

# Nucleic Acids

*Nano 501*

*April 18, 2007*

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Pharmacology*

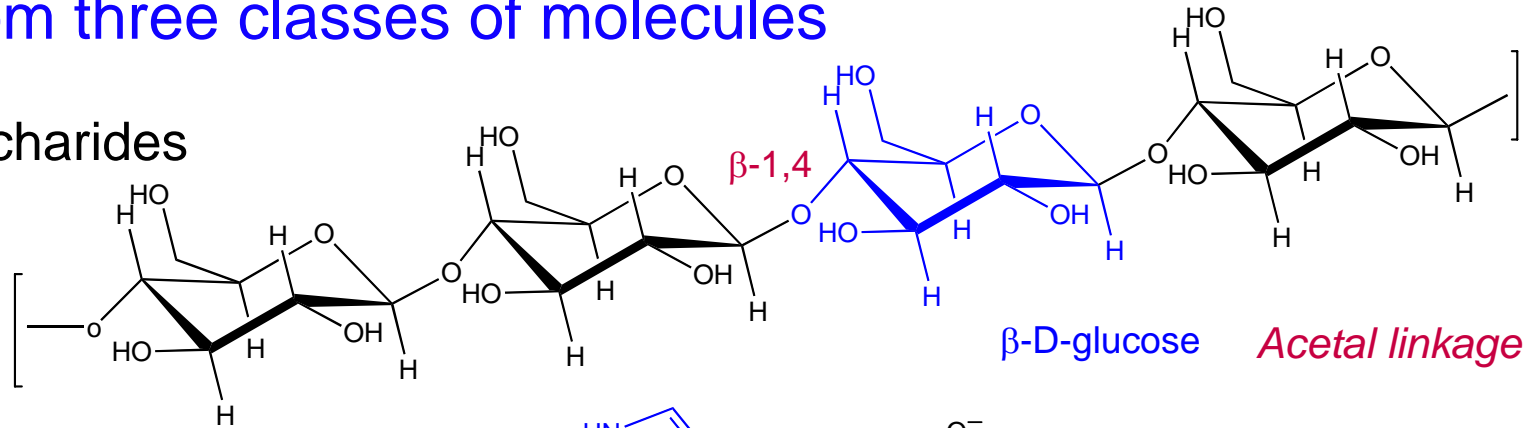
*Birck Nanotechnology Center*

# Nucleic Acids

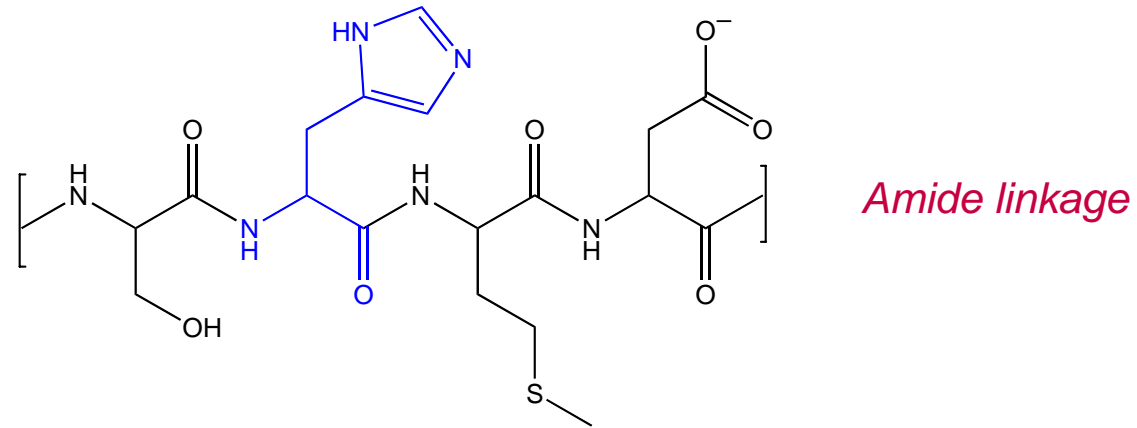
- I. Structure
- II. Biological Function
- III. Modification of RNA
- IV. Genesis of Adenine
- V. Material Science

