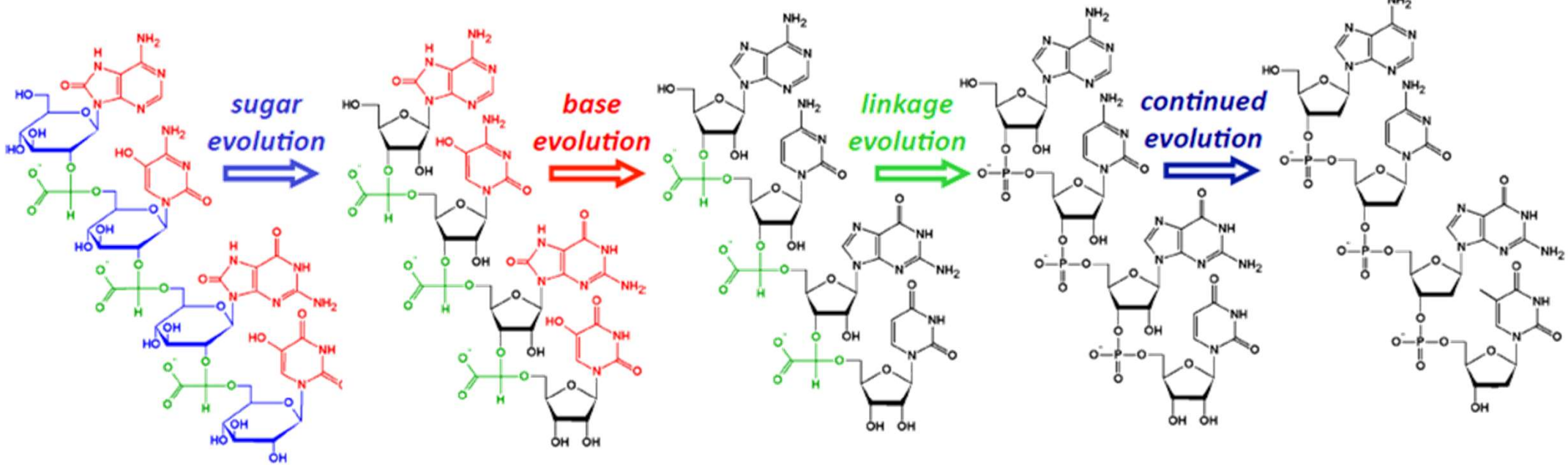


„RNA-second“

proto-RNA

RNA

DNA



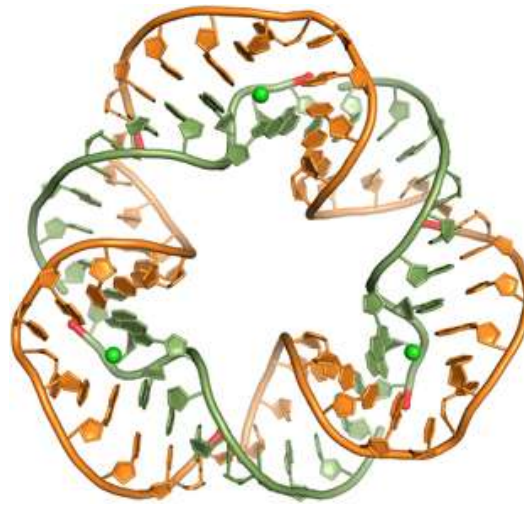
Easy to assemble



Functionally superior

Proto-RNA evolution: According to the protoRNA theory, each of the components of RNA — sugar, base and phosphate backbone — may have originally taken different forms.

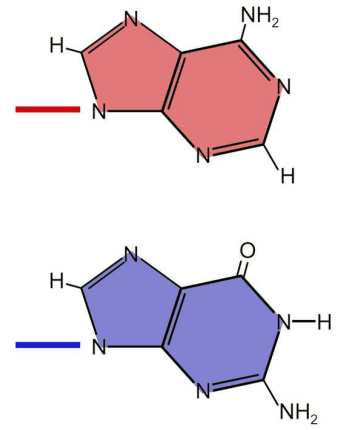
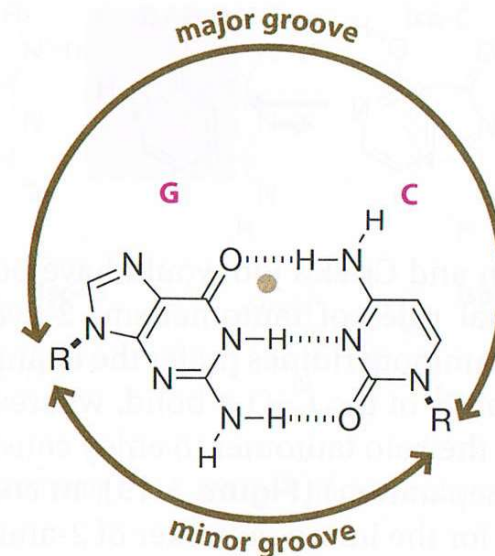
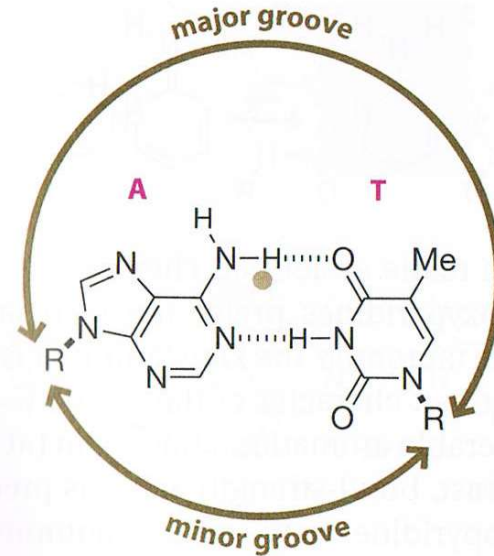
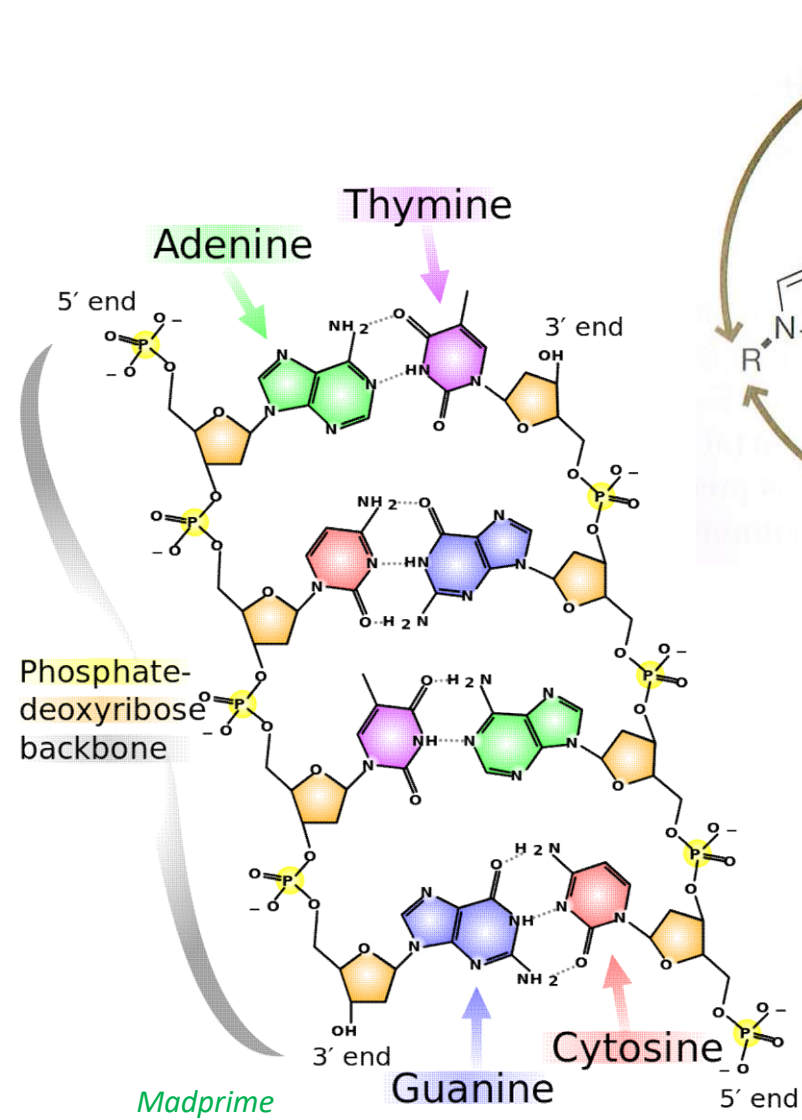
CHAPTER 1



