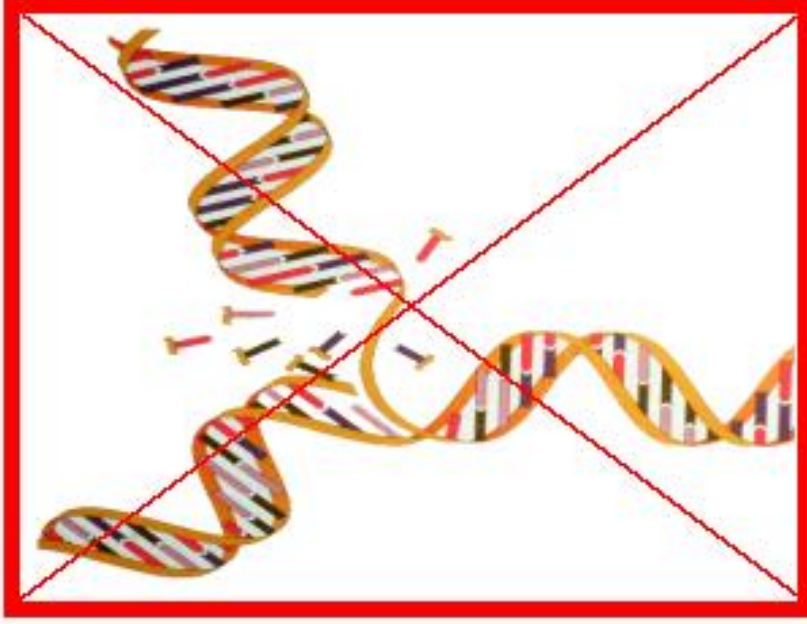
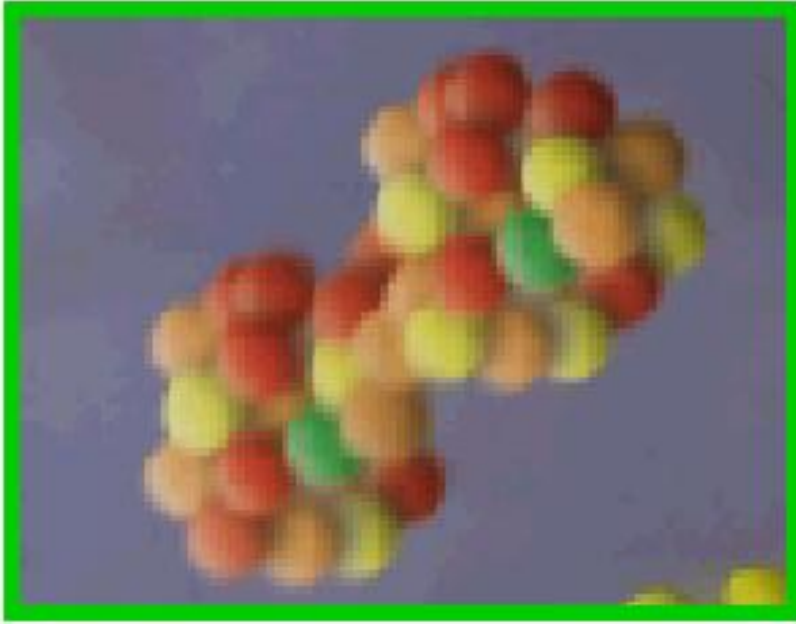




Origin of life without biopolymers

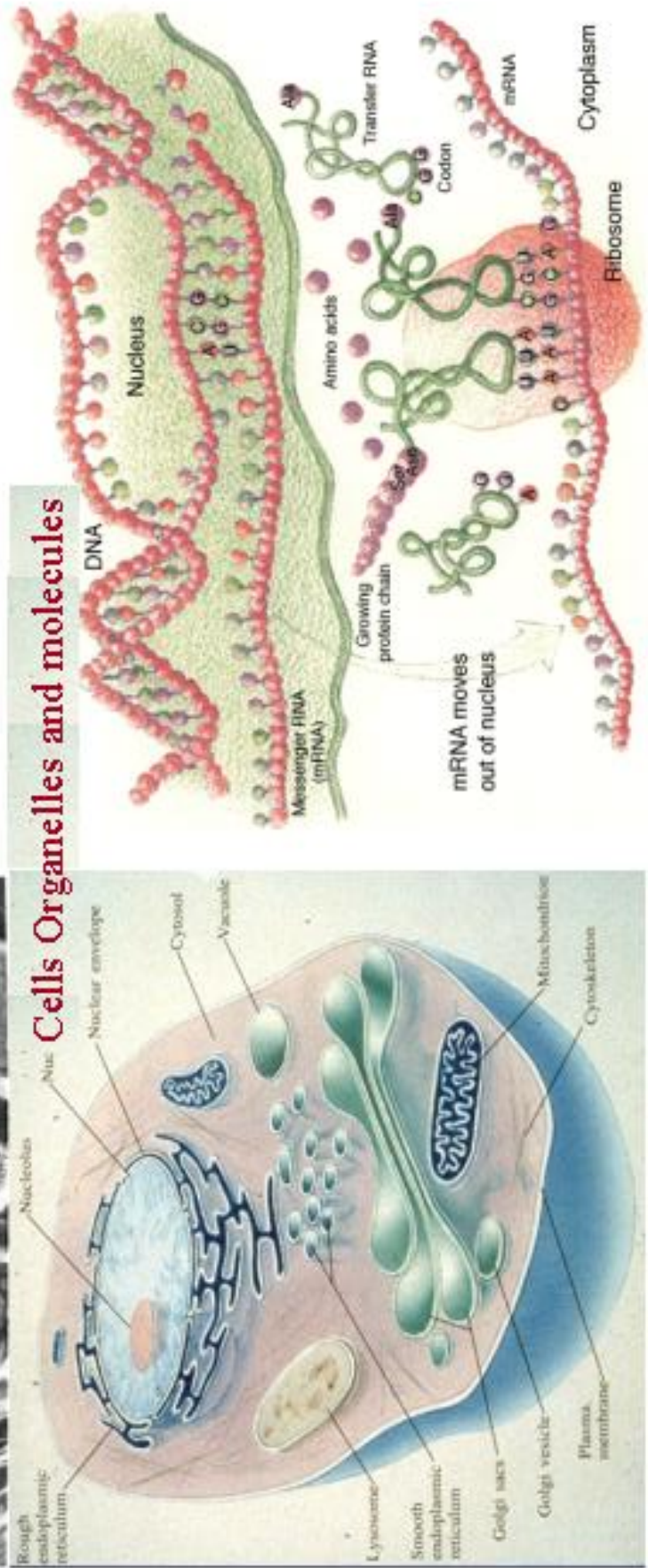


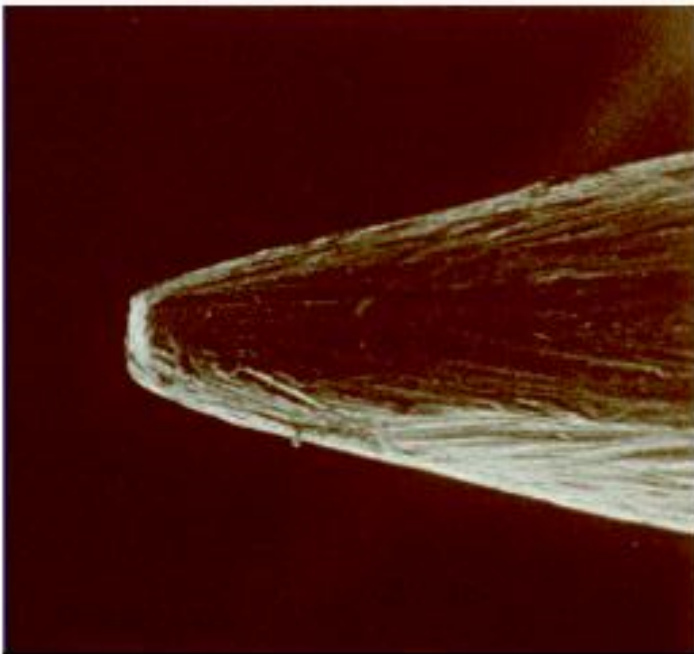
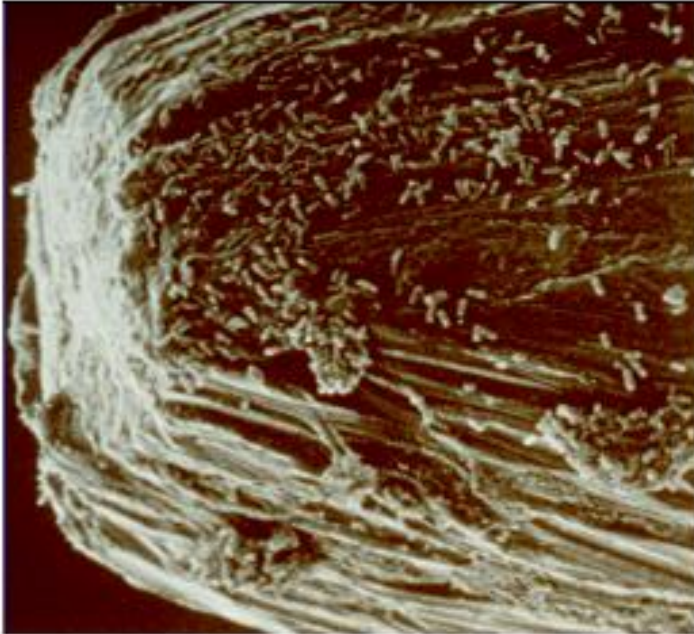
Doron Lancet, Crown Human Genome Center,
the Weizmann Institute of Science, Rehovot, Israel

[Http://ool.weizmann.ac.il](http://ool.weizmann.ac.il)



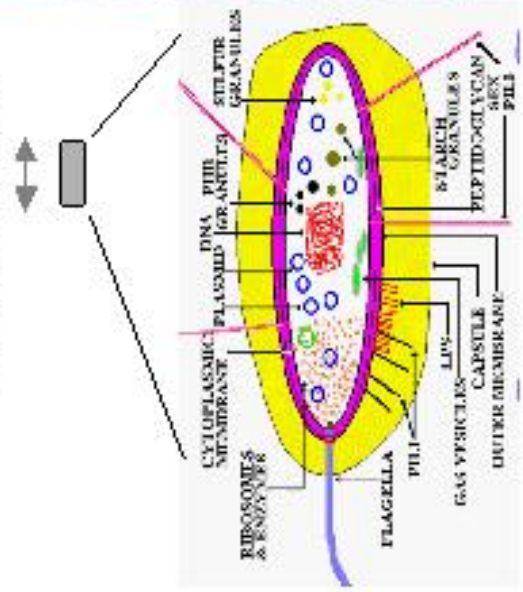
Life is complex!
How did it emerge?



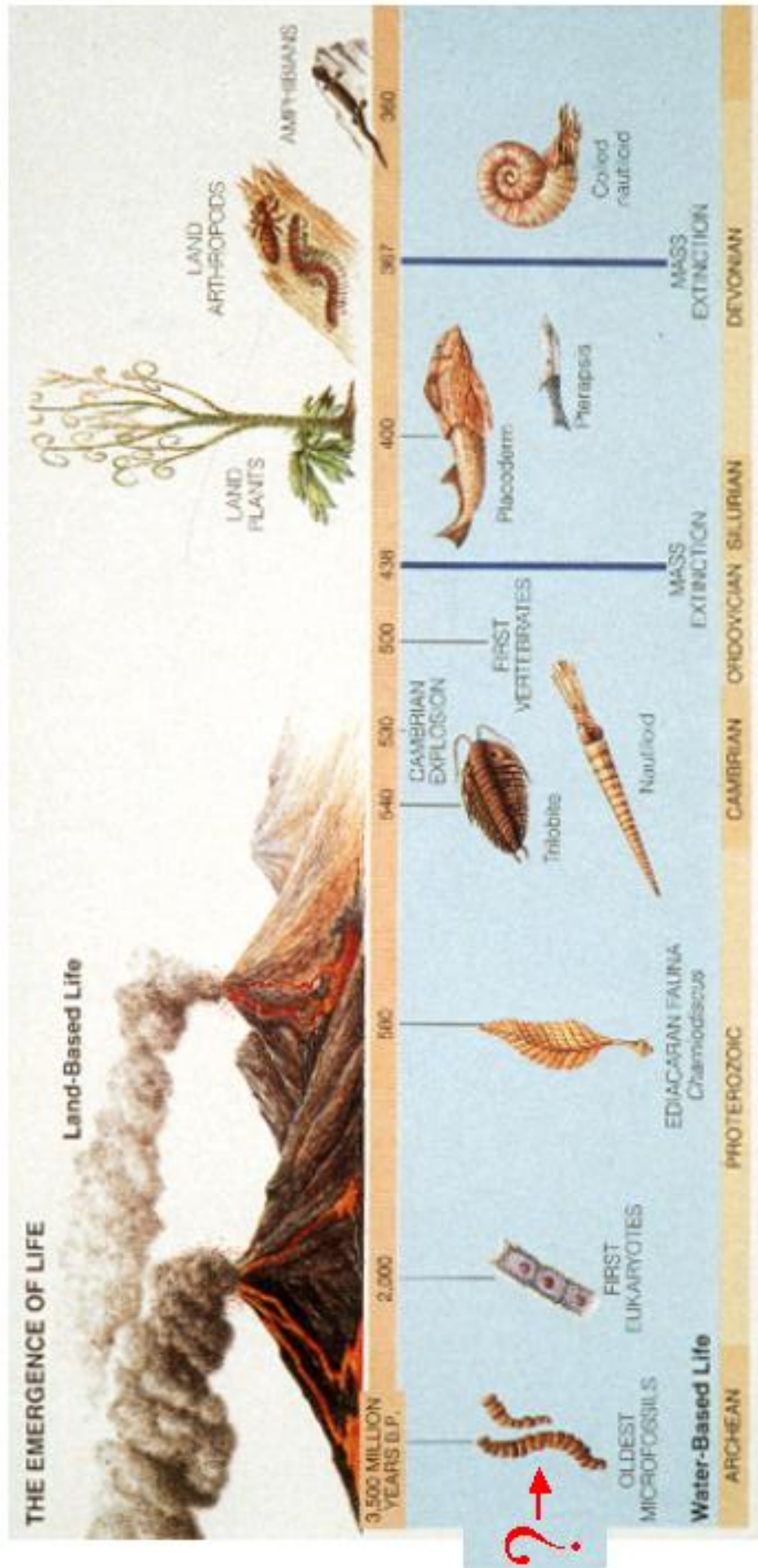


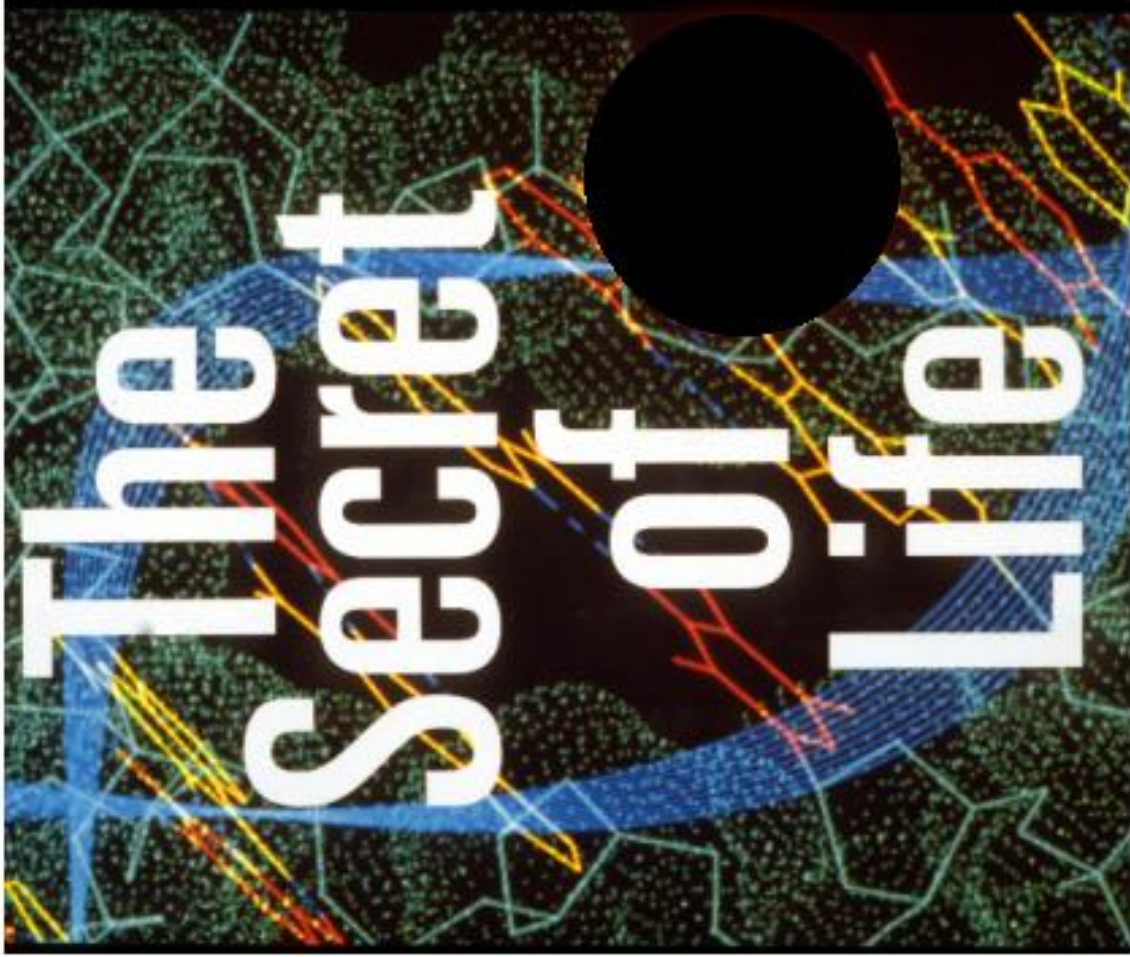
Life:
a haystack
at the point
of a needle

~3 micrometers



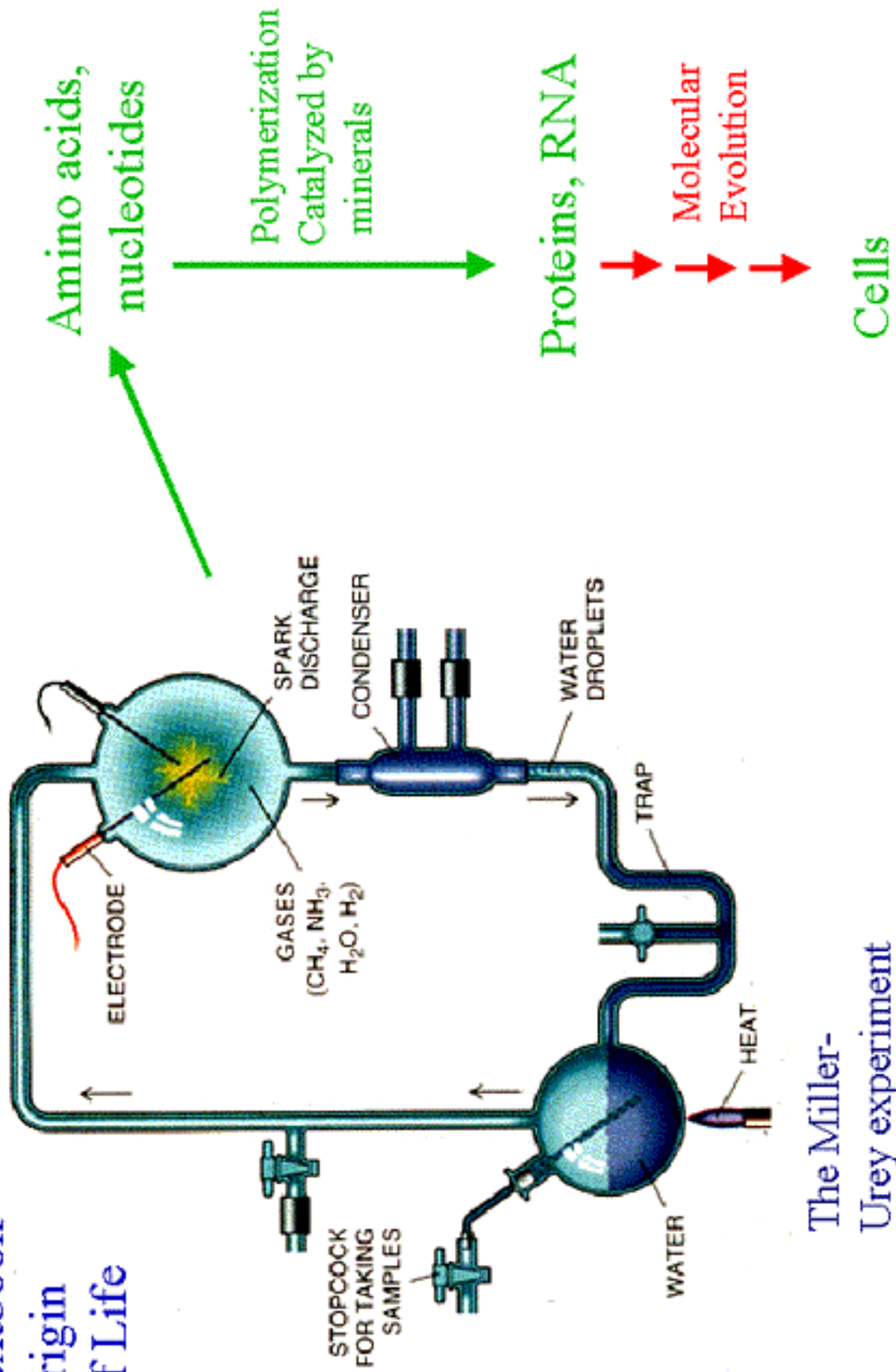
How to reconstruct the events preceding the fossil record?





Finding out
about the
Origin of Life
may lead to a
much better
understanding
of present-day
life!

Textbook
Origin
Of Life



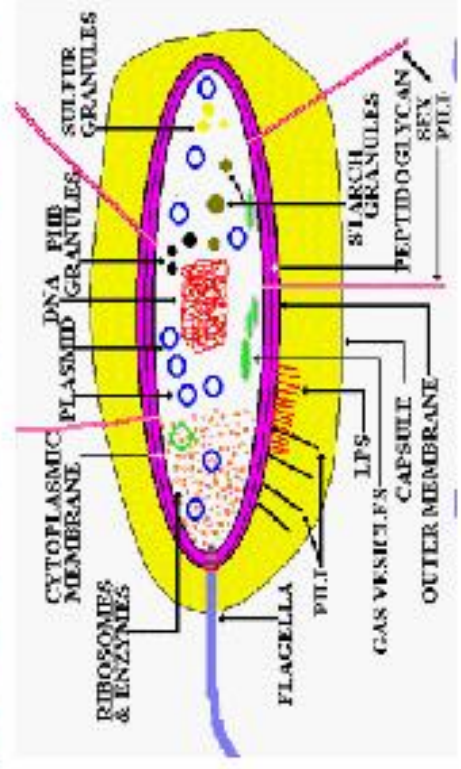
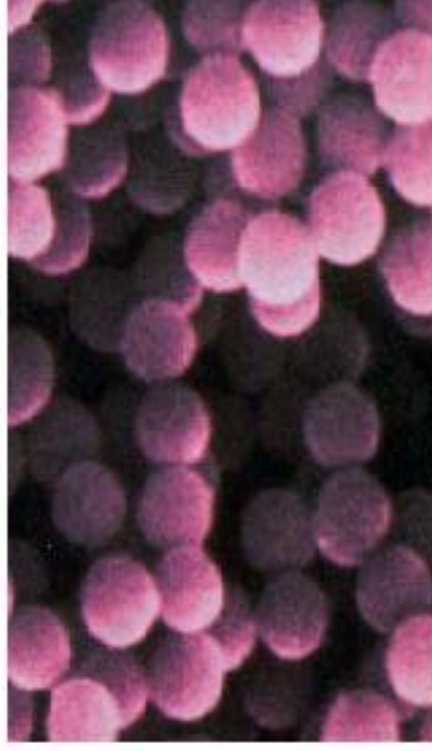
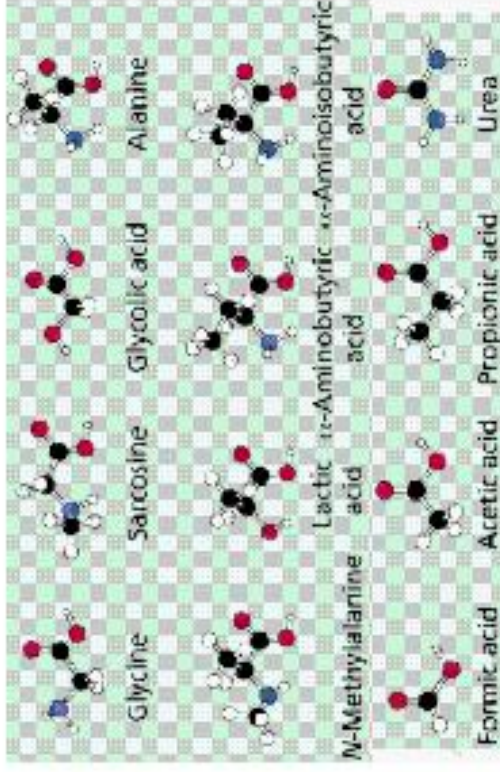
Three proposed stage in the Origin of Life

1) Origin of organic compounds
Understood in principle, need more details

2) Origin of protocells (Morowitz, 1992)
Possible to model by general
physicochemical reasoning

3) Origin of the Last Universal
Common Ancestor (LUCA)