# Nature builds nanoscale materials largely from three classes of molecules

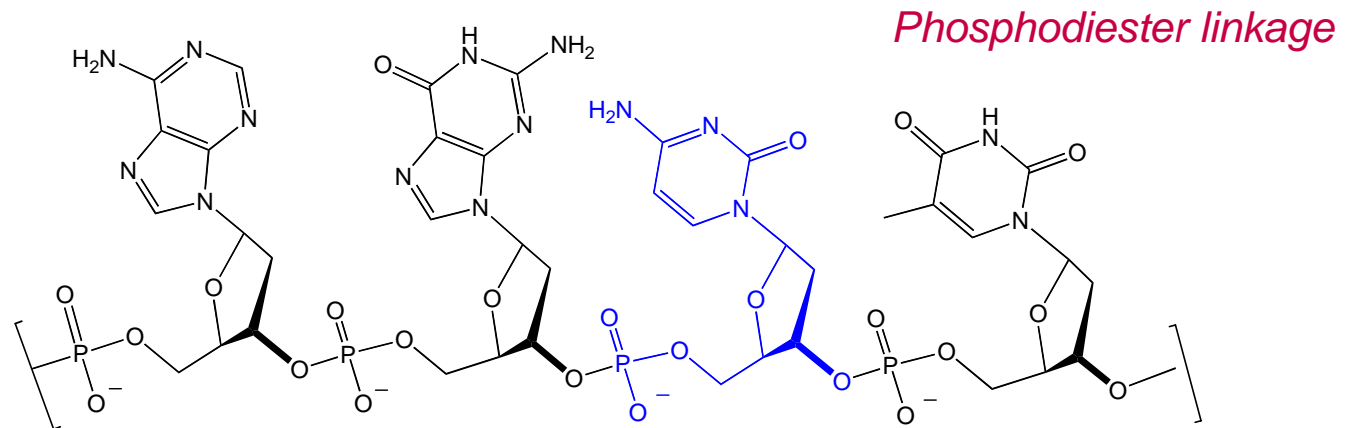
## Polysaccharides



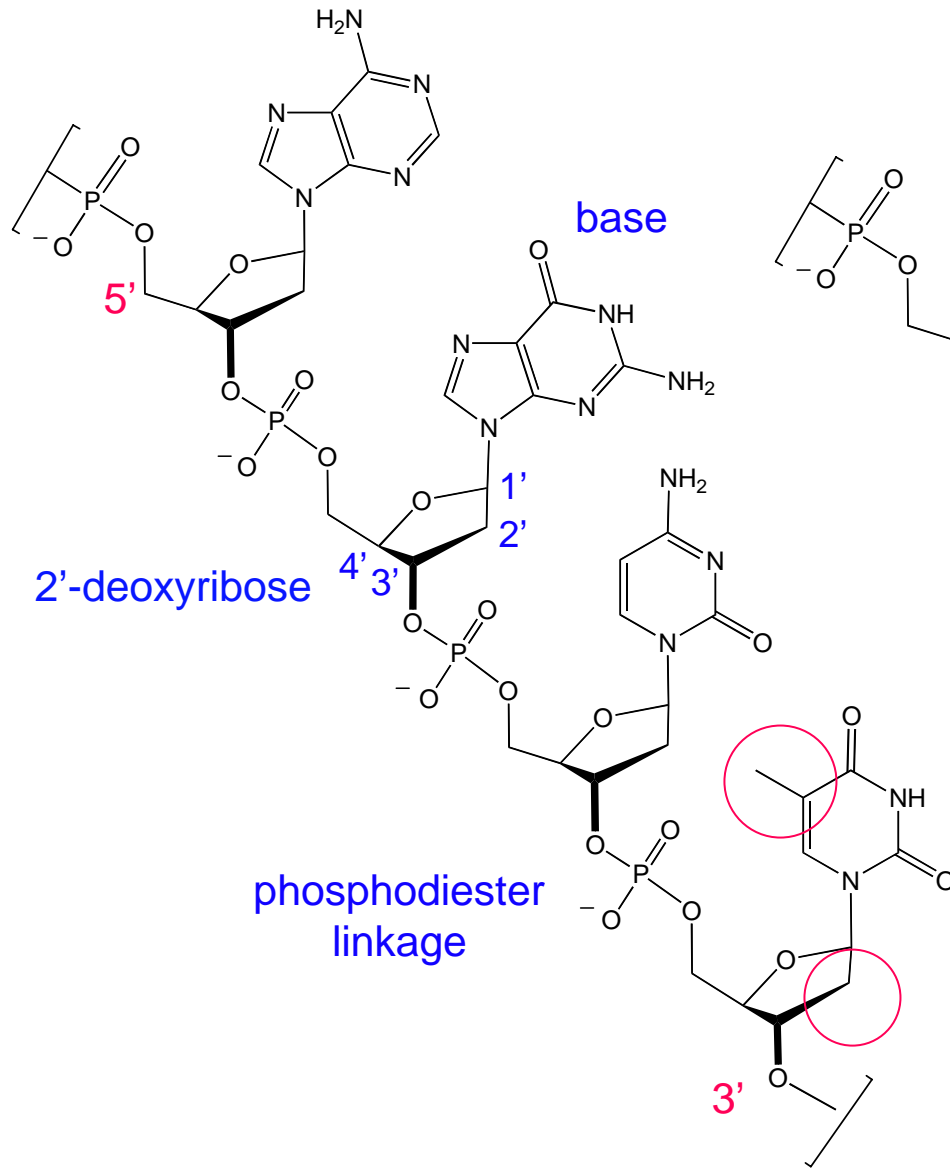
## Polypeptides



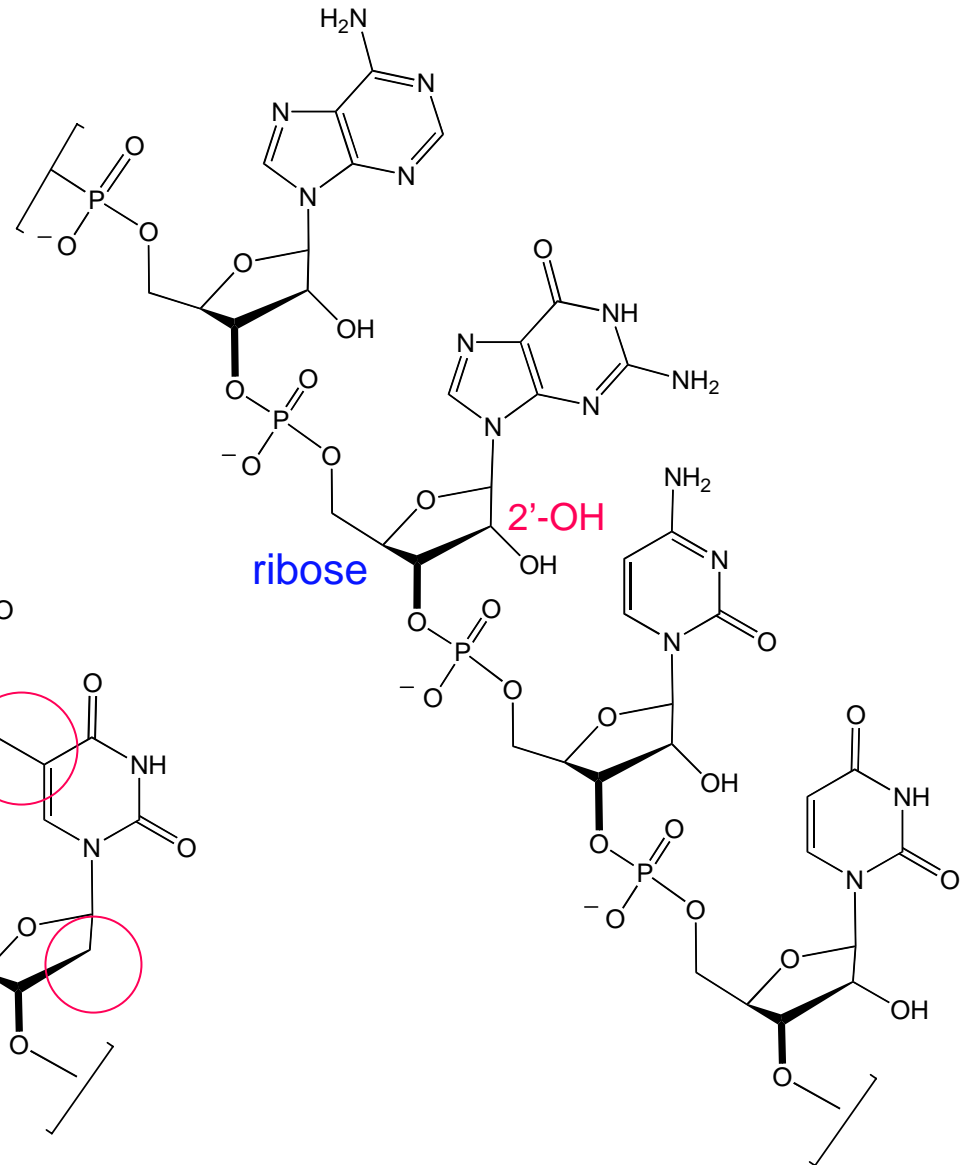
## Polynucleotides



## Deoxyribonucleic acid (DNA)

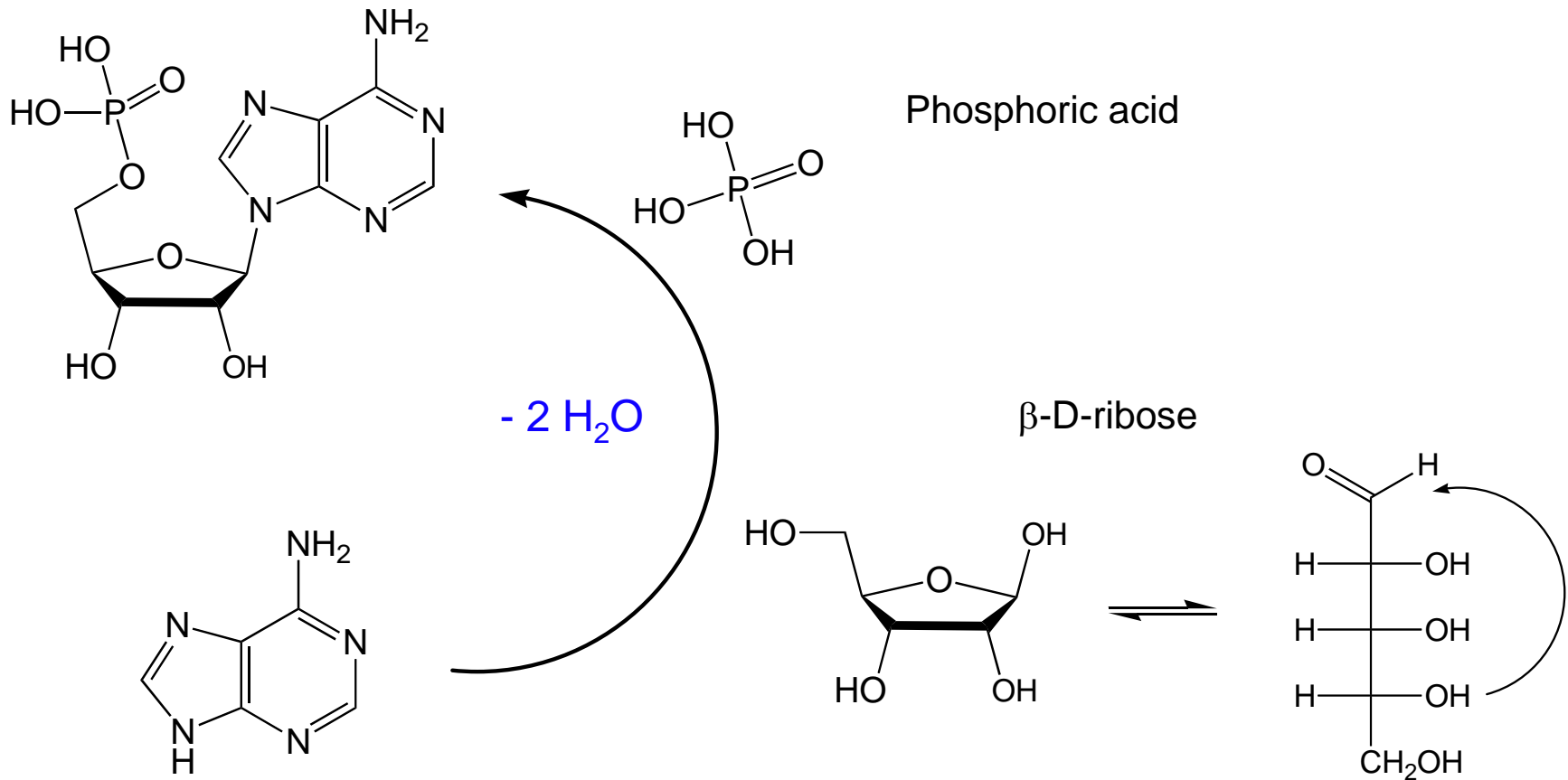


## Ribonucleic acid (RNA)



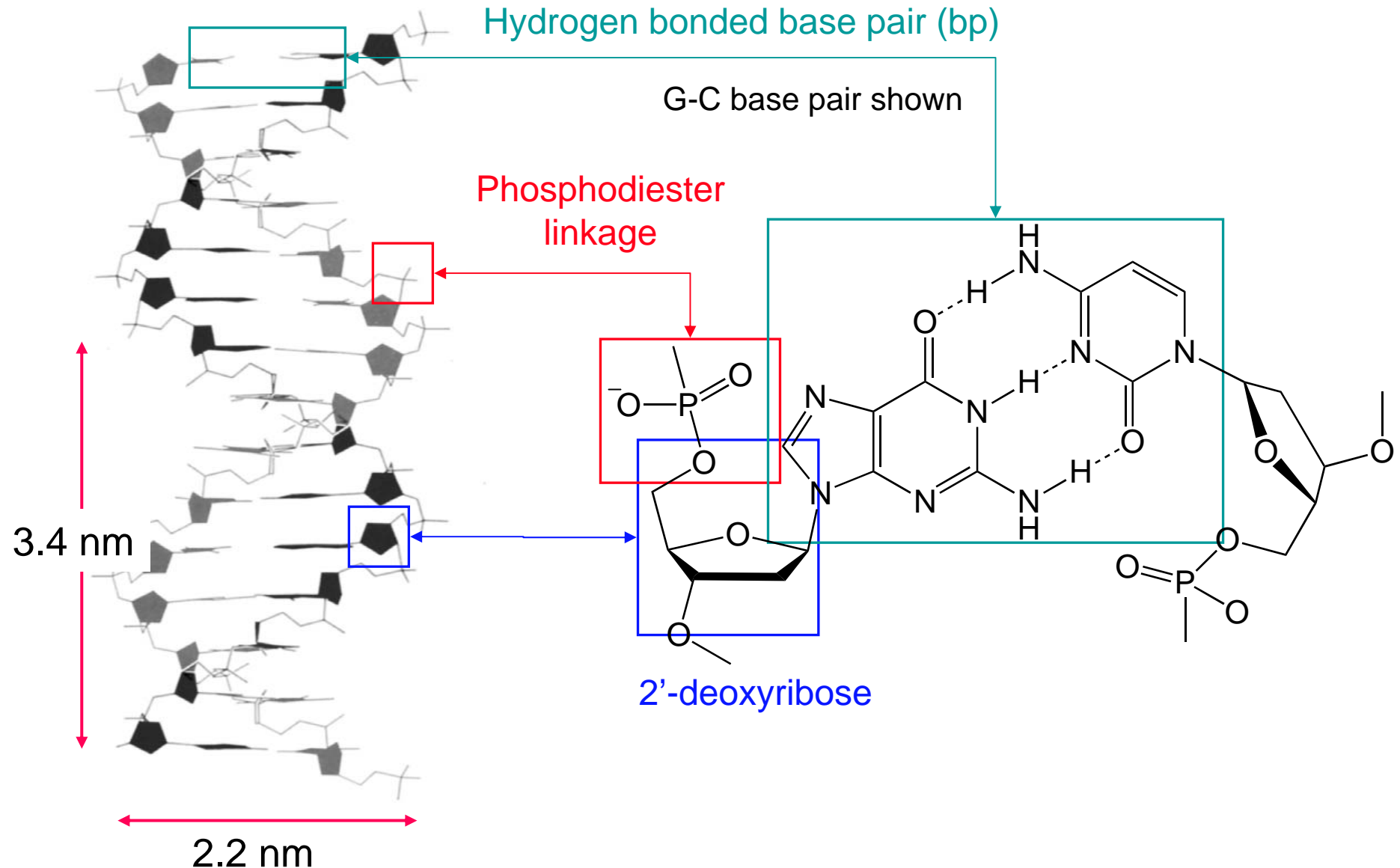
# DNA monomeric unit

Adenosine monophosphate (AMP)

