Using the maximum entropy production principle to understand and predict microbial biogeochemistry

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Reductionist Modeling Approach, Marine Biogeochemistry



- Aggregation causes errors
- Huge number of parameters typically
- Insufficient information
- What happens when the community composition changes?
- How about the Rare Biosphere?

Focus on functions that dissipate energy instead



Pathways are distributed across phyla

No centralized control

Which pathways are upregulated?

Which are downregulated?

How do resources limit?

Many degrees of freedom

Does "who's there" matter?

How to predict protein allocation?

Use optimization-base approach!

Maximum Entropy Production (MEP)

Steady state nonequilibrium systems with many degrees of freedom will likely organize to maximize the rate of entropy production.

Examples: Fire, Hurricanes, and Living systems.

Entropy?

- Dispersal (spreading out) of energy
- Order is usually a small and unimportant component of entropy
- Order must contend with Boltzmann's constant: k=1.3806488 \times 10⁻²³ J K⁻¹
- Living organisms are not low entropy structures.

Easier to think about entropy as the destruction of *free* energy⁺

See: Ziegler 1963 Paltridge 1975 Dewar 2003, 2005 Martyushev & Seleznev 2006

+ Energy is conserved, free energy (or exergy) is not.

Maximum Entropy Production (MEP)

Or: Systems organize to maximize dissipation rate of energy potentials







Autotrophs Nitrifiers hv NH₄⁺ Heterotrophs OM О, NO₂-Sulfate reducers OM SO² hv NO₂ Fe³⁺ 0, 0, Oxidizing Reducing OM Fe²⁺ OM CH₄ CO, Ν, FeS₂ CO₂ H₂S NH₄⁺ CO, Sulfur bacteria Pyrite Iron cyclers Η, Syn. NO₂ OM Methanogens Denitrifiers Anoxygenic phototrophs Anammox DNRA Fermentors

MEP, Which Entropy Term?

Entropy Balance Equation:

$$\frac{dS}{dt} = \sum_{i} J_i \hat{S}_i + \sum_{j} \frac{\dot{Q}_j}{T_j} + \dot{\sigma}$$

where $\dot{\sigma}$ is internal entropy production from irreversible processes. Second Law says $\dot{\sigma} \ge 0$



MEP concerns $\dot{\sigma}$ not S

For Chemical Reaction

$$aA + bB \rightarrow cC + dD$$

Gibbs free energy

$$\Delta_r G(T, P) = \Delta_r G^{\circ}(T, P) + RT \log\left(\frac{(\gamma_c[C])^c (\gamma_d[D])^a}{(\gamma_a[A])^a (\gamma_b[B])^b}\right)$$

$$CH_4 + 2O_2 \rightarrow CO_2 + 2H_2O \quad \Delta_r G^\circ = -818 \ kJ \ mol^{-1}$$

If free energy is not stored, then: $\dot{\sigma} = -\frac{V}{T}r\Delta_r G(T, P)$

Maximum Entropy Production for non-steady state systems

MEP only derived for SS systems (Dewar 2003, Martyushev & Seleznev 2006)



Difference between abiotic (fire) and Life:

•Abiotic: maximizes instantaneous $\dot{\sigma}(t)$ •Life: maximizes time-averaged $\langle \dot{\sigma}(t) \rangle$

Temporal strategies are a hallmark of biology:

circadian rhythm, resource storage, dormancy, life cycles, anticipatory control, etc.



Coordination Over Space

Entropy production can also be increased with coordination over space (see Vallino 2010)



$$\langle \dot{\sigma} \rangle_{\Omega} \ge \frac{1}{V} \iiint_{\Omega} \max \dot{\sigma}(x, y, z) \, d\Omega$$



Cable bacteria

Living systems evolve to maximize energy dissipation over the greatest possible spatial and temporal scales

"Paradigm shift, from 'we eat food' to 'food has produced us to eat it'" (Lineaweaver&Egan 2008)

Diel vertical migration



Stigmergy



MEP approach to modeling biogeochemistry

- Represent biogeochemistry as a distributed metabolic network focused on redox reactions
- Allocate catalysts (protein) to metabolic pathways that maximize entropy production over time and space
- Optimization replaces need to understand how communities assemble (aka, climate verses weather modeling)
- Replace parameters with optimization variables

Example: Siders Pond "Laboratory"



Vertical Gradients



Large spatial gradients (over m not mm)

Distributed metabolic network

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Reaction Kinetics (adaptive Monod)



