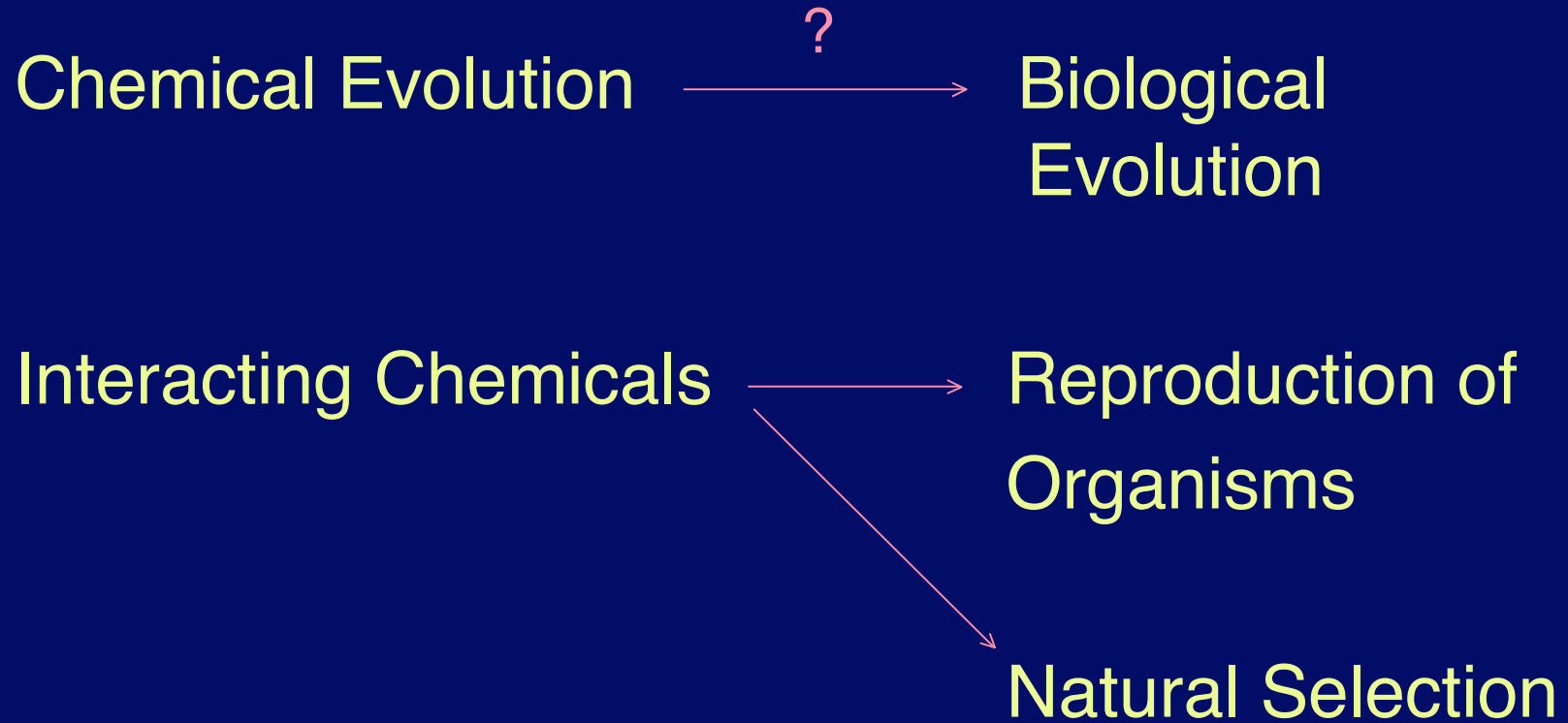


# The Transition to Life

# The Transition to Life



## Based on Simplest Life Now:

Need:

- |                                |                                       |
|--------------------------------|---------------------------------------|
| 1. Nucleic Acids               | Replicable Information                |
| 2. Proteins                    | Enzymes (Catalysts)                   |
| 3. Lipids                      | Membranes (Enclosure)                 |
| 4. Carbohydrates<br>(Pigments) | Energy Storage<br>(Energy Conversion) |

Too much to ask of chemical evolution

□ Protolife?

# Protolife

1. “Virus” Free living but equivalent in complexity

Protein + Nucleic Acid + Supply by Environment



Genetic Code

2. Protein Protolife

Protein  $\longrightarrow$  Self Replication?

