

# Autocatalytic replicators embedded in metabolic networks

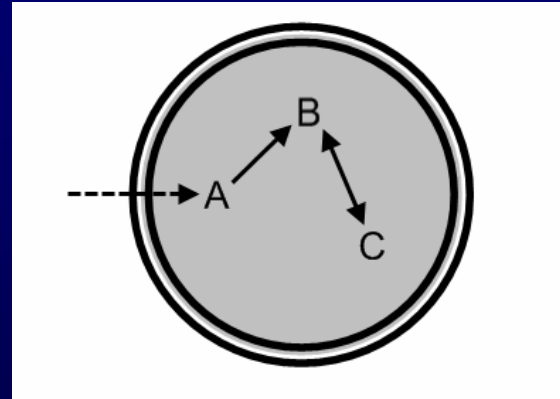
**Balázs Papp**

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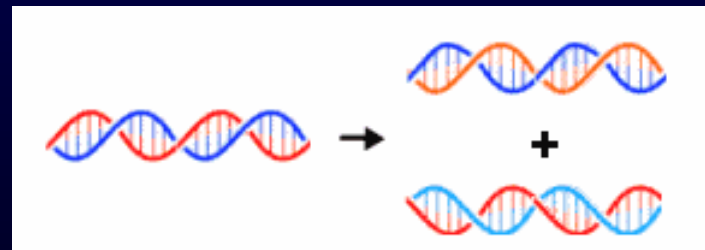
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## Two fundamental features of life:

- Metabolism



- Replication (heredity)



# What replicates at the molecular level?

- Template based replication is well established (DNA, RNA)
- But other replicators also exist:
  - Epigenetic chromatin markings
  - Membranes

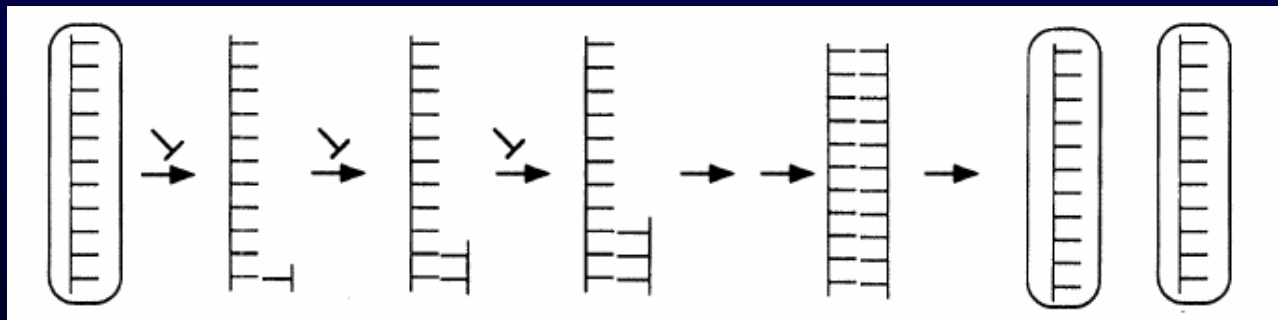
→ The essence of replication is **autocatalysis**

# Autocatalysis

A compound (A) catalyses its own formation:

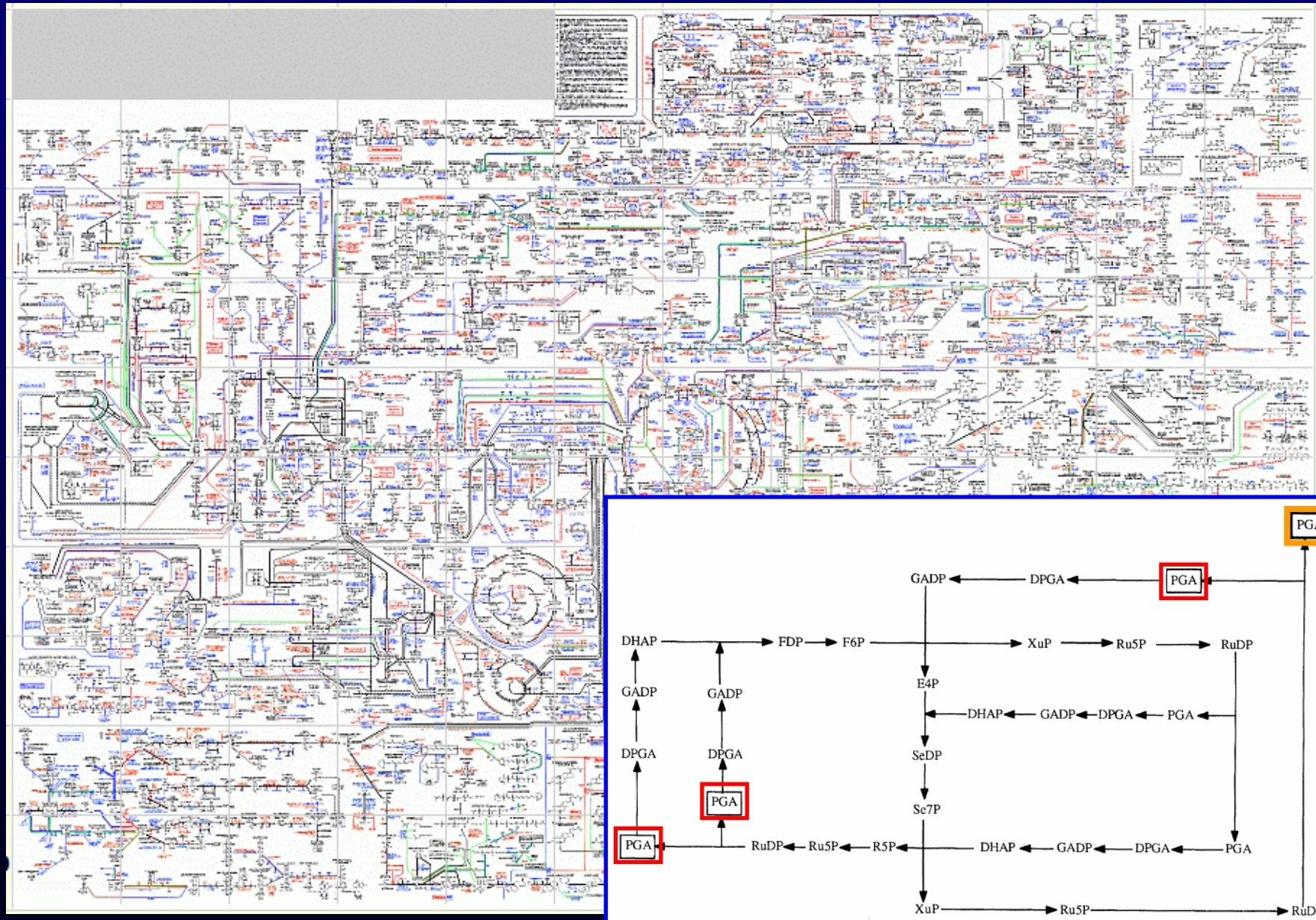


Autocatalysis in nucleic acid replication:



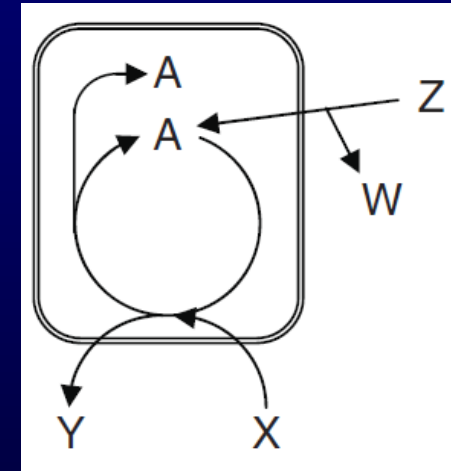


# Autocatalytic cycles are embedded in large networks



# Is the whole metabolic network necessarily autocatalytic?

- Not, if the cycle intermediates can be reconstructed via other routes



A can be also synthesized from Z

- How about heterotrophic organisms? Do they also contain autocatalytic cycles?

