

Extracting plankton physiological parameters from batch culture data

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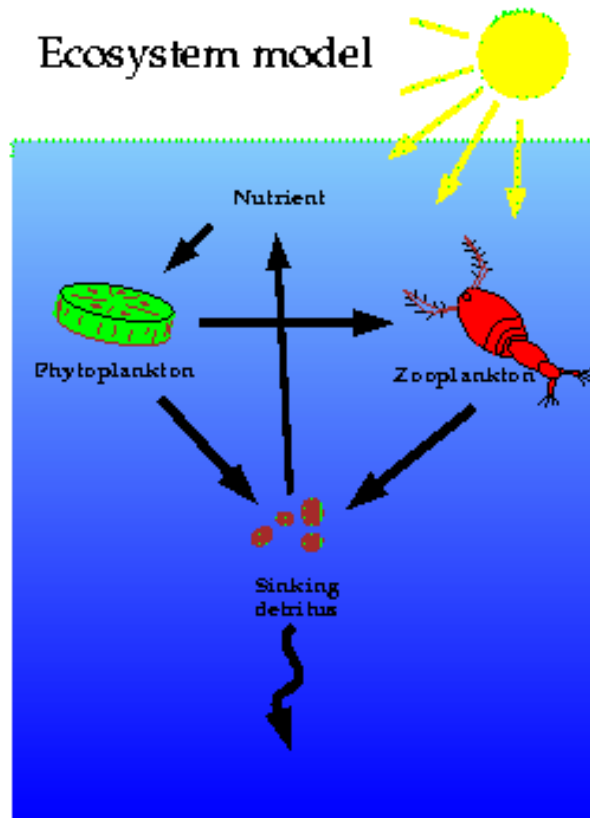
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SIMONS FOUNDATION