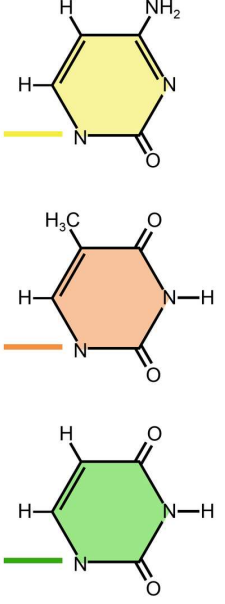
OLIGONUCLEOTIDES

Part 1 – modified canonical nucleobases

Canonical nucleobases and Watson-Crick pairing in DNA



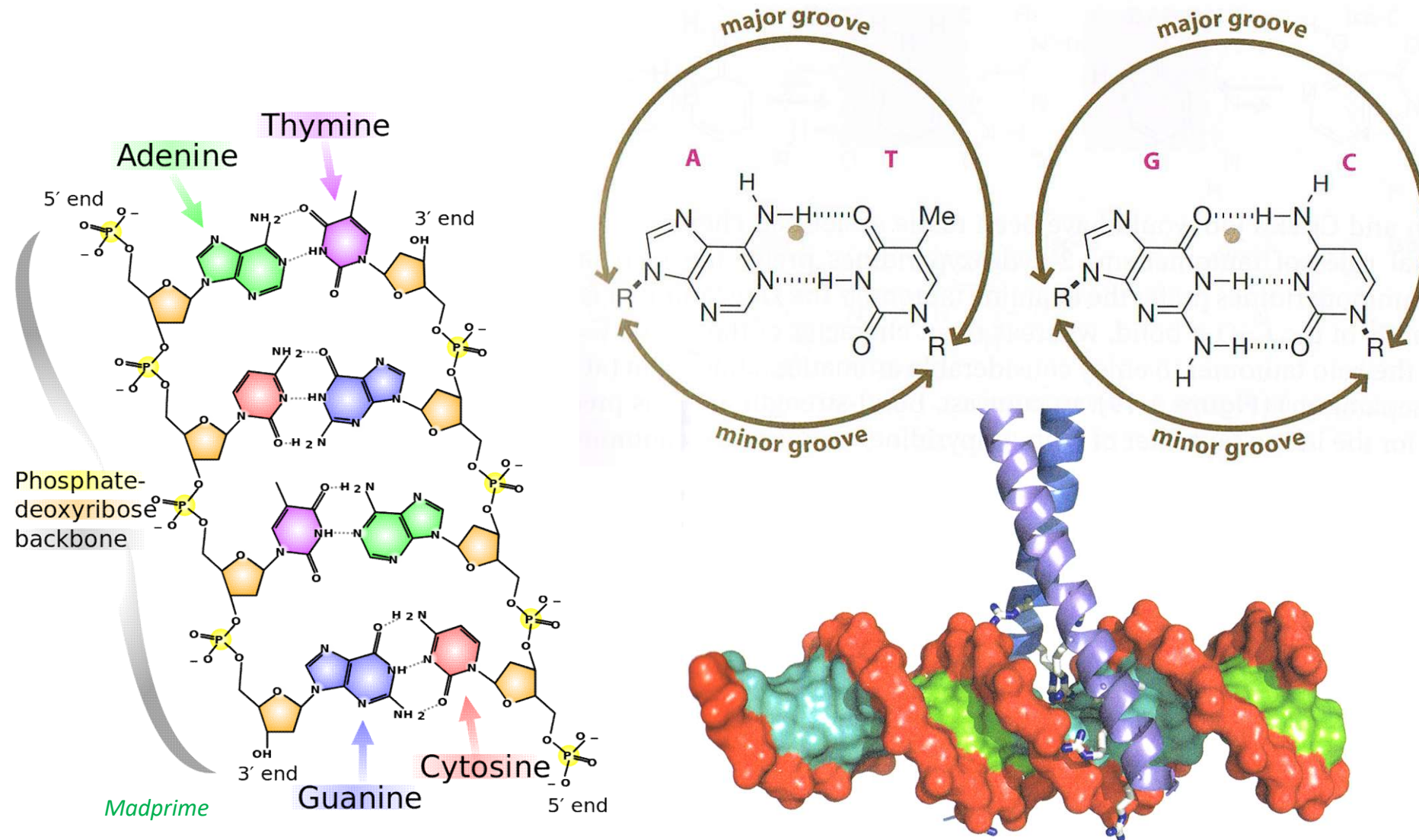
Purines



Pyrimidines

Bruce Blaus

Canonical nucleobases and Watson-Crick pairing in DNA



Hoogsten base pairing of canonical DNA nucleobases

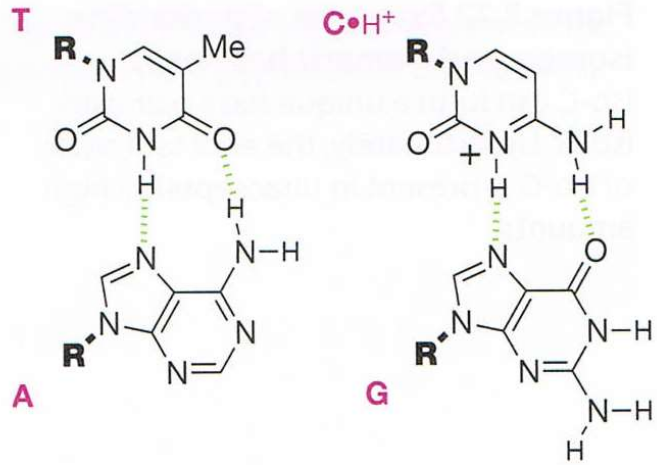
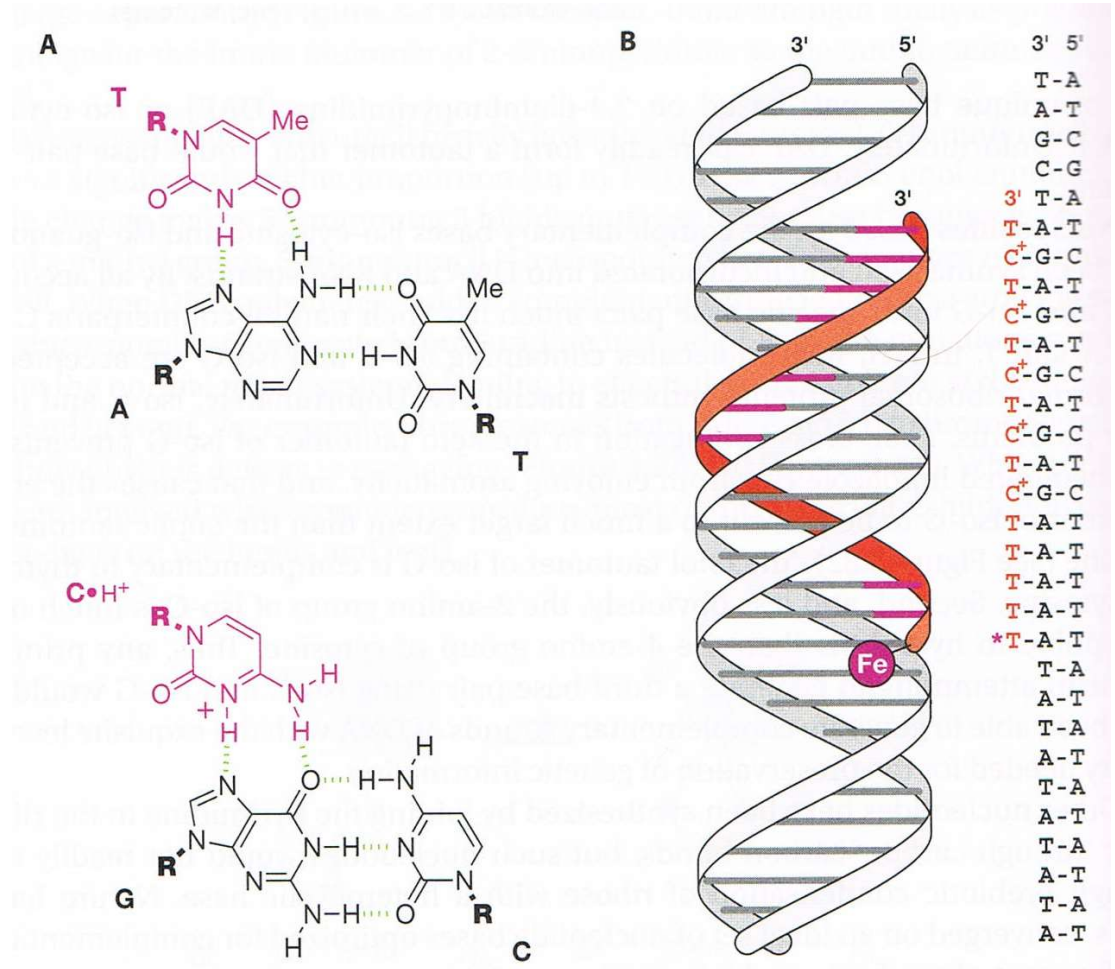
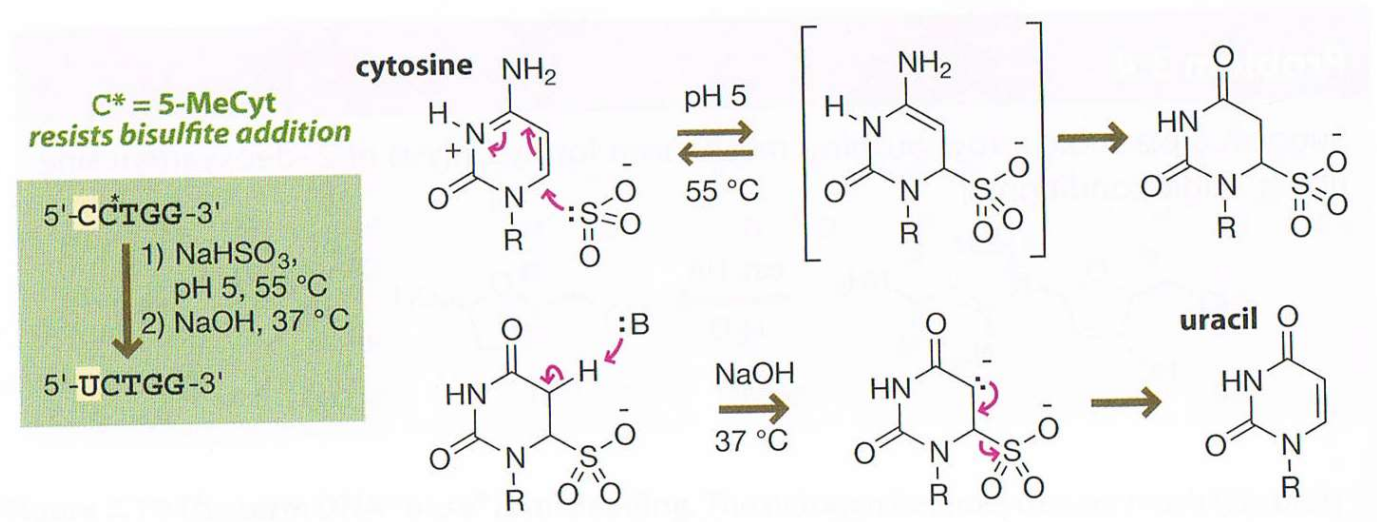
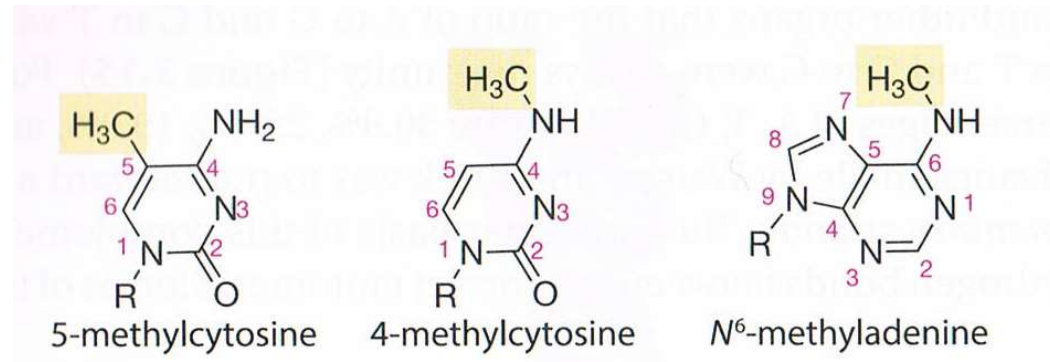


Figure 3.24 Hoogsteen base pairs. Hoogsteen base pairs use a different edge of the purine from a Watson–Crick base pair.

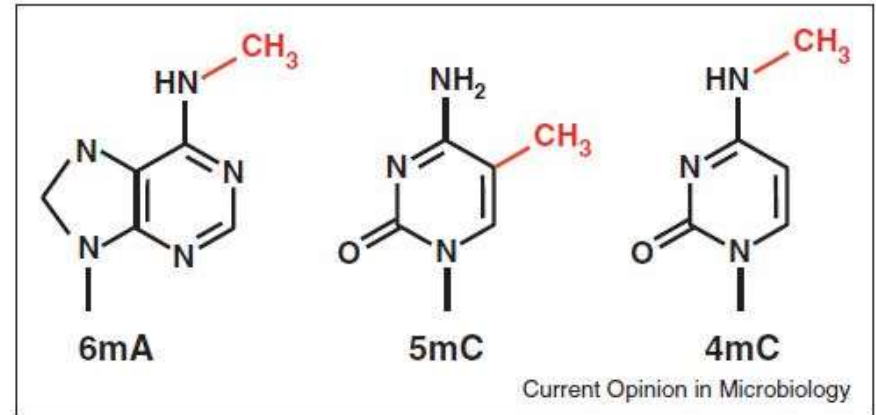
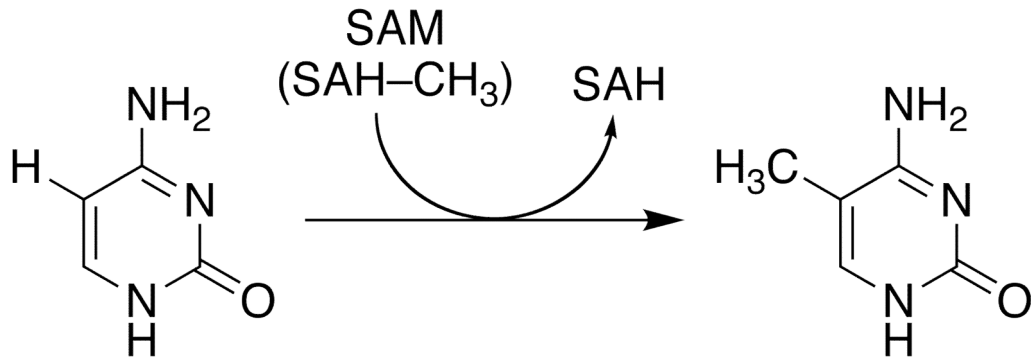


H. E. Moser, P. B. Dervan *Science* **1987**, *238*, 645-650

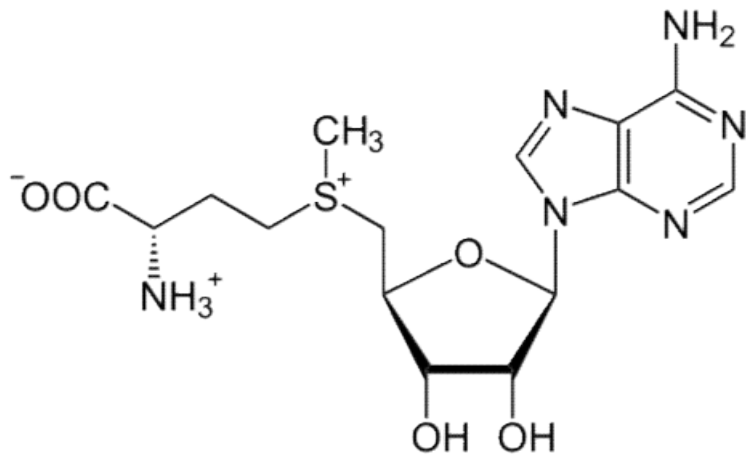
Modifications of nucleobase structures tolerated by polymerases



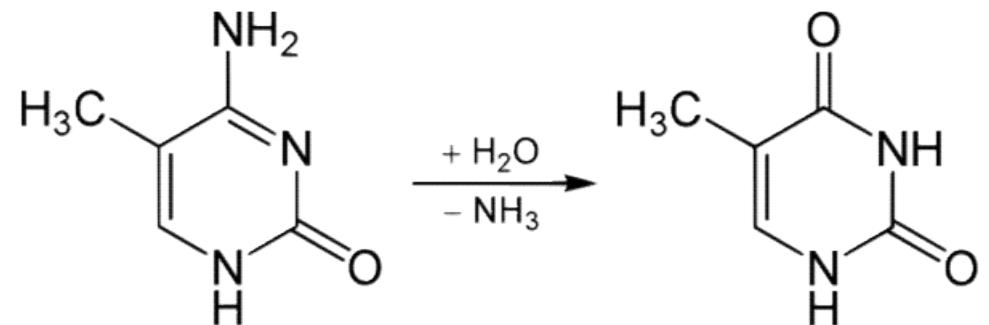
Modifications of nucleobases



Chemical structures of common modified bases generated by DNA methyltransferases.

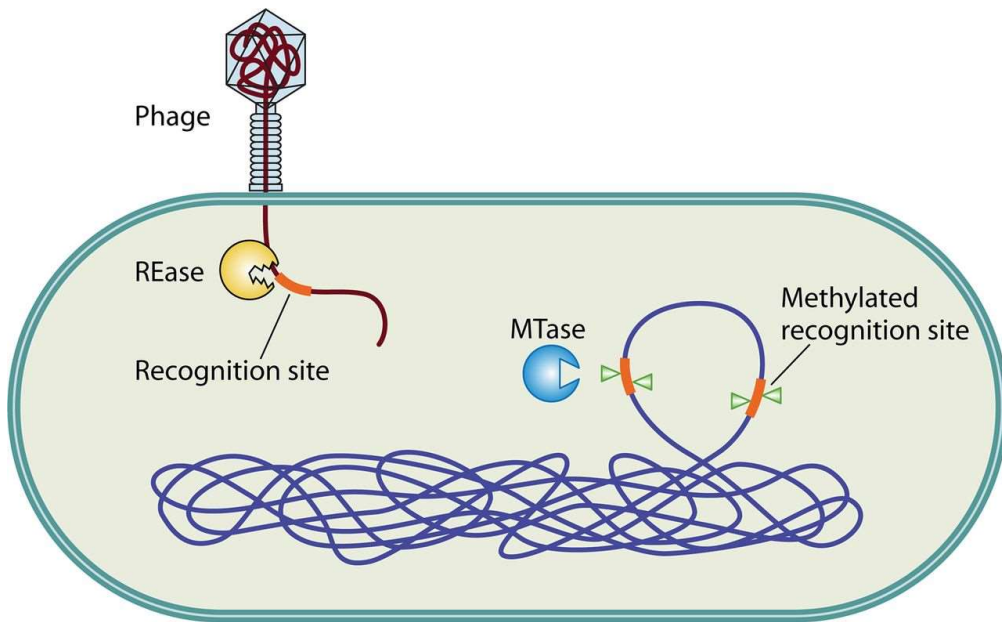


S-Adenosylmethionine (SAM)



