



Maximum entropy & maximum entropy production in biological systems: survival of the likeliest?

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Maximum Entropy

(MaxEnt)

Maximum (relative) entropy:

Maximise $H = -\sum_i p_i \ln \frac{p_i}{q_i}$ w.r.t. p_i subject to

$$\left\{ \begin{array}{l} \sum_i p_i x_i = X \\ \sum_i p_i = 1 \end{array} \right.$$

$$\therefore p_{i|C} = \frac{q_i e^{-\beta x_i}}{\sum_i q_i e^{-\beta x_i}}$$



constraints (C)

What is the rationale for MaxEnt ?

- information theory (least biased $p_{i|C}$)
- combinatorics of sample frequencies (most likely $p_{i|C}$)

MaxEnt: the combinatorial rationale

N independent observations

M possible outcomes for each observation, $i = 1 \dots M$

q_i = prior probability of outcome i

Outcome i observed n_i times \rightarrow frequency distribution $p_i = \frac{n_i}{N}$

$$\Pr(\{n_i\}) = N! \prod_{i=1}^M \frac{q_i^{n_i}}{n_i!}$$

$$\lim_{N \rightarrow \infty} \frac{1}{N} \ln \Pr(\{n_i\}) = - \sum_{i=1}^M p_i \ln \frac{p_i}{q_i} = \text{relative entropy of } p_i \text{ and } q_i$$

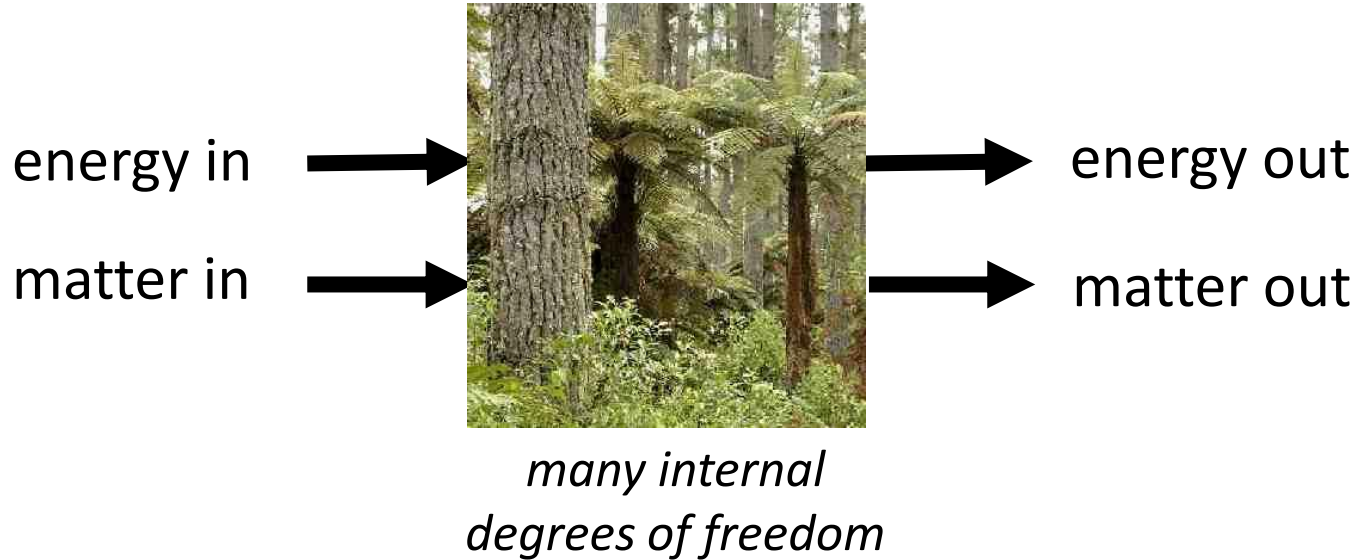
- \therefore The MaxEnt distribution $p_{i|C}$ is **by far the most likely** long-term frequency distribution of outcomes, among all those distributions consistent with given constraints C

(transparent connection to observations)



The prediction challenge in biology:

biological systems are complex, open, non-equilibrium



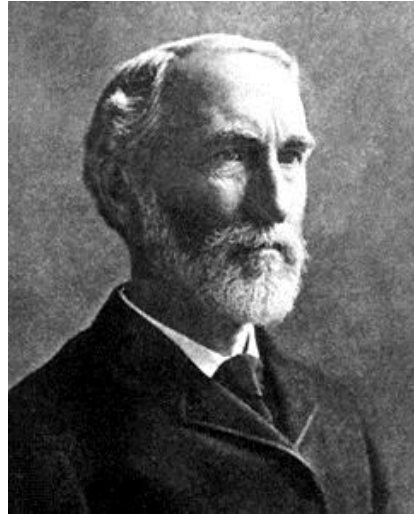
Statistical mechanics:

Some (many?) details of the underlying dynamics are irrelevant for making predictions at larger scales

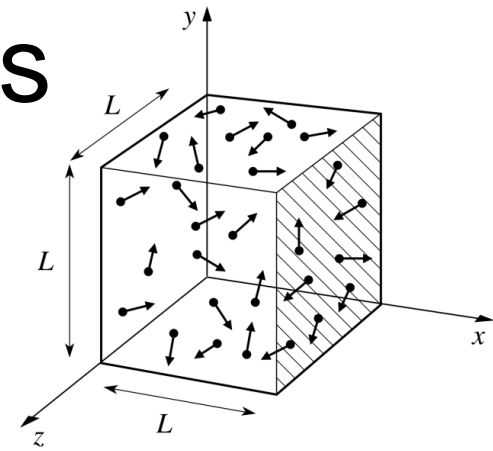
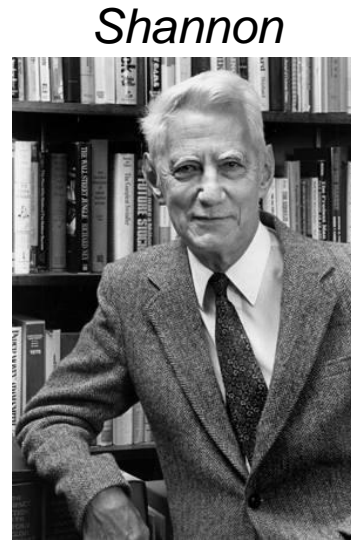
statistical mechanics



Boltzmann



Gibbs



Jaynes

maximum (relative) entropy



~~detailed dynamics $x(t)$~~

most likely
 $p(x \mid \text{key dynamical constraints})$

Maximum (relative) entropy:

Maximise $H = -\sum_i p_i \ln \frac{p_i}{q_i}$ w.r.t. p_i subject to

$$\left\{ \begin{array}{l} \sum_i p_i x_i = X \\ \sum_i p_i = 1 \end{array} \right.$$

$$\therefore p_{i|C} = \frac{q_i e^{-\beta x_i}}{\sum_i q_i e^{-\beta x_i}}$$



constraints (C)

What do the constraints represent ?