3. Nucleic Acid Protolife

RNA  $\longrightarrow$  Self Catalysis?

## 4. Something Else

Minerals

Clay Layers

Mineral - Molecule

Pyrite

Thioesters

Genetic Takeover

? → RNA → DNA

# Protein-Based Protolife

## 1. Proteinoid microspheres - Sidney Fox



Protocells

Protolife?

Can Add Proteinoid

Split

Bud

Form Chains

(Look like life)

Grow

Divide } "Reproduce"

Bud }

Like Bacteria

But "Reproduction" not exact

Later incorporate Nucleic Acids

**Proteinoid**  $\longrightarrow$  **Cells**  $\longrightarrow$  **Genes**

Problem: How to incorporate Nucleic acids?

## Nucleic Acid Based Protolife

RNA → Genes → Protein → Cells

Self-replicating RNA molecules

Experiment by Sol Spiegelman

RNA from Q<sub>β</sub> Virus - parasite on bacteria

Injects RNA - Bacterium makes replicase

↑  
Enzyme to Replicate RNA

RNA multiplies, using activated nucleotides in

Bacterium → new viruses

In Test Tube: Template RNA, Replicase,  
Activated Nucleotides (ATP, CTP, GTP, UTP)

□ RNA copied without machinery of cell

Variation: No template RNA

Replicase made RNA from nucleotides

↑  
Protein



Manfred Eigen - further experiments with RNA  
in test tube:

Mutant RNA strands compete

Degrade to smallest (~ 200 nucleotides)

RNA that replicase could recognize

(Monster - Selfish RNA)

RNA can do self-catalysis in some cases

Could this have led to self replication?

## Eigen scenario

1. A replicating RNA molecule forms by chance (random replicator - not a gene)  
ribozyme (catalyst, made of RNA)
2. Family of **similar** RNA's develops (quasispecies)
3. Connection to proteins  
(quasispecies specialize to make parts of protein)

