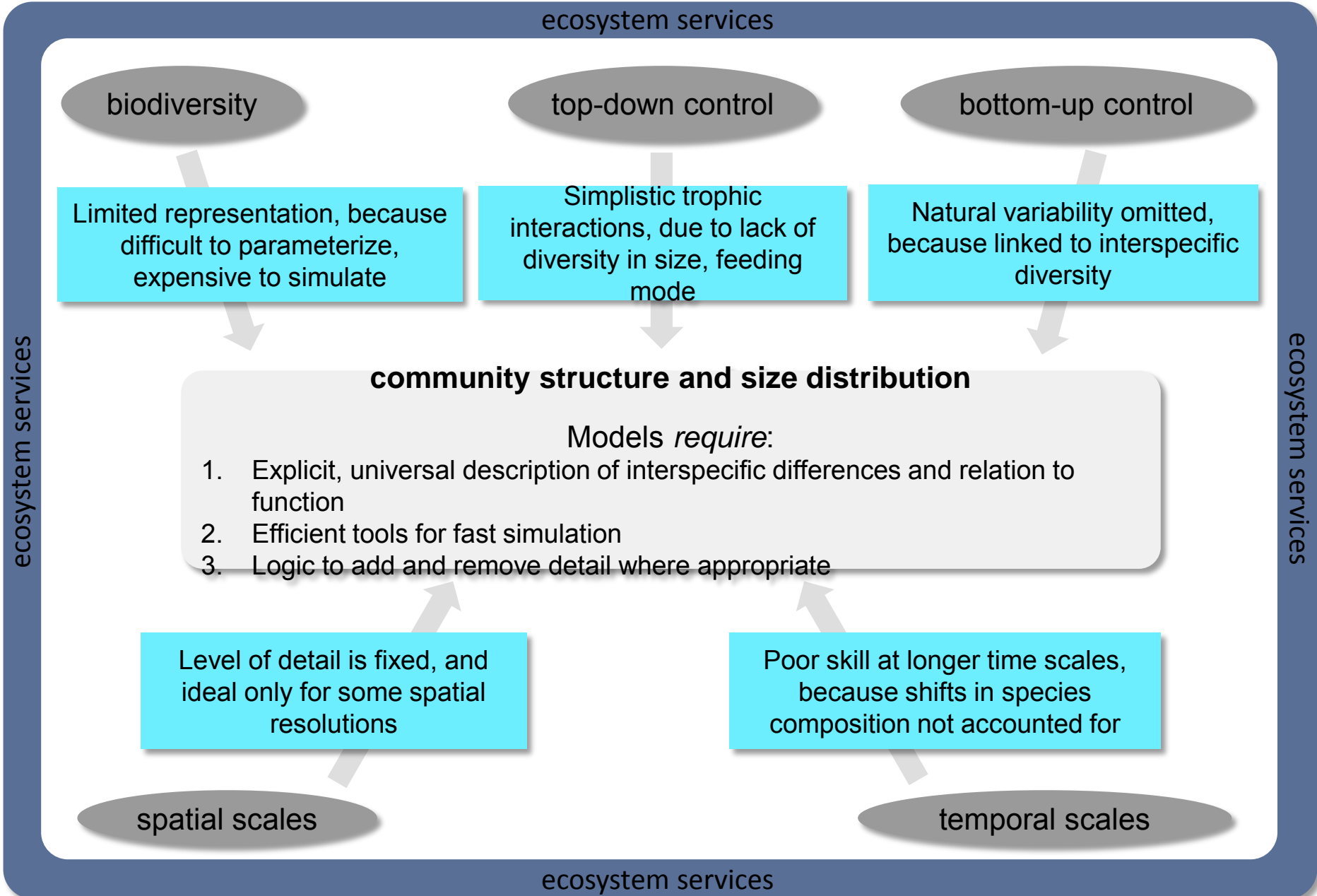
- Celtic Deep (4°80 W, 51°14 N)
- Celtic Sea - new (9°00 W, 49°50 N)

• **Model evaluation:**

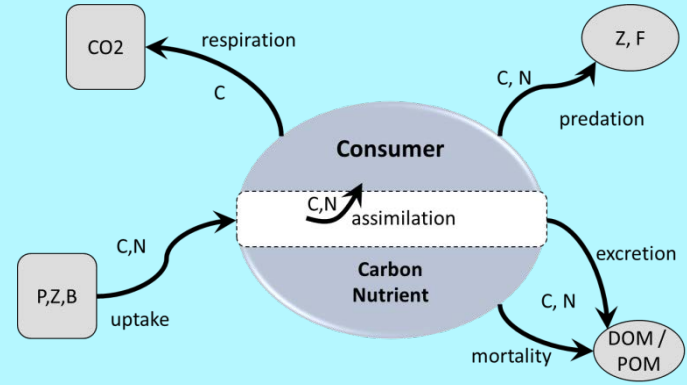
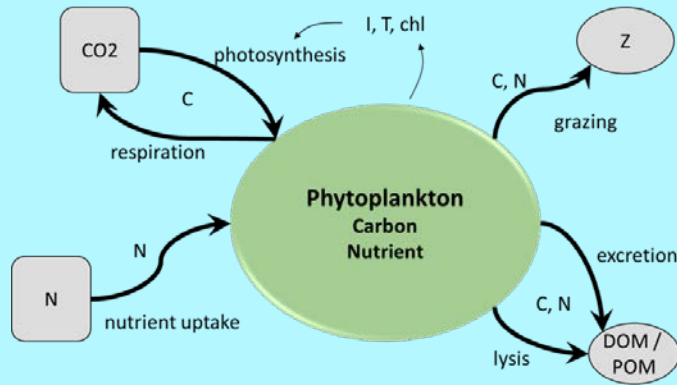
- OPEC benchmarking tool
  - Basic univariate statistics for assessing model skill.
  - Taylor & Target diagrams.
  - Uses current and new data to be collected during the course of the SSB project.



