

Synthetic life

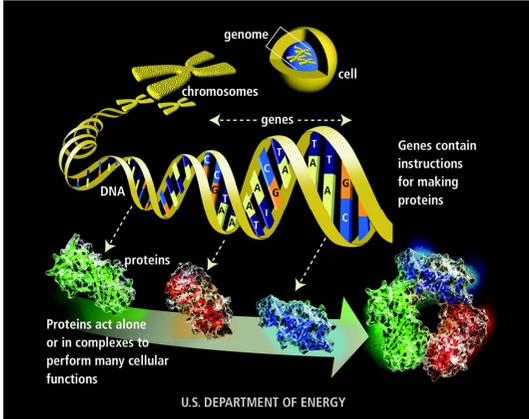


WiSe 2017/18

Zbigniew Pianowski

NaturalNews.com

From DNA to proteins



<https://www.youtube.com/watch?v=gG7uCsUOrA>

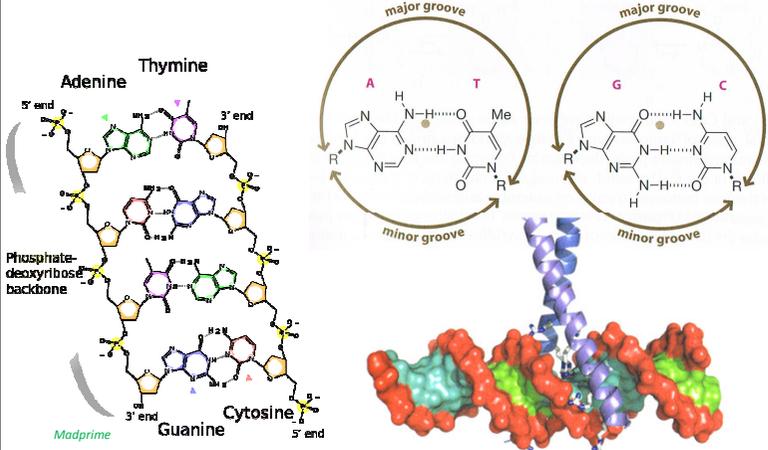
U.S. DEPARTMENT OF ENERGY

CHAPTER 1

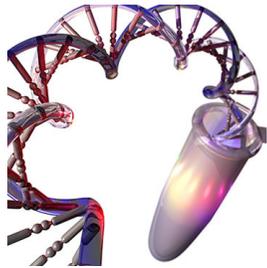


OLIGONUCLEOTIDES

Canonical nucleobases and Watson-Crick pairing in DNA

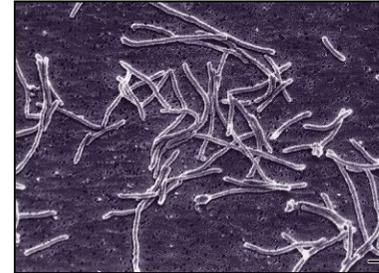


PCR – Polymerase Chain Reaction



National Library of Medicine,
National Institutes of Health

PCR – Polymerase Chain Reaction

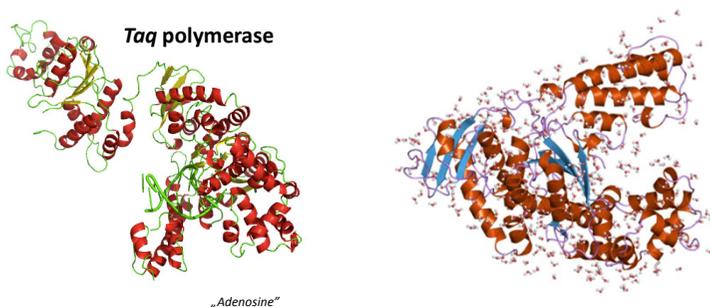


Brian W. Schaller, Yellowstone Park

Thermus aquaticus is a thermophilic bacteria from hot springs in Yellowstone Park
70°C – optimum, living range: 50-80°C

It is a source of thermostable enzymes

PCR – Polymerase Chain Reaction



Taq polymerase

„Adenosine“

Taq polymerase withstands denaturing conditions (hot temperatures) detrimental for most enzymes. Activity optimum: 75-80°C, half-life at 95°C > 2.5 h

1990 – Kary Mullis optimized the PCR technique with *Taq* polymerase (1993 Nobel Prize)

PCR – Polymerase Chain Reaction

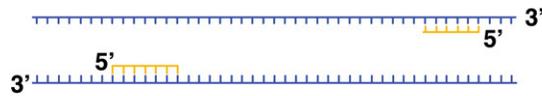


We begin with a single molecule of DNA.

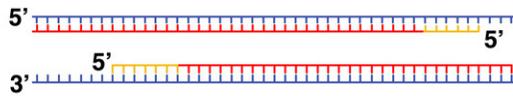


We can melt the DNA (break the hydrogen bonds holding the helix together) by heating it to 98 degrees.