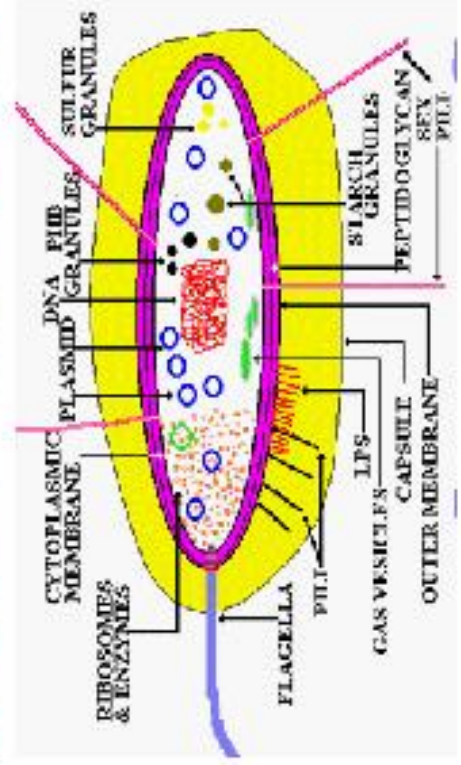
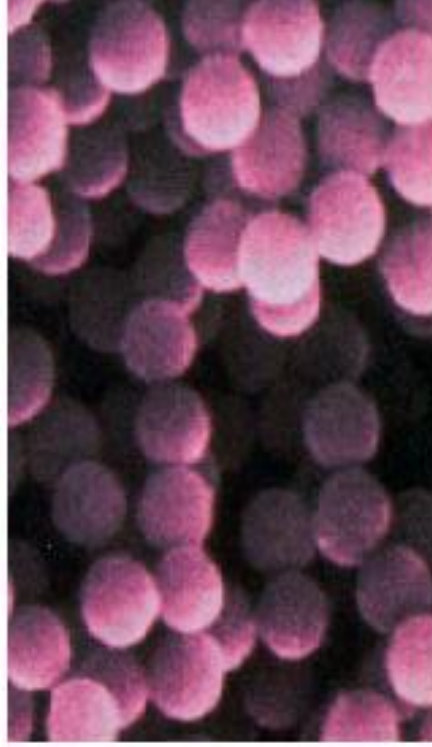
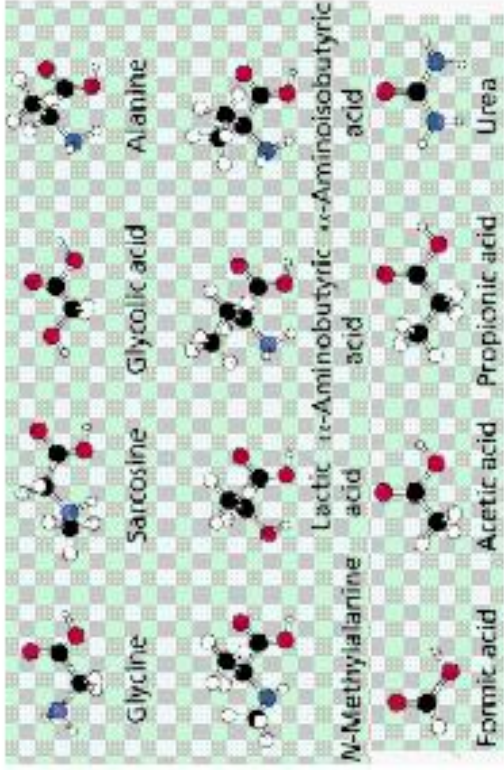
Much more difficult, requires detailed
historical molecular-evolution scenarios



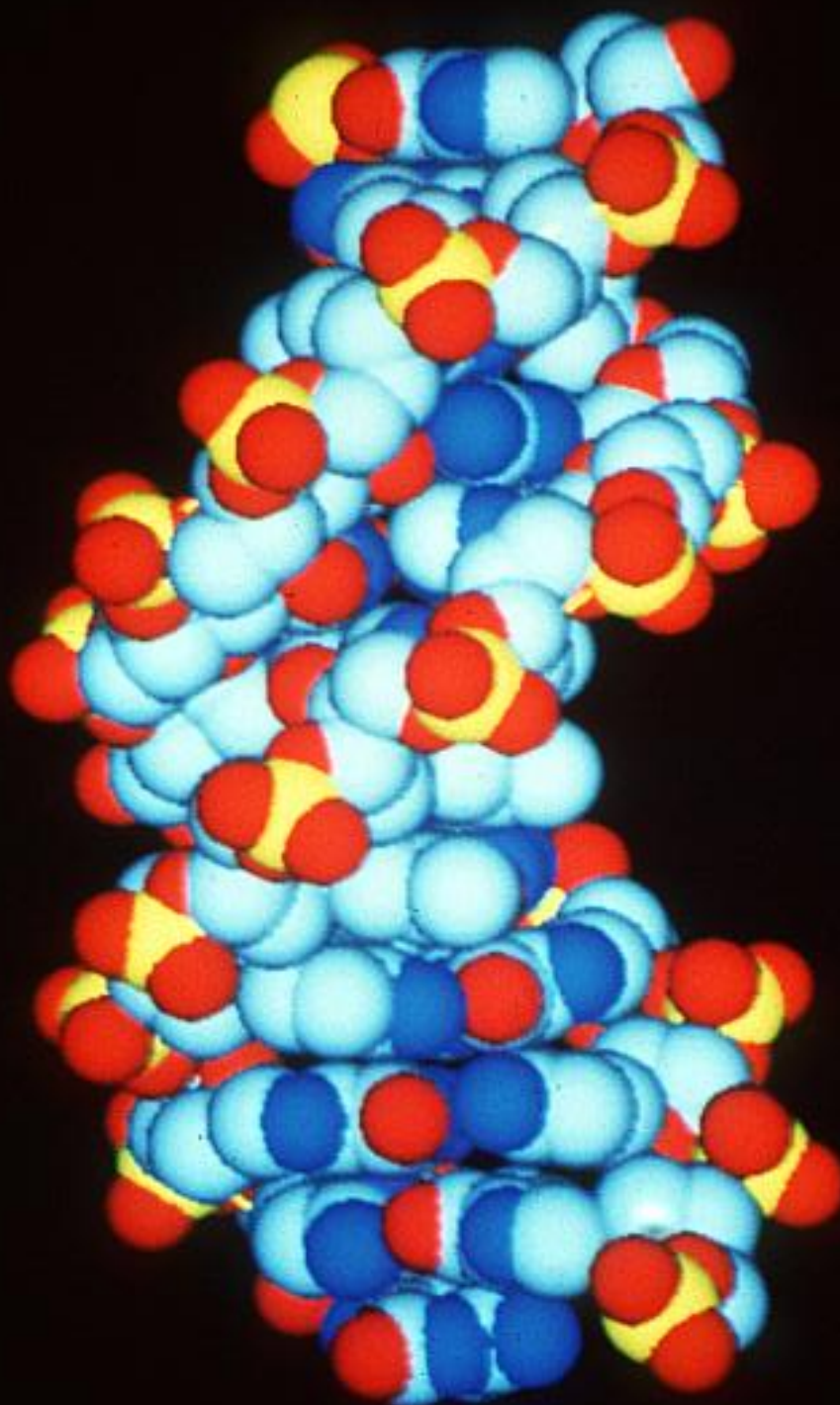
Organic compounds: form naturally
by many different processes: UV,
sparks, volcanoes, comets, meteorites

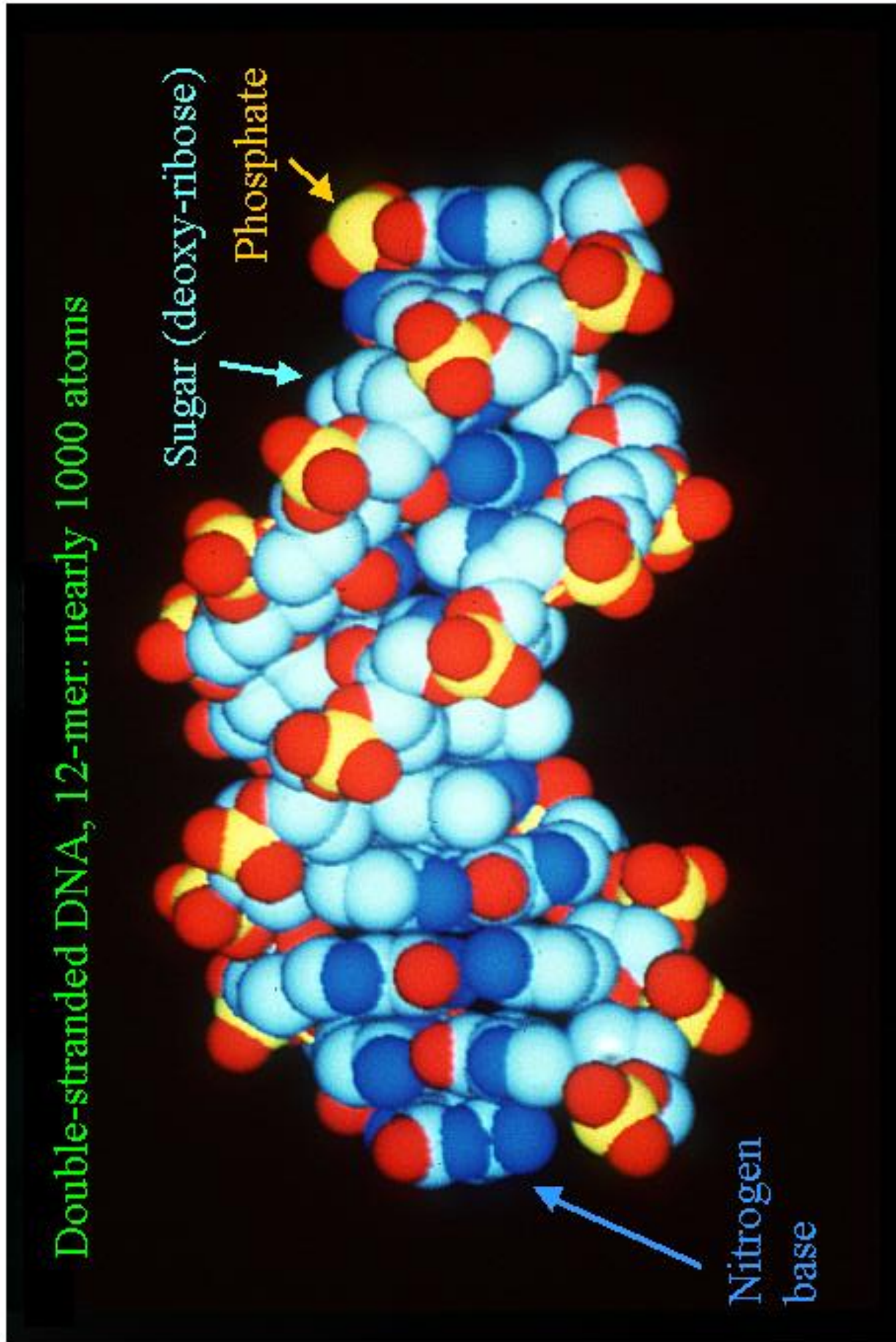
Protozell: must have
rudimentary self-replication,
free energy harvesting and
capacity to evolve, but not
necessarily RNA or proteins

LUCA: has DNA, RNA, tRNA,
ribosomes, proteins, membrane,
elaborate metabolism, photosynthesis



What is that?

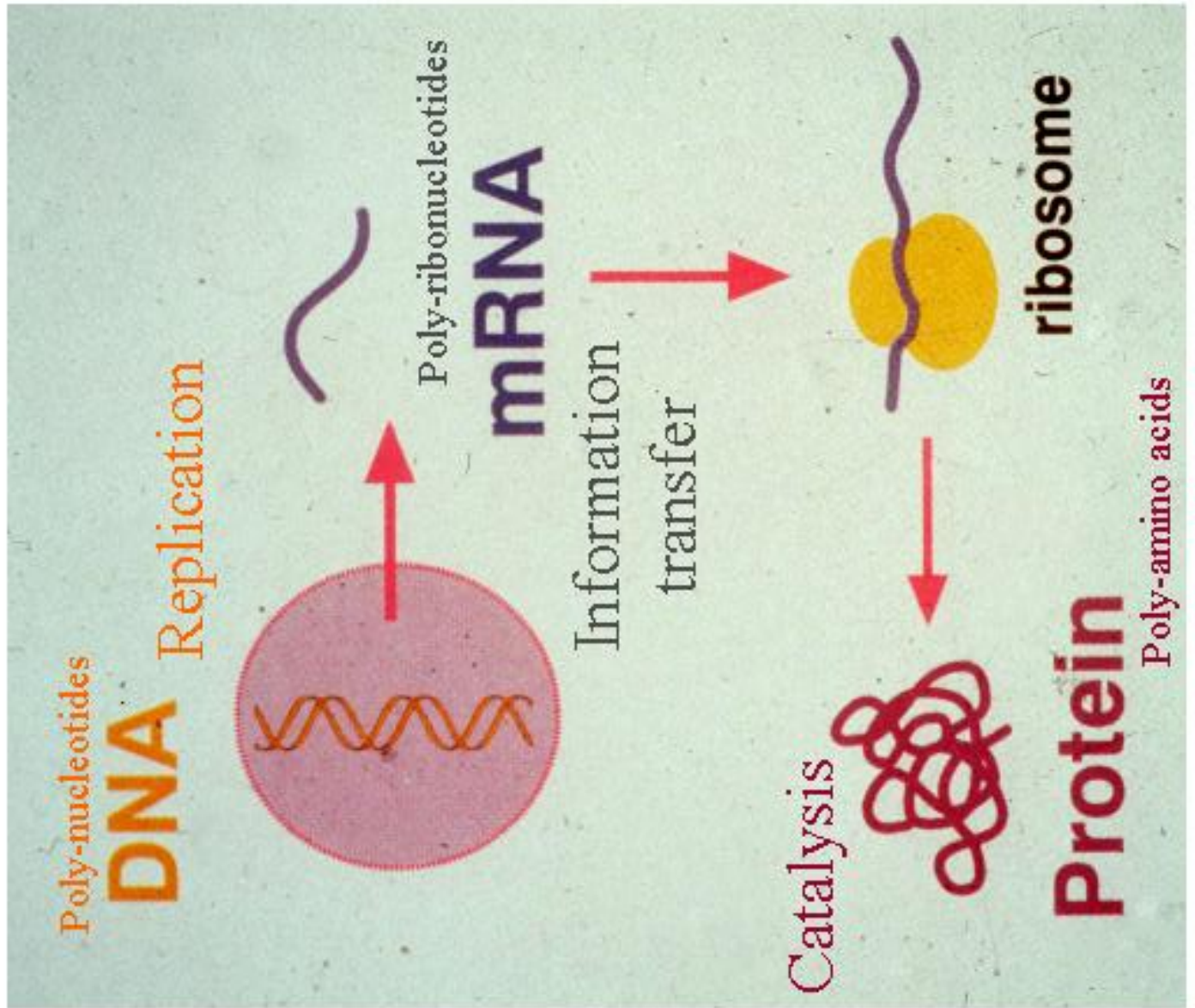




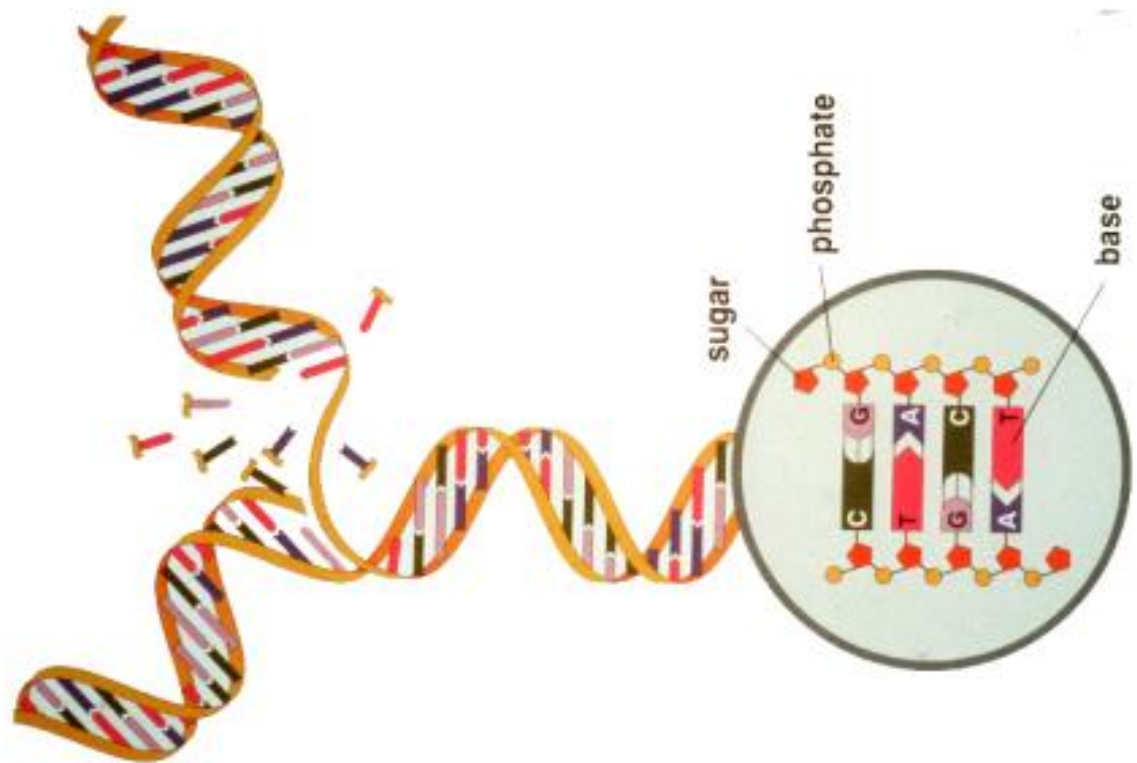
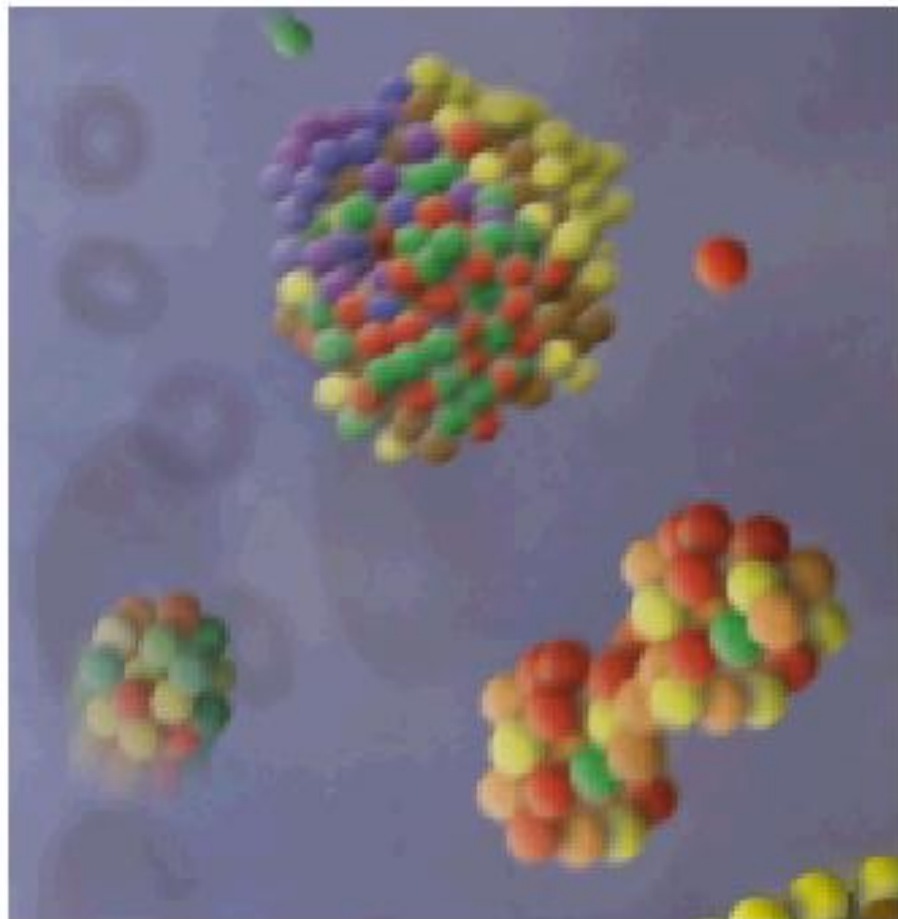
Polymers are too large, too complex. Need to focus on smaller molecules! ("monomers")

Present-day life
is totally based on
polymers.

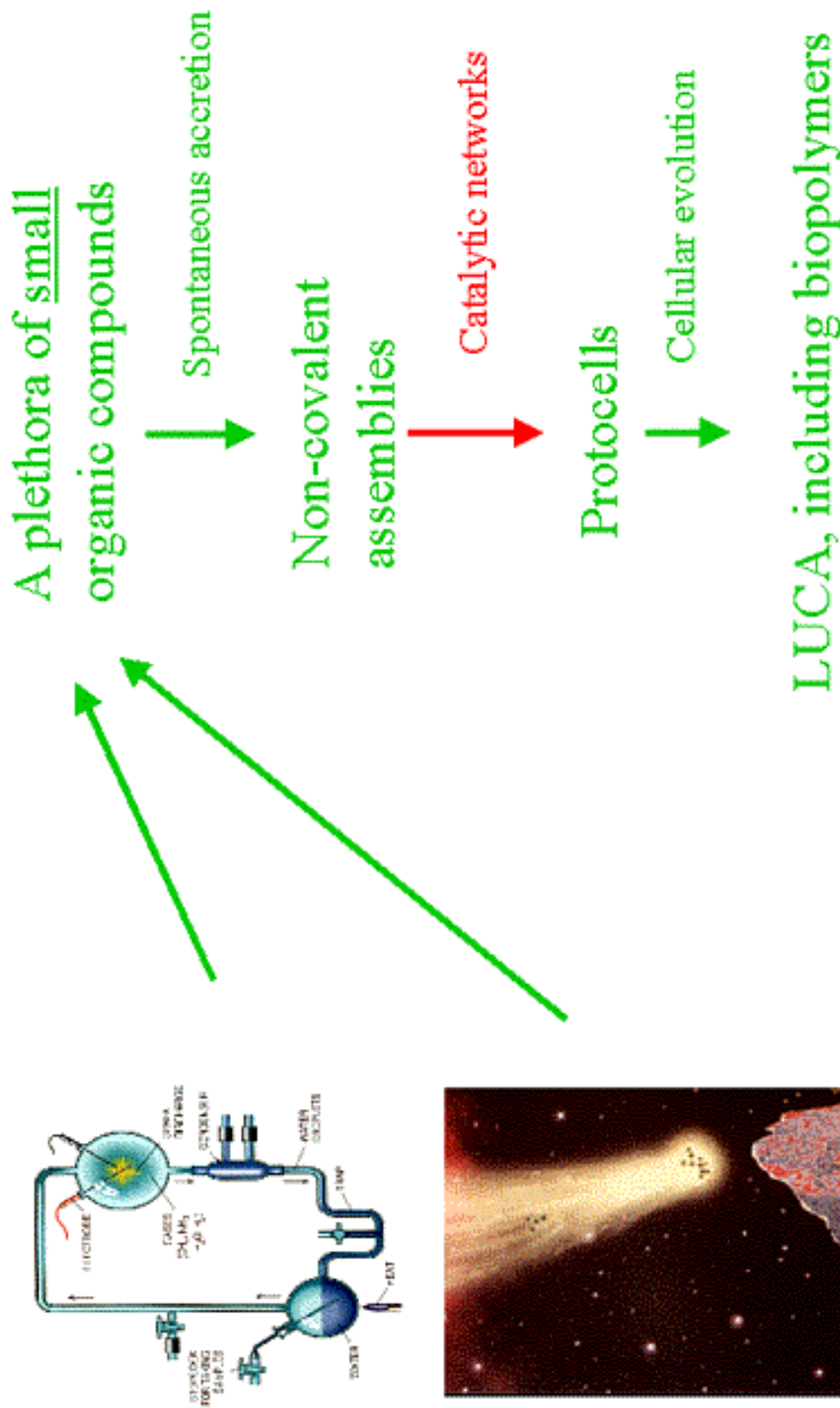
But: did it start
this way?



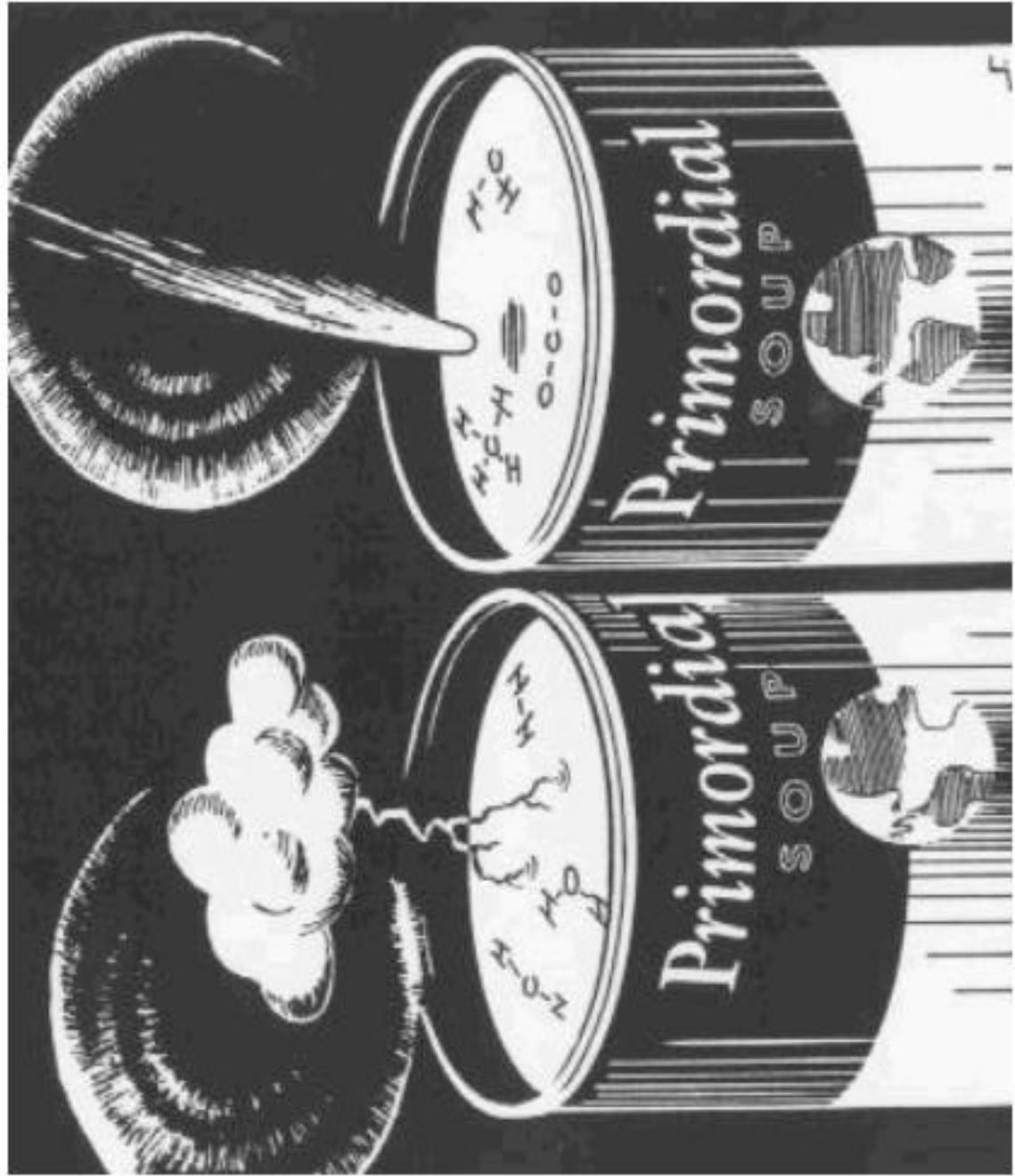
Non-covalent assemblies: the simple alternative.
But many questions need to be answered!