4. Complex interactions (hypercycles)
5. Use lipids to make protocells
6. Competition leads to biological evolution

EXON1 INTRON EXON2

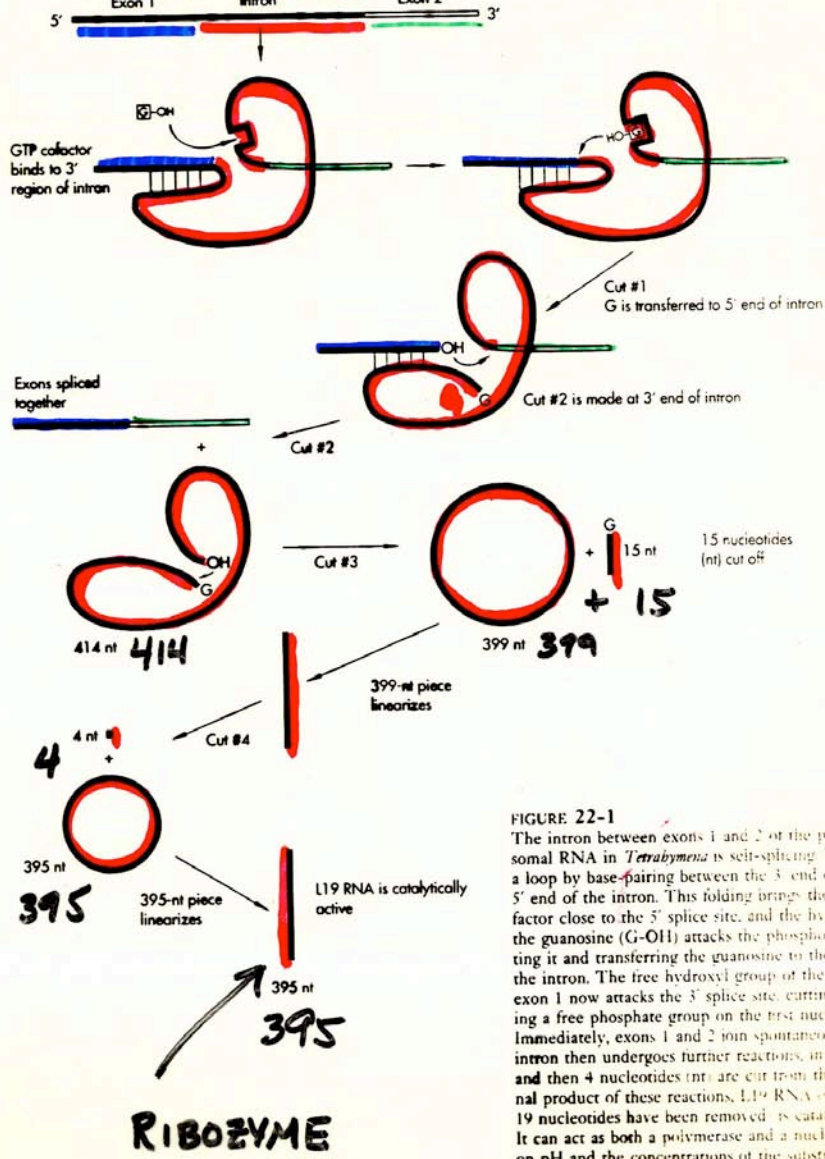


FIGURE 22-1

The intron between exons 1 and 2 of the pre-mRNA in *Tetrahymena* is self-splicing. The lariat is formed by base-pairing between the 3' end of the intron and the 5' splice site. This folding brings the guanosine (G-OH) close to the 5' splice site, and the hydroxyl group of the guanosine attacks the phosphate group of the 5' splice site, forming a phosphodiester bond and transferring the guanosine to the 5' end of the intron. The free hydroxyl group of the last exon 1 now attacks the 3' splice site, cutting it off and leaving a free phosphate group on the first nucleotide. Immediately, exons 1 and 2 join spontaneously. The intron then undergoes further reactions, in which 15 nucleotides are cut from the 5' end and 4 nucleotides are cut from the 3' end. The final product of these reactions, L19 RNA (395 nucleotides), is catalytically active. It can act as both a polymerase and a nuclease. The activity is dependent on pH and the concentrations of the substrates.

RIBOZYME

HOW AN "INTRON" (NOT A GENE) CAN CUT, SPLICE & BECOME A CATALYST

# Problems with Nucleic Acid First Scenario

1. Hard to get monomers
2. Unlikely to link correctly
3. Need existing proteins and lipids
4. Hypercycles subject to instabilities

N = size of molecular population

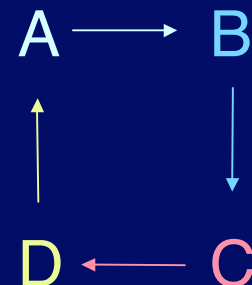
If N small

Population Collapse

If N large

Selfish RNA

Short Circuit



If B → D Short Circuit

□ Only narrow range of sizes works

# The Origin of the Genetic Code

- We need more than **either** protein or RNA protolife
- Need interaction via genetic code
- Need **translation**
- Consider first a scenario by R. Shapiro

# Shapiro's Fable

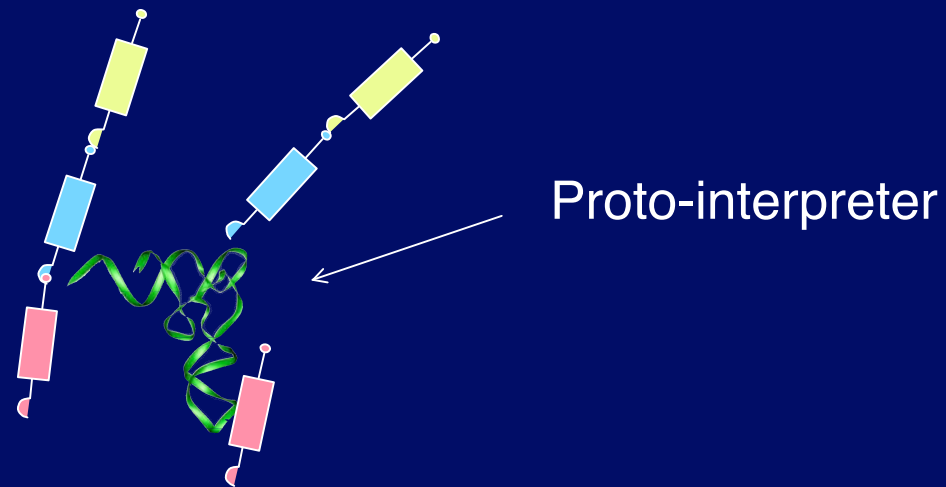
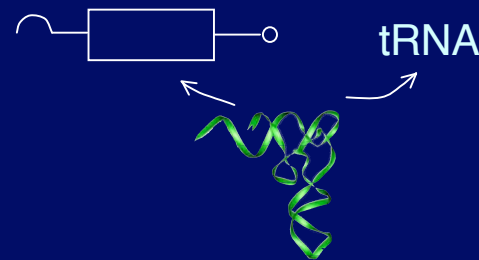
The case for the “chicken”

