



מכון ויצמן למדע
WEIZMANN INSTITUTE OF SCIENCE

Compositional Lipid Assemblies as Evolving Protocells

Omer Markovitch and Doron Lancet

Department of Molecular Genetics,

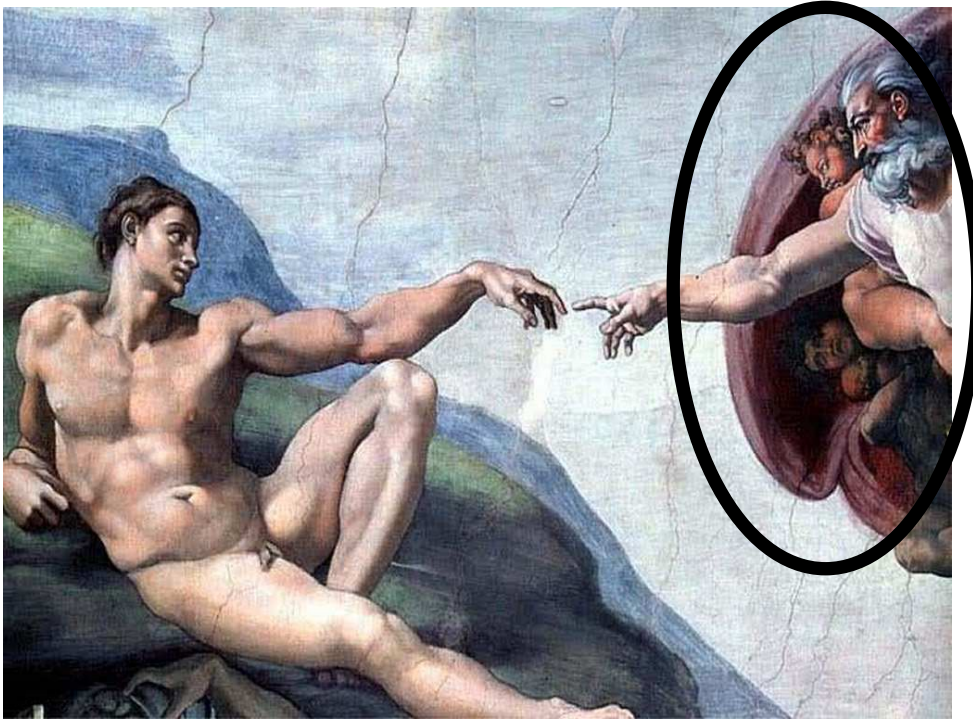
Weizmann Institute of Science,

Rehovot, Israel

<http://sites.google.com/site/omermar>

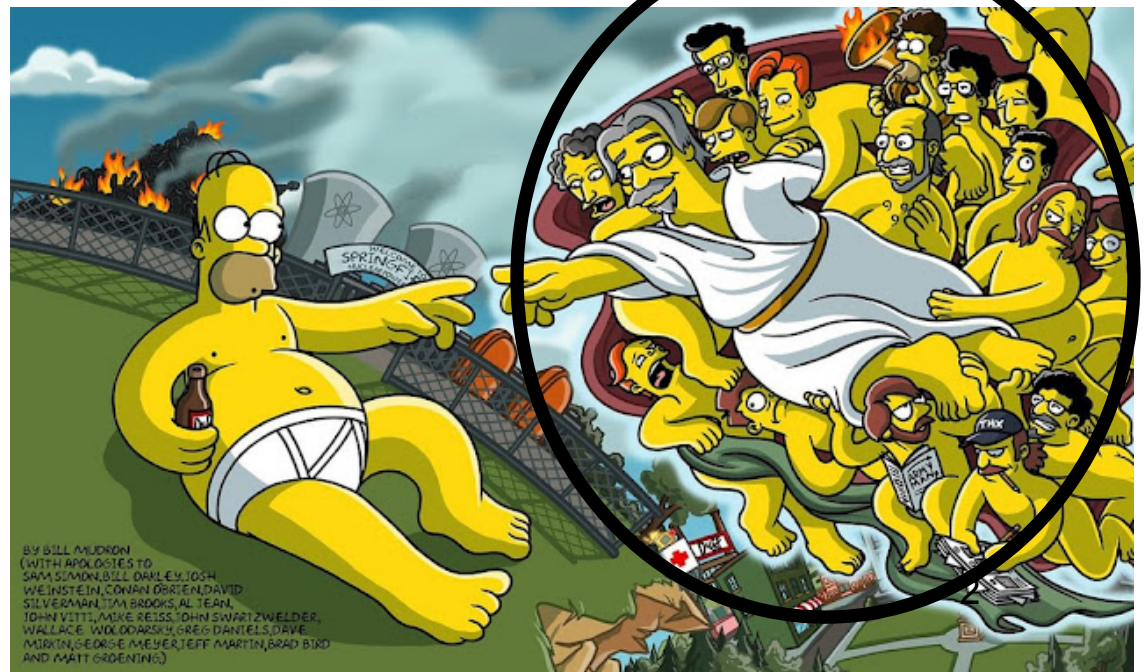
Origin of life workshop, ELSI, June 2013

ELSI EARTH-LIFE SCIENCE
INSTITUTE
地球生命研究所



Life

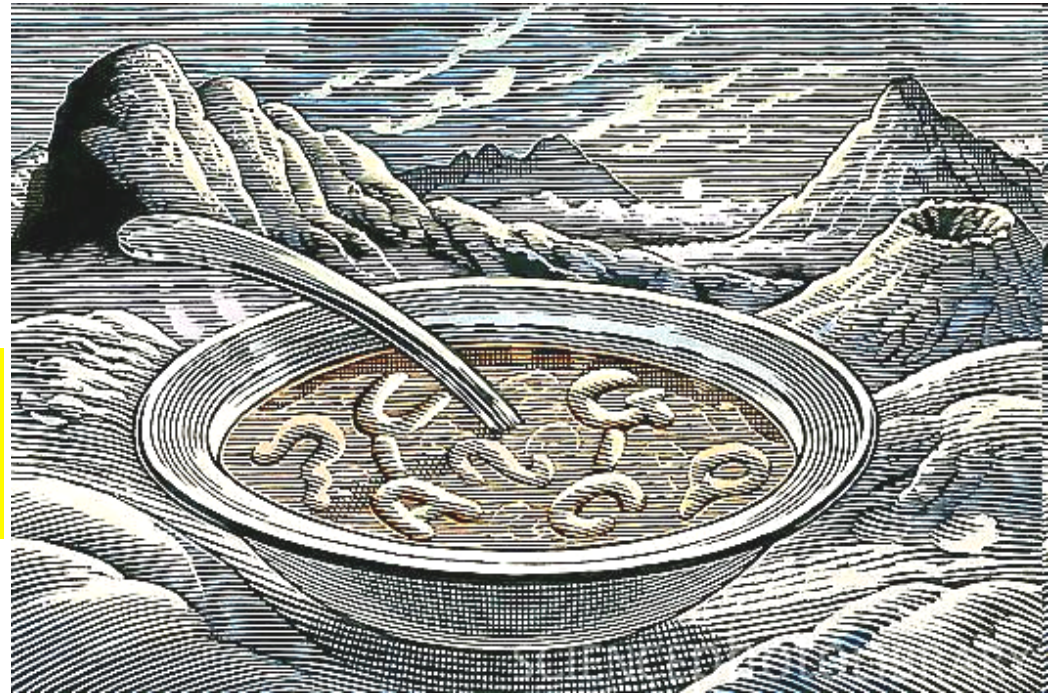
... is complex



“Prebiotic Soup”

~4,000,000,000 years ago

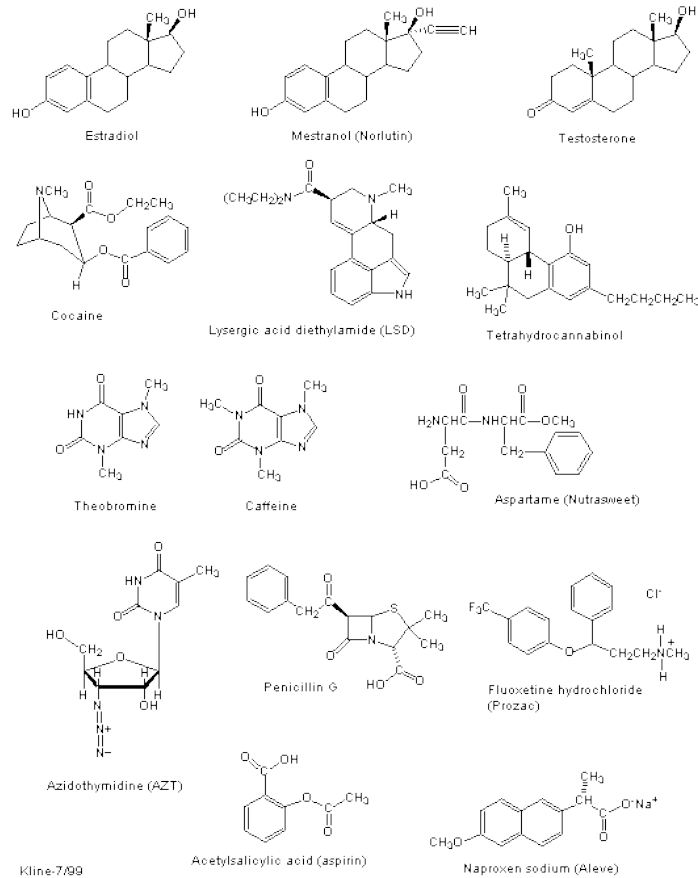
The emergence of the first cell-like entity, the *Protocell*.



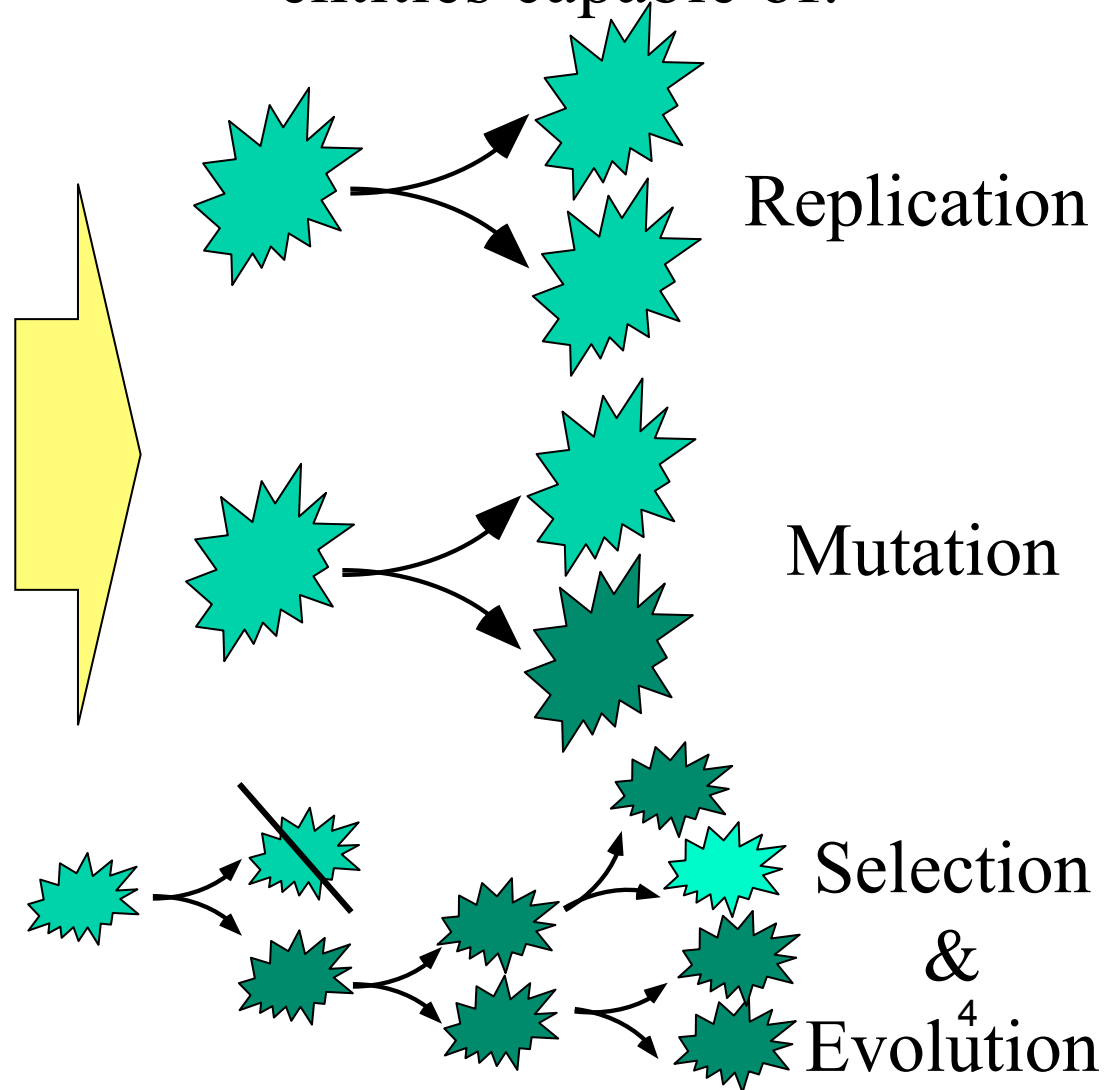
Life is a self-sustaining system capable of undergoing Darwinian evolution.

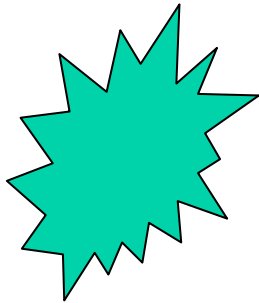
The problem

Organic molecules



More complex chemical entities capable of:

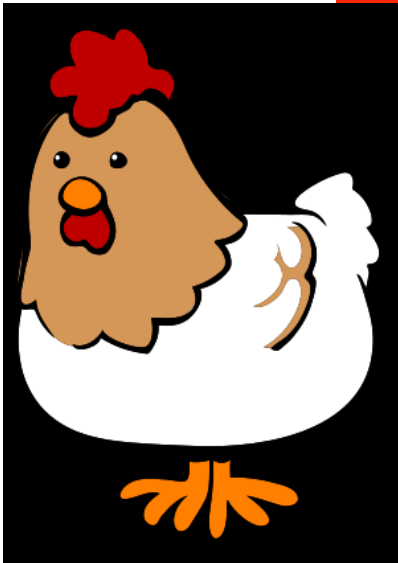


What is  ?