PCR – Polymerase Chain Reaction

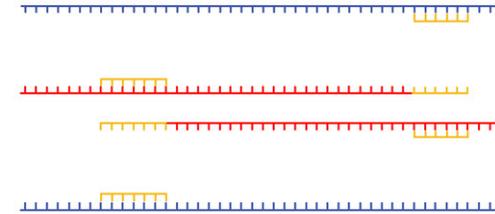


Two DNA primers (18-22 bp, T_m : 50-60°C) are designed to anneal to a known sequence. The primers are separated in the sequence that we are targeting by a few hundred base pairs. Cooling the reaction from 98°C to a more moderate temperature allows annealing to take place.



Now we have two primed templates. With dNTPs and DNA polymerase in the reaction mixture, new DNA is synthesized.

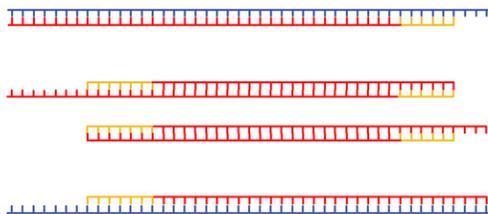
PCR – Polymerase Chain Reaction



The DNA is molten for another cycle. Because there is a vast molar excess of primers, when we cool the mixture, we again anneal primers

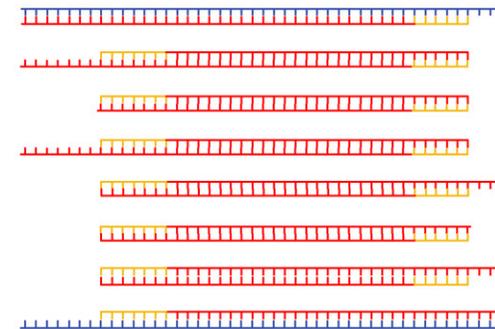
PCR – Polymerase Chain Reaction

New DNA is synthesized



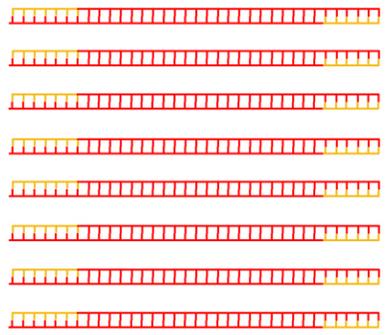
PCR – Polymerase Chain Reaction

In the next cycle, we begin to see DNA molecules whose ends are defined by the primers



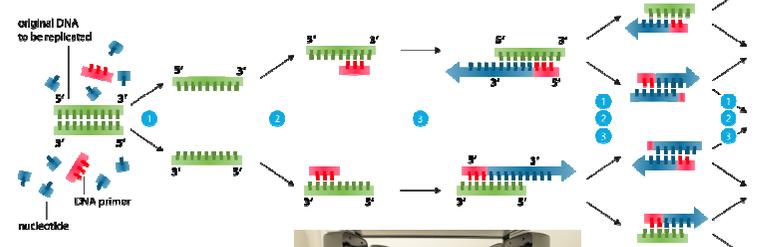
PCR – Polymerase Chain Reaction

After many cycles of melting, annealing, and replication, the overwhelming majority of DNA molecules in the mixture have ends defined by the primers

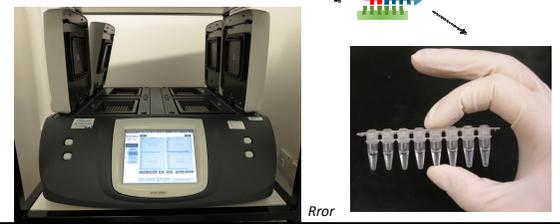


<https://www.dnalc.org/view/15475-The-cycles-of-the-polymerase-chain-reaction-PCR-3D-animation.html>

Polymerase chain reaction - PCR



- 1 Denaturation at 94-96°C
- 2 Annealing at 50-65°C
- 3 Elongation at 72 °C



Rror

DNA sequencing

DNA SEQUENCING

NHGRI FACT SHEETS
genomes.gov

Genomic Sequences into Sequencing Machine

COMPARATIVE GENOMICS

NHGRI FACT SHEETS
genomes.gov

Researchers choose the appropriate time-scale of evolutionary conservation for the question being addressed.

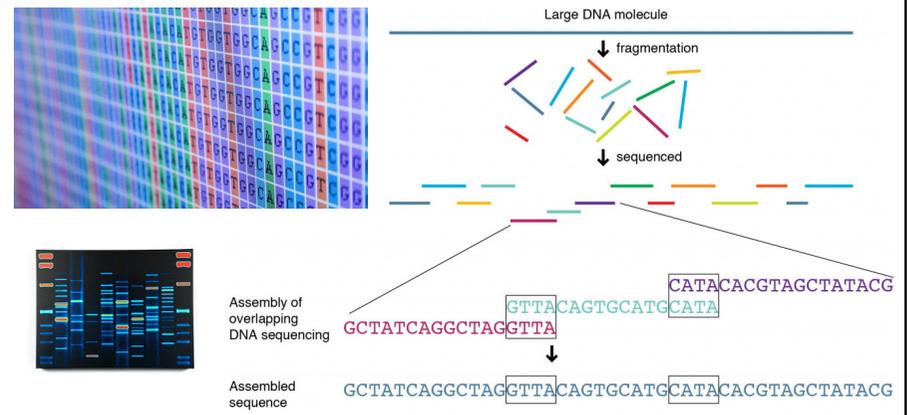
Common features of different organisms such as humans and fish are often encoded within the DNA evolutionarily conserved between them.