- What are the relative roles of **top down and bottom up control** processes and to what extent do impacts of environmental changes cascade through marine food webs and affect ecosystem services?
- As many processes are inherently scale-dependent, and **scale-dependence** is poorly understood, what are the most appropriate approaches to quantify the large-scale impacts on ecosystem services of changes at small spatial scales (e.g. marine conservation zones); and *vice versa*?
- How does **functional diversity affect the way marine food webs regulate ecosystem services**? This is potentially important because there is growing evidence that the loss of biodiversity from marine ecosystems can adversely impact ecosystem function.

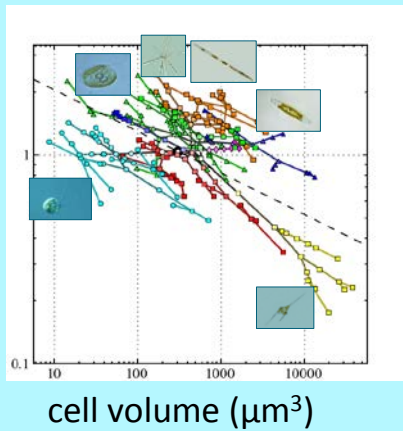


## 1. Define *standard organism*

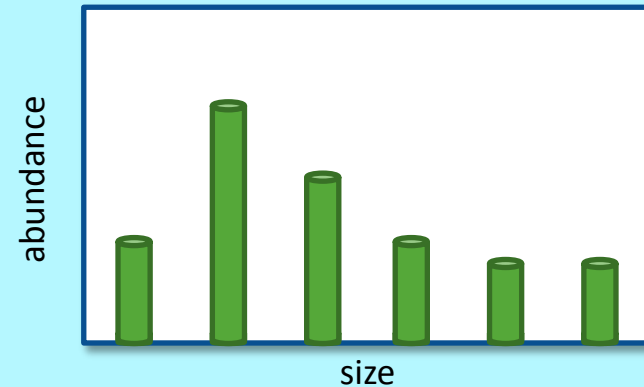


## 2. Select *traits*, link to function via *trade-offs*

maximum growth rate ( $d^{-1}$ )

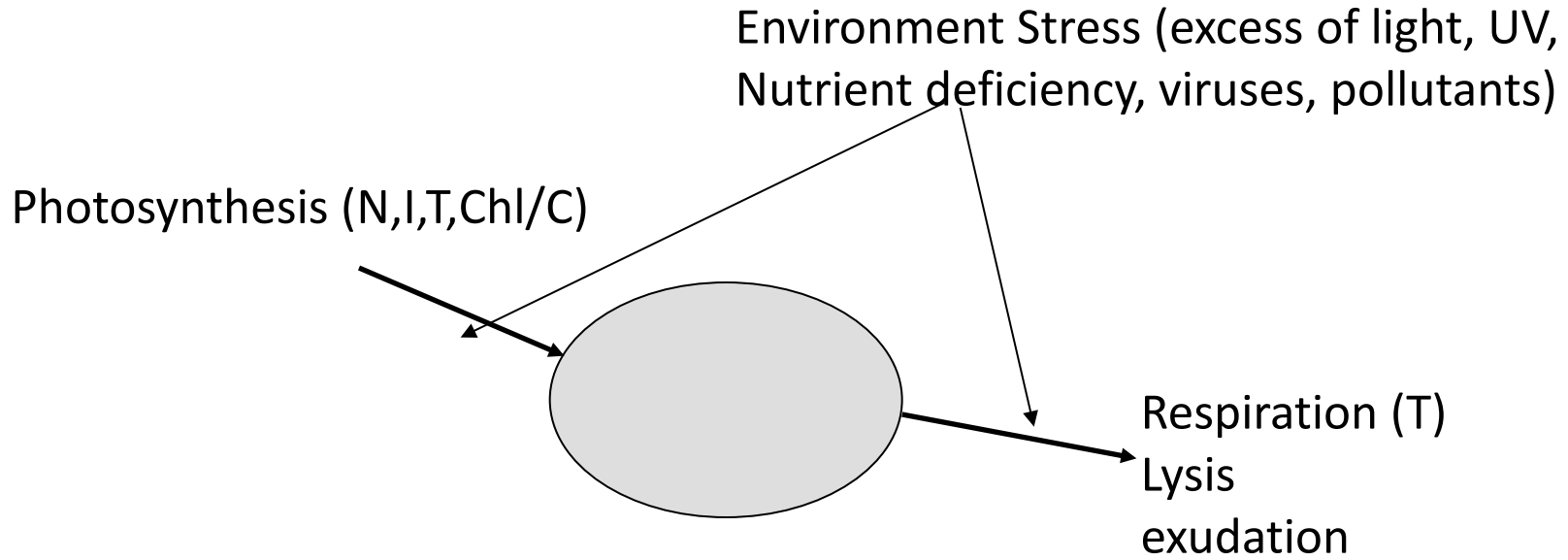


## 3. Community *emerges* from random initial assemblages



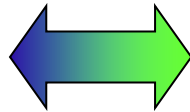


In “traditional” phytoplankton models, the cell is a sort of “black box” with an income and outcome of carbon.