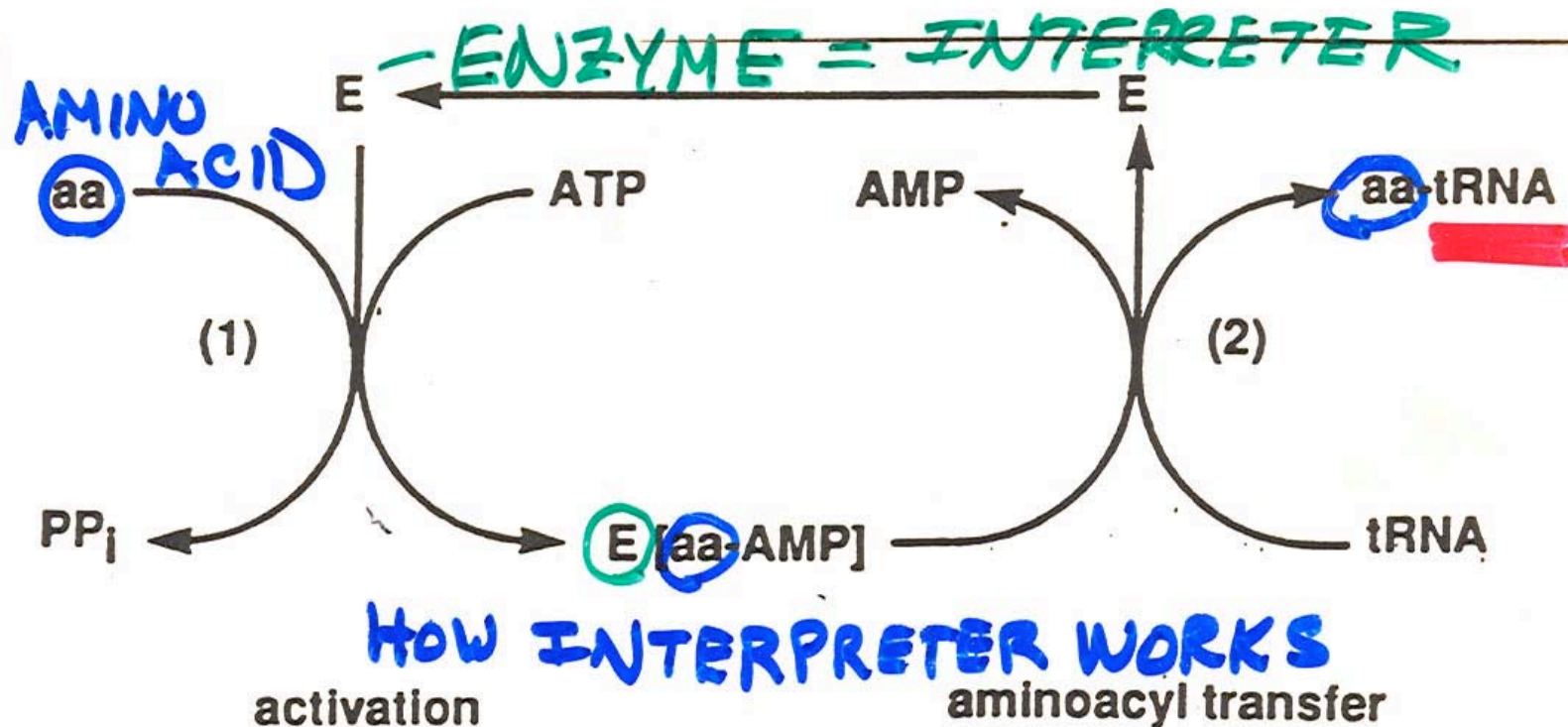
Protein first □ replication problem

“interpreters” aminoacyl tRNA synthetases

Match tRNA &  
Amino acids

Could an earlier version have copied proteins directly?





amino acid recognition

tRNA recognition

Fig. 1 The two steps of the reaction catalysed by aminoacyl-tRNA synthetases. The tRNA is recognized in the second step (through the features designated paracodon in this article) by the enzyme carrying a bound aminoacyl-AMP intermediate. Participation of the aminoacyl group in the recognition process is thus an attractive possibility.



