Marine Biogeochemical Modeling: **Basic Philosophy and Methods**

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OCB Ocean Acidification Short Course 2009

Talk Outline Science Motivation Philosophy of Modeling Model Construction •Evaluation Against Data

Supported by:











<u>Ocean Carbon Cycle &</u> <u>Ocean Acidification</u>



Anthropogenic CO₂ uptake currently controlled by ocean circulation; but in future, what will be role of climate & biology?

For ocean acidification may want models to address many different aspects:

-patterns & trends in seawater chemistry -population biology of individual species -food-web & ecological interactions

-biogeochemical feedbacks

-socio-economic effects on fisheries & ecosystem services



"I am never content until I have constructed a mechanical model of the subject I am studying. If I succeed in making one, I understand; otherwise I do not."

- Lord Kelvin

"People don't understand the earth, but they want to, so they build a model, and then they have two things they don't understand,"

-Gerard Roe in "The Whale and the Supercomputer" by C. Wohlforth





The Interdisciplinary Conundrum

Physics

(relatively) simple set of (mostly) known governing equations (Navier-Stokes) => complex phenomenon
unresolved scales

Chemistry

mass balance equations
chemical fields integrate over time/space variability
don't uniquely identify process/mechanism

Biology

information as stories and conceptual pictures
historical/evolutionary contingency
no "biological" Navier-Stokes
aggregate complicated dynamics across multiplescales (genes, cells, populations, ecosystems)



Why do we build/use models?

-Quantitative dynamical framework •Are different data sets, rate estimates consistent?

-Design of experiments & observing systems •What, where and when do we sample?

-Hypothesis testing •If we add or change X, what happens?

-Forecasting

•What will the ocean look like at some point in the future?

Solving for unknown parameters and rates
 Given things we can measure, can we estimate can we estimate properties that are difficult to measure?





What is a model?

-regression curves

variable Y is a function of other variables X and regression parameters p: Y=f(variables, parameters)

 $ChI = p_0 + p_1T$

-forward models (often time dependent) Integrate forward in time to find Y: dY/dt = circulation + f(variables, parameters, forcing) d Phyto /dt = circulation + μ*Phyto*(1-e^{-αE/μ}) -λPhyto

-inverse models

Invert the problem to find parameters from data: Parameters = f(circulation, data, forcing)



What is a model? (continued)

-diagnostic versus prognostic models In diagnostic models, some variables may be prescribed based on observations: e.g., satellite chlorophyll => ecosystem model (no equation for dChl / dt)

-data assimilation

Combine model equations and observations in a dynamically consistent fashion e.g., weather prediction analysis = f(model forecast, observations)

Prognostic, forward models needed to project into the future



From Word Problem to Equations

Phytoplankton levels depends on nutrient inputsPerturbations relax back to some stable background level



Simple NPZ Model

$$\frac{dP}{dt} = \mu_0 \left(\frac{N}{k_N + N} \right) \left(1 - e^{\alpha E / \mu 0} \right) P - g \left(\frac{P}{k_P + P} \right) Z - m_P P$$

Nutrient Light limitation

$$\frac{dZ}{dt} = ag\left(\frac{P}{k_P + P}\right)Z - m_Z Z$$

$$\frac{dN}{dt} = -\mu_0 \left(\frac{N}{k_N + N} \right) \left(1 - e^{\alpha E/\mu 0} \right) P + (1 - a)g \left(\frac{P}{k_P + P} \right) Z + m_P P + m_Z Z$$

Three coupled ordinary differential equationsMass conservation



Discretization & Numerical Integration

$$\frac{dy}{dt} = f(t, y)$$

Discrete form $\Delta y = f(t, y) \Delta t$

numerical method: Euler's Method $y^{n+1} = y^n + f(t^n, y^n)\Delta t$



Subdivide time into discrete time-steps
"Integrate" forward using ∆y/∆t approximation
Numerical methods introduce errors



Discretization & Numerical Integration



•use a "trial step" to find gradient at mid-point •even with 2x larger Δt , more accurate integration



<u>How do you estimate parameters and</u> <u>functional forms?</u>

Laboratory & field incubations •P-E curves; nutrient uptake curves •elemental stochiometry Comparative analysis •allometric relationships Tuned or optimized against field data •mismatch between parameters and data •cross-site comparison Previous models



Adding Circulation



$$\frac{\partial P}{\partial t} + \vec{u} \cdot \nabla P - \kappa \nabla^2 P = RHS$$

advection diffusionbiologicalsource/sinkterms



Models have time/space scale limits



- •Computational costs scale as (length)³ to (length)⁴ and (time)
- •Typically get ~2-3 decades in space (more in time)
- •Can not resolve all scales; parameterize sub-grid scale



Coupled "Eco-biogeochemical" Elements

Physics (flow field; mixing)

equations for resolved flow; level of approximation (e.g., primative equations; quasi-geostrophy)
forcing (winds, heat & freshwater fluxes, light, tides)
parameterization of unresolved scales (mixing)
model architecture (e.g., horizontal vs. isopycnal)

Chemistry (CO_2 , O_2 , nutrient fields)

air-sea gas exchange
elemental stoichiometries
trace metal deposition and scavenging

Biology

•primary production, respiration, remineralization

- community structure and succession
- bio-optics

•etc.