Looking at closely related species such as humans and chimpanzees shows which genomic elements are unique to each.

Genetic differences within one species such as our own can reveal variants with a role in disease.

National Library of Medicine, National Institutes of Health

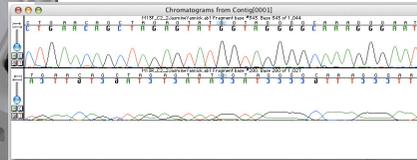
Genome sequencing



Genome sequencing



Electropherograms are commonly used to sequence portions of genomes



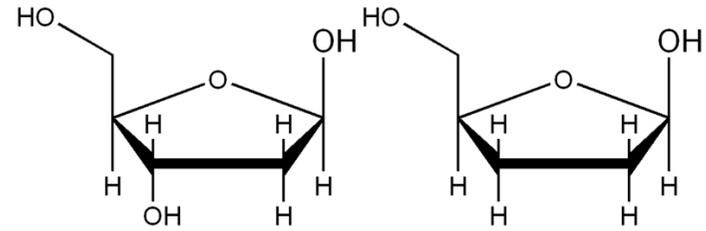
Tom David

Mark Pellegrini

An ABI PRISM 3100 Genetic Analyzer. Such capillary sequencers automated the early efforts of sequencing genomes.

Sanger sequencing

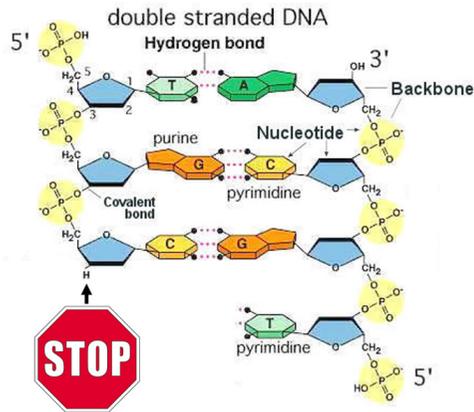
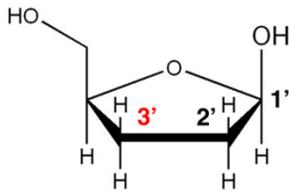
What good is dideoxyribose?



deoxyribose

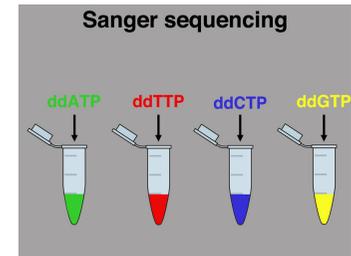
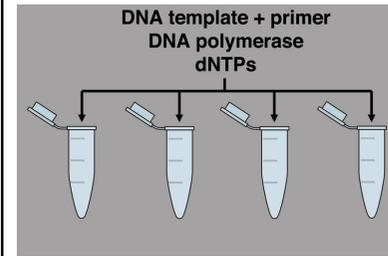
dideoxyribose

Sanger sequencing



Sanger sequencing

primer
 5' 3'
 TACGT
 ATGCATTAGGGCCTGGCTCTTT
 3' 5'
 template



Sanger sequencing

Fluorescent dideoxynucleotides

standard

chemically cleavable (reduction)

J. Ju et al. PNAS 2005, 102 (17), 5926-5931

Sanger sequencing Sequencing Gel

TACGTAA
ATGCATTAGGGCCTGGCTCTTT

TACGTAA
ATGCATTAGGGCCTGGCTCTTT

TACGTAATCCCGGA
ATGCATTAGGGCCTGGCTCTTT

TACGTAATCCCGGACCGA
ATGCATTAGGGCCTGGCTCTTT

A T C G
G C A T

Sanger Sequencing

PCR containing fluorescent, chain-terminating dideoxynucleotide triphosphates

Detector
Chromatogram
120 130
ACTGCTTG CAGCA

G C A T

Sanger sequencing uses ddNTPs (dideoxynucleotide triphosphates) which do not have a free 3' OH mixed in with dNTPs. Whenever the DNA polymerase incorporates a ddNTP it won't be able to add any other nucleotides. Then gel electrophoresis is used to separate the DNA.