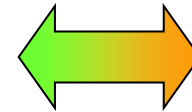


No mechanistic description of what happens inside the cell when exposed to stress  
 In other words, there is no link between physiology, biogeochemistry and ecology

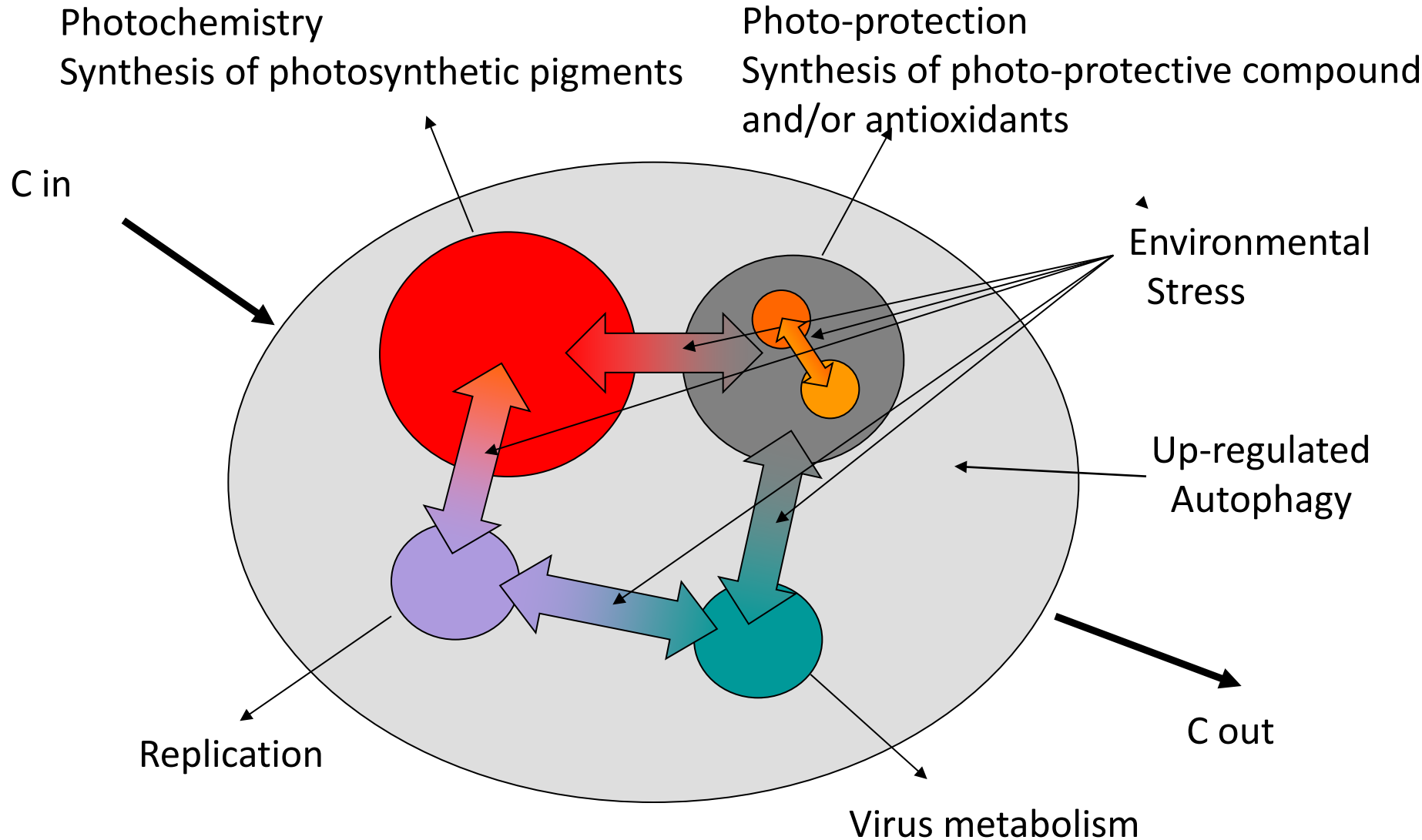
**Physiology**



**Biogeochemistry**



**Ecology**

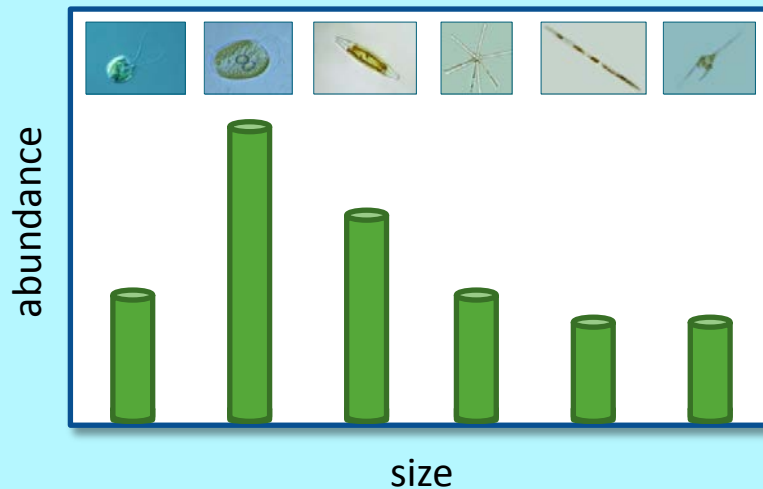


Development of new methods enabling

- Calibration and sensitivity studies