Ocean ecosystem models: the olden days

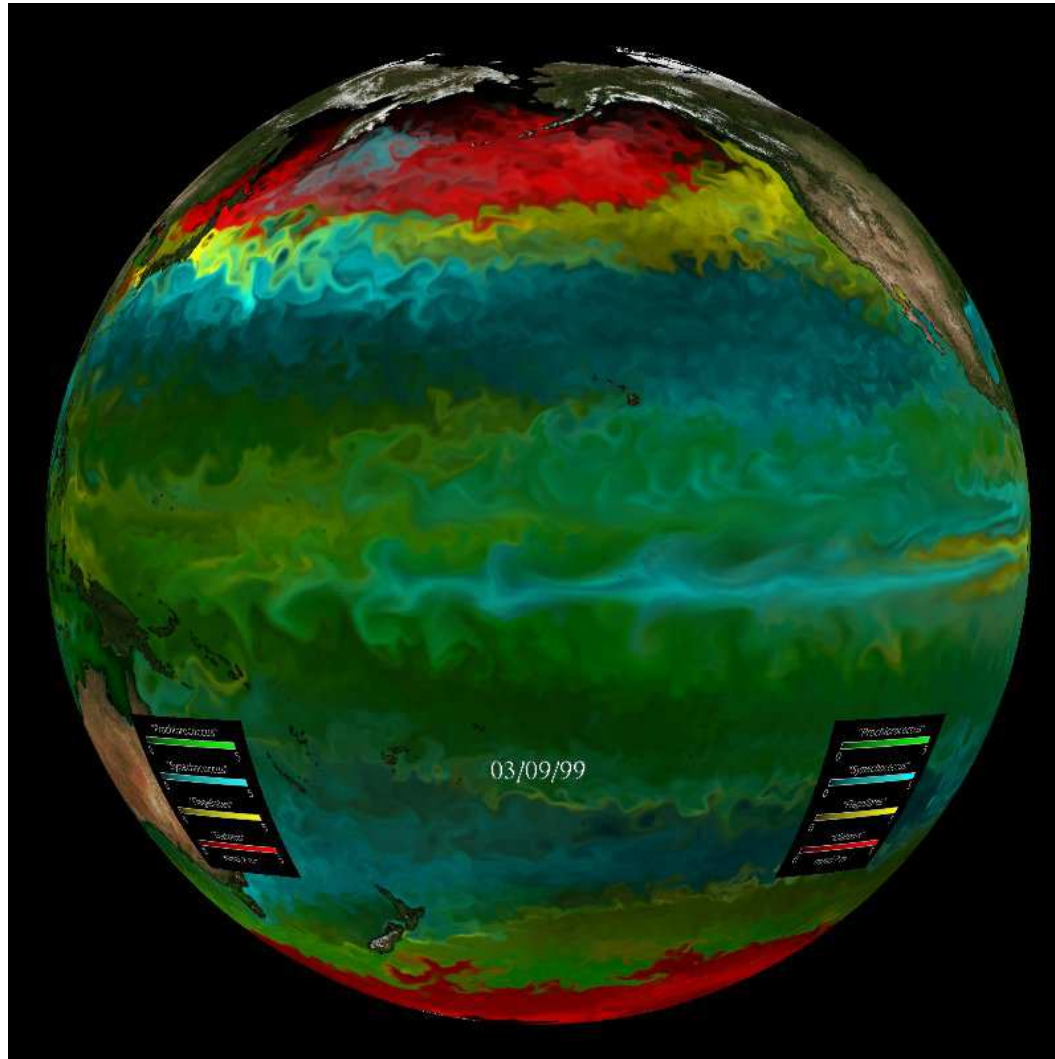


<http://www.geos.ed.ac.uk/homes/hkettle>

Only 2 plankton types (Phytoplankton & Zooplankton)

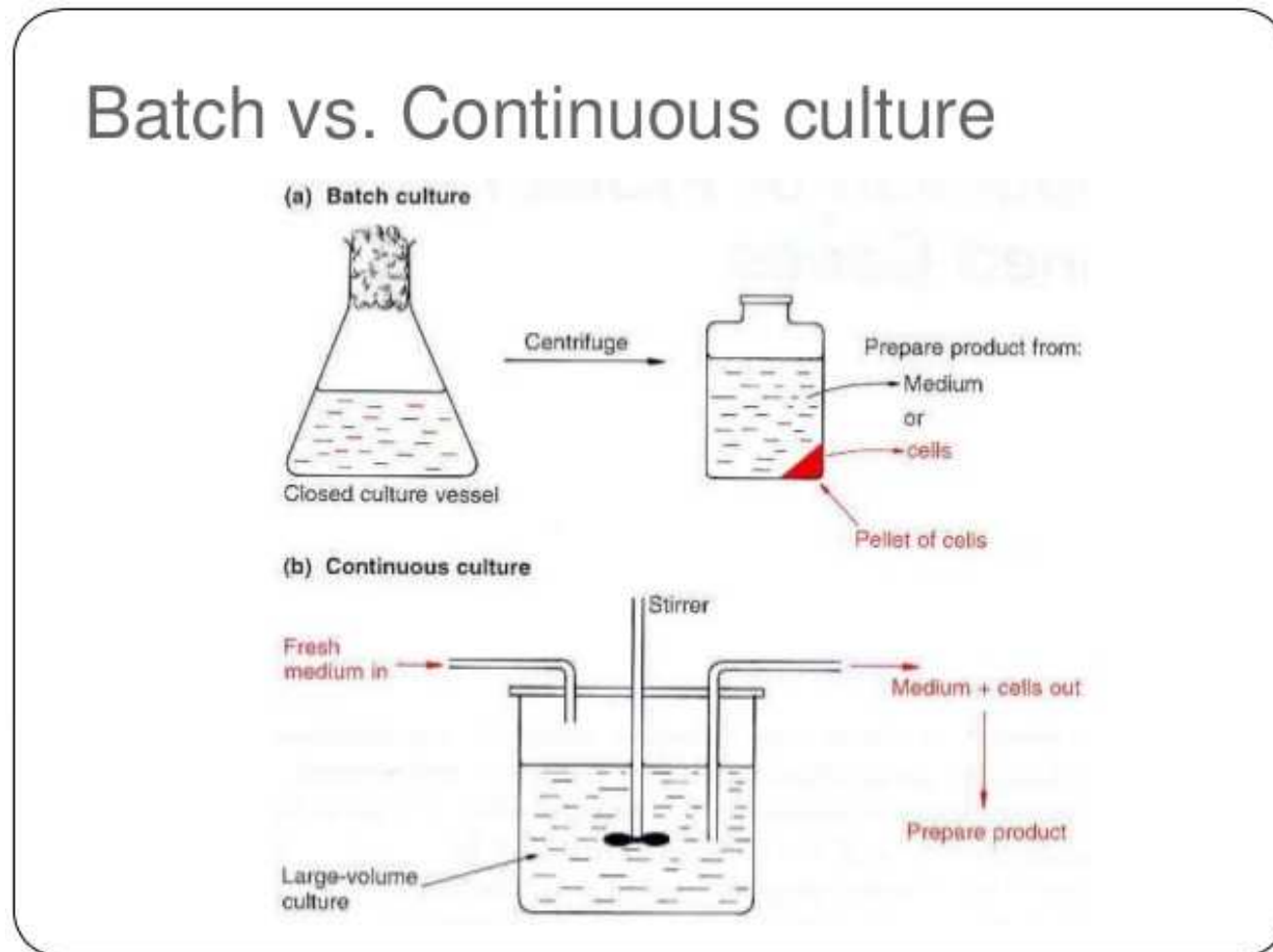
Parameter values not that critical:
Ecosystem will consume nutrients that are supplied