<https://www.youtube.com/watch?v=ONGdehkB8jU> (from 0:50)

Sanger Sequencing

- Reaction mixture
 - Primer and DNA template
 - DNA polymerase
 - dNTPs with fluorochromes = dNTPs (dATP, dGTP, dCTP, and dTTP)

Primer
Template

Primer elongation and chain termination

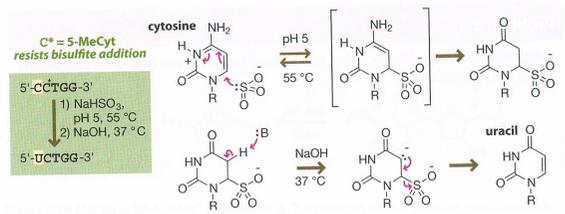
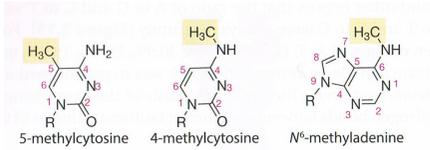
Capillary gel electrophoresis separation of DNA fragments

Laser
Detector

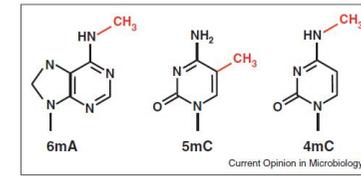
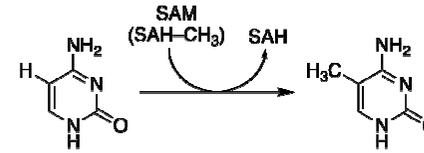
Laser detection of fluorochromes and computational sequence analysis

Chromatogram

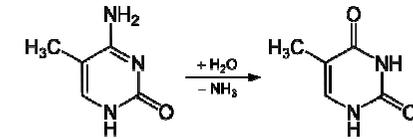
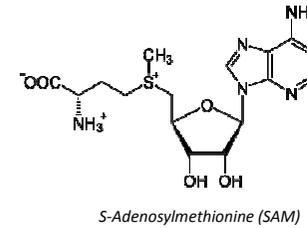
Modifications of nucleobase structures tolerated by polymerases



Modifications of nucleobases

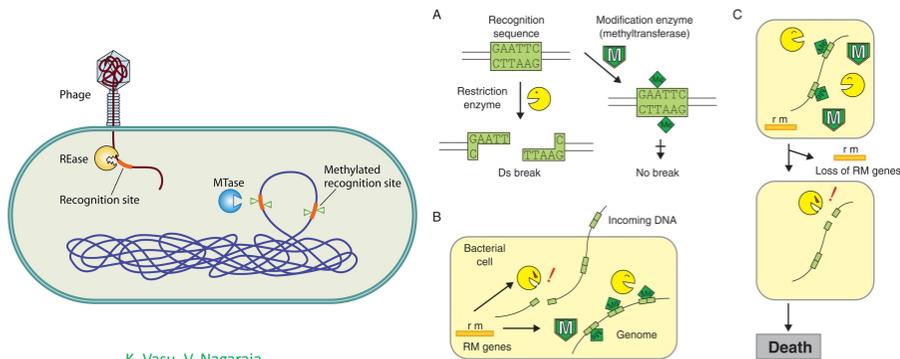


Chemical structures of common modified bases generated by DNA methyltransferases.



Restriction modification system

„Immune system“ of bacteria and archaea against attacking viruses

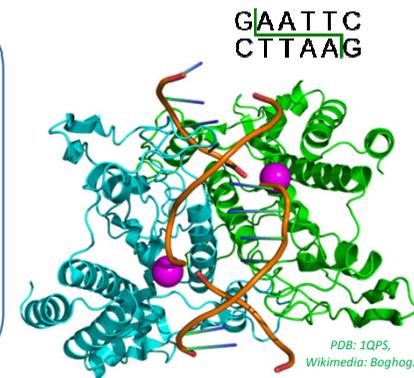
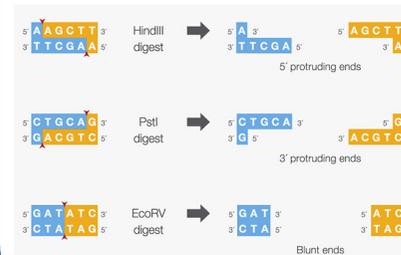