- Investigation of optimal level of detail
- High-resolution 3D simulation

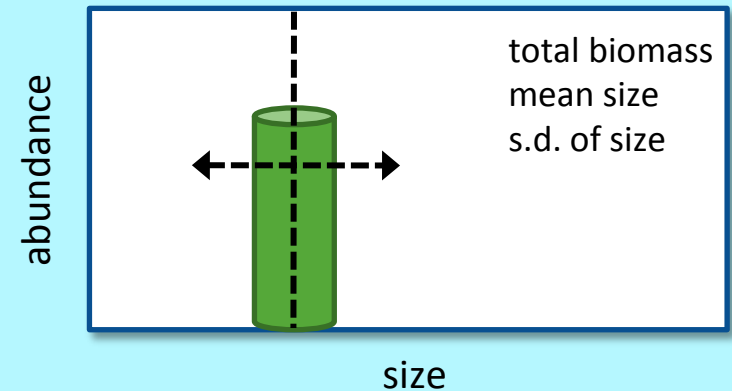
## Full model – $N$ species

DivERSEM



## Adaptive dynamics

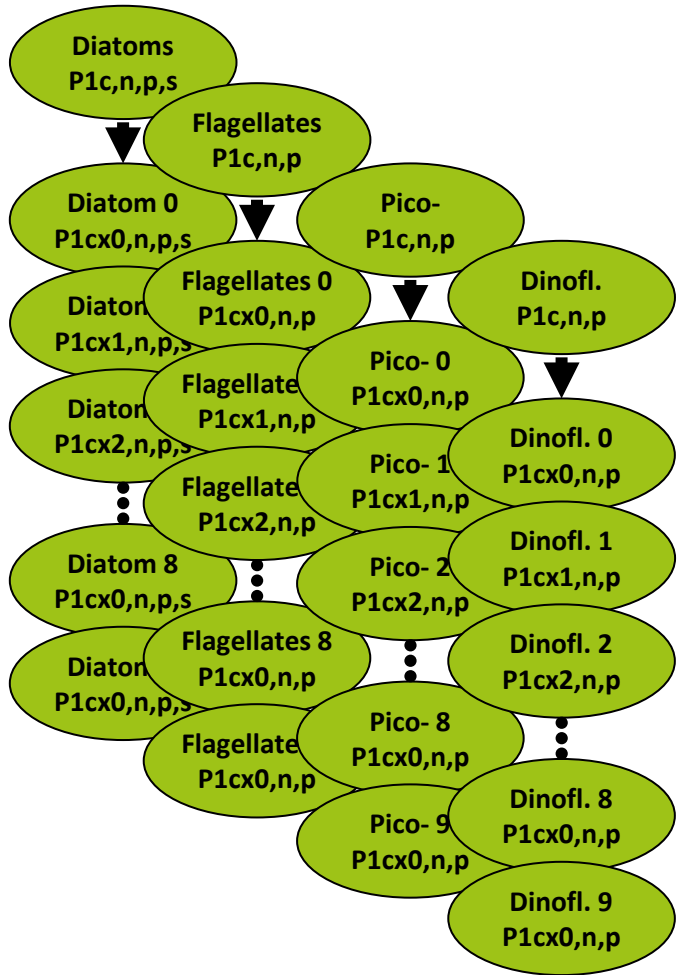
Summarize in terms of aggregate statistics



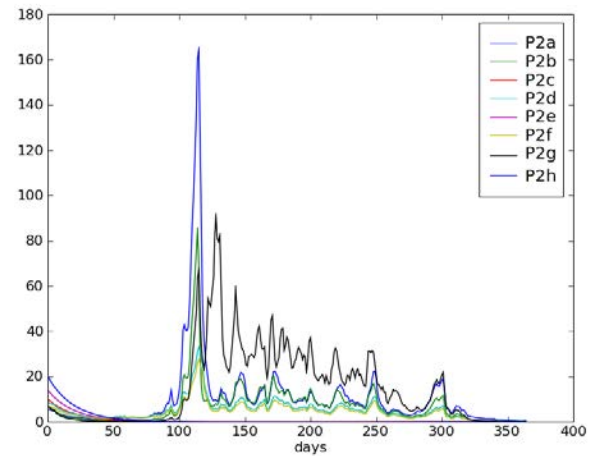
## Optimized advection

- Mixing schema (1 master variable,  $N-1$  subservient variables)
- Schema that scale better with increasing variable number