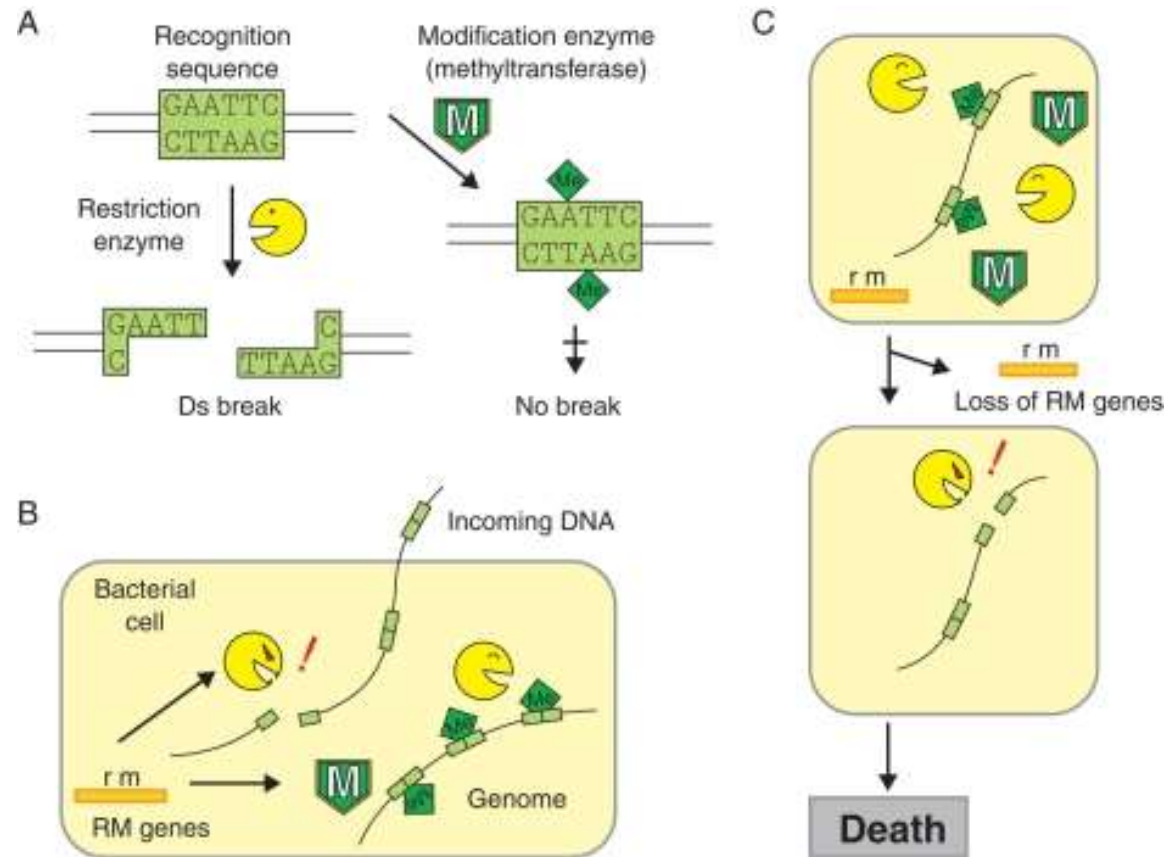
Restriction modification system

„Immune system“ of bacteria and archaea against attacking viruses



K. Vasu, V. Nagaraja

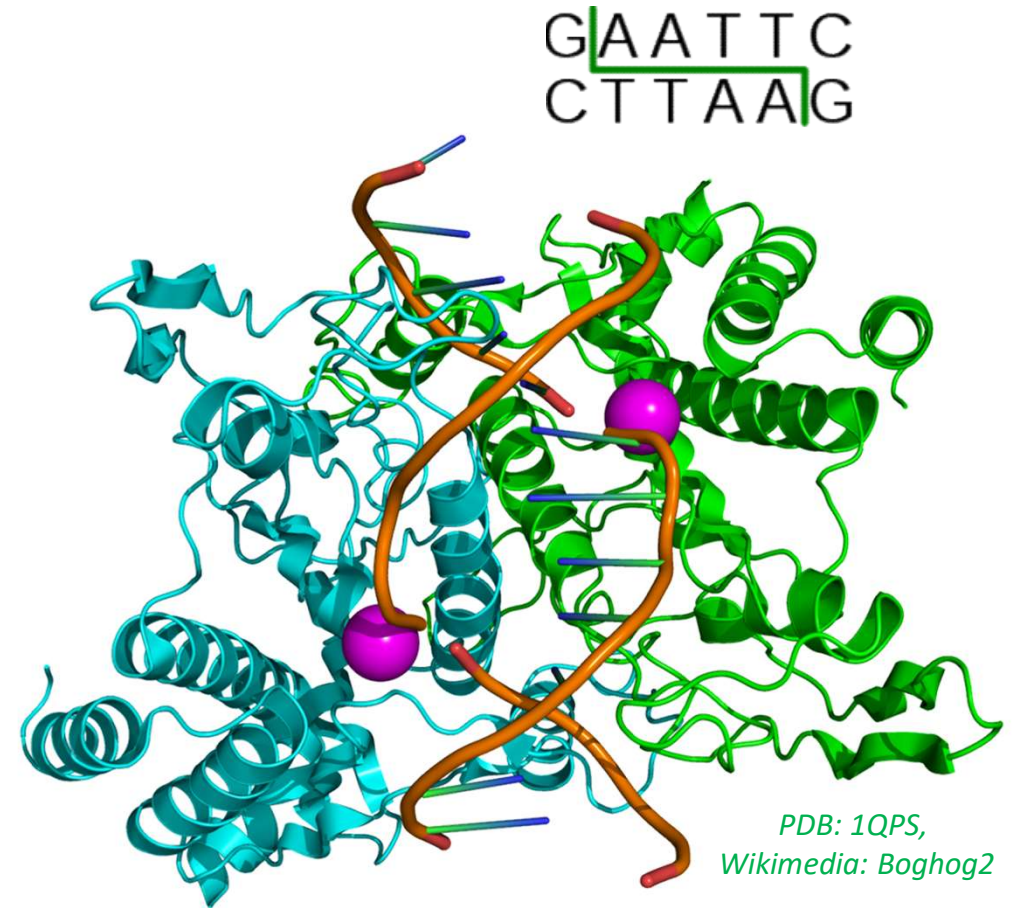
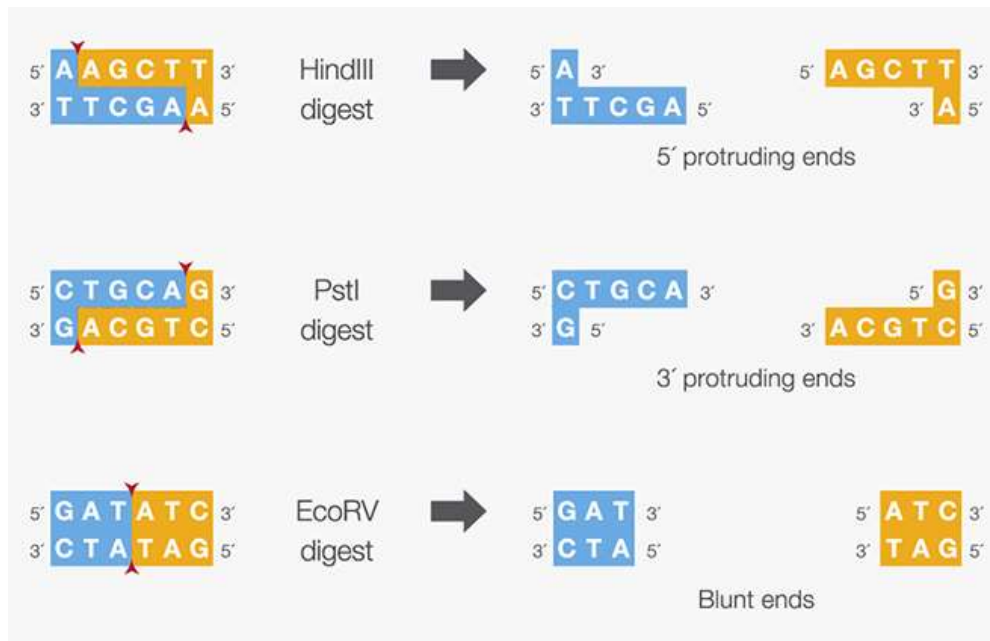
Microbiol. Mol. Biol. Rev. **2013**, 77(1), 53-72



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

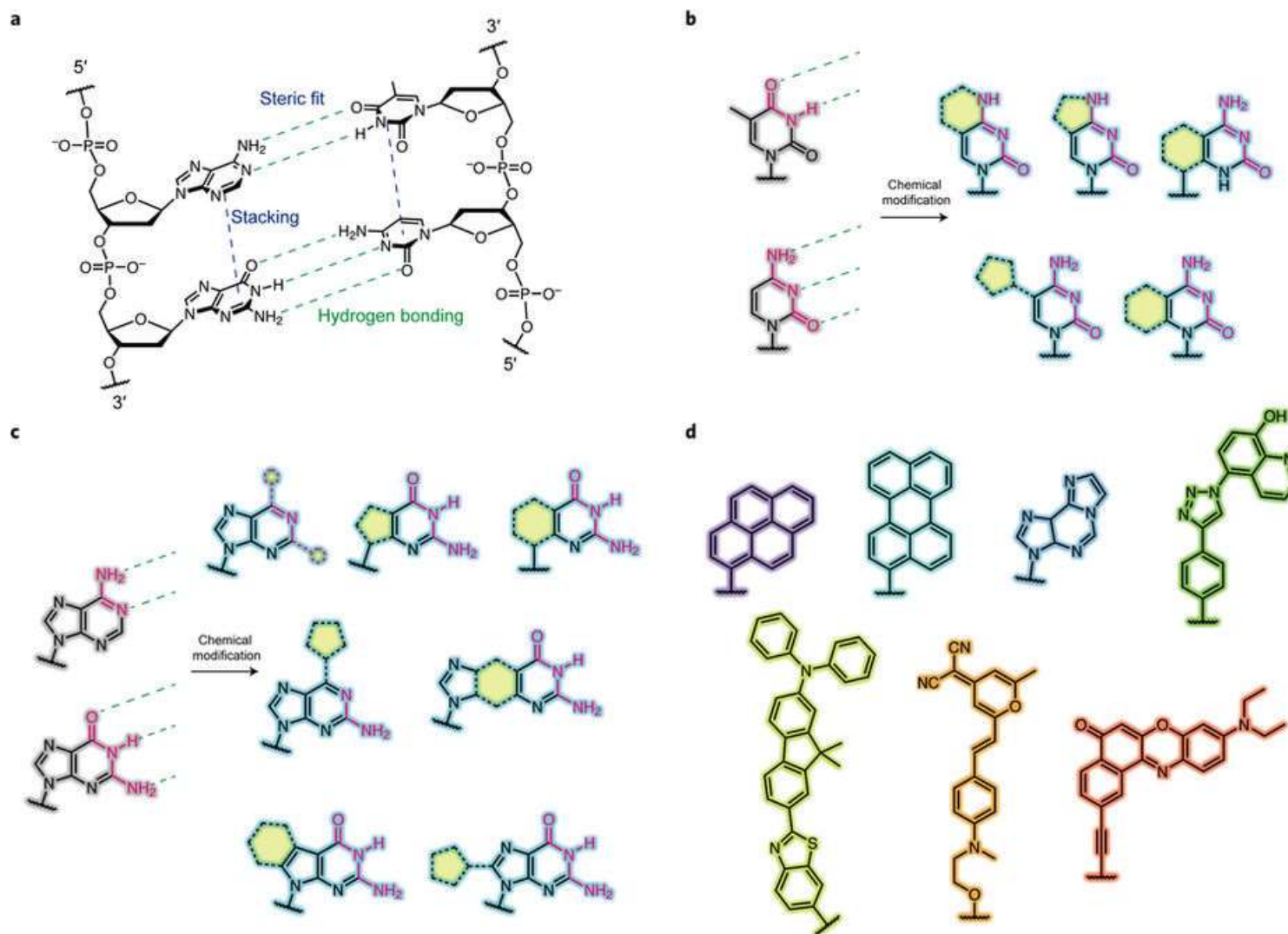
EcoI – a typical restriction enzyme

Products of restriction enzymes



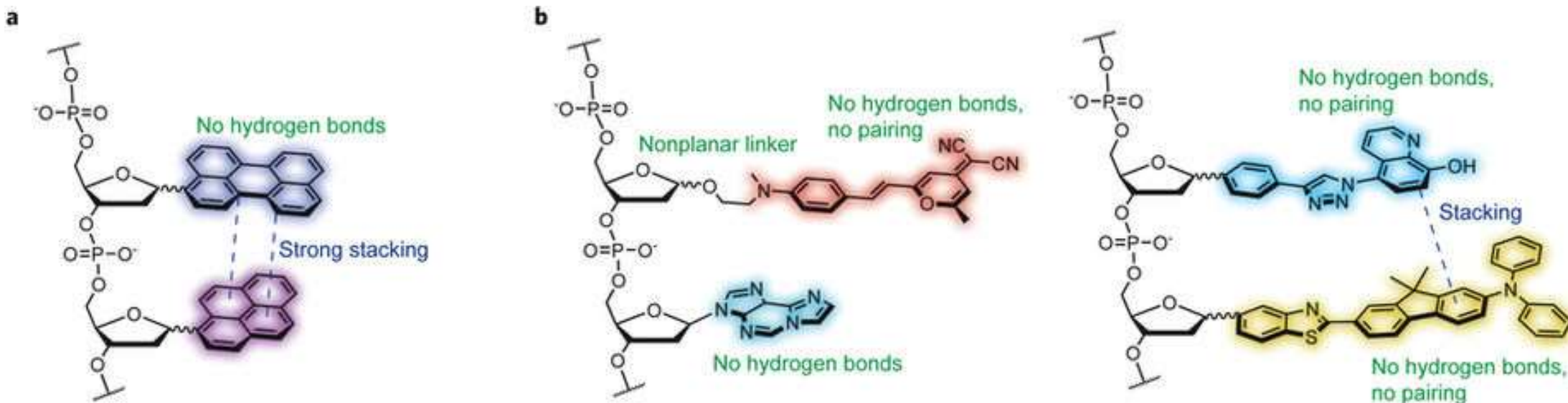
Structure of the homodimeric restriction enzyme EcoRI (cyan and green cartoon diagram) bound to double stranded DNA (brown tubes). Two catalytic magnesium ions (one from each monomer) are shown as magenta spheres and are adjacent to the cleaved sites in the DNA made by the enzyme (depicted as gaps in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

Non-canonical fluorescent nucleobases



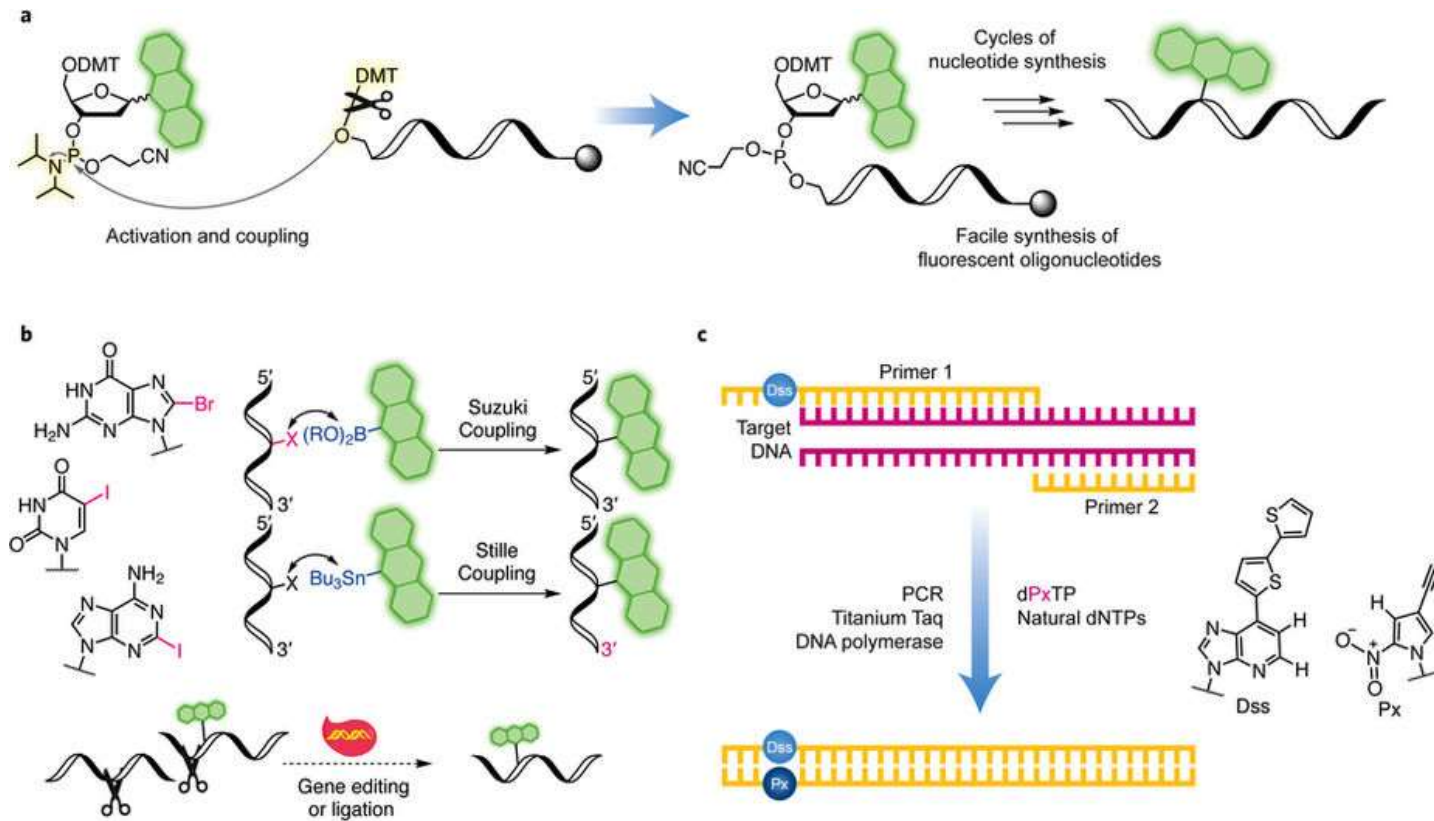
W. Xu, K. M. Chan, E. T. Kool *Nature Chem.* **2017**, *9*, 1043-1055

Non-canonical fluorescent nucleobases



a, Nucleobases composed of aromatic hydrocarbons. **b**, Nucleobases composed of planar heterocyclic fluorophores. The lack of hydrogen bonding and weaker π -stacking are compensated by versatile energy states brought from the heteroatoms. These fluorophores contribute to a broader spectrum of emission wavelengths. Functional groups can be added to expand functionality, such as metal binding.