K. Vasu, V. Nagaraja
Microbiol. Mol. Biol. Rev. **2013**, *77*(1), 53-72

K. Ishikawa et al. DOI: 10.1093/dnares/dsq027

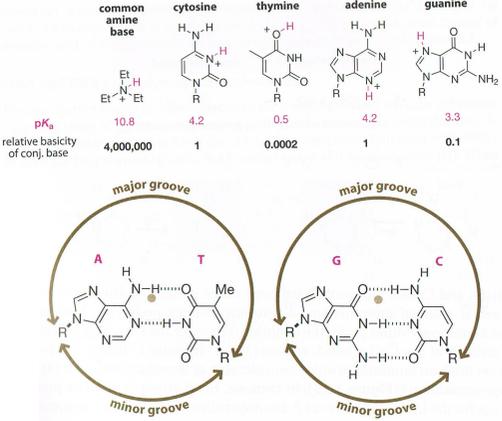
EcoI – a typical restriction enzyme

Products of restriction enzymes

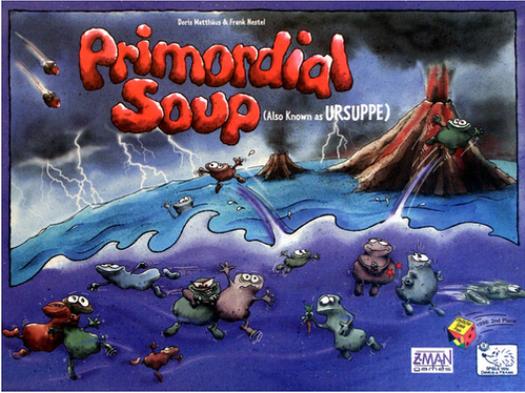


Structure of the homodimeric restriction enzyme EcoRI (cyan and green cartoon diagram) bound to double stranded DNA (brown tubes). Two catalytic magnesium ions (one from each monomer) are shown as magenta spheres and are adjacent to the cleaved sites in the DNA made by the enzyme (depicted as gaps in the DNA backbone).

Why are A, C, G and T the letters of genetic alphabet.