1. Early Evolution: Start with 4-6 amino acid types, gradually add more  
enzymes increase in size and catalytic power
2. First use of phosphate as energy? (ATP)  
or sugar-phosphate chains for construction  
(Teichoic acids in membranes of some bacteria)  
(partial Q<sub>1</sub> replicase)
3. Bases added for structure  
Support for protein synthesis → ribosome

4. Begin to copy RNA (Full Q<sub>1</sub> replicase)  
Natural selection → better ribosome
5. Specialized, Short RNA aided attachment of amino acids to proteins; became tRNA
6. Then mRNA - to align tRNA's  
now a separate genetic system that evolves
7. DNA developed from RNA

Shapiro dates last step to prokaryote -eukaryote split (different ways of storing DNA info)

### Tests:

1. Synthesize in lab? Not possible yet.
2. Molecular archaeology - vestigial ability of interpreters to recognize amino acids in proteins
3. Survivors of protein era? prions?

## Support for the “chicken”

1. 1988 discovery that interpreter does not use tRNA codon to recognize correct tRNA (in some cases)  $\sim 1/2$ 
  - instead a single base pair at the other end of tRNA
    - simpler, older code  
second genetic code
    - connection of interpreter and tRNA  
more primitive than current code

## 2. Dyson modeling of molecular “populations”

Transition from disorder to order

(non-life)      (life)

Finds number of monomer types likely to be  
9 - 11 (ok if used ~ 1/2 of modern proteins)