Strategies for incorporating fluorescent nucleobases

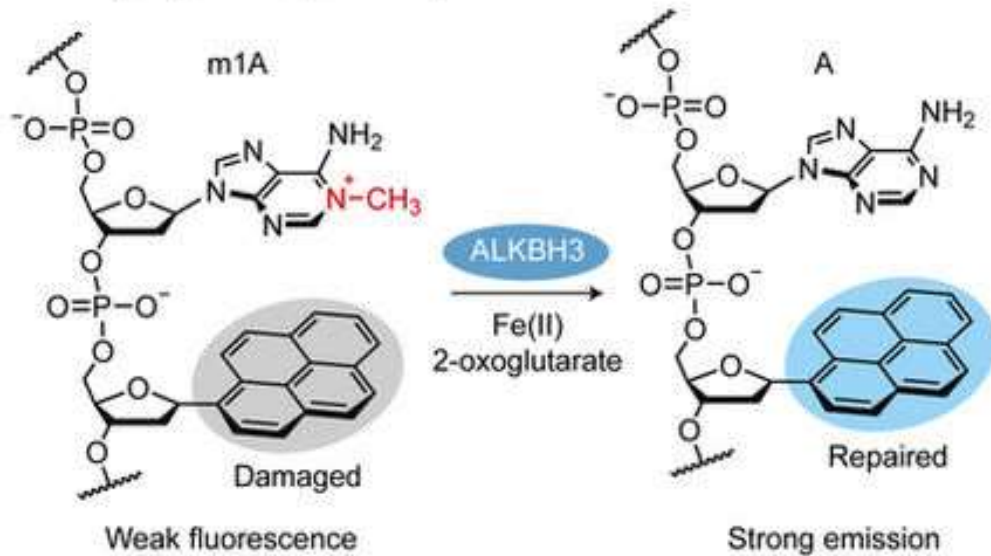


a, Direct oligonucleotide synthesis via synthesizer and phosphoramidite chemistry. **b**, Post-synthesis modification using mild coupling methods or gene-editing methods. Gene-editing and ligation methods enzymatically join smaller labelled strands to make longer ones. **c**, Direct enzymatic incorporation using fluorescent nucleoside triphosphate derivatives. When the fluorescent nucleobases are labelled in the primers or supplied as free nucleobases in the pool, polymerases that recognize them can incorporate the fluorescent nucleobases into DNA sequences.

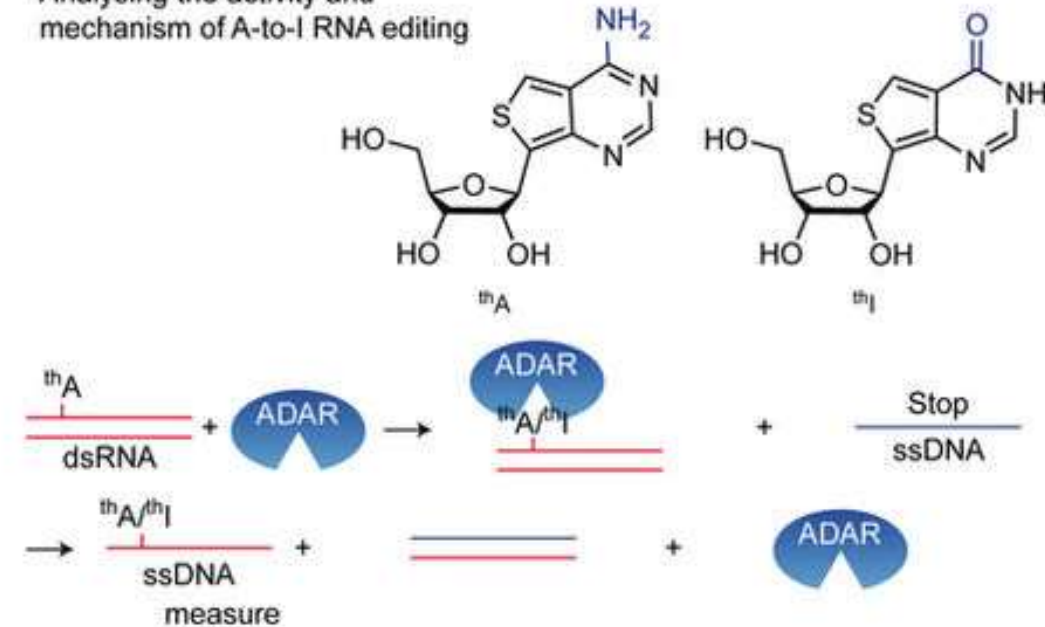
W. Xu, K. M. Chan, E. T. Kool *Nature Chem.* **2017**, *9*, 1043-1055

Fluorescent nucleobases for studying DNA and RNA

a Probing alkylation repair activity



b Analysing the activity and mechanism of A-to-I RNA editing



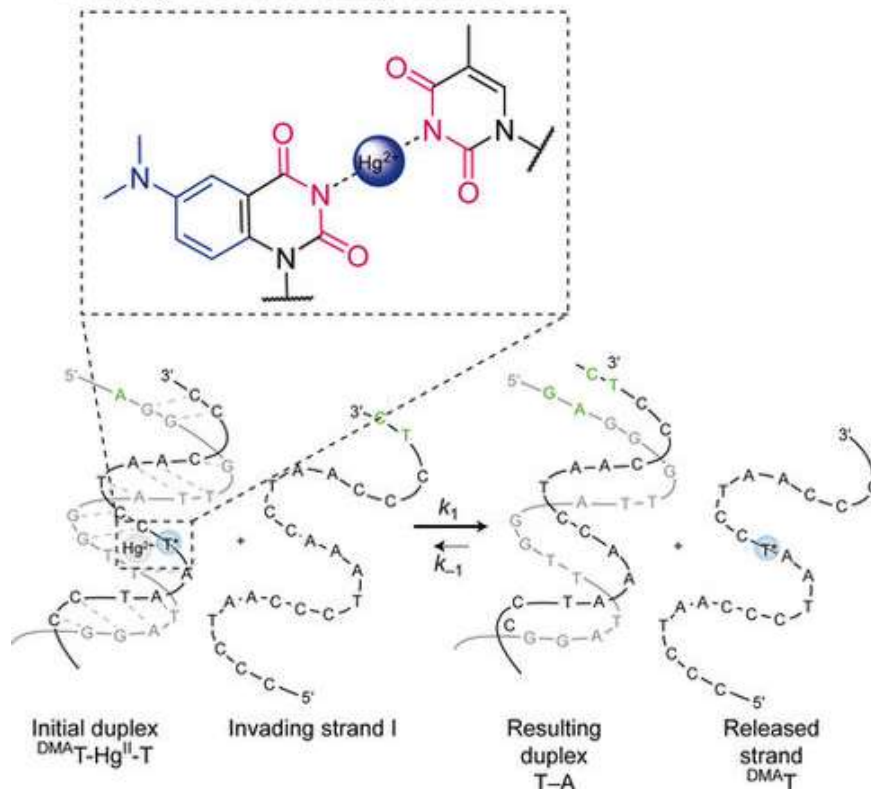
a, Fluorogenic sensing of a demethylation enzyme, ALKBH3. The emission of pyrene is initially quenched by the positive charge of 1-methylated adenine (m1A). When ALKBH3 demethylates m1A, the quenching effect is removed and a signal is generated.

b, Fluorogenic analysis of adenine-to-inosine RNA editing enzyme. The emission maxima of the thiolated adenine (thA) and inosine (thI) are different. Hence by measuring the intensity of thA and thI at their respective maximal wavelengths, the activity of the A-to-I enzyme can also be measured.

W. Xu, K. M. Chan, E. T. Kool *Nature Chem.* **2017**, *9*, 1043-1055

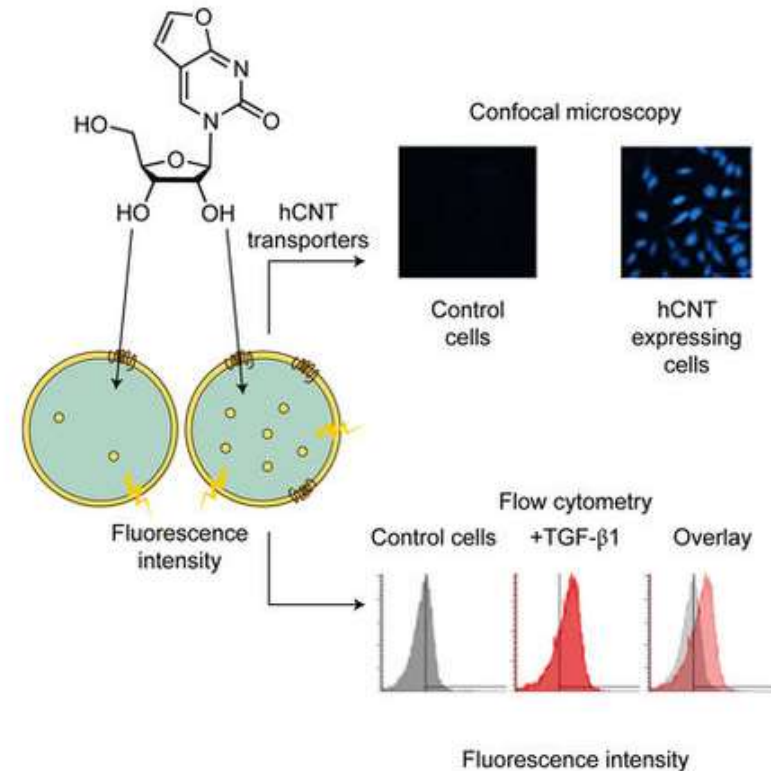
Fluorescent nucleobases for studying DNA and RNA

c Analysing the effects of mercury metabolism



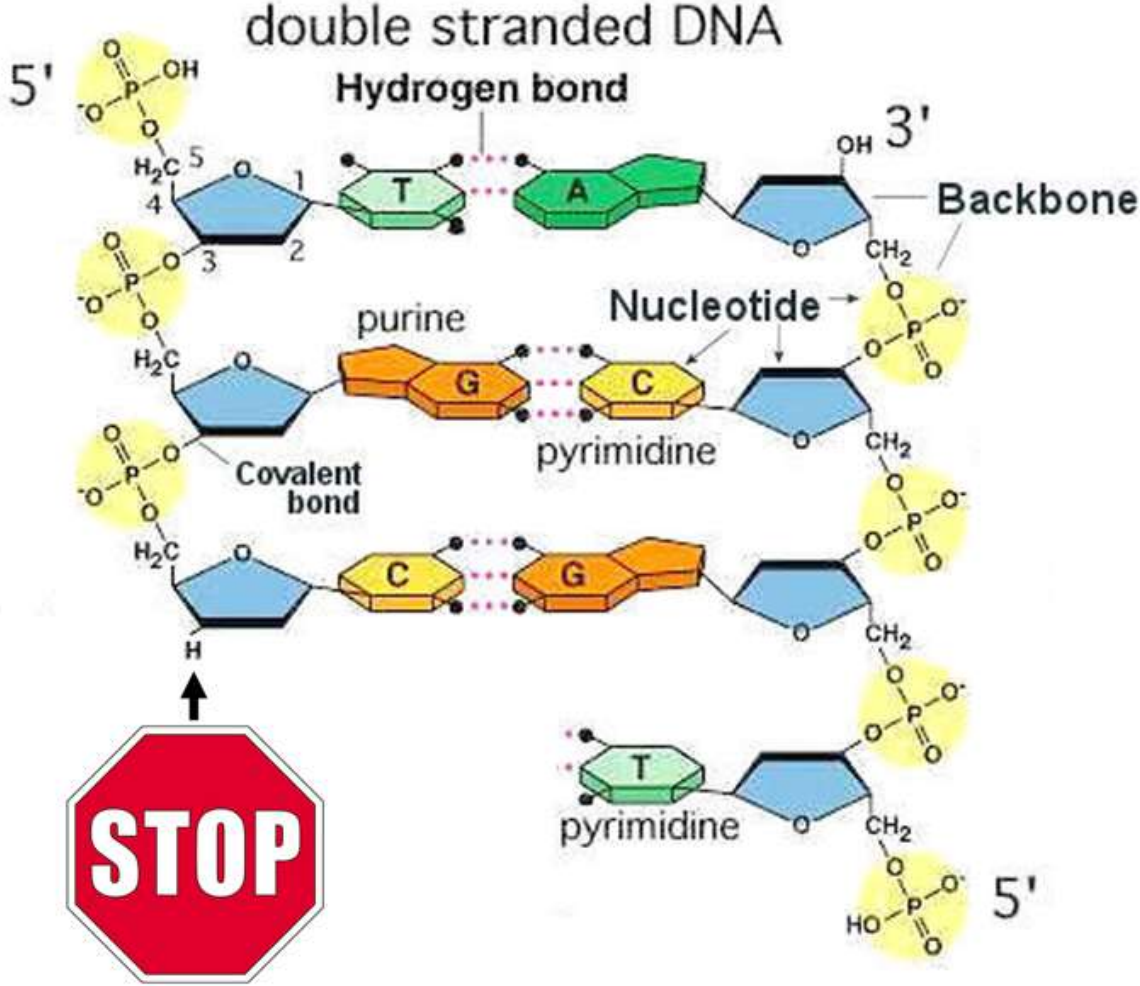
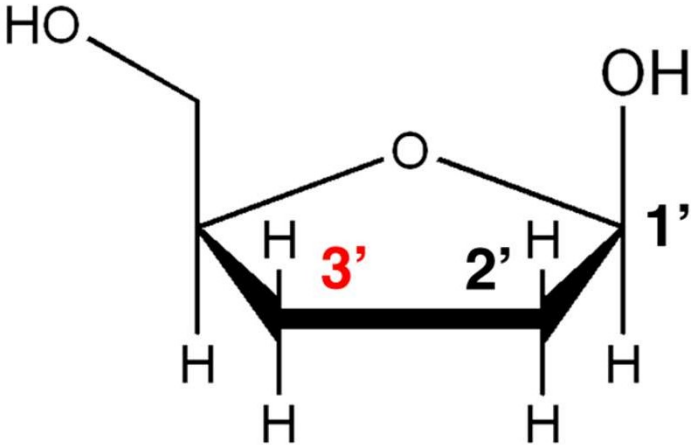
c, Kinetic and thermodynamic investigation of the effects of mercury on DNA metabolism. The fluorescent thymine can chelate mercury with another thymine ring and link DNA strands. This can be used to probe mercury metabolism *in vivo* and to study the effects of mercury on DNA status.