-Aggregate into trophic levels/functional groups

- -Rates/processes from limited culture/field studies
- -Many aspects empirically based
- -Data poor for validation (rates, grazing, loss terms)

Multi-scale models

-cellular models

•gene expression, metabolism, energetics

-population models

individual based or continuous dist.
cell-cell interactions (LES & DNS)
-simulate ecological functions

•"genotype" => "phenotype"

abandon "boxes"

-ecological/evolutionary rules for ecosystem assembly

- maximize resiliance or energy flow
 emergent behavior & selection
- selection and niche adaptation
 Physiological plasticity & constraints
 micro/macroevolution

metabolic & energetic models Individual-based & discrete multi-scale models I microbes Continuous Phytoplankton Zooplankton distribution models n size size Size-structured & functional group models Phytoplankton Herbivore Higher trophic levels Nutrient Carnivore DOM & Detritus Bacteria

Cellular genomic,

<u>Cell</u> Physiology/ Genomics

Chitin

UDP-GlcNac

GIcNac-1P

GICNac

6P

Pyrimidine

biosynthesis

HCO₃

Glycolysis

Gluconeogenesis glucose

fructose

6P

GAP

PEP

GIC 6P

OAA

soprenoid succinate 3PGA glutamine biosynthesis glycolate malate pyruvate glyoxylate Photo pyruvate 2P respiration carbamoy|-P Calvin malate fumarate GAR DOX-P 02 cycle CO₂ pyruvate UMP co, MEP acety| CoA CO, NH4 RUBP OAA UDP acety CoA citrate UTP TCA β-carotene cycle Fatty Acid Fatty CTP Biosynthesis zeaxanthin CO2 Acid (FASII) xanthophy Oxidation cycling FA FA antheraxanthin carnitine carnitine FA CoA glutamate 16:0,18:0 fatty acids diatoxanthin carbamoy| P <---CO2+ NH4 ·· > _ diadinoxanthin ornithine citru||ine glutamine cycling carnitine carnitine diadinoxanthin Urea NH, + -- NO cycle fucoxanthin proline ornithine citrulline urea soprenoid NO putrescine argino NO₂ biosynthesis arginine Purine succinate biosynthesis acety| CoA long-chain polyamines NO₃ PRPP guanidino spermidine HMG COA acetate IMP meva onate creatine spermine XMP SAMP anostero creatine=P ergosterol, cholesterol GMP AMP phytostero 3

02 + H_0 - H_02

Glyoxylate

cycle

acetyl CoA

Silicic

acid

PUFA

FA

CoA

Fatty Acid

Oxidation

HCO₃

glycerate

PO4

SO4 NO3

glutamate

glutamine

NH4⁺ Urea Sugars Amino

urea

NH4

+

CO2

s

Diatom Genome Armbrust et al., Science, 2004

IC INSTITUTION

1930



Ocean ecology and biogeochemistry are (still) data-driven sciences



How do we avoid the trap of: "false models tested by inadequate data"

John Steele







20 February 2009

188N-0124-7963



Special Joans, Skill Assessment for Coupled Biological (Physical Models of Harine Systems

edited by Daniel R. Lynch, Dennis J. McGillicuskiy, 3t and Francisco E. Werner

JOURNAL OF MARINE SYSTEMS

J. Marine Systems Special Issue on Skill Assessment for Coupled Biological / Physical Models of Marine Systems Vol. 76, Issue 1-2, 2009



4) AE – the average error (bias)

$$AE = \frac{\sum_{i=1}^{n} (P_i - O_i)}{n} = \overline{P} - \overline{O}$$

 r – the correlation coefficient of the model predictions and observations:

Correlation

Bias

$$r = \frac{\sum_{i=1}^{n} \left(O_i - \overline{O}\right) \left(P_i - \overline{P}\right)}{\sqrt{\sum_{i=1}^{n} \left(O_i - \overline{O}\right)^2 \sum_{i=1}^{n} \left(P_i - \overline{P}\right)^2}},$$

2) RMSE – the root mean squared error (also referred to as root mean squared difference): rms Error

$$\mathsf{RMSE} = \sqrt{\frac{\sum\limits_{i=1}^{n} \left(P_i - O_i\right)^2}{n}},$$

Doney et al. J. Mar. Systems 2009 Stow et al. J. Mar. Systems 2009



Figure 2. Diagram for displaying pattern statistics. The radial distance from the origin is proportional to the standard deviation of a pattern. The centered RMS difference between the test and reference field is proportional to their distance apart (in the same units as the standard deviation). The correlation between the two fields is given by the azimuthal position of the test field.