# Darwinian ERSEM



- P2a
- P2b
- ~~P2c~~
- P2d
- ~~P2e~~
- P2f
- P2g
- ~~P2h~~

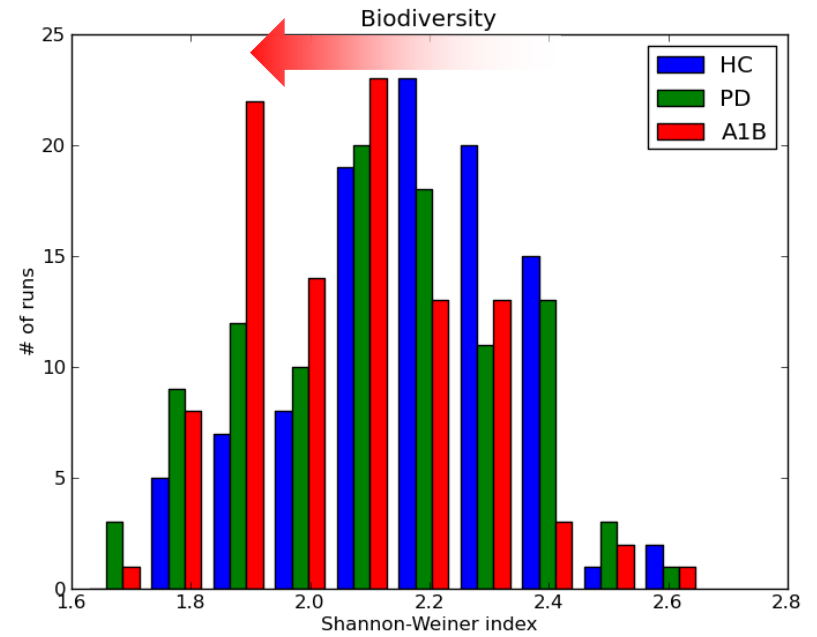
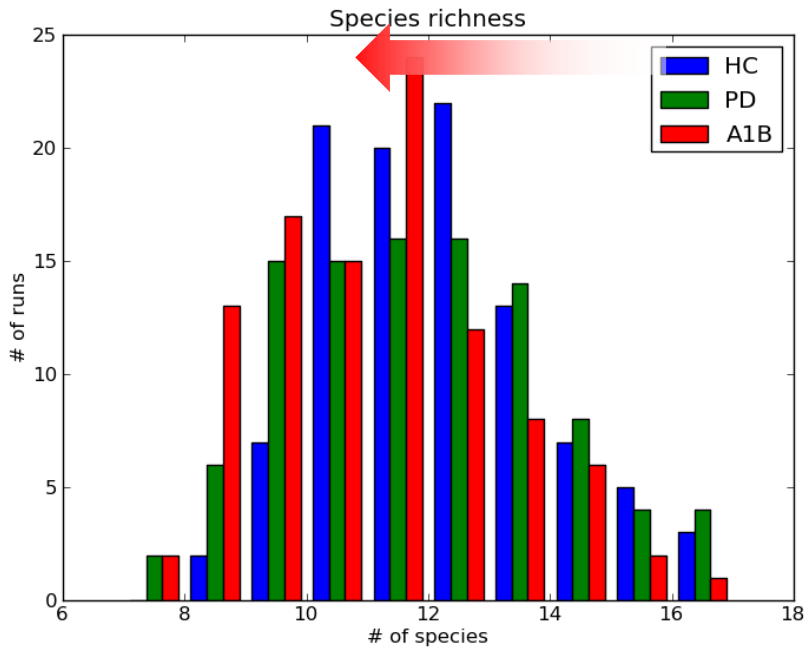


X 5 years

- P2a
- P2b
- 
- P2d
- 
- P2f
- P2g
- P2h



# Biodiversity



Shannon index calculated using biomass

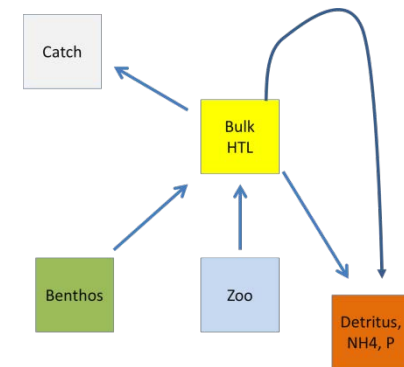
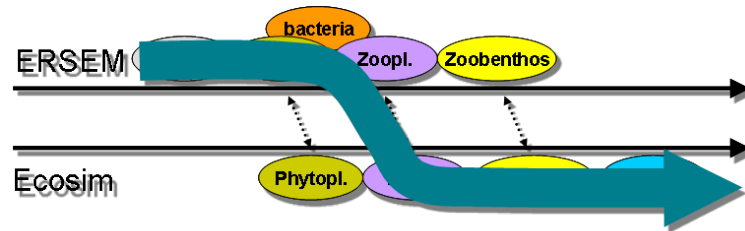


