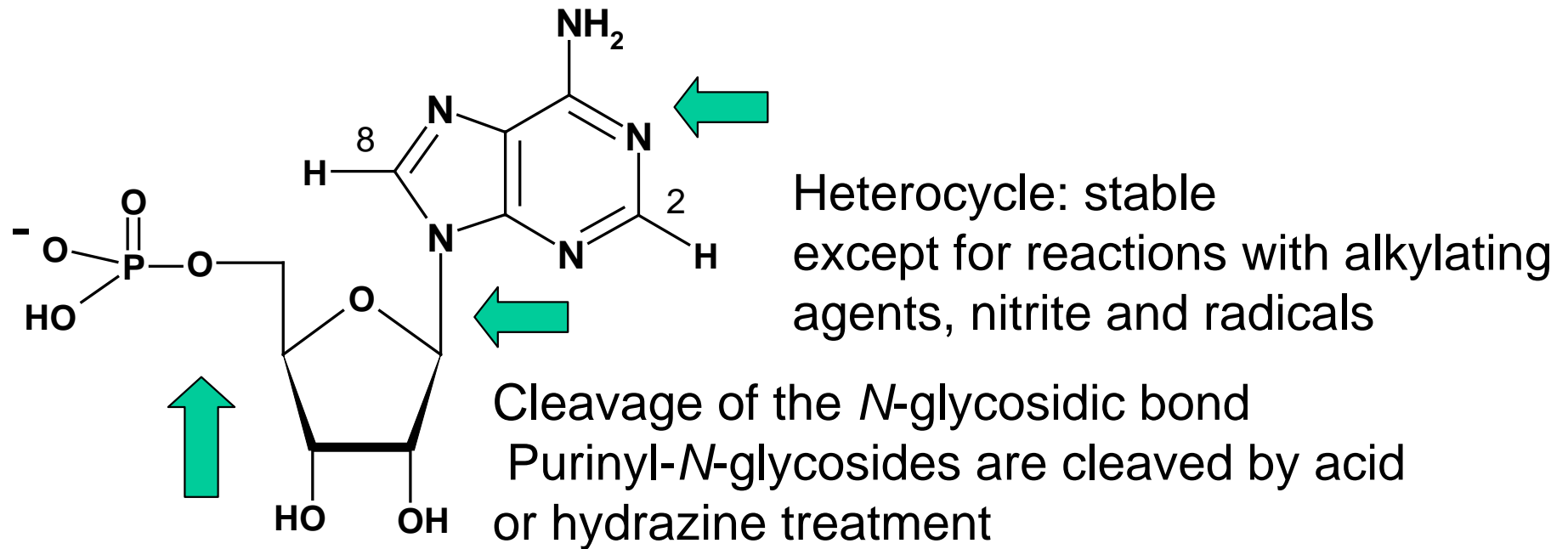


4. Chemical stability of nucleic acids



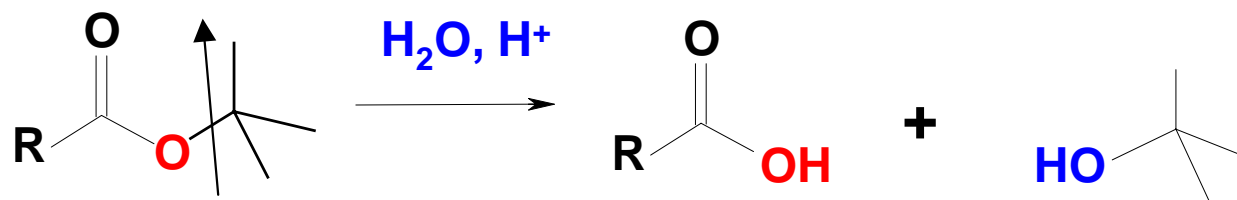
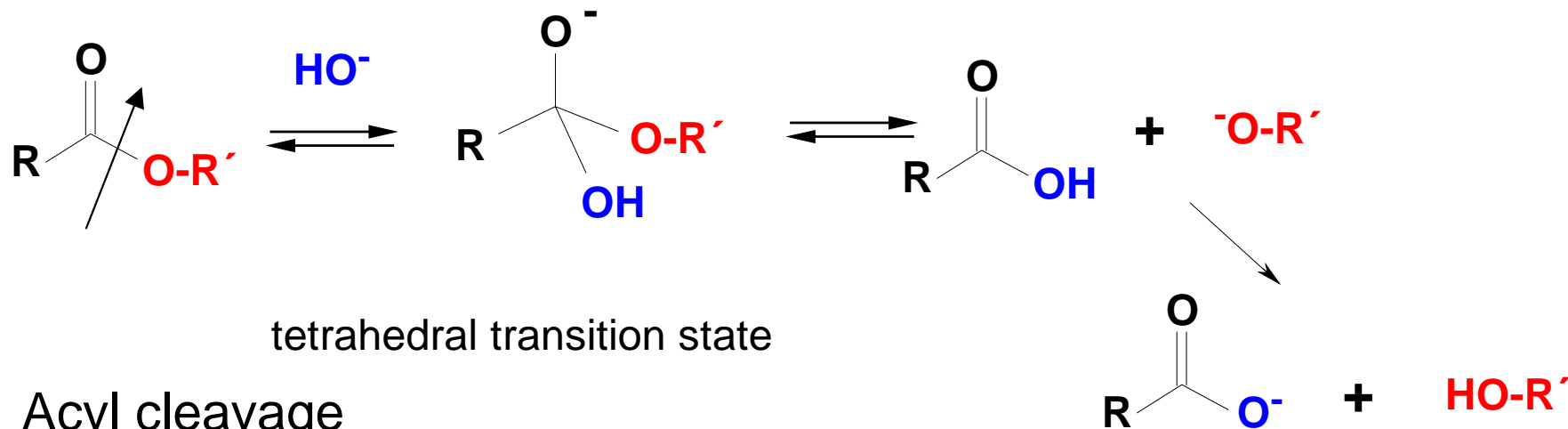
Cleavage of the phosphodiester linkage:

RNA: hydrolyzed by 0.3 M KOH

DNA: stable

Chemical hydrolysis of phosphoesters

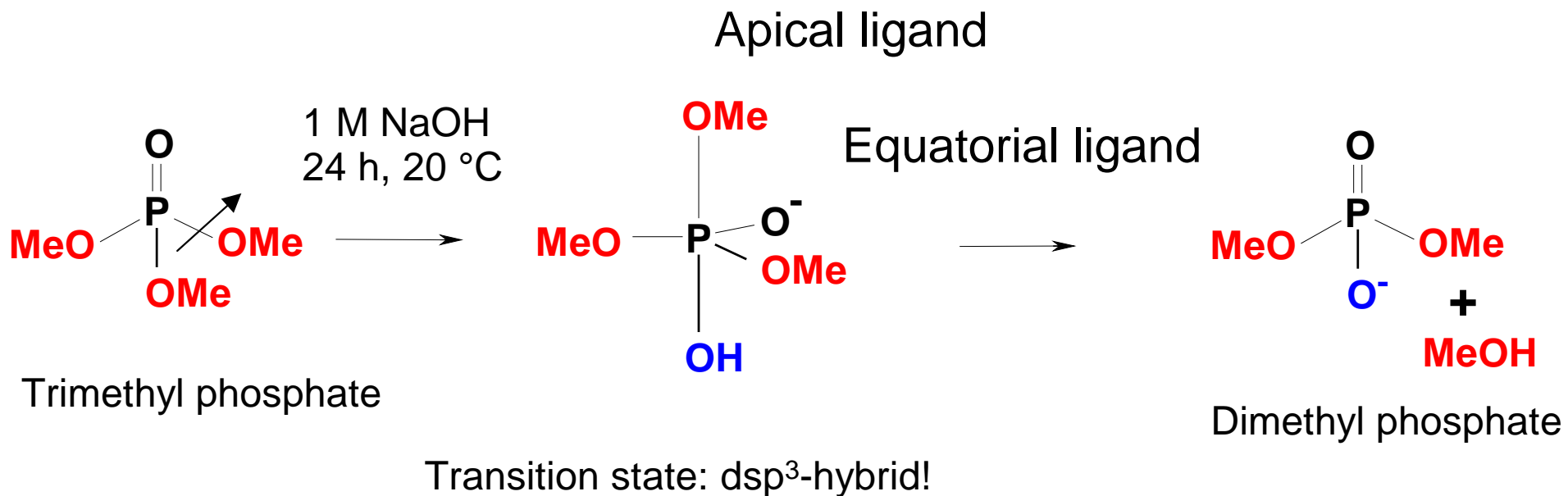
Comparison: Ester hydrolysis of carboxylic acid



Alkyl cleavage in specific cases only

Chemical hydrolysis of phosphoesters

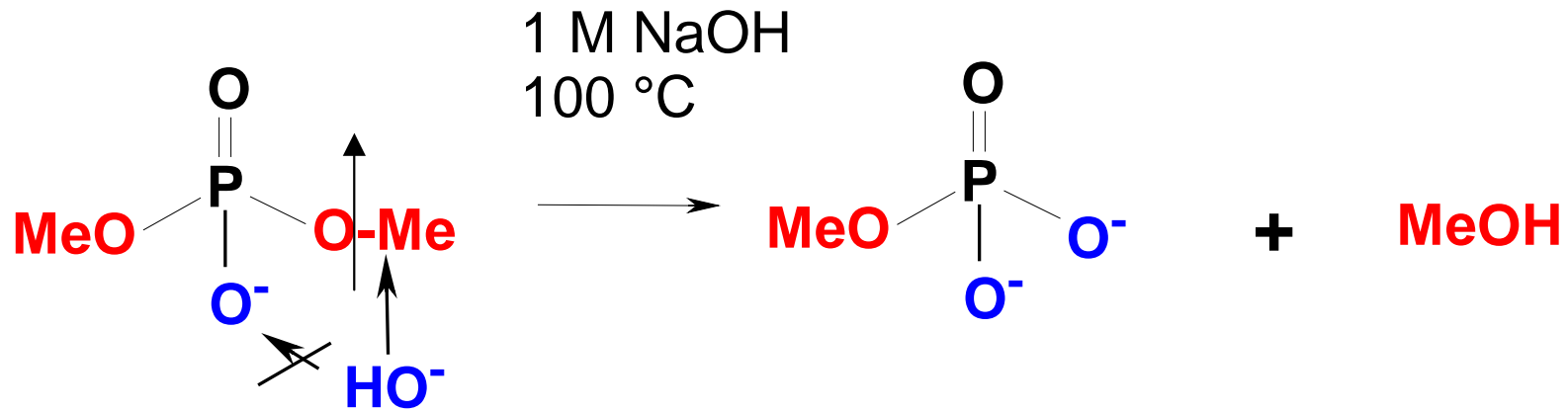
1. Phosphotriester



Phosphoryl cleavage

Chemical hydrolysis of phosphoesters

2. Phosphodiester



Dimethyl phosphate

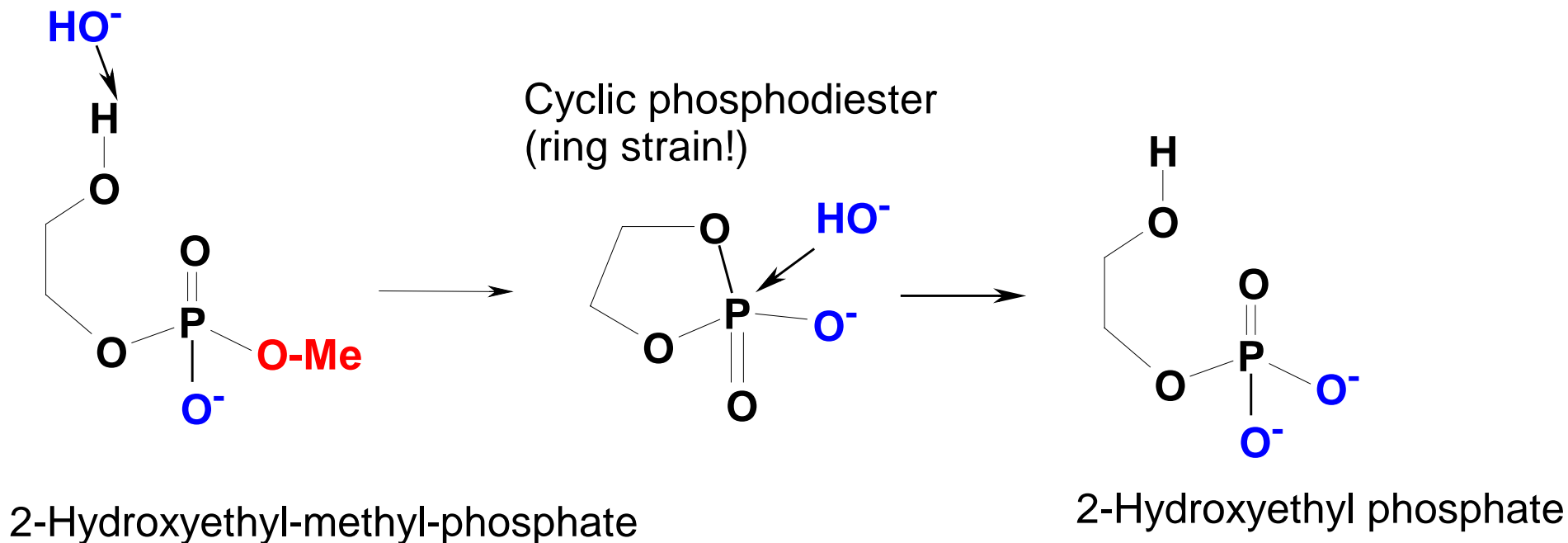
Methyl phosphate

Alkyl cleavage

Half life time: 16 days!