Ocean ecosystem models: nowadays



Competition between many plankton types: \Rightarrow
Need to know parameter values for different plankton species

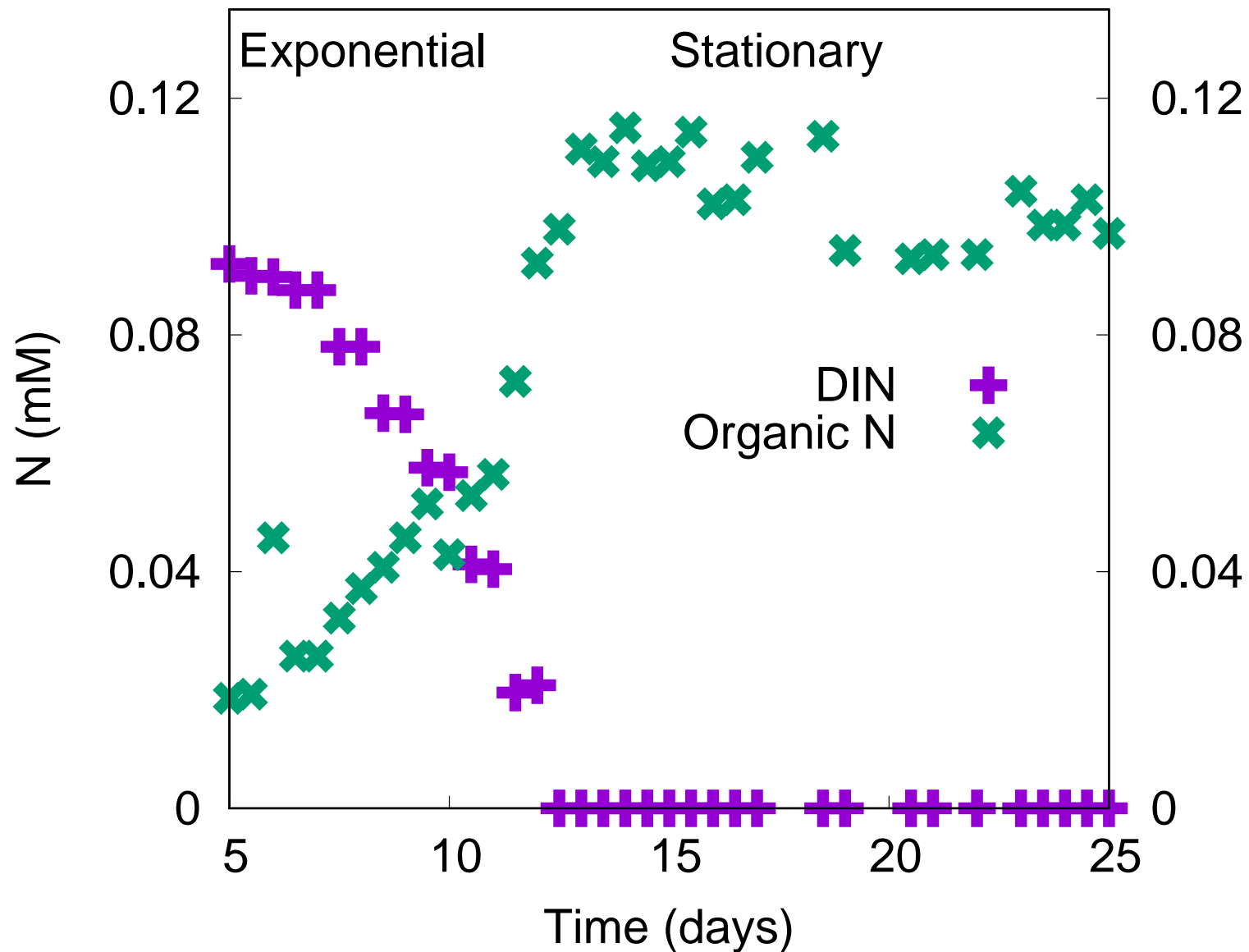
Data to constrain plankton parameters