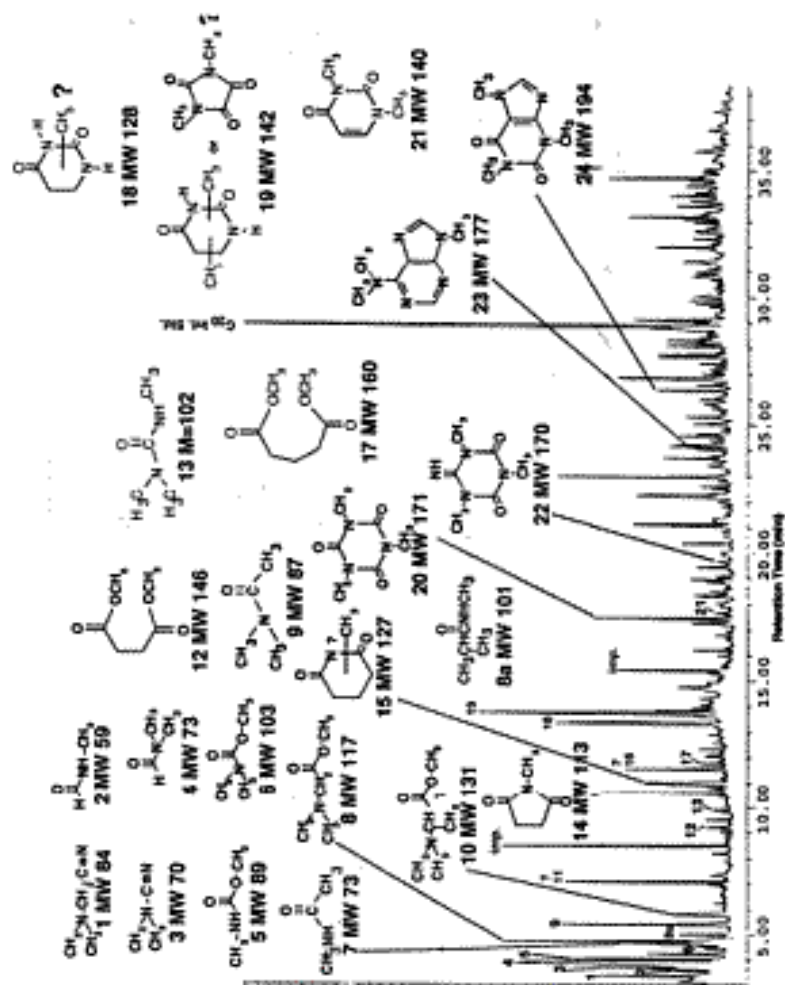


An alternative scenario: Origin of Life without polymers



The origin of life: a planet-scale random chemistry experiment

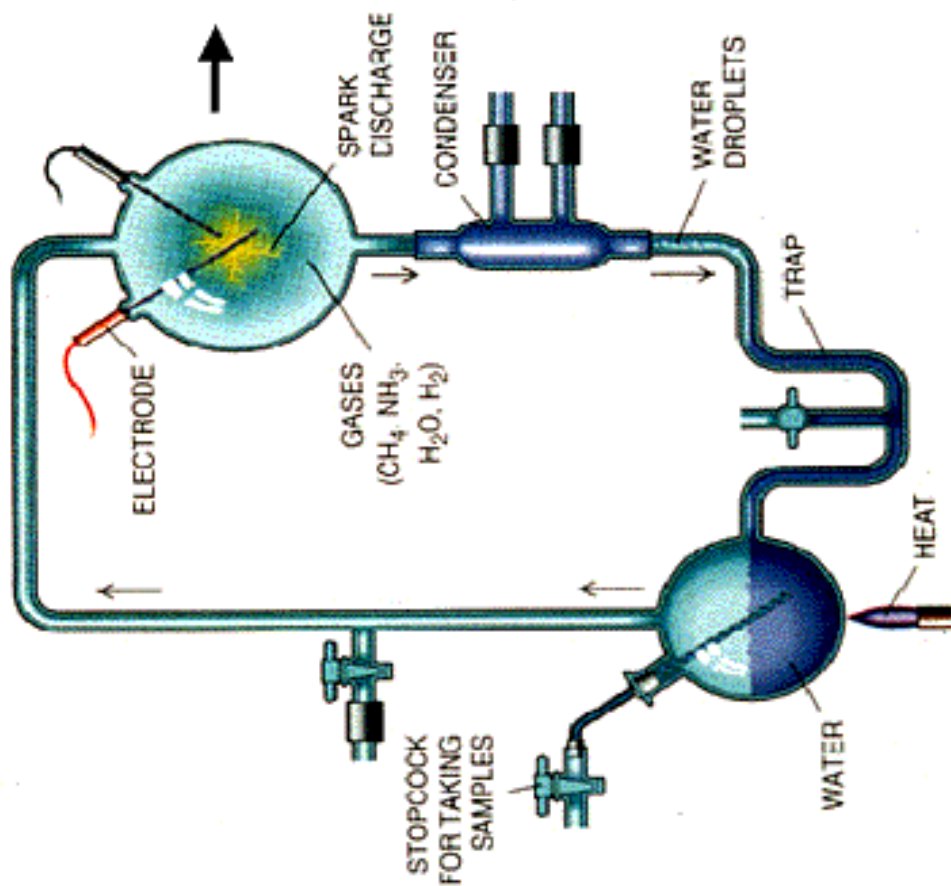


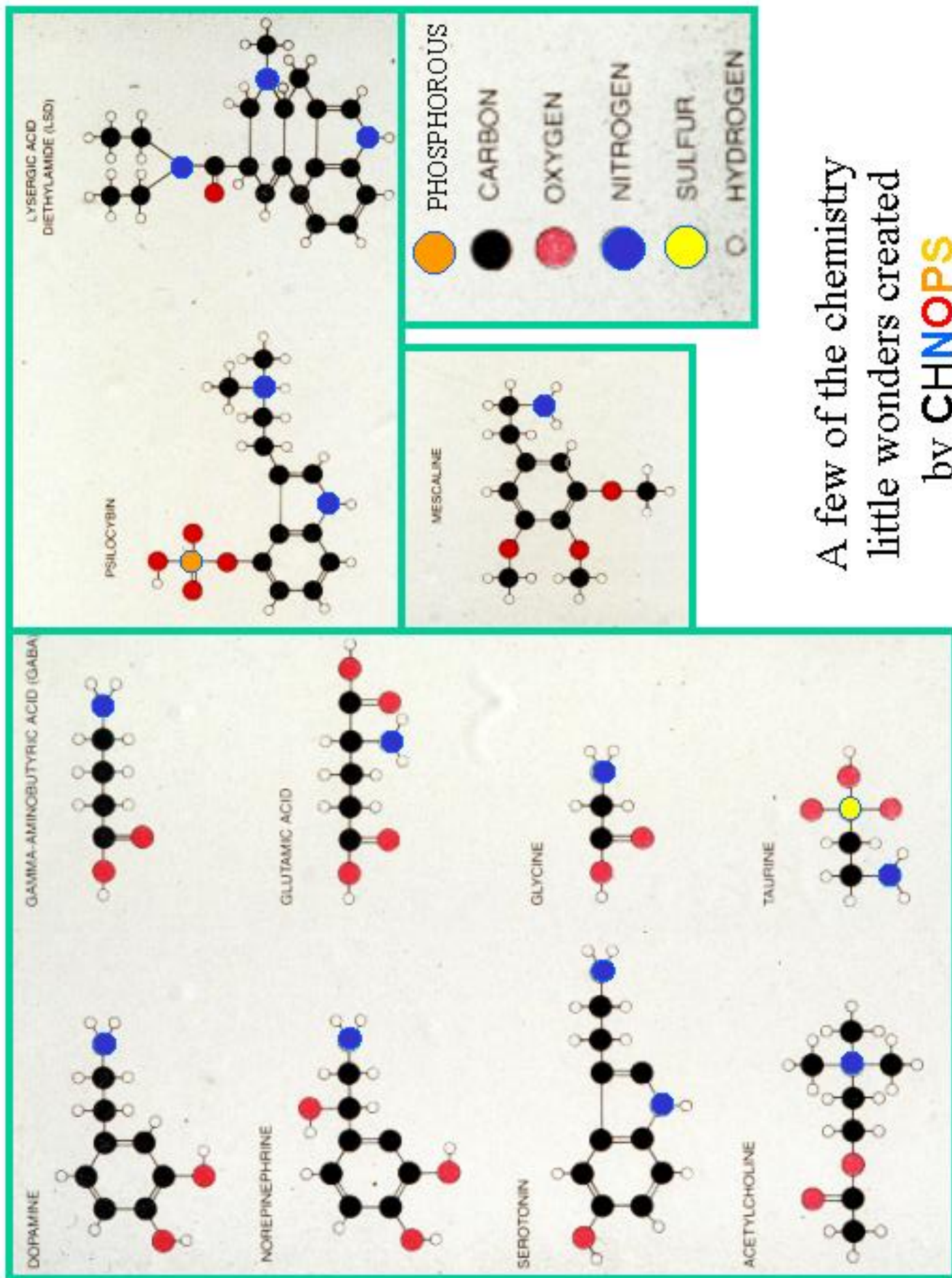


Two different questions:

1) Are compounds of life as we know it present in the mixture?

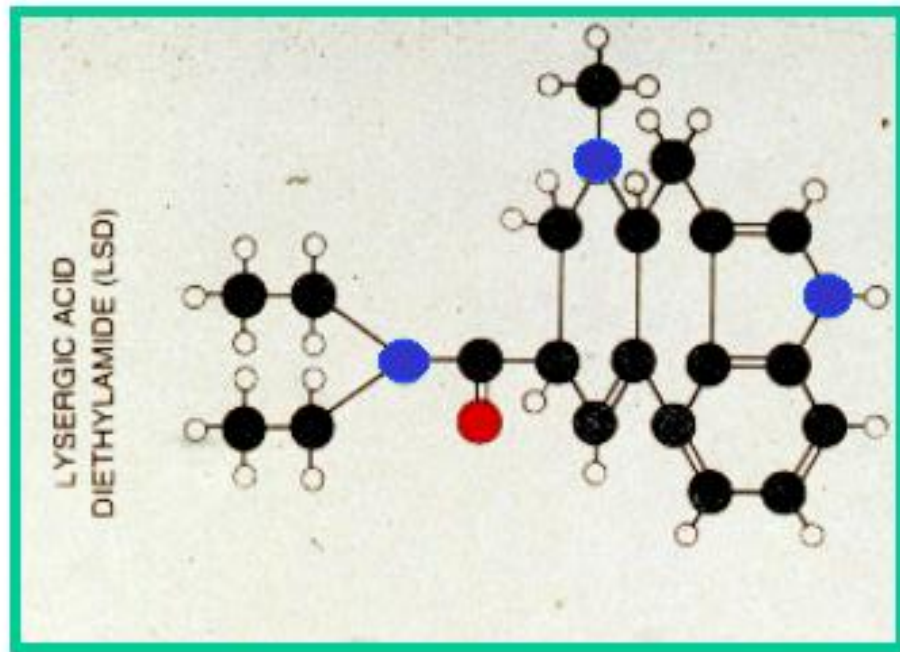
1) What is actually formed and how to get from such a mixture to the narrower assortment of present life forms?



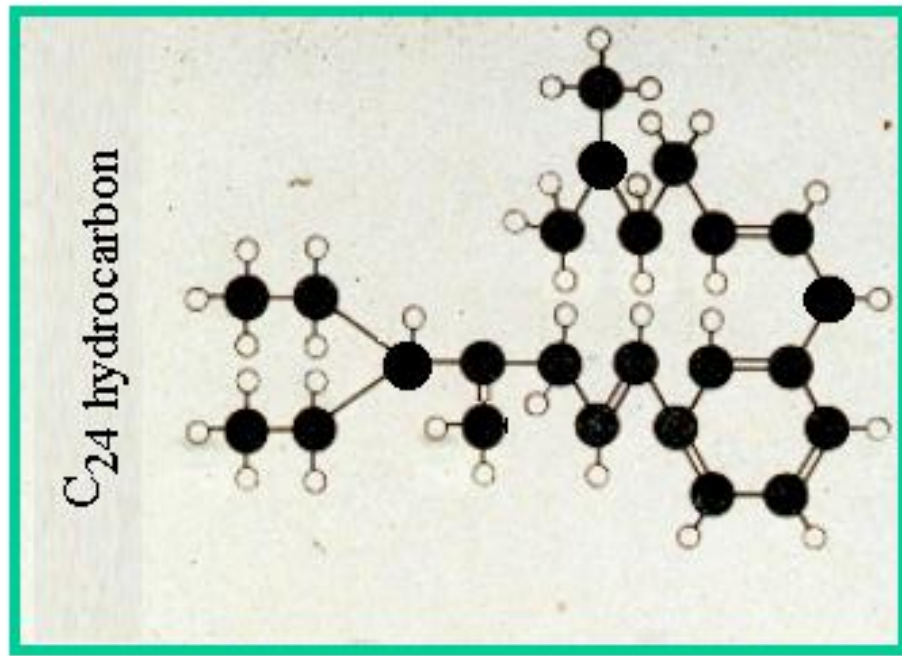


A few of the chemistry
little wonders created
by **CHNOPS**

24 non-hydrogen atoms



24 carbon atoms



This has about
10,000,000
isomers!

In general,
A C_n Compound
has $\sim 0.01e^n$
isomers

With heteroatoms - at least 10^9 isomers

If let loose, **CHNOPS** can generate billions of compounds!

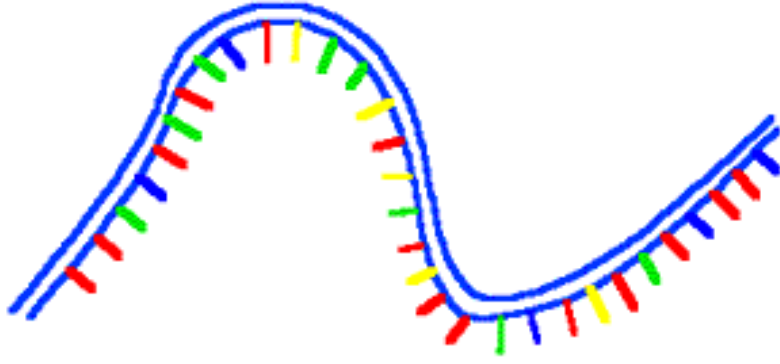
Which substances played the planet-scale random chemistry?



Lipid World:

Easy monomers

Easy non-covalent “polymers”

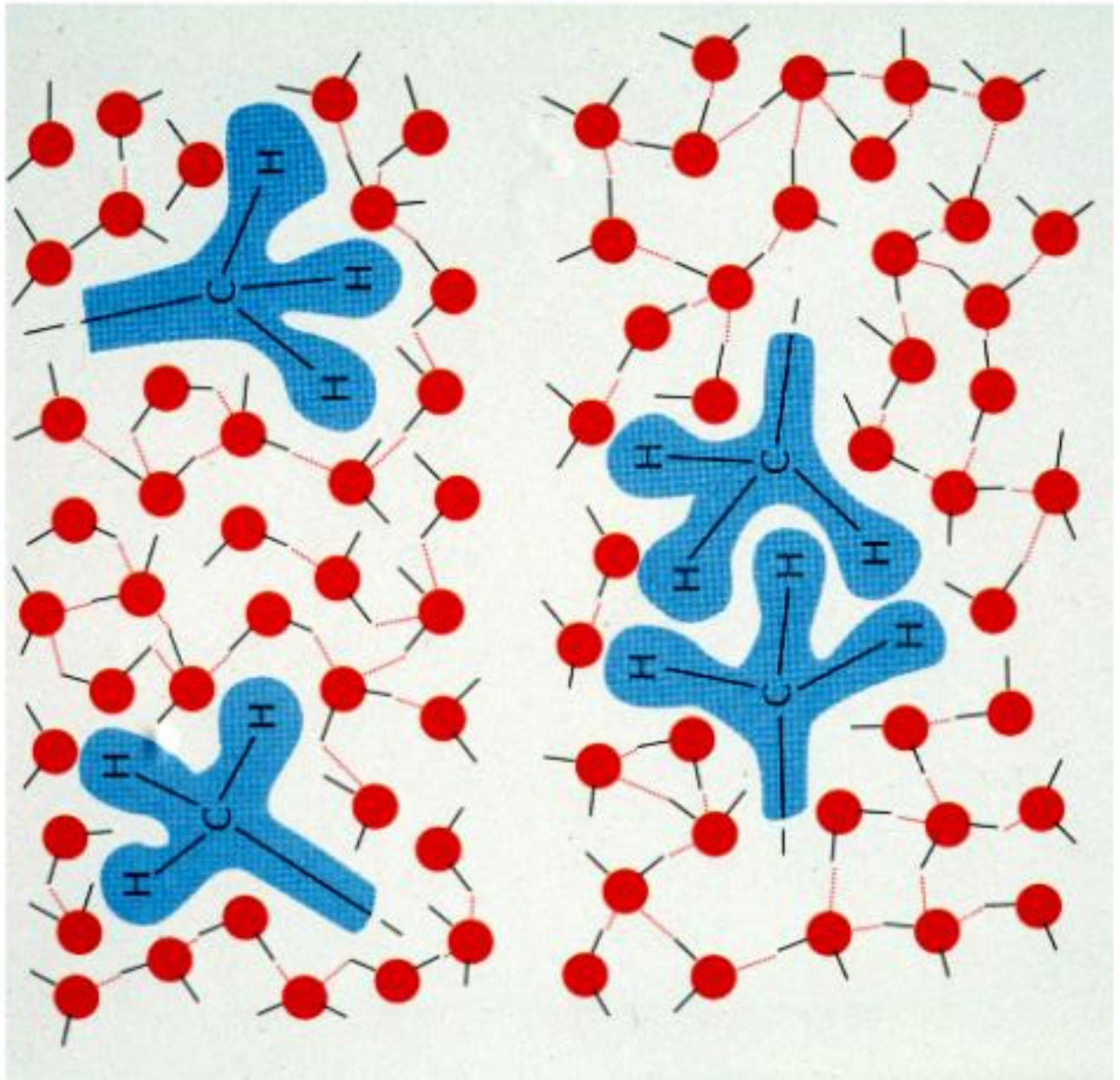


RNA World:

Rare monomers

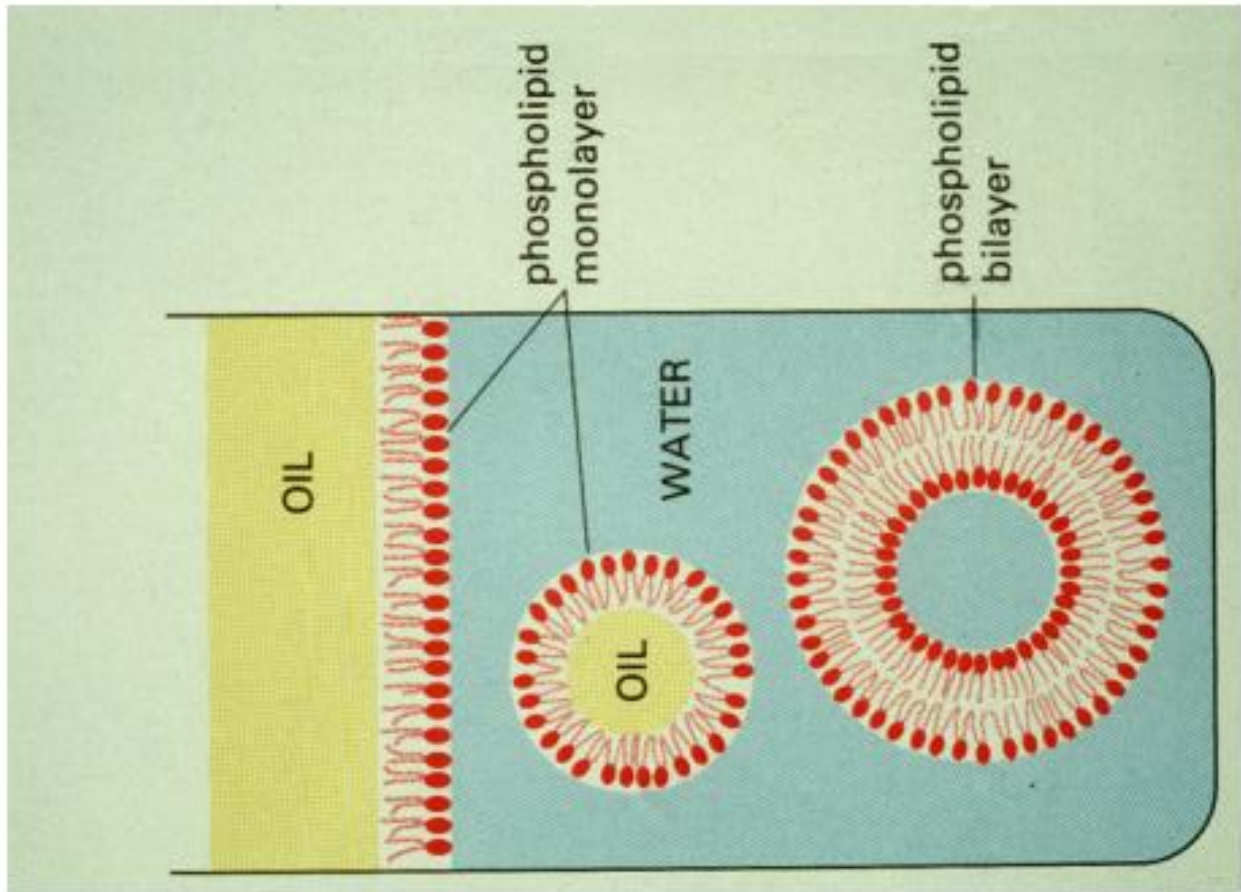
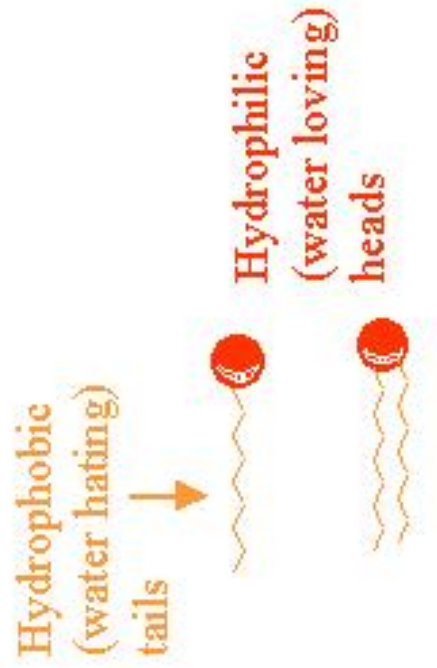
Difficult covalent
polymerization