Chemical hydrolysis of phosphoesters

2.Phosphodiester

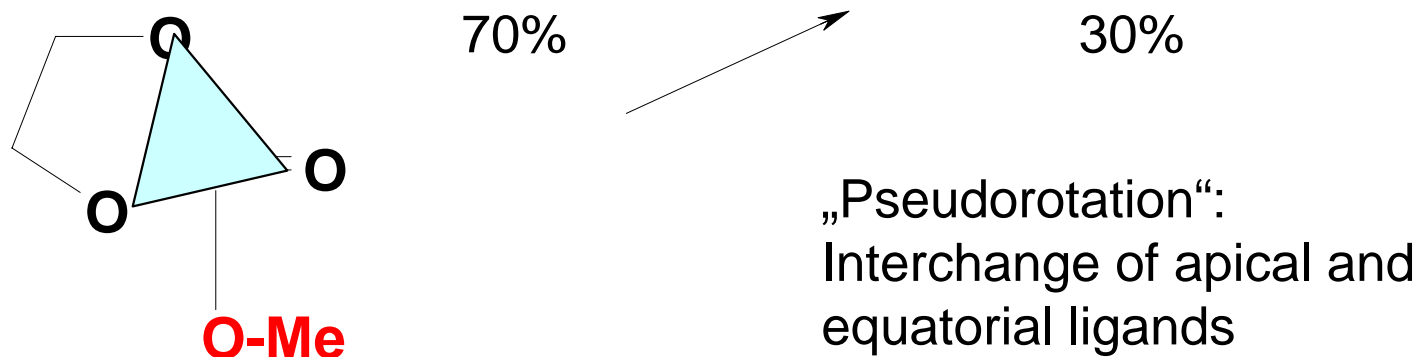
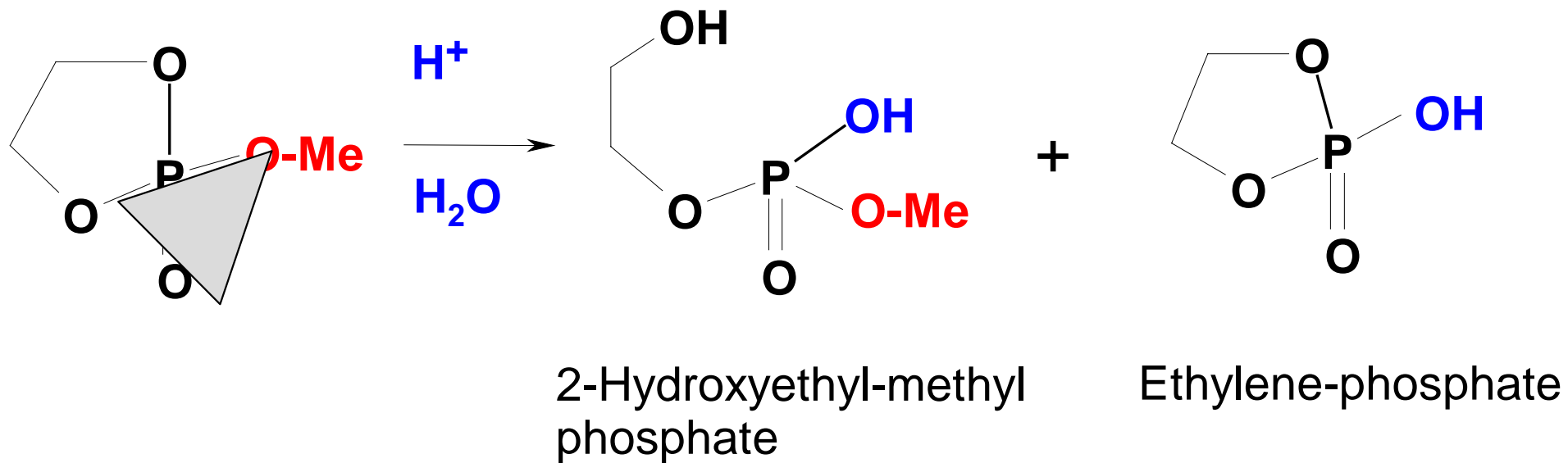


$$\Delta H = -26.8 \text{ kJ/mol}$$

Cf: Dimethyl phosphate: -7.5 kJ/mol

Half life time at 25°C 25 min!

Neighboring group participation
Anchimeric assistance

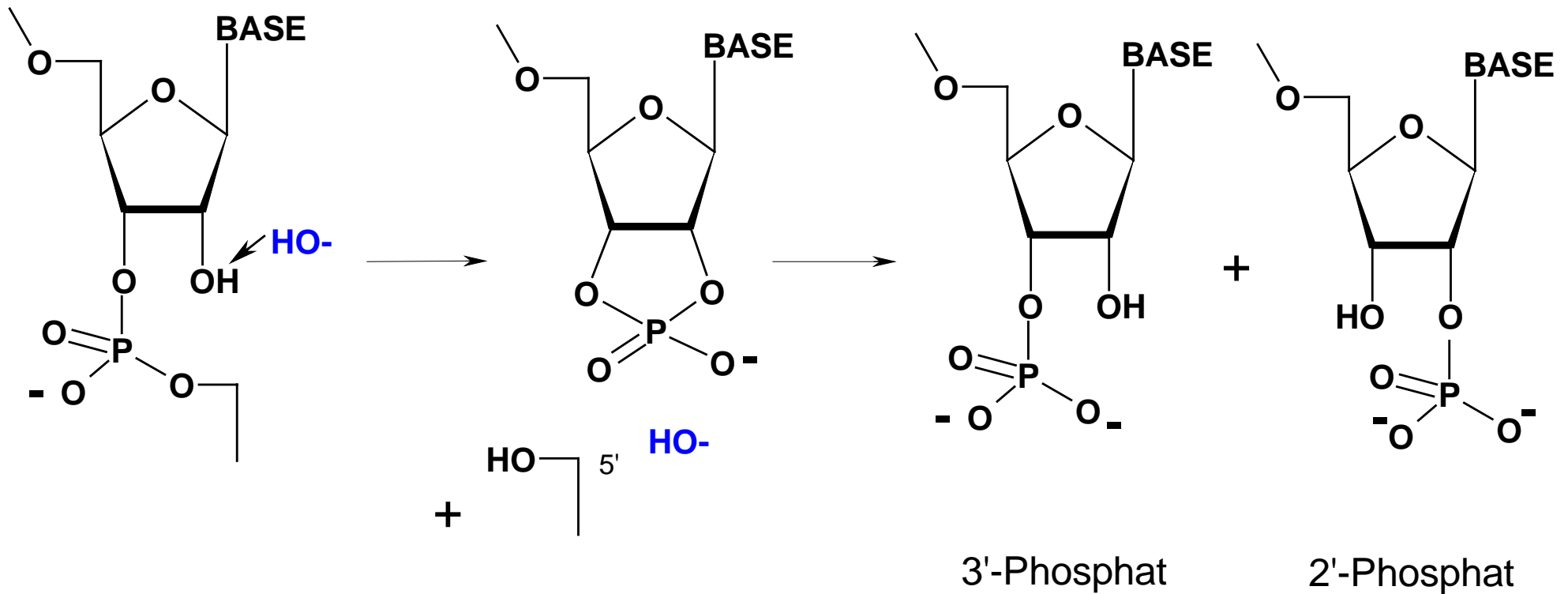


Leaving group may only be displaced from an apical position (longest bond)

