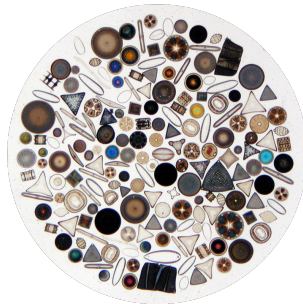


# An efficient inverse model of the ocean's coupled nutrient cycles.

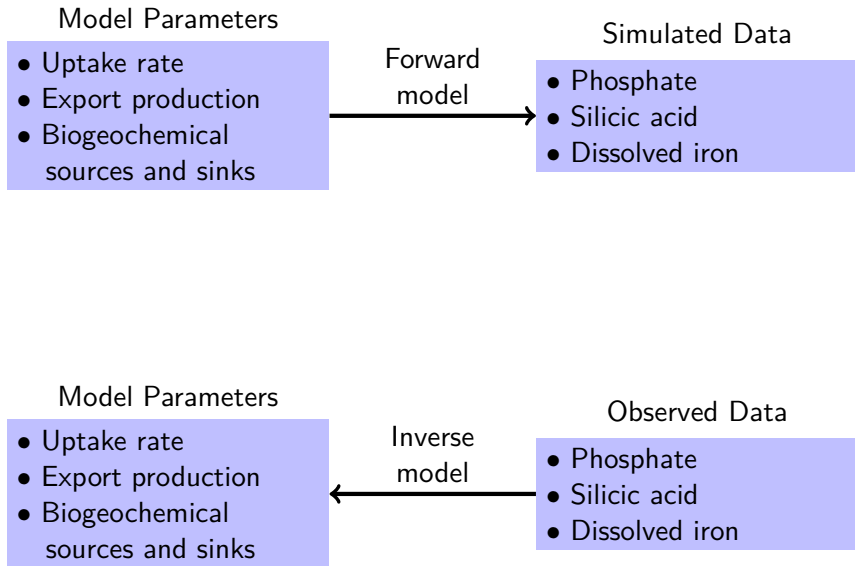
Benoît Pasquier  
Ph.D. student

Under the supervision of Mark Holzer  
School of Mathematics and Statistics UNSW



10 microns

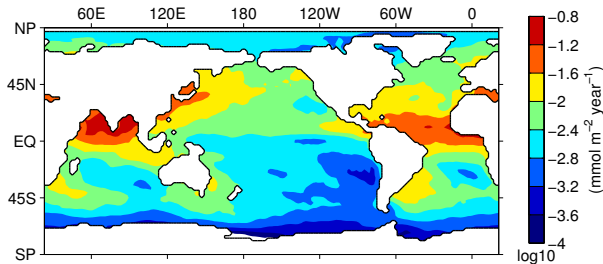
# What is inverse modeling?



# Motivation

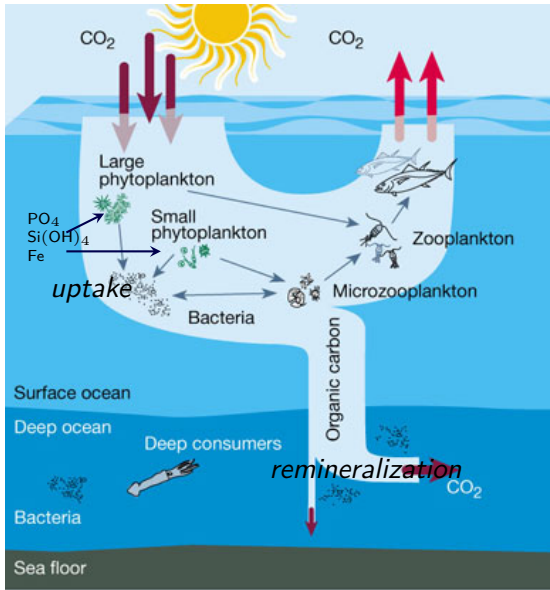
1. Understand and quantify how nutrient cycles are coupled
2. Quantify the pathways and timescales of the global response to large-scale iron perturbations
3. Quantify the role of marine diatoms and the silicon cycle in mediating the response
4. Quantify the sensitivity of elemental ratios (e.g., Si : P) to iron perturbations

Atmosphere-to-Ocean flux of soluble iron  
(dust, combustion deposits)