HYDROPHOBIC FORCES



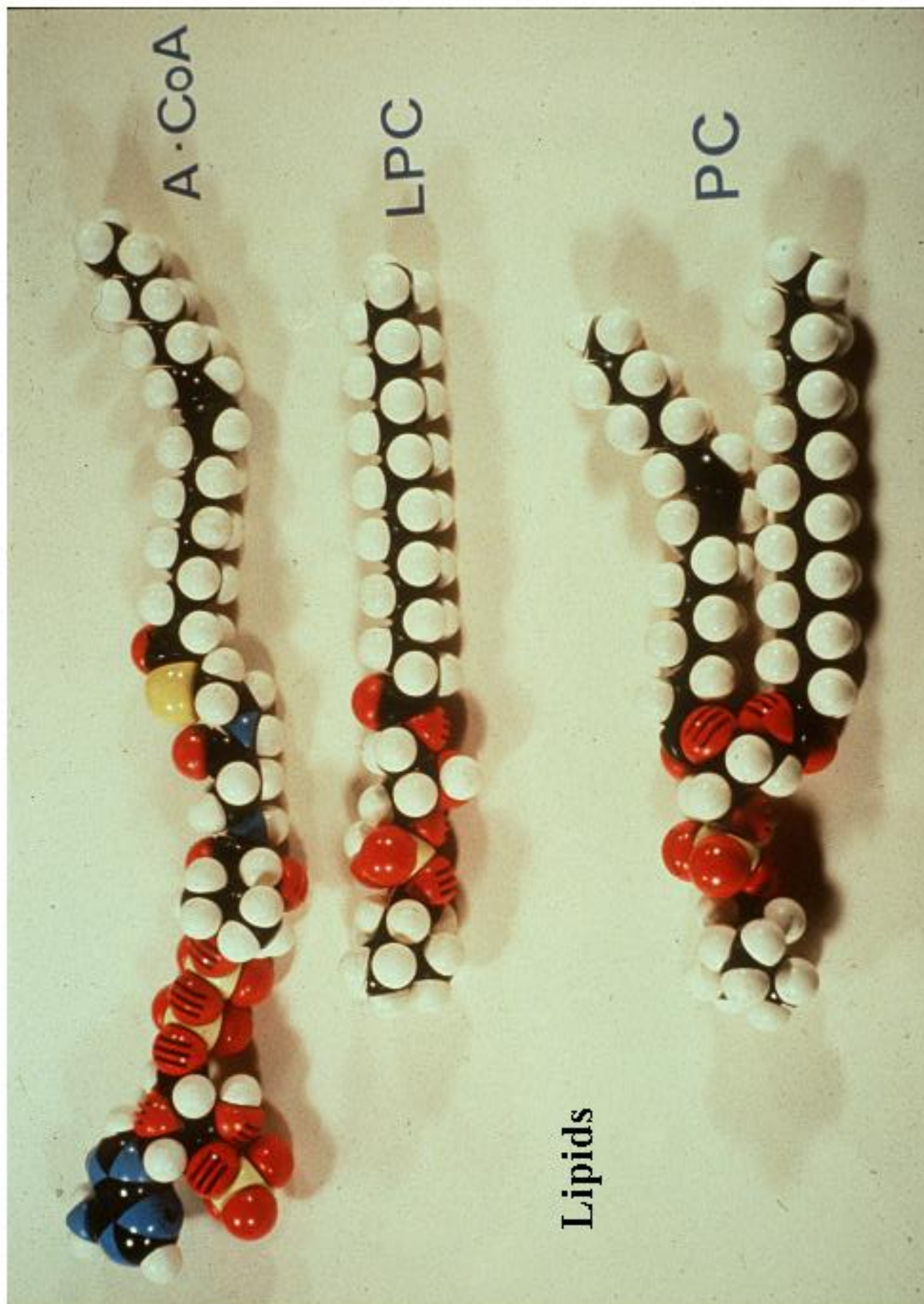
These interactions
is driven by the
increased entropy,
due to disordering
of **water molecules**,
as they get released
from the interface
between two “oily”
molecules

Amphiphilic Molecules (e.g. phospholipids)

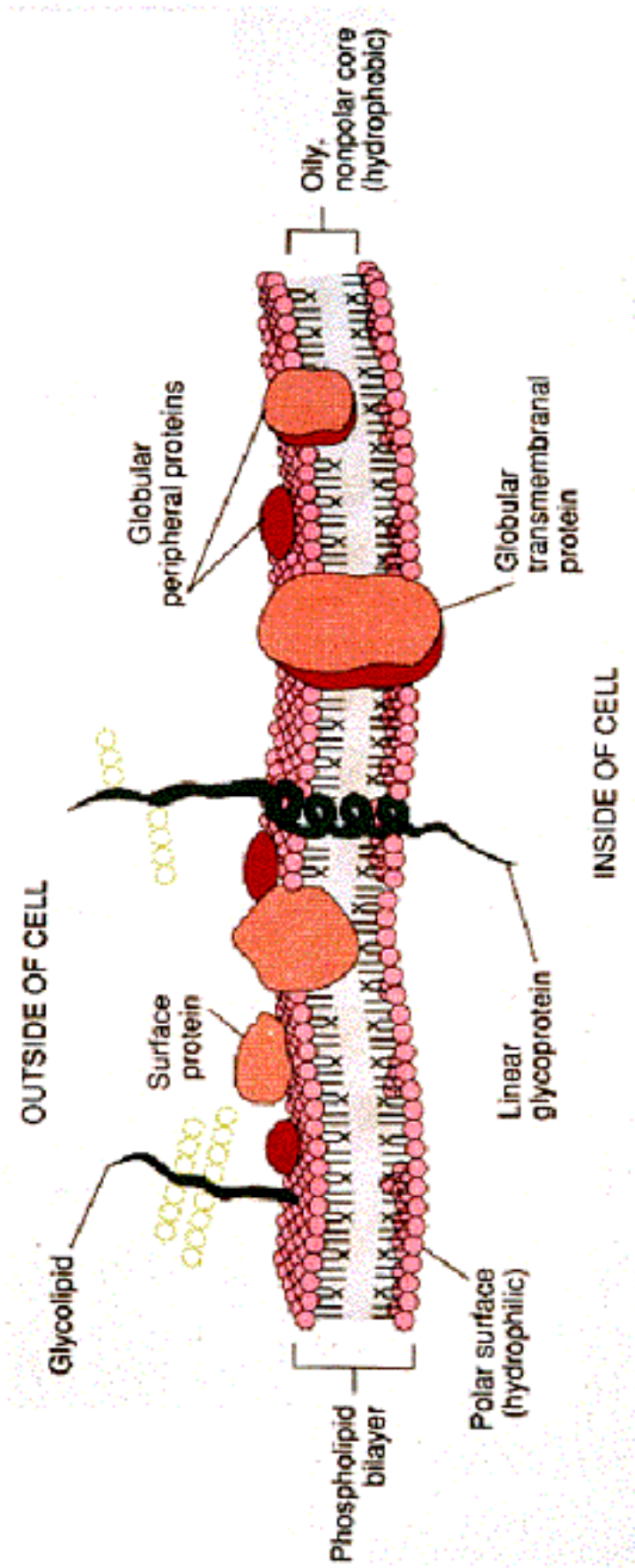


Micelle

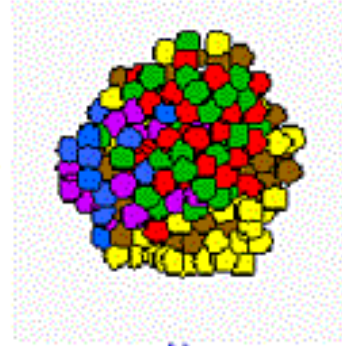
Vesicle



Present-day membranes are bustling places, where most of the business is carried out by **proteins** within or near the **lipid bilayer**



But could it have begun like this?



A complex aggregate of lipid-like molecules with diverse chemical groupings

Lipids are
abiotically
available

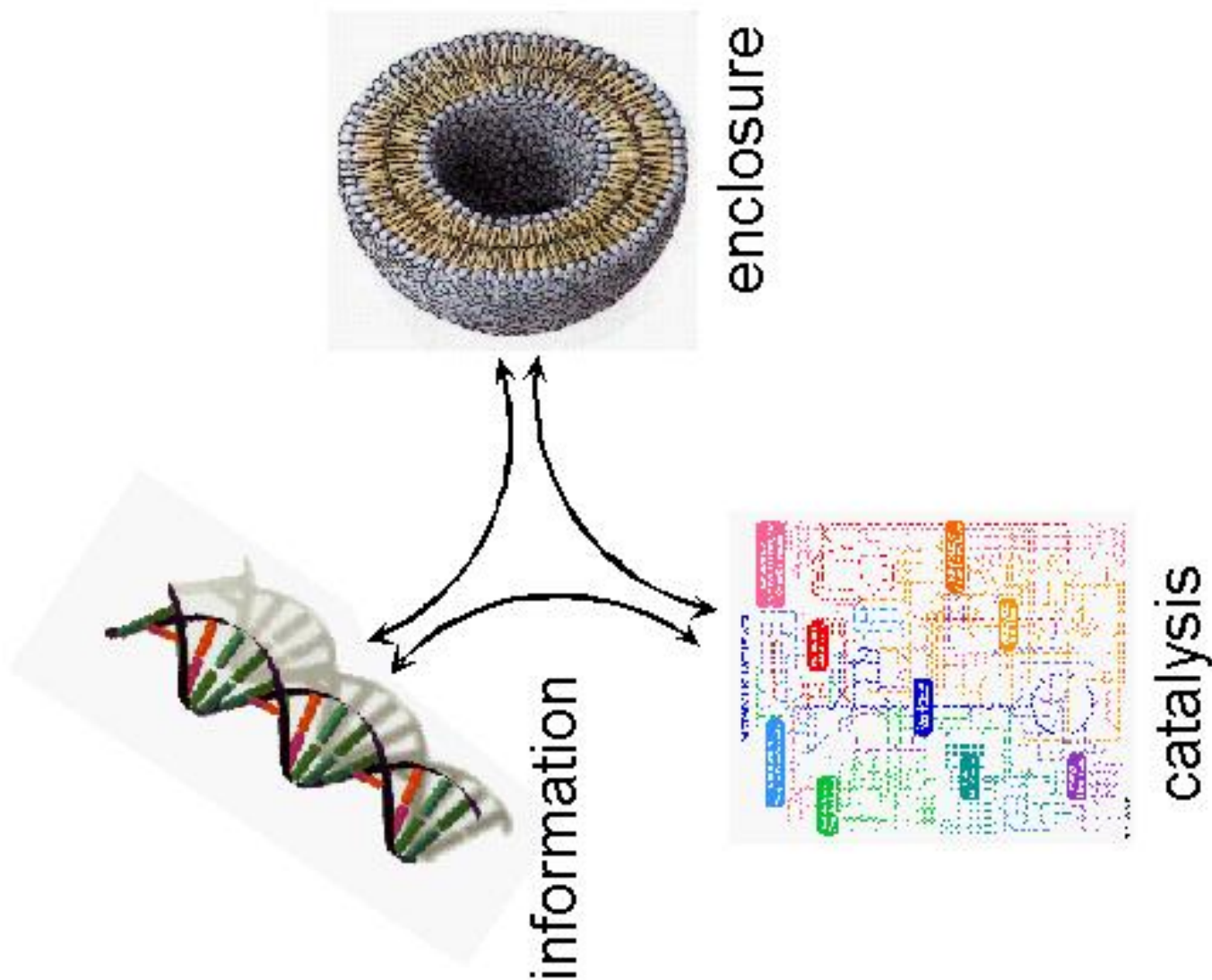
**AMPHIPHILIC COMPONENTS OF THE
MURCHISON CARBONACEOUS CHONDRITE:
SURFACE PROPERTIES AND MEMBRANE FORMATION**

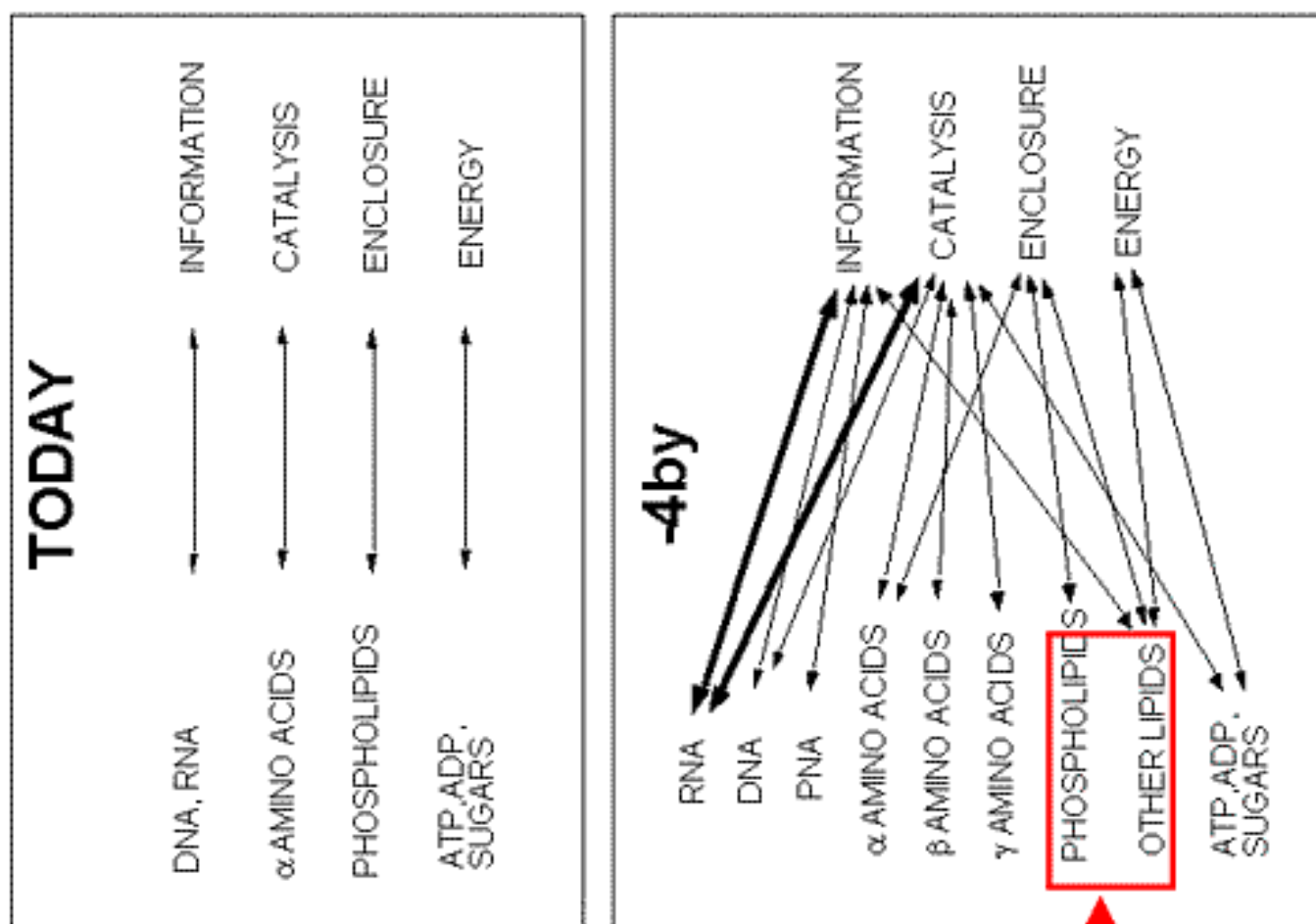


D. W. DEAMER R. M. PASHLEY

Origins of Life & Evolution of the Biosphere, Volume 19 (1989), pp. 21-38.

Primordially,
lipids could do
much more than
they do in
present-day
cells

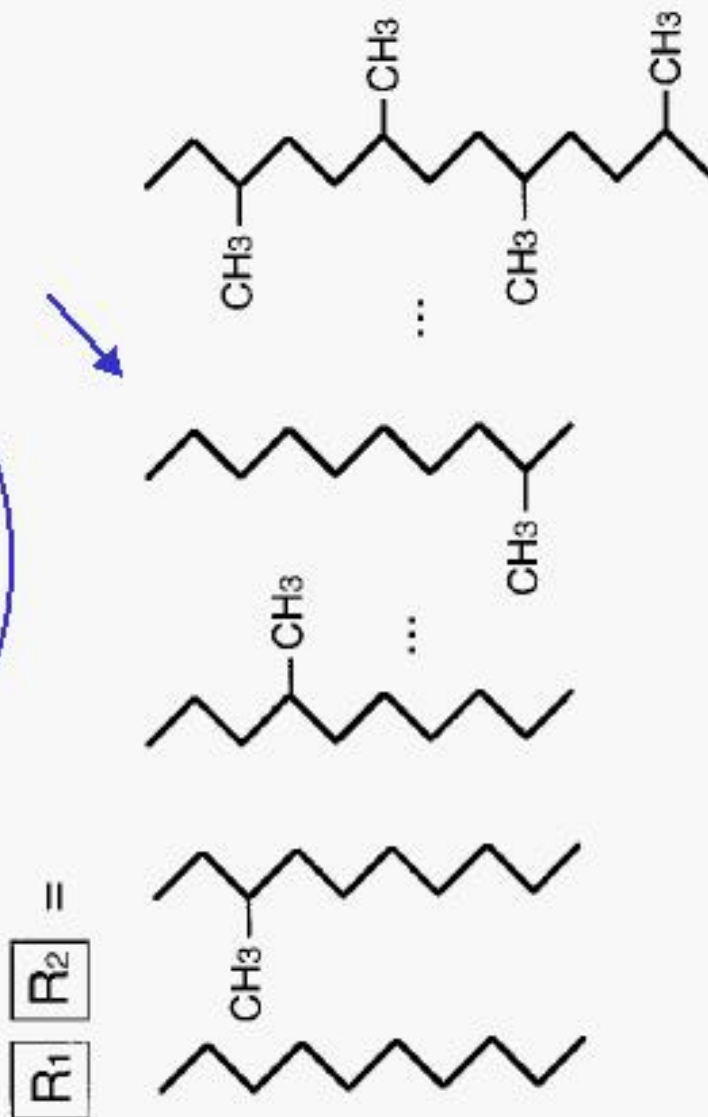
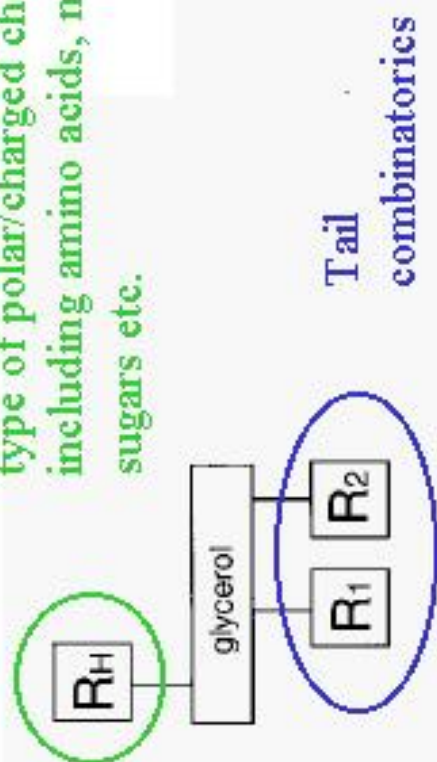




Amphiphiles:
spontaneous
assembly!

Combinatorial lipids

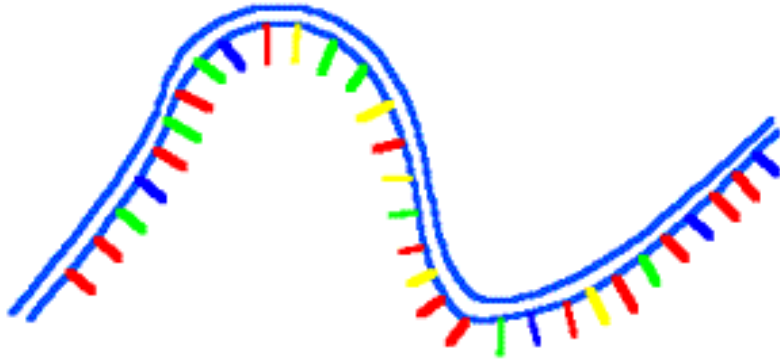
The head group can be any type of polar/charged chemical moiety, including amino acids, nucleotides, sugars etc.



Where does information reside?

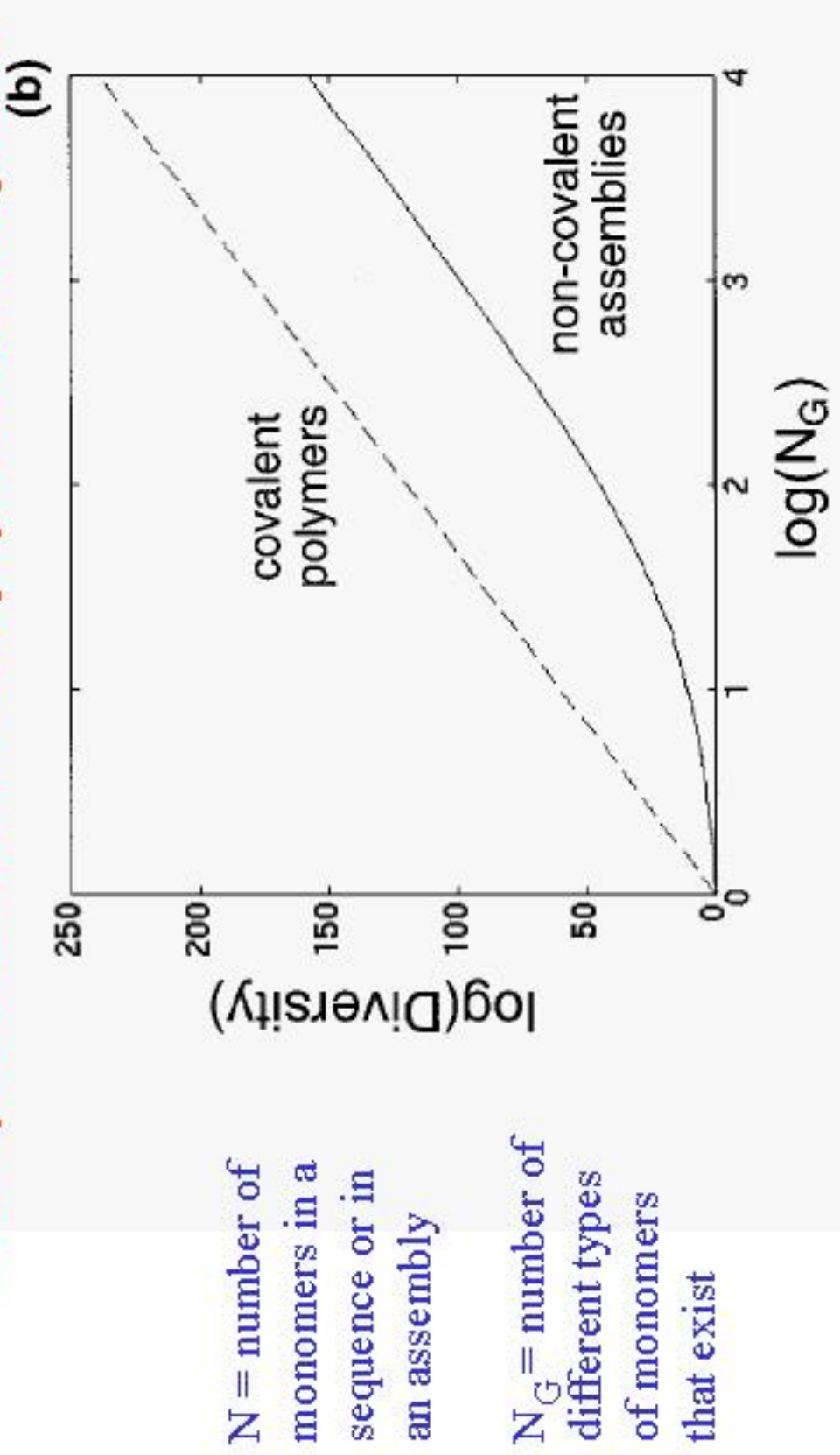


Lipid World:
Compositional diversity!
(A novel concept)



RNA World:
Sequence diversity

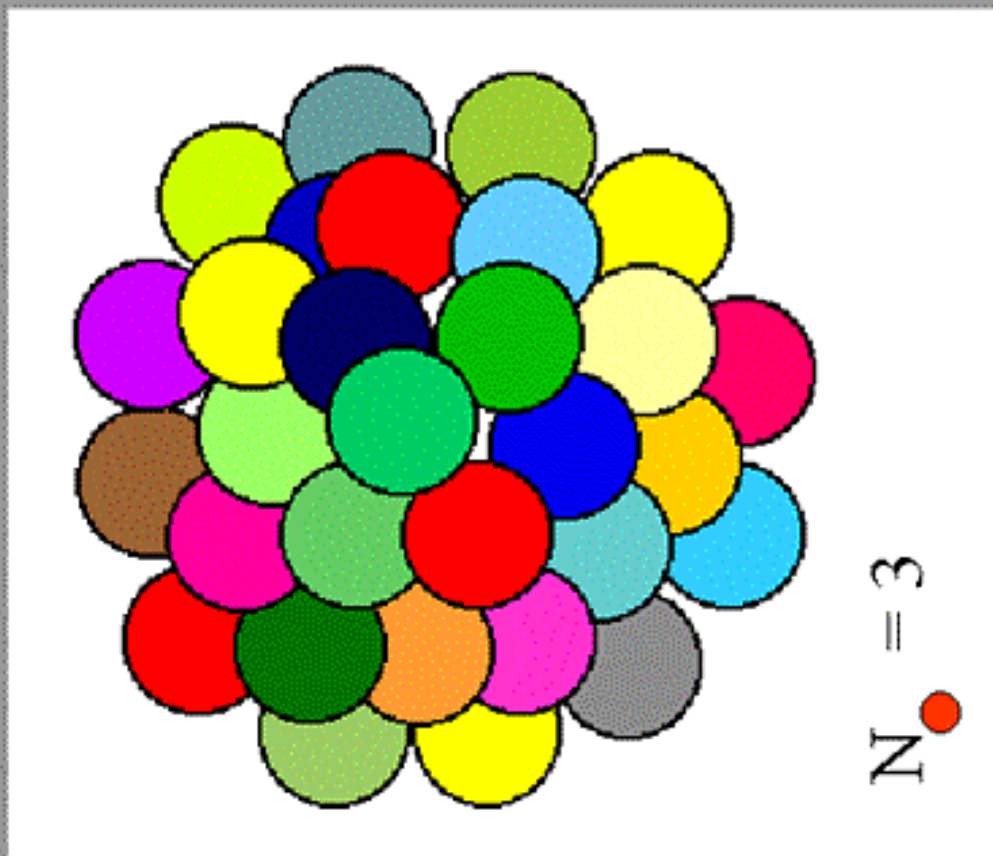
Both diversity (D) functions have the same limiting slope
Diversity is the number of different polymers/assemblies possible