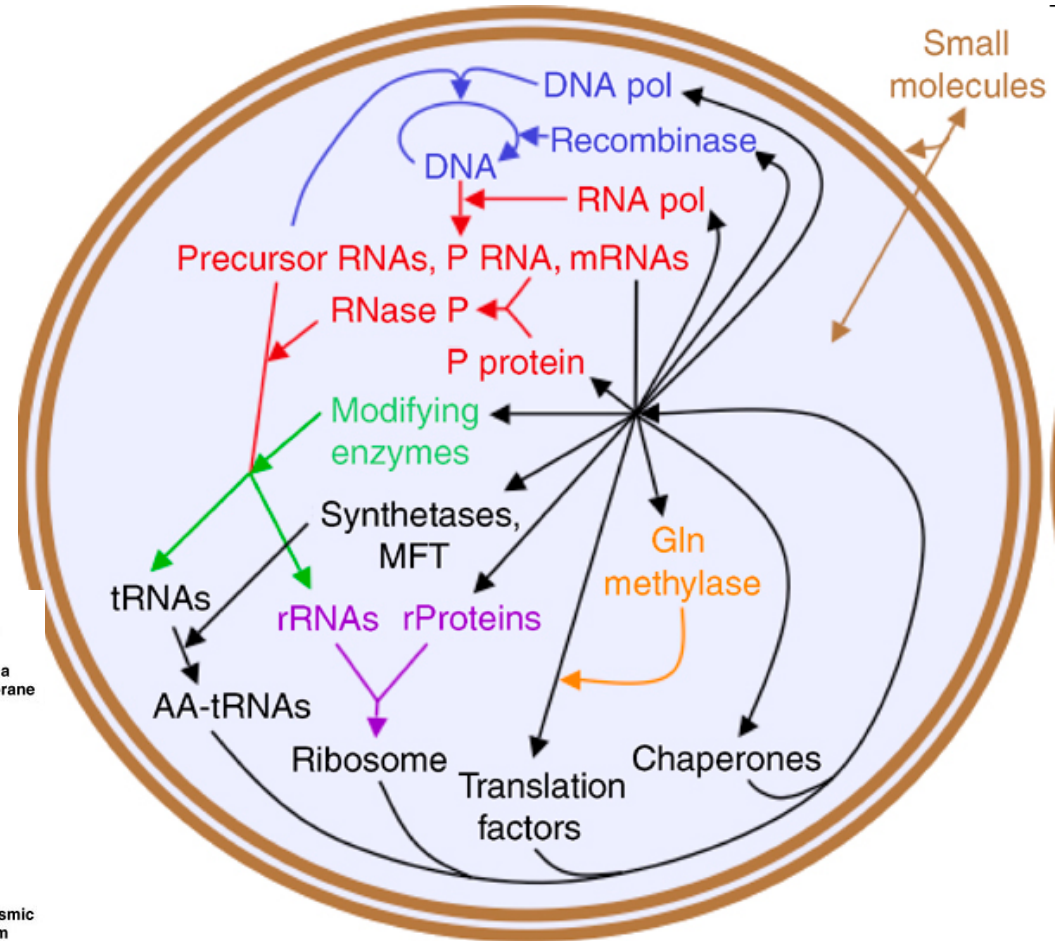
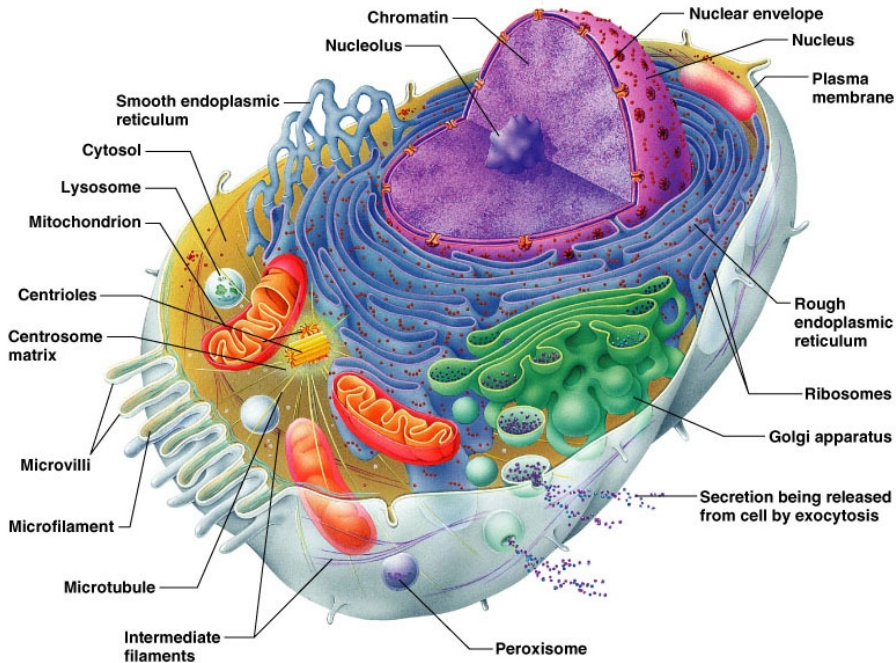
Occam's razor –
simple

Life requires **information**.
Information undergoes evolution.

Which came first: chicken or egg?
Chicken + Egg = Chegg 😊



The Cell as a Complex Network



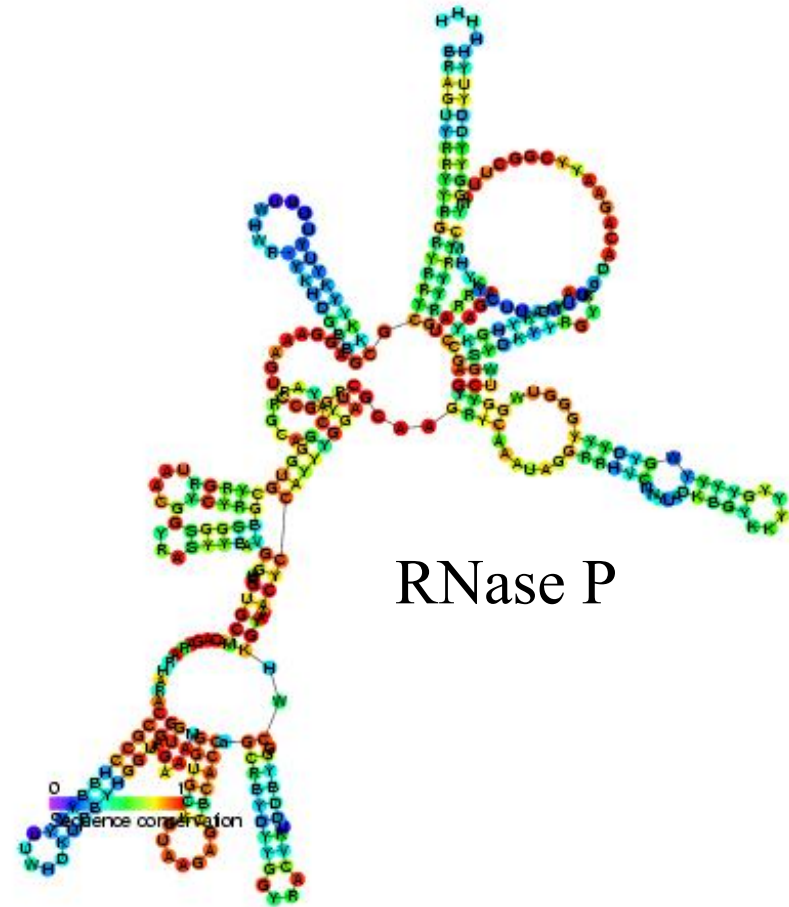
RNA World / Replicator-First

Altman and Cech – self-cleaving Ribozyme (Nobel prize in Chemistry, 1989).

Could  be a ribozyme?

Strongly depends on **covalent** bonds

- Difficult to form
- Mutation = breaking and remaking two **covalent** bonds



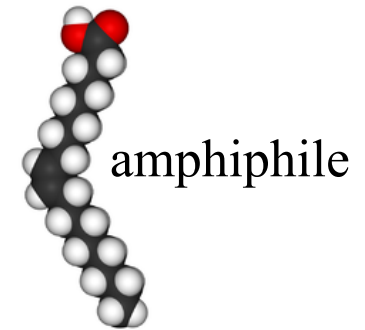
The Lipid World

Could  be a Micelle?

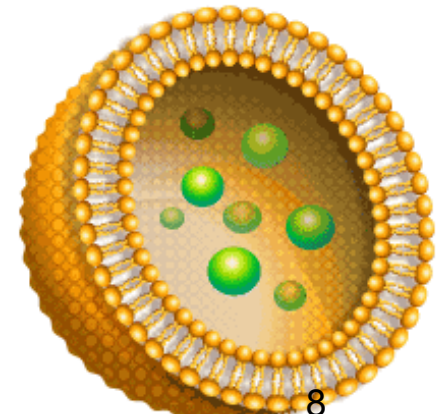
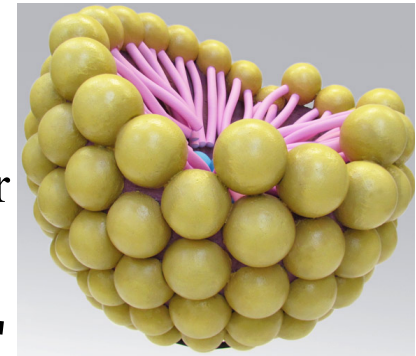
Held together by **non-covalent** bonds.

- Forms spontaneously
- Mutation = “random access” lipid entry/exit

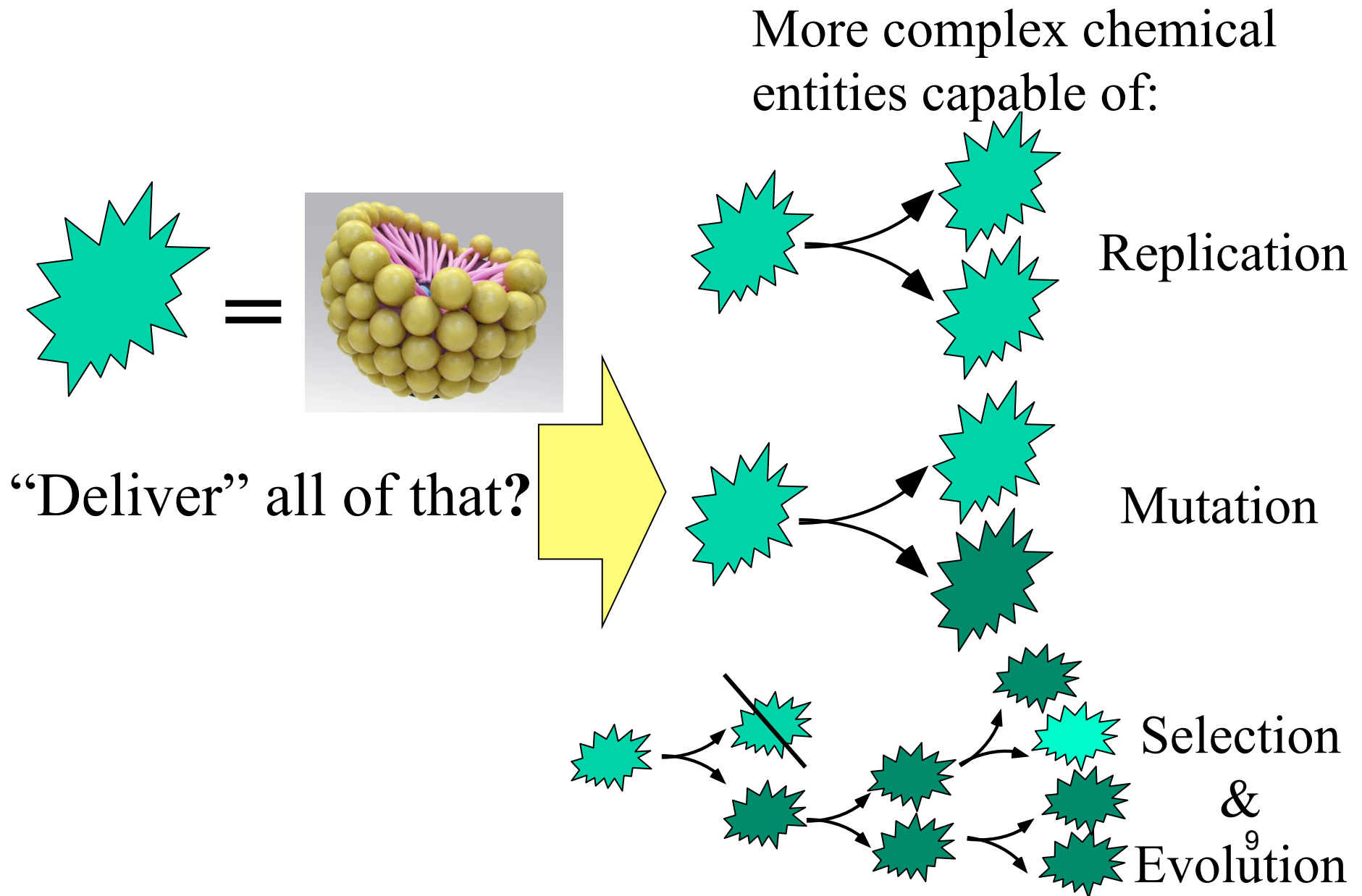
Much simpler!