Alkaline hydrolysis of DNA and RNA

DNA: stable (1 h at 100 °C, 1 M NaOH)

RNA: hydrolyzed at RT in 0.1 M NaOH



Enzymatic hydrolysis of RNA

Ribonucleases

Non-specific cleavage or specific cleavage of bases

Ribonuclease A: Pyrimidin-Nucleotides (C,U)

Ribonuclease T₁ (*Aspergillus oryzae*): Guanine (in *syn*-conformation)

Ribonuclease P (Ribozymes)

Ribonuklease A

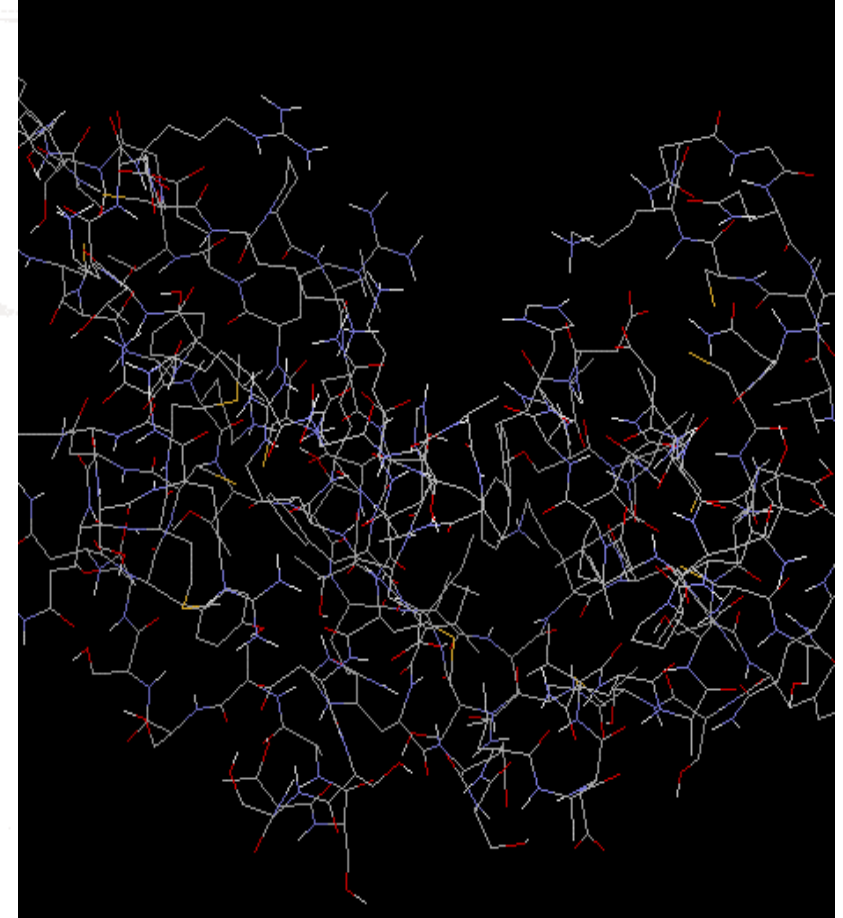
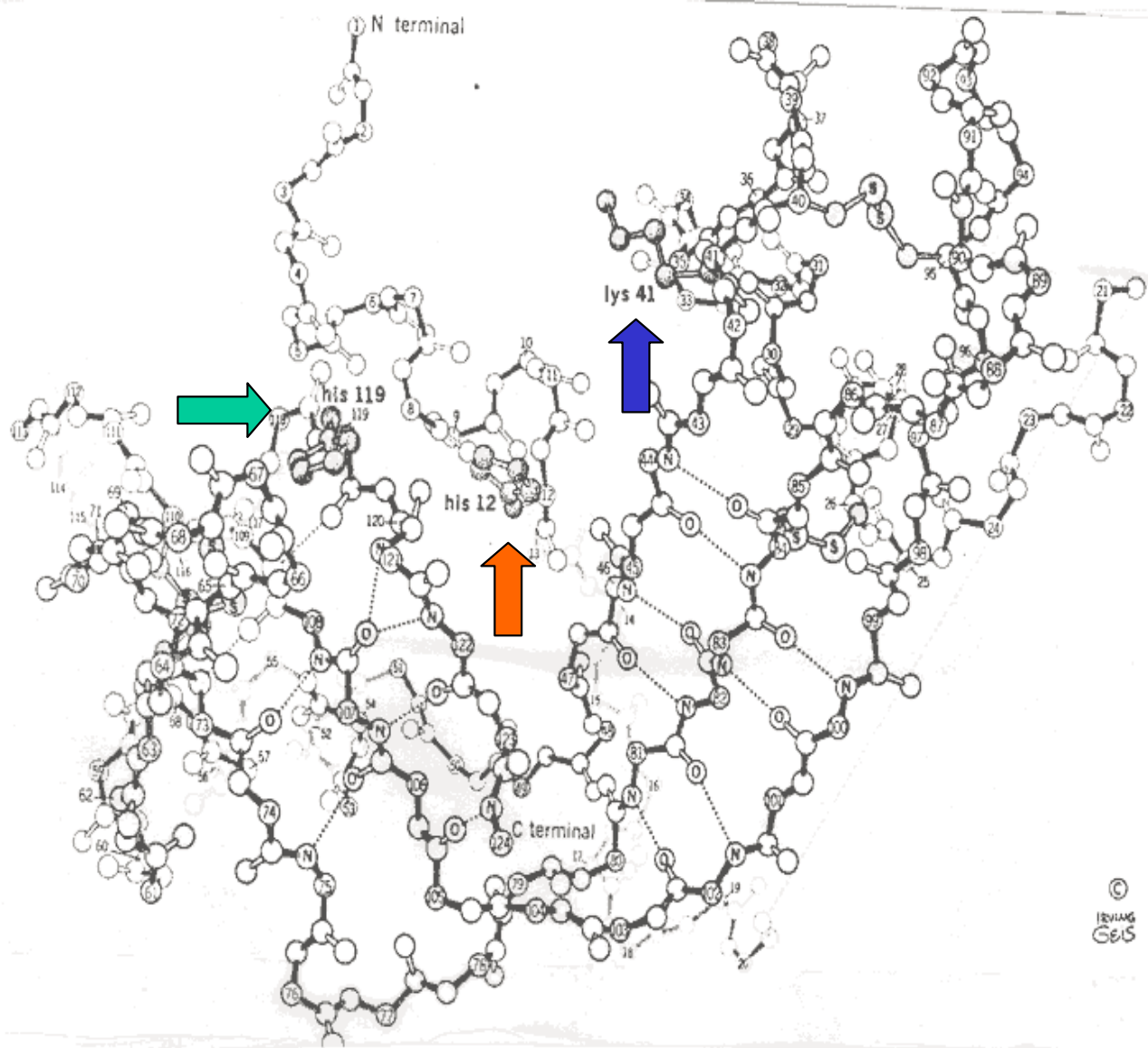
Isolation from bovine pancreas