# Research questions

Are the metabolites of a network accessible just from the food molecules or do we need to add compounds from the network itself to bootstrap the metabolism?



Do 'bootstrapping' compounds differ between different species?

# Method

- We searched for autocatalytic compounds in various species and in an inferred minimal metabolism
- Only high-quality network reconstructions were used → with information on food molecules that can be imported

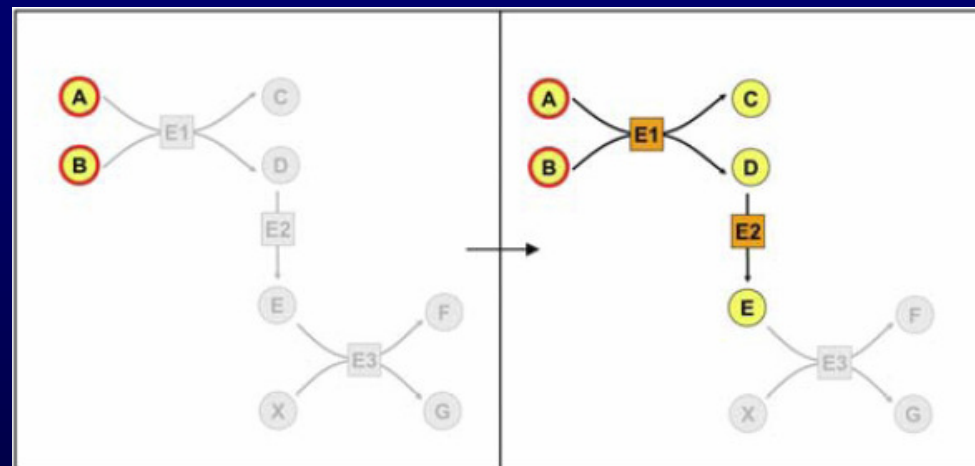
	Total number of metabolites	Number of food molecules
<i>Escherichia coli</i>	761	143
<i>Helicobacter pylori</i>	485	74
<i>Staphylococcus aureus</i>	644	83
<i>Saccharomyces cerevisiae</i>	672	101
<i>Lactococcus lactis</i>	508	92
<i>Streptomyces coelicolor</i>	601	104
<i>Mycobacterium tuberculosis</i>	830	87
<i>Methanosarcina barkeri</i>	628	70
<i>Geobacter sulfurreducens</i>	541	41
<i>Synechocystis</i> <sup>†</sup>	879	18
Minimal metabolism	68	11

# Method

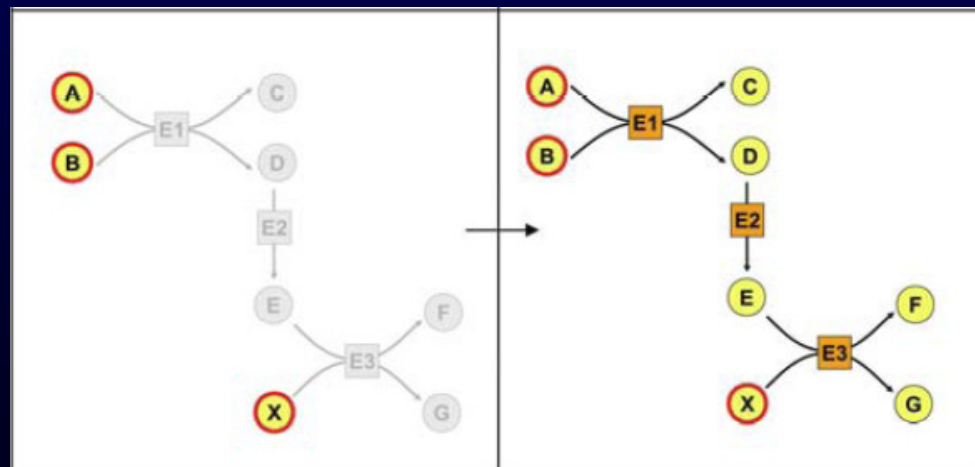
Scope analysis\*: identifying compounds that are not accessible from food molecules

A, B: food molecules

X, F and G are not accessible!