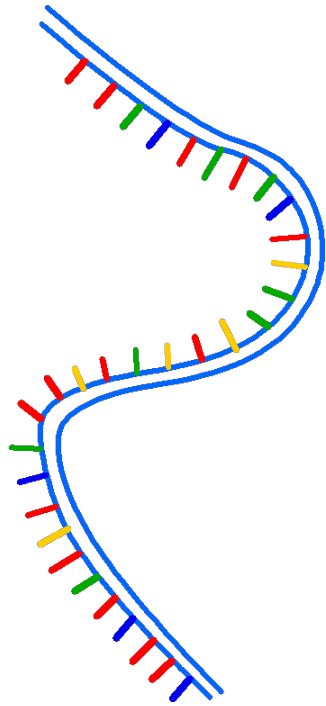


Micelle: 10^{-8} meter

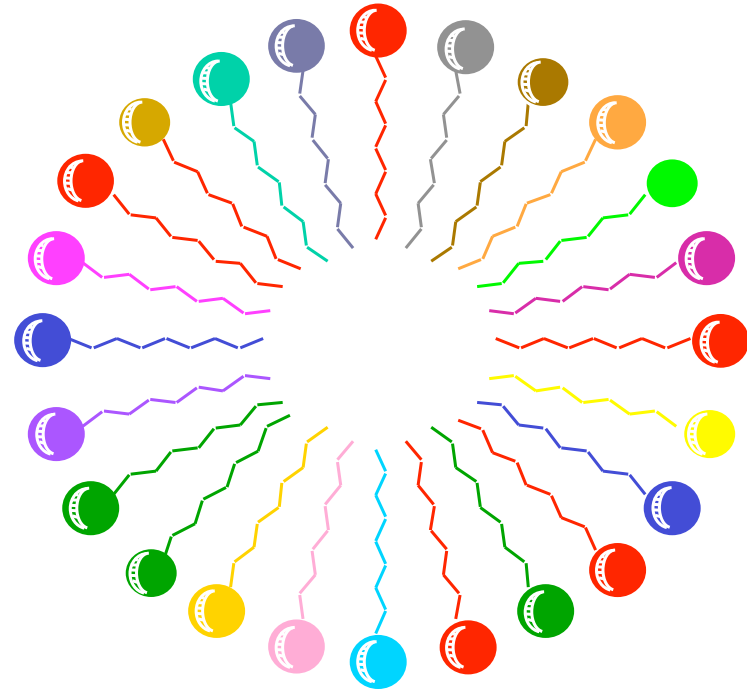


The Lipid World





DNA / RNA / Polymers →
Sequence



Assemblies / Clusters /
Vesicles / Membranes →
Composition

Sequential vs. Compositional Information

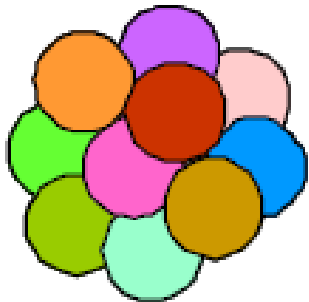


Alphabet: 20 amino acids



10-letters long polymer

$$\log_2(20^{10}) \approx 43 \text{ bits}$$

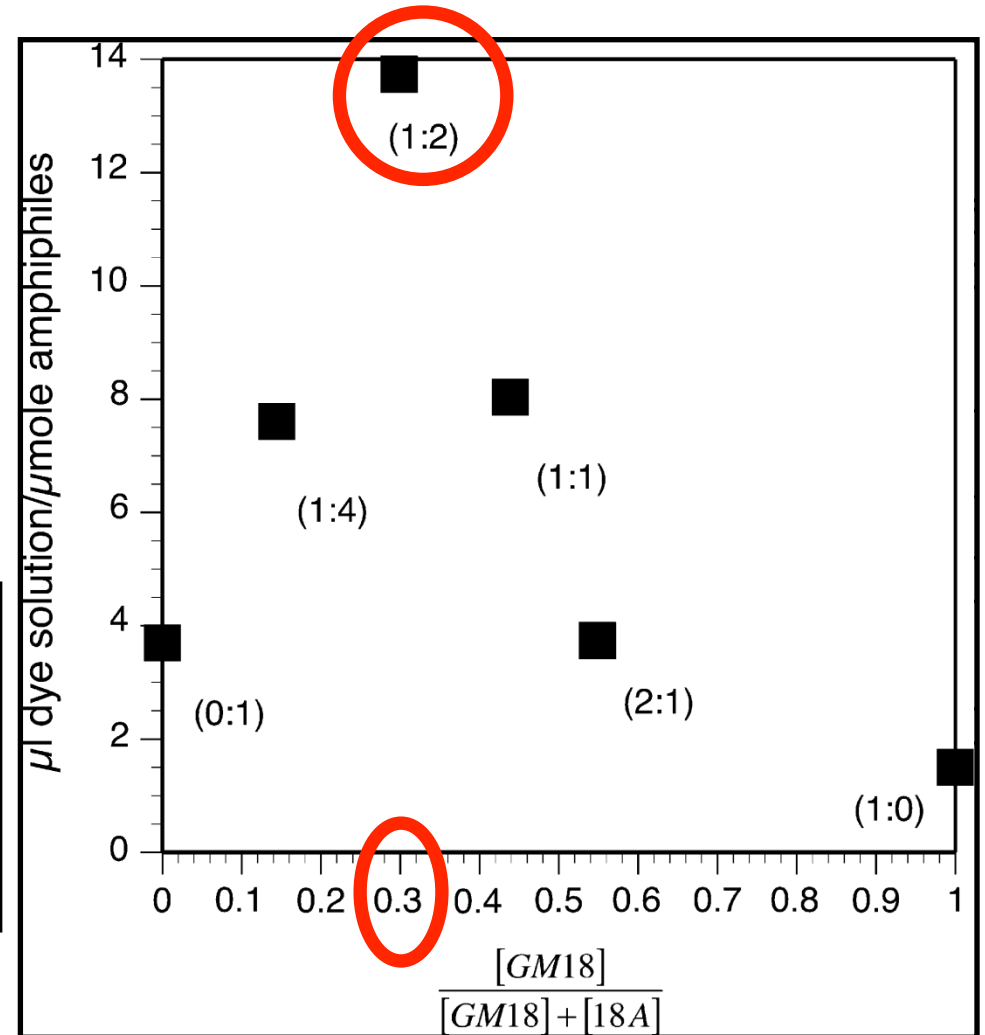
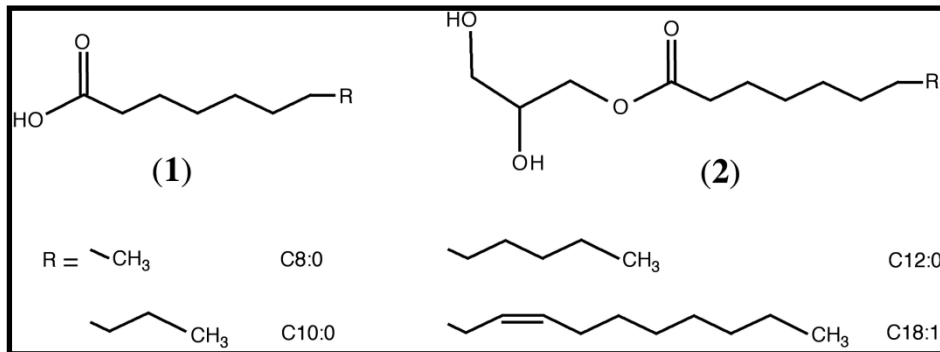


10 molecules
assembly

$$\log_2\left(\frac{(20 + 10 - 1)!}{10!}\right) \approx 27 \text{ bits}$$

Composition effect vesicle encapsulation-efficiency

Phenotype

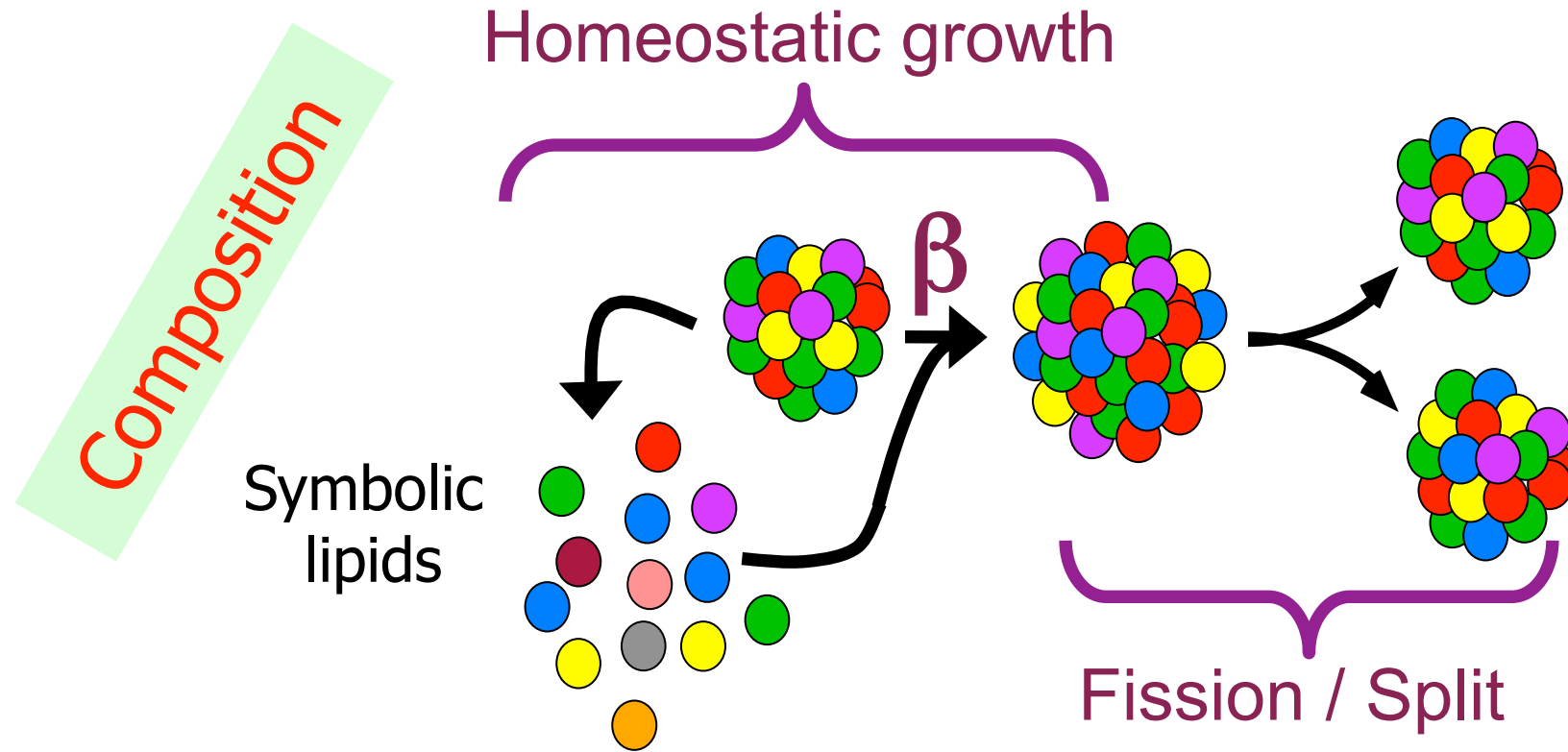




The evolution of authority



GARD model (Graded Autocatalysis Replication Domain)



$$\frac{dn_i}{dt} = (k_f \rho_i N - k_b n_i) \left(1 + \sum_{j=1}^{N_G} \beta_{ij} \frac{n_j}{N} \right)$$

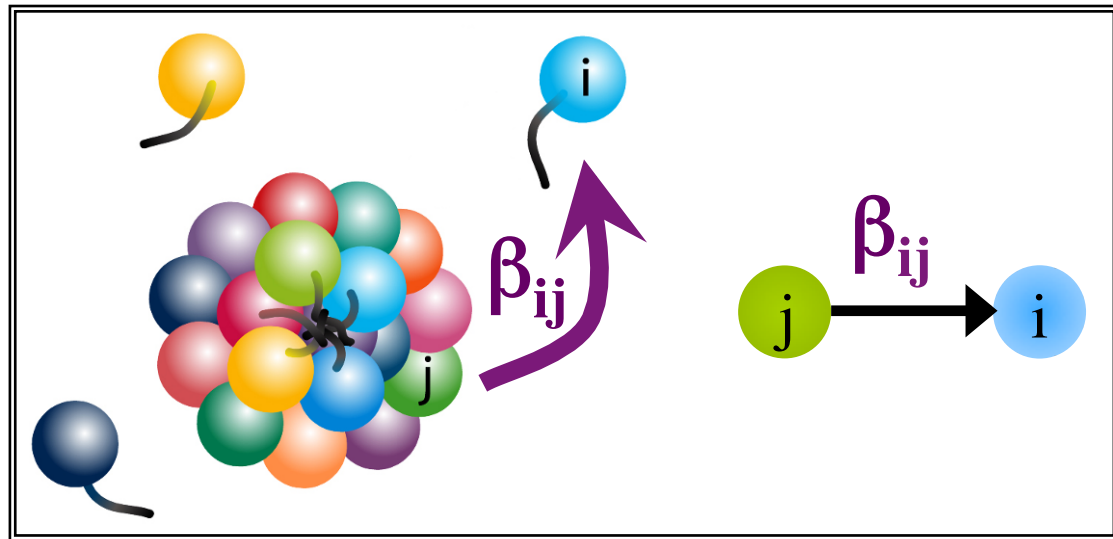
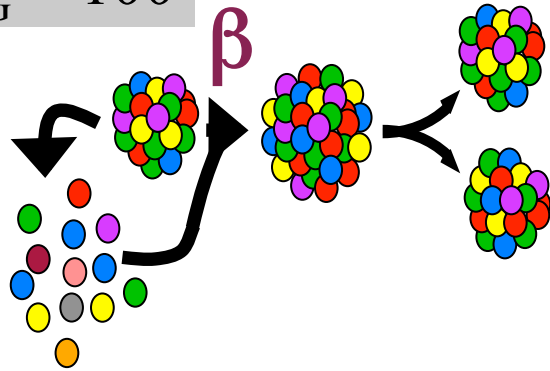
$$(i = 1..N_G)$$

Rate enhancement

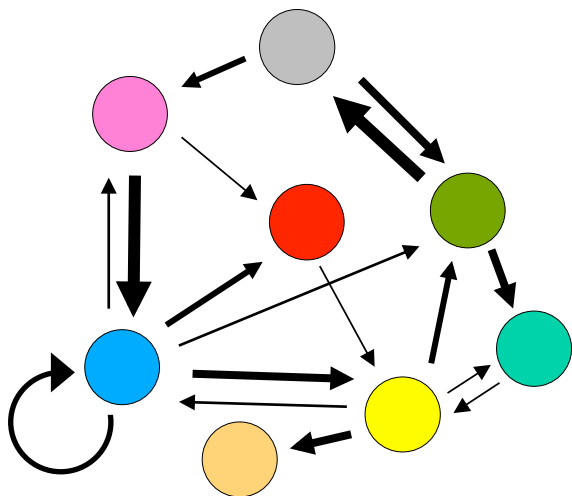
Molecular repertoire

β ; Catalytic Network (environmental chemistry)

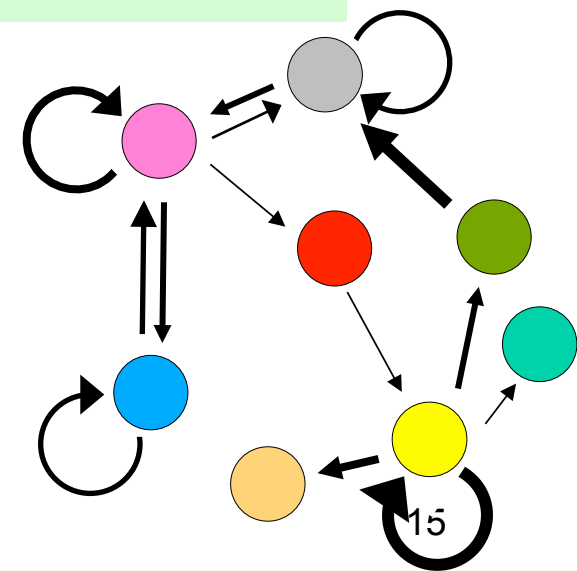
$N_G = 100$



More mutualistic



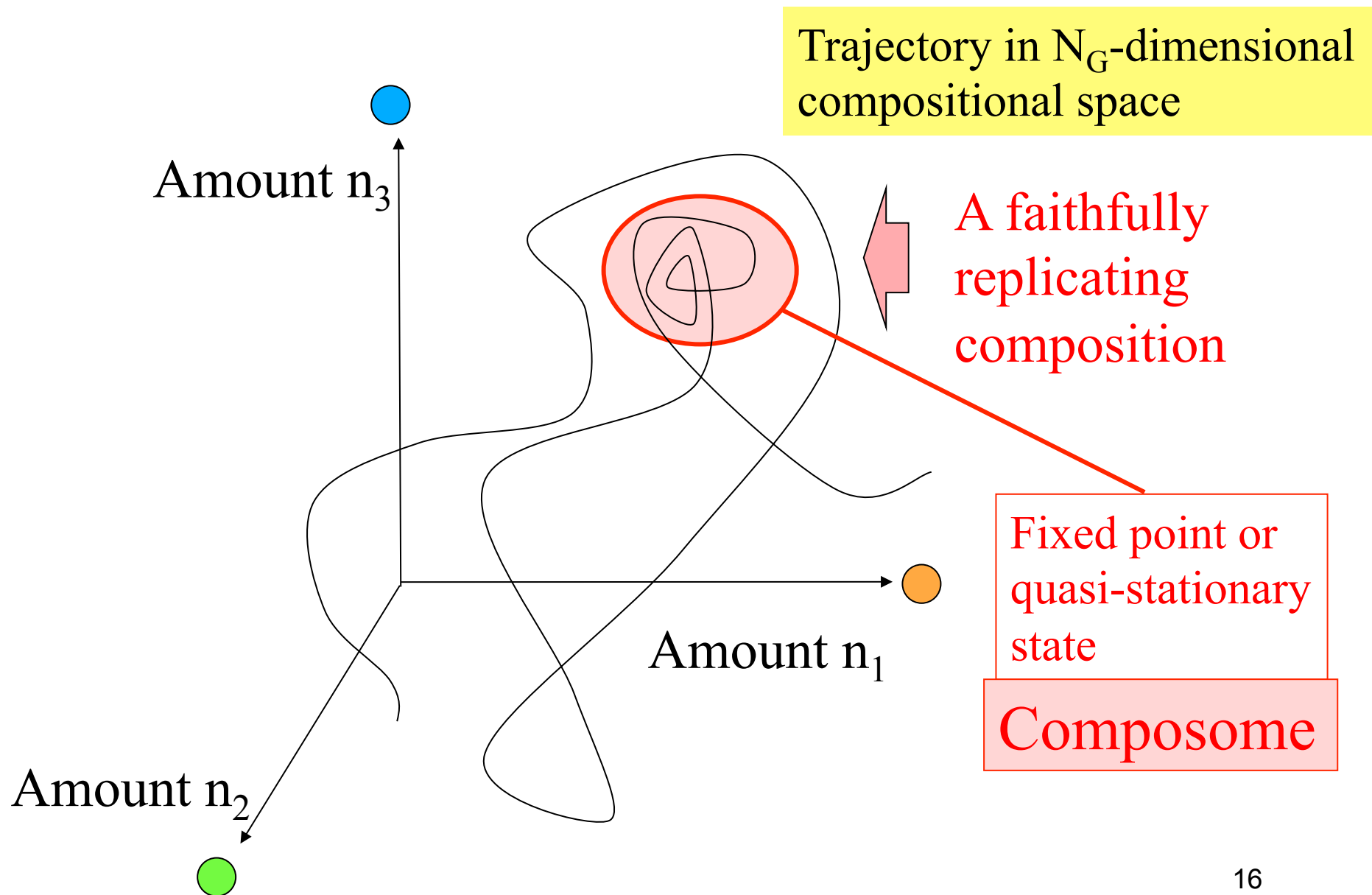
More selfish



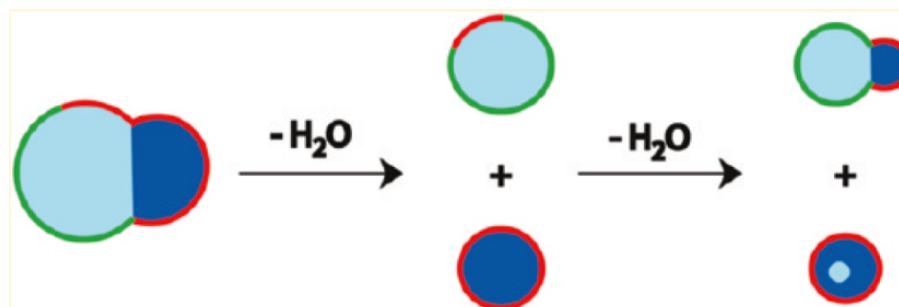
*Self-catalysis is the chemical manifestation of self-replication [Orgel, Nature 358 (1992)]