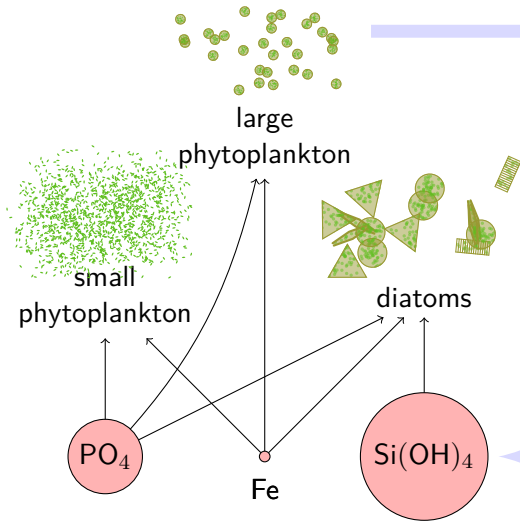
*Luo et al. [2008]*

# Nutrient cycle and coupling



- complex ecosystem
- large amount of species
- multiple nutrients
- poorly understood mechanisms (e.g. iron scavenging)
- simulation can be costly
- large parameter space
- lack of data (e.g. iron)
- poorly constrained parameters

# Distill essentials and simplify



- Zooplankton grazing embedded in phytoplankton mortality
- plankton not explicitly transported as tracers
- only nutrients tracers
- data constrained
- embedded in a data-assimilated global circulation
- numerically highly efficient; allows for optimization and novel diagnostics *Pasquier et al.* (in prep.)

# Plankton population model

- population local equation, a modified logistic equation, for each phytoplankton class

$$\partial_t b(t) = \mu b(t) - \lambda \left( \frac{b(t)}{b^*} \right)^{\frac{1}{\alpha}} b(t)$$

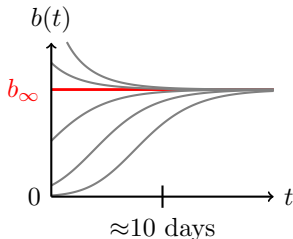
- assumed fast evolution to

$$b_{\infty} = b^* \left( \frac{\mu}{\lambda} \right)^{\alpha}$$

- associated nutrient uptake is population  $\times$  growth rate

$$J = b_{\infty} \mu = b^* \left( \frac{\mu}{\lambda} \right)^{\alpha} \mu$$

- $b(t)$  plankton concentration
- $\mu$  growth rate
- $\lambda$  mortality rate
- $b^*$  pivotal concentration sets the scale of grazing



*Galbraith et al.* [2010]

*Dunne et al.* [2004]

*Armstrong* [1999, 2003]

# A versatile coupling approach

- mortality rate dependencies

$$\lambda = \lambda_0 F_T$$

- growth rate dependencies

$$\mu = \mu_0 F_T F_I \min(m_P, r_{Fe} m_{Fe}, \dots)$$

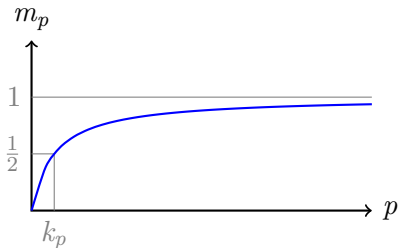
- $F_T = e^{\kappa T}$ , temperature dependence *Eppley* [1972]
- $F_I$ , irradiance dependence
- $m_X$ , nutrient  $X$  deficiency
- $P$ , phosphate concentration
- $Fe$ , iron concentration

*Galbraith et al.* [2010]

- Michaelis-Menten kinetics

$$m_P = \frac{P}{P + k_P}$$

$$m_{Fe} = \frac{Fe}{Fe + k_{Fe}}$$



# Coupled nutrient cycling and biogenic transport embedded in global circulation

- total uptake is summed over phytoplankton classes  $X$

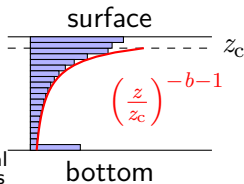
$$J_{\text{up}}(P, Fe, Si) = \sum_X J_X(P, Fe, Si)$$