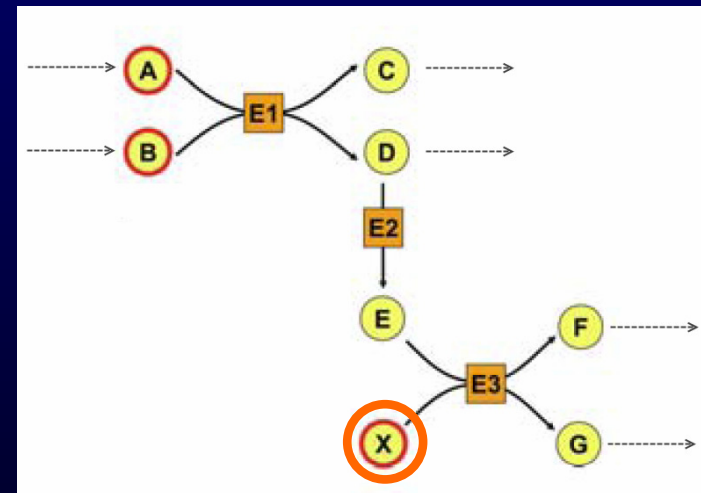


All metabolites become accessible when X is provided



# Method

- We identified the smallest set of compounds that have to be added to the network to make all metabolites accessible
- Not all these compounds are autocatalytic, some are simply dead-end

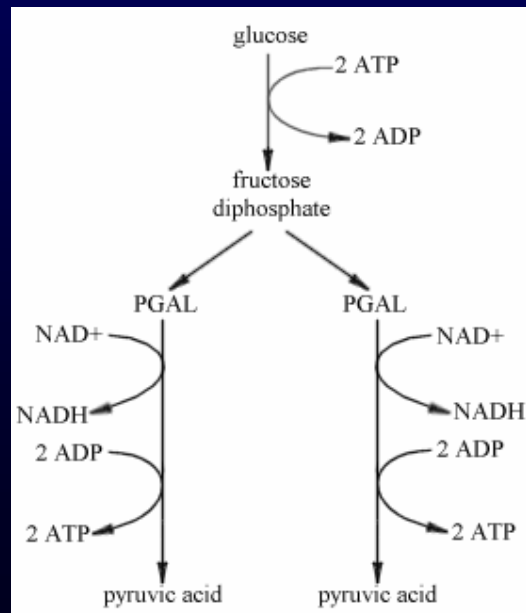


X is a dead-end metabolite

# Results

- All analysed metabolic networks show autocatalytic behaviour, even the minimal metabolism
- At least one molecule has to be always added to kick-start the metabolism:

**ATP** synthesis is universally autocatalytic



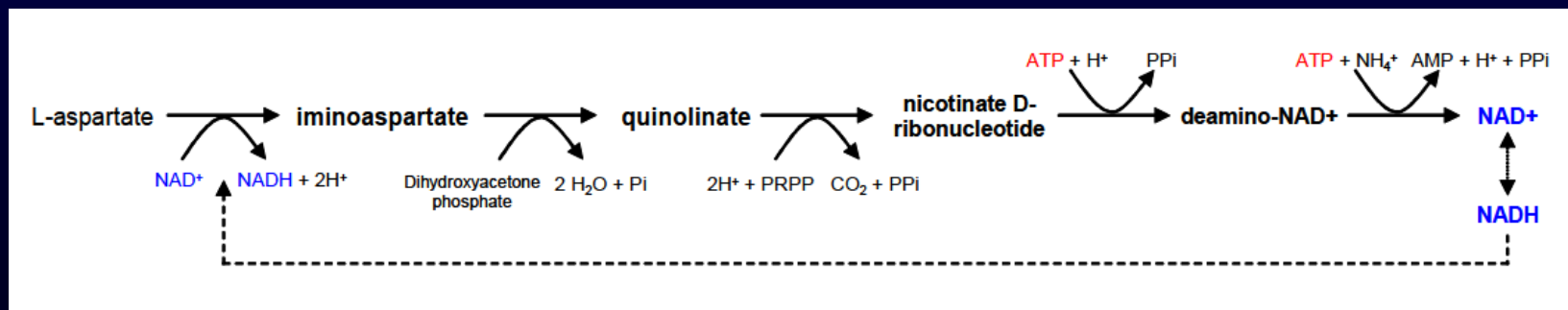
E.g.: glycolysis

# Results

- Sugar synthesis is autocatalytic in *Synechocystis* when growth is photosynthetic ( $\rightarrow$  Calvin cycle), but not when the food set includes organic compounds
- Some organisms contain other autocatalytic cofactors in addition to ATP:  $\text{NAD}^+$ , CoA, THF

Example:

$\text{NAD}^+$  biosynthesis in *Geobacter sulfurreducens*



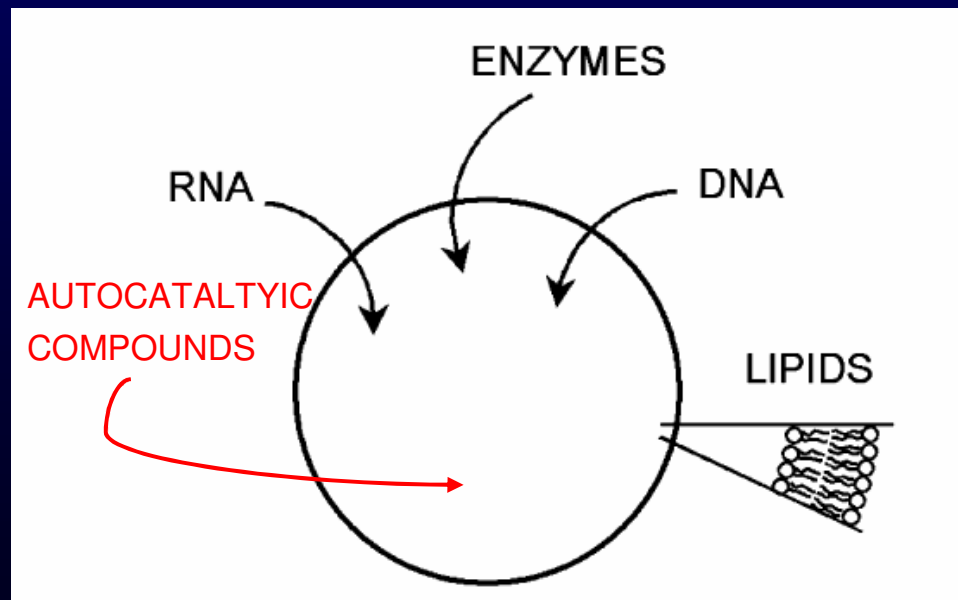


# Conclusions

- Metabolism is universally autocatalytic at the level of small molecules, even when organic compounds are available for uptake
- Even a hypothetical minimal network is autocatalytic
- There are different metabolic replicators: ‘alleles’

# Implications for synthetic biology

- There are projects to synthesize minimal cells from scratch
- ‘bootstrapping’ compounds have to be considered when designing synthetic cells



# Implications for evolutionary biology

- Metabolic replicators provide a limited form of inheritance

DNA: number of possible types  $\gg$  number of DNA molecules

metabolic replicators: only few possible types

- The autocatalytic nature of coenzyme biosynthesis might be an ancestral feature
- Origin of life: are non-enzymatic autocatalytic cycles possible (apart from formose reaction)?

# Acknowledgements

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Human Frontier Science Program

## Open postdoctoral positions

[www.brc.hu/~sysbiol](http://www.brc.hu/~sysbiol)

# Hypothetical minimal metabolism based on endosymbiont genomes (Gil et al. 2004)