Dyson, Gánti, Kauffman, Varela

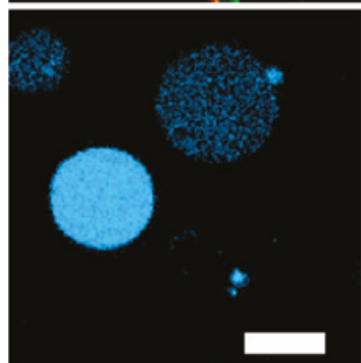
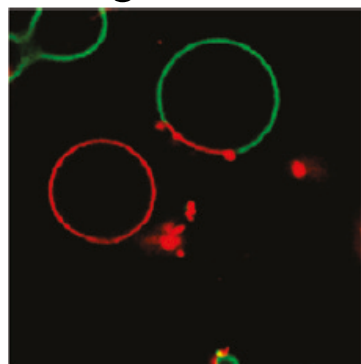
GARD Dynamics



Experimental Vesicle Heredity



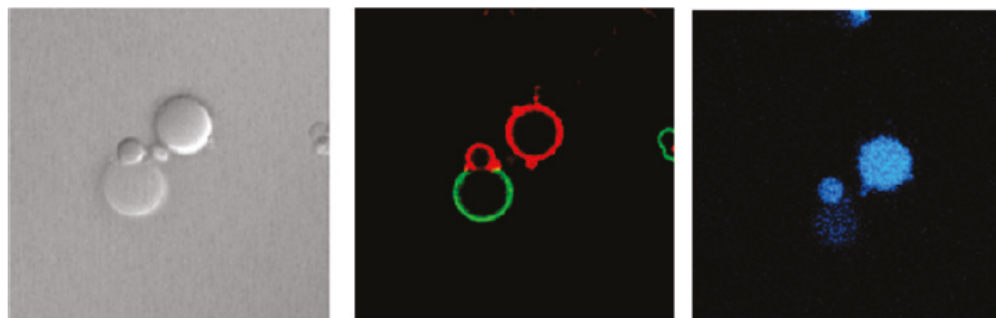
First generation



10^{-6} meter

Membrane composition was inherited by daughter vesicle, and affected daughter fission.

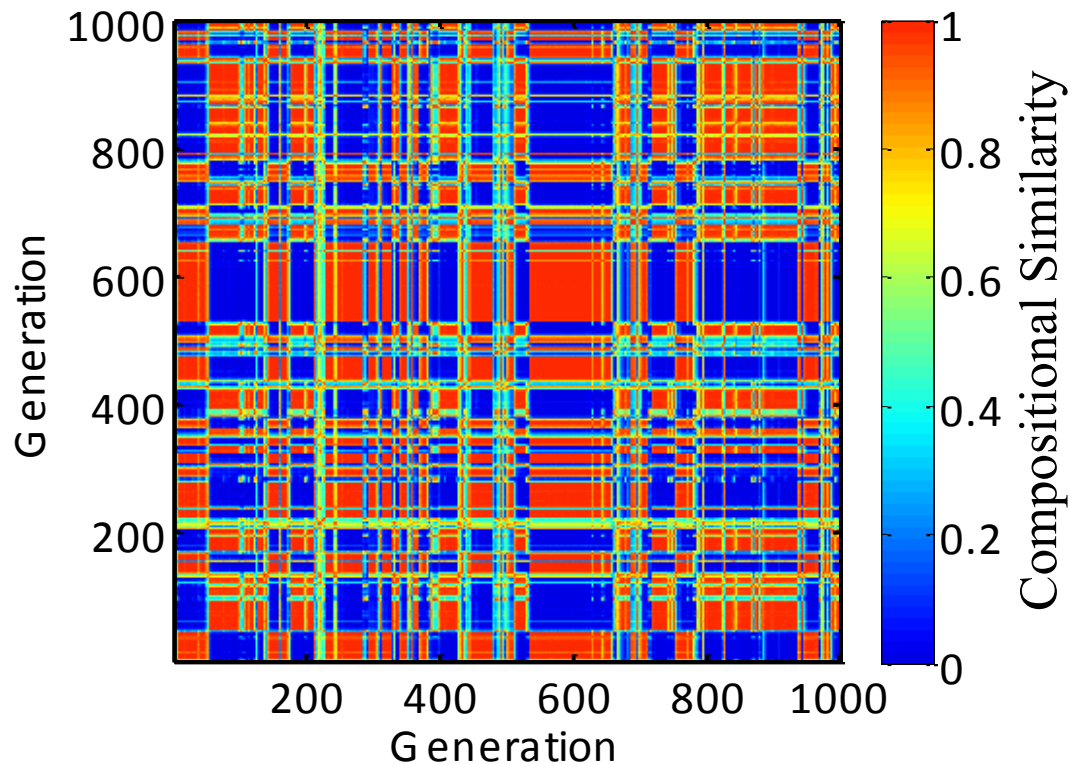
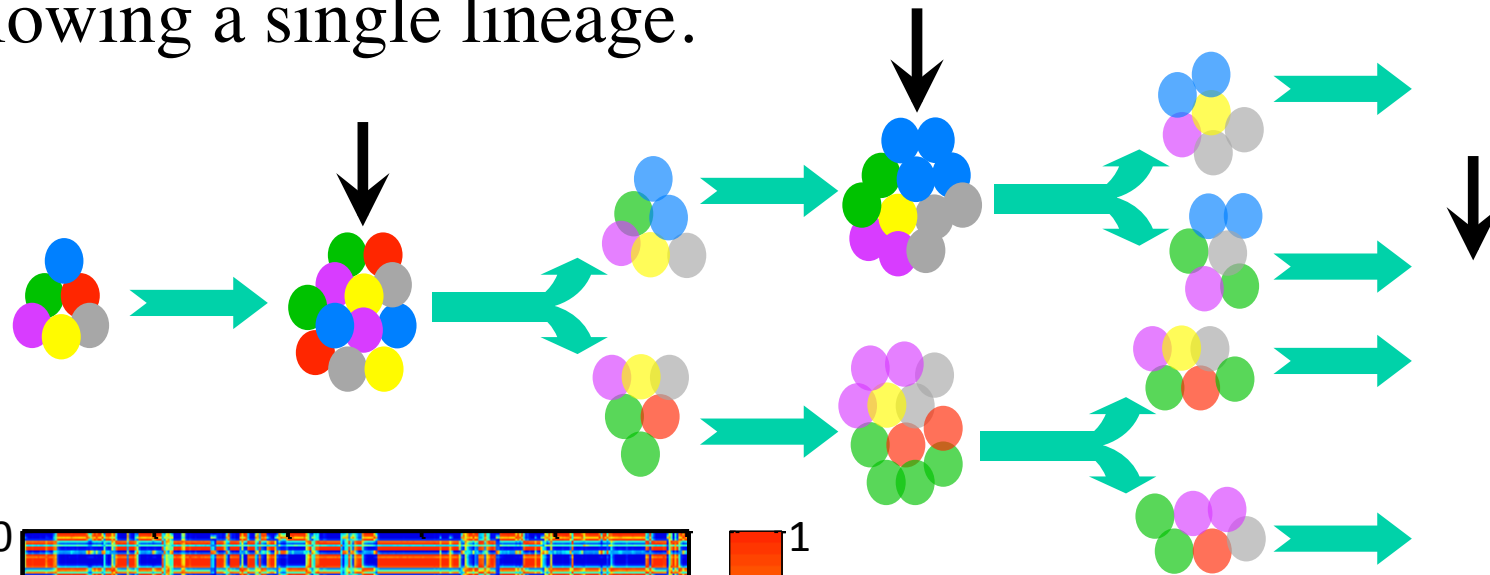
Second generation



Andes-Koback and Keating, JACS 133¹⁷ (2011)

Example of GARD Similarity ‘Carpet’

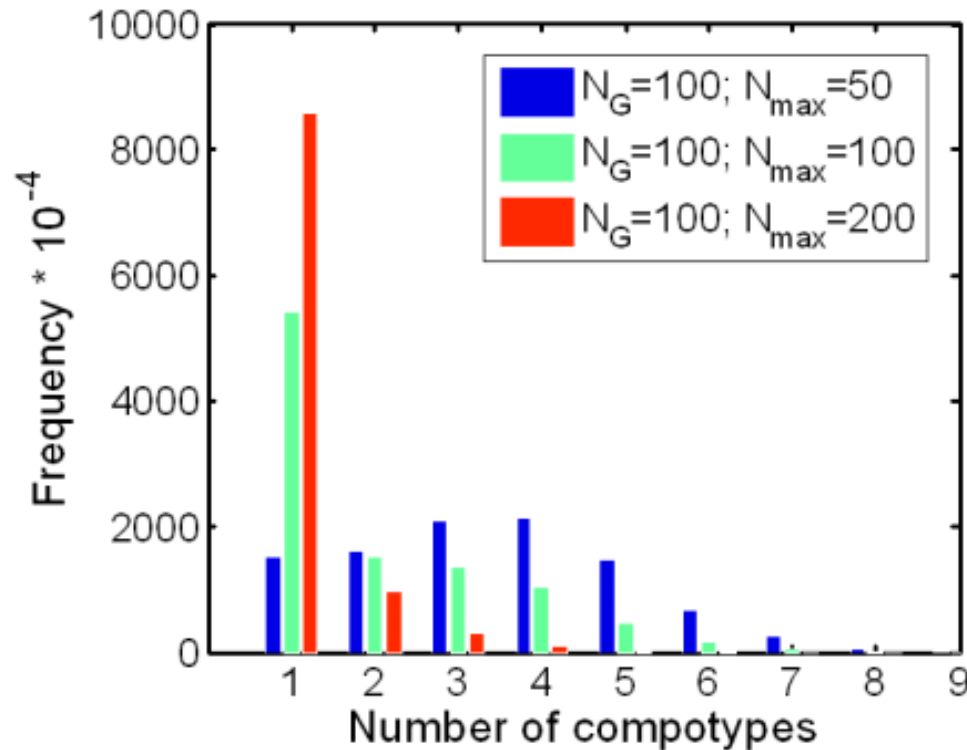
Following a single lineage.



Compotype, a cluster of similar composomes (**quasi species**)

$$H(\chi, \delta) = \frac{\bar{n}^\chi \cdot \bar{n}^\delta}{|\bar{n}^\chi| \cdot |\bar{n}^\delta|}$$

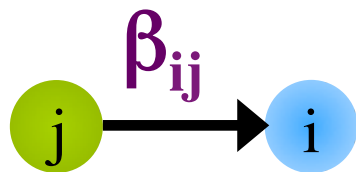
GARD Simulations Show Multiple Compotypes



Different β networks give rise to different dynamics.

Number of compotypes: range 1-7, average ~ 2 .

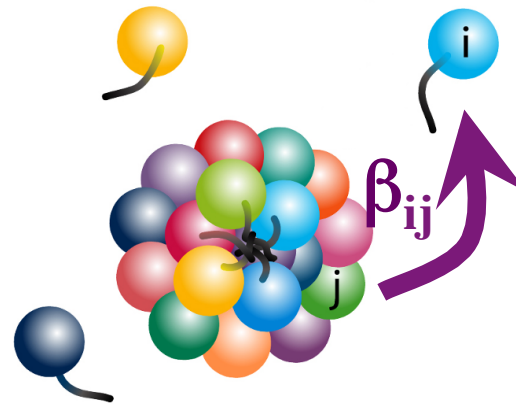
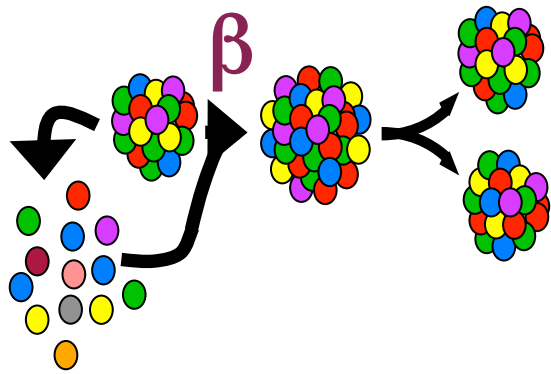
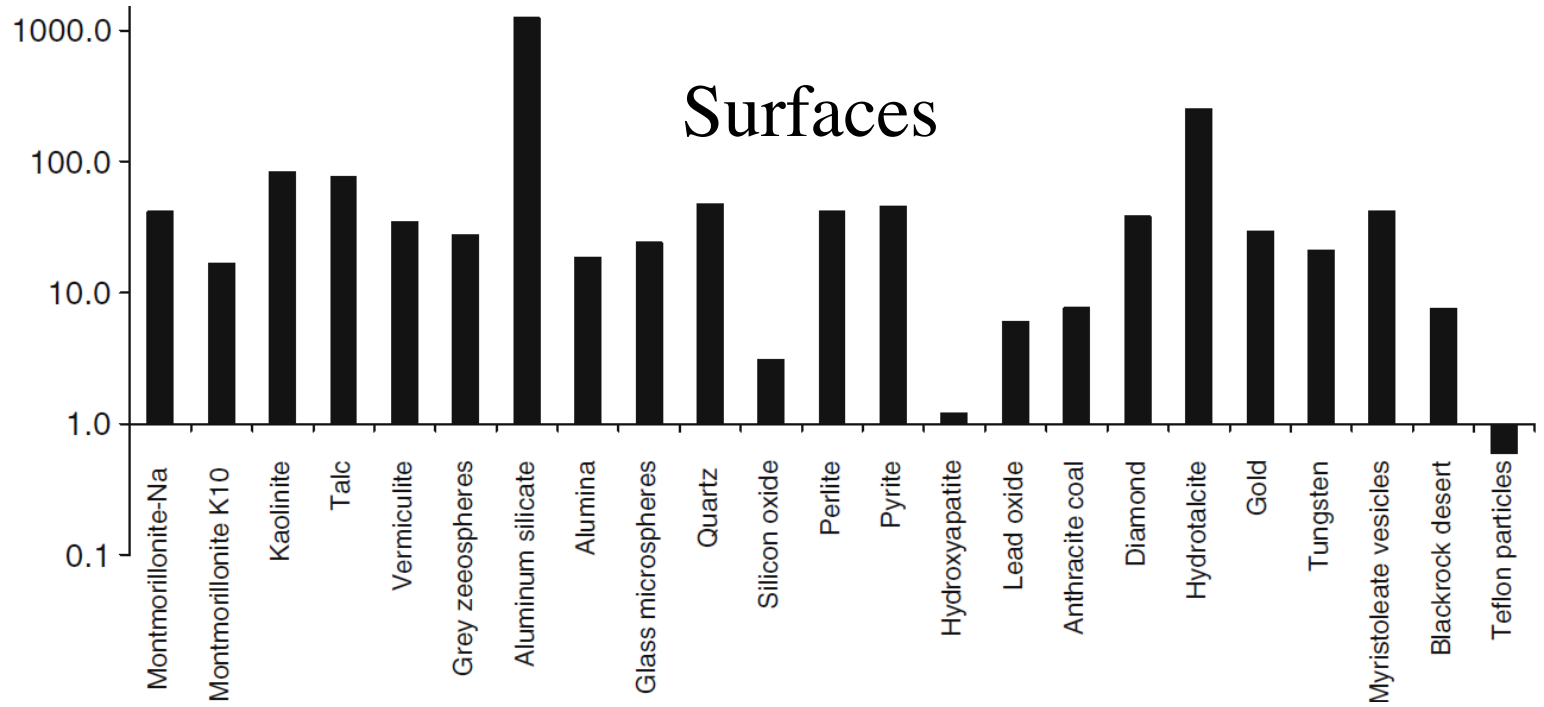
β 's \leftrightarrow environmental chemistries



values are drawn from a lognormal distribution ('graded network')

β 's \leftrightarrow environmental chemistries

Vesicle formation
rate enhancement





sometimes, evolution sucks

Selection in GARD

- Can a network of chemical reactions undergo Darwinian evolution?
- Are metabolism first & lipid world even worth to consider as protocells?