- information theory: $C =$ what we know
- statistical mechanics: $C =$ the relevant dynamics (key resource constraints, steady-state balance ...)

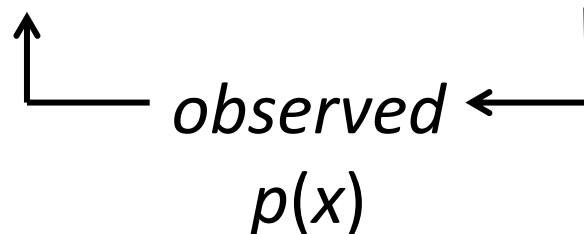
The role of MaxEnt in statistical mechanics

- MaxEnt as a statistical selection principle

Known constraints $C \rightarrow$ most likely $p(x | C)$

- MaxEnt as a tool for identifying the relevant dynamics (C)

Guess constraints $C \rightarrow$ most likely $p(x | C)$



↑
**fraction of time
system is in state x**

Combining mechanism and drift* in ecology

Bertram & Dewar (2015)

Relevant dynamics C + MaxEnt \rightarrow most likely $p(x | C)$



mechanism



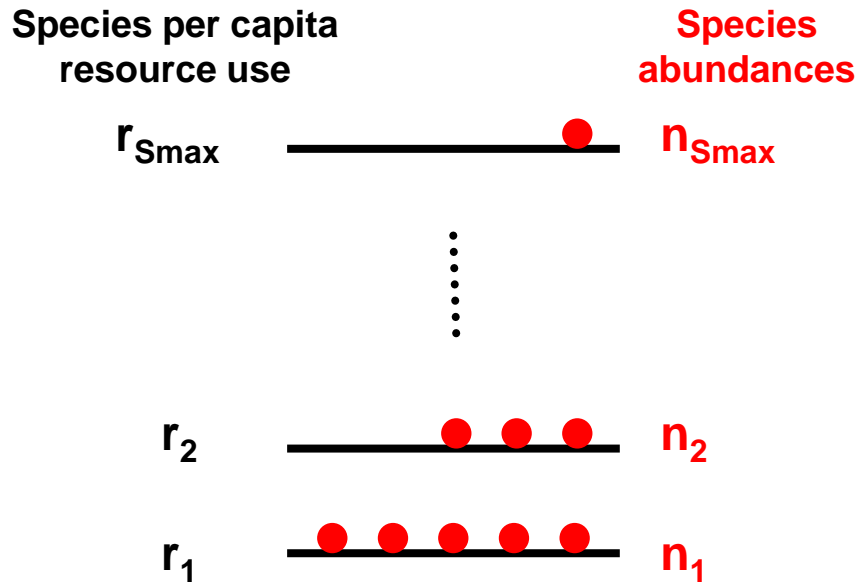
drift*

*** in the biological sense!**

MaxEnt: a non-neutral, resource-based approach

Dewar & Porté (2008), Bertram & Dewar (2013, 2015)

mechanism



Mean annual resource balance : $\sum_{i=1}^{Smax} \bar{n}_i r_i = \bar{R}$
(mean use = mean supply)

Area constraint (hard) : $\sum_{i=1}^{Smax} n_i a_i = A$

drift

MaxEnt



$p(n_1, n_2 \dots n_{Smax} | R)$

(\approx Bose-Einstein)



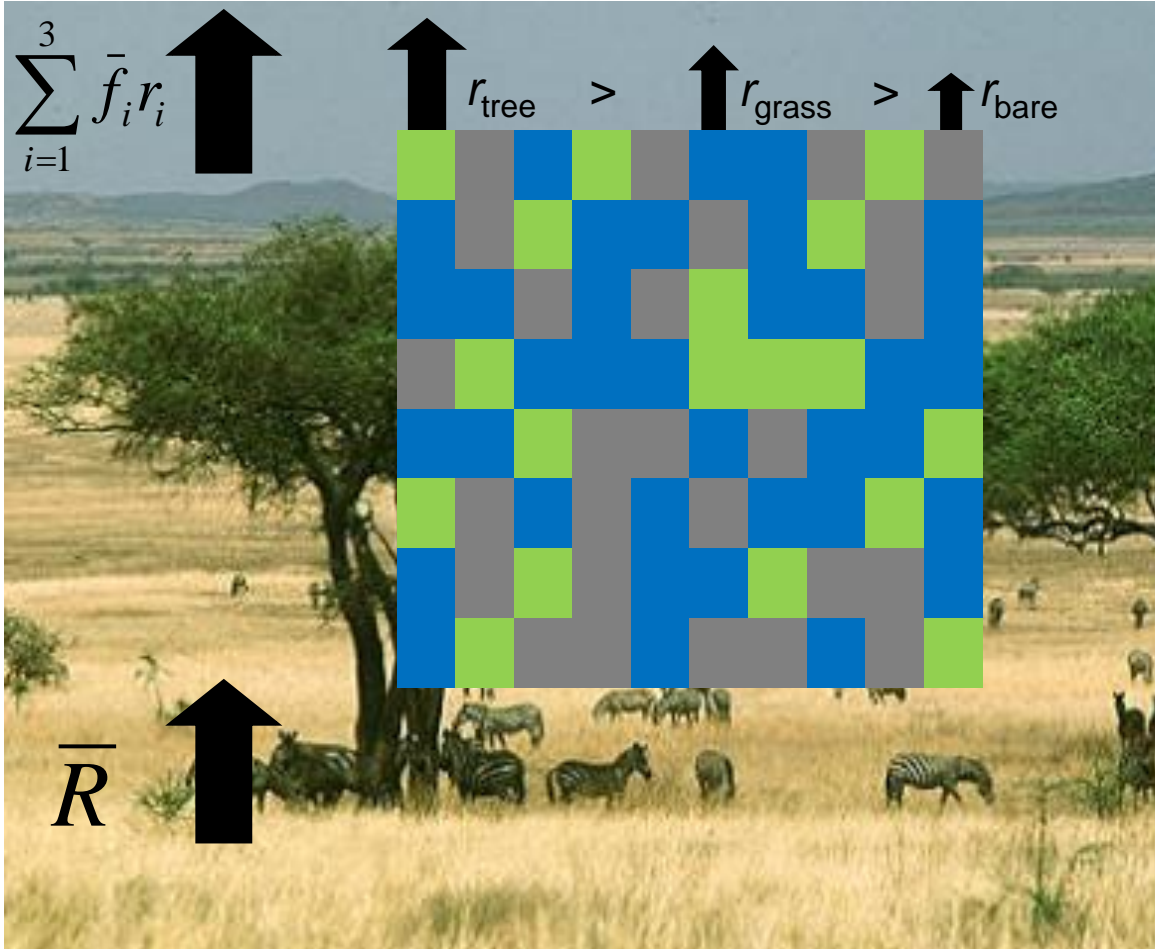
unifies different macroecological patterns:

SAD's

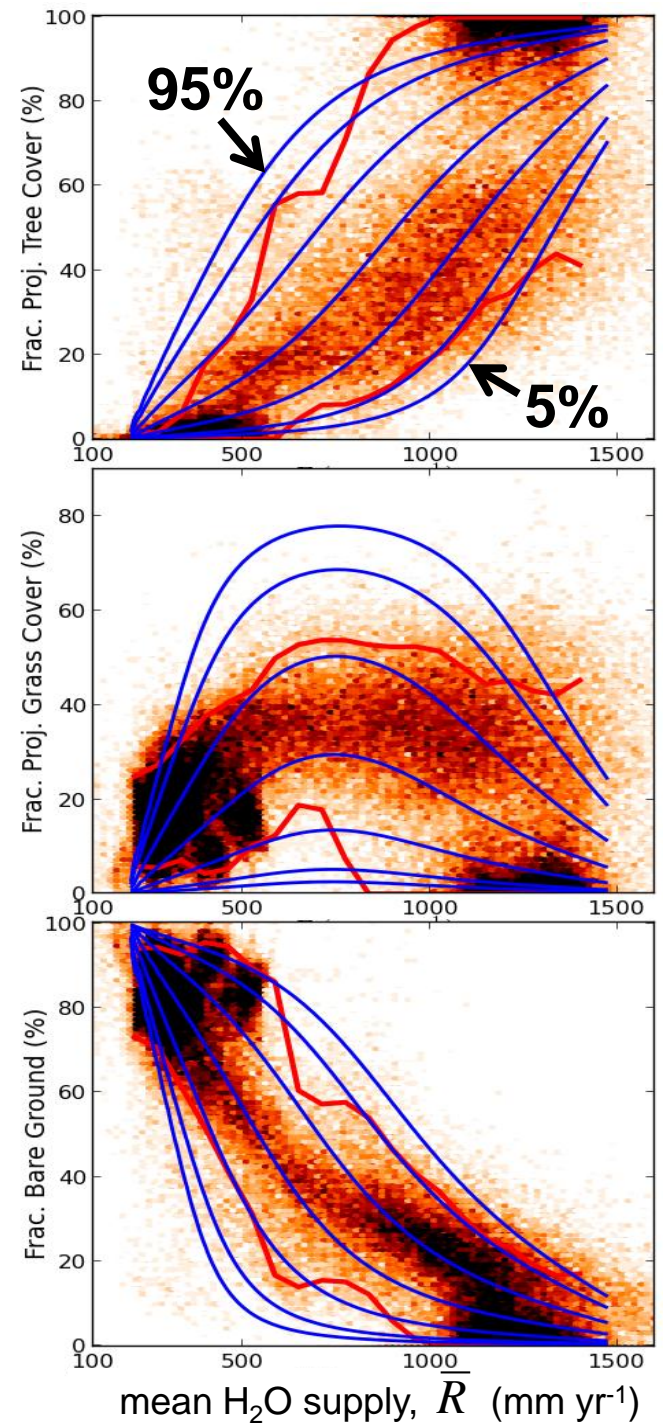
energy equivalence
diversity-productivity
diversity-stability

The statistical mechanics of savannas

Bertram & Dewar (2013)



For given \bar{R} what is most likely $p(f_{\text{tree}}, f_{\text{grass}})$?



Maximum Entropy Production (MEP)

Understanding Complex Systems

Springer:
COMPLEXITY

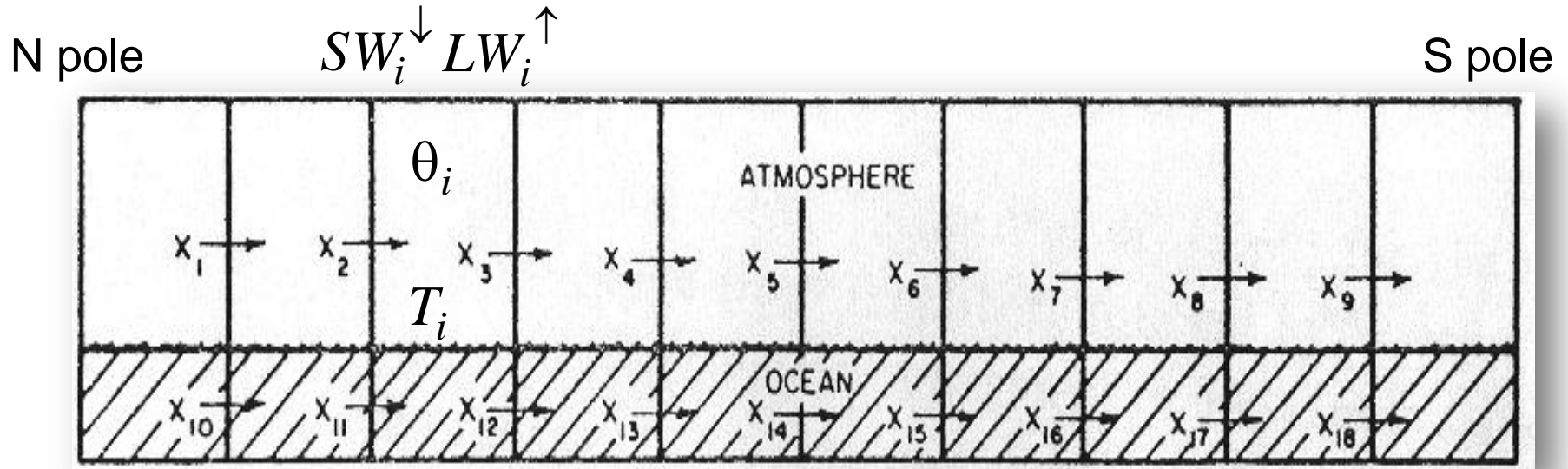
Roderick C. Dewar
Charles H. Lineweaver
Robert K. Niven
Klaus Regenauer-Lieb *Editors*