Many crystal structures (124 entries, e.g. pdb-codes: 1AFK, 1H1H, 1QHC, 1RND)

First „NMR-structure“, 4 Nobel prizes

M: 13.680, 124 amino acids

Chemical syntheses: 1969 Merrifield (0.4 mg), 1979 Yajima (3 mg)



RNase A

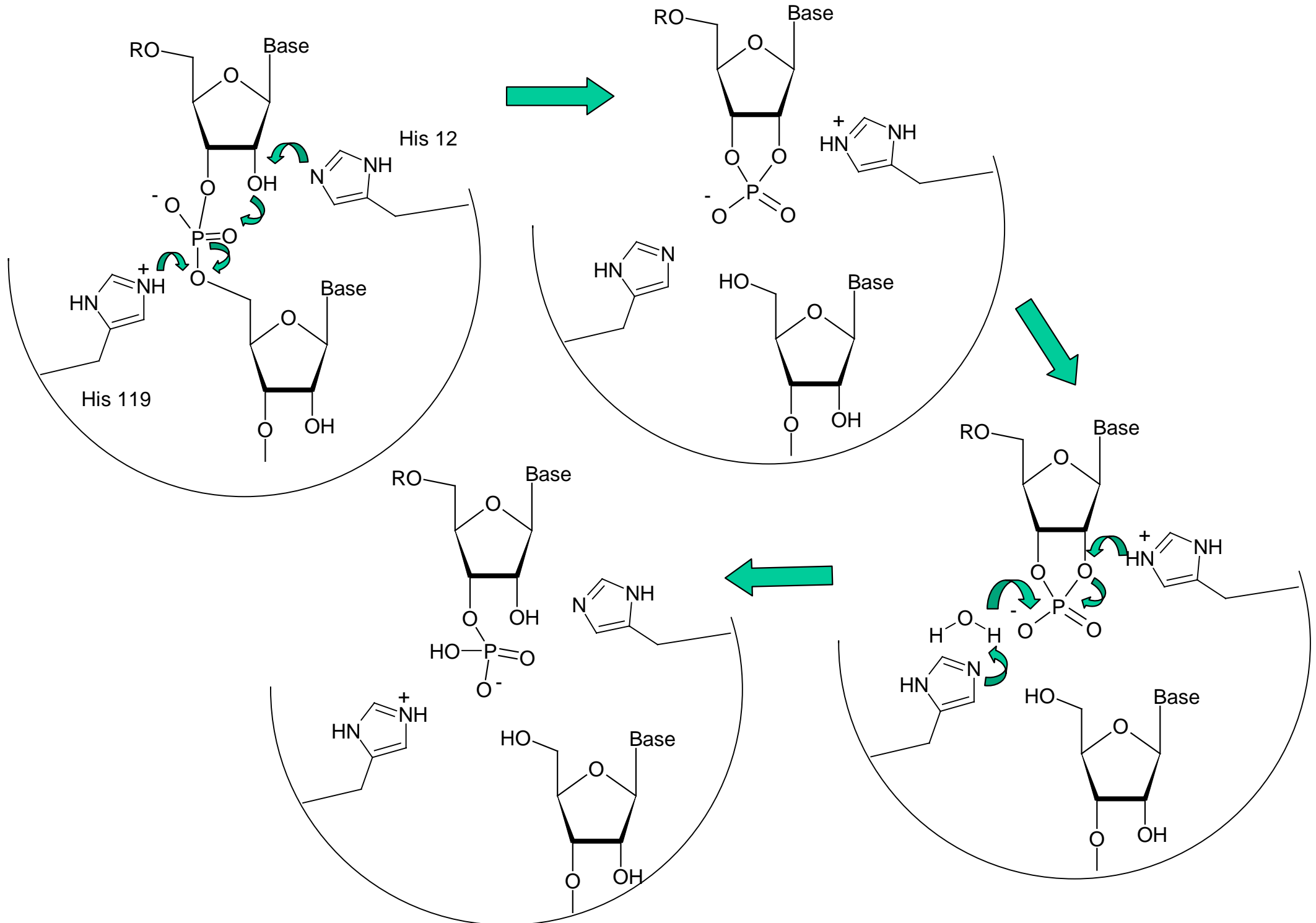
Catalytic centre:

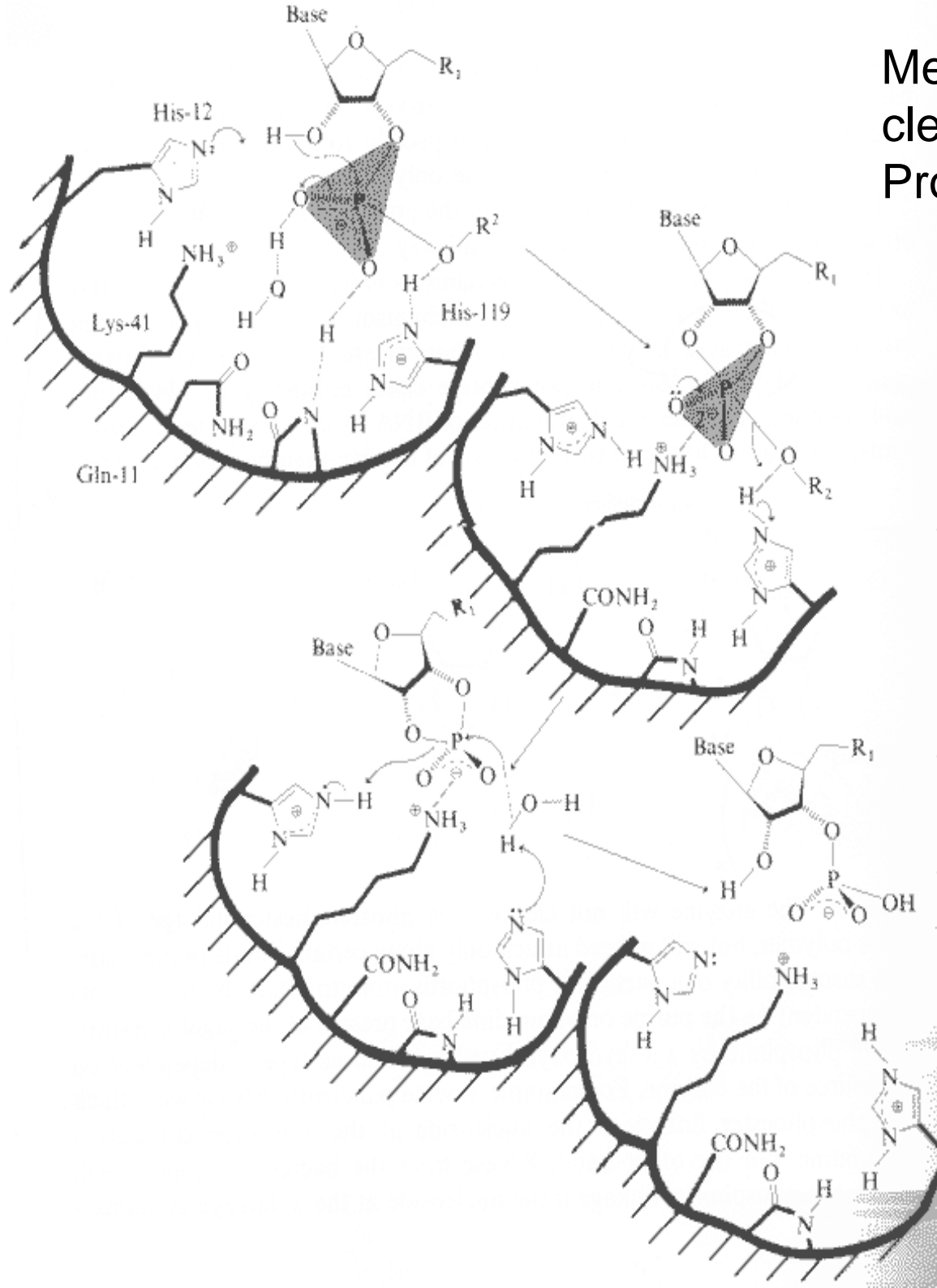
His 12

His 119

Lys 41

Mechanism of ribonuclease A reaction





Mechanism of phosphodiester-cleavage
 Product: 3`-Phosphate

Major steps:
 Transesterification - hydrolysis

General acid-base catalysis (histidine)

Ribozymes – catalytically active ribonucleic acids - metalloenzyme

Discovered in 1982

2 Groups

35 – 155 Nucleotides: Hammerhead, Hairpin, Hepatitis Delta
Formation of 2',3'-cyclo-phosphates and 5'-OH (analogous to RNase A)

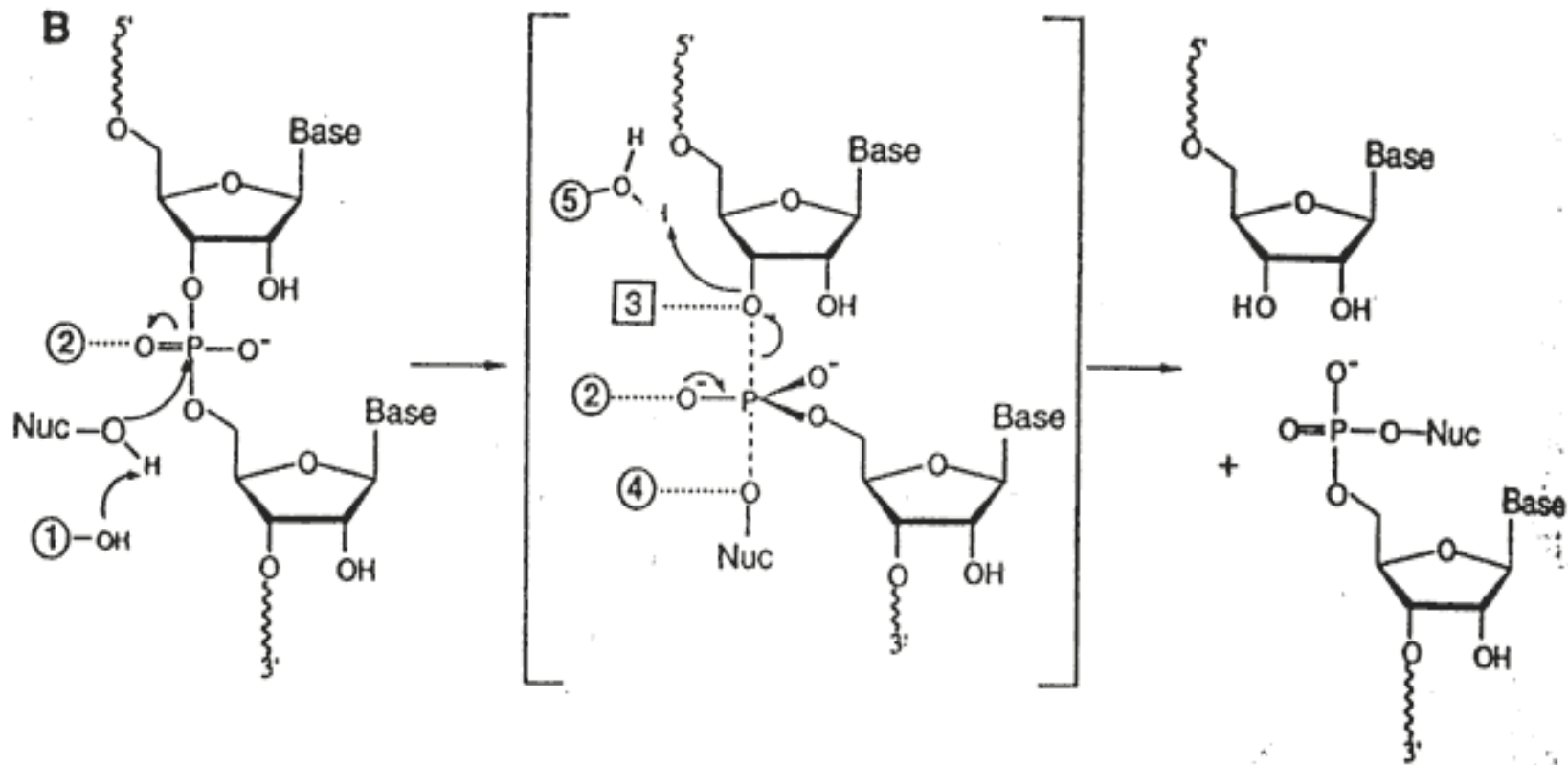
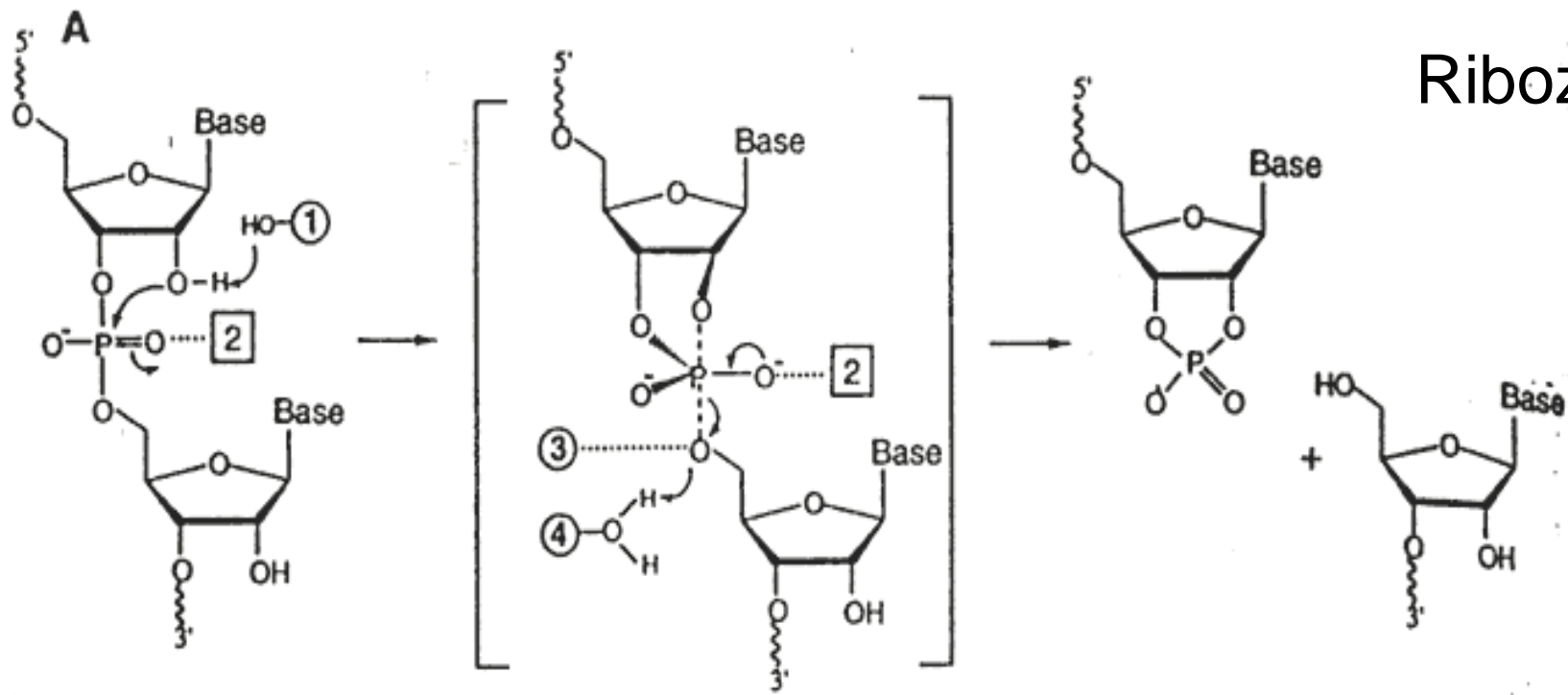
100- 3000 Nucleotides: RNase P, Group I and II Introns
Formation of 5'-phosphate (and 3'-OH)