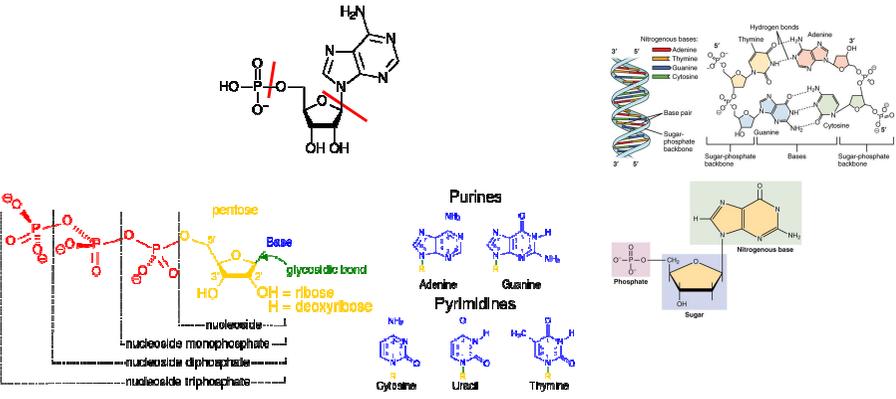
Prebiotic synthesis of nucleotides



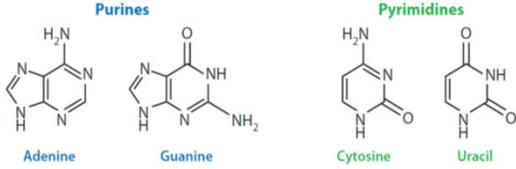
State of the art

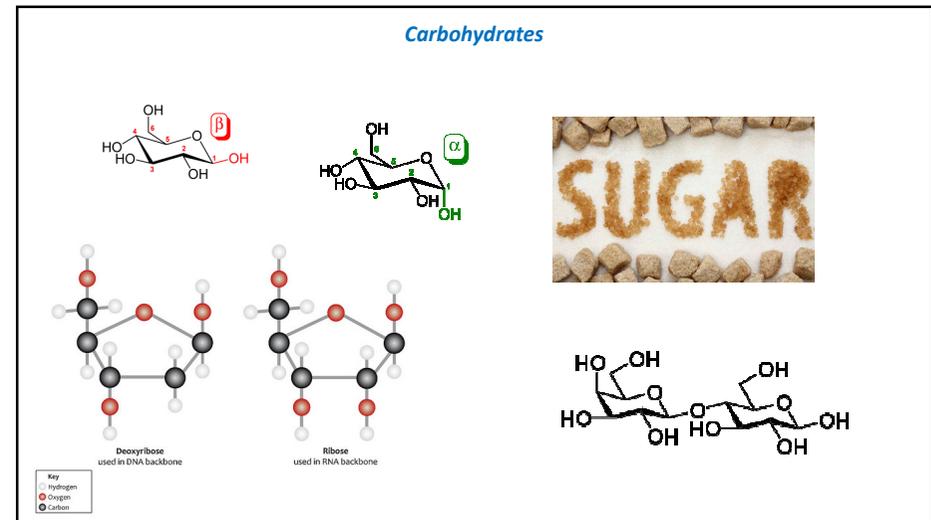
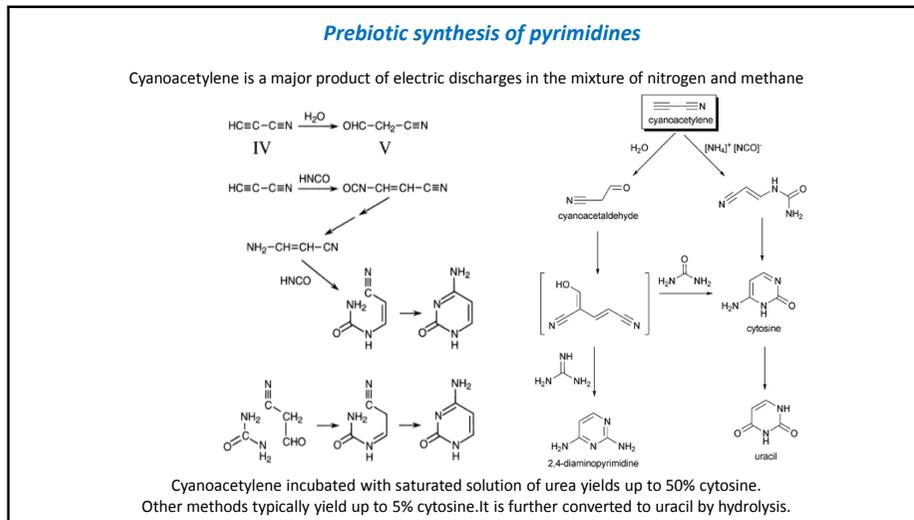
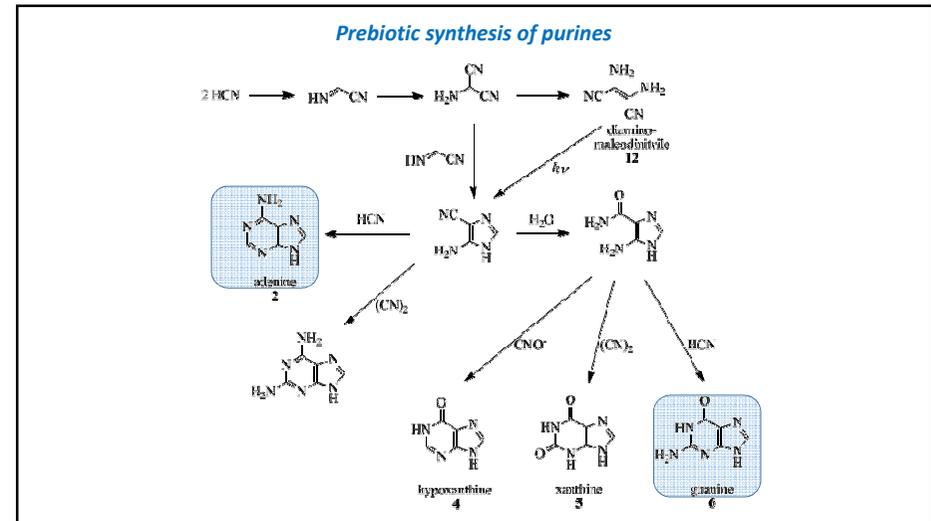
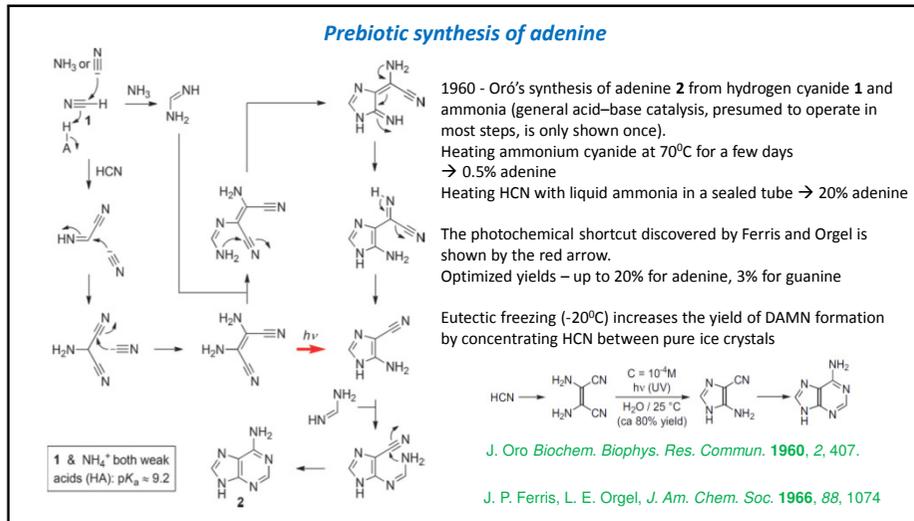
Nucleotides - components

RNA – most likely evolutionarily older („RNA World“) than DNA → prebiotic origin of ribose + A, C, G, and U nucleobases



Prebiotic synthesis of nucleobases





Formose reaction

The reaction begins with two formaldehyde molecules condensing to make glycolaldehyde 1 which further reacts in an aldol reaction with another equivalent of formaldehyde to make glyceraldehyde 2. An aldose-ketose isomerization of 2 forms dihydroxyacetone 3 which can react with 1 to form ribulose 4, and through another isomerization ribose 5. Molecule 3 also can react with formaldehyde to produce tetrosule 6 and then aldoltetrose 7. Molecule 7 can split into 2 in a retro-aldol reaction.