d Probing nucleoside transporter activity *in vivo*

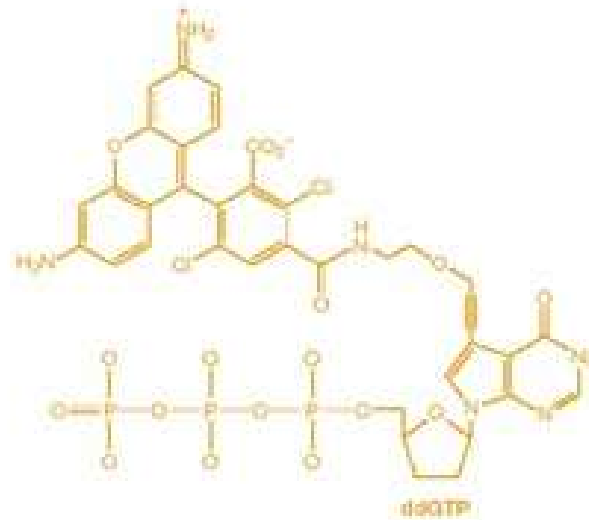
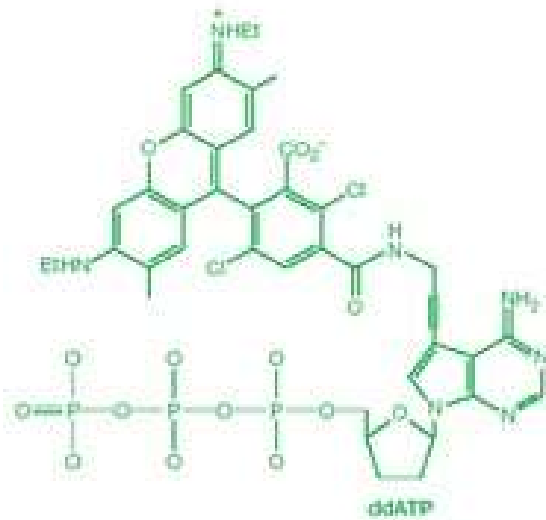
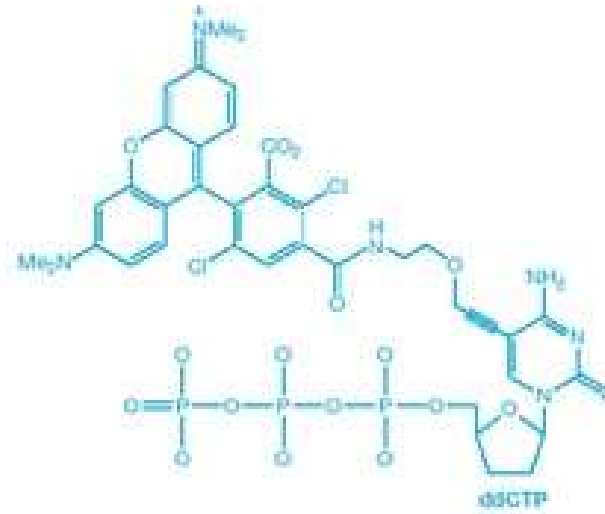
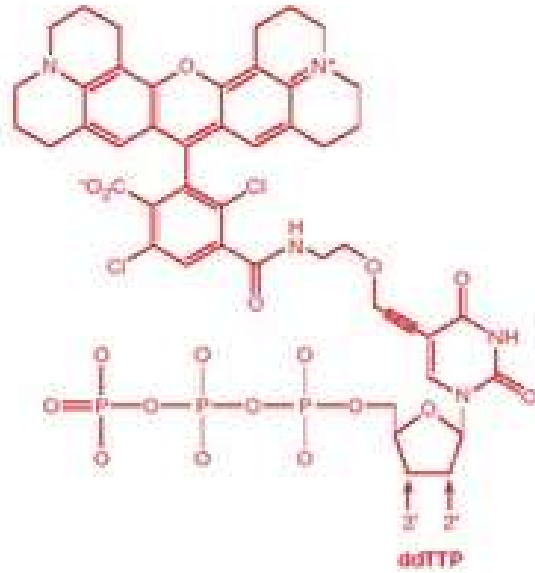


d, Visualization and analysis of human concentrative nucleoside transporters (hCNTs). The fluorescent nucleoside can enter the plasma membrane through the transporters, thus allowing the measurement of the transport activity. TGF-β1, transforming growth factor β1. [W. Xu, K. M. Chan, E. T. Kool *Nature Chem.* 2017, 9, 1043-1055](#)

Sanger sequencing



Sanger sequencing



Sanger sequencing

primer

5' 3'

TACGT

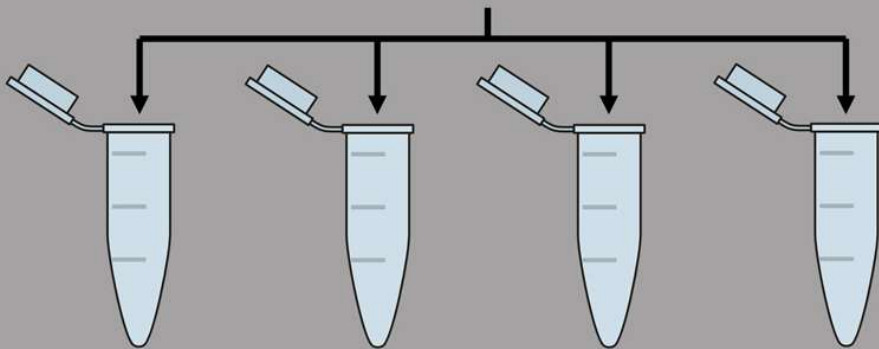
ATGCATTAGGGCCTGGCTCTTT

3'

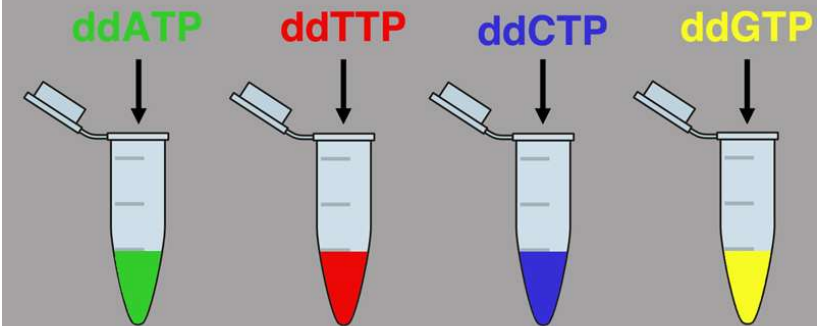
5'

template

DNA template + primer
DNA polymerase
dNTPs



Sanger sequencing



Sanger sequencing

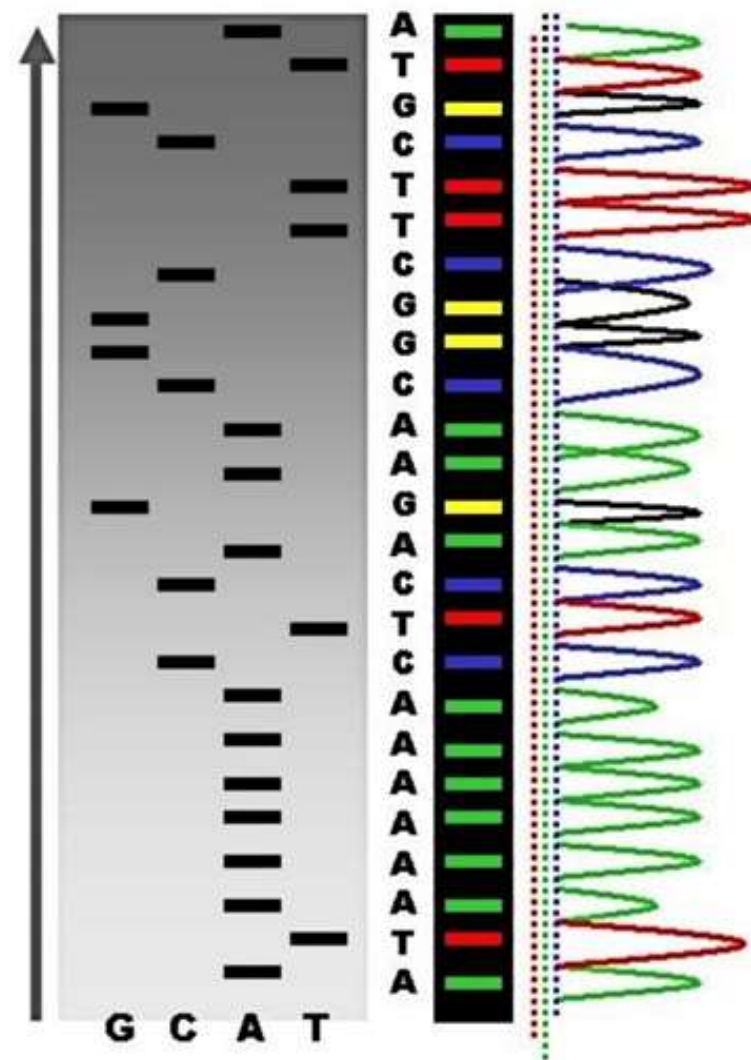
TACGTA**A**
ATGCATTAGGGCCTGGCTCTTT

TACGTA**A**
ATGCATTAGGGCCTGGCTCTTT

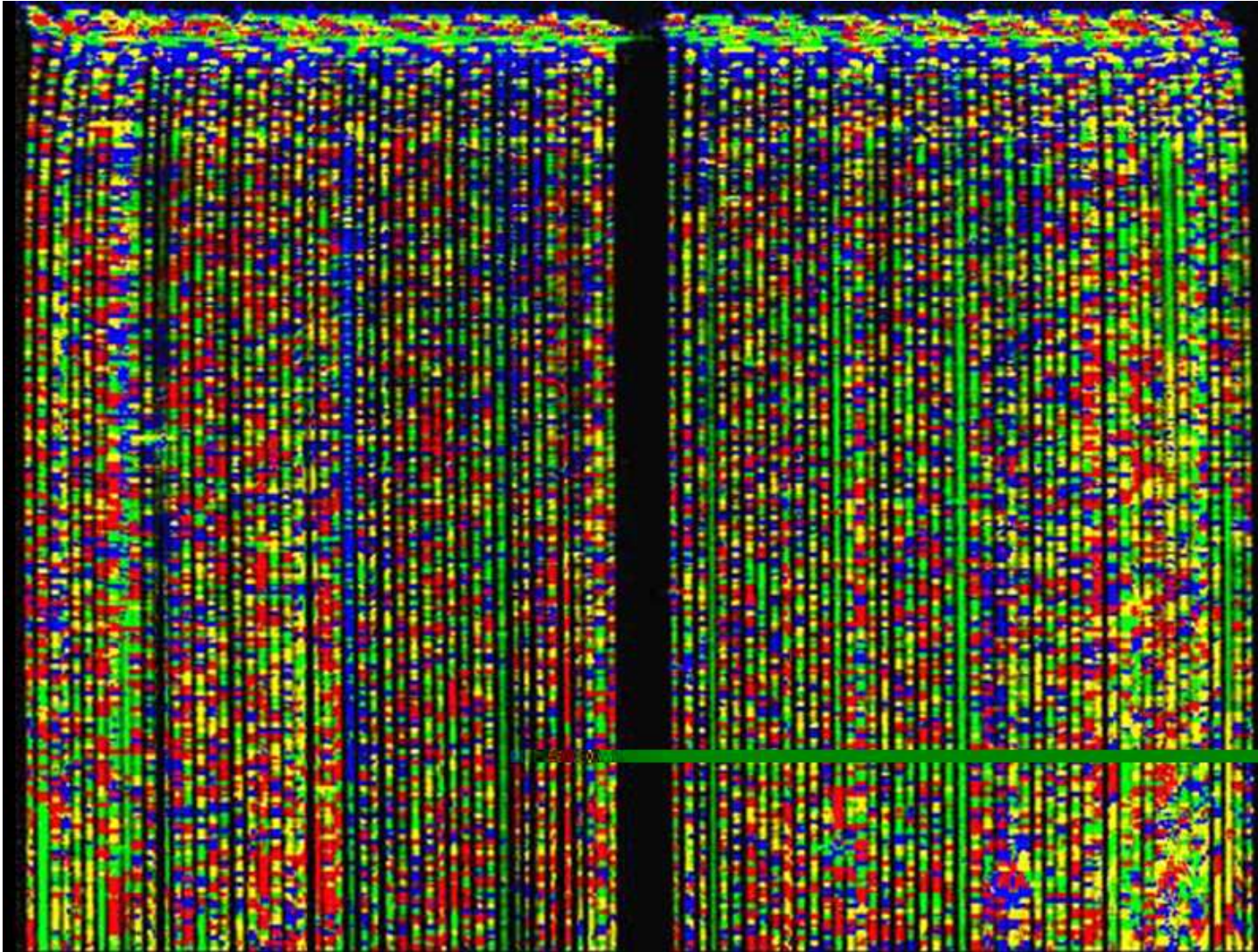
TACGTAATCCCGG**A**
ATGCATTAGGGCCTGGCTCTTT

TACGTAATCCCGGACCG**A**
ATGCATTAGGGCCTGGCTCTTT

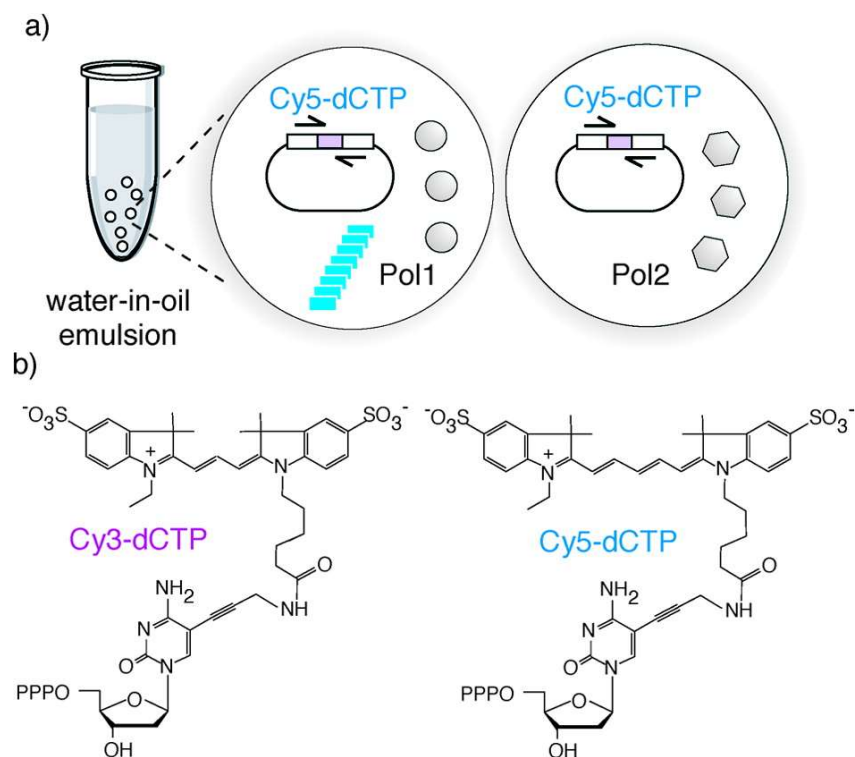
Sequencing Gel



Sanger sequencing



CyDNA – synthesis and replication of highly fluorescently-labelled DNA



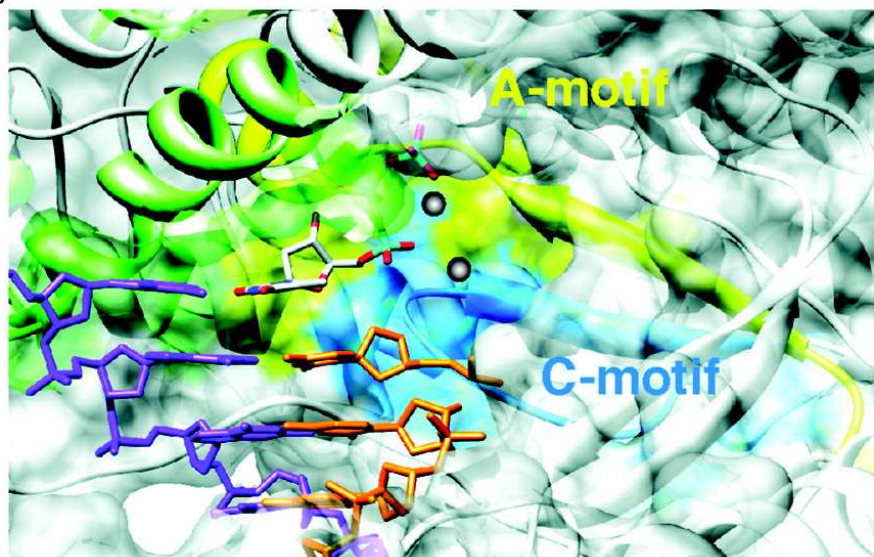
DNA Polymerase is evolutionarily optimized by mutagenesis to accept fluorescent Cy3 or Cy5 nucleotides instead of simple dCTP. Here, the polymerase replicates a short segment of its own encoding gene. Water/oil compartmentalization allows testing many independent mutations in parallel

Polymerases (Pol1 (left compartment)) that are capable of utilizing Cy5-dCTP are able to replicate, i.e., produce “offspring”, while polymerases like Pol2 (right compartment) that are unable to utilize it disappear from the gene pool.