Beyond the Second Law

Entropy Production
and Non-equilibrium Systems

 Springer

Paltridge (1978): 10-zone energy balance model



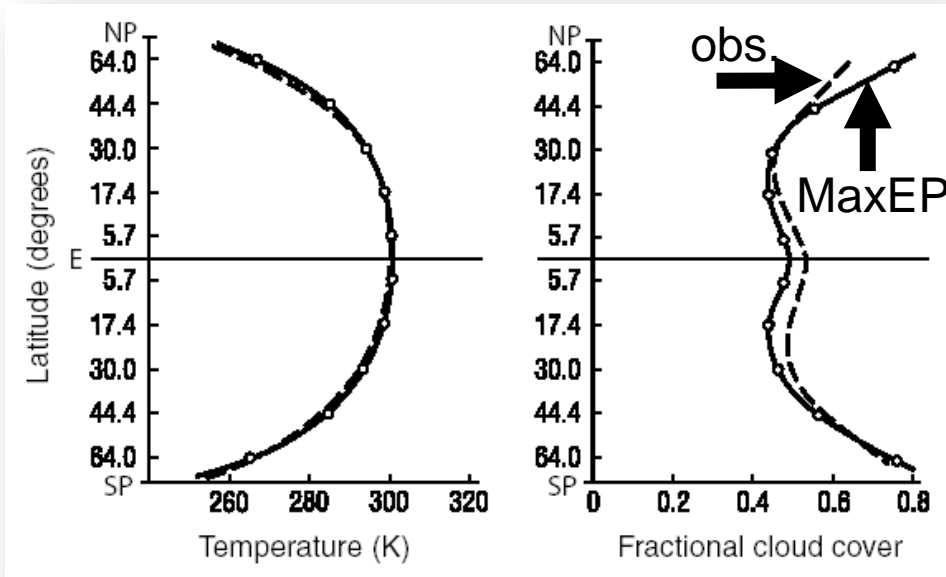
Maximize

$$EP \equiv \sum_{i=1}^{10} \frac{LW_i^\uparrow - SW_i^\downarrow}{T_i} = \sum_{i=1}^9 X_i \left(\frac{1}{T_{i+1}} - \frac{1}{T_i} \right)$$

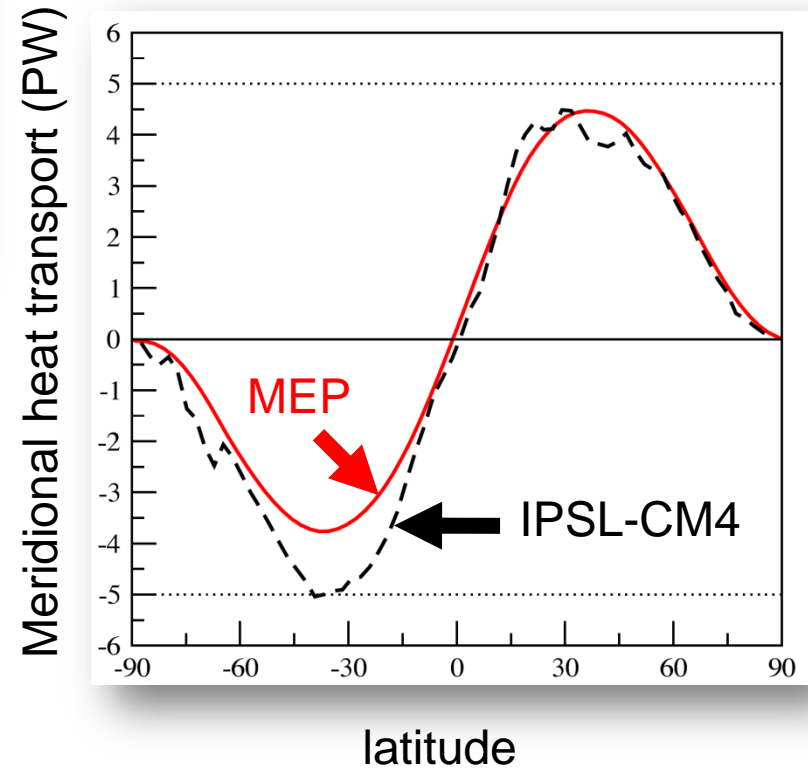
‘entropy production’
(it’s not!)

subject only to steady-state energy balance

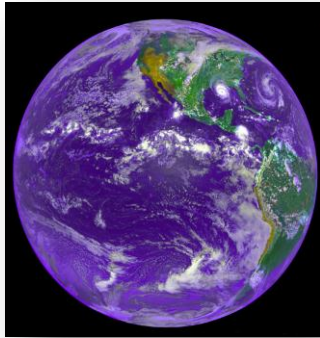
Paltridge (1978): 1D model



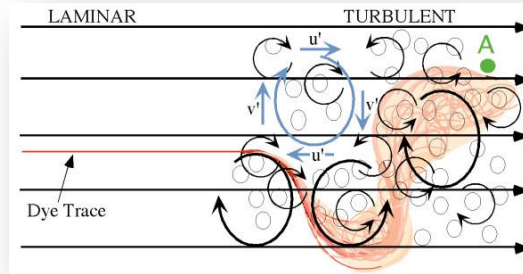
Herbert & Paillard (2013): 2D model



MEP applications across physics & biology



Paltridge (1978) ...

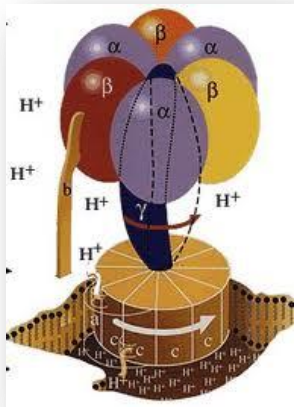


Malkus (2003) ...



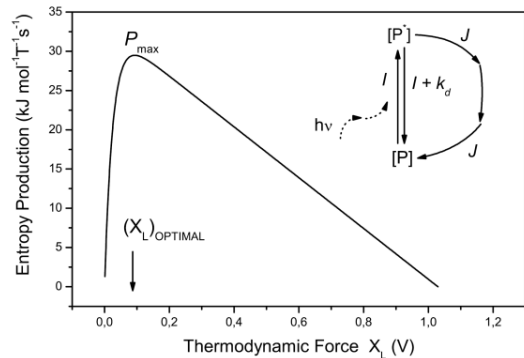
Main & Naylor (2008)

some intriguing successes,
but what does it mean?



Dewar et al (2006)

Juretić et al (2003)



Dewar (2010)



Martyushev et al (2000)

Some potentially misleading statements about Maximum Entropy Production

- MEP is a corollary to the second law ($dS_{\text{universe}}/dt \geq 0$) which states that S_{universe} increases as fast as possible
- MEP means that, over time, EP approaches a maximum in the steady-state