Polymers: $D = (N_G)^N$

Assemblies: $D = (N_G + N - 1)! / (N!(N_G - 1)!)$

Non-covalent assemblies have compositional information



n_i is the number of molecules of type i in the assembly

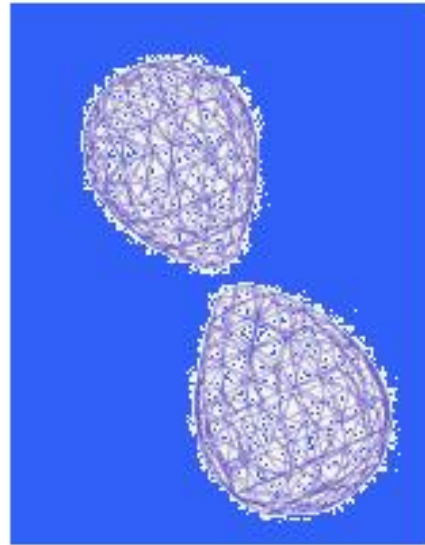
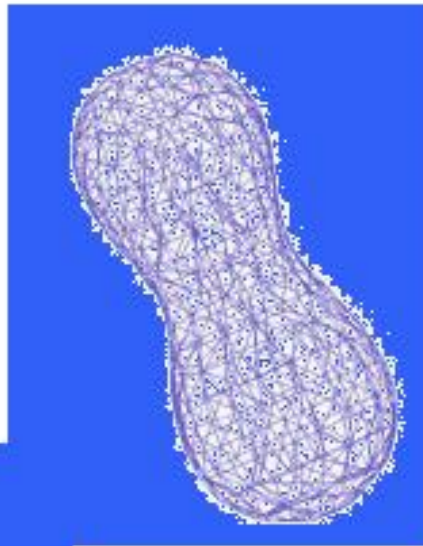
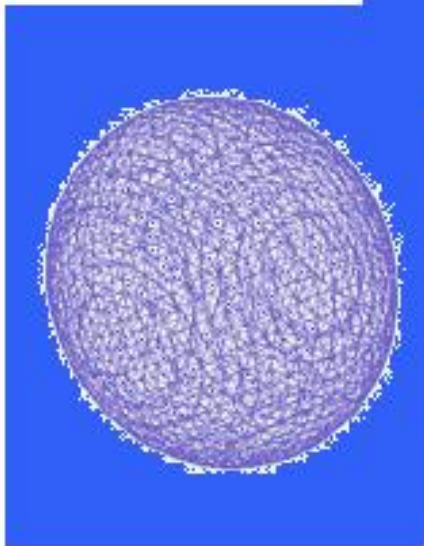
The vector \mathbf{n} defines the Assembly's composition

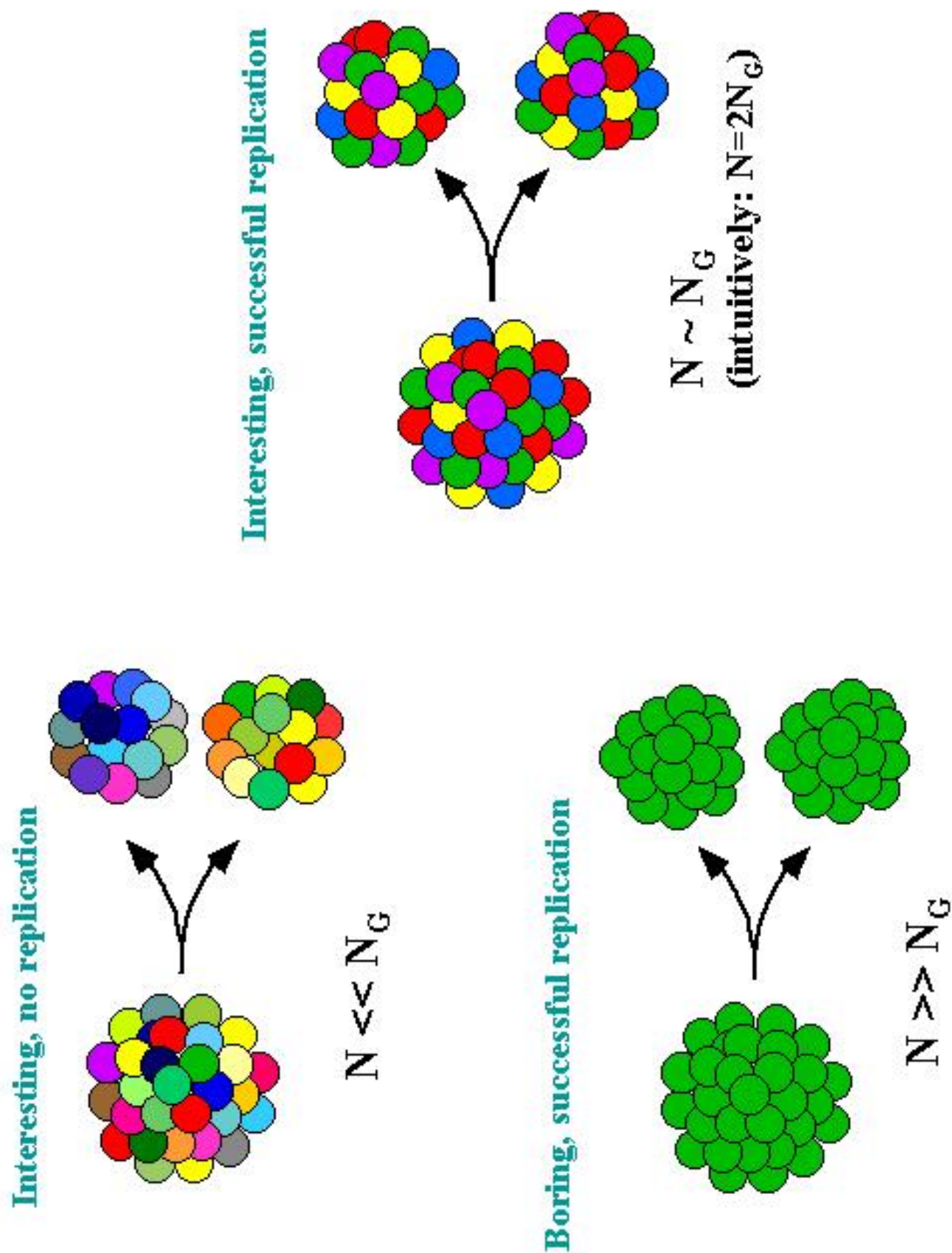
The scalar product

$$H = \mathbf{n}' \cdot \mathbf{n},$$

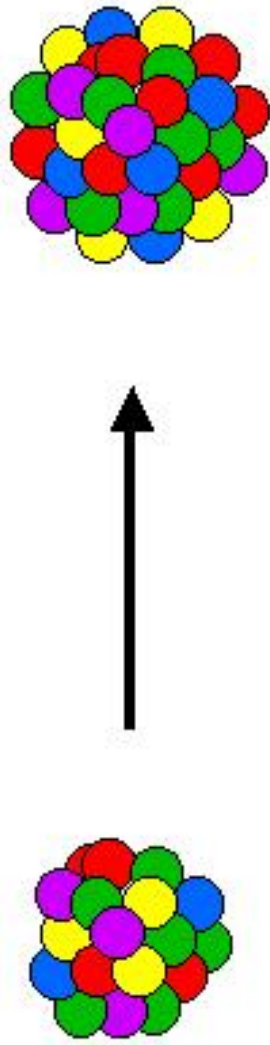
is the distance between compositions

Splitting is easy and natural

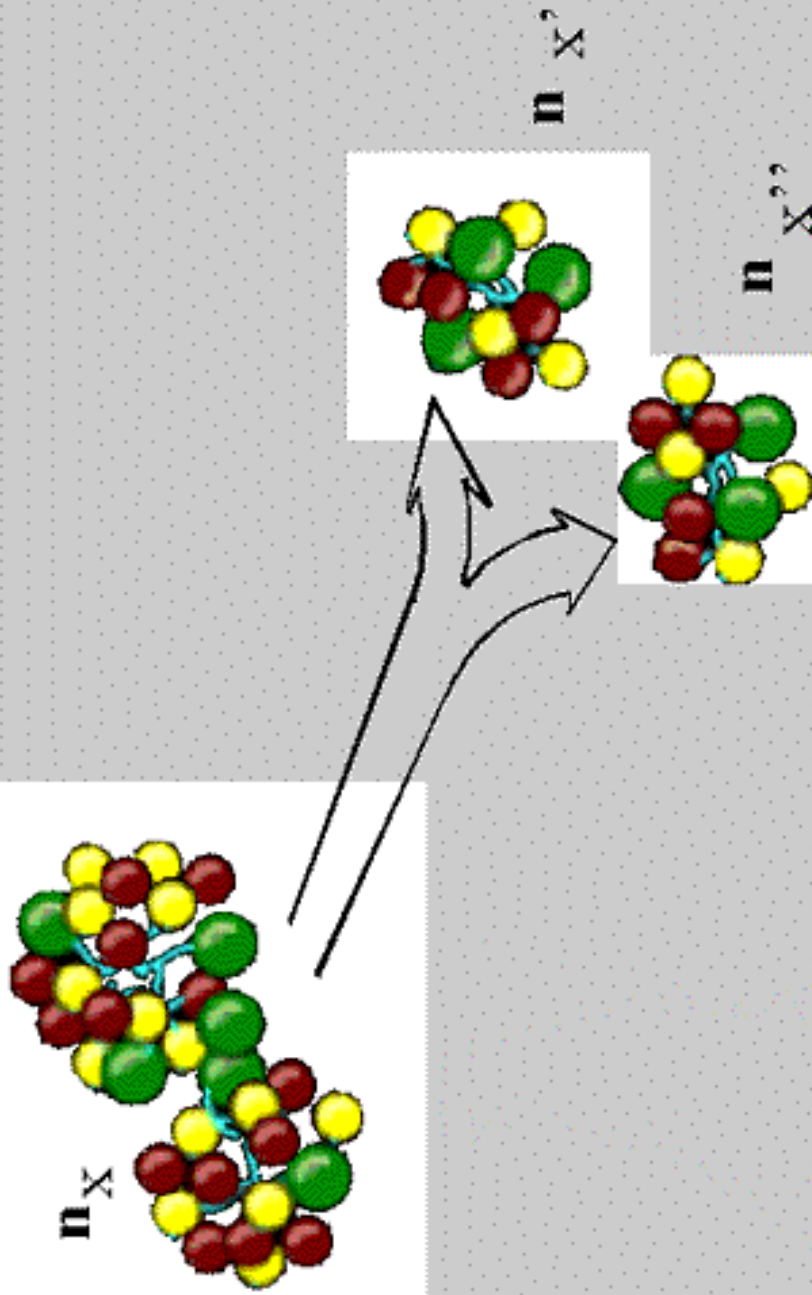




How to achieve replication capacity?
By homeostatic growth! (time-invariable concentrations)



This may be mediated by mutually catalytic networks!
(see Freeman Dyson, Stuart Kauffman)

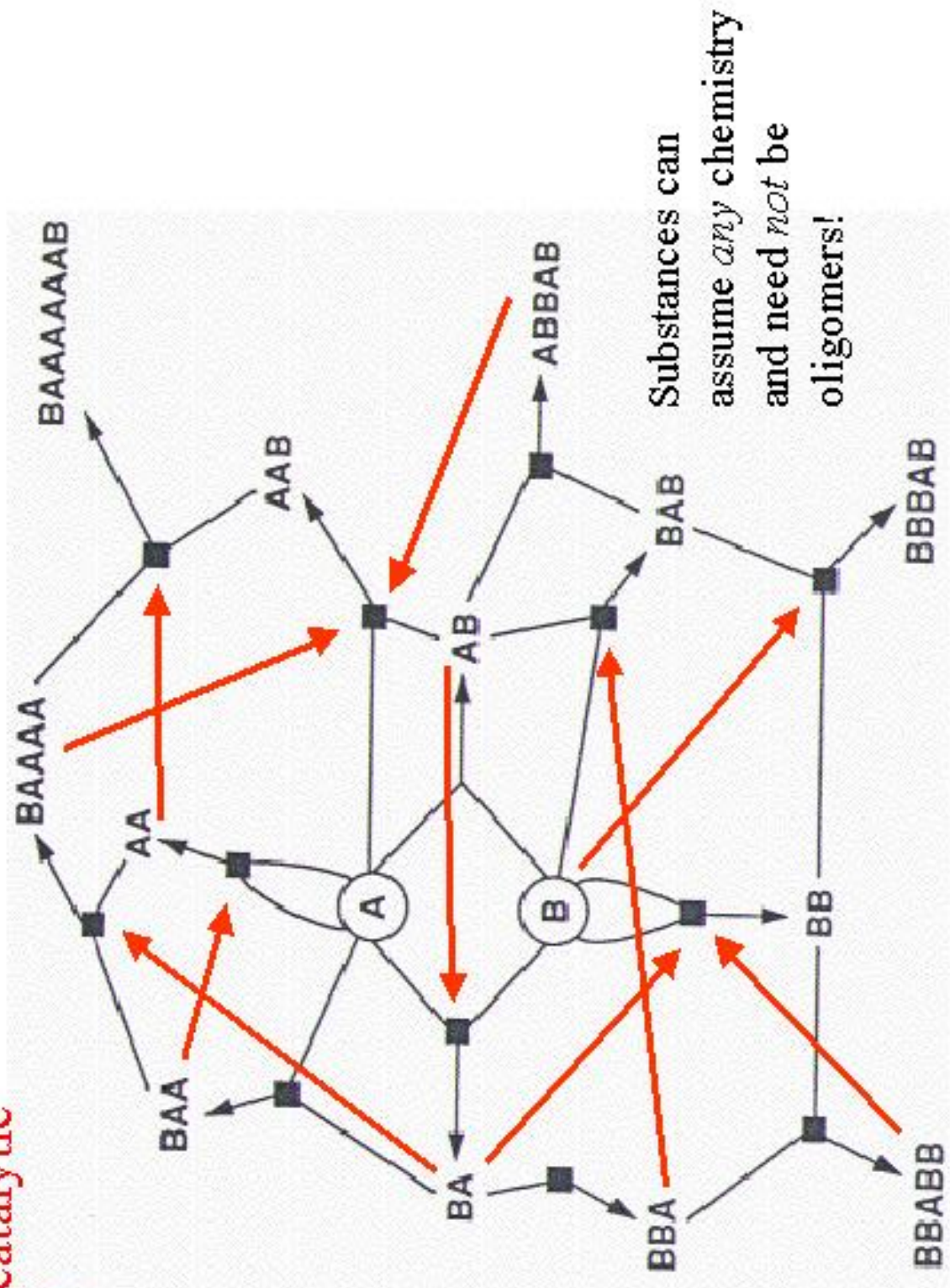


Requirements for successful replication:

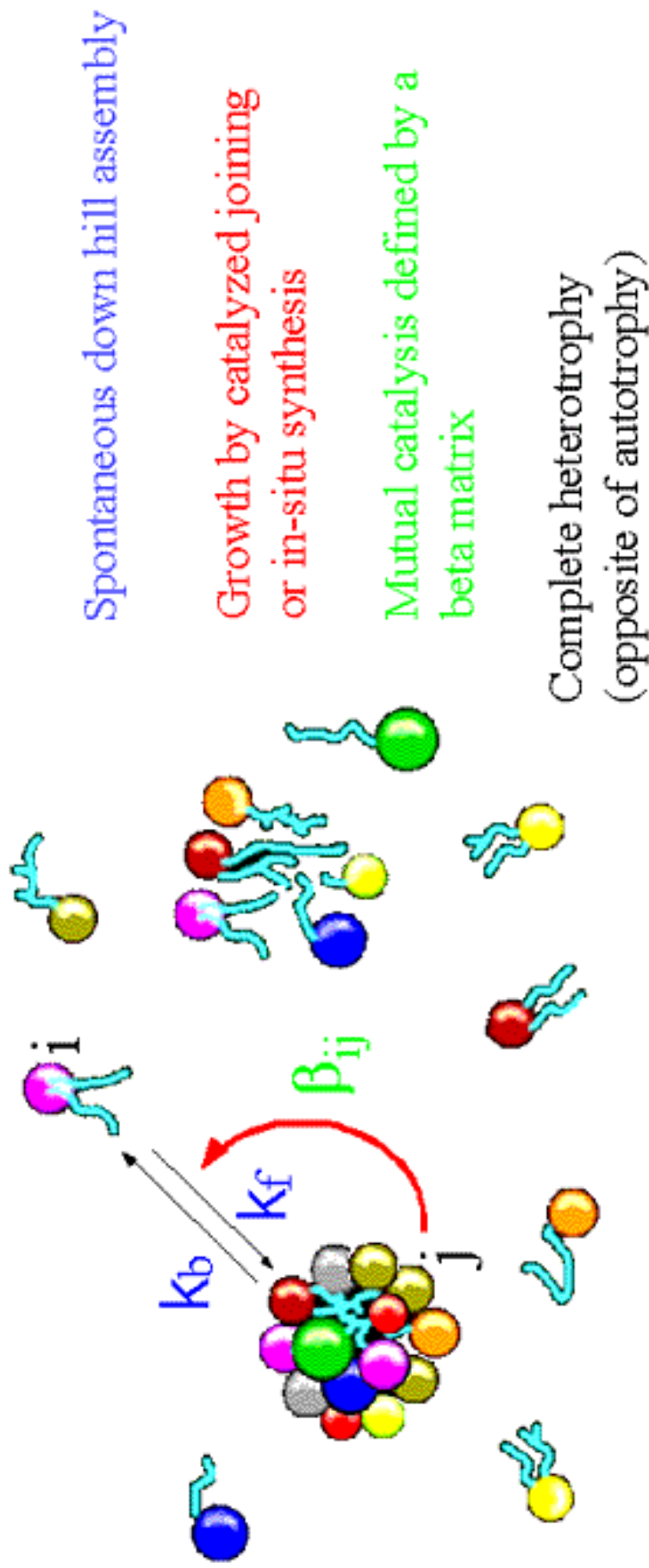
- 1) Efficient catalyzed growth
- 2) Homeostatic preservation of composition
- 3) High inheritance fidelity upon splitting

$$n_X \sim n_{X'} \sim n_{X''}$$

Kauffman/Dyson:
Mutually catalytic
networks

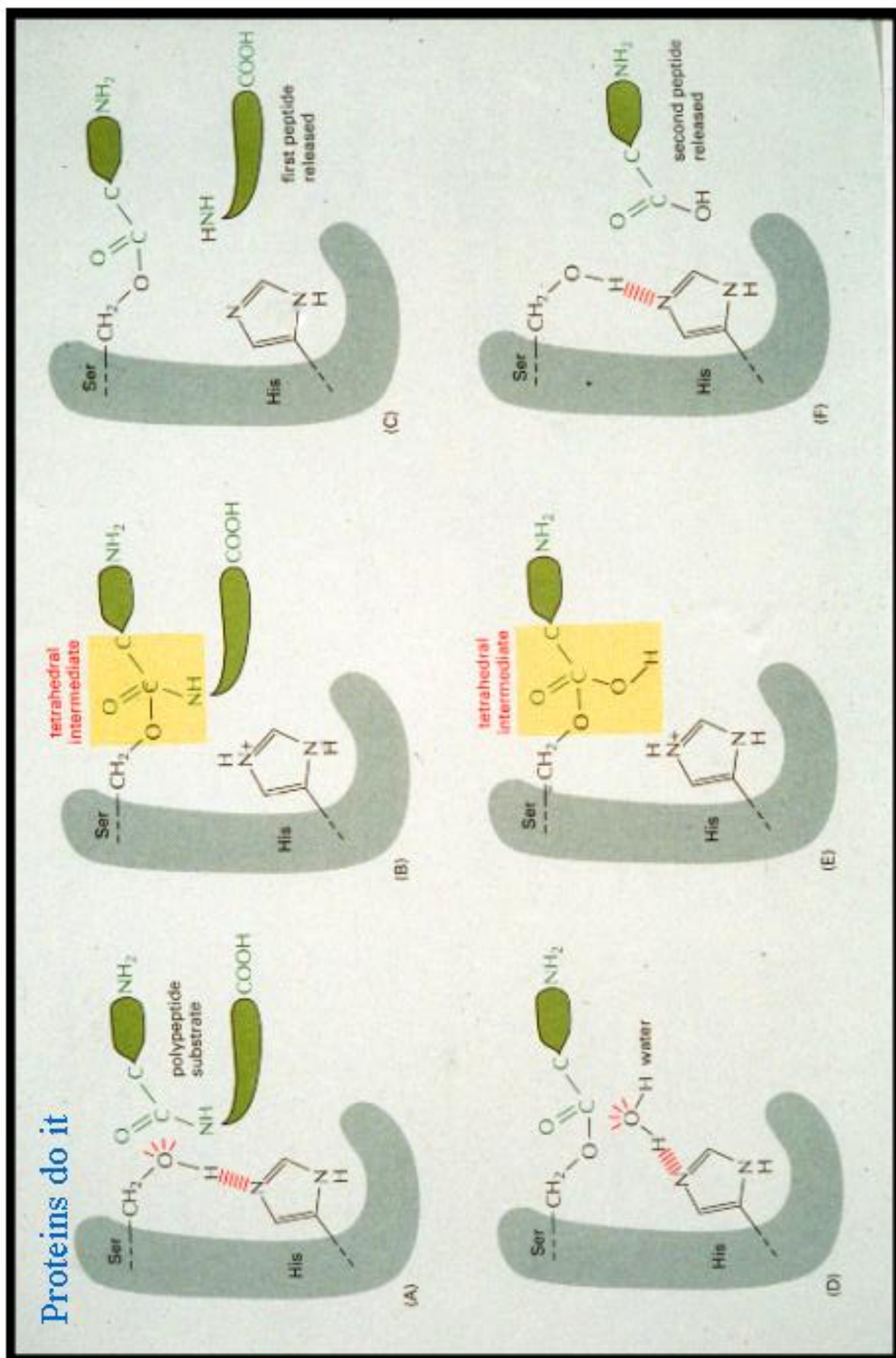


GARD: the Graded Autocatalysis Replication Domain model (Segre, Ben-Eli and Lancet, PNAS 2000)



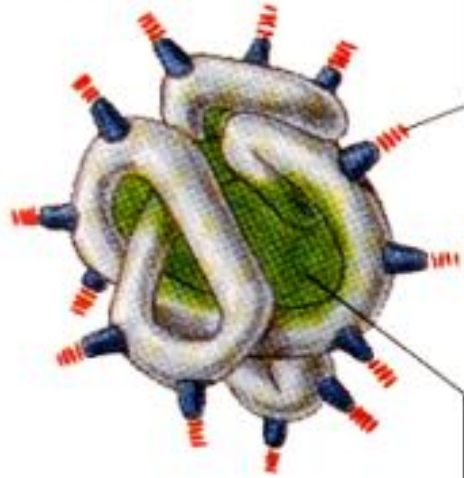
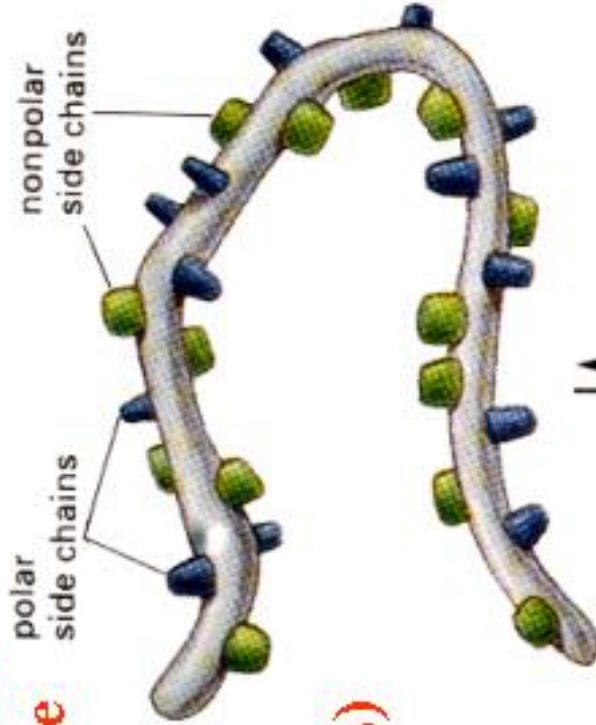
But, are lipids catalytic at all?