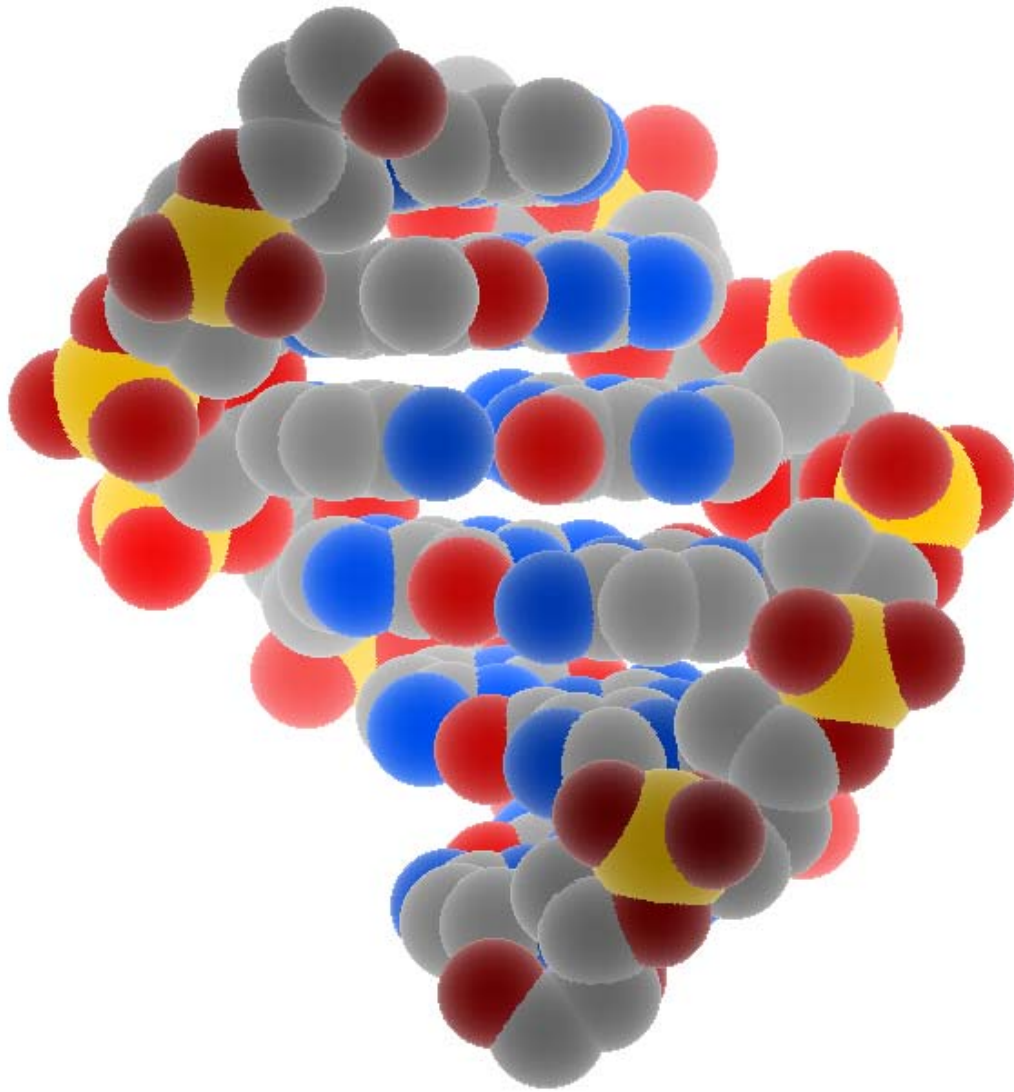


# B-DNA Structure

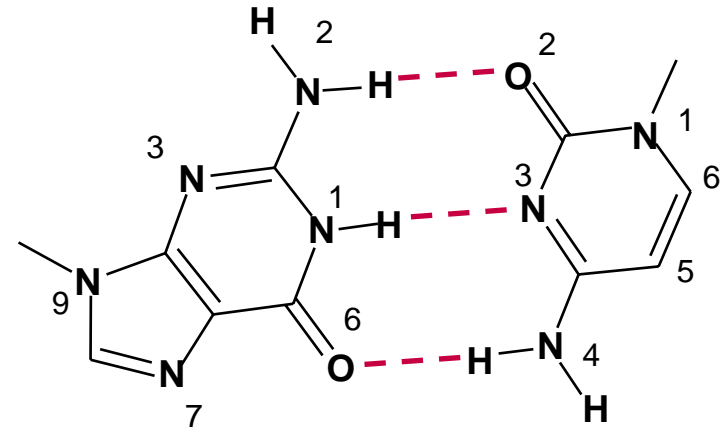
10 bp / helix turn



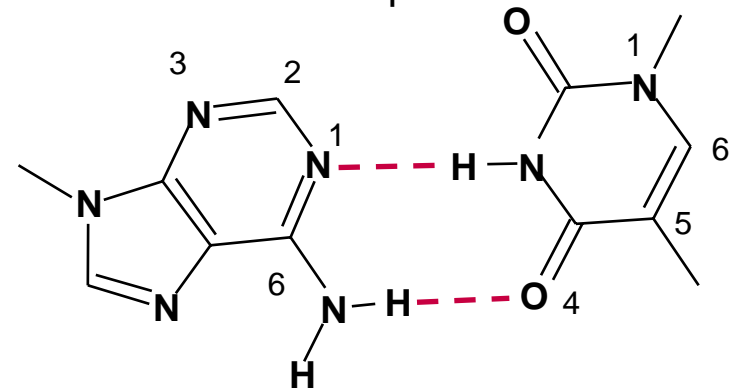
# Space filling model of B-DNA structure



G-C base pair



A-T base pair



# Central Dogma of Molecular Biology

DNA



Transcription

RNA



Translation

Proteins

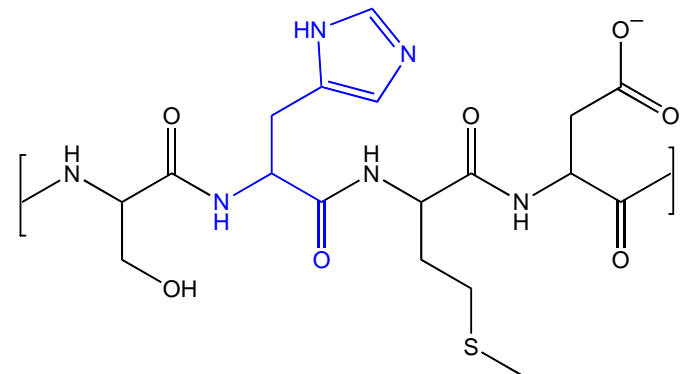
5'....TCA CAT ATG GAT....3'  
: : : : : : : :  
Template strand 3'....AGT GTA TAC CTA....5'



mRNA 5'....UCA-CAU-AUG-GAU....3'



...Ser-His-Met-Asp...





# The Human Genome

## Human nuclear DNA

3.1 billion bp

~ 24,000 genes (3% of genome)

## Genes in the human code for the construction of:

Proteins ~ 23,000

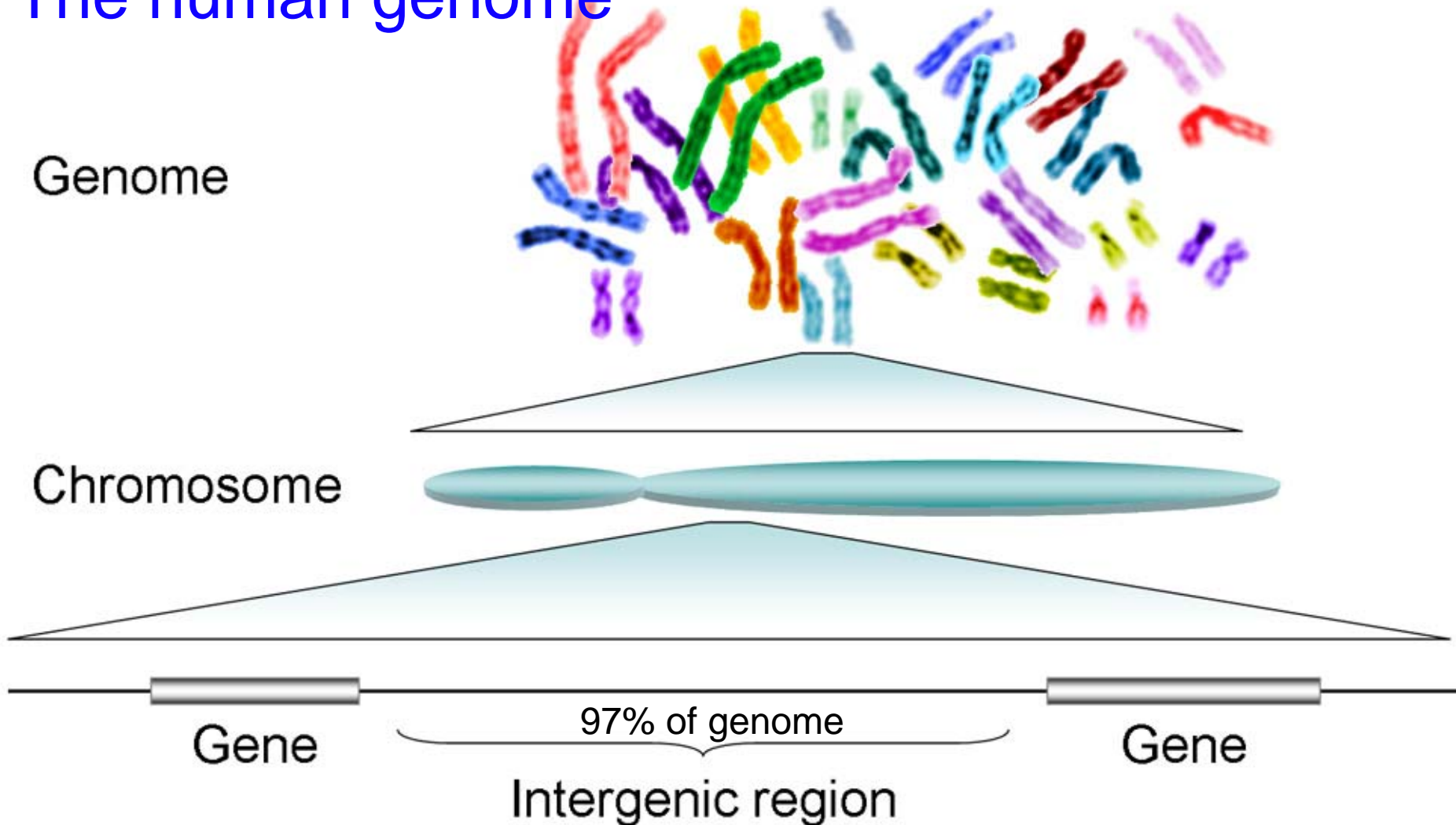
miRNA ~ 400

rRNA ~ 7

snRNA ~ 170

tRNA ~ 60

# The human genome



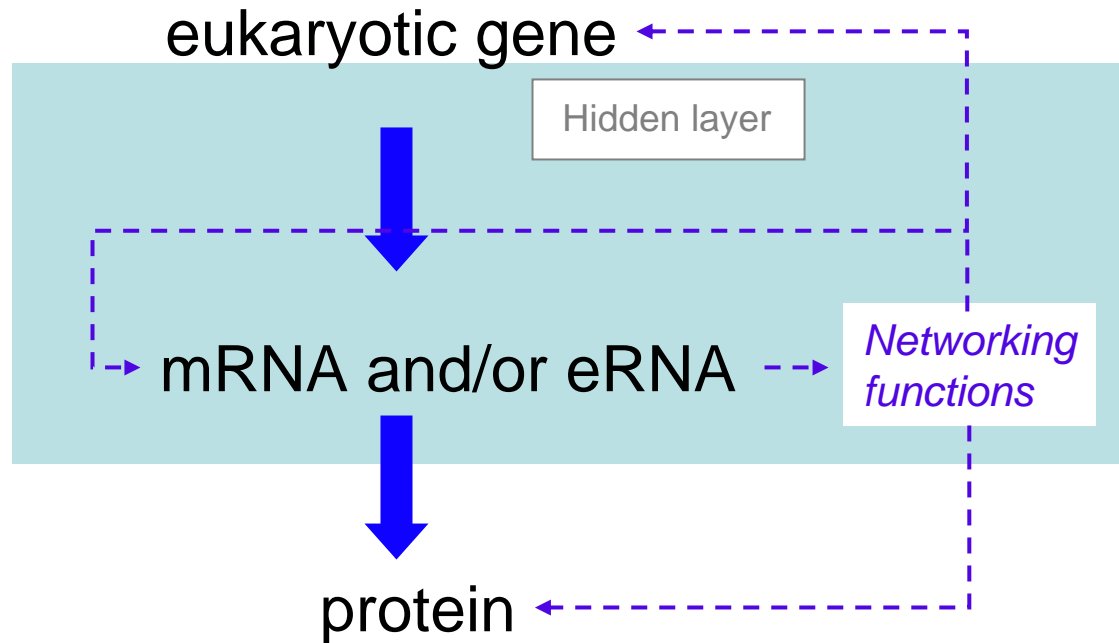
The human genome is composed of 23 pairs of **chromosomes** (46 *in total*), each of which contain hundreds of **genes** separated by *intergenic regions*. Intergenic regions may contain **regulatory sequences** and non-coding DNA. [http://en.wikipedia.org/wiki/Human\\_genome](http://en.wikipedia.org/wiki/Human_genome)