Look at the magnitude & structure in model-data residuals

Ducklow et al. Ann. Rev. Mar. Res. 2009

Log-Normal Variables (e.g., chlorophyll)

 $X = \log(\chi)$

$$\langle \chi \rangle_{\rm G} = \sqrt[N]{\prod_{i} \chi_i} = \exp(\langle X \rangle)$$

The geometric bias:

 $\varepsilon_{\text{bias}}^{\text{G}}(\chi) = \exp(\langle X_{\text{P}} \rangle \neg \langle X_{\text{O}} \rangle)$

$$\varepsilon_{\rm rms}^{\rm G}(\chi) = \sqrt{\exp\left(\frac{1}{N}\sum(X_{\rm P}-X_{\rm O})^2\right)}$$

geometric mean (7)

(6)

(8)

geometric bias (no bias =>1)

geometric rms error (~normalized to (9) typical data value)

Doney et al. J. Mar. Systems 2009 Stow et al. J. Mar. Systems 2009 3) RI – the reliability index

$$RI = \exp\sqrt{\frac{1}{n}\sum_{i=1}^{n} \left(\log \frac{O_i}{P_i}\right)^2},$$

6) MEF – the modeling efficiency

$$MEF = \frac{\left(\sum_{i=1}^{n} (O_i - \overline{O})^2 - \sum_{i=1}^{n} (P_i - O_i)^2\right)}{\sum_{i=1}^{n} (O_i - \overline{O})^2},$$

 $MEF = 1 - RMSE^2/s^2$

 $\chi_{\upsilon}^2 = 1/\upsilon \sum_i (P_i - O_i)^2 / \varepsilon_i$

average factor model differs from data

predictions relative to observed mean

MEF = 1 great MEF = 0 no better than obs. mean MEF < 0 worse than obs. mean

Reduced Chi squared => ~1



Assessing Model Skill

Relationships between the truth, model and data

(adapted from the ideas of Dan Lynch)

a)



Stow et al. J. Mar. Systems 2009





Ocean carbonate system determined by temperature, salinity & 2 of 4 parameters (pH, total carbon, alkalinity, pCO₂) Add sensors to autonomous platforms (AUVs, gliders & floats)

Some Issues to Ponder

Representativeness of data y_0

"footprint" of observation & mismatch with model grid
local heterogenity or point sources
aliasing of unresolved frequencies/wavenumbers (e.g., diurnal cycle)
data selection (i.e., exclude "unrepresentative" observations)

$$R = R_{instrument} + R_{representativeness}$$

$$y_{obs} = y_{true} + \mathcal{E}_{obs} \qquad \mathcal{E}_{obs} = \mathcal{E}_{random} + \mathcal{E}_{systematic}$$
$$E\left[\mathcal{E}_{obs}^{1}, \mathcal{E}_{obs}^{2}\right] \neq 0$$



Modeling Methods for Marine Environments

David M. Glover, William J. Jenkins & Scott C. Doney

-data analysis
-modeling techniques
-ocean examples and applications
-MATLAB based demos and code
-detailed web notes (and perhaps some day a book)

(http://eos.whoi.edu/12.747/)



<u>Matlab Primer</u>

```
-can run from Matlab command window or "scripts" (m-files)
-use help & lookfor commands
```

-define variables (case sensitive) & standard functions:

a = 7.3e-7b = -log10(a)

(follow with ";" if don't want the answer echoed back)

-vector mathematics



<u>Matlab Primer</u>

```
-for-loops to cycle over common set of commands
       for i=1:n
              F(i) = exp(-i*lambda)
       end
-call user-written functions or subroutines
       C = convert to centigrade(F)
-hands-on demonstration (m-files)
       -Euler vs. 2<sup>nd</sup> order Runge Kutta
       -"simple" phytoplankton model
-Ordinary Differential Equation (ODEs) integrators
       - find y(t) from y(t0) and equation for dydt=f(y,t,p)
       -define "function" to integrate e.g. "dydt" (m-file)
       [T,Y] = ode12s('dydt',T,Y0)
CO<sub>2</sub> thermodynamics code [pH,pCO2, ...]=f(DIC,ALK,T,S...)
```