Enzyme catalysis = right chemical groupings in a correct, rigid orientation.
This allows concerted chemical action that lowers the activation energy



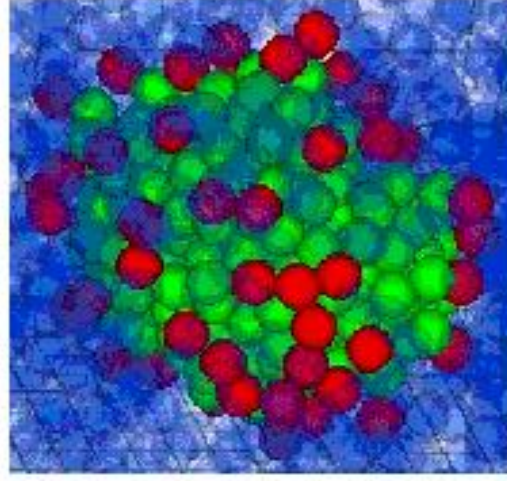
**But are proteins
the only ones?
no, there are
enzyme-
mimetics
(e.g. cage
Compounds)**

unfolded polypeptide



folded conformation in aqueous environment

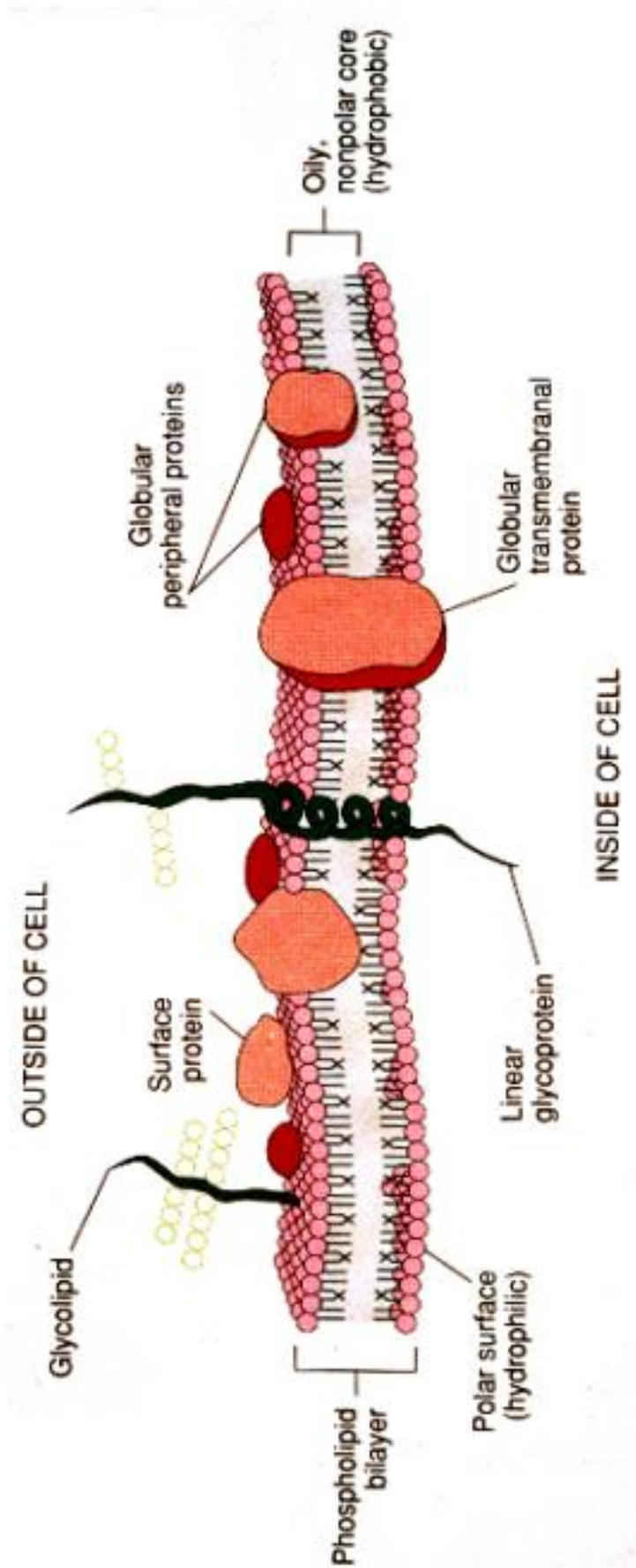
A heterogeneous lipid micelle may be viewed as a “disconnected” folded globular protein



(Lipowsky)

hydrogen bonds can form to polar side chains on the outside of the molecule

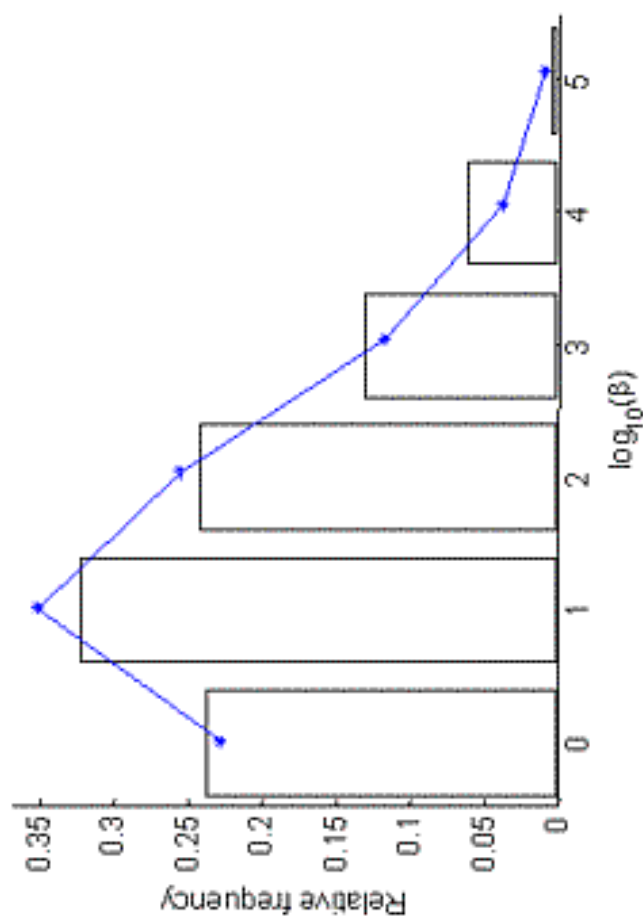
It could have enzyme mimetic properties



Lipids modulate enzymatic reactions in membranes

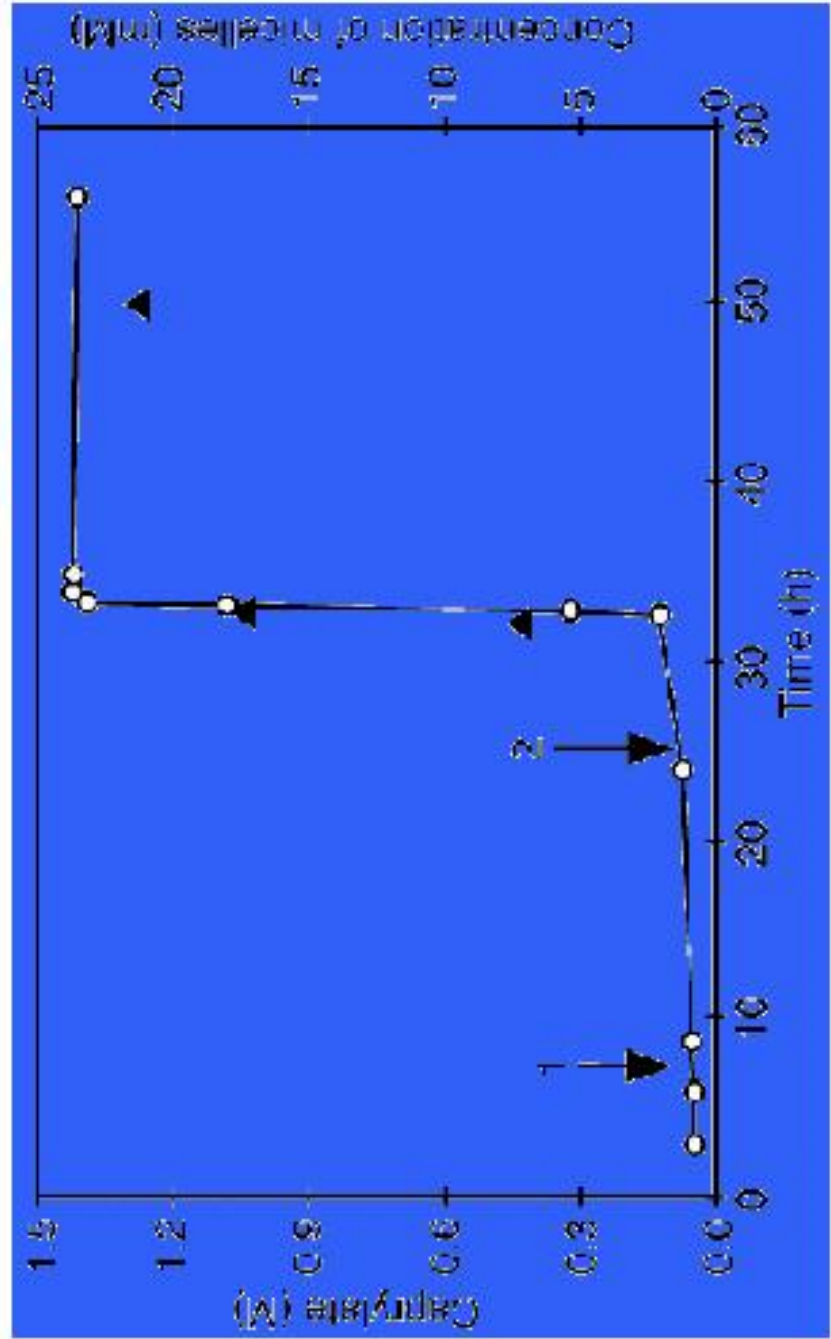
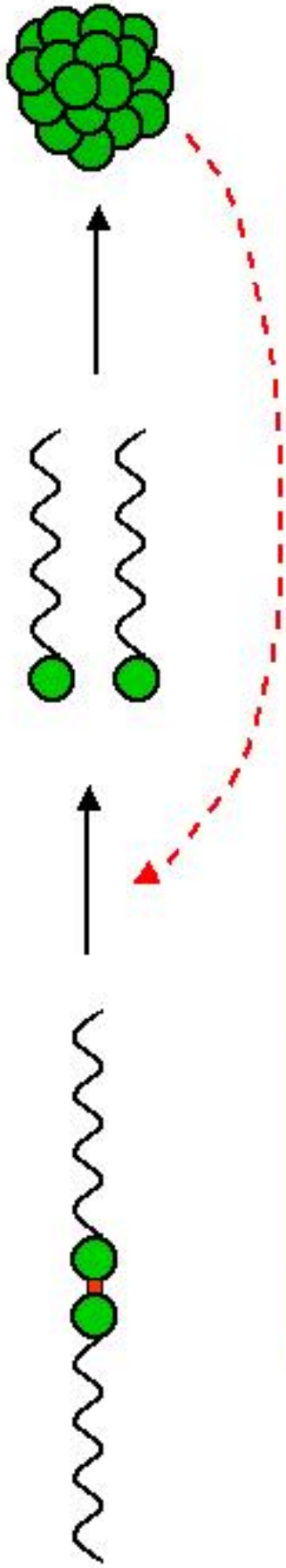
Lipid synthesis, insertion,
exchange and leaflet flipping may be catalyzed

Lipid catalysis



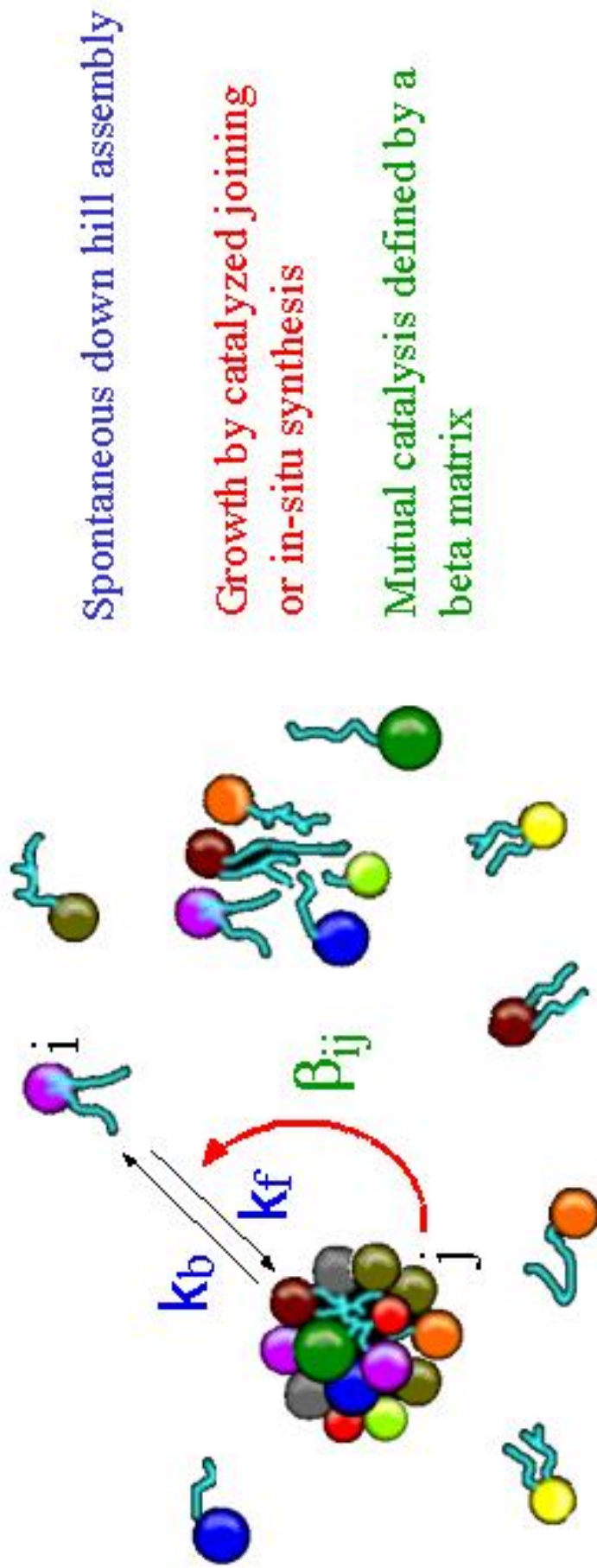
Histogram of lipid catalytic rate enhancement values (from Fendler, Membrane Mimetic Chemistry)

Lipid assembly may catalyze their own replication

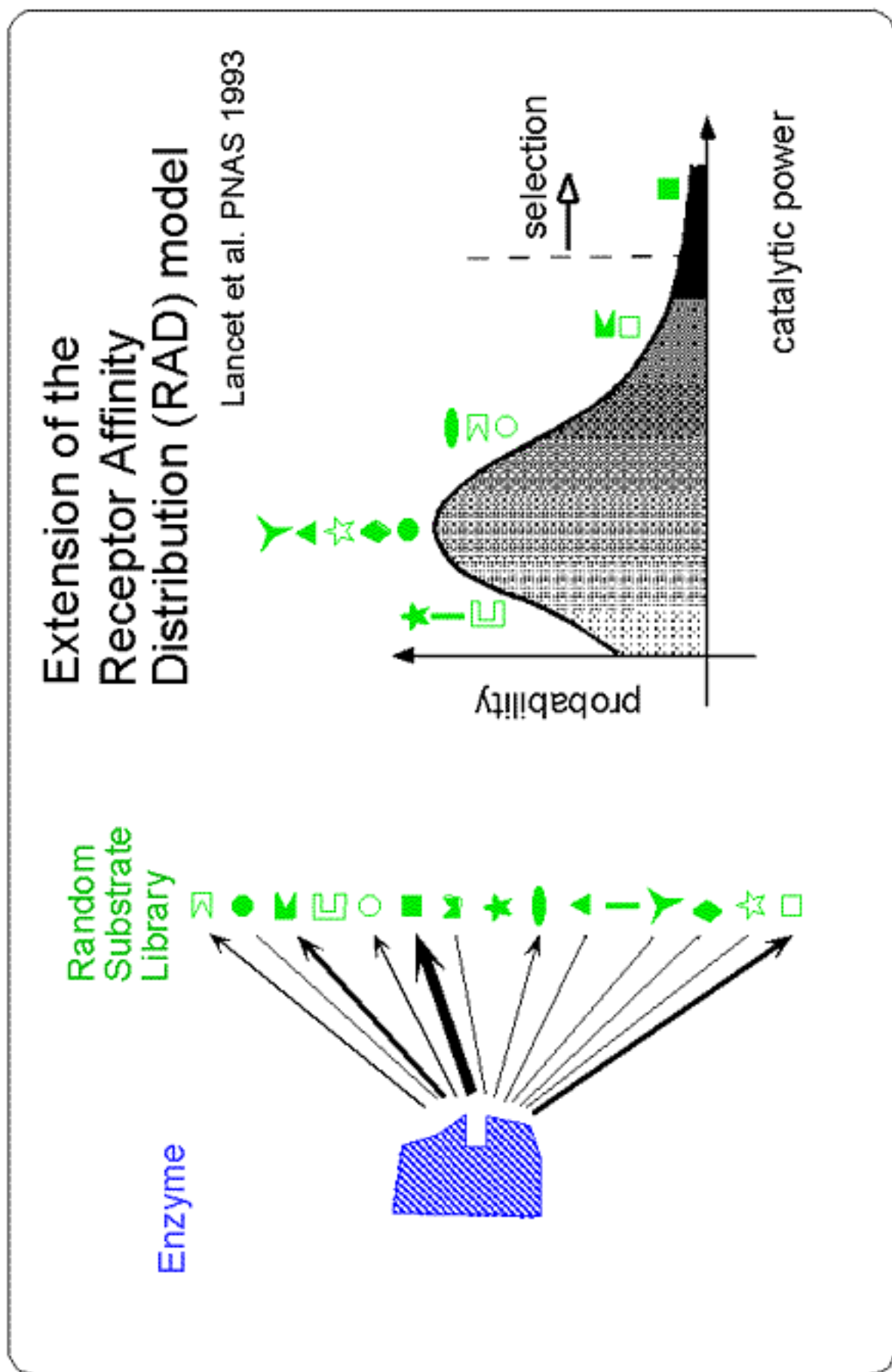


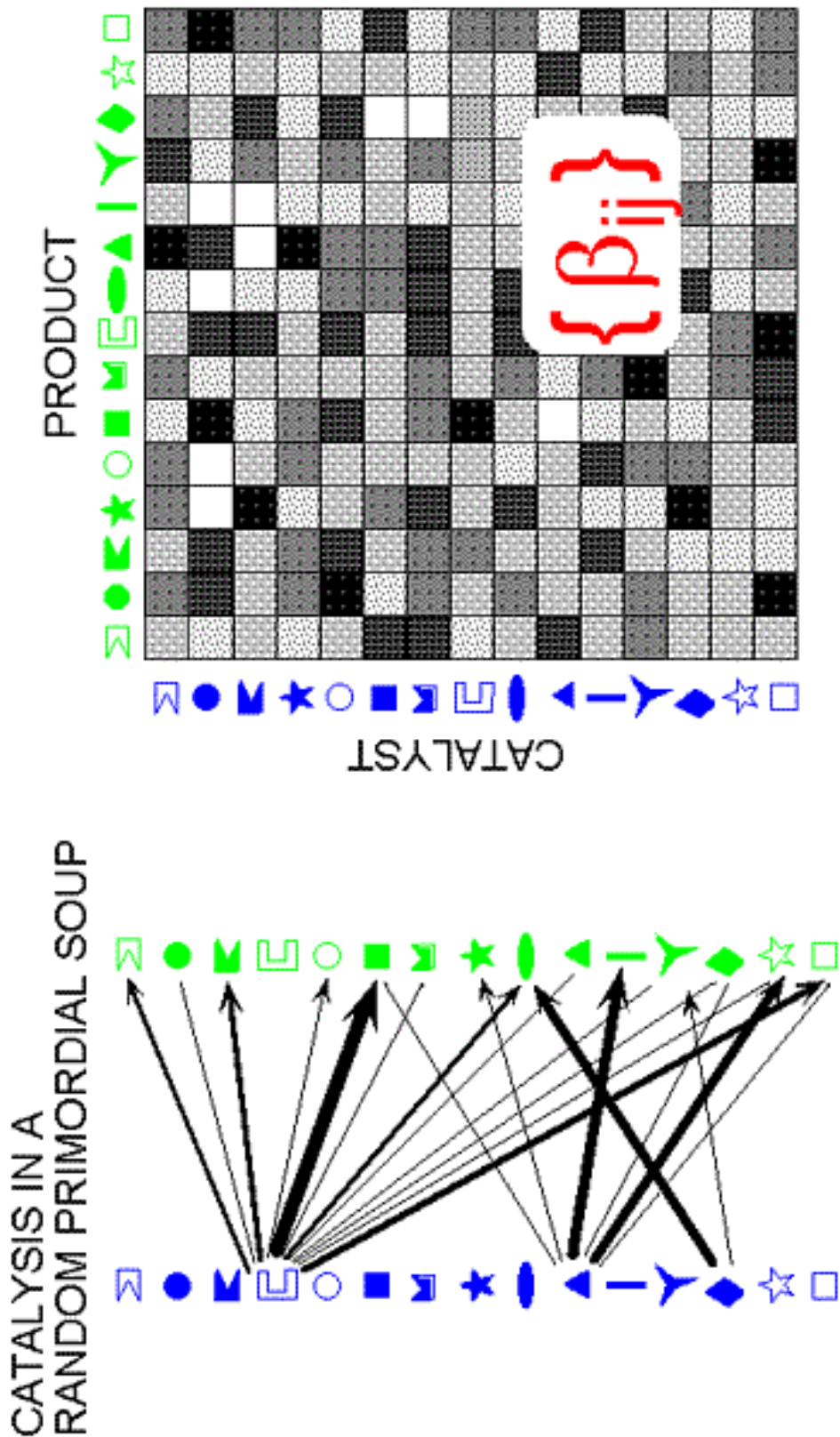
P.L. Luisi, Nature 1992

GARD: the Graded Autocatalysis Replication Domain model

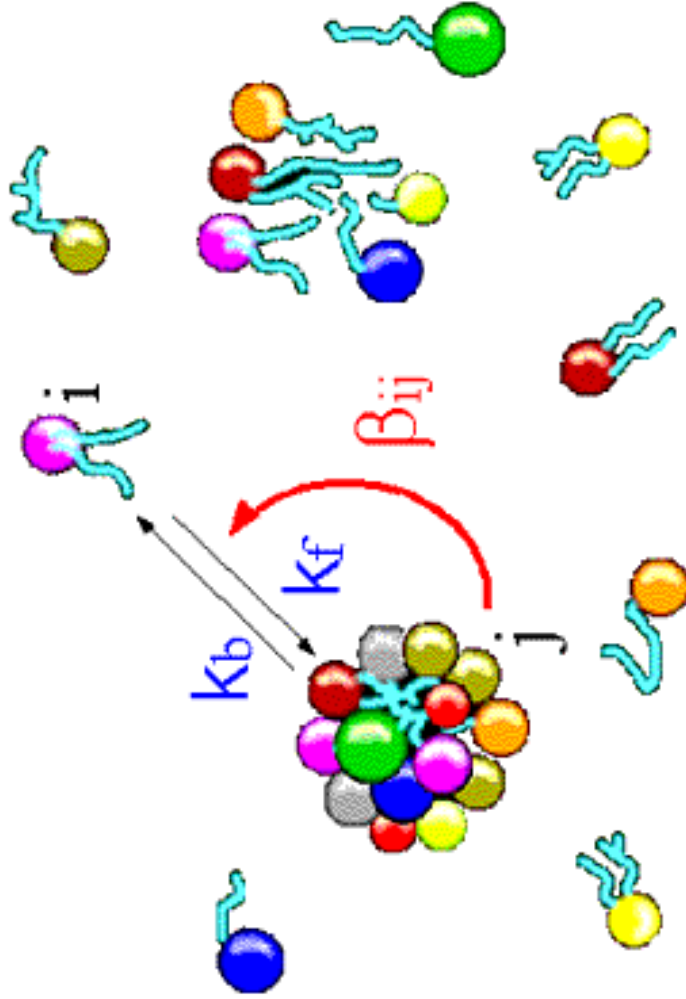


But how can one obtain the beta values for simulations?





A kinetic model for the catalyzed growth of heterogeneous noncovalent assemblies



N_G = number of
different kinds

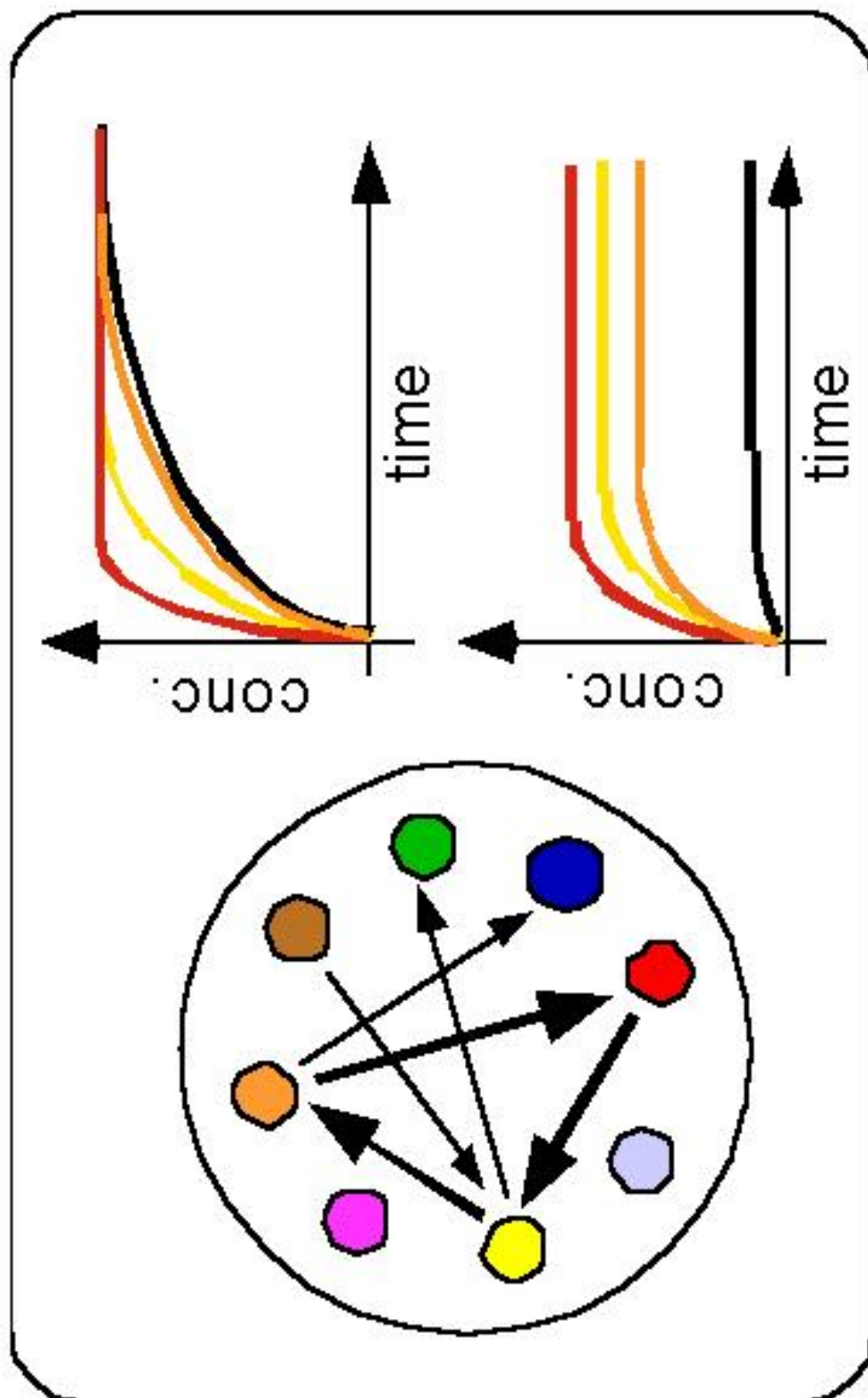
n_i = count of molecules of
kind i in the assembly