# RNA

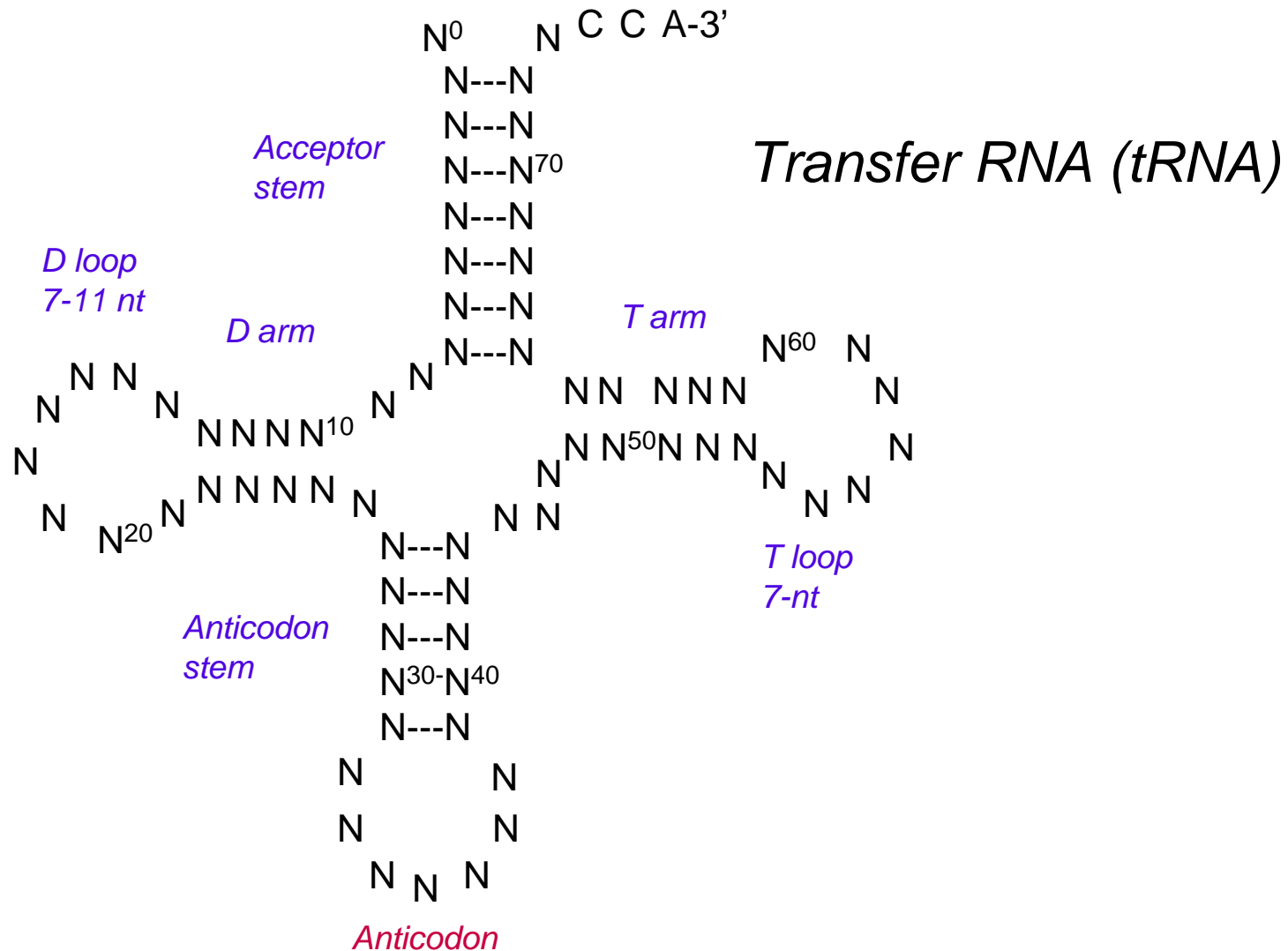
	Name	Size	Function
miRNA	microRNA	21-23 nt	regulation of gene expression
mRNA	messengerRNA	1400 nt (average)	code for protein synthesis
snRNA	small nuclear RNA	40-50 nt	pre-mRNA splicing
tRNA	transfer RNA	73-93 nt	protein synthesis
ncRNA	non-coding RNA	small	Unknown function
rRNA	ribosomal RNA	5000 nt (varies)	organization of protein synthesis machinery

# RNA Regulatory Networks

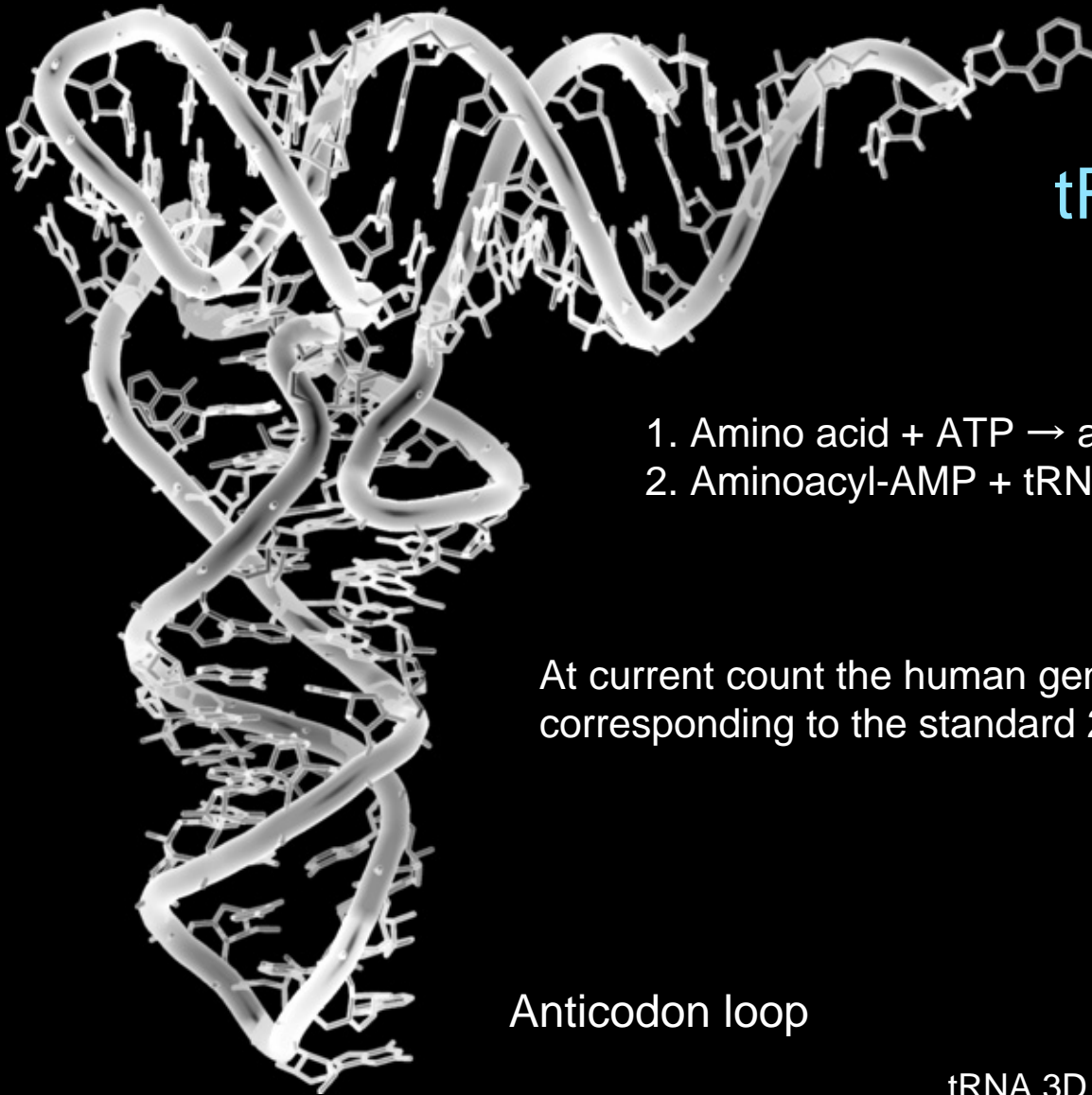
Undiscovered dark matter of the eukaryotic cell



# RNAs typically have extensive secondary structure



# RNAs are more complex than DNA



tRNA 3D structure

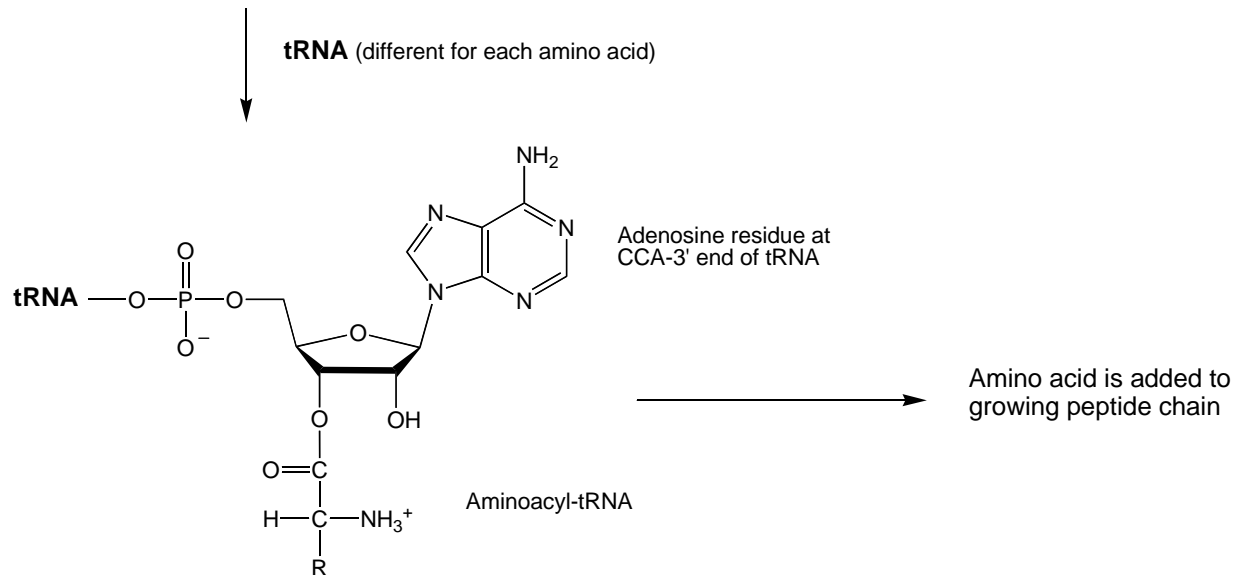
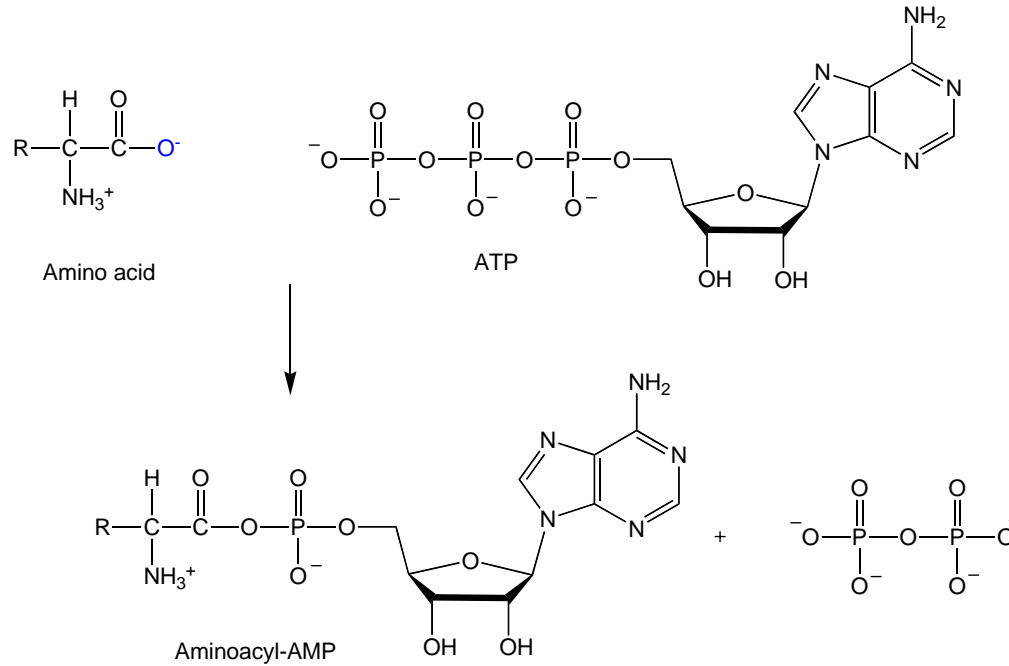
1. Amino acid + ATP  $\rightarrow$  aminoacyl-AMP + P<sub>pi</sub>
2. Aminoacyl-AMP + tRNA  $\rightarrow$  aminoacyl-tRNA + AMP

At current count the human genome contains 448 genes corresponding to the standard 20-amino acid decoding tRNAs