Alexander Butlerov (1828-1886)
St. Petersburg, Kazan, Russia



Ronald Breslow (1931-)
Columbia University, USA

Formose reaction as an autocatalytic process

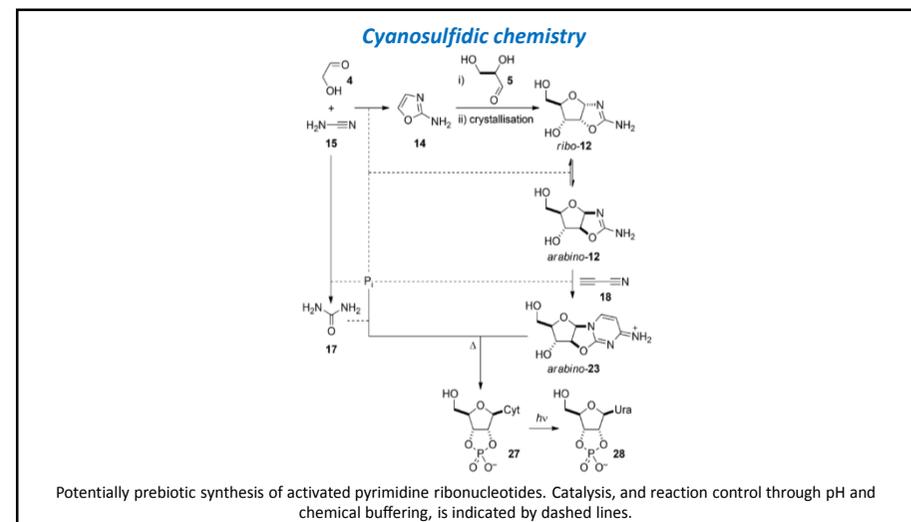
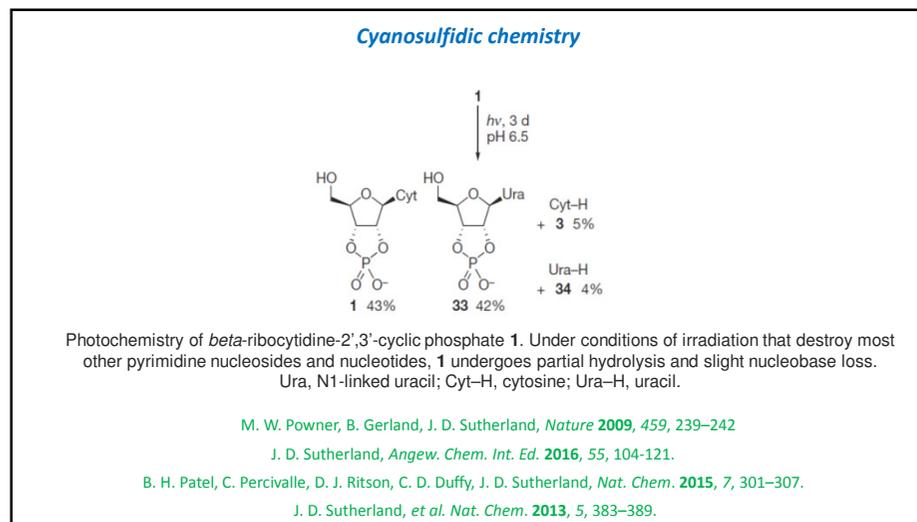
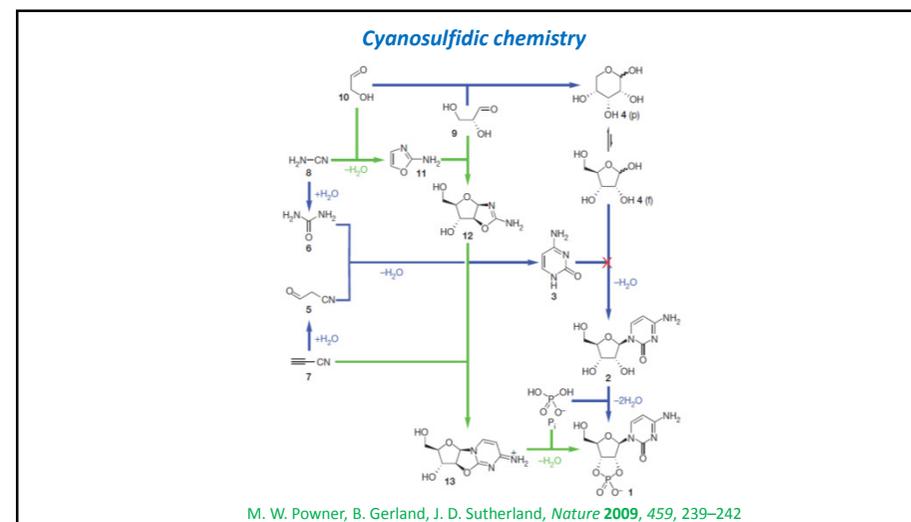
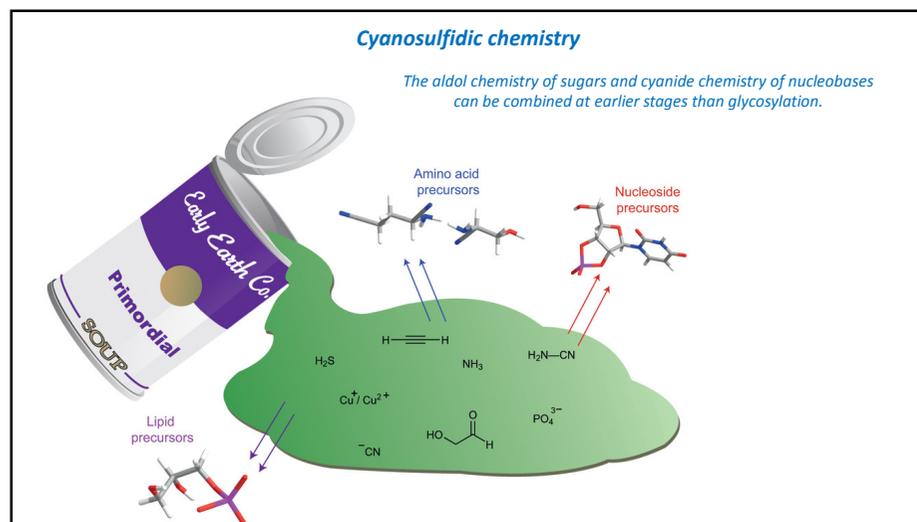
Formose reaction starts in concentrated alkaline aqueous solutions of formaldehyde alkali are typically calcium, magnesium or lead