## 1. Top closure ERSEM

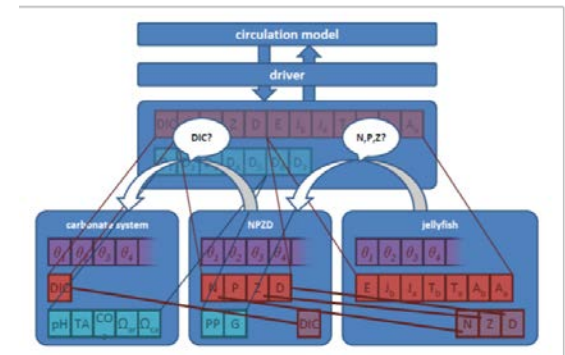
- Density dependent mortality
- Bulk HTL model (see right)
- Two way coupled dynamic size spectra (Phase II)

## 2. Coupling to HTL models

- Predation fields from offline model /data
- 2 way coupling to HTL model



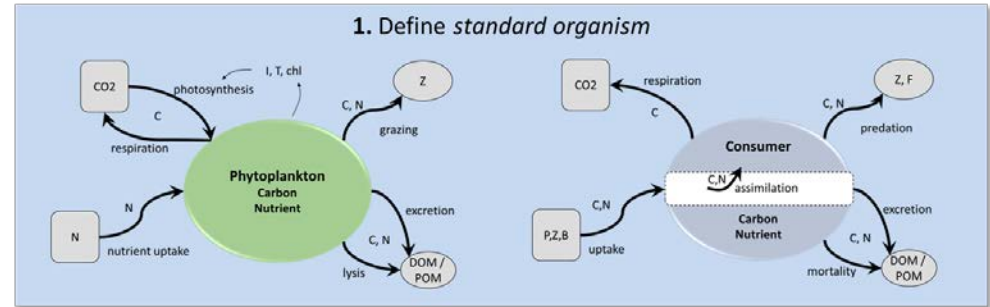
Generic Coupler



*"a thin layer of code for communication and data exchange, enveloped by explicit programming interfaces through which a physical host and any number biogeochemical models can pass information"*

## 1. Standard Organisms:

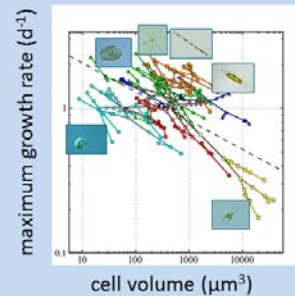
- Conceptual frameworks,
- Allometric and metabolic scaling rules to simplify the parameterisations.
- Trophic interactions.



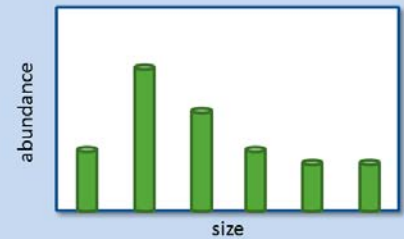
## 2. Traits:

- trait definition e.g. diatoms, macroalgae, zooplankton and macrobenthos,
- feeding strategies,
- trade offs,
- trophic interactions,
- mortality.

### 2. Select *traits*, link to function through *trade-offs*



### 3. Community *emerges* from random initial assemblages



## 3. Diversity:

- Parameter ranges

## 1. Bulk Properties

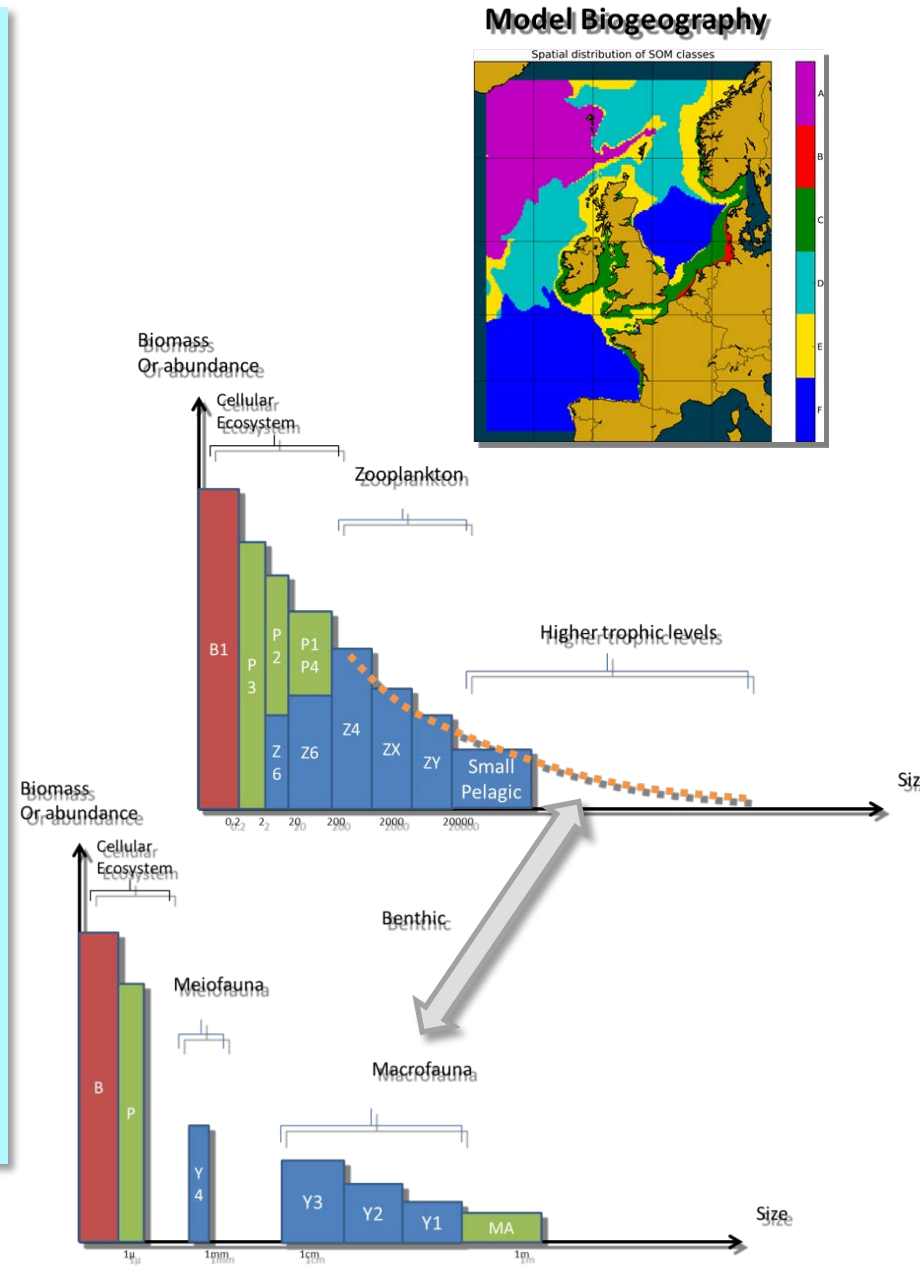
- Habitats (T, S, O<sub>2</sub>, pH)
- Biogeochemistry (N, P, Si)
- Chlorophyll, PP, SP
- Zooplankton and benthos
- Trait based biomass