Anticodon loop

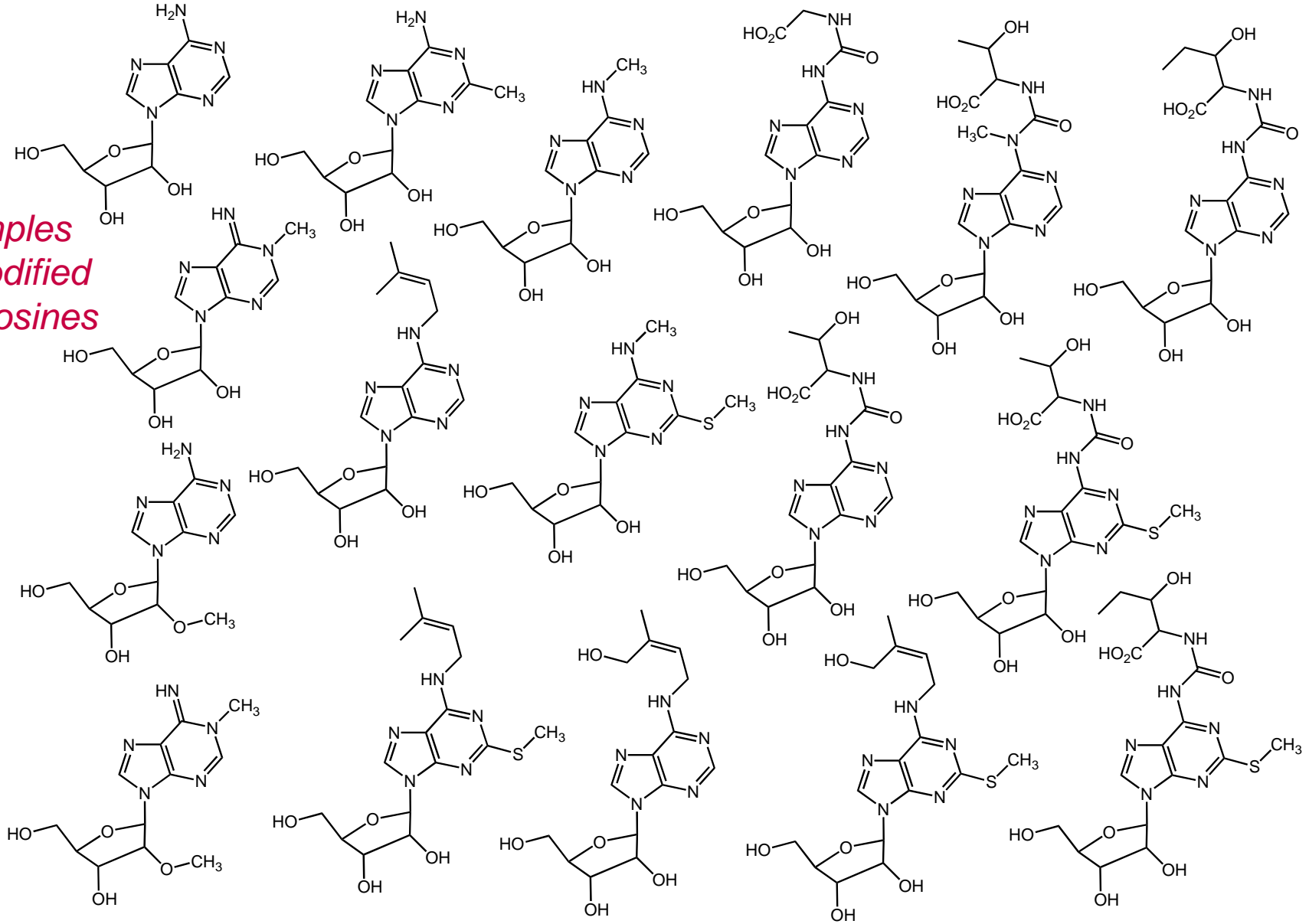
tRNA 3D structure -1974 Alexander Rich,  
Aaron Klug & coworkers- *Wikipedia*

# tRNAs are critical intermediates in protein synthesis



## Nucleotides in tRNA are often modified

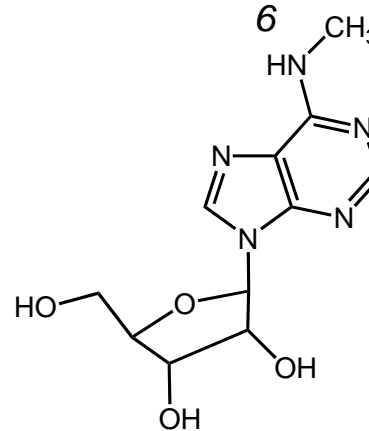
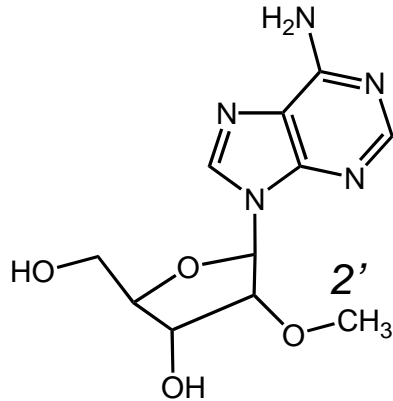
## Examples of modified adenosines





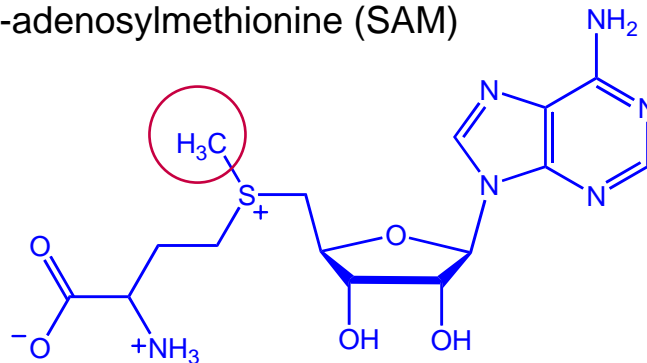
# Nucleotide Methylation

The most common modification in tRNA, mRNA, and rRNA



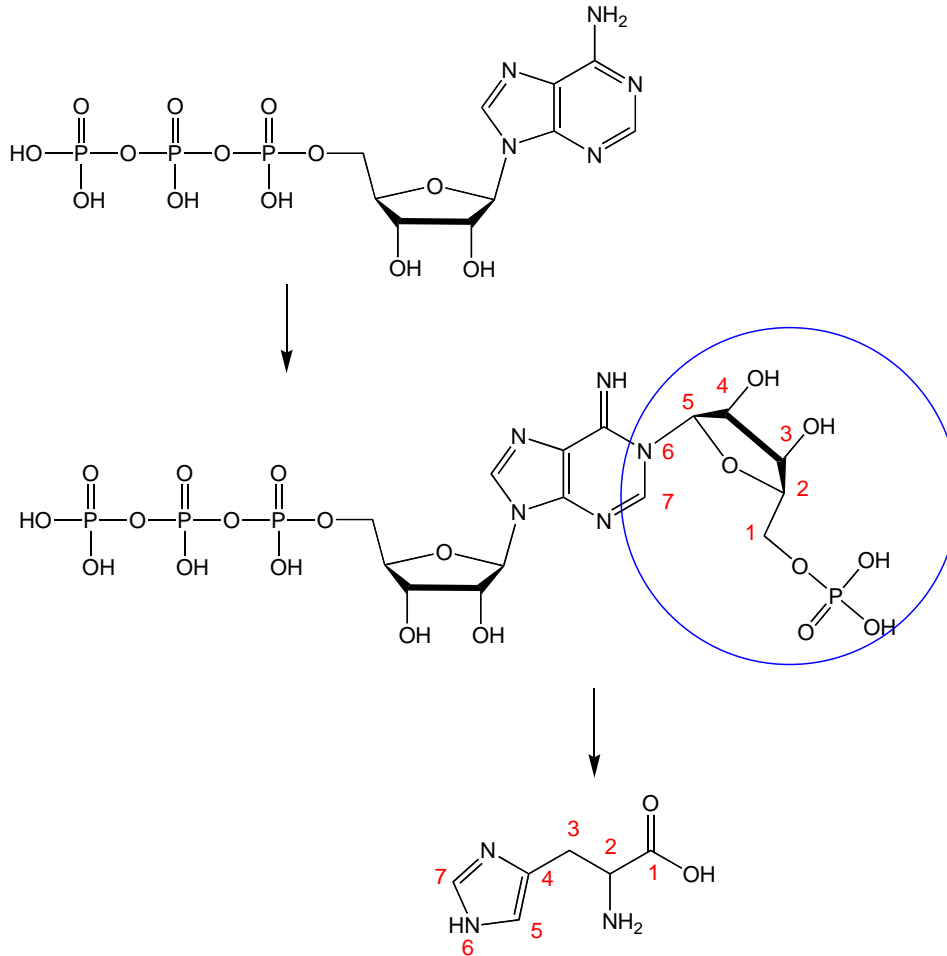
*N-6 is sometimes also modified in DNA*

The co-enzyme that facilitates adenosine methylation is S-adenosylmethionine (SAM)

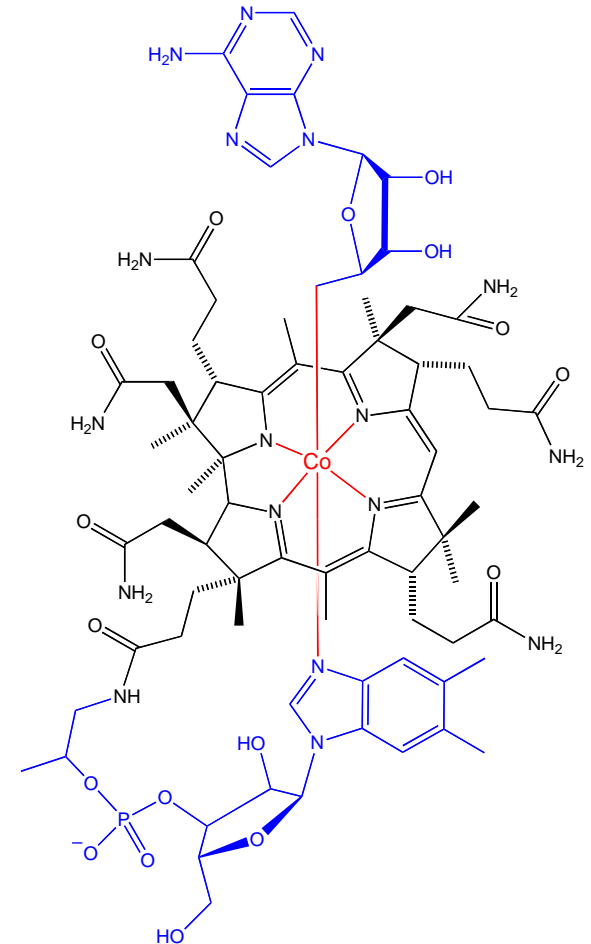


# Adenosine is responsible for the biosynthesis of DNA from RNA

1) ATP is the biosynthetic precursor of histidine. C7 of histidine becomes the C5 methyl group of thymine