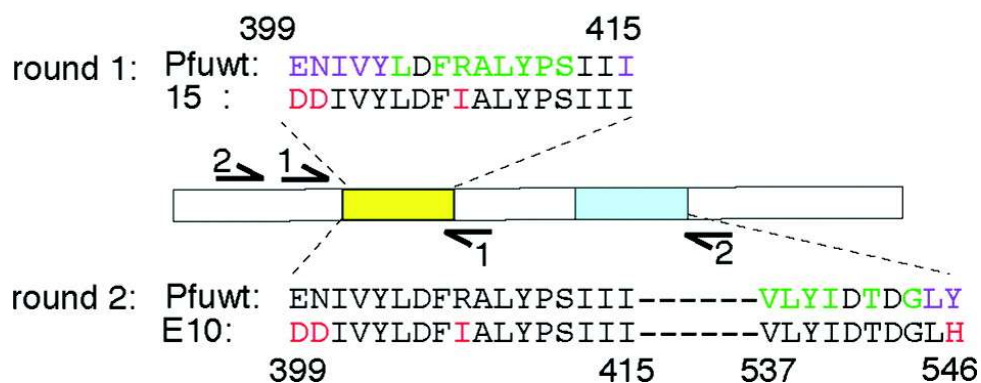
P. Holliger *et al.* *J. Am. Chem. Soc.* **2010**, *132*, 5096-5104

CyDNA – synthesis and replication of highly fluorescently-labelled DNA

a)



b)



(a) Structural model of the active site of a polB-family polymerase (RB69, PDB: 1IG9). The polypeptide chain is shown as a ribbon diagram overlaid with a transparent surface model. Primer and template strands are shown in orange and purple, incoming nucleotide triphosphate in elemental colors, and gray spheres represent the two catalytic Mg^{2+} ions. Regions corresponding to the A-motif are colored yellow, and those for the C-motif cyan. Also shown is the conserved B-motif (green).

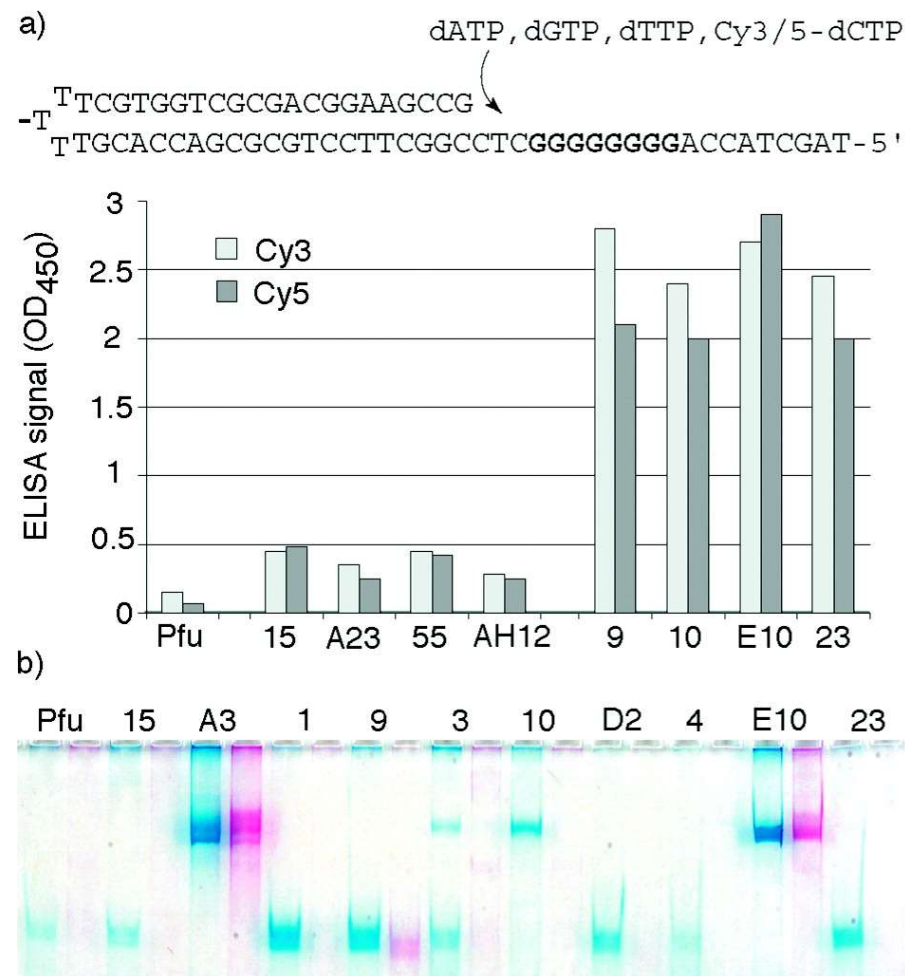
(b) For round 1 selection, diversity was focused on the A-motif and vicinity (399–415) comprising random mutation spike at the core of the A-motif (lime) as well as phylogenetic diversity in the adjacent sequences (purple). For round 2, successful clones from round 1 (e.g., 15) were diversified in the C-motif region (cyan), and selected for replication of A- and C-motif (399–546) yielding polymerase E10 (selected mutations in red)

CyDNA – synthesis and replication of highly fluorescently-labelled DNA

380 selected variants from round 1 were screened by Polymerase-ELISA and ranked for their ability to incorporate 4 consecutive Cy5-dCTPs or Cy3-dCTPs. Polymerase-ELISA identified 4 mutant polymerases with significantly enhanced ability to incorporate either Cy3-dCTPs or Cy5-dCTPs compared with wild-type Pfuexo-:

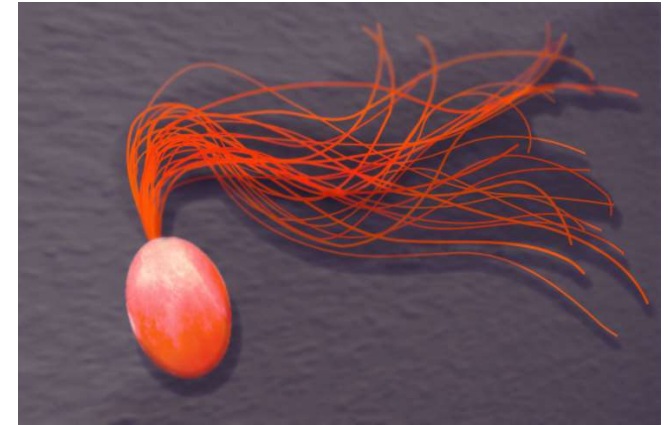
- A23 (N400D, I401L, R407I),
- AH12 (E399D, N400G, I401L, V402A, R407I, Q572H),
- 55 (N400G, R407I), and in particular
- 15 (V337I, E399D, N400G, R407I).

- a) activities of round 1 clones (15, A23, 55, AH12) and round 2 clones (9, 10, E10, 23). Clones were chosen i.a. on the their ability to incorporate both Cy3- and Cy5-dCTPs with comparable efficiency.
- b) PAGE gel of a 0.4kb PCR amplification (70% GC) with complete replacement of dCTP with either Cy5- or Cy3-dCTP comparing selected clones to wild-type Pfuexo- (Pfu). Only E10 (and to a lesser extent A3) are able to perform PCR amplification with both dyes.



CyDNA – synthesis and replication of highly fluorescently-labelled DNA

The polymerase fidelity in the selected mutant „E10” was not significantly compromised as compared to the starting Pfu DNA polymerase (*Pyrococcus furiosus*)



Fulvio314

Pyrococcus furiosus – an extremophilic *Archaeon* from marine sediments
Optimal life temperature 100°C

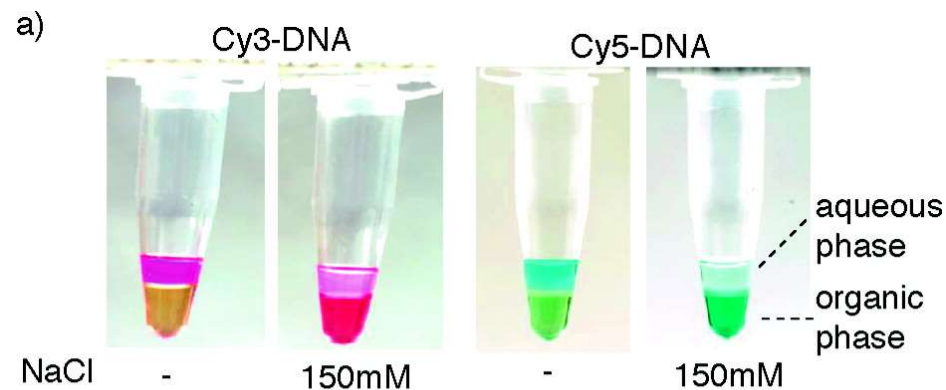
Table 1. Polymerase Fidelity

polymerase	PCR substrates	mutations/kb	mutation rate/bp/doubling ^a
Pfuexo-	dNTPs	1.1	4.4×10^{-5} (4.7×10^{-5}) ^b
E10	dNTPs	0.4 (2.6 without additives ^c)	1.6×10^{-5} (1.04×10^{-4}) ^c
Pfuexo-	dNTPs ^b	2.7	6.0×10^{-5}
E10	dATP, dGTP, dTTP, Cy3-dCTP	4.3	9.6×10^{-5}
E10	dATP, dGTP, dTTP, Cy5-dCTP	4.9	1.1×10^{-4}

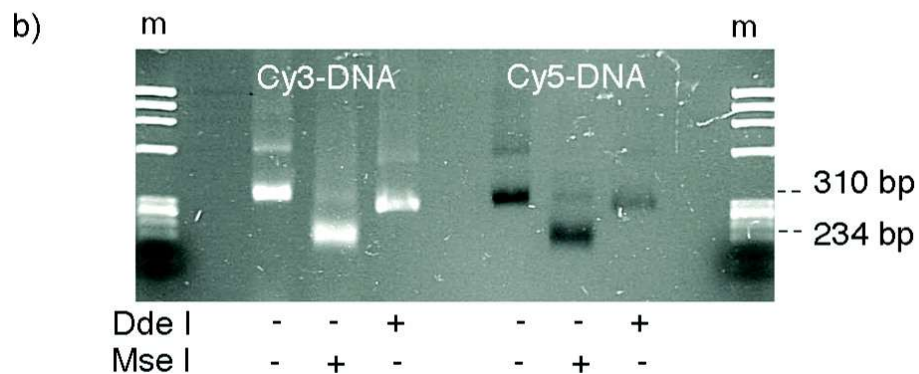
^a Corrected for the number of doublings (PCR cycles). ^b As determined by a lacZ reversion assay.⁴⁴ ^c In the absence of additives (1% formamide, 10% glycerol, 10 µg/mL RNase, 1 mM DTT).

P. Holliger *et al.* *J. Am. Chem. Soc.* **2010**, *132*, 5096-5104

CyDNA – synthesis and replication of highly fluorescently-labelled DNA



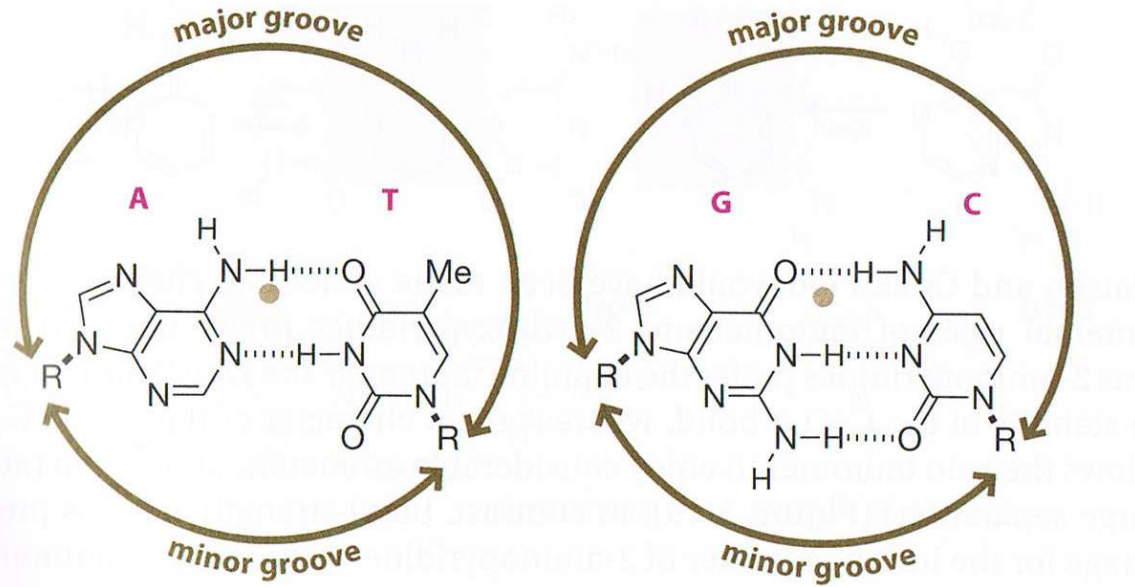
(a) Organic phase partitioning of CyDNA is shown for Cy3-DNA (left) and Cy5-DNA (right). Essentially 100% partitioning occurs in the presence of 150 mM NaCl (the yellow color of the phenol phase is due to addition of 8-hydroxyquinoline to prevent oxidation).