- modeled  $P$  and  $Fe$  obey tracer equation

$$\partial_t P + \mathcal{T}P = \overbrace{-J_{\text{up}}}^{\text{sink}} + \overbrace{\mathcal{S}J_{\text{up}}}^{\text{source}}$$

$$\partial_t Fe + \mathcal{T}Fe = \overbrace{R_{Fe:P}(Fe)}^{\text{uptake ratio}}(-J_{\text{up}} + \mathcal{S}J_{\text{up}}) + \overbrace{J_{\text{ext}}(Fe)}^{\text{external sources and sinks}}$$

- $\mathcal{T}$  data-assimilated transport operator  
*Primeau et al. [2013]*
- $\mathcal{S}$  (biogenic transport) instantly redistributing  $J_{\text{up}}$  as the sinking particle flux divergence



*Martin et al. [1990]*

- similar equation for Si



# Discretized equations

An expression of the form  $\partial_t \mathbf{x} = \mathbf{f}(\mathbf{x})$  where the variable  $\mathbf{x}$  is

$$\mathbf{x} = \begin{bmatrix} \mathbf{p} \\ \mathbf{f} \\ \mathbf{s} \end{bmatrix} \quad \text{with } \mathbf{p} = \begin{bmatrix} P_1 \\ P_2 \\ \vdots \\ P_n \end{bmatrix}, \mathbf{f} = \begin{bmatrix} Fe_1 \\ Fe_2 \\ \vdots \\ Fe_n \end{bmatrix}, \text{ and } \mathbf{s} = \begin{bmatrix} Si_1 \\ Si_2 \\ \vdots \\ Si_n \end{bmatrix}.$$

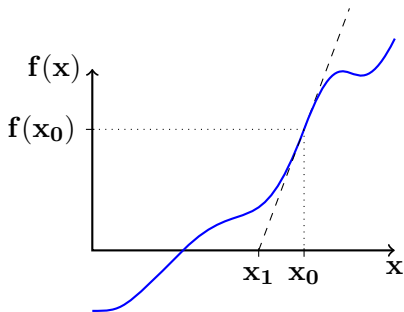
and the right-hand side function is

$$\mathbf{f}(\mathbf{x}) = - \overbrace{\begin{bmatrix} \mathbf{T} & & \\ & \mathbf{T} & \\ & & \mathbf{T} \end{bmatrix}}^{\text{linear}} \mathbf{x} + \overbrace{\begin{bmatrix} (\mathbf{S} - \mathbf{I})\mathbf{J}_{\text{up}}(\mathbf{x}) \\ (\mathbf{S} - \mathbf{I})\text{diag}(\mathbf{R}_{\mathbf{f}:\mathbf{p}})\mathbf{J}_{\text{up}}(\mathbf{x}) + \mathbf{J}_{\text{ext}}(\mathbf{Fe}) \\ (\mathbf{S} - \mathbf{I})\text{diag}(\mathbf{R}_{\mathbf{s}:\mathbf{p}})\mathbf{J}_{\text{up}}(\mathbf{x}) + \mathbf{J}_{\text{ext}}(\mathbf{Si}) \end{bmatrix}}^{\text{nonlinear}}.$$

# Newton PDE solution

- steady state:  $\partial_t \mathbf{x} = \mathbf{f}(\mathbf{x}) = \mathbf{0}$
- use Newton's Method (generalized zero search)  
linear approximation:

$$\mathbf{f}(\mathbf{x}_1) = \mathbf{f}(\mathbf{x}_0) + \mathbf{Df}(\mathbf{x}_0) (\mathbf{x}_1 - \mathbf{x}_0) + \mathbf{o}(\|\mathbf{x}_1 - \mathbf{x}_0\|)$$

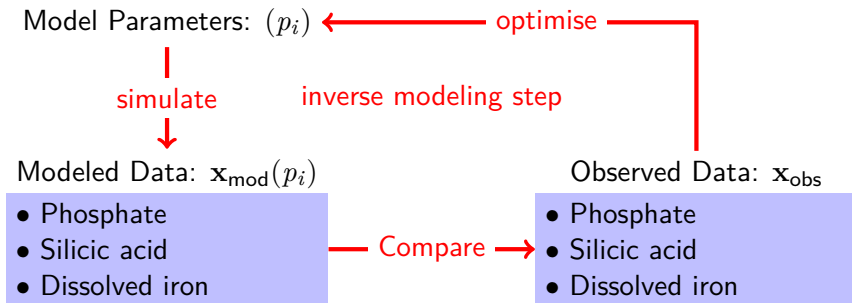


where  $\mathbf{Df}$  is the Jacobian,  
a  $n \times n$  sparse matrix  
where  $n \approx 200000k$   
for  $k$  nutrients!