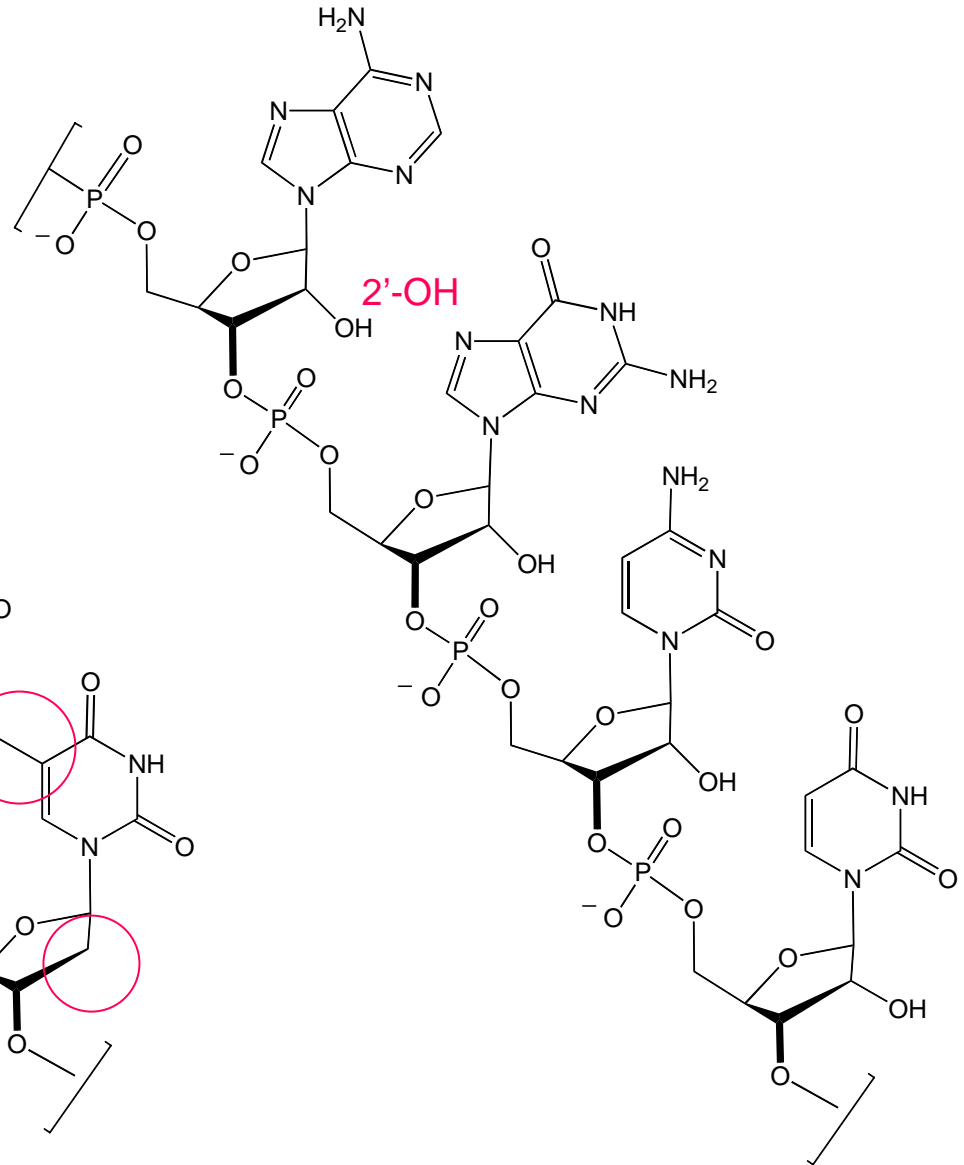
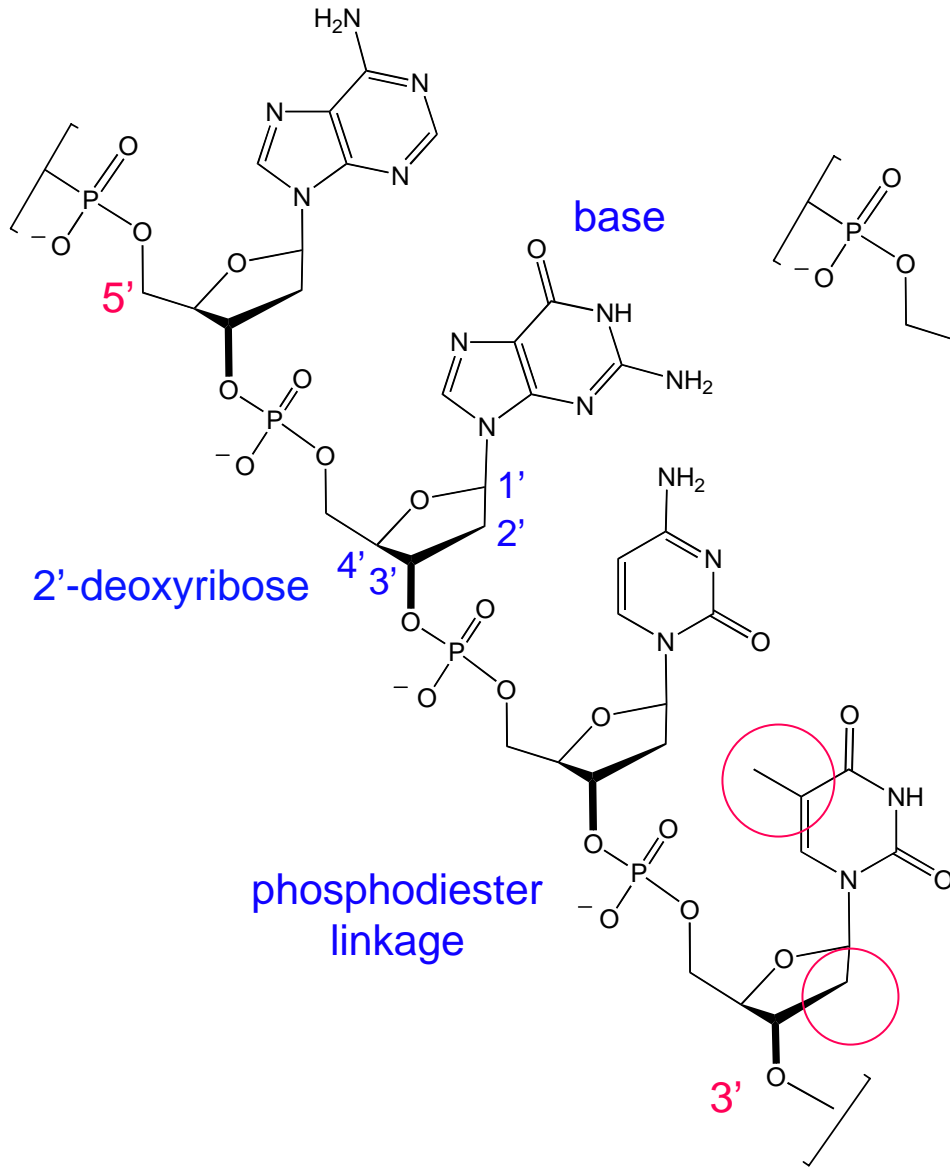


2) The coenzyme adenosylcobalamin participates in the reduction of the 2'-OH to 2'-H



## Deoxyribonucleic acid (DNA)

## Ribonucleic acid (RNA)



## Why did DNA evolve from RNA?

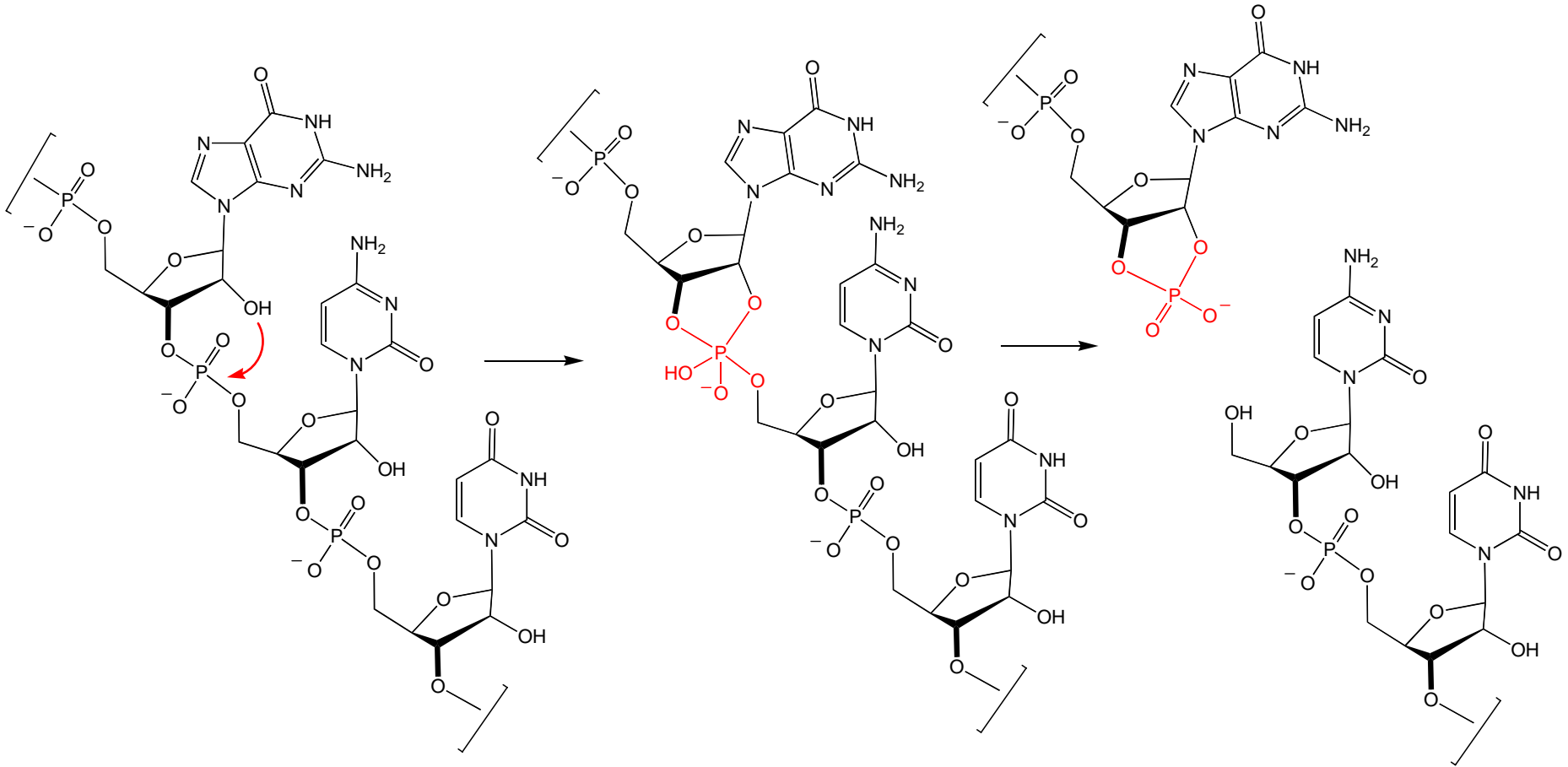
The 2'-OH in RNA participates in a relatively facile degradation pathway to monomers

*Removal of the 2'-OH eliminates this degradation pathway -  
As a result DNA is a more stable polymer.*

Cytosine is prone to spontaneous deamination in water to yield uracil

*Addition of the C5 methyl provides a mechanism for efficient repair of spontaneous cytosine to uracil mutations since uracil can be distinguished from thymine.*