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We consider batch culture data on the prymnesiophyte
Isochrysis galbana (Flynn et al., 1994)

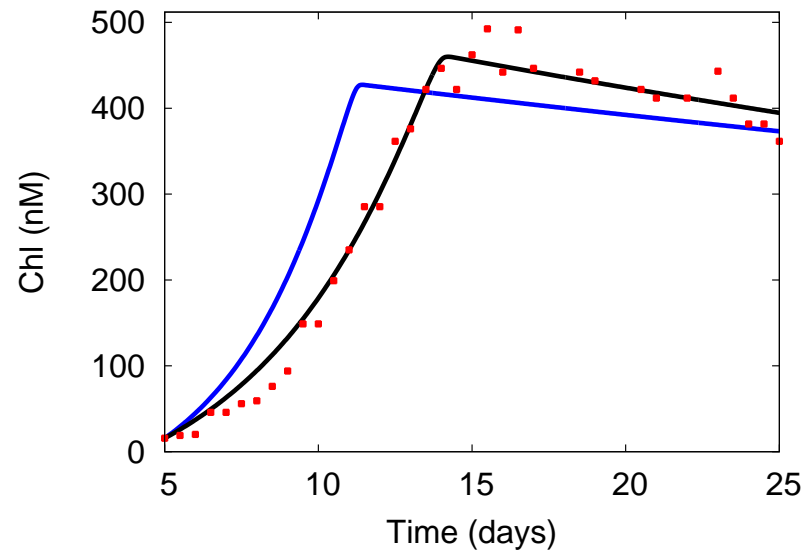
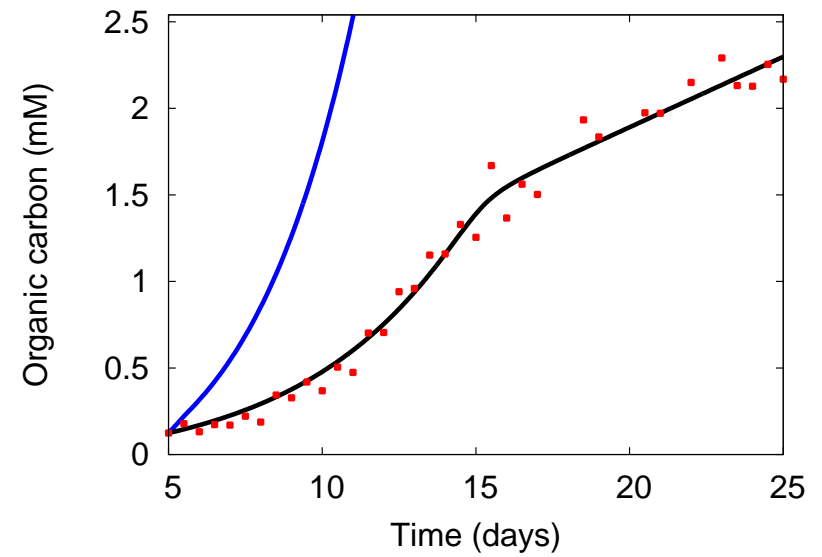
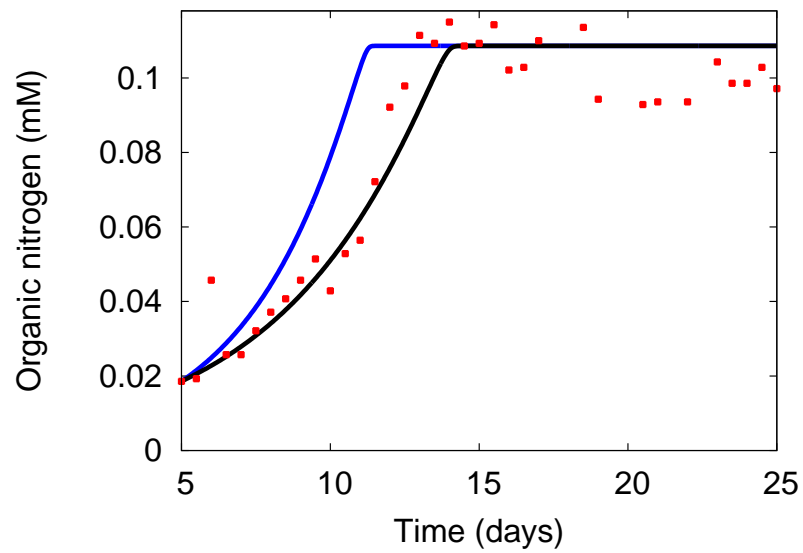
Flynn et al. (1994) batch culture data