Selection of GARD assemblies towards a target compotype.

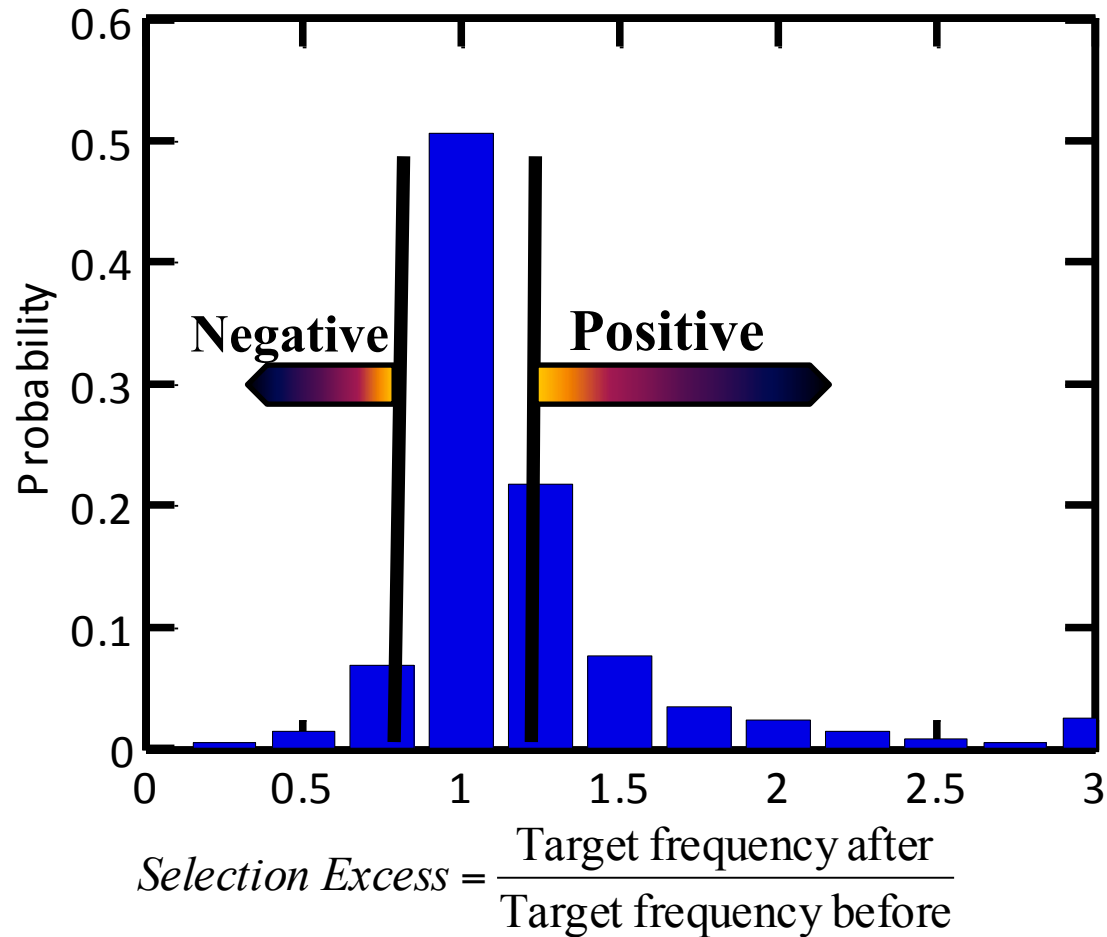
- 1) Identify most frequent compotype (= target).
- 2) Rerun the same simulation while modifying the β_{ij} values at each generation, biasing the growth rate towards the target.

$$\beta'_{ij} = \left\{ \begin{array}{ll} \beta_{ij} & i \text{ or } j \notin \text{Current} \\ 1.1H\beta_{ij} & i \text{ and } j \in \text{Current} \end{array} \right\}$$

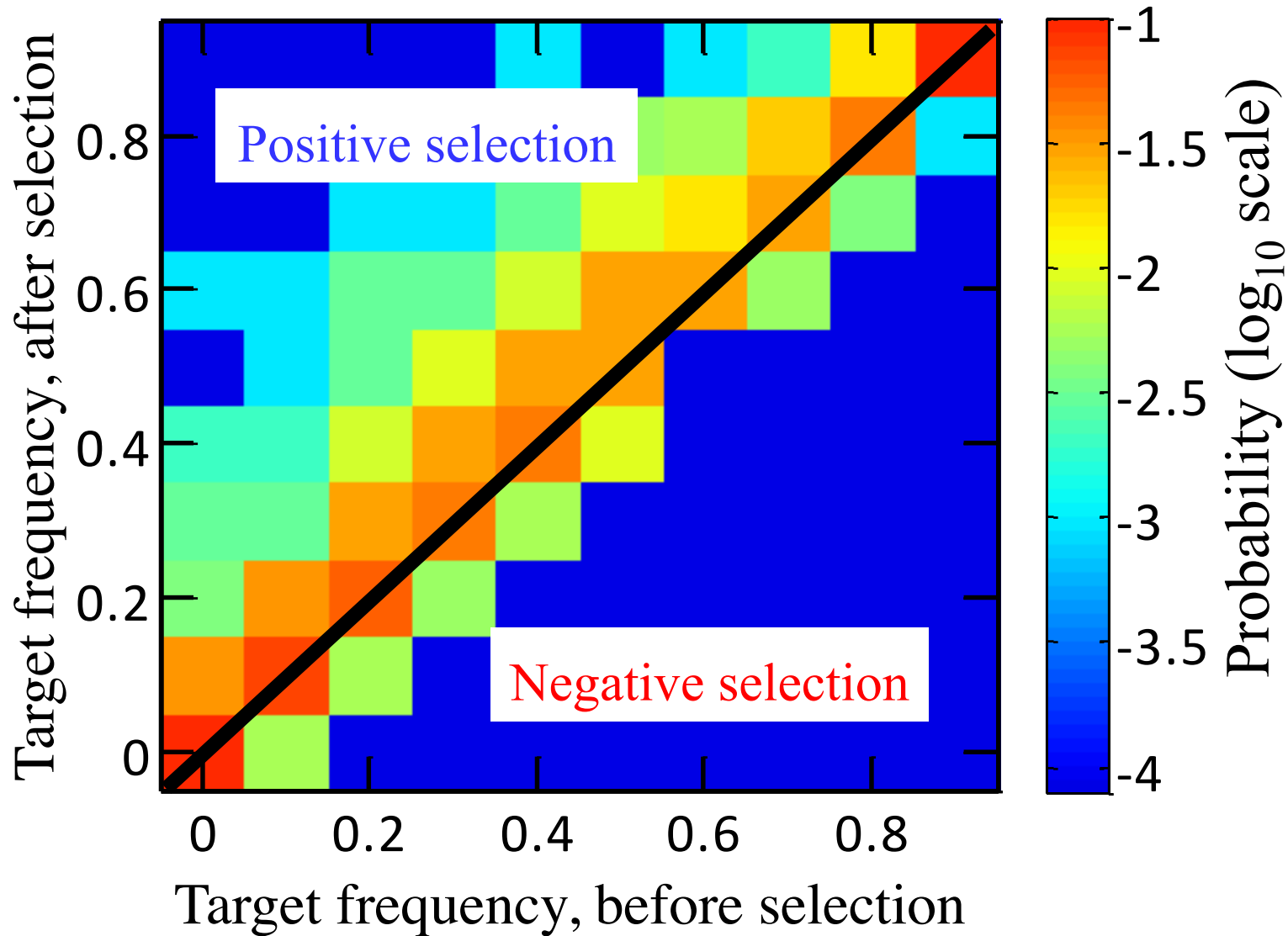
H : compositional similarity between **current** and **target**.

$$\textit{Selection Excess} = \frac{\text{Target frequency after}}{\text{Target frequency before}}$$

Selection in GARD

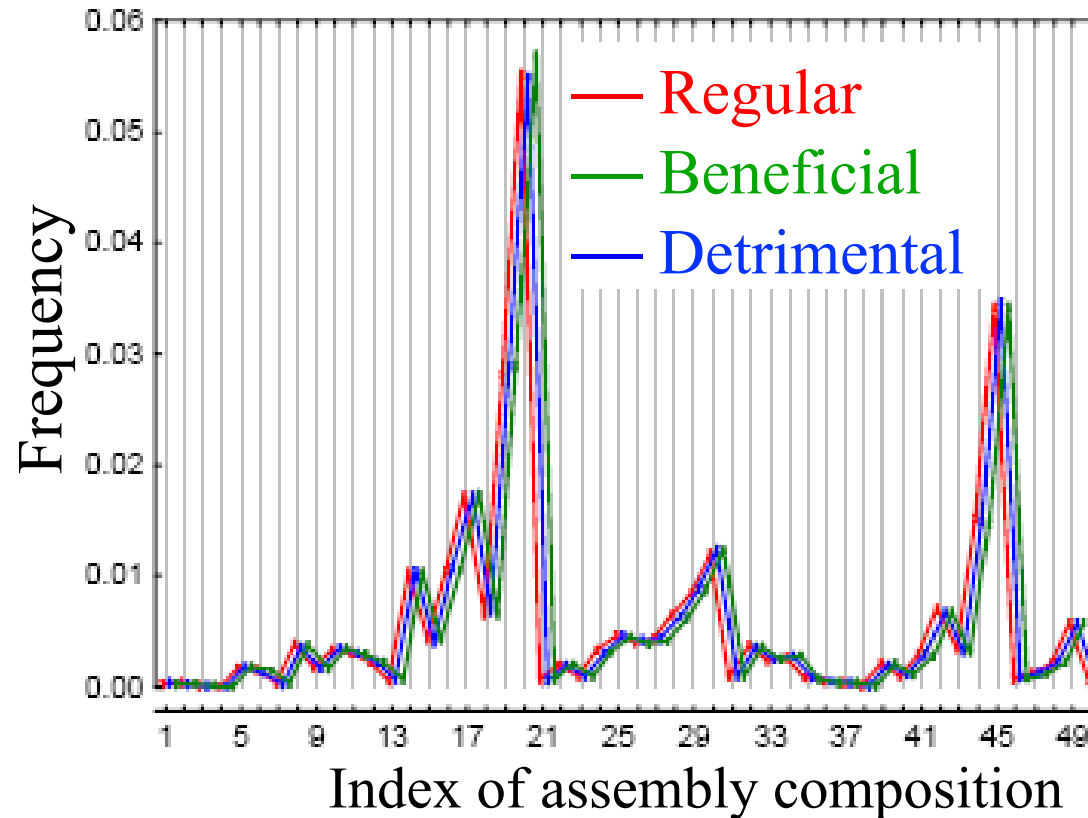


Selection in GARD



Lack of selectivity in GARD? *NO.*

Vasas, Szathmary & Santos, PNAS 107, 1470-1475 (2010): Imposing Darwinian selection in GARD has, at most, negligible effect...

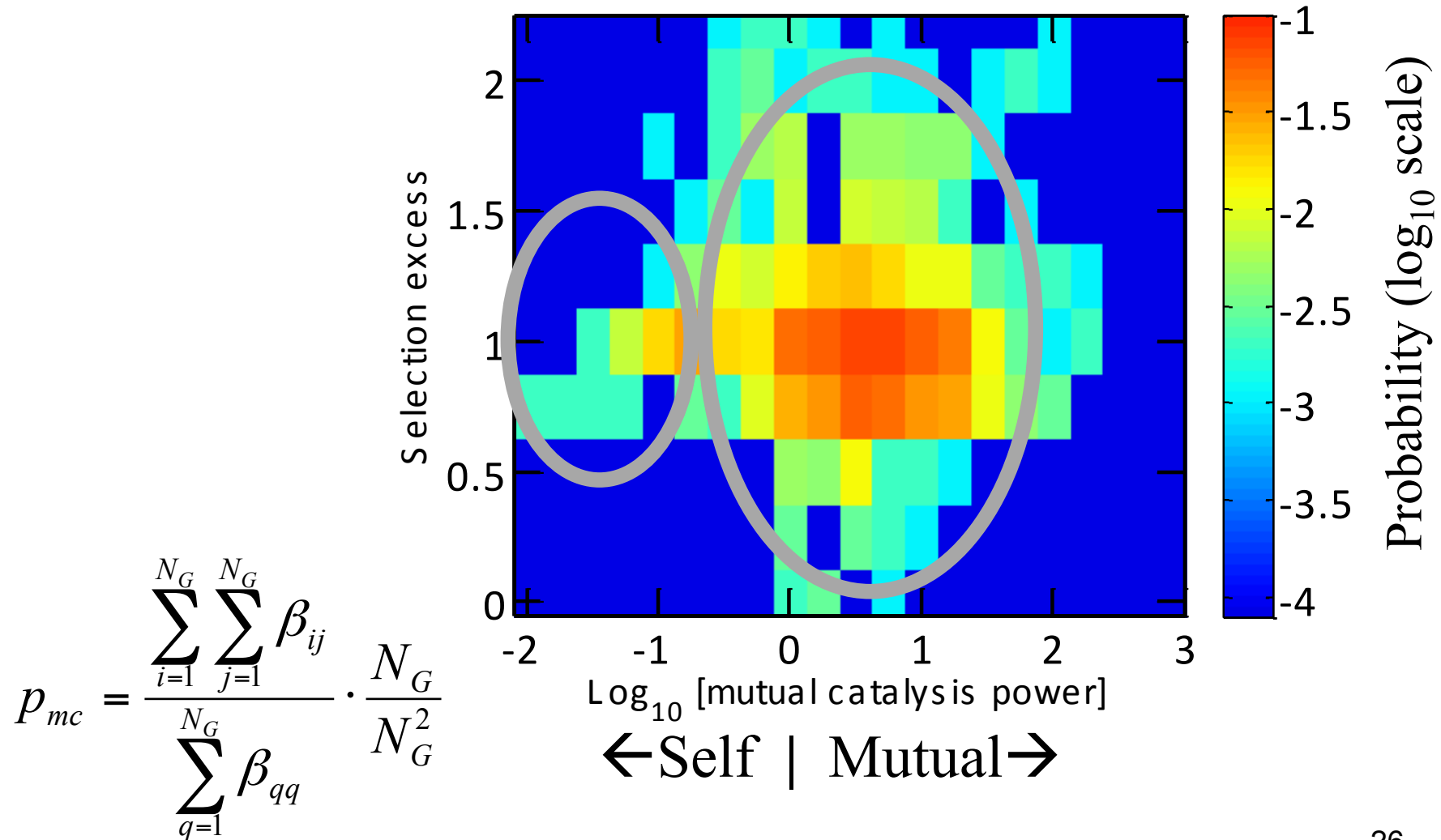


Their weak points:

- (1) Target is not a composome.
- (2) Only a single simulation performed.
- (3) Small repertoire ($N_G=10$) and assembly size ($N_{\max}=6$).
- (4) Arbitrary fitness threshold.