To get  $\mathbf{f}(\mathbf{x}_1) \approx \mathbf{0}$ ,  
we take

$$\mathbf{x}_1 = \mathbf{x}_0 - \mathbf{Df}(\mathbf{x}_0)^{-1} \mathbf{f}(\mathbf{x}_0)$$

# Objective parameter determination



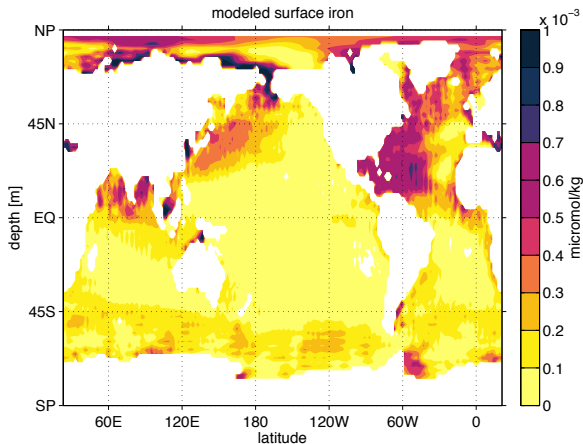
- Choosing appropriate weights  $\mathbf{w}$ , we build an objective function of the quadratic concentration mismatch:

$$c(p_i) = (\mathbf{x}_{\text{mod}} - \mathbf{x}_{\text{obs}})^T \text{diag}(\mathbf{w})(\mathbf{x}_{\text{mod}} - \mathbf{x}_{\text{obs}})$$

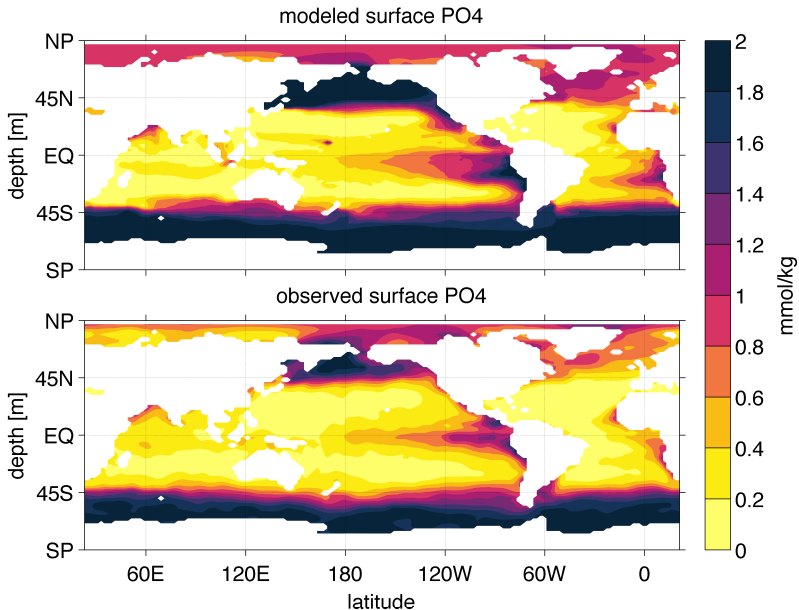
- nonlinear w.r.t. parameters
- optimisation only possible with efficient simulation