MEP as a statistical stability criterion for non-equilibrium stationary states



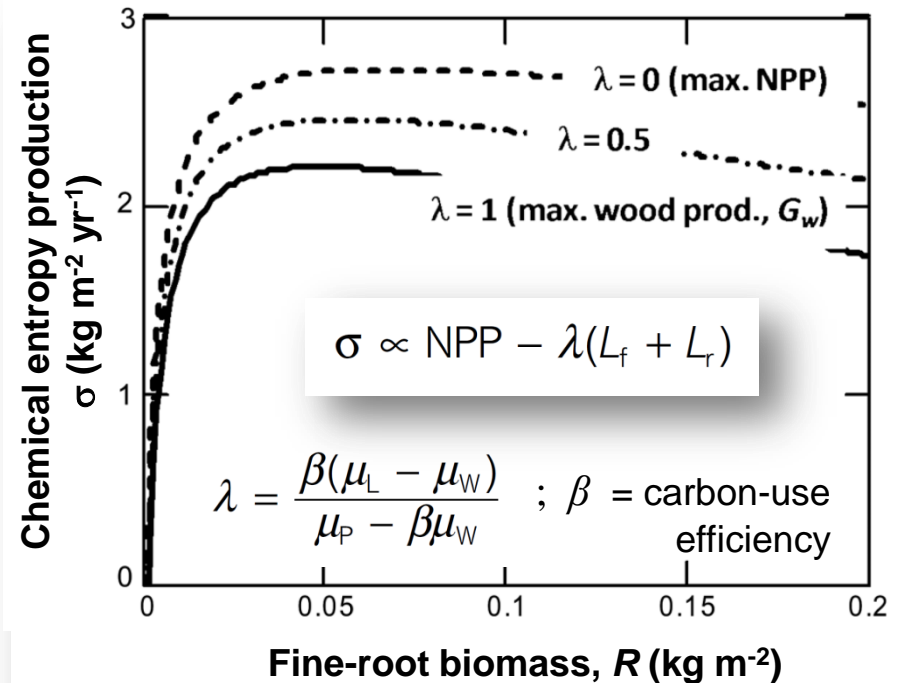
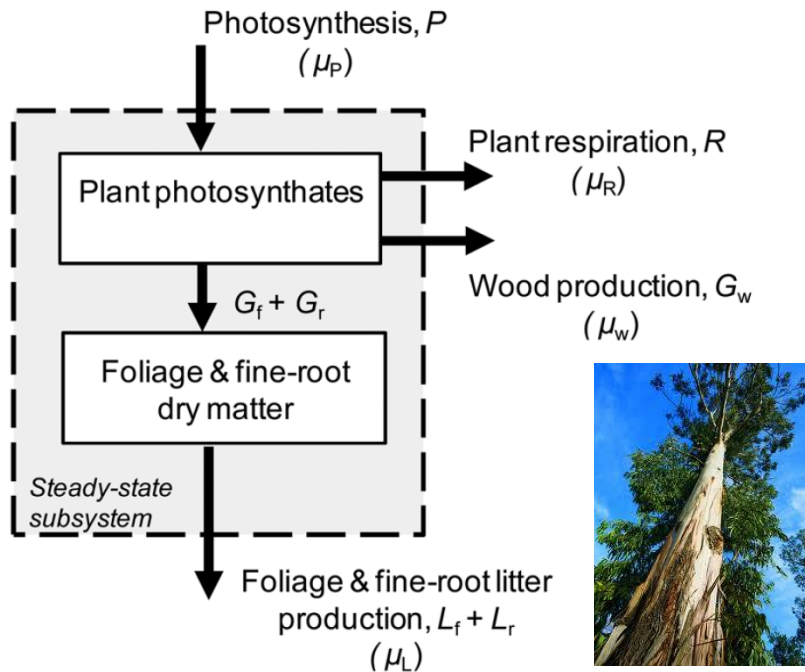
$$D = \sum_{\Gamma} p_{\Gamma} \ln \underbrace{\frac{p_{\Gamma}}{p_{\Gamma_R}}}_{d_{\Gamma}} \geq 0 \quad \text{with equality iff } p_{\Gamma_R} = p_{\Gamma}$$

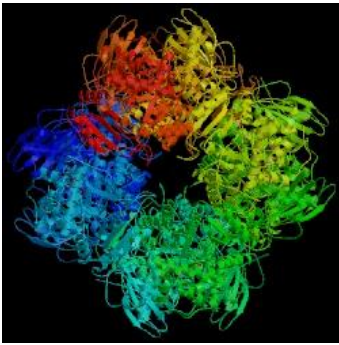
- D is a generic measure of irreversibility / time-reversal symmetry breaking / distance from equilibrium
- Fluctuation theorem (follows from definition of d_{Γ}): $\frac{p(d)}{p(-d)} = e^d$
- If $p(d) = N(D, \sigma^2)$: $\sigma^2 = 2D \quad \therefore \text{CV} = \frac{\sigma}{D} \propto \frac{1}{\sqrt{D}}$ is minimal when D is maximal
- Use MaxEnt to construct p_{Γ} : D depends on the constraints C

Modeling carbon allocation in trees: a search for principles

Oskar Franklin^{1,7}, Jacob Johansson^{1,2}, Roderick C. Dewar³, Ulf Dieckmann¹, Ross E. McMurtrie⁴, Åke Brännström^{1,6} and Ray Dybzinski⁵
Tree Physiol. **32**, 648-666 (2012)