How the β network effects selection ?

→ Too much self-catalysis is not good → ‘Dead-End’.



Self vs. Mutual catalysis

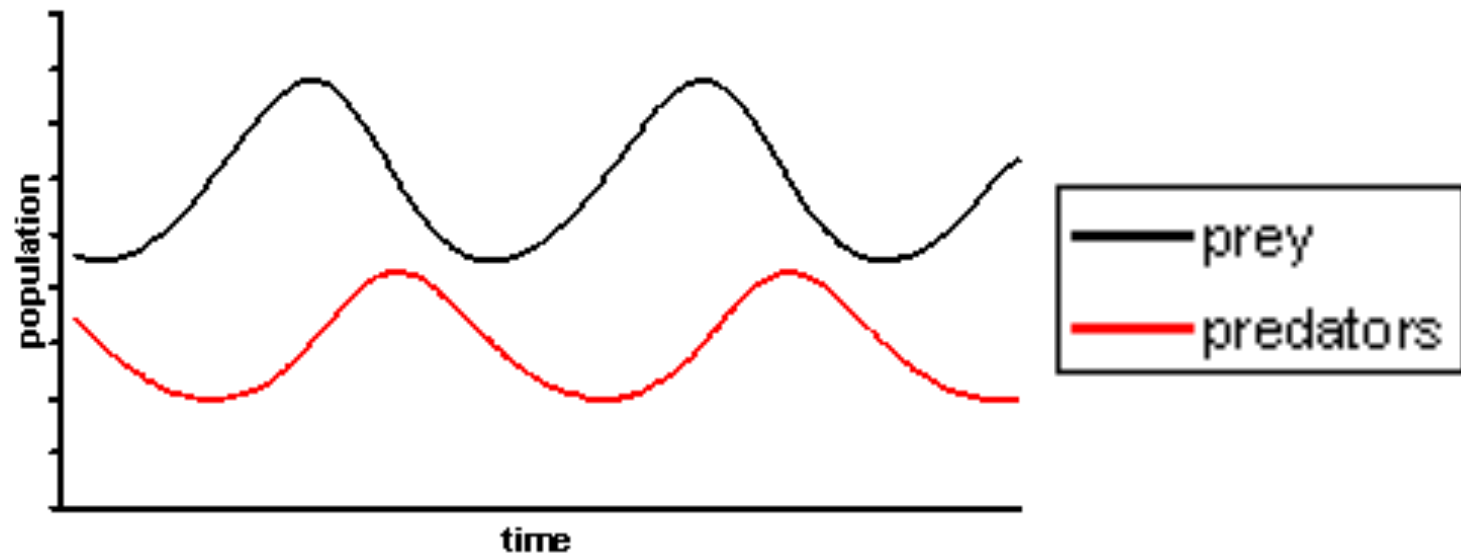
Experimental hints that excess mutual catalysis is required for effective evolvability:

- In an abstract chemistry model, if self-replication is inhibited then self-maintaining organizations arise [Fontana & Buss, PNAS (1994); see also Szathmary, Proceedings: Biological Sciences (1995)].
- RNA fragments (of *Azoarcus group I ribozyme*) that are mutually interacting outcompete selfish yet efficient individual fragments [Vaidya & Lehman Nature (2011)].
- A particular ribozyme (R3C) is capable of only 2 slow doublings, yet a conversion into two cross-replicating ribozymes allows for many fast doublings [Lincoln & Joyce, Science (2009)].
- A mutualistic network of replicating peptides is adaptable to physiochemical conditions (pH, salt) [Dadon et al, Angew. Chem. Int. Ed. (2008)].
- Mutualism is also needed for effective contagion [Ugander et al, Proc. Natl. Acad. Sci (2012)].

Ecology



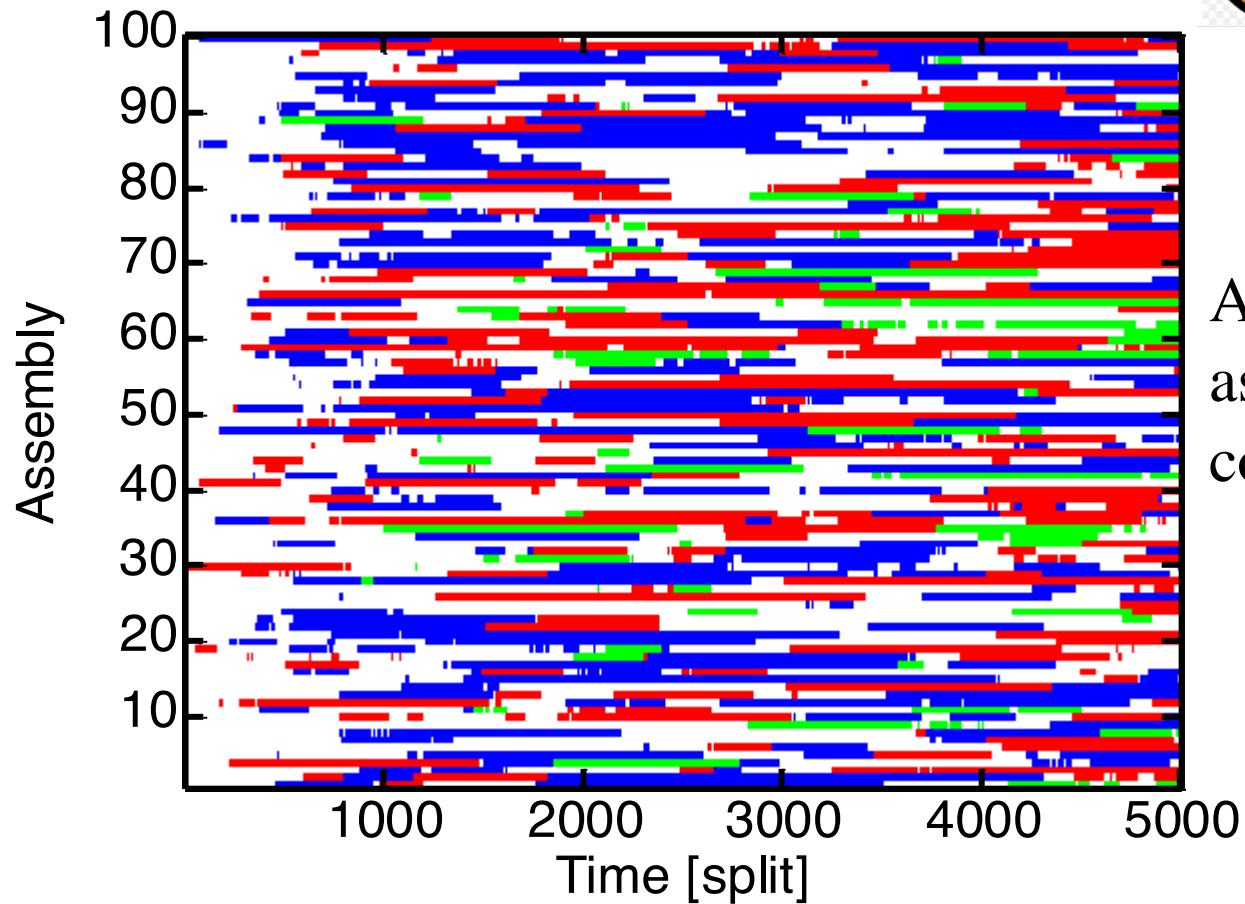
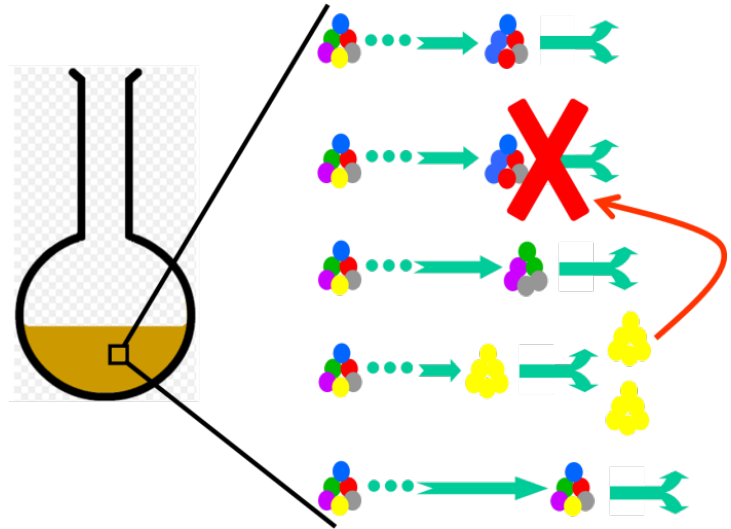
- ❖ Predator–Prey
- ❖ r–K relations



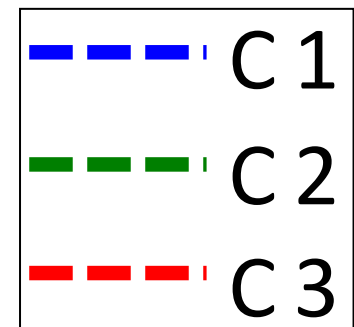
- ❖ Lotka & Volterra
- ❖ MacArthur
- ❖ Malthus & Verhulst

Population Dynamics in GARD

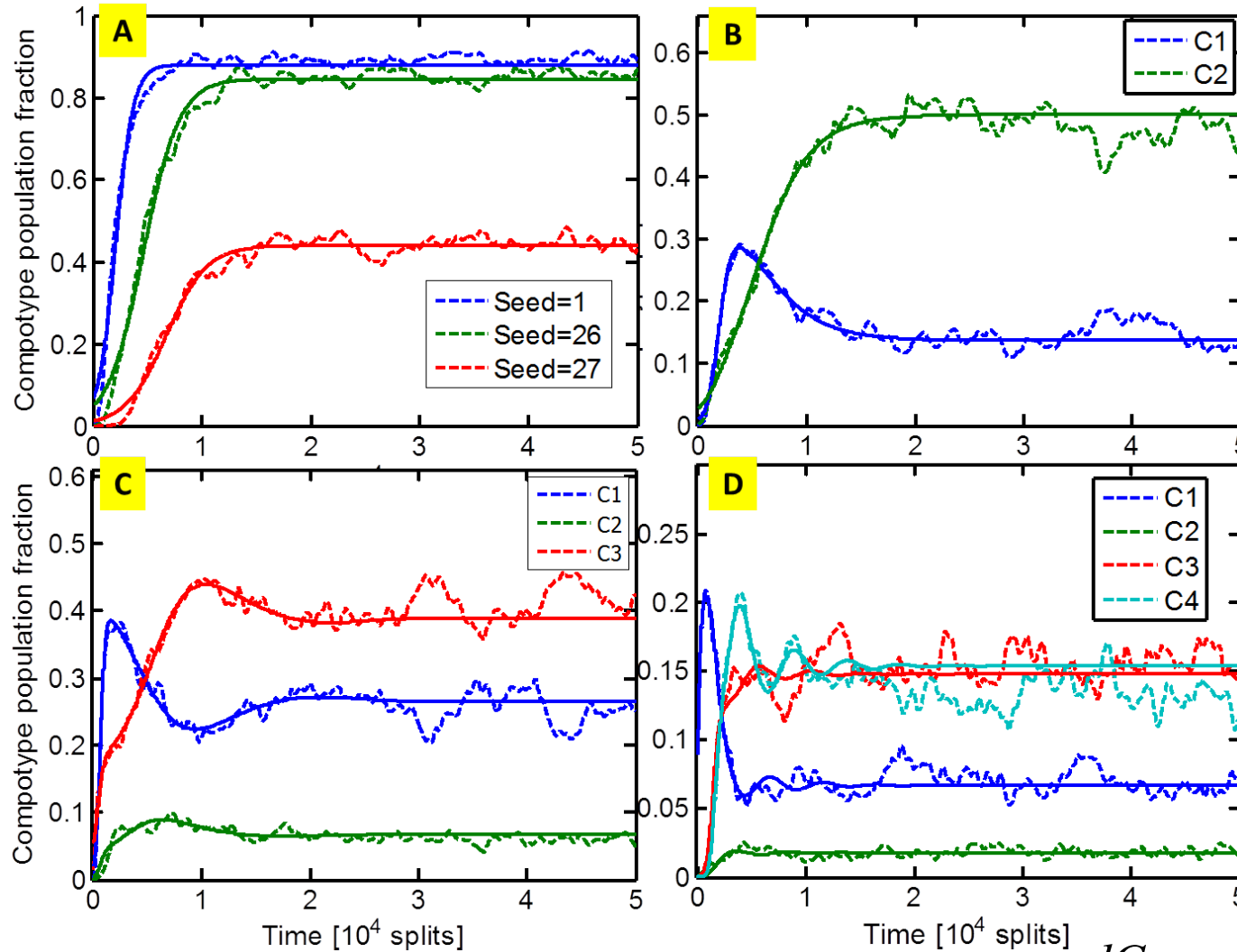
Fixed population size.



At each time point, each assembly is colored by its compotype.



Population Dynamics in GARD



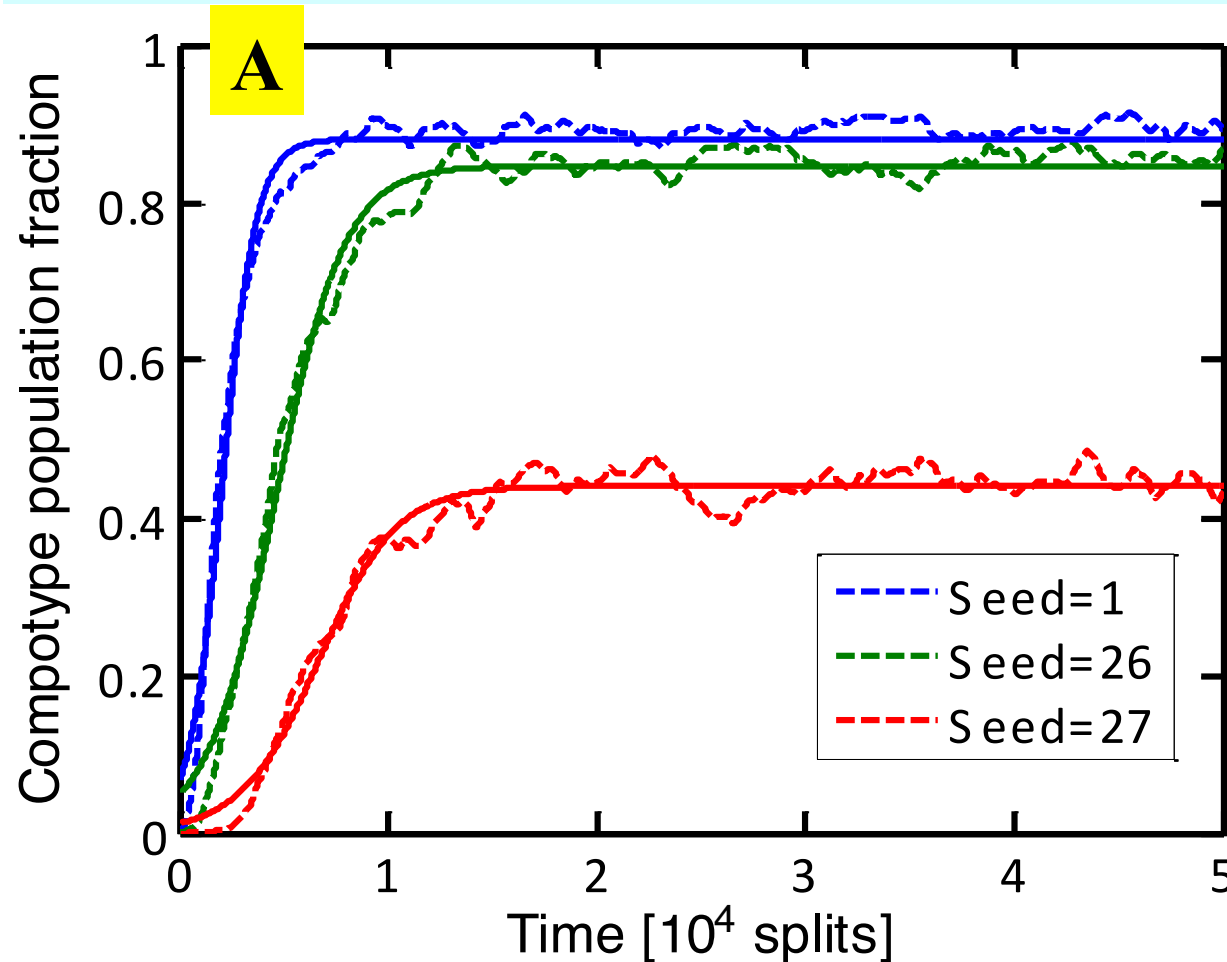
Species = comptypes

Logistic growth:

$$\frac{dC_i}{dt} = r_i C_i \left(1 - \frac{C_i + \sum_{j=1, j \neq i}^{N_c} \alpha_{ij} C_j}{K_i} \right)$$

30

Population Dynamics in GARD



3 examples with single compotype (different β).