## 2. Scaling relationships

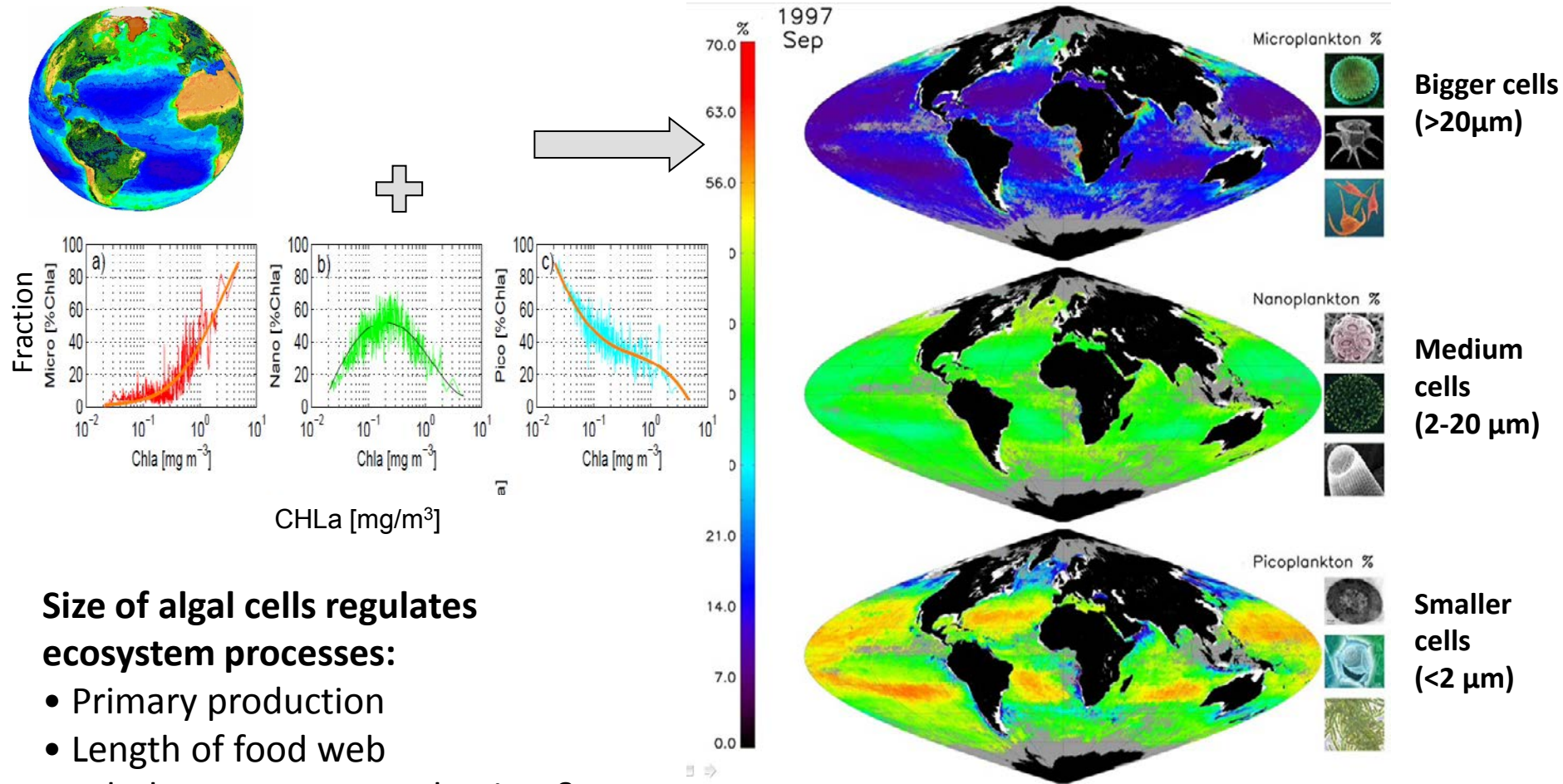
- Size spectra
- Biogeographic
- Trophic level relationships
- Trophic transfer
- Connectivity
- Diversity

## 3. Expert knowledge

- Is the model behaviour plausible?