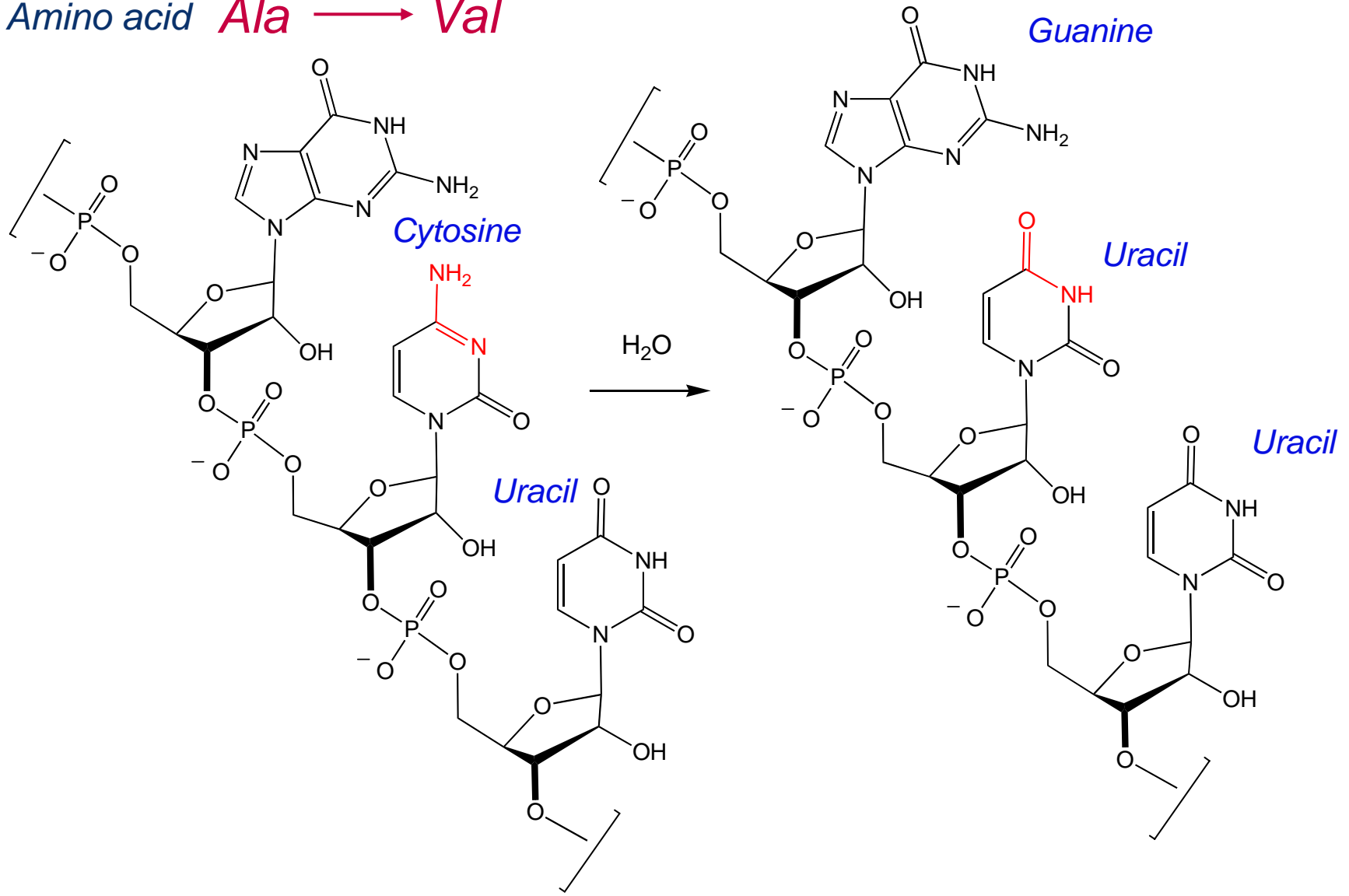
## 2'-OH mediated RNA degradation



# Deamination of cytosine to uracil in RNA is mutagenic

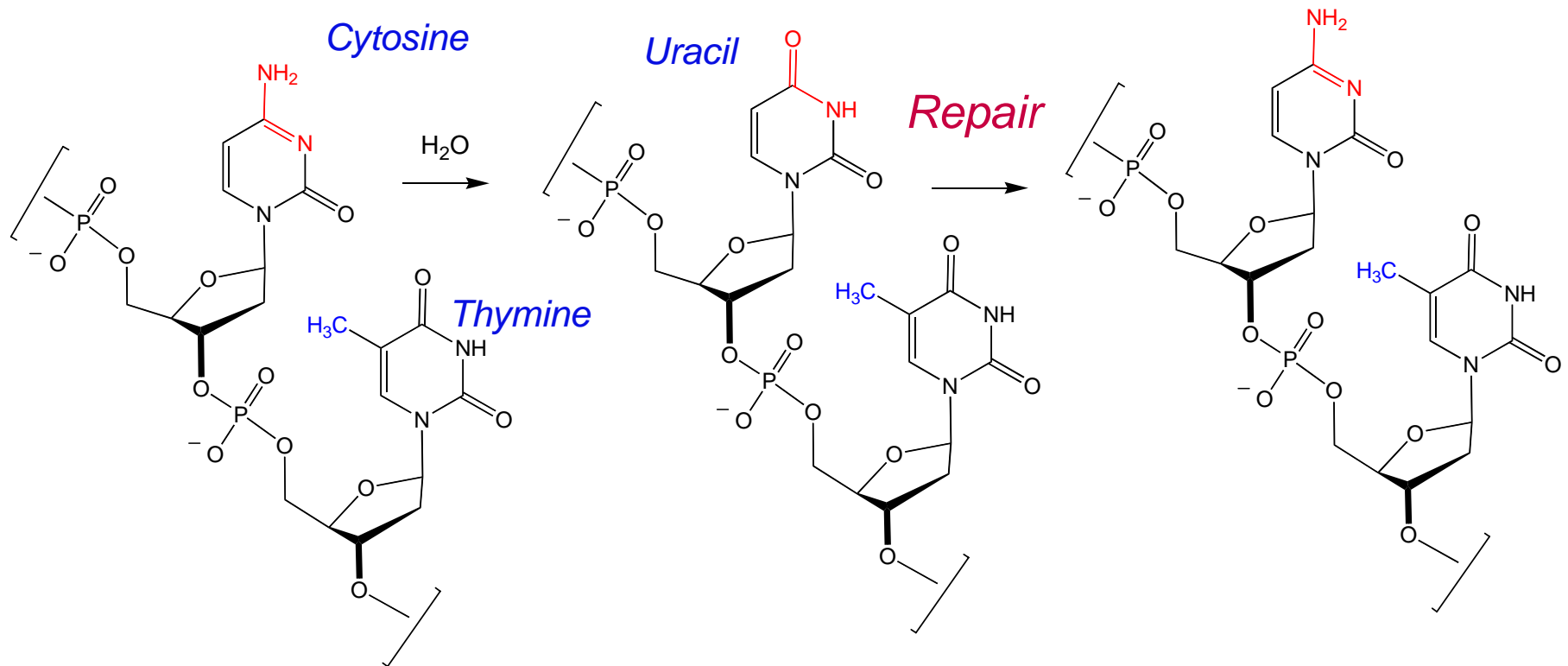
Codon  $GCU \longrightarrow GUU$

Amino acid  $Ala \longrightarrow Val$



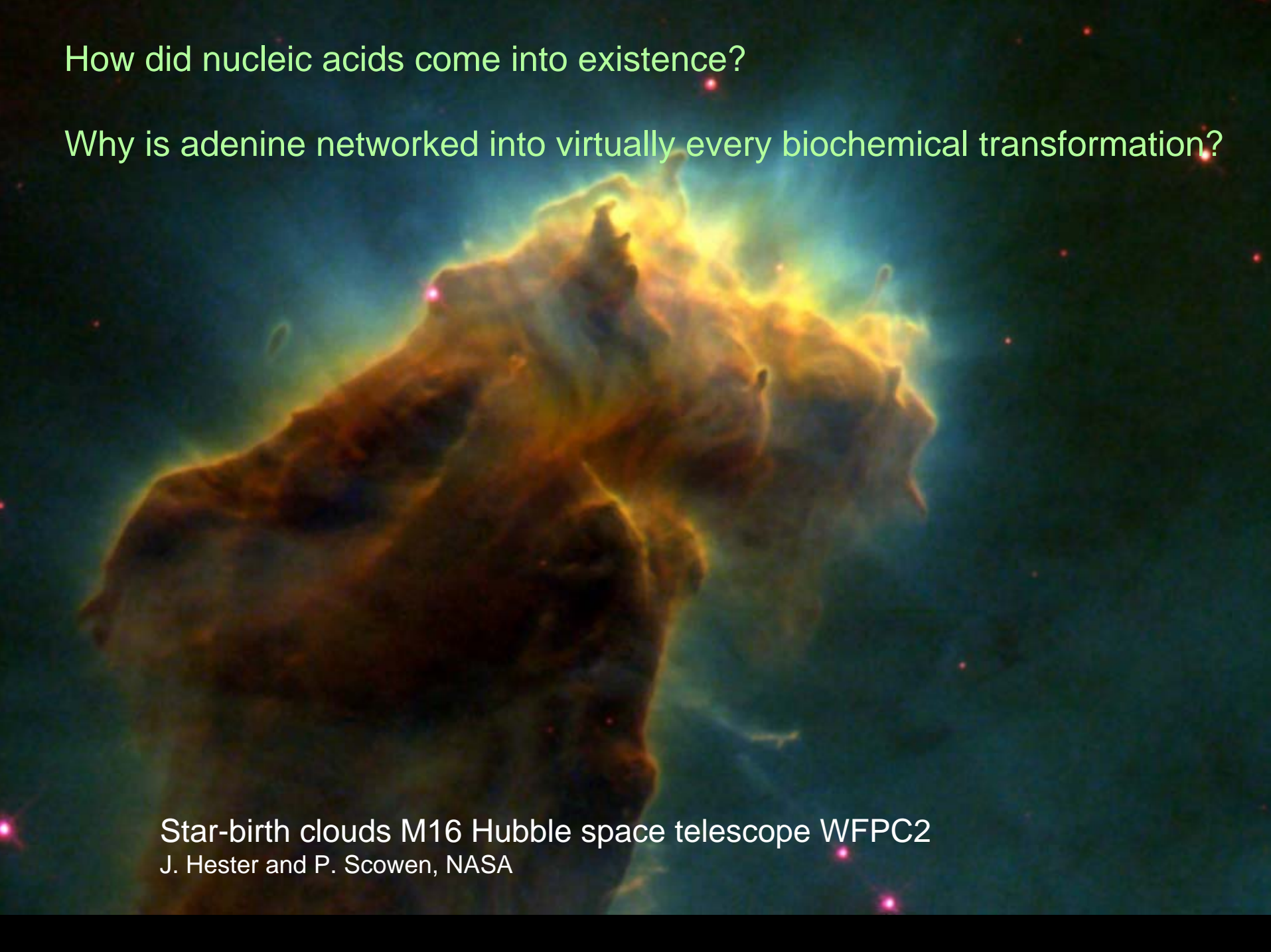
# Deamination of cytosine to uracil in DNA is repaired

*Repair enzymes distinguish uracil from thymine*



How did nucleic acids come into existence?

Why is adenine networked into virtually every biochemical transformation?

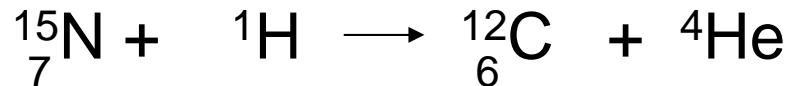
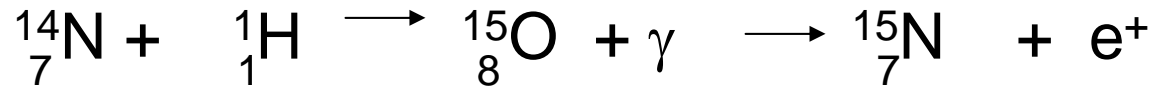
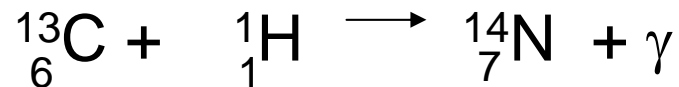
A vibrant image of the M16 star-forming region, also known as the Eagle Nebula. It shows a massive, dark, and turbulent cloud of gas and dust. The cloud is illuminated from within and by nearby stars, creating a complex pattern of bright yellow and orange light against a deep blue background. Several bright, point-like stars are visible, some with prominent diffraction spikes. The overall scene captures the raw energy and chaotic beauty of a stellar nursery.

Star-birth clouds M16 Hubble space telescope WFPC2  
J. Hester and P. Scowen, NASA



# *Stellar generation of the elements*

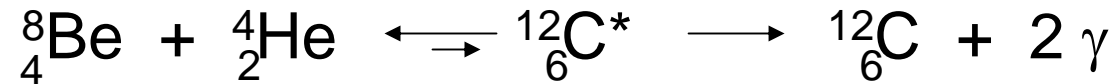
CNO cycle     $T \sim 3 \times 10^7 \text{K}$



The reaction cycle is autocatalytic, which is also a fundamental characteristic of biological processes

## Triple alpha process (Red-Giant Core)

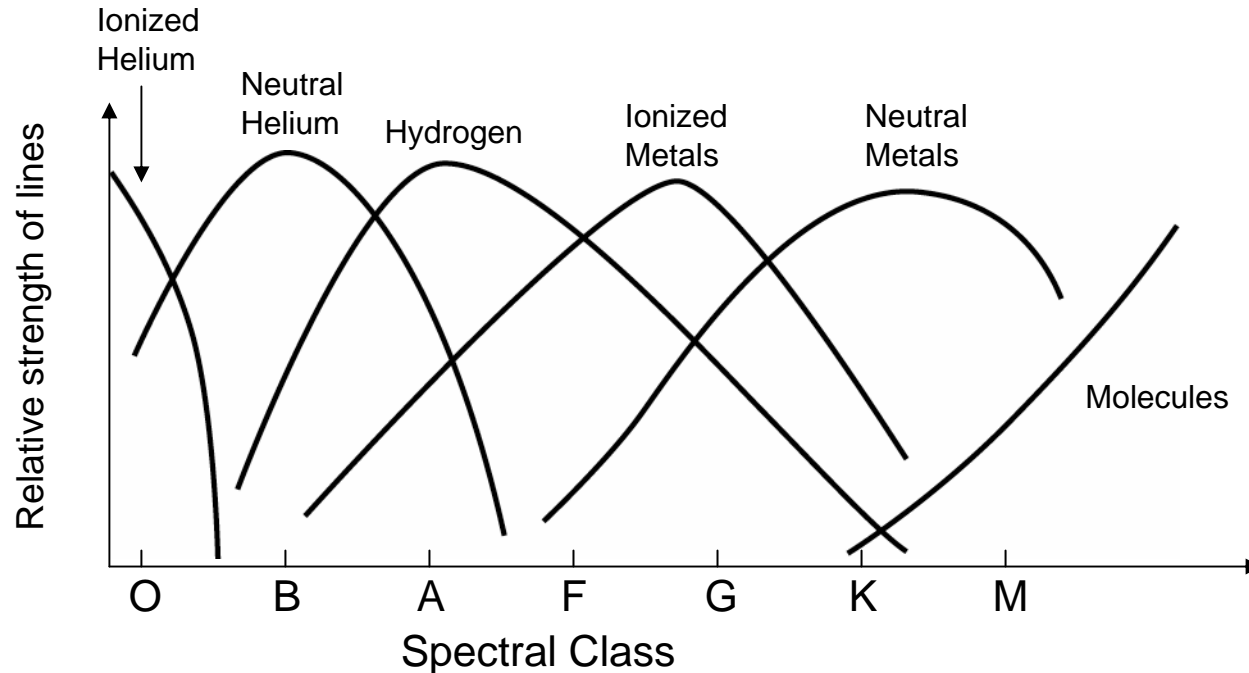
$$T \sim 10^8 \text{K} \quad 2 \text{ } ^4_2\text{He} \rightleftharpoons \text{ } ^8_4\text{Be} \quad T_{1/2} = 7 \times 10^{-17} \text{ s}$$