Why plateau doesn't reach 1.0 ?

Species = comptypes

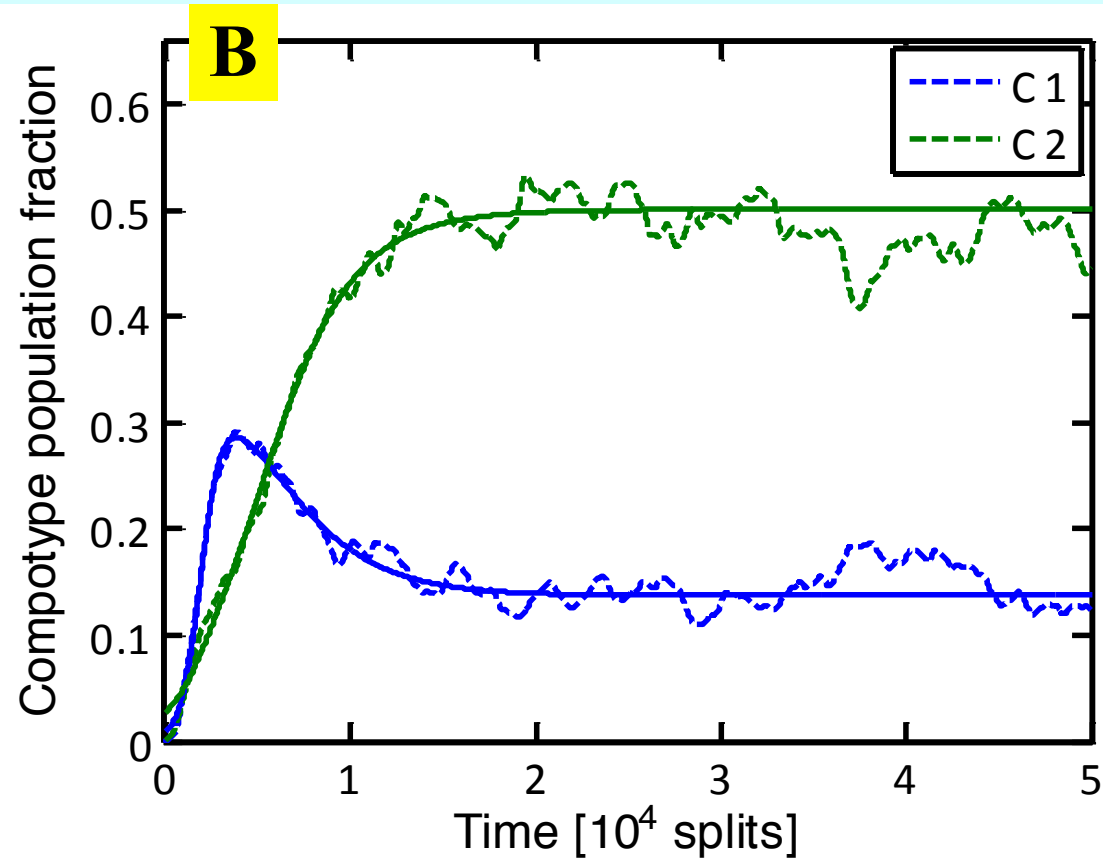
Logistic growth:

$$\frac{dC_i}{dt} = r_i C_i \left(1 - \frac{C_i + \sum_{j=1, j \neq i}^{N_c} \alpha_{ij} C_j}{K_i} \right)$$

31

Population Dynamics in GARD

“**Takeover**” of a fast-rising compotype by a slower one.



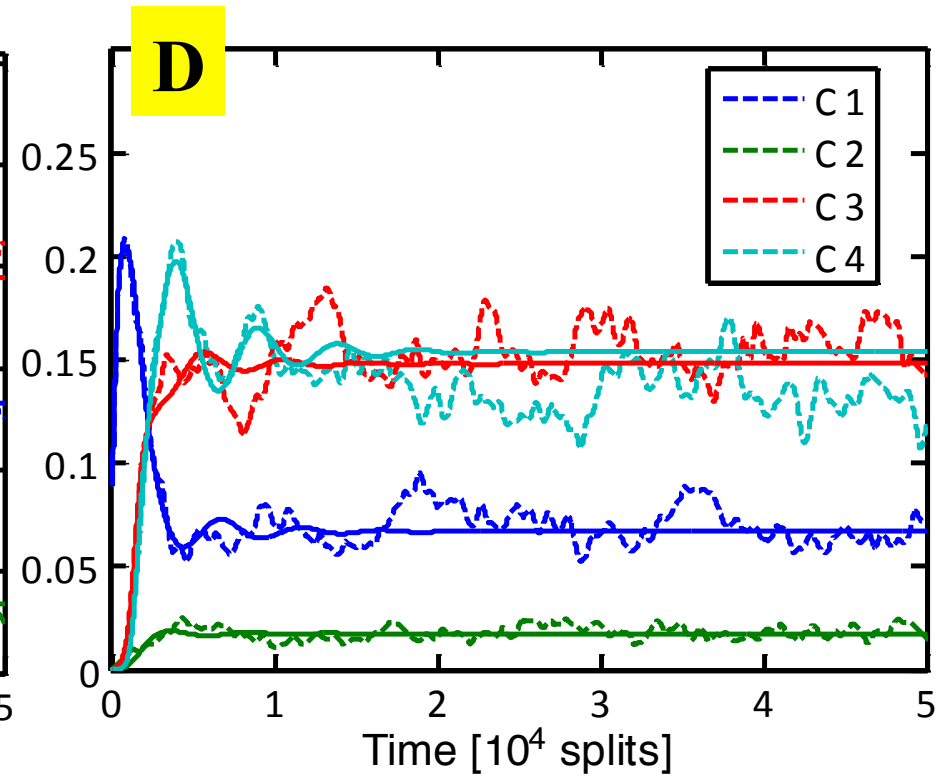
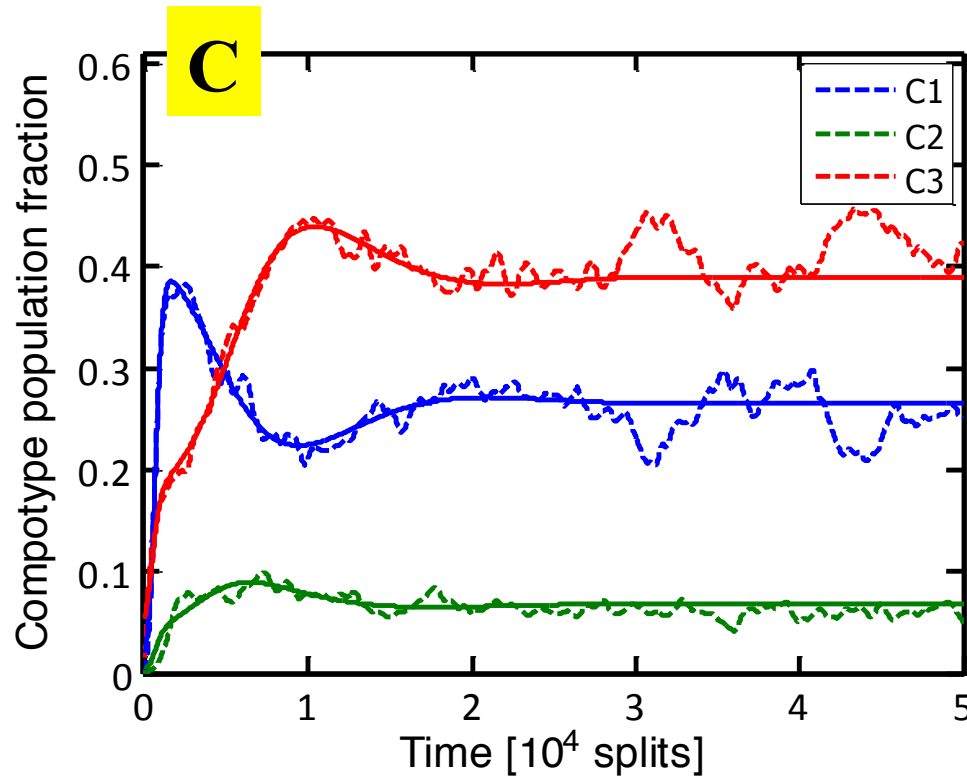
Species = compotypes

Logistic growth:

$$\frac{dC_i}{dt} = r_i C_i \left(1 - \frac{C_i + \sum_{j=1, j \neq i}^{N_c} \alpha_{ij} C_j}{K_i} \right)$$

32

Population Dynamics in GARD



Intricate food-web (α_{ij} values).

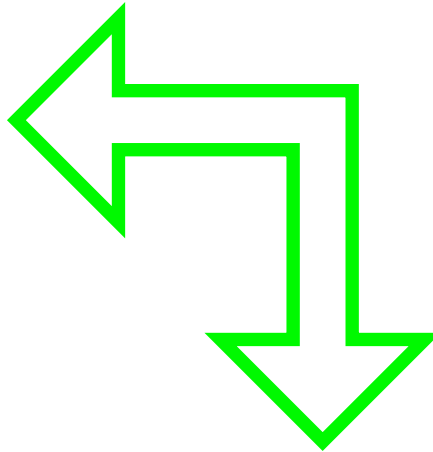
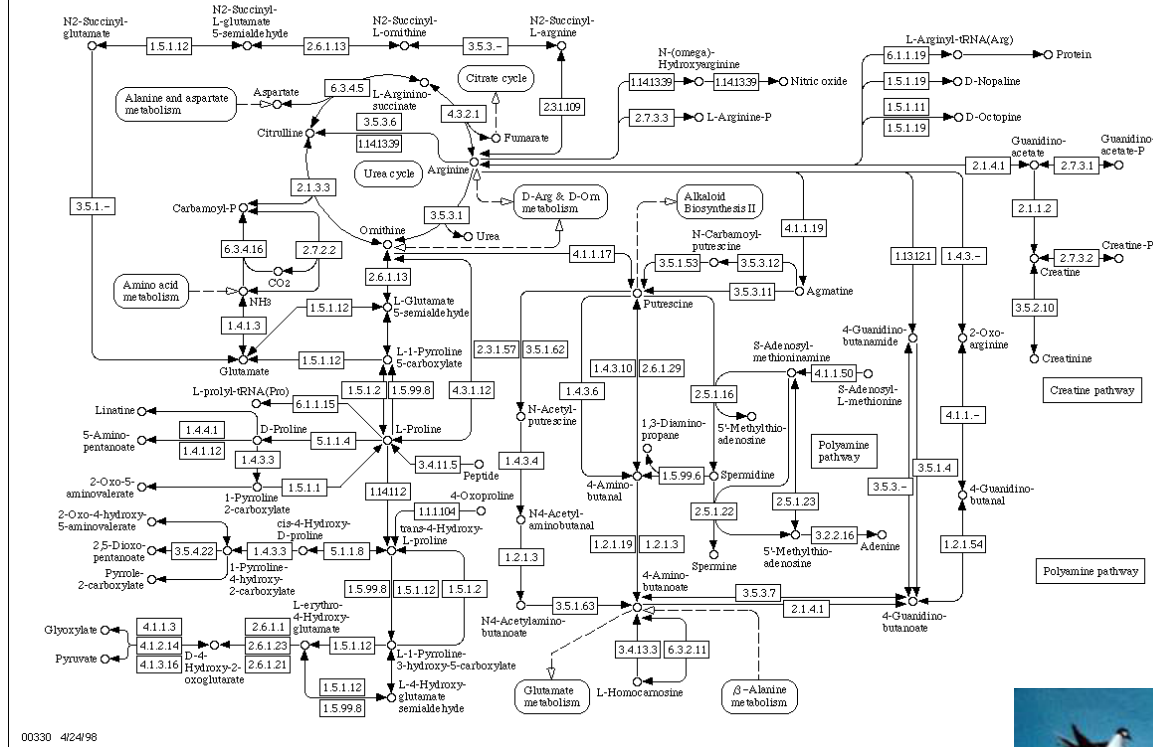
Species = compotypes

Logistic growth:

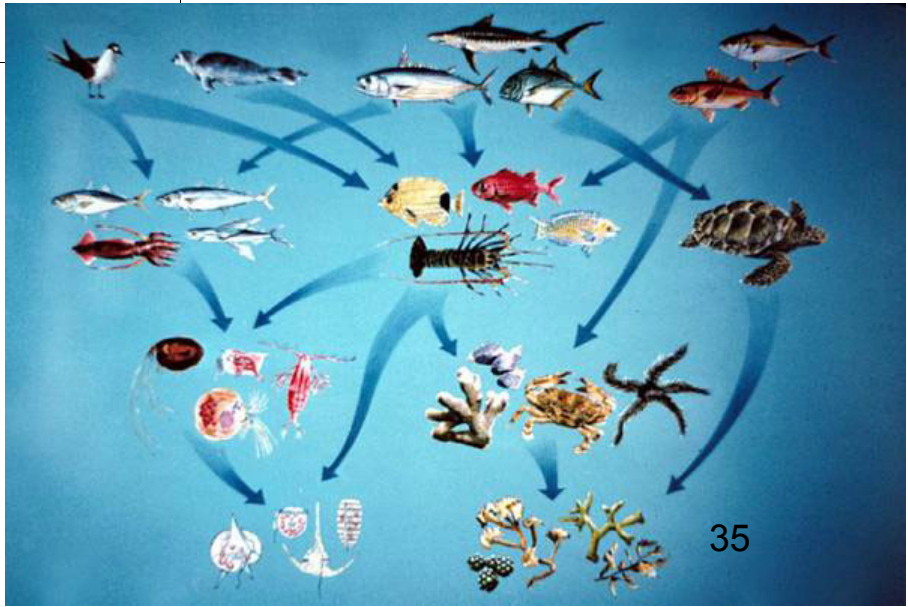
$$\frac{dC_i}{dt} = r_i C_i \left(1 - \frac{C_i + \sum_{j=1, j \neq i}^{N_c} \alpha_{ij} C_j}{K_i} \right)$$