Self modification, except RNase P (processes t-RNA Precursor)

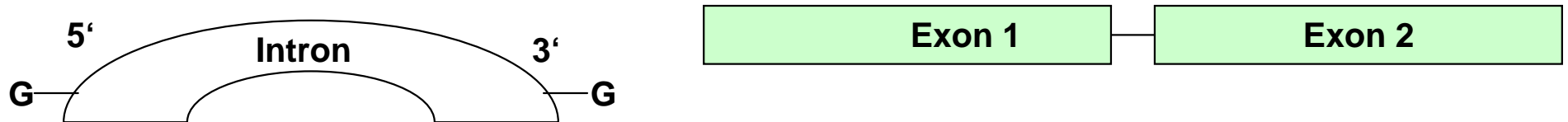
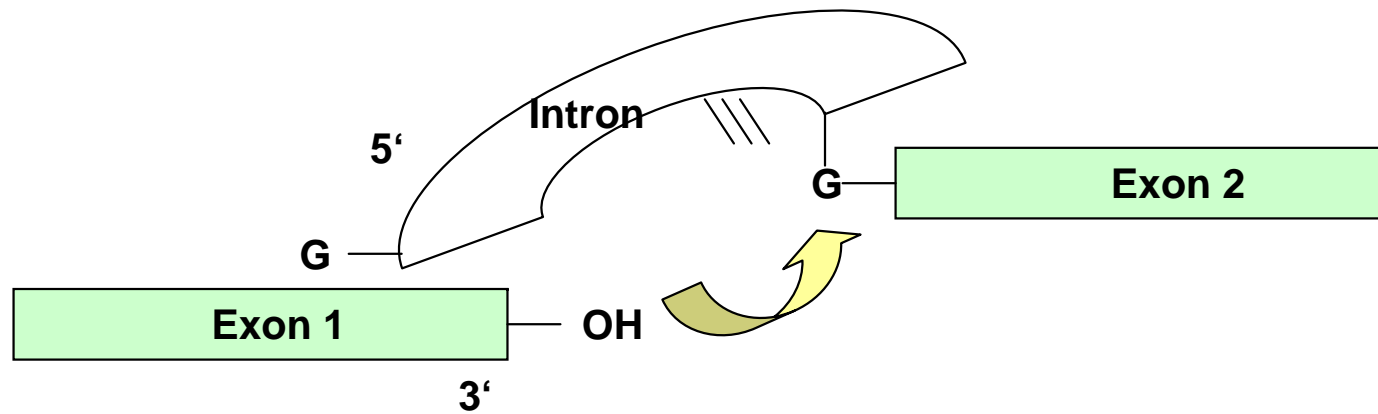
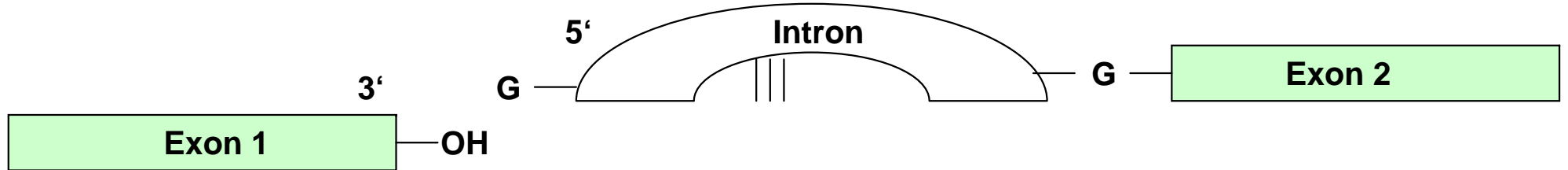
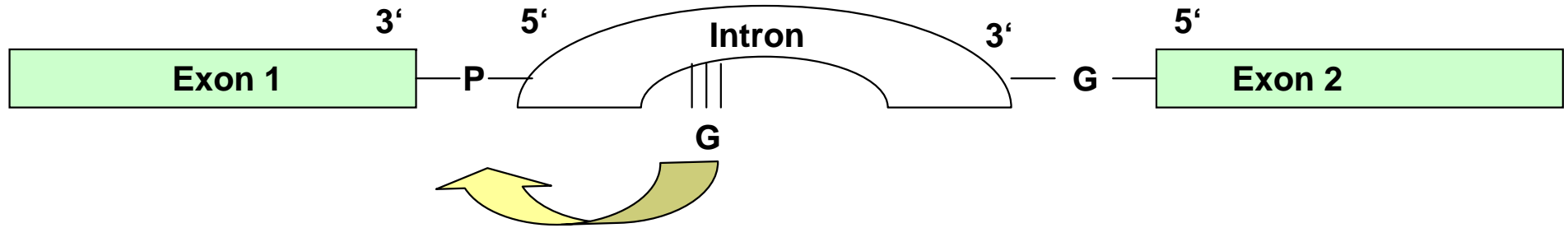
Reaction rate enhancement: $\sim 10^{11}$