MEP mimics traditional maximum fitness models





Evolutionary optimisation of Rubisco: Earth's most abundant protein

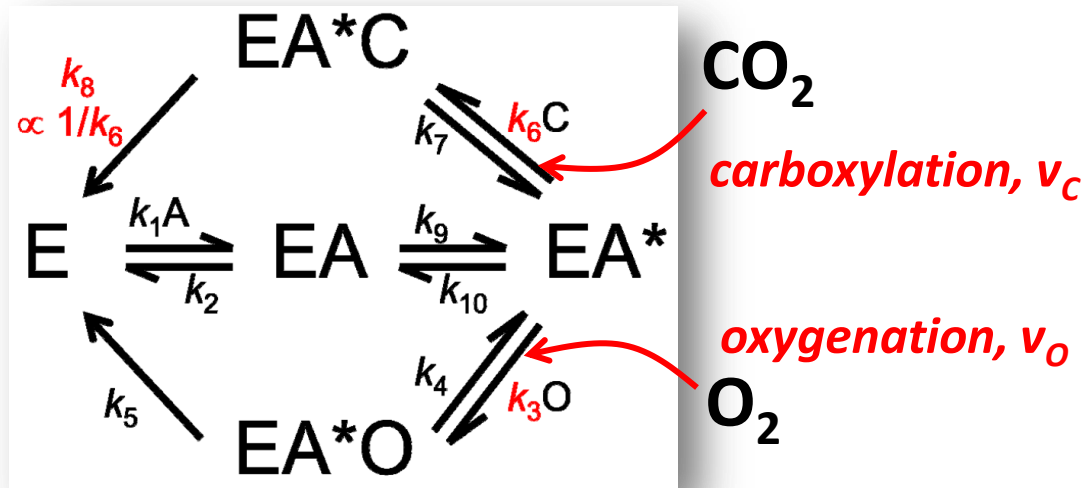
Enzyme state i

Kinetic design $\mathbf{k} = (k_1, k_2 \dots k_{10})$

Apply MaxEnt to $p(i, \mathbf{k}) = p(i|\mathbf{k})p(\mathbf{k})$

$$\therefore p(\mathbf{k}) \propto e^{H(\mathbf{k})}$$

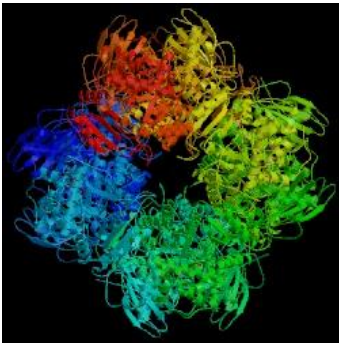
Enzyme kinetics



$$H(\mathbf{k}) = -\sum_i p(i|\mathbf{k}) \ln p(i|\mathbf{k})$$

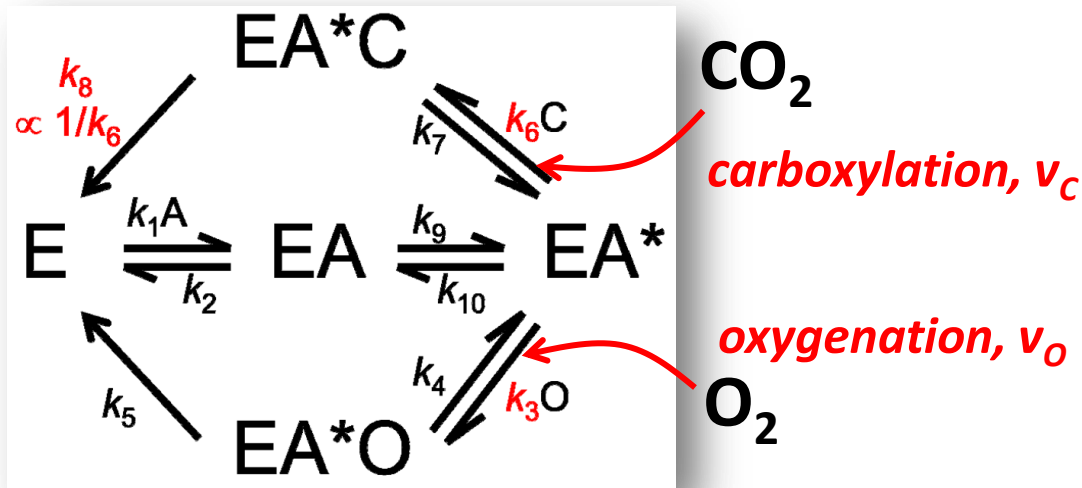
$$EP(\mathbf{k}) = v_c \ln \frac{v_c^+}{v_c^-} + v_o \ln \frac{v_o^+}{v_o^-}$$

Maximise simultaneously
w.r.t. k_6 & k_3

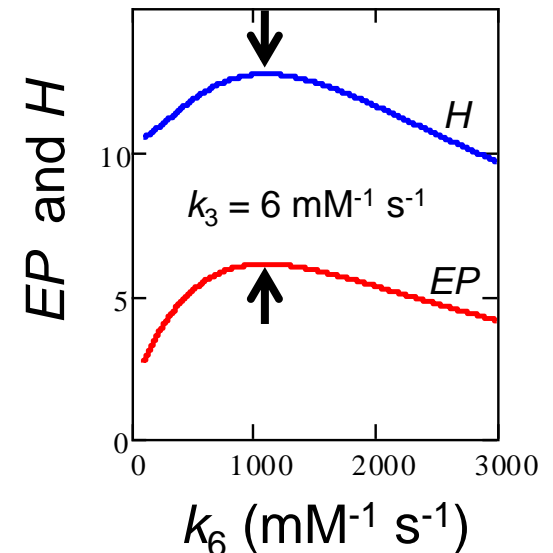


Evolutionary optimisation of Rubisco: Earth's most abundant protein

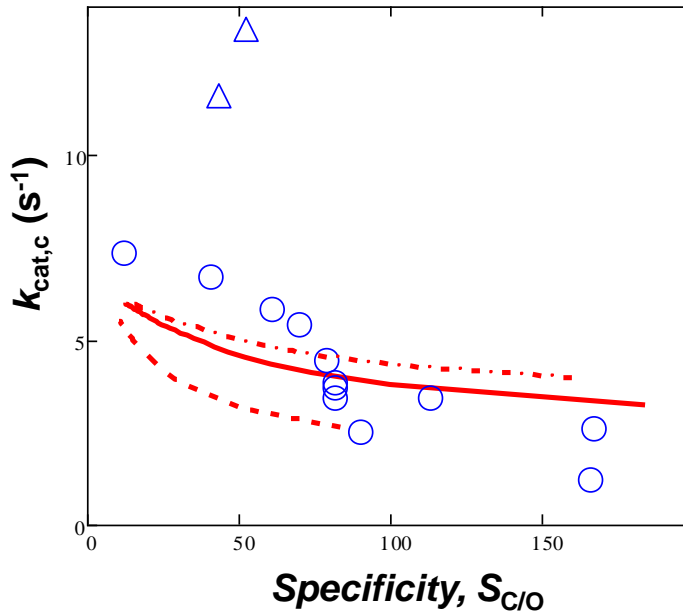
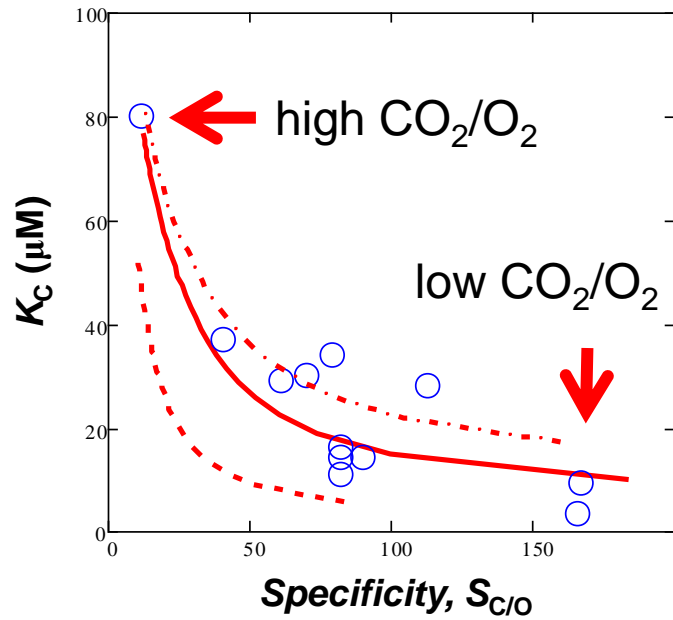
Enzyme kinetics