$$N = \sum n_i \quad (\text{assembly size})$$

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

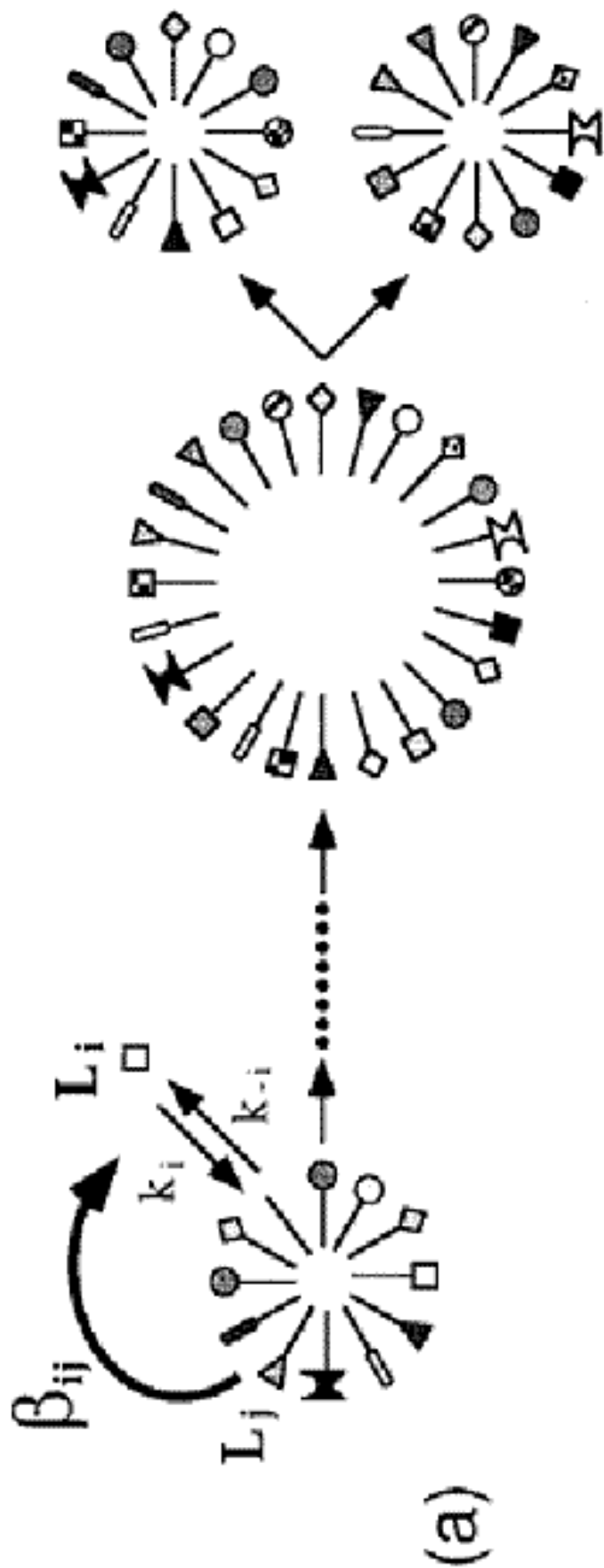
Forward reaction
Backward reaction

Catalytic
enhancement

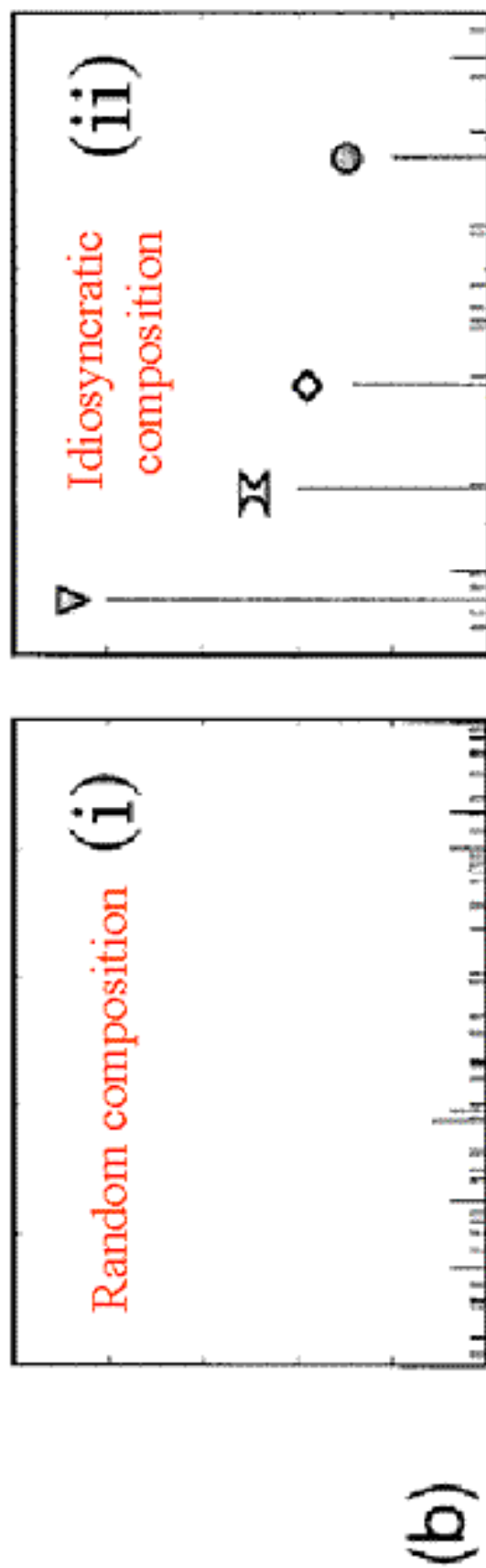


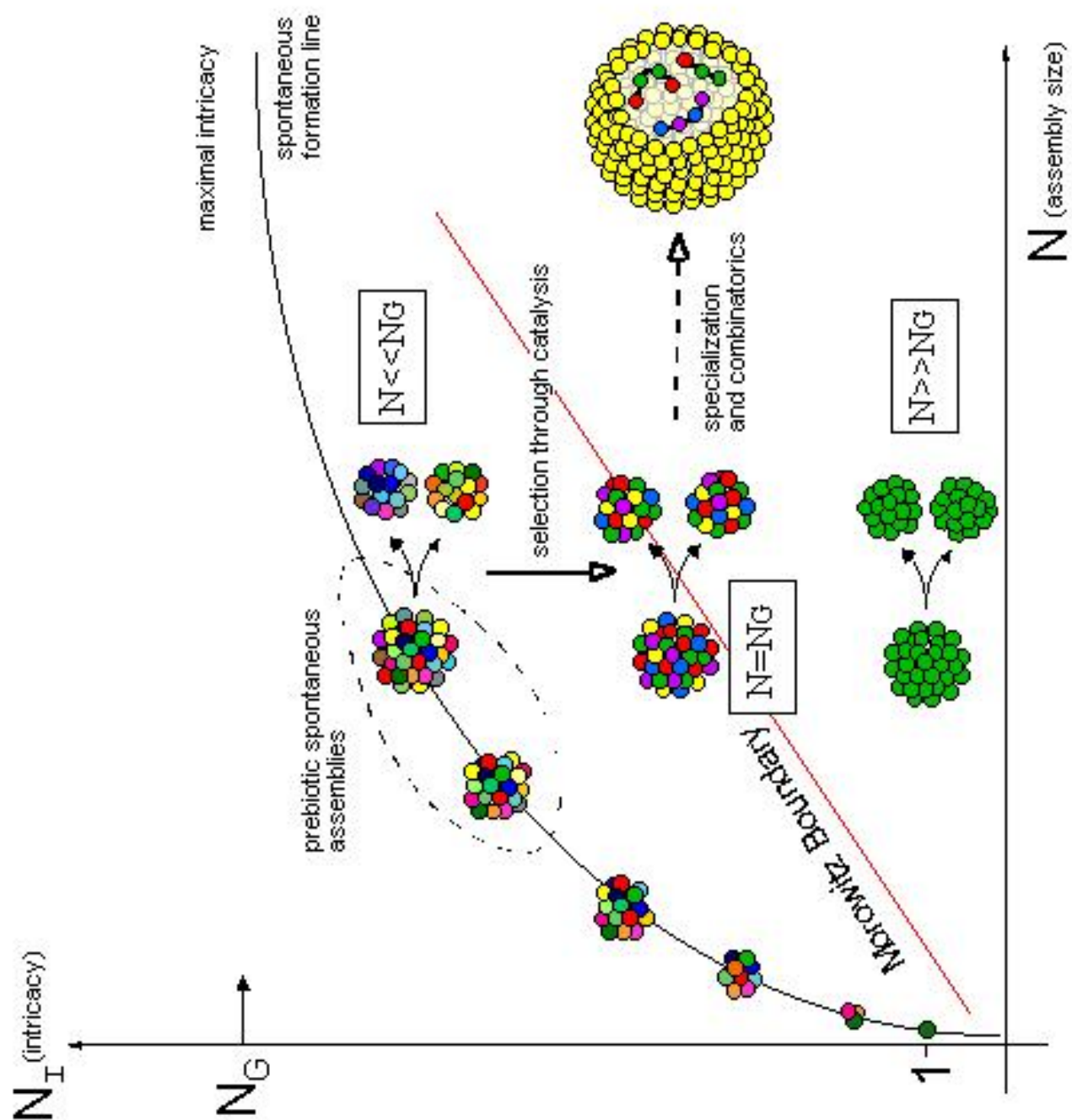
Network is established
(thicker arrows)

Lower panel – out of equilibrium

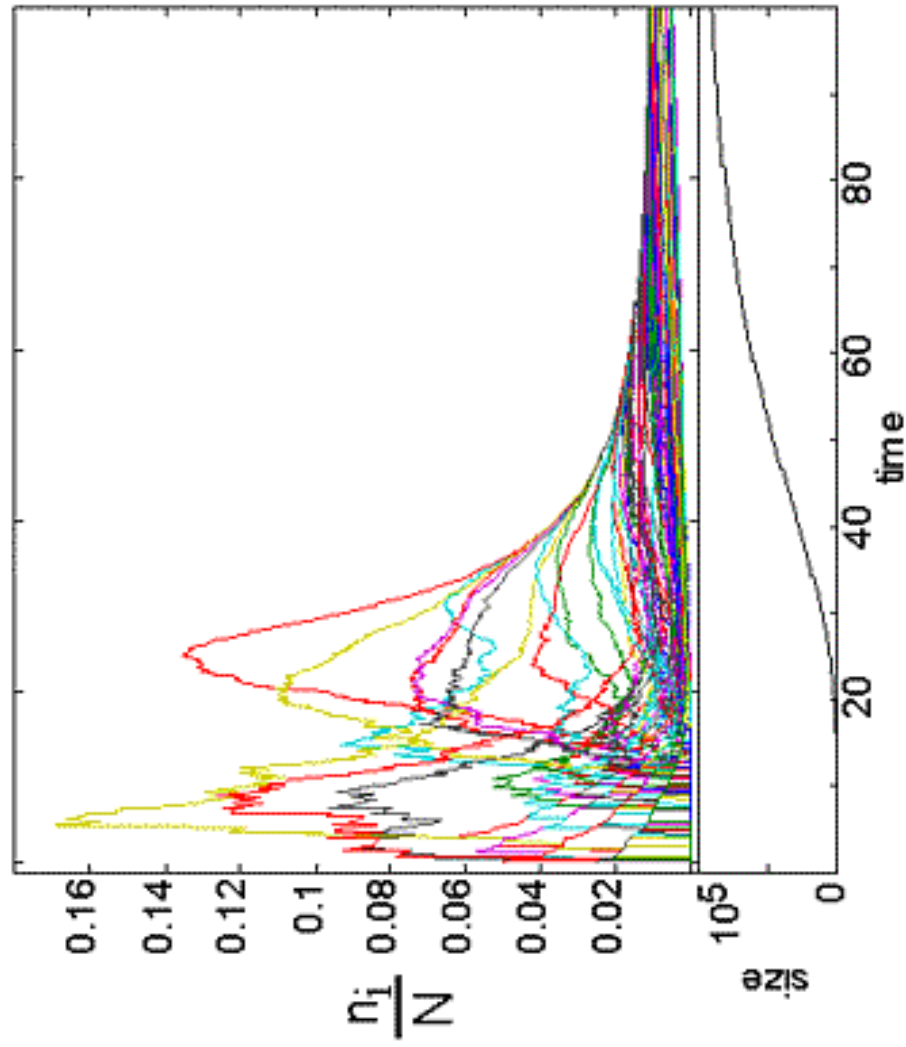


Compositional diagrams

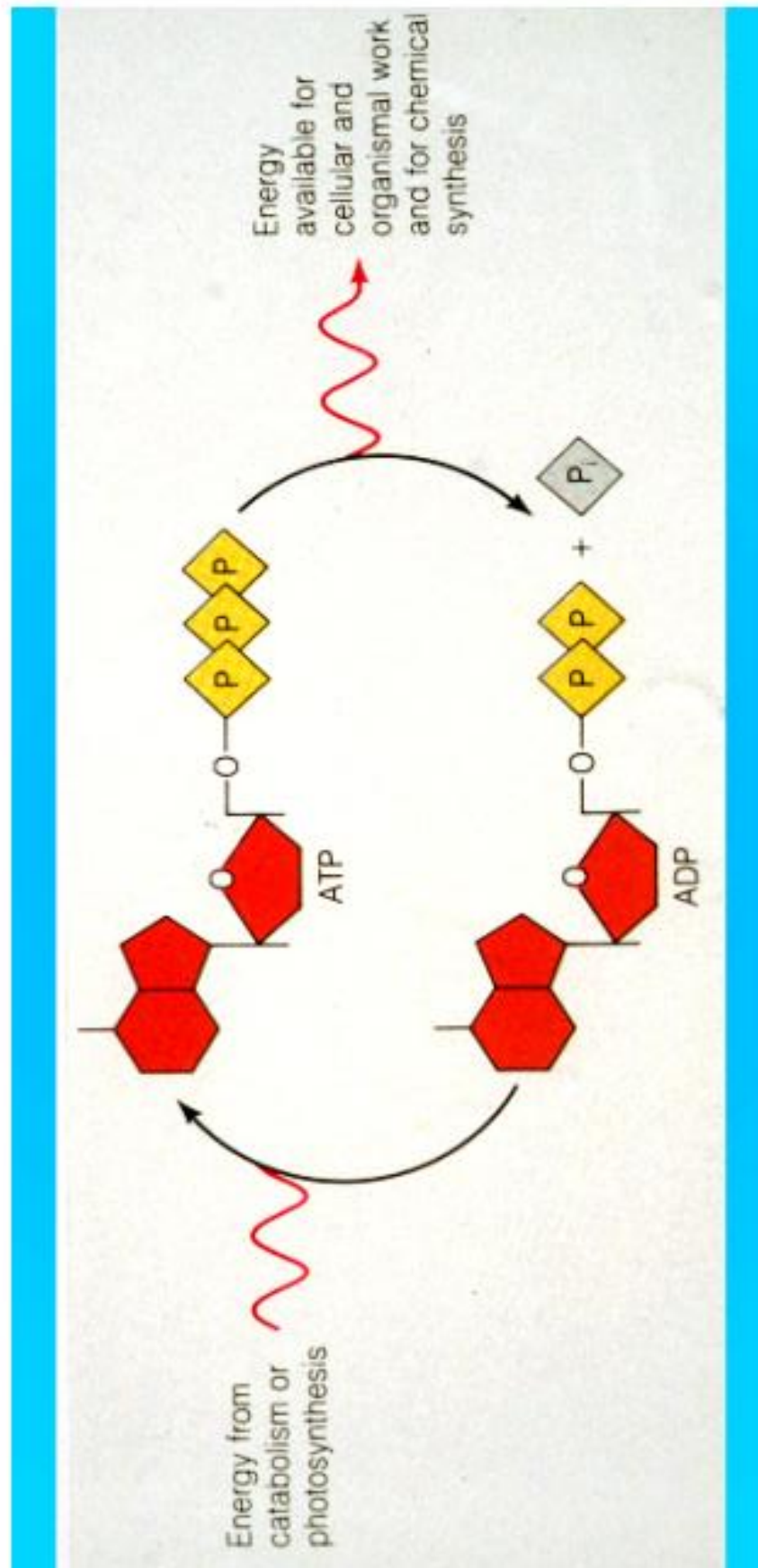




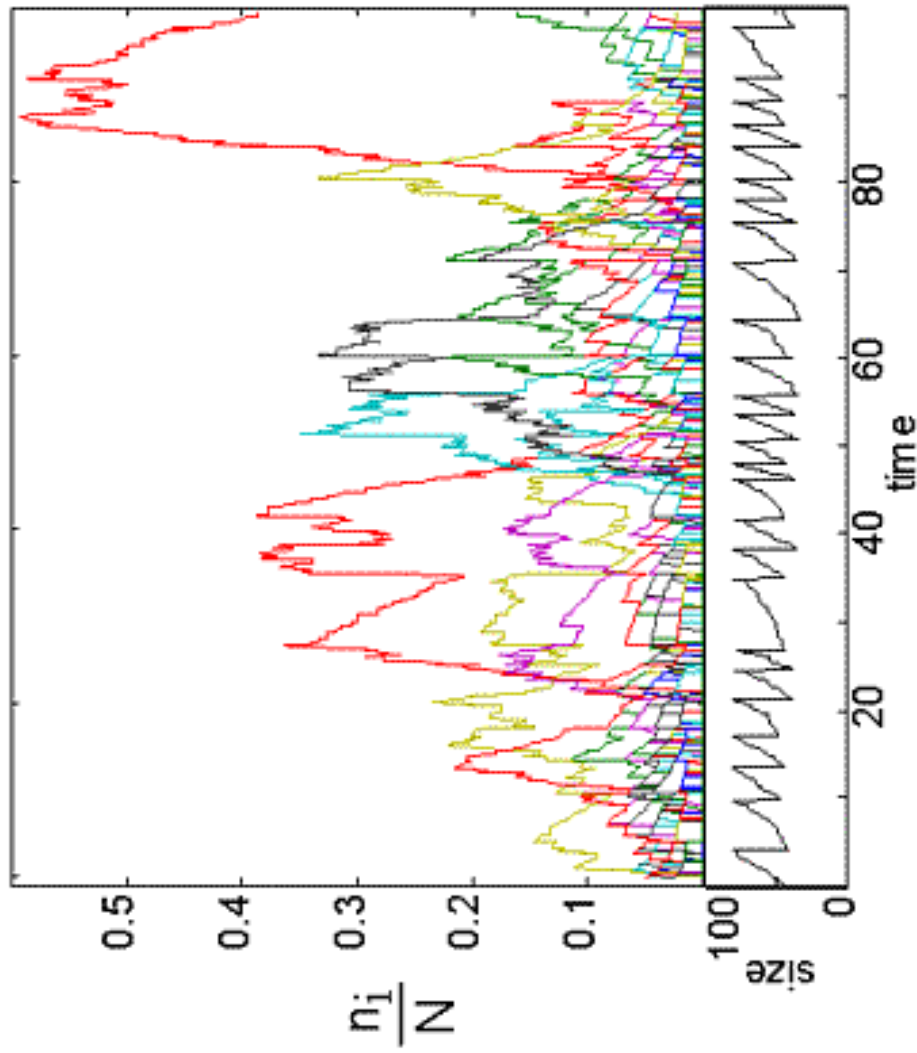
Simulated behavior (limited "food" supply):
 - complex transient
 - homogeneity at equilibrium



$N_G = 100$
 $N(t=0) = 40$
 $\Phi(\log \beta_{ij}) = \text{Gauss}(-4, 4)$



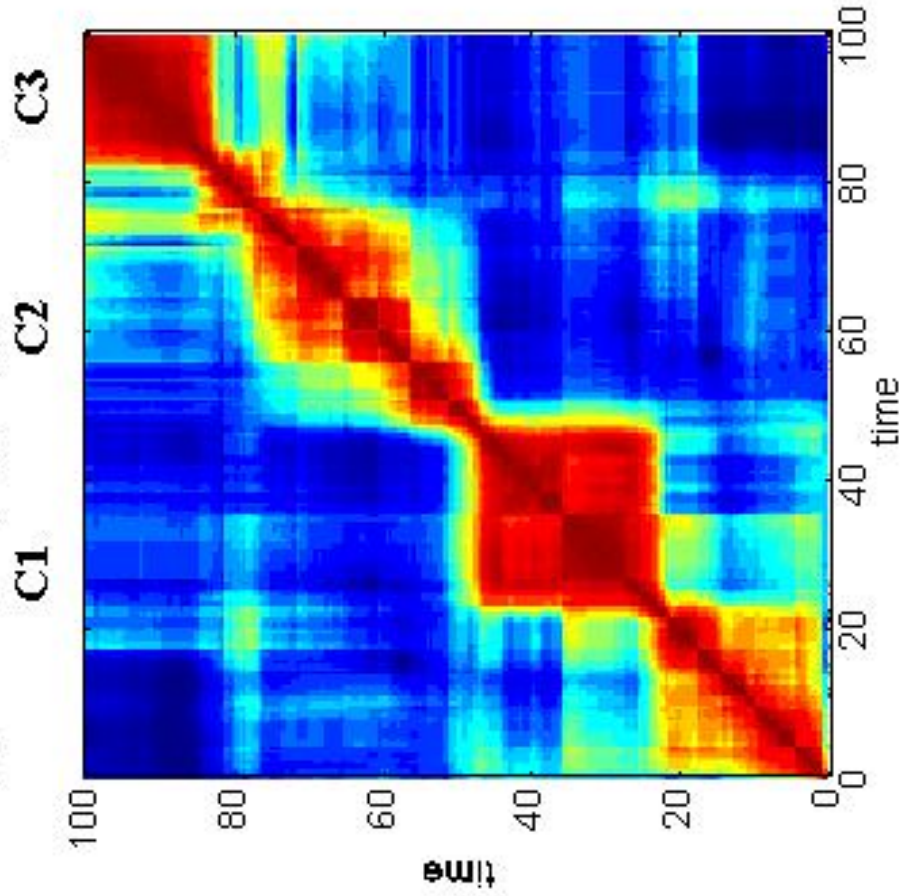
Splitting & Constant Population
 Free energy input \rightarrow far from equilibrium
 Multiple stationary states ?



$N_G = 100$
 $N_{\min} = 40$
 $\Phi(\log \beta_{ij}) = \text{Gauss}(-4, 4)$

DETECTION OF COMPOSITIONAL METASTABLE STATES (COMPOSOMES) IN COMPUTER SIMULATIONS OF THE LIPID WORLD

Metastable compositional states ("composomes") are detected by plotting $H(\mathbf{n}(t), \mathbf{n}(t'))$ for all t, t' :



$$H(\mathbf{n}, \mathbf{n}') = \frac{\mathbf{n}}{|\mathbf{n}|} \cdot \frac{\mathbf{n}'}{|\mathbf{n}'|}$$