## Phytoplankton Community Structure



### Size of algal cells regulates ecosystem processes:

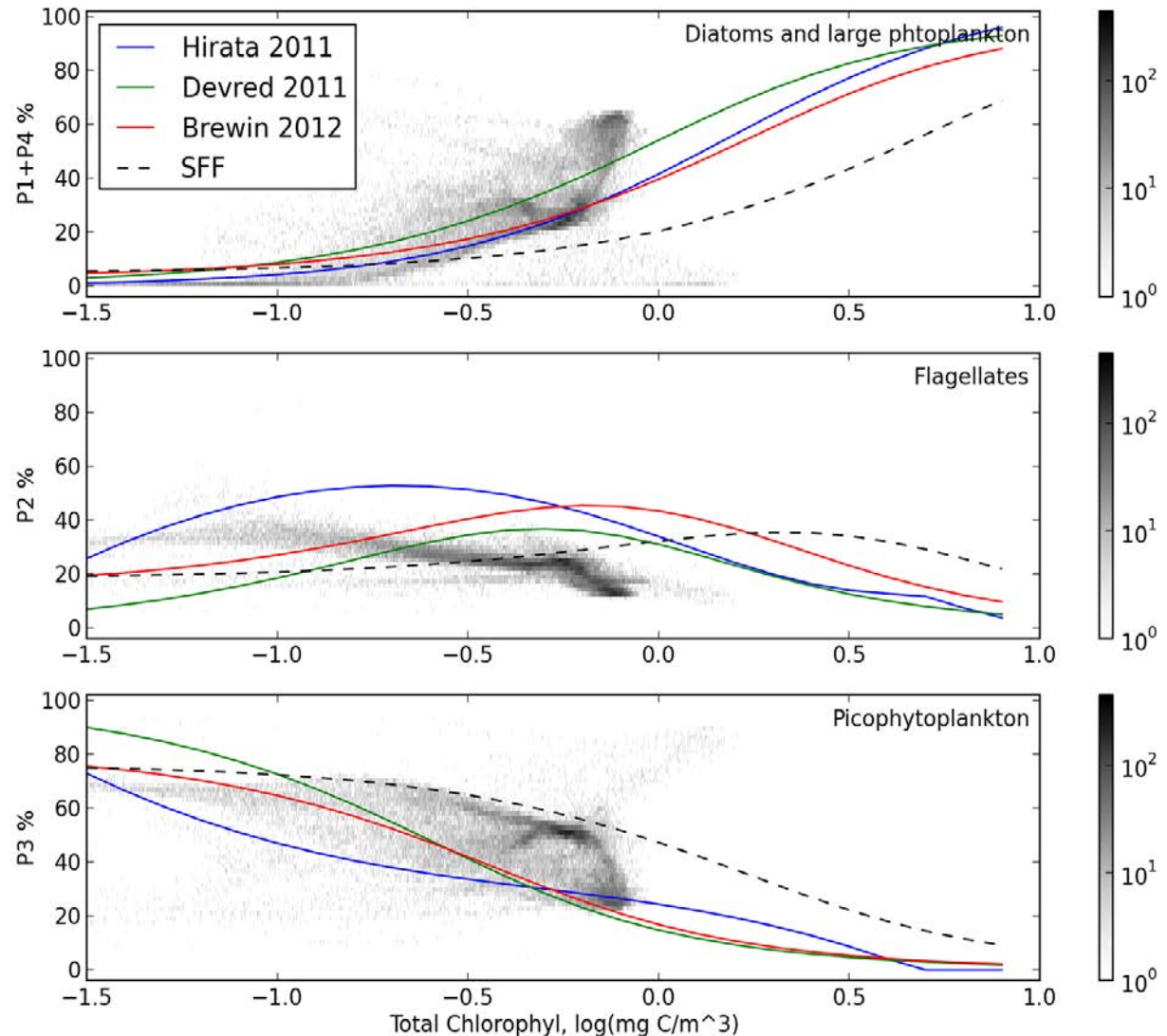
- Primary production
- Length of food web
- Whole ecosystem production & respiration
- Carbon dioxide drawdown

Hirata et al., 2008, 2011

Brewin et al., 2010 a,b,c, 2011

- **Pigment models:** Hirata 2001, Brewin 2012, Devred 2011
- **Size based models:** Size fractionated filtration.

Surface Phytoplankton community structure- 2001





1. Trophic structure in terms of organism size and function (here we refer to high level ecosystem function, i.e. autotrophy, heterotrophy, decomposition).
2. Within size / functional class diversity, by subdividing by biological traits (e.g. feeding strategy, motility, physiology).
3. Within trait diversity whereby intra- and inter-specific competition is described by defining a set of species within each trait type, stochastically drawing parameters from a rule based parameter space.