Siders Pond Transport (1D approximation)

$$\frac{\partial \mathbf{c}(t,z)}{\partial t} = D(z) \frac{\partial^2 \mathbf{c}(t,z)}{\partial z^2} + \left(\frac{D(z)}{A(z)} \frac{\partial A(z)}{\partial z} + \frac{\partial D(z)}{\partial z} - \frac{q(z)}{A(z)}\right) \frac{\partial \mathbf{c}(t,z)}{\partial z} - \frac{\mathbf{c}(t,z)}{A(z)} \frac{\partial q(z)}{\partial z} + \frac{q_L(z)\mathbf{c}_L(z)}{A(z)} + \mathbf{r}(t,z,\varepsilon,\Omega),$$

$$BC: \left. \left. \frac{\partial \mathbf{c}(t,z)}{\partial z} \right|_{z=0} = 0 \qquad \left(-D(z)A(z) \frac{\partial \mathbf{c}(t,z)}{\partial z} + q(z)\mathbf{c}(t,z) \right)_{z=d} = q(d)\mathbf{c}_B$$

Dispersion coefficient, D(z), was determined by fitting predicted to observed salinity vertical profile

Volumetric flow: q(z) and lateral inputs: $q_L(z)$, $c_L(z)$ obtained from observations and assuming:



1D, Local MEP Model Setup

One fine grid for PDE solution, $\mathbf{c}(t, z)$ (space and time adaptive, BACOLI95) Two grids: One course grid for Control Variables, $\varepsilon_i(z, t)$ and $\Omega_{i,i}(z, t)$.

Receding Horizon Optimal Control Problem:

Receding Horizon Optimal Control Problem:

$$\underset{\varepsilon,\Omega}{\overset{t_n+\Delta t_{\infty}}{\longrightarrow}} \int_{t_n}^{t_n+\Delta t_{\infty}} \dot{\sigma}_T(\tau, z_j) e^{-k_W \tau} d\tau \quad \forall j \text{ in the control grid } [z_1, z_2, \dots, z_m] \\
\text{where:} \quad \dot{\sigma}_T(\tau, z_j) \approx -\frac{A(z)}{T(z)} \left(\frac{\Delta I_j \Delta G_{\gamma}}{\eta_{\gamma}} + \sum_k r_k(\tau, z_j) \Delta r_k G_k \right) \\
= t_n \quad t_n + \Delta t_l \quad t_n + \Delta t_l \quad t_n + \Delta t_{\infty} \quad t_{n+1} + \Delta t_{\infty} \quad t_{n+1} + \Delta t_l \quad t_{n+1} + \Delta t_{\infty} \quad t_{n+1} + \Delta t_{\infty}$$

$$\frac{\partial \mathbf{c}(t,z)}{\partial t} = D(x)\frac{\partial^2 \mathbf{c}(t,z)}{\partial z^2} + \left(\frac{D(z)}{A(z)}\frac{\partial A(z)}{\partial z} + \frac{\partial D(z)}{\partial z} - \frac{q(z)}{A(z)} - \nu_s\right)\frac{\partial \mathbf{c}(t,z)}{\partial z} - \left(\frac{1}{A(z)}\frac{\partial q(z)}{\partial z} + \frac{\nu_s}{A(z)}\frac{\partial A(z)}{\partial z}\right)\mathbf{c}(t,z) + \frac{q_L(z)\mathbf{c}_L(z)}{A(z)} + \mathbf{r}(t,z,\varepsilon,\Omega)$$

Note, this is a Control of PDE problem.

Subject to.

Short (0.25 d) vs Long (3 d) optimization: PAR & DO

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Short (0.25 d) vs Long (3 d) optimization: Phytoplankton



Breathing with Sulfate (sulfate reducing bacteria)



dsr: dissimilatory sulfite reductase

*(3 d Interval Optimization)

ATP-citrate lyase (anaerobic photosynthesis)



^{*(3} d Interval Optimization)

Summary

- Maximizing entropy production (destruction of free energy) produces results that are similar to observations
 - Abiotic systems maximize instantaneous entropy production, while biotic system maximize entropy production over time using information.
 - Systems that coordinate information over space can increase global entropy production via coordination of function unless energy is degraded quickly abiotic (e.g., light)
- Model function not individuals (metabolic network)
- > Replace parameters with control variables as much as possible
- The Control of PDE problem is computationally challenging
 - Faster solution approaches?
 - Different model formulation: Trait-based model optimized by MEP?

Acknowledgements

Collaborators:

Mick Follows Ashley Bulseco Nuria Fernández González Chris Algar Jeremy Rich Anne Giblin Aboozar Tabatabai **Gretta Serres** Amy Smith

Students:

Hilary Smith Amy Snyder Alice Carter George Allen Joe Hakam

Meg Yang Petra Byl

RA's:

Suzanne Thomas

Rich McHorney

Jane Tucker

Stef Strebel

Stef Oleksyk

Funding