Essential: Mg^{2+}

Ribozyme



Group I Intron-Splicing



Magnesium:

Coordination of O only (Mn²⁺ also with N)

6 Ligands (Ca²⁺ > 6)

pKa of water lowered to 11.4

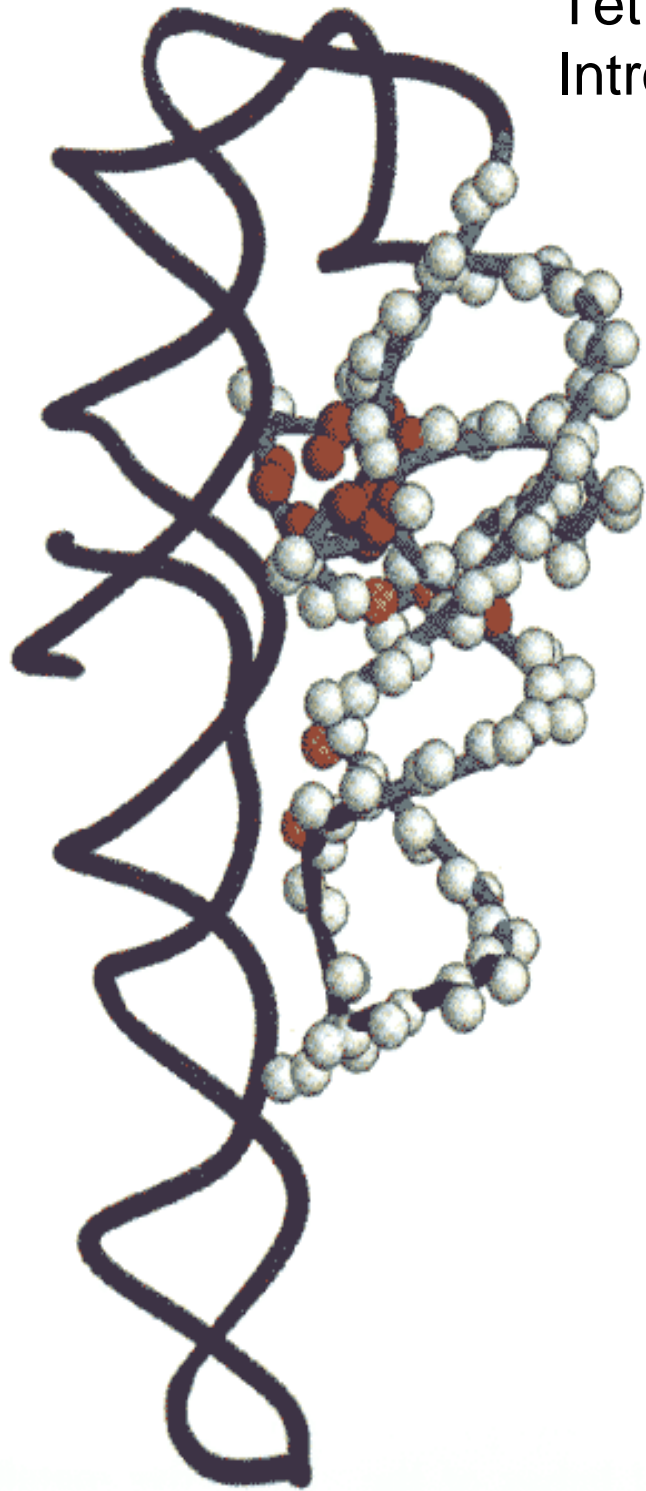
Smallest catalytically active RNA:

UUU

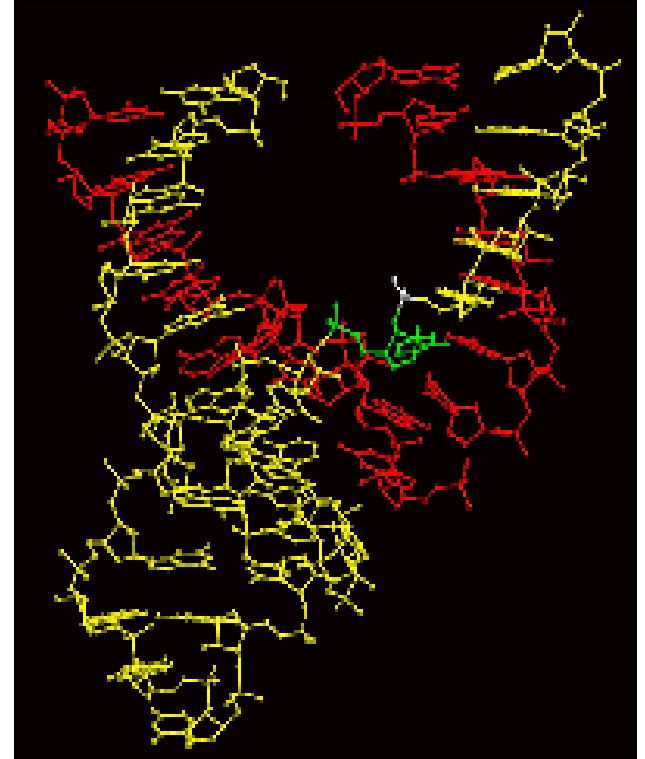
Hydrolysis of G[↑]AAA in the presence of Mn²⁺

Weakest codon-anticodon binding interaction

Tetrahymena Group I Intron



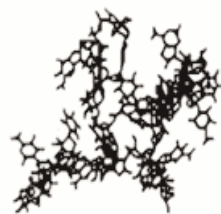
Hammerhead-RNA



Simulated Annealing

nOe Constraints On

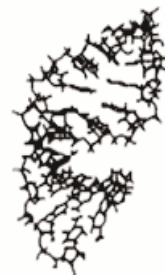
Electrostatic Off
VdW Off



Random starting structure

Refinement

Dihedral constraints On



Global fold

Minimization

(Electrostatic On)
VdW On



Refined structure



Final structure