Omta et al. (2017) model formulation

$$\begin{aligned}\frac{dC}{dt} &= (C_{syn} - E) P, E = \frac{m_{ex} \left(1 + \tanh\left(\frac{C}{P} - r_{ex}\right)\right)}{2} \\ \frac{dP}{dt} &= P_{syn} P, P_{syn} = \mu \left(\frac{N}{N + K}\right) \\ \frac{dN}{dt} &= -\frac{dP}{dt} \\ \frac{dr}{dt} &= \frac{r_0 - r}{\tau}, r_0 = b \frac{P}{C}\end{aligned}$$

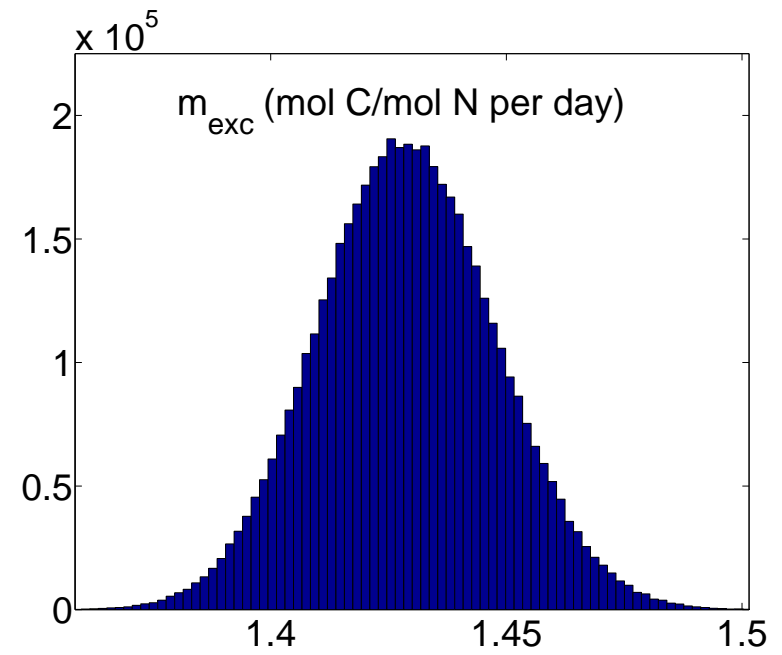
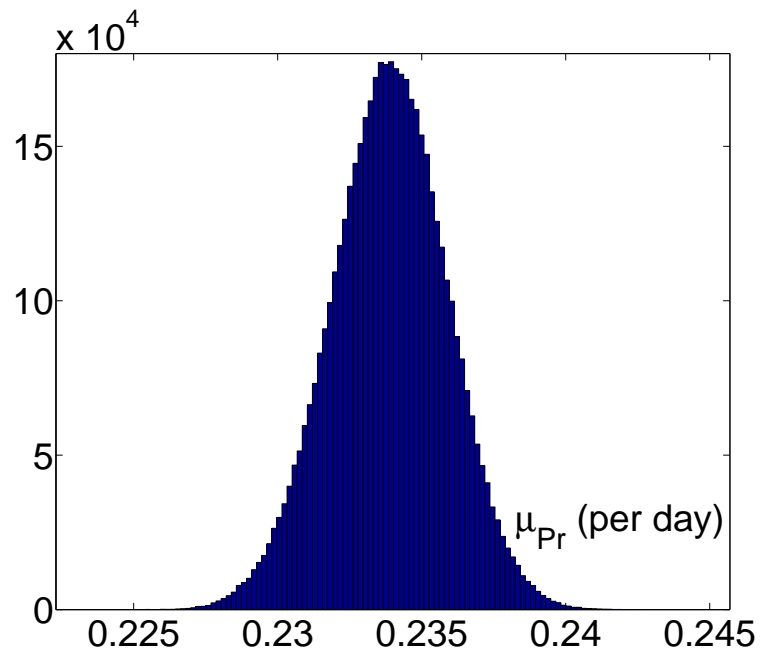
Model fits