MaxEnt & MEP

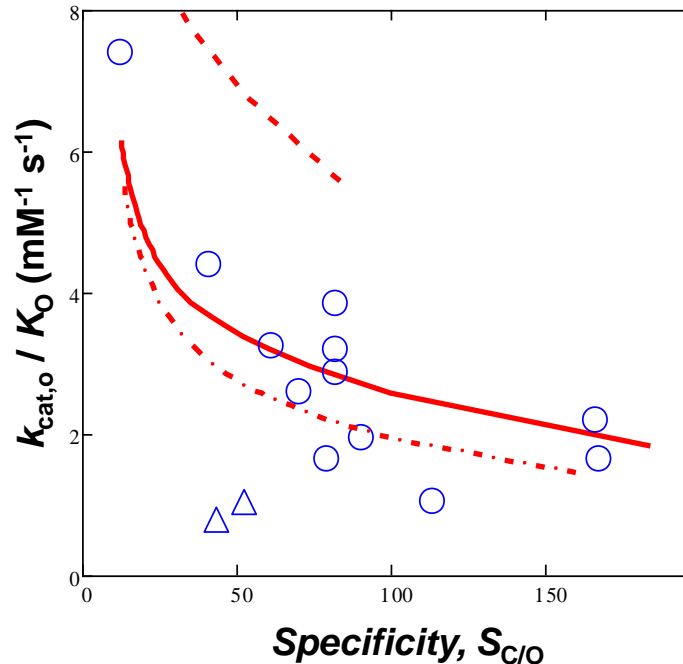
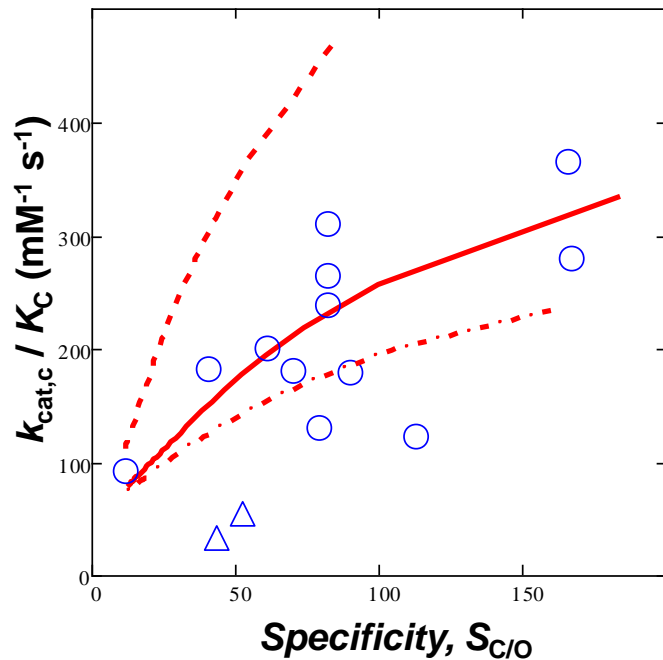


Rubisco adaptation to different CO_2/O_2 environments



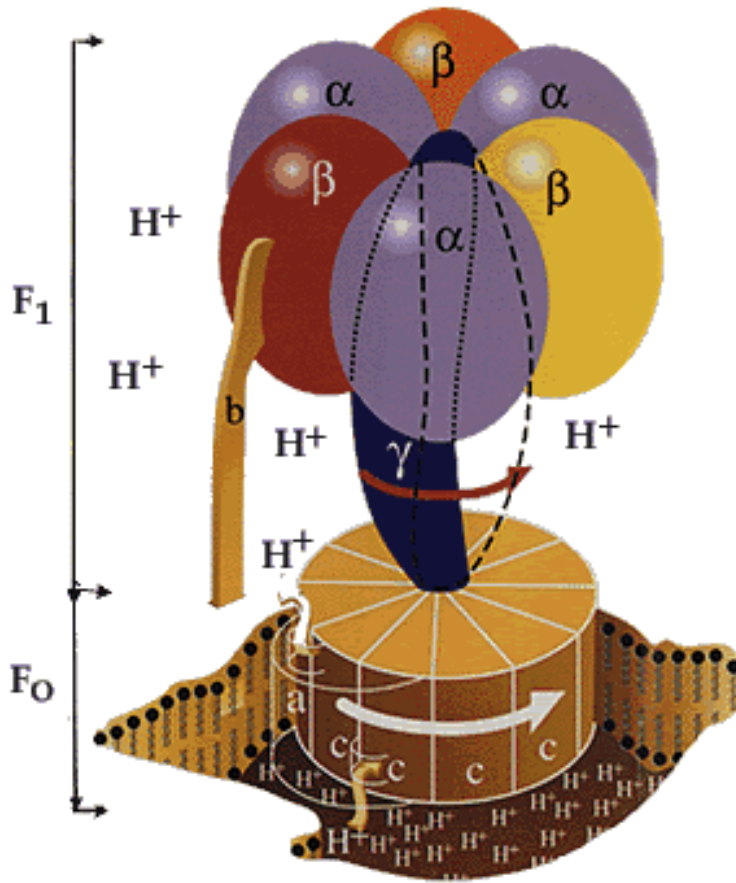
MaxEnt & MEP
predictions

- default parameters
- - - $k_7 \times 1/10$
- · - $k_4 \times 1/10$

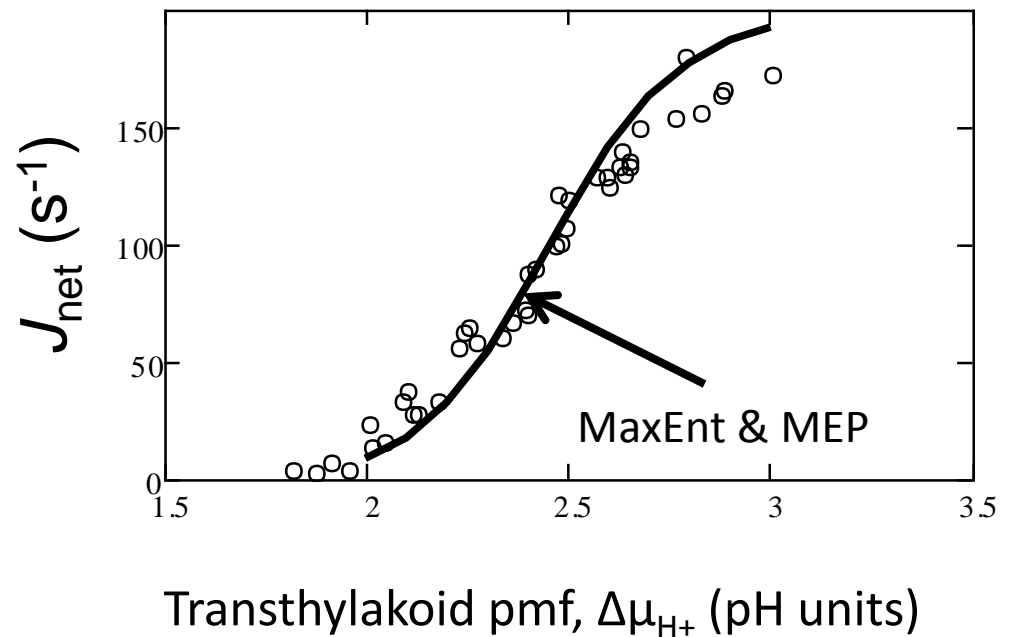


Dewar et al.
(in prep.)