But nucleotides (only 4) - not enough

Favors protein first

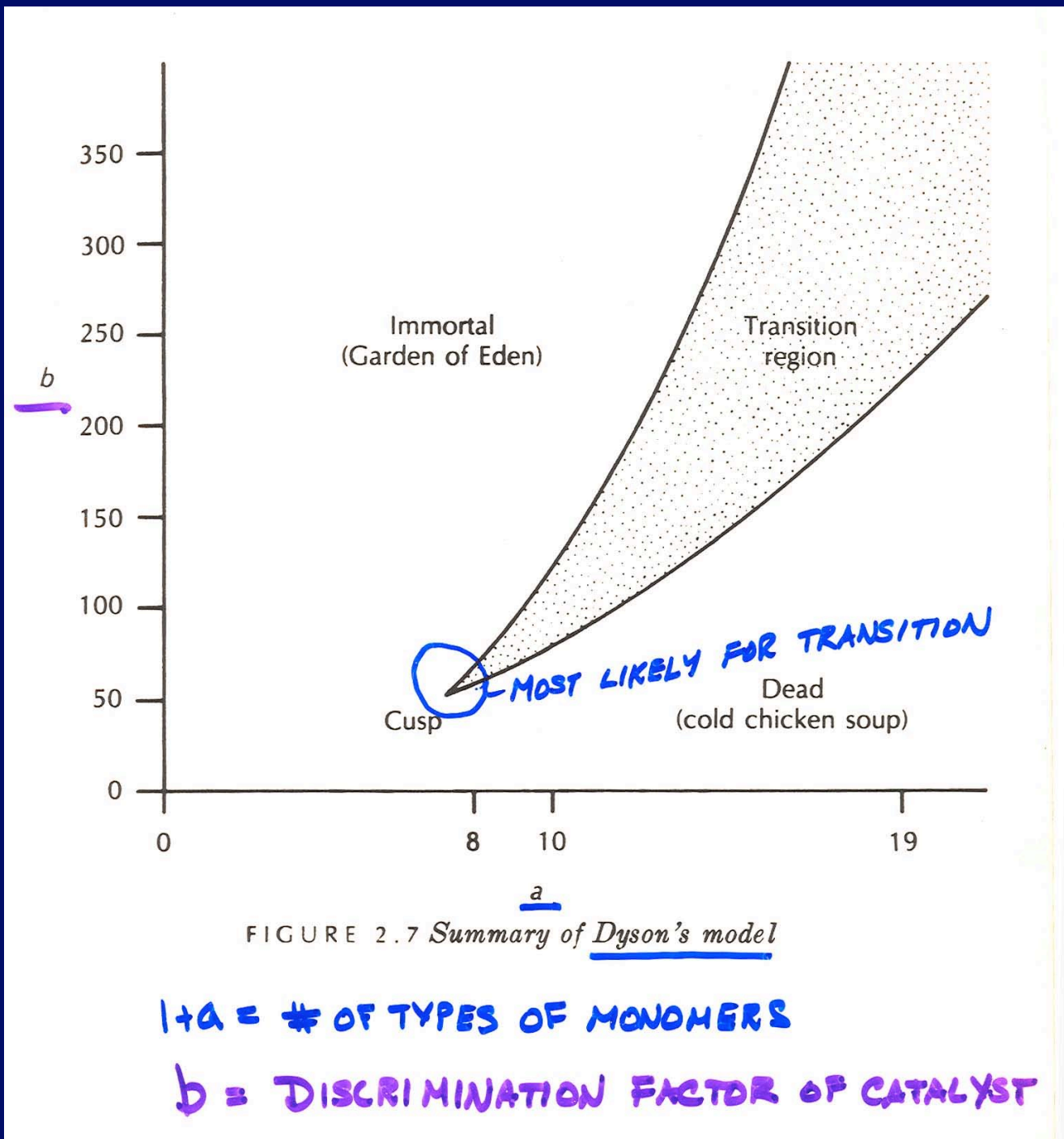


FIGURE 2.7 Summary of Dyson's model

$1+a$  = # OF TYPES OF MONOMERS

$b$  = DISCRIMINATION FACTOR OF CATALYST

## The Egg Strikes Back

Recent work shows some RNA can catalyze  
Non-RNA reactions

1. RNA in ribosome appears to be what catalyzes peptide bond formation

Noller, et al. 1992, *Science*, **256**, 1416

2. RNA “ribozyme” catalyzes reactions between amino acids and tRNAs

—> First “interpreter” may have been RNA

Piccirilli, et al. 1992, *Science*, **256**, 1420

# Origin of the Genetic Code

Crucial step in any theory

Allows communication

Nucleic Acids  $\longleftrightarrow$  Proteins

Early versions probably coded fewer amino acids - less specific

G - C more stable



Some evidence for RNY

Purine      Either  
|      /  
RNY  
|      \  
Pyrimidine

□	4 codons	GGC	glycine	} Common in Miller Urey and Meteorite
		GCC	alanine	
		GAC	aspartic acid	
		GUC	valine	

Others added later

# Evolution of Genetic Code

Gaining specificity

If early tRNAs carries more than 1 kind of amino acid

e.g.

Glycine or alanine



CGG  
GCC  
-----  
mRNA

Mutation  
→

Glycine



CCG  
GGC  
-----  
mRNA

Evidence that code has evolved  
Freeland, et al. Tested  $10^6$  other codes

Only one better at minimizing bad effects of mutations

□ Natural Selection

**Still Evolving**

Some organisms have slightly different codes in mitochondria or in nucleus

## Other Ideas

- Neither the chicken nor the egg came first
- Transitional forms that were later discarded

## Or was it the “egkin”?

Recent experiments with peptide nucleic acid (PNA).

PNA: Peptide backbone with bases

Can act as template for polymerization of RNA  
From activated nucleotides

(Böhler, et al., *Nature*, **376**, 578  
& comments by Piccirilli, pg. 548 } 17 Aug. 1995

PNA could be simpler to form under prebiotic conditions  
Main point is that a simpler thing (not necessarily PNA)  
could have preceded RNA

# Membranes

- Membranes provide enclosure
  - Also fundamental for metabolism
- Membranes never arise from scratch
  - Always passed down and added to
  - All derived from ancestral cell
- T. Cavalier-Smith proposes membranes
  - Plus nucleic acid formed “ob-cell”
  - Merger of 2 ob-cells formed first cell

# Thioester World

C. deDuve

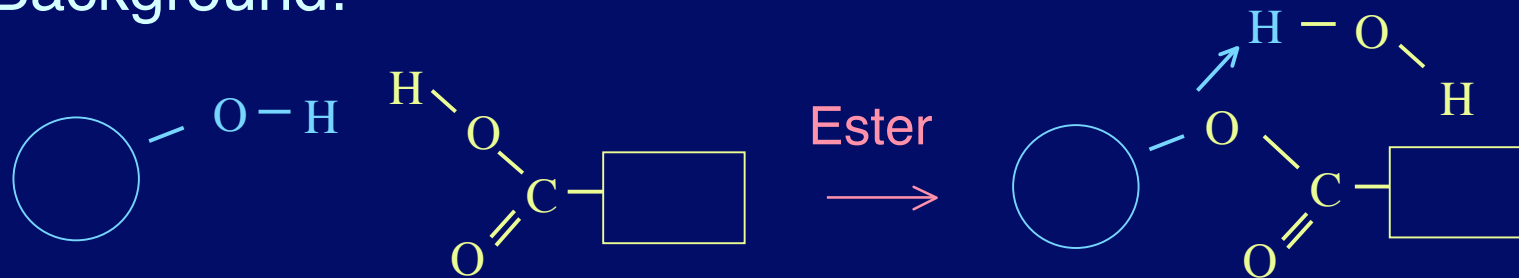
In Vital Dust

1. Need precursor to RNA world

2. Need energy conversion

Protometabolism

Background:



Hydroxyl + Carboxyl



Thiol + Carboxyl

Thiols involved in metabolism, particularly in ancient pathways

Also can catalyze ester formation by group transfer  
Reactions

e.g. peptide bonds



# Catalytic Multimers

C. deDuve

“Multimer” short peptides and esters

(NH<sub>2</sub>)

(OH)

of amino acids and hydroxy acids

Will form from thioesters assume some catalytic  
ability → protometabolism

## Energy Sources

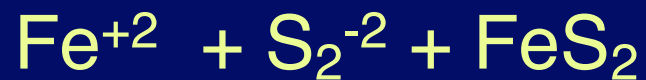
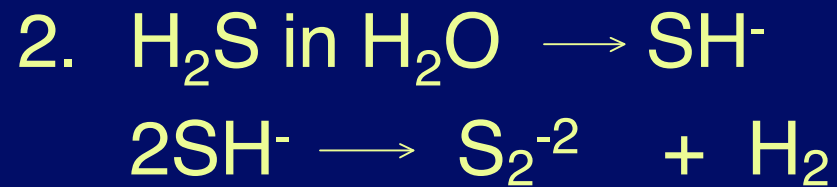
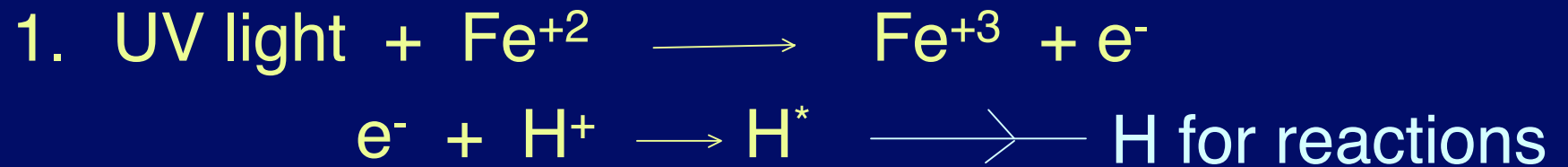
Basic need is hydrogen atoms  
(or electrons in excited states)

In pure water  $\frac{H^+ + OH^-}{H_2O}$  more if acidic

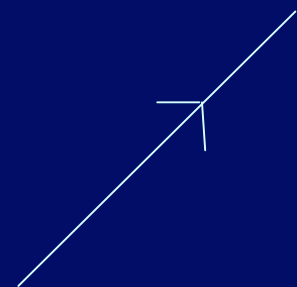


Now chlorophyll + sunlight

Then?