Formose reaction under standard basic catalysis – Ca(OH)₂

Escosura

Formose reaction in presence of borates

Escosura



How else could it end up?

common amine base	cytosine	thymine	adenine	guanine	
pK _a	10.8	4.2	0.5	4.2	3.3
relative basicity of conj. base	4,000,000	1	0.0002	1	0.1

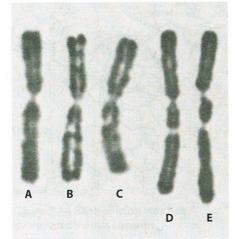
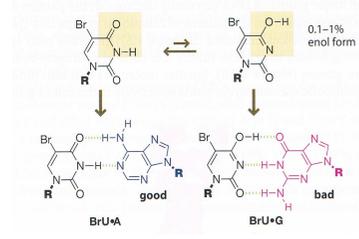
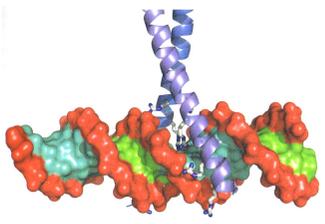
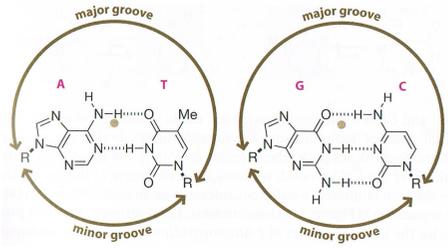
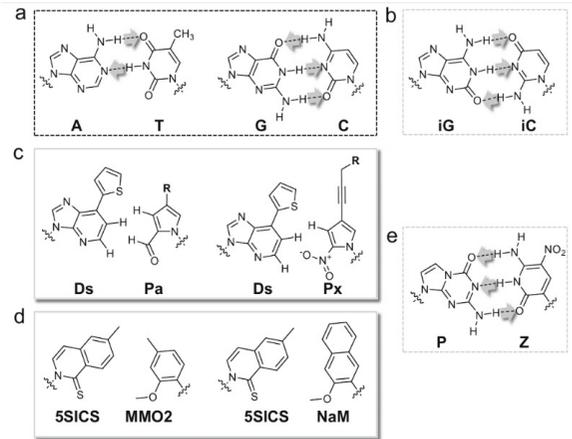
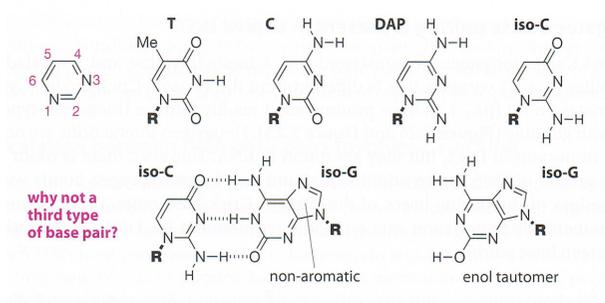


Figure 3.20 Chromosome 1 from hamster cells exposed to bromodeoxyuridine. (A) Normal chromosome. (B-E) Aberrant chromosomes. (From T.C. Hsu and C.E. Somers, *Proc. Natl. Acad. Sci. USA* 47: 396-403, 1961. With permission from the MD Anderson Cancer Center.)

Natural and non-natural base pairs that function in polymerase reactions



Alternative base pairs – synthetic biology



DAP – one tautomer forms a base pair with guanine

iso-C/iso-G
 - specificity (the enol tautomer of iso-G, stabilized by aromatization, complementary to thymine)
 - the 2-amino group of iso-C hydrolyses easily to uracil