# Preliminary model setup

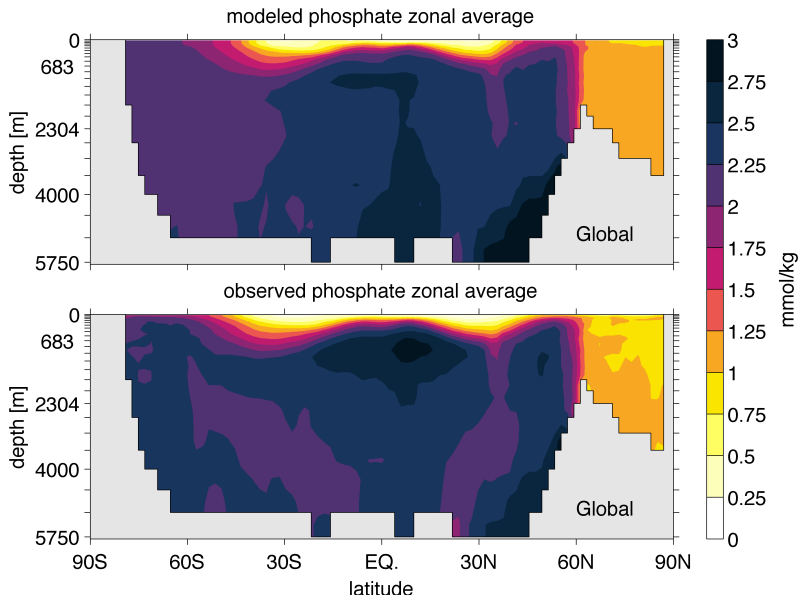
- prescribed iron from *Frantz et al.* (in prep.)
- some data interpolated from different grid
- only 2 parameters optimised



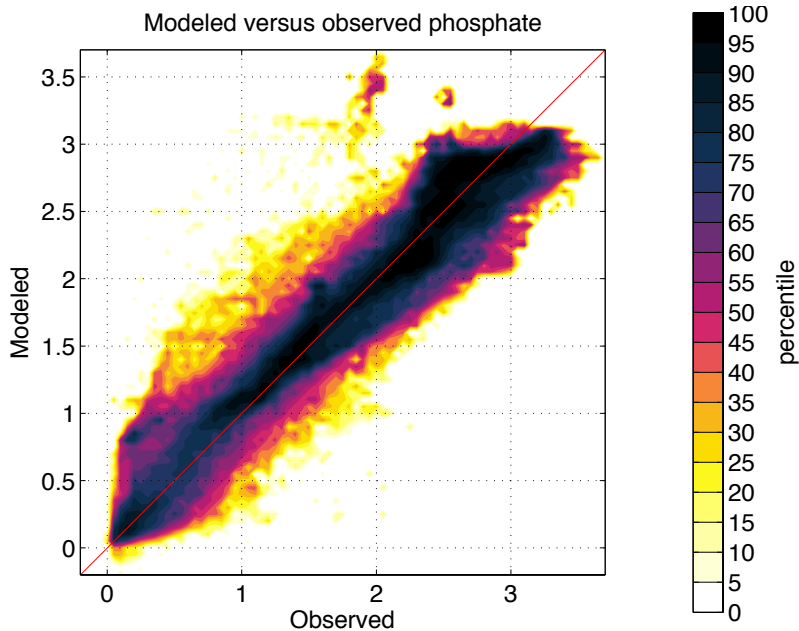
# Model vs. observed Phosphate



# Model vs. observed Phosphate

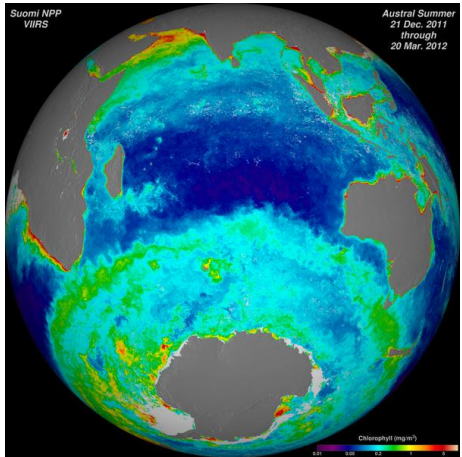


# Model vs. observed Phosphate



# Future outlook and conclusion

- from prescribed iron to coupled iron
- refine coupling parameters
- optimise larger set of parameters
- formulate iron perturbations
- diagnose the teleconnections of the response using Green functions forward and adjoint models *Pasquier et al.* (in prep.)
- include silicon cycle
- elucidate elemental ratios





# Future outlook and conclusion

- from prescribed iron to coupled iron
- refine coupling parameters
- optimise larger set of parameters
- formulate iron perturbations
- diagnose the teleconnections of the response using Green functions forward and adjoint models *Pasquier et al.* (in prep.)
- include silicon cycle
- elucidate elemental ratios