Estimated parameter values

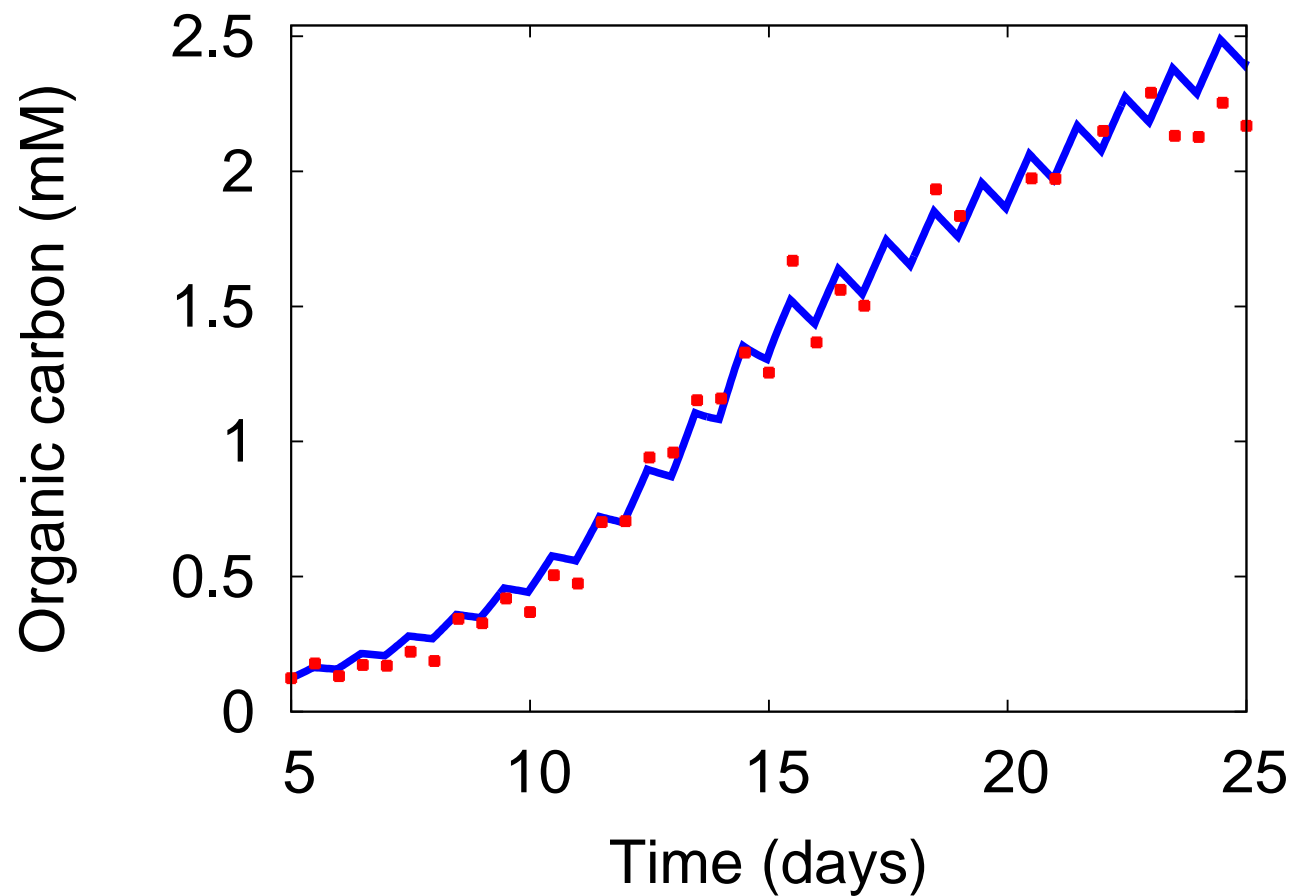
Parameter	Description	Estimated value	Units
μ	Maximum protein synthesis rate	0.234 ± 0.002	d^{-1}
C_{syn}	Maximum photo-synthesis rate	2.19 ± 0.02	$\text{mol C}/(\text{mol N}) \text{d}^{-1}$
m_{ex}	Maximum C exudation rate	1.43 ± 0.02	$\text{mol C}/(\text{mol N}) \text{d}^{-1}$
r_{ex}	Carb:protein ratio above which exudation occurs	13.49 ± 0.11	$\text{mol C}/(\text{mol N})$
τ	Photoacclimation time	9.59 ± 0.13	d
b	Photoacclimation parameter	57.1 ± 0.2	$\mu\text{mol Chl mol C}/(\text{mol N})^{-2}$

Posterior parameter distributions



Model extension: diurnal cycle

$$I = \begin{cases} 100 & 12 \text{ hrs/d} \\ 0 & 12 \text{ hrs/d} \end{cases}$$



Possible model extension: Multiple limiting nutrients

Minimum formulation most straightforward:

$$P_{syn} = \mu * \min \left(\frac{N}{N + K_N}, \frac{I}{I + K_I} \right)$$

But tricky in Stan...

Synthesizing Unit provides an alternative:

$$P_{syn} = \mu \frac{1}{1 + \frac{K_N}{N} + \frac{K_I}{I} - \frac{1}{\frac{N}{K_N} + \frac{I}{K_I}}}$$

Possible model extension: Pahlow/Smith optimal uptake kinetics

Optimal allocation between surface uptake sites and internal enzymes leads to slightly modified uptake kinetics. Instead of:

$$P_{syn} = \mu \frac{N}{N + K_N}$$

we obtain:

$$P_{syn} = \mu \frac{N}{N + 2\sqrt{K_N N} + K_N}$$

Possible model extension: Different exudation formulation

Perhaps make C exudation linearly proportional to Carb:protein ratio?

Instead of:

$$E = \frac{m_{ex} \left(1 + \tanh \left(\frac{C}{P} - r_{ex} \right) \right)}{2}$$

We could simply have:

$$E = m_{ex} \frac{C}{P}$$