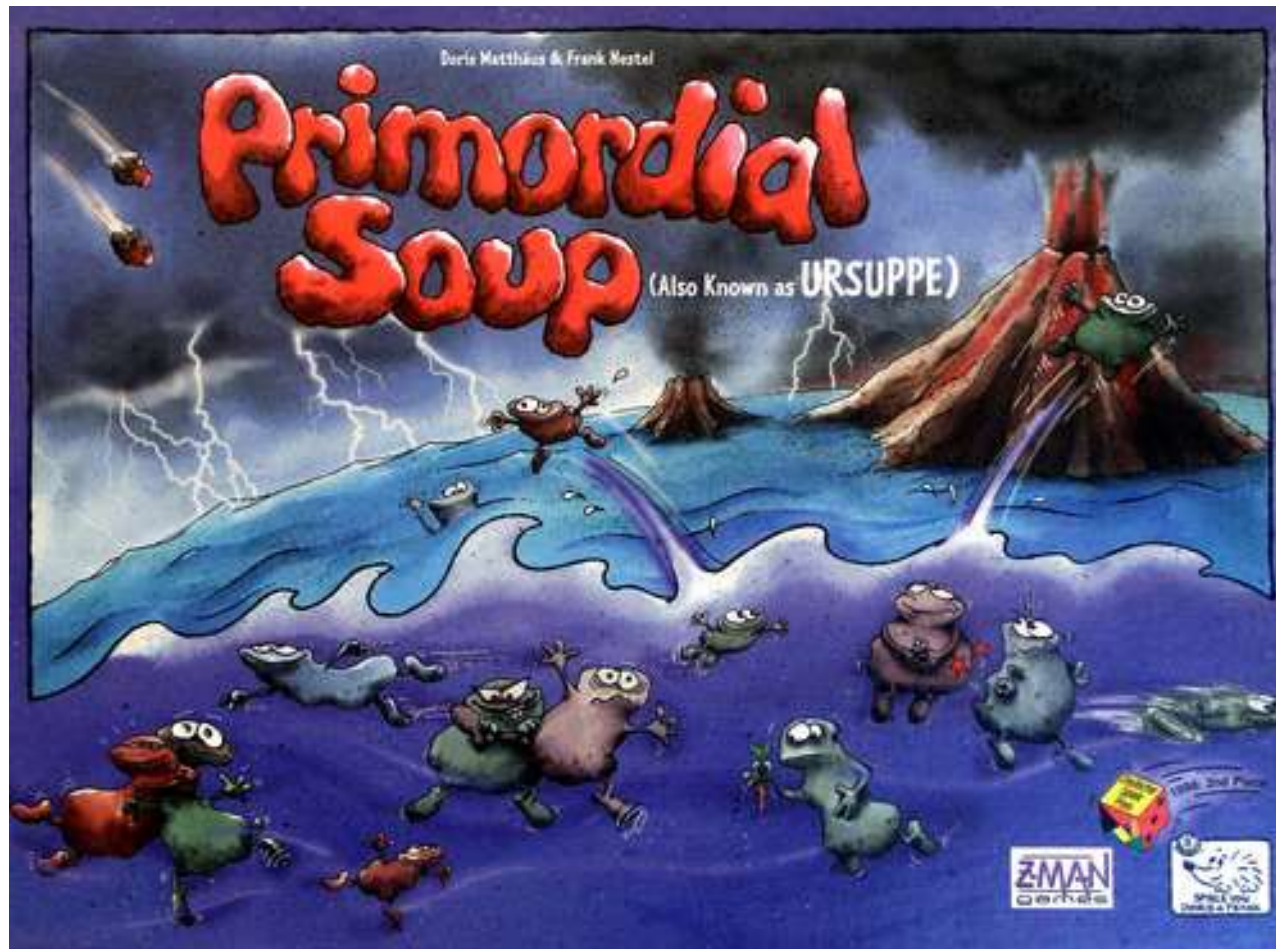
(b) Agarose gel electrophoresis of CyDNA restriction digests. Restriction endonucleases are sensitive probes of noncanonical DNA conformations such as those which occur under torsional strain. Both Cy3- and Cy5-DNA are resistant to cleavage by the restriction endonuclease DdeI (C'TNAG) but are cut by MseI (T'TAA). This indicates that at least the local regions of AT-sequence in Cy-DNA adopt a canonical B-form conformation.

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

	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



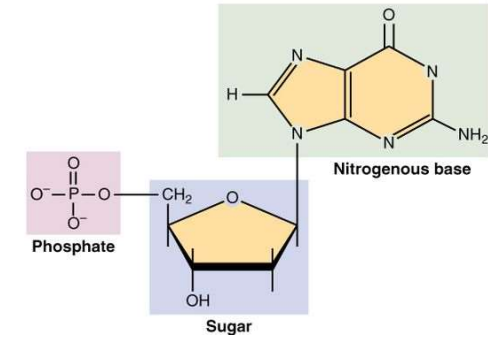
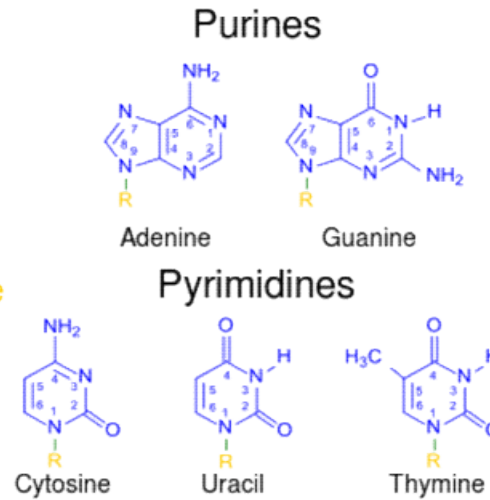
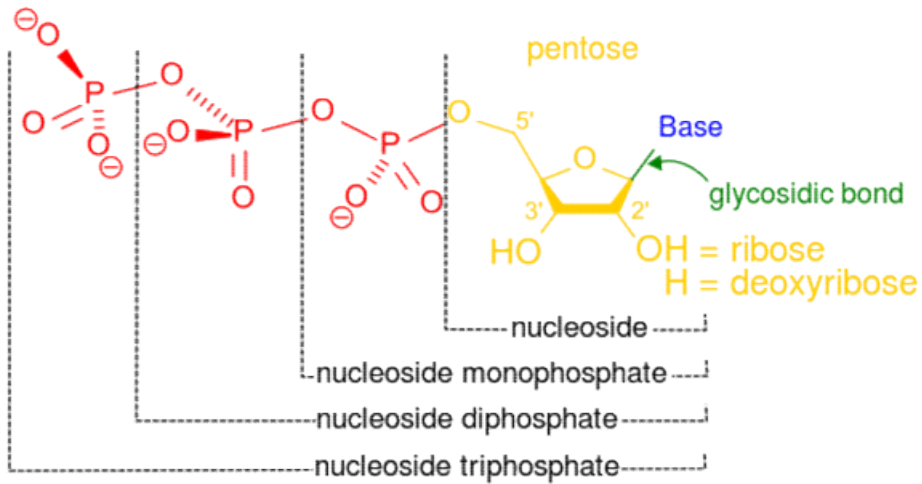
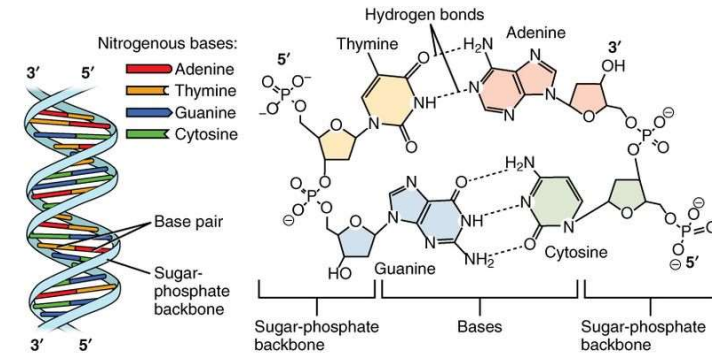
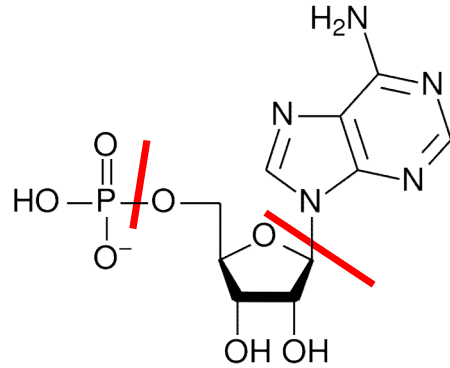
Prebiotic synthesis of nucleotides