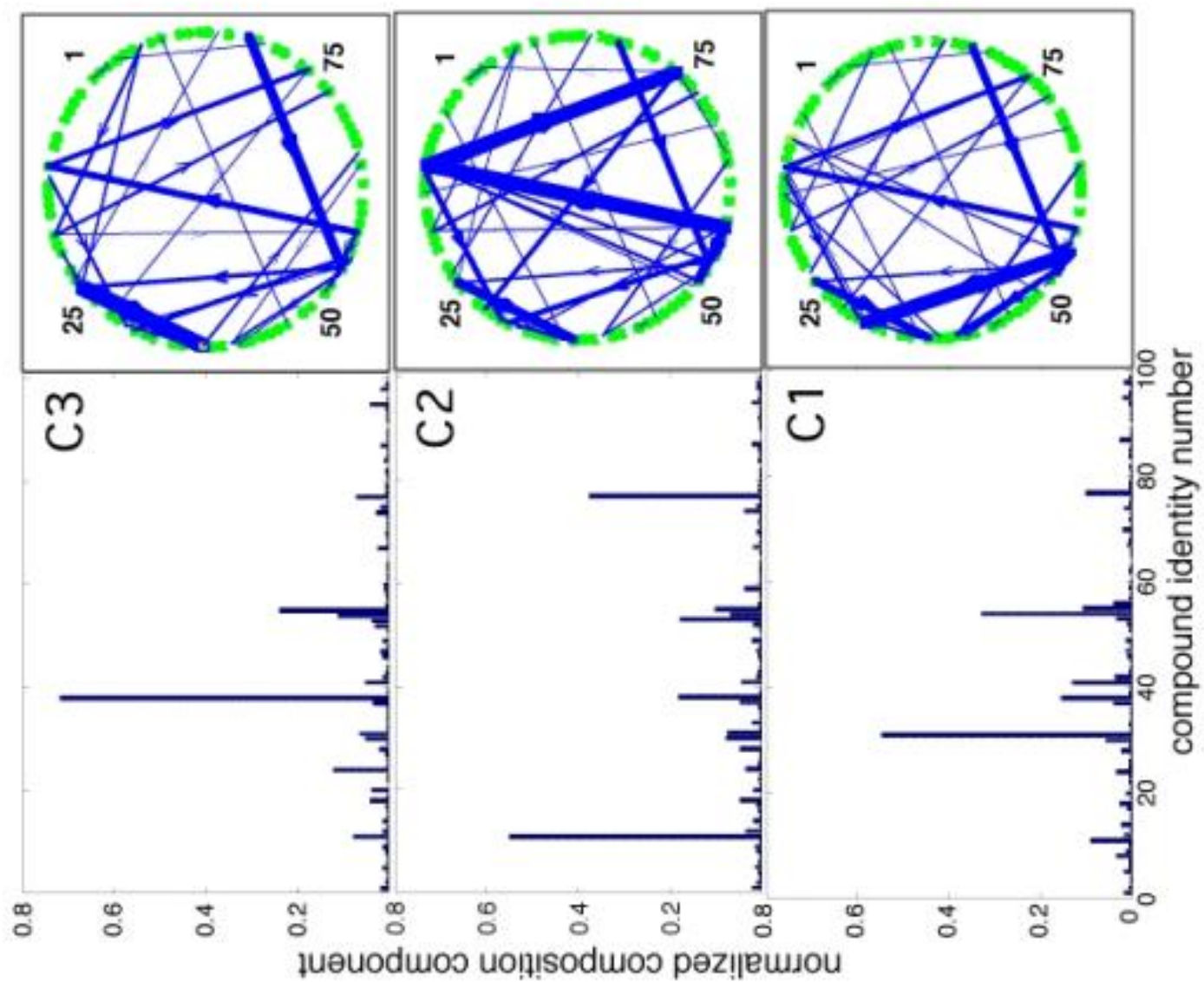


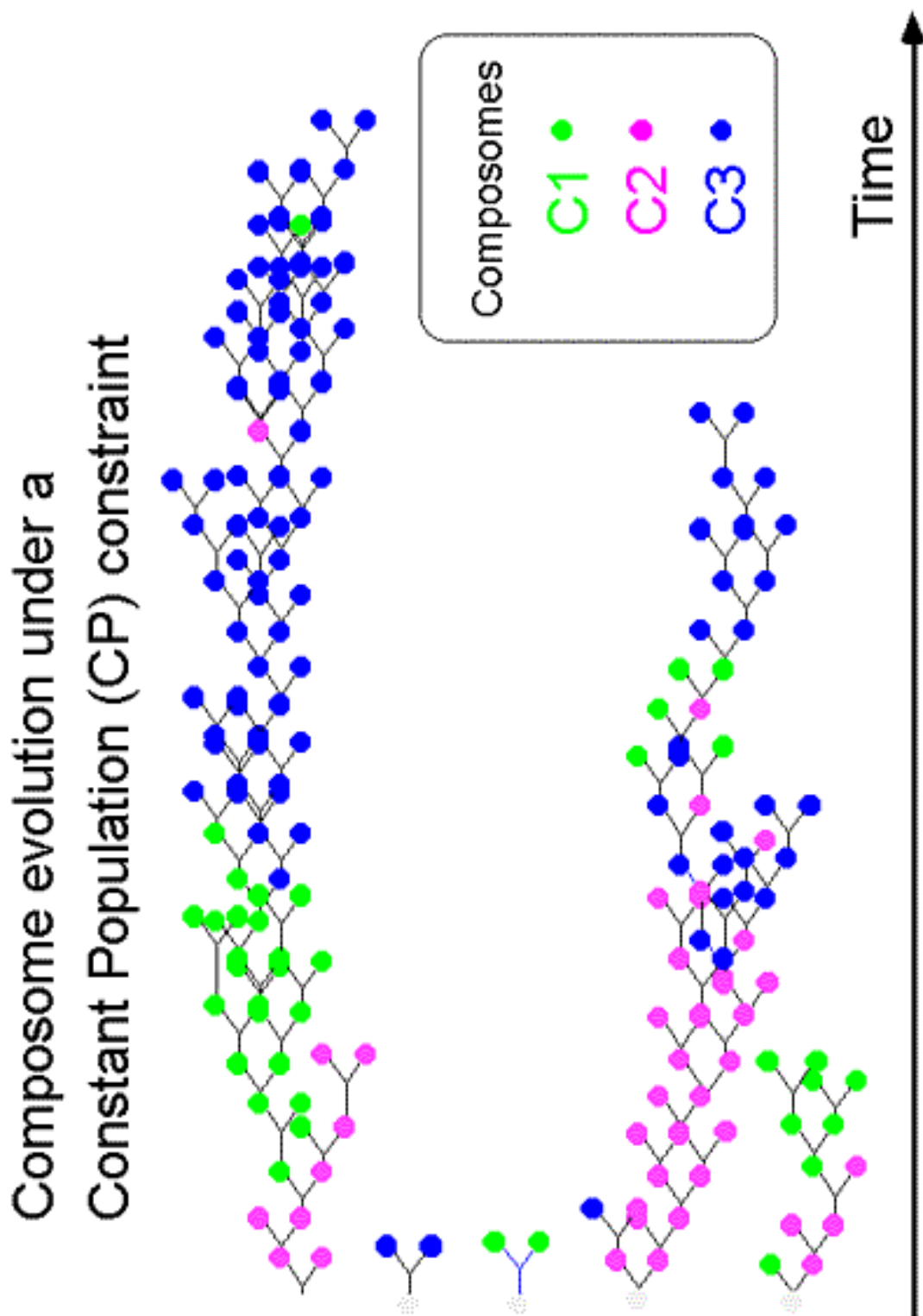
H=1



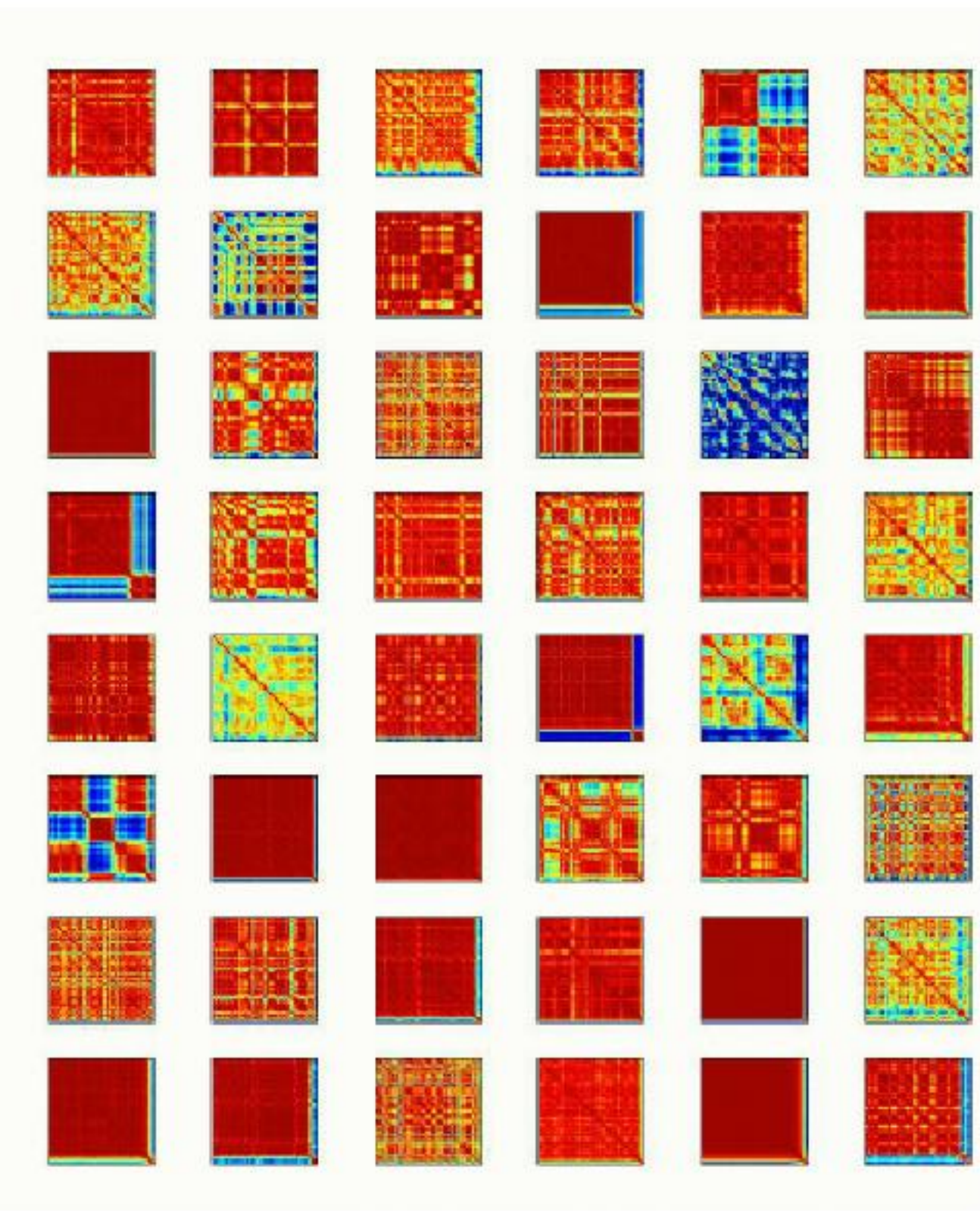
H=0

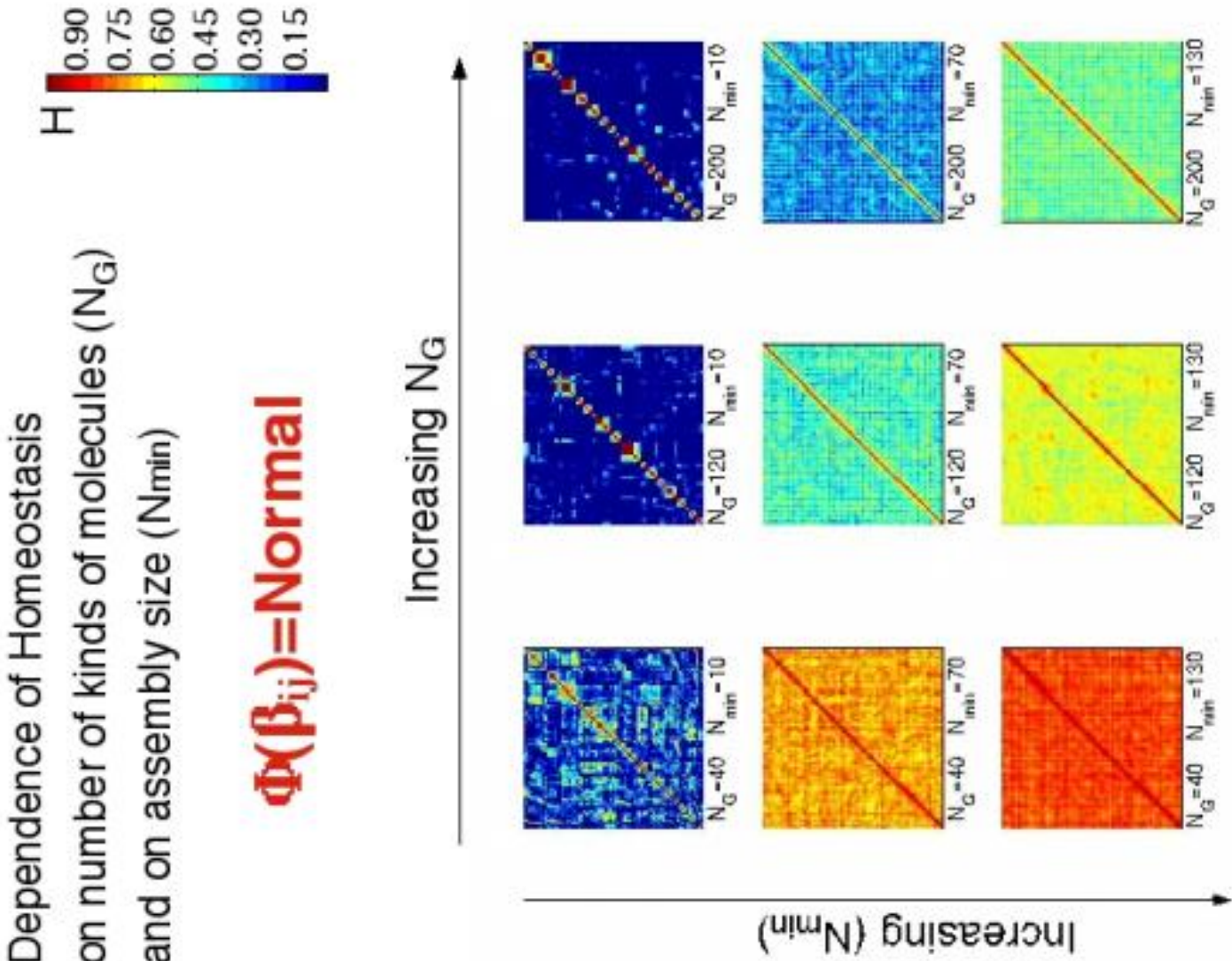
Segre, Ben-Eli and Lancet
PNAS 97:4112-7 (2000)

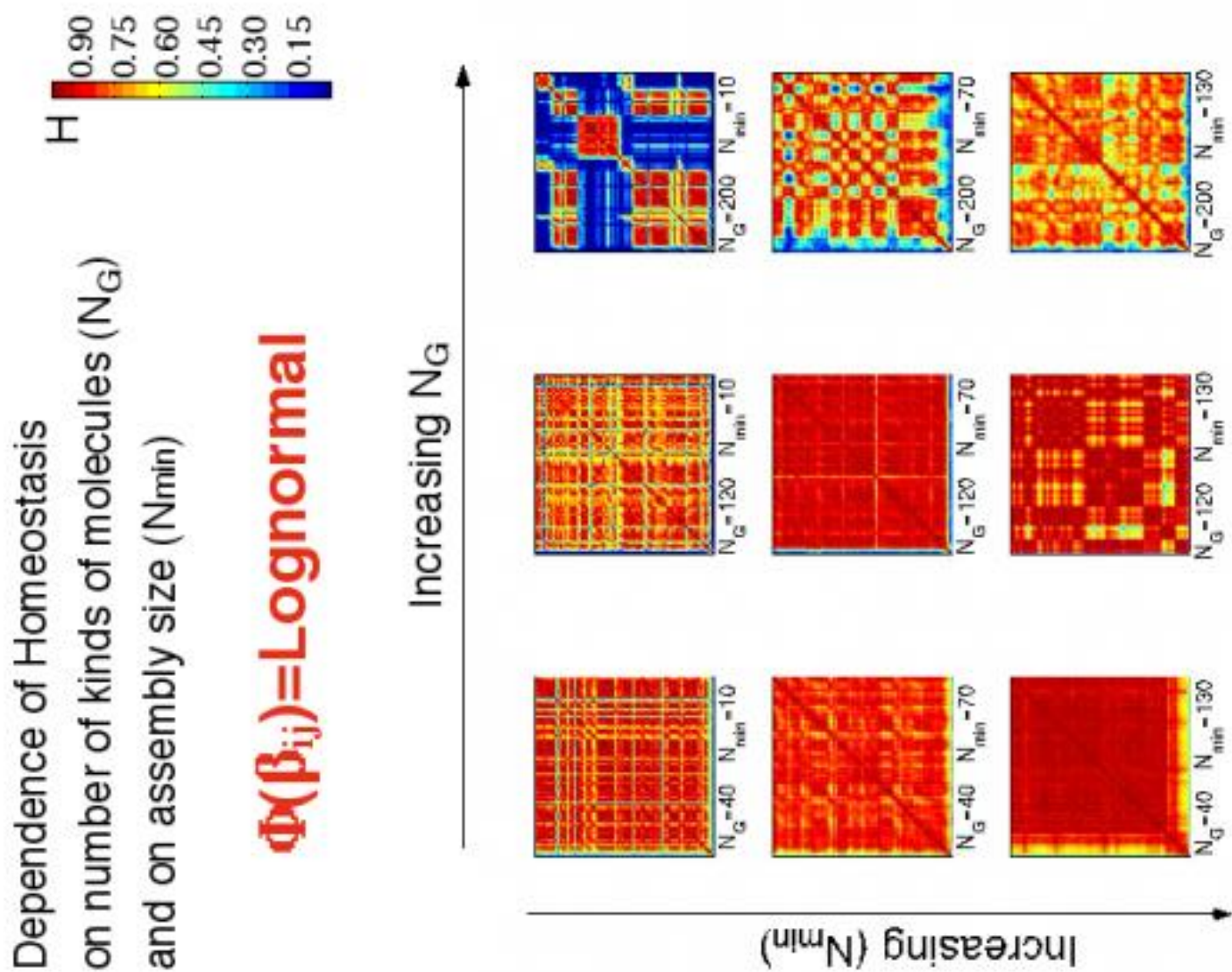


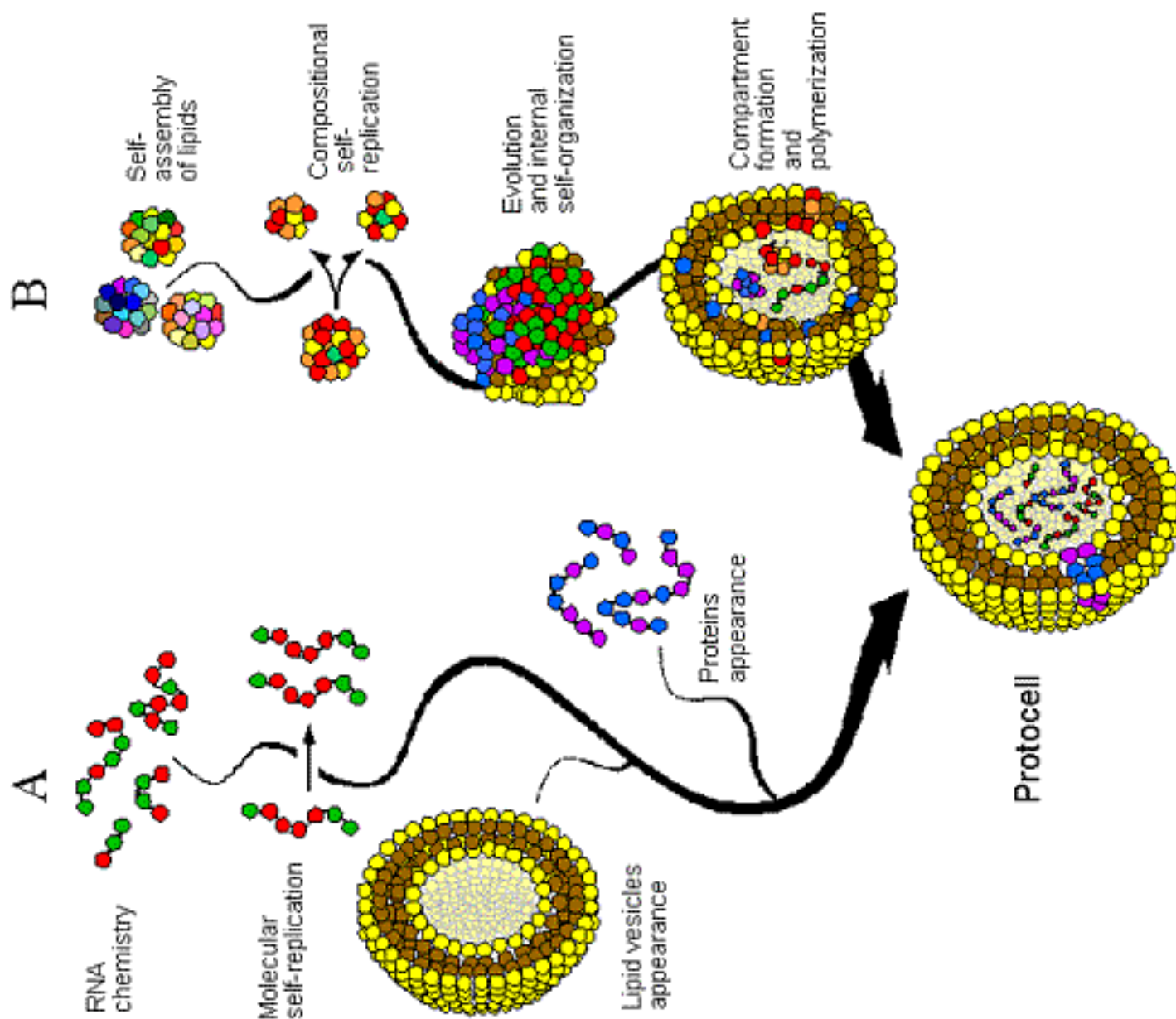


Dynamic behavior for different beta matrices (based on the same global parameters)



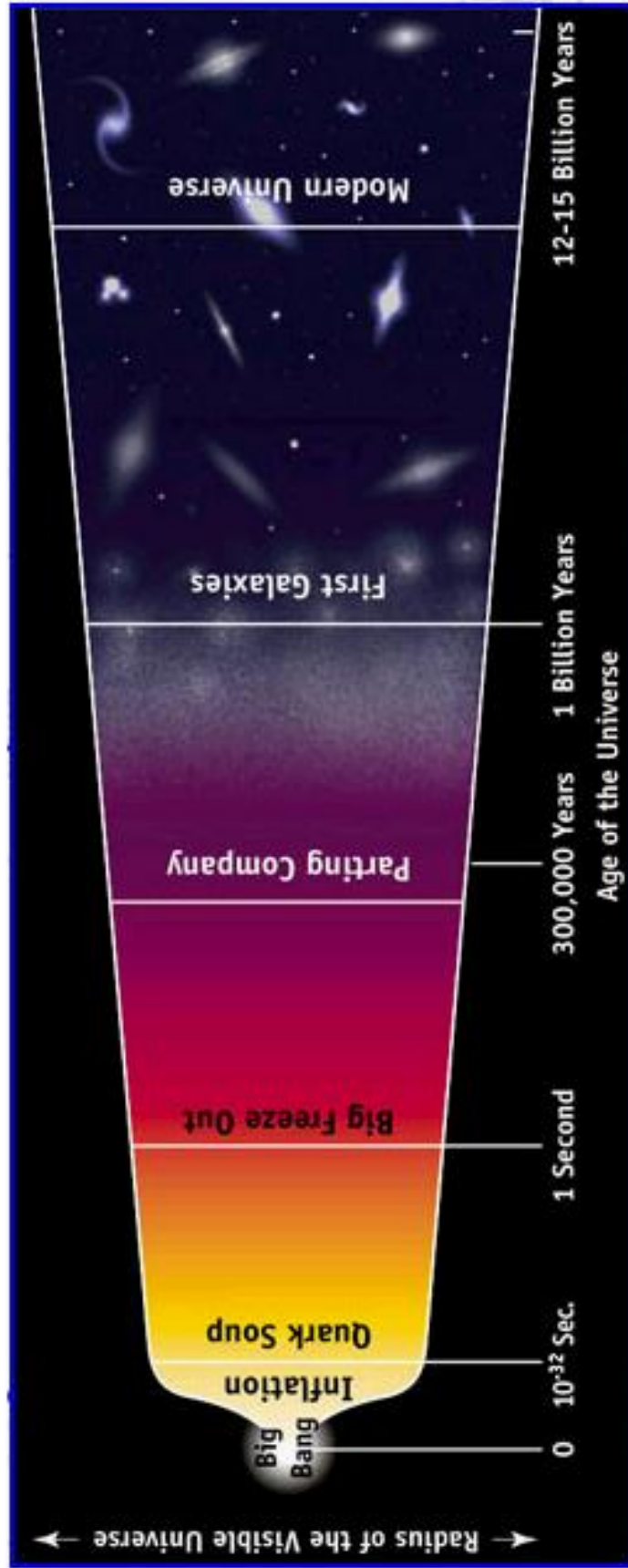






Experimental verification of the model:

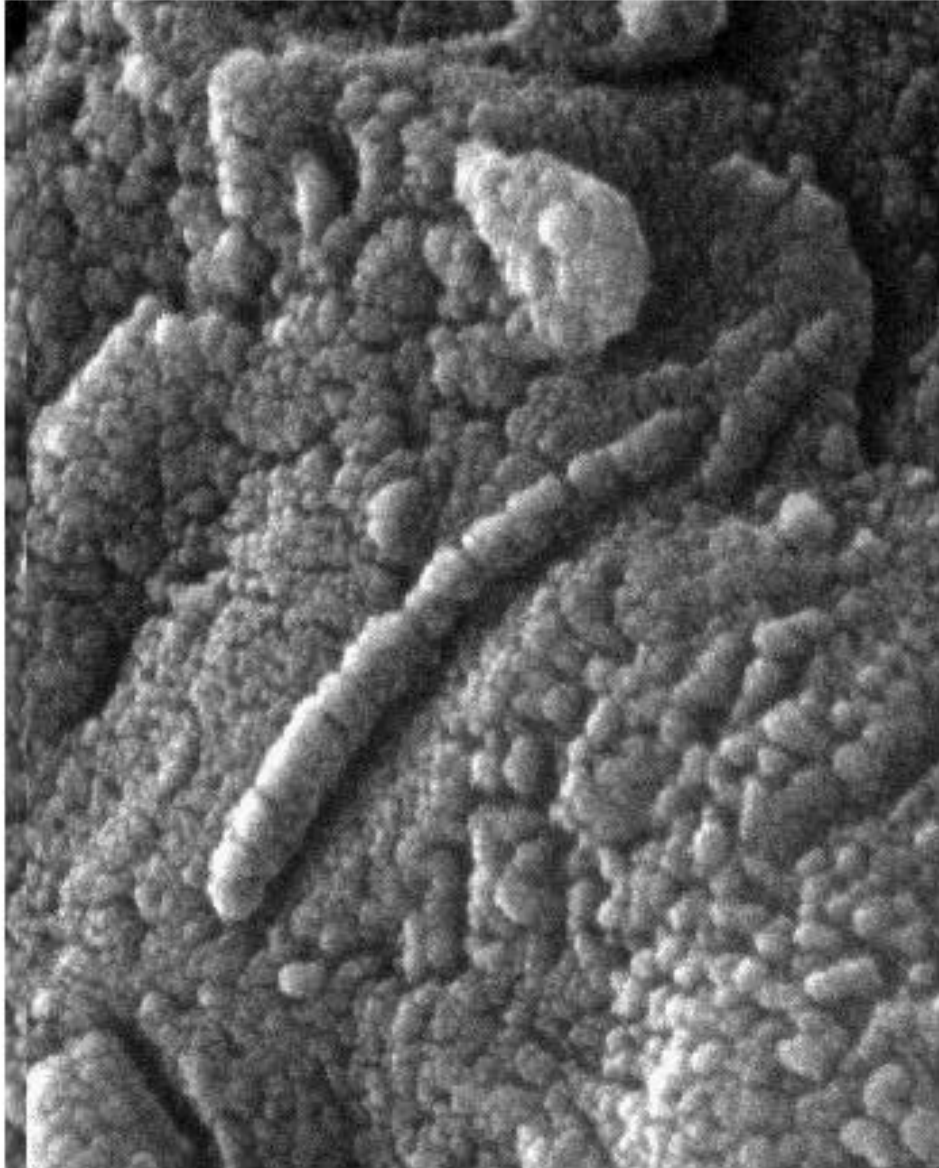
1) Large scale computer simulations. The Origin of life is a planet-scale process that took millions of years to unravel. Shouldn't we attack it with tools similar to those used to study the origin of the universe or of galaxies and suns?



2) Microanalysis of individual assemblies in a very large scale/duration experiment

3) Galactic travel - know what to expect!

GARDobes on Mars??



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Origin of Life

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