F_0F_1 -ATP synthase : Nature's smallest rotary motor



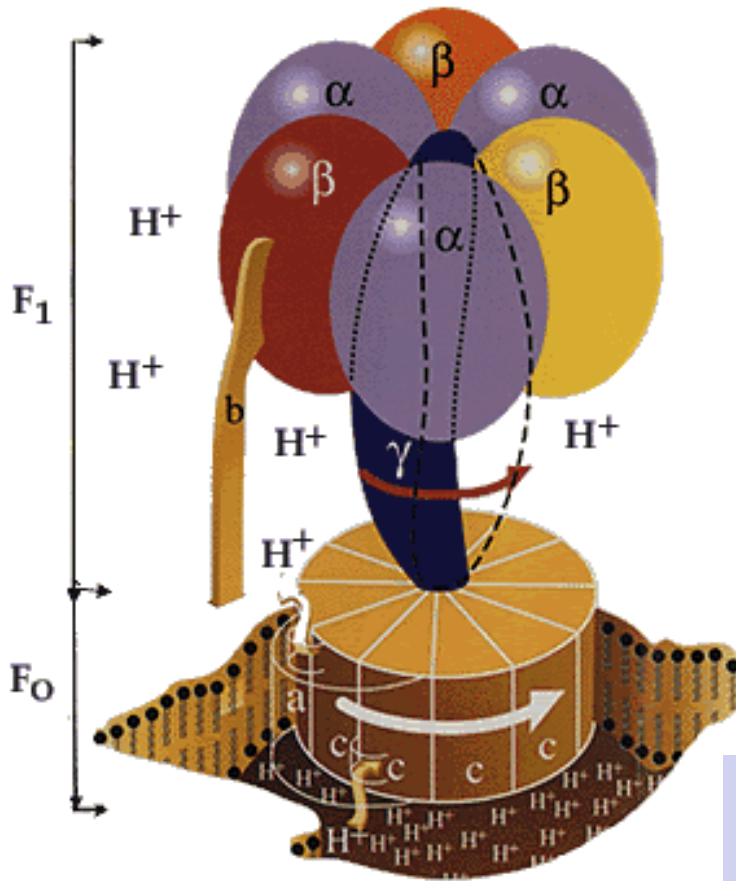
Net ATP synthesis vs. proton driving force



Dewar, Juretic & Zupanovic (2006)

Evolution of ATP-synthase kinetics

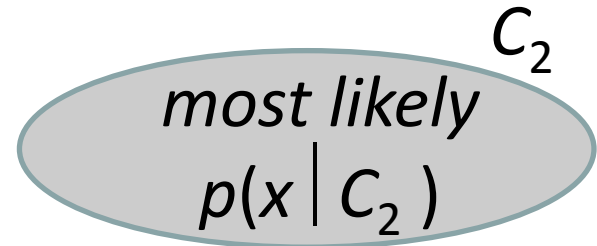
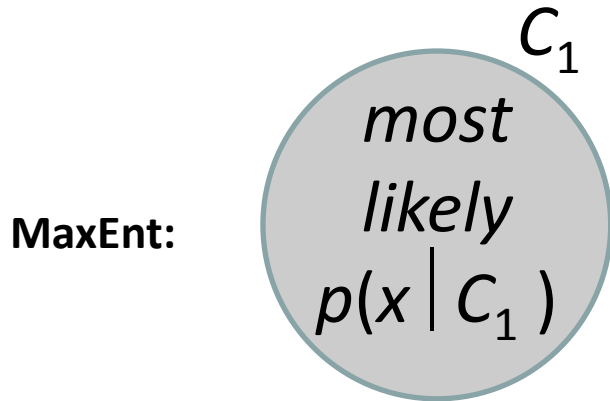
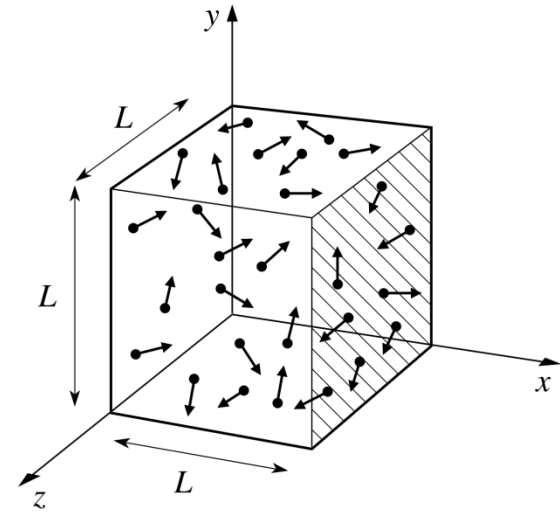
MaxEnt & MEP predict ...



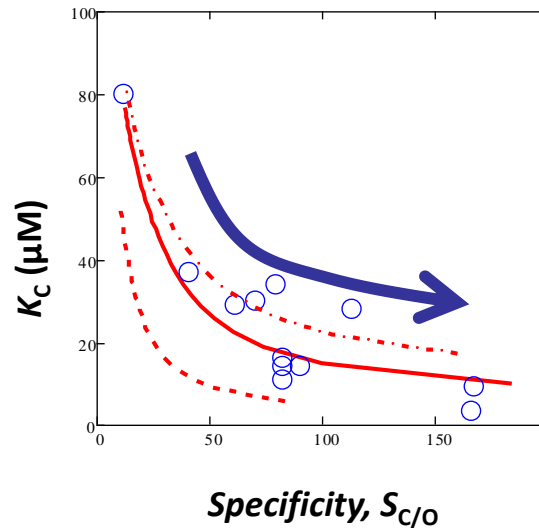
- ❑ optimal angular position for ATP synthesis close to observed (0.6)
- ❑ optimal gearing ratio $(\text{H}^+/\text{ATP}) \propto 1 / \text{pmf}$
- ❑ J_{net} maximally sensitive to pmf
- ❑ high free-energy conversion efficiency (69%) within the experimental range (50 – 80%)

→ observed kinetic design of ATP synthase consistent with most likely design

Optimal trait adaptation to changing constraints ($C_1 \rightarrow C_2$): survival of the likeliest



MEP: max stability ?



(cf. P - V - T gas laws)

Where next?

Theory

- Basis of MEP
- Non-stationary behaviour



Applications

- Ecological interactions: $r_i \rightarrow r_{ij}$
- Food webs
- Plant adaptations (e.g. leaf stomatal responses)