iron pyrite



# Transition to Phosphate

Energy currency in life now is ATP

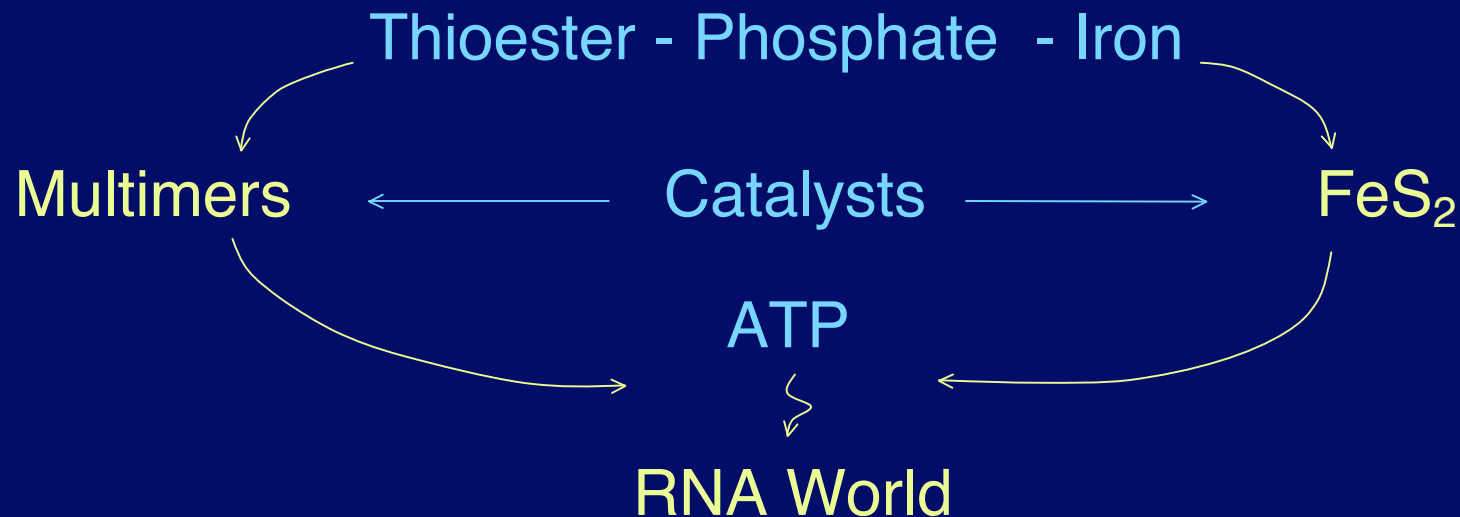
Adenosine Triphosphate

used to make bonds, remove H<sub>2</sub>O

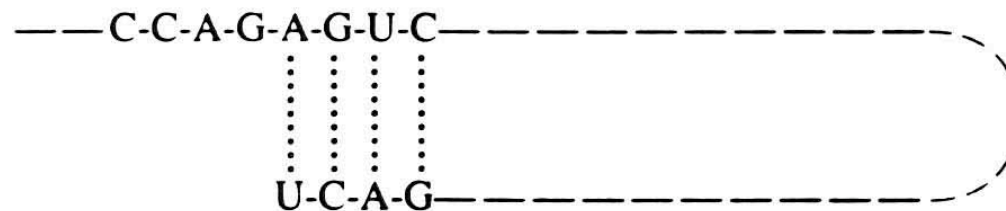
Earlier, inorganic phosphate

p-p diphosphate or polyphosphate

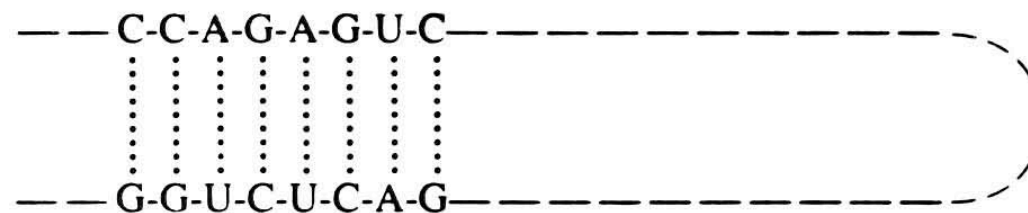
still involved in ATP reactions



age—will be followed by GUC. This AGUC sequence is complementary to the terminal sequence written in antiparallel fashion, and will cause the chain to double up as follows:



Assume now that this folded chain is subject to elongation, by the addition of new nucleotides, from right to left, to the U end. The presence of G next to the A paired with the terminal U is likely to favor the addition of a complementary C over that of the other three possible nucleotides. Repeat the process and you get U added opposite A, G opposite C, G again opposite the next C, and so on. What you get is the formation of a stretch complementary over all its length to the other end of the molecule:



# Summary of Proto-Life Development

<u>Stage</u>	<u>Proteins</u>	<u>Halfway # 1</u> Peptide Nucleic Acids	<u>Halfway # 2</u> RNA Ribozyme	<u>Nucleic Acids</u>
Monomers	Amino Acids	Bases Amino Acids	Ribose Sugars Bases Phosphates Amino Acids	Ribose Sugars Bases Phosphate
Polymerization	Proteinoids	Short strands of PNA's	Short strands of RNA + amino acids	Short strands of RNA
Replication	?	Affinity for complementary bases + ease of peptide bonding	Affinity for complementary bases	Affinity for complementary bases
Pre-life	Proteinoids + RNA?	Separation of proteins and nucleic acids	Separation of nucleic acids and protein parts	RNA adapts proteinoids as needed
Life	Proteins	Disappears	Disappears	DNA and RNA