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Metabolism



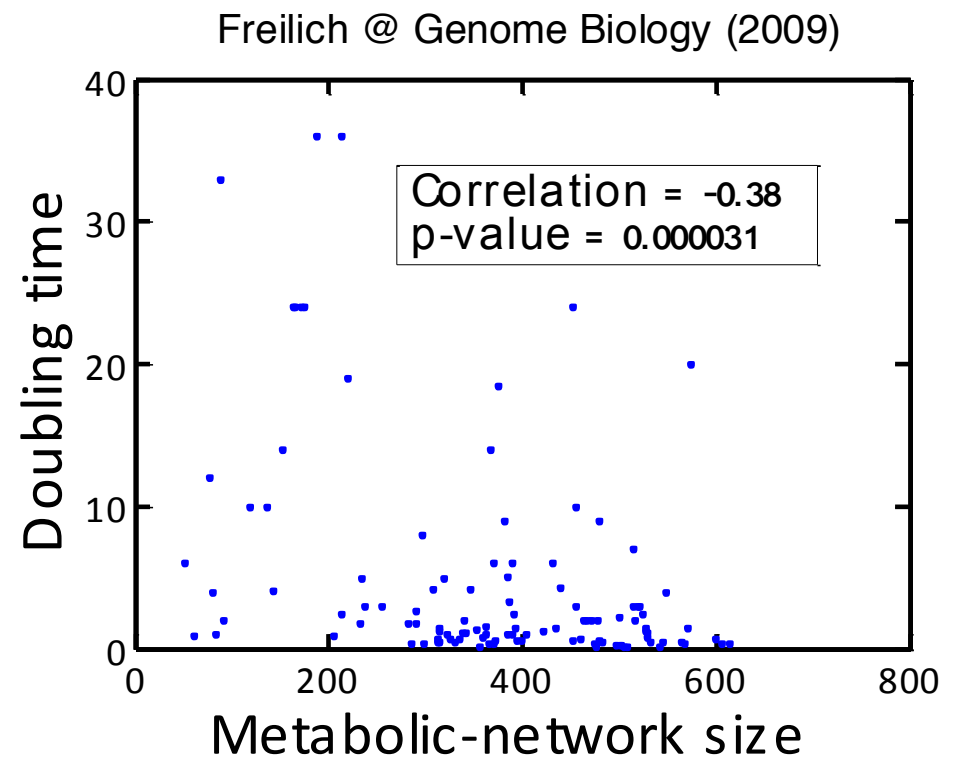
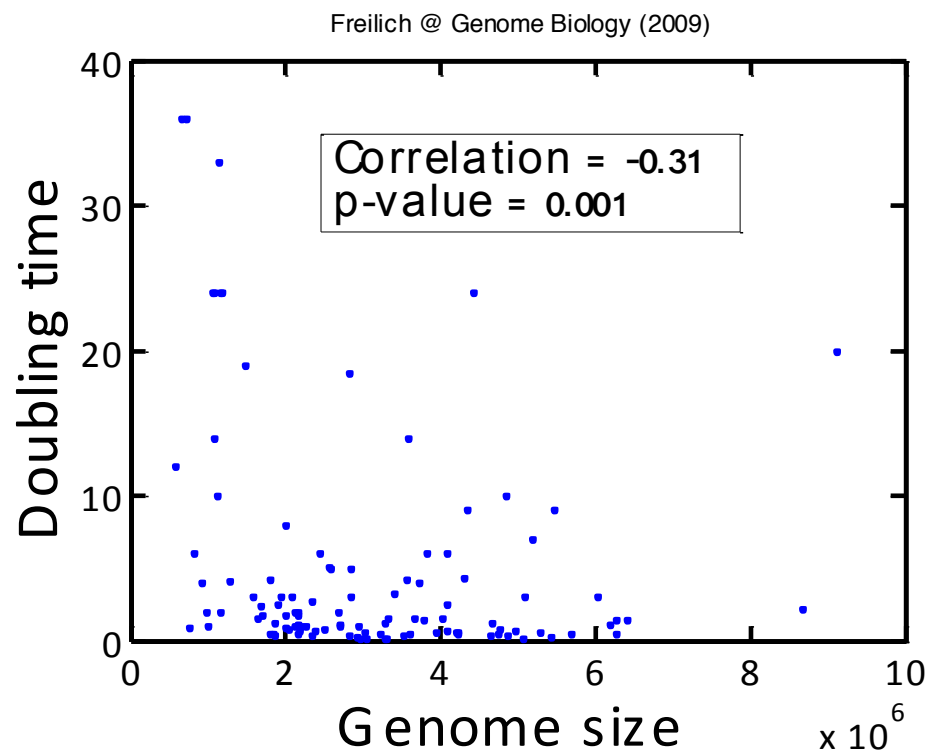
Prebiotic Ecology From molecules to Ecosystem



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GARD's Ecology

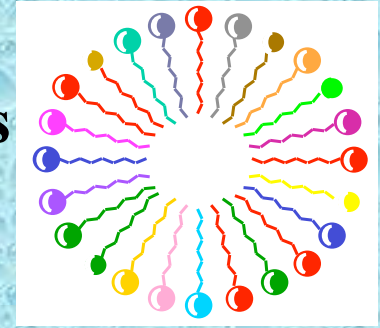
Based on experimental data of 111 bacteria.



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Lipid-world & GARD model: compositional assemblies

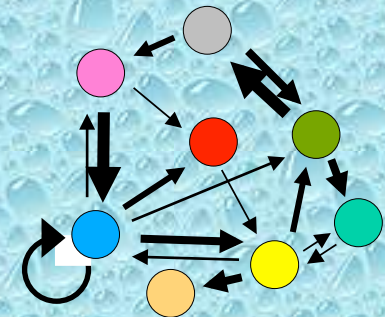
➤ NOT JUST CONTAINER !



Compotypes (clusters of faithfully replicating compositions)

➤ Darwinian selection

Mutual catalysis is required for effective evolvability



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Acknowledgments:

Doron Lancet.

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Natalio Krasnogor (U. Nottingham, UK).

Lancet group.



Omer Markovitch

Funding:

*** Minerva Center for Life Under Extreme Planetary Conditions, at Weizmann Institute.**

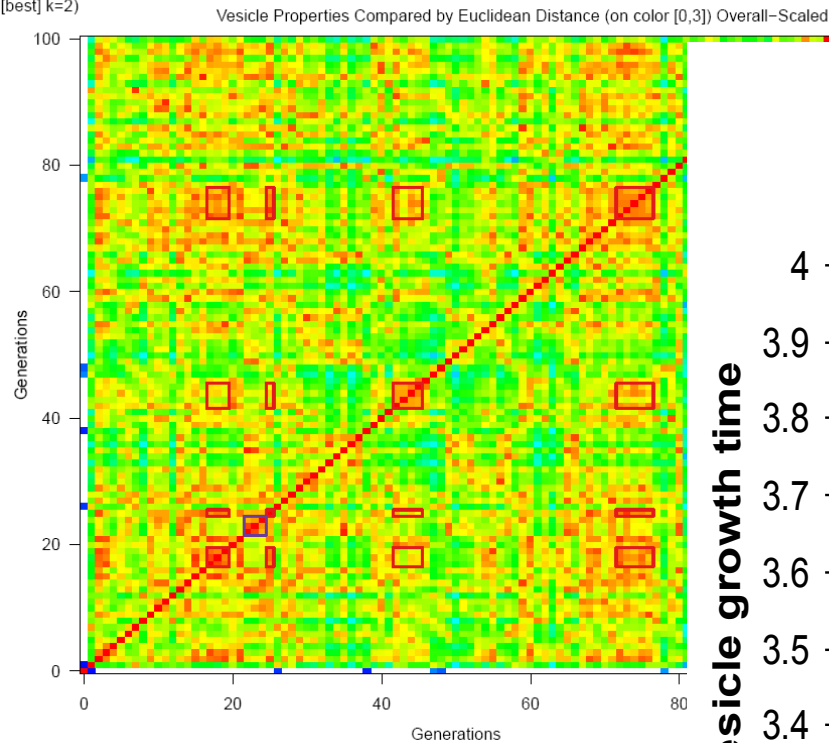
*** E.U. FP7 “MATCHIT”**

Real GARD – Raphael Zidovetzki, U. California Riverside

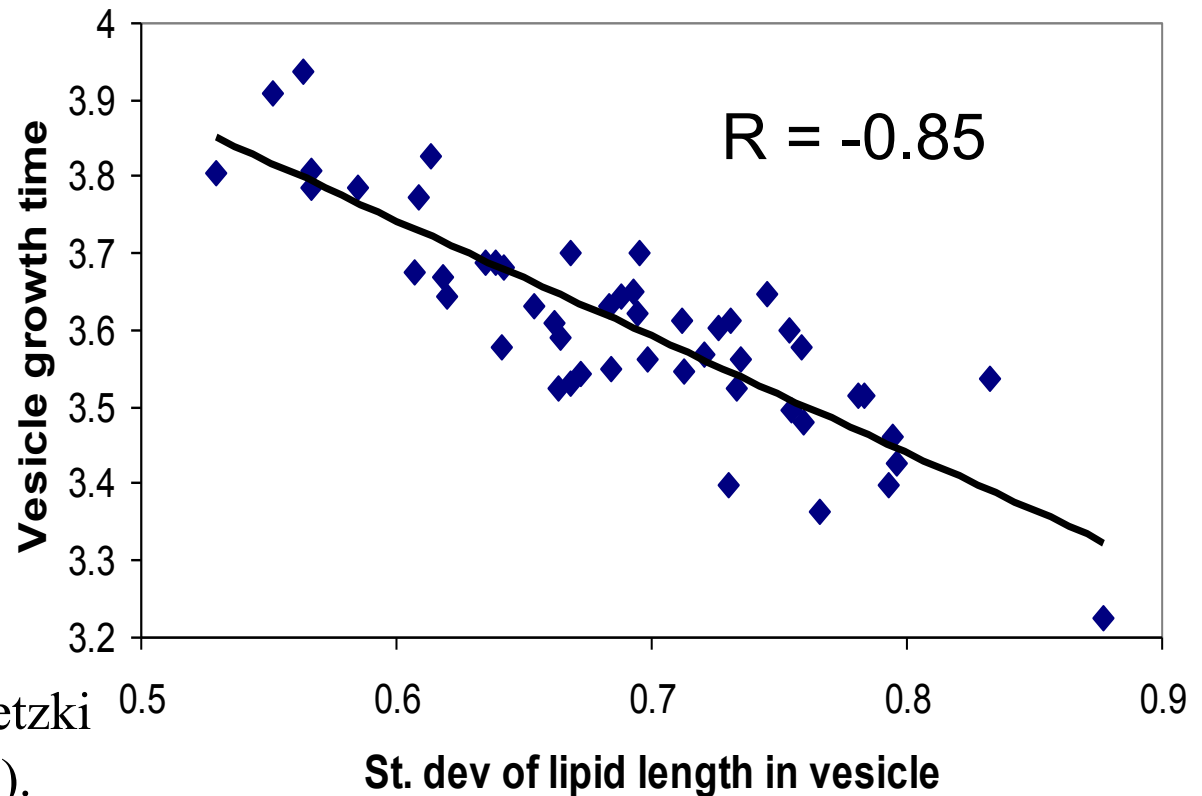
Real lipids: phosphate-idyl-(serine / amine / choline), sphingo-myelin and cholesterol.

Actual physical properties (charge, length, unsaturation).

No run description [ool_paper_runs/ool_runs_2010_08_25_paper_step_7_r635/run_03_comp_20_size_100] (kmeans proptypes [best] k=2)

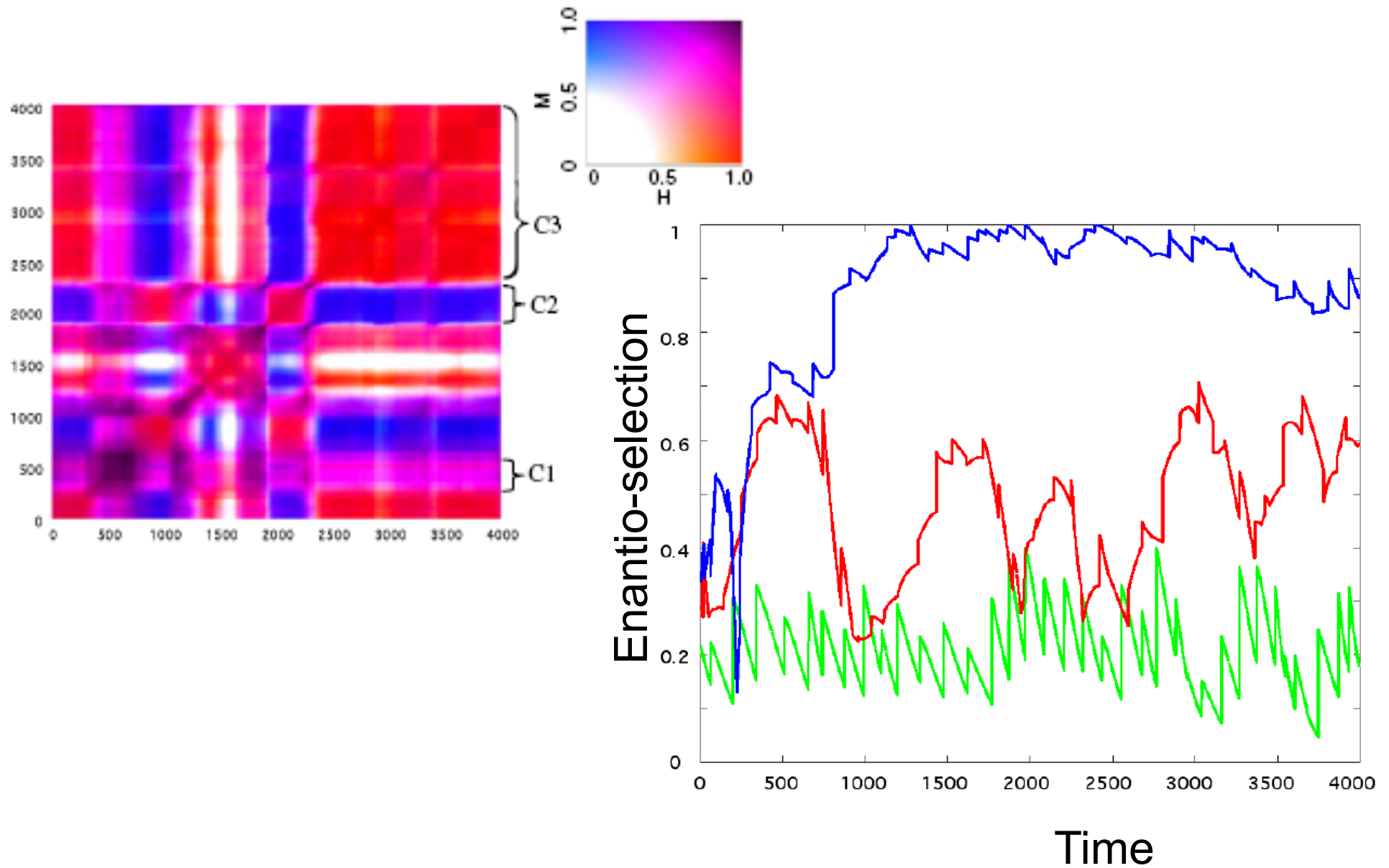


Variability of lipid lengths in vesicle is highly correlated to vesicle replication time



Armstrong, Markovitch, Zidovetzki and Lancet, Phys. Biol. 8 (2011).

Chiral GARD – origins of biochirality



Same behavior in other evolutionary parameters

Comptype diversity

Evolvability Score:

$$ES = \tau(1 - H_0)$$

$1/\tau \sim$ compositional mutation rate.

Compositional diversity