State of the art

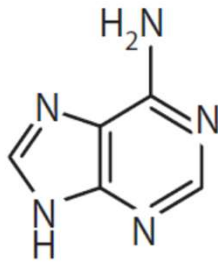
Nucleotides - components

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

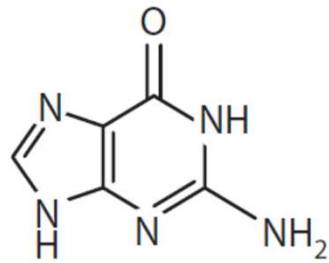


Prebiotic synthesis of nucleobases

Purines

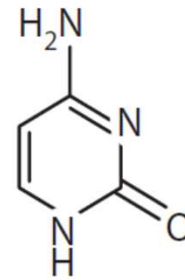


Adenine

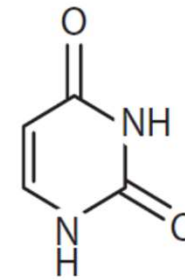


Guanine

Pyrimidines

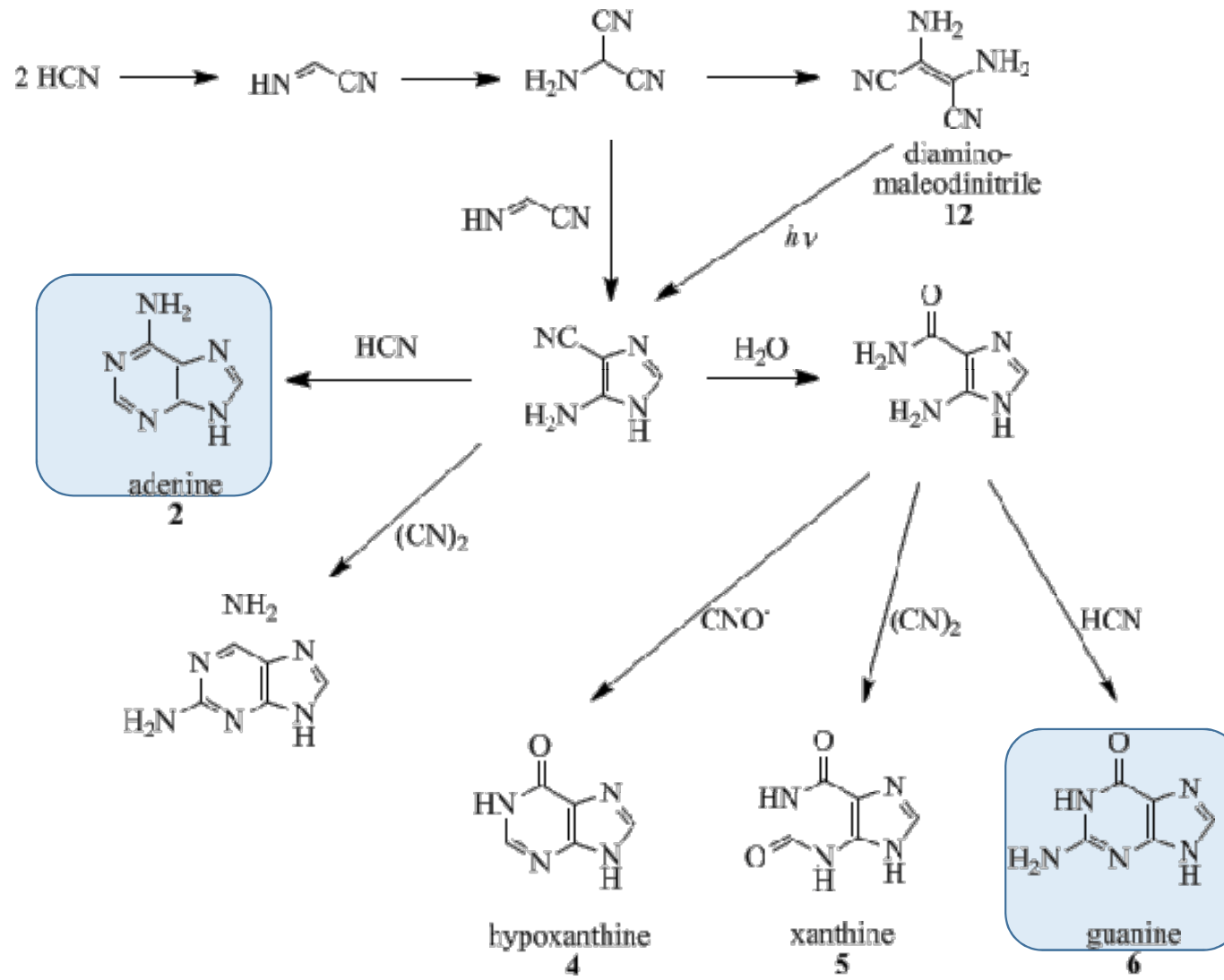


Cytosine



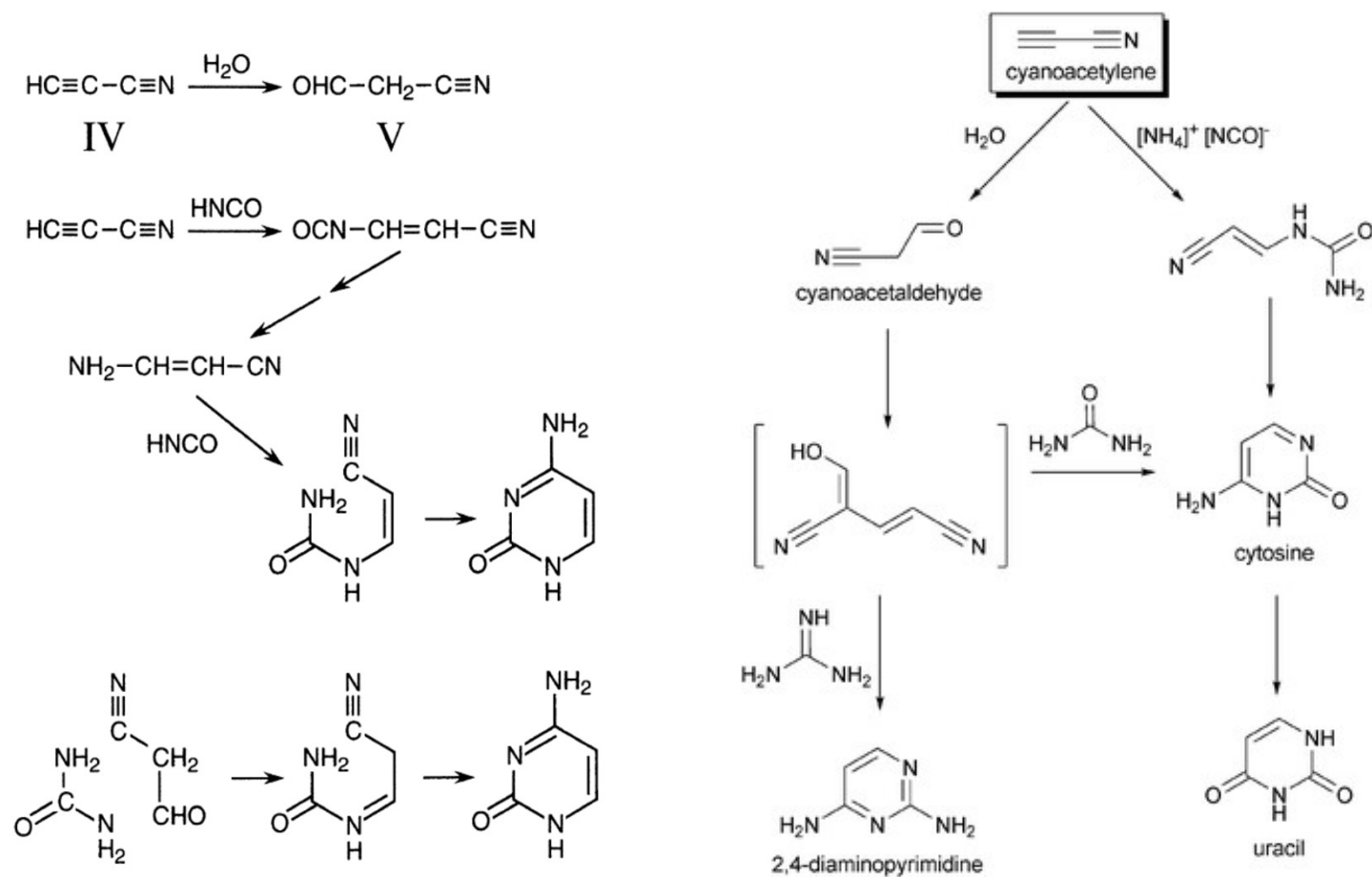
Uracil

Prebiotic synthesis of purines



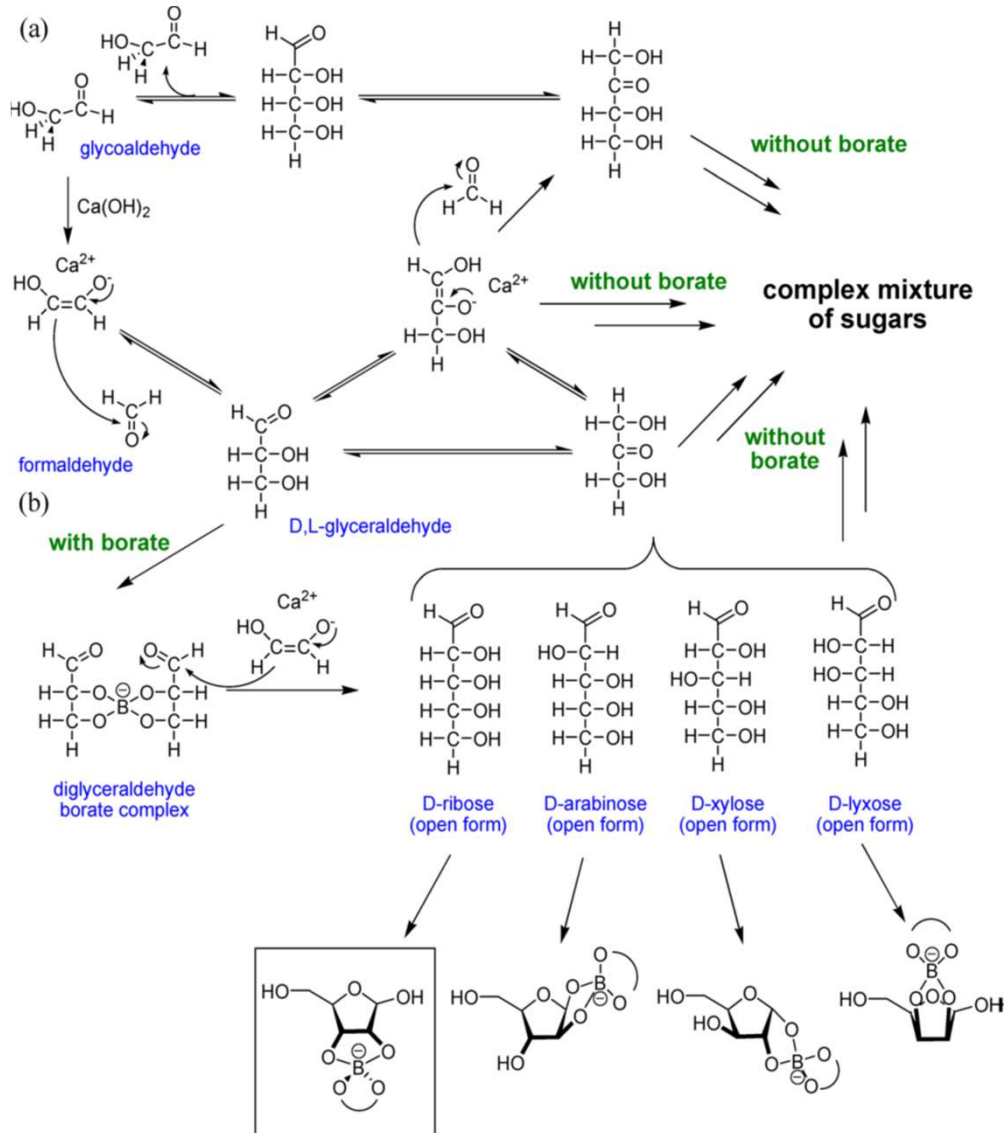
Prebiotic synthesis of pyrimidines

Cyanoacetylene is a major product of electric discharges in the mixture of nitrogen and methane



Cyanoacetylene incubated with saturated solution of urea yields up to 50% cytosine. Other methods typically yield up to 5% cytosine. It is further converted to uracil by hydrolysis.

Formose reaction in presence of borates

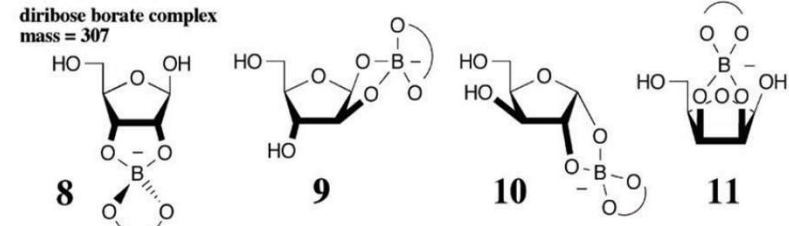


Pentose formation in the presence of borate

With borate (left)
Without borate (right)
Colemanite (background)



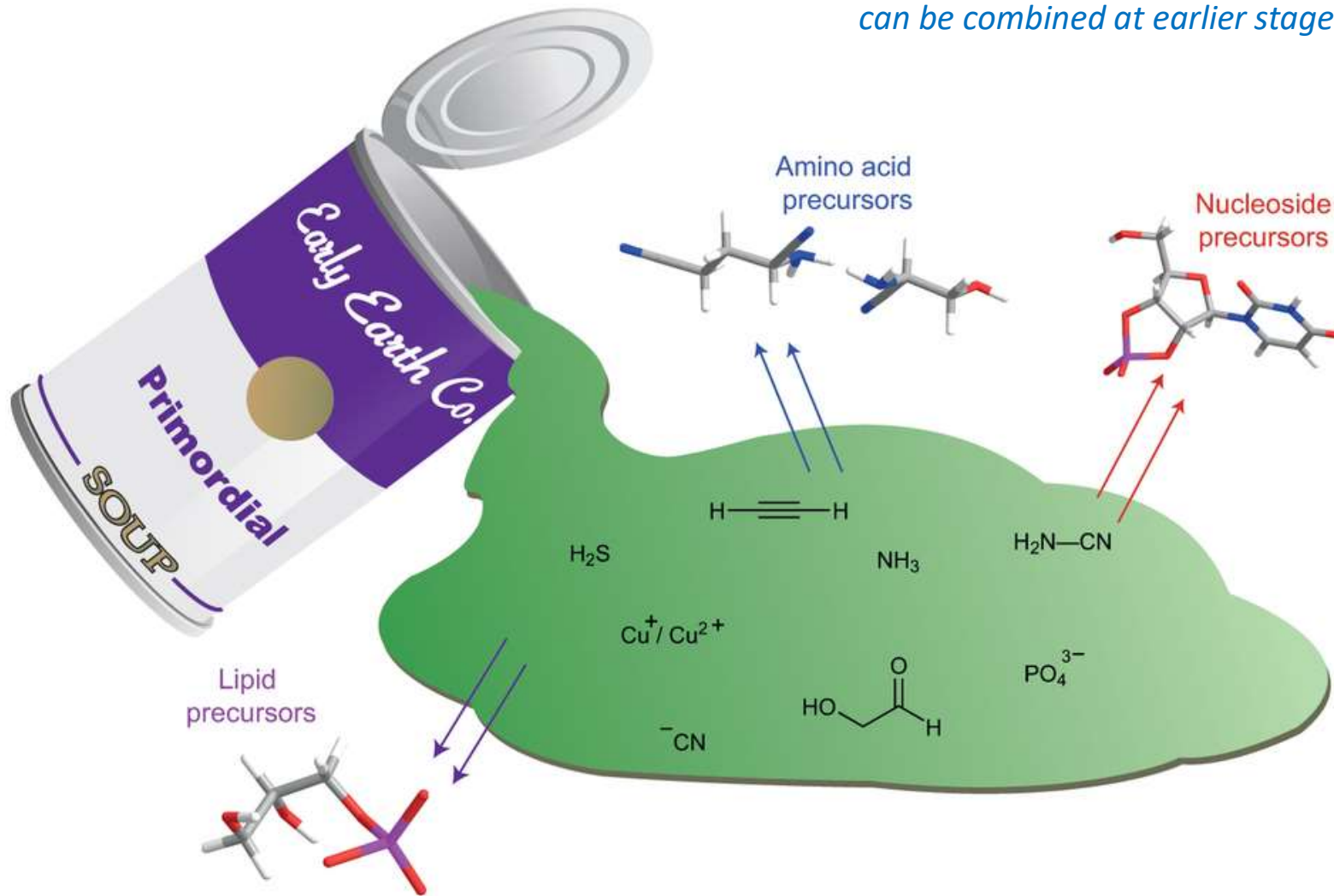
By NMR, the ribose borate complex **8** has the structure shown; cyclic structures for other pentoses are speculative.



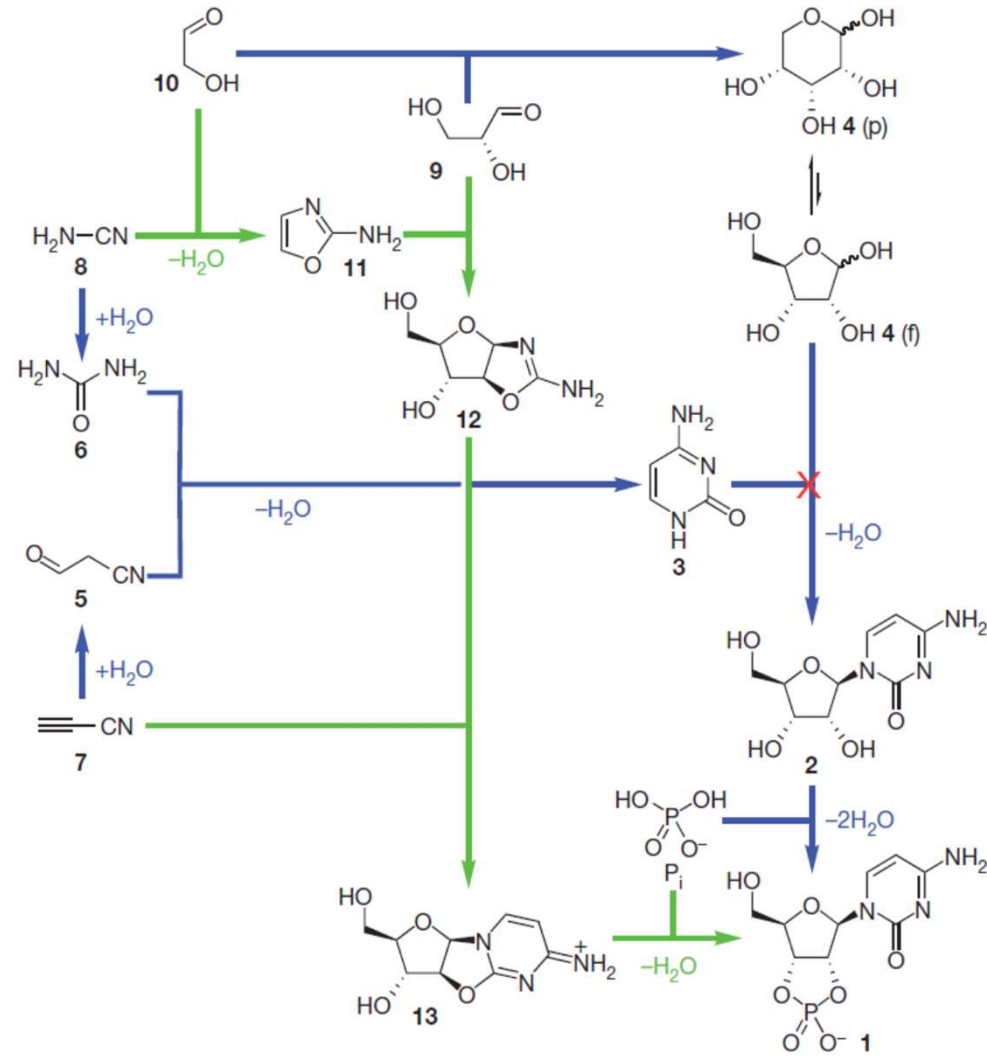
A. Ricardo, M. A. Carrigan, A. N. Olcott, S. A. Benner
Science **2004**, 303, 196

Cyanosulfidic chemistry

The aldol chemistry of sugars and cyanide chemistry of nucleobases can be combined at earlier stages than glycosylation.

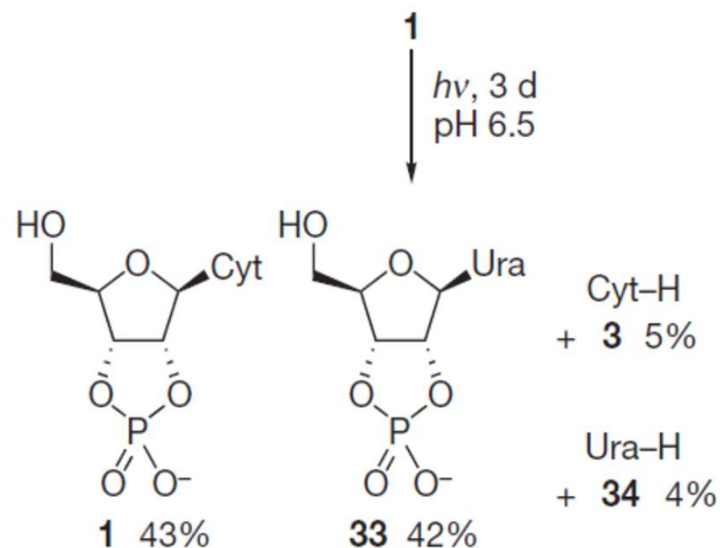


Cyanosulfidic chemistry



M. W. Powner, B. Gerland, J. D. Sutherland, *Nature* **2009**, *459*, 239–242

Cyanosulfidic chemistry



Photochemistry of *beta*-ribocytidine-2',3'-cyclic phosphate **1**. Under conditions of irradiation that destroy most other pyrimidine nucleosides and nucleotides, **1** undergoes partial hydrolysis and slight nucleobase loss. Ura, N1-linked uracil; Cyt-H, cytosine; Ura-H, uracil.

M. W. Powner, B. Gerland, J. D. Sutherland, *Nature* **2009**, *459*, 239–242

J. D. Sutherland, *Angew. Chem. Int. Ed.* **2016**, *55*, 104-121.

B. H. Patel, C. Percivalle, D. J. Ritson, C. D. Duffy, J. D. Sutherland, *Nat. Chem.* **2015**, *7*, 301–307.

J. D. Sutherland, *et al. Nat. Chem.* **2013**, *5*, 383–389.