# Harvard Spectral Classification

Overseas [Bulletin/Broadcast]: A Flash! Godzilla Kills Mothra! (Rodan Named Successor)

<http://www.shef.ac.uk/physics/people/pacrowther/obafgkmrns.html>



Harvard Spectral Classification created by Annie Jump Canon 1918-1924

[www.shef.ac.uk/physics/people/pacrowther/spectral\\_classification.html](http://www.shef.ac.uk/physics/people/pacrowther/spectral_classification.html)

O

Violet

> 28,000K

B

Blue

10,000-28,000K

A

Blue

7,500-10,000K

F

Blue-white

6,000-7,500K

G

Yellow-white

5,000-6,000K

K

Orange-Red

3,500-5,000K

M

R N

Red

< 3,500K

## Low temperature processes

$T \sim 10^4 \text{K}$       Hydrogen exists as  $\text{H}^+$

$T \sim 5 \times 10^3 \text{K}$       Hydrogen exists as  $\text{H}^\bullet$

$T \sim 5000 - 6000 \text{K}$  CH star

G-type giant (G5 to K5) with strong molecular bands for CH

$T < 3500 \text{ K}$  Carbon star (red giant)

Spectra indicate abundance of  $\text{C}_2$  and  $\text{CN}$  as well as other carbon compounds

$T > 1000 \text{ K}$  C star (giant)(R and N in Harvard Classification)

Spectra show  $\text{C}_2$ ,  $\text{CN}$ , and  $\text{CH}$ , but no metallic oxides

$T > 1000 \text{ K}$  CN strong star (giant)

Spectra for  $\text{CN}$  as well as metallic oxides

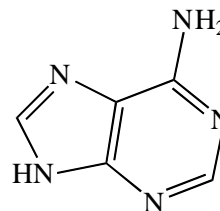
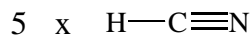


Orion Nebula where new stellar systems are being created; many are surrounded by clouds of HCN or H<sub>2</sub>CO. NASA/JPL-Caltech

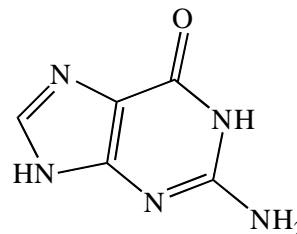
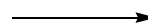
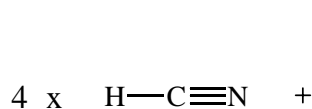
The nucleic acid heterocyclic bases are formally polymers of interstellar CHNO species

## Interstellar CHNO species

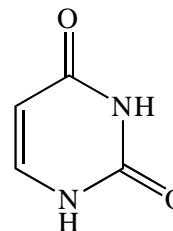
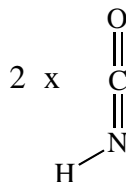
## RNA bases



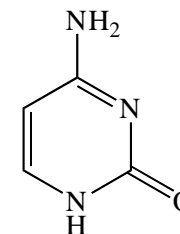
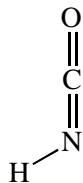
Adenine



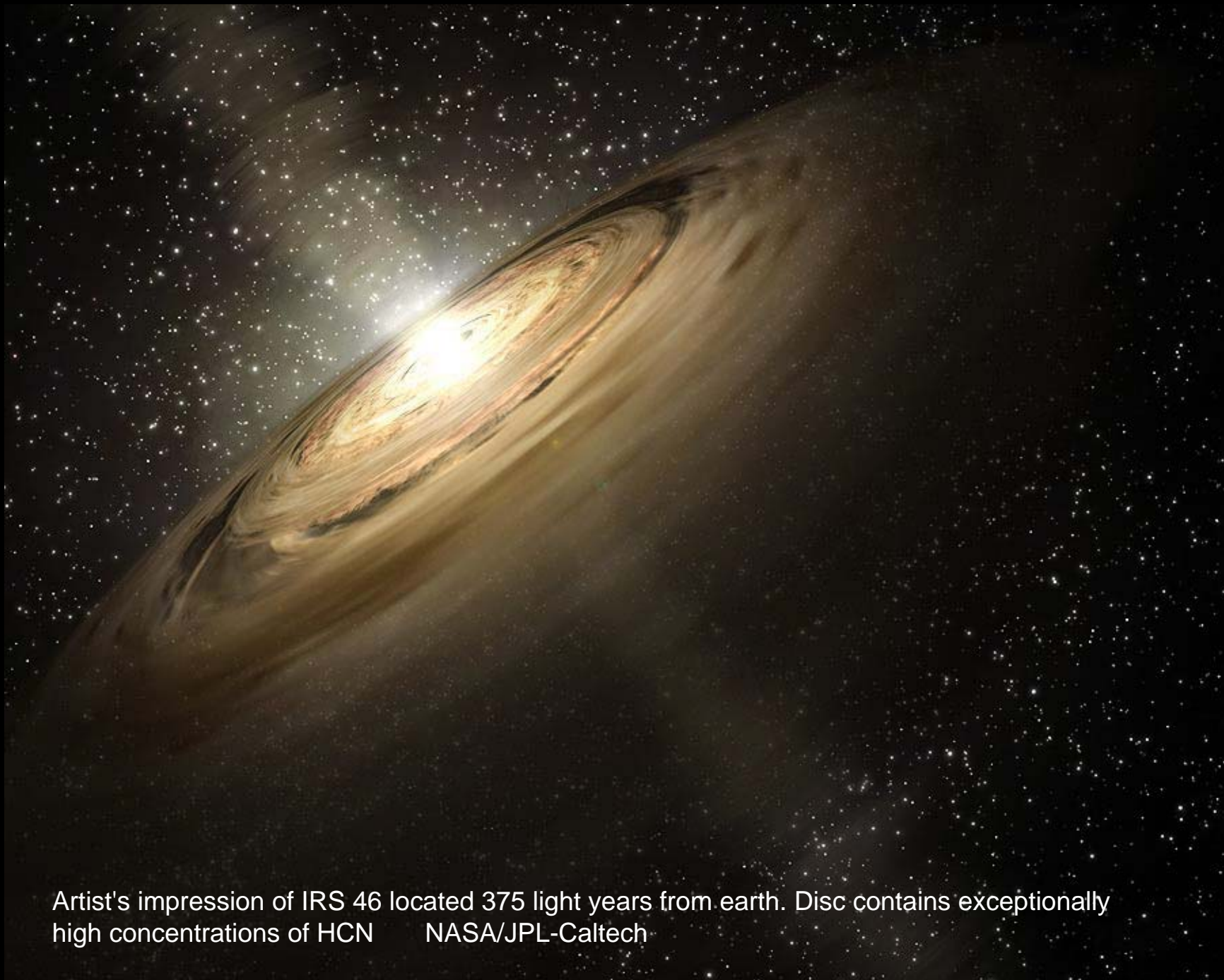
Guanine



Uracil



Cytosine



Artist's impression of IRS 46 located 375 light years from earth. Disc contains exceptionally high concentrations of HCN      NASA/JPL-Caltech

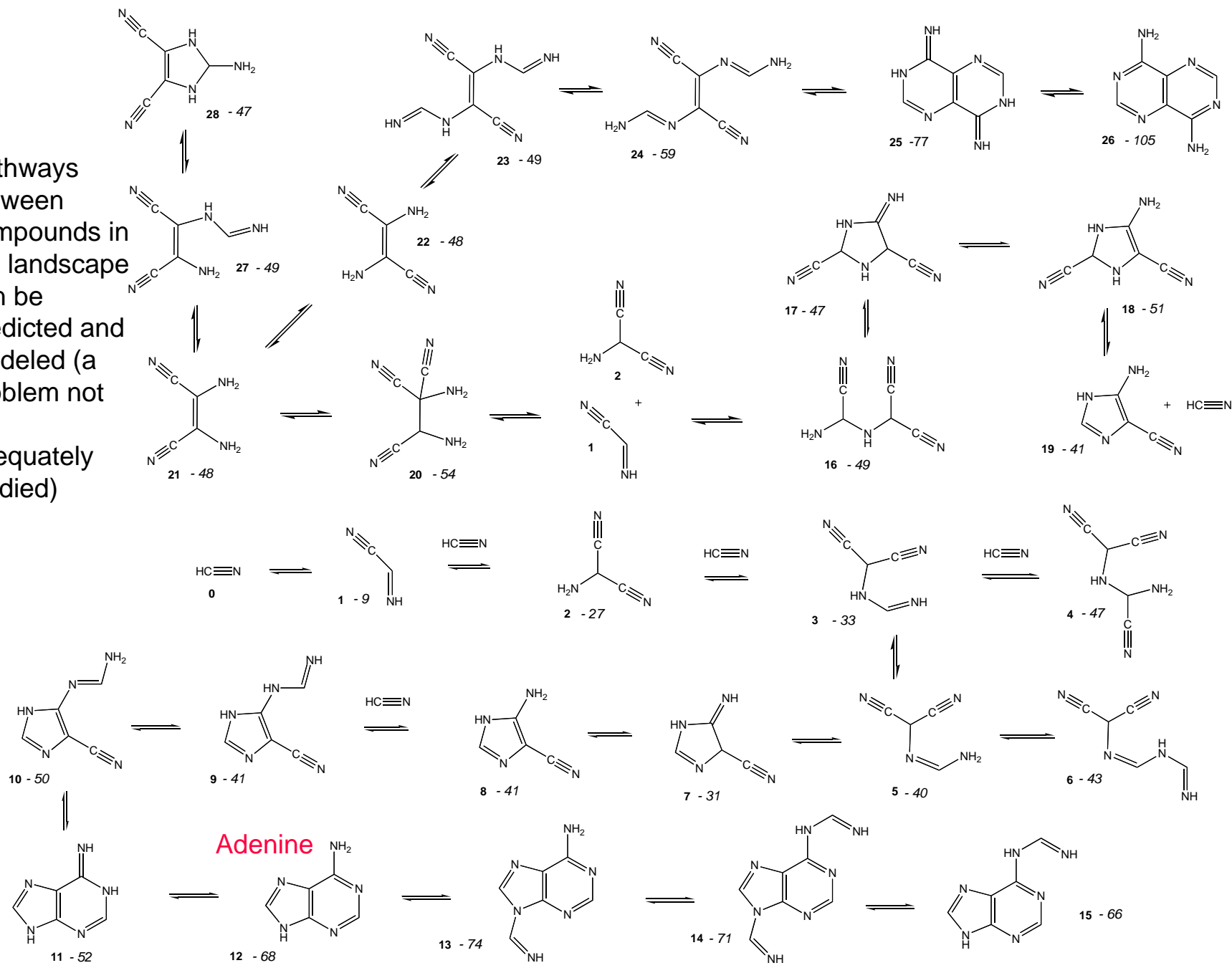
## HCN polymerization

*At sufficiently high concentrations HCN will polymerize to a complex mixture of products. The products and pathways will be influenced by the presence of other elements and catalysts as well as temperature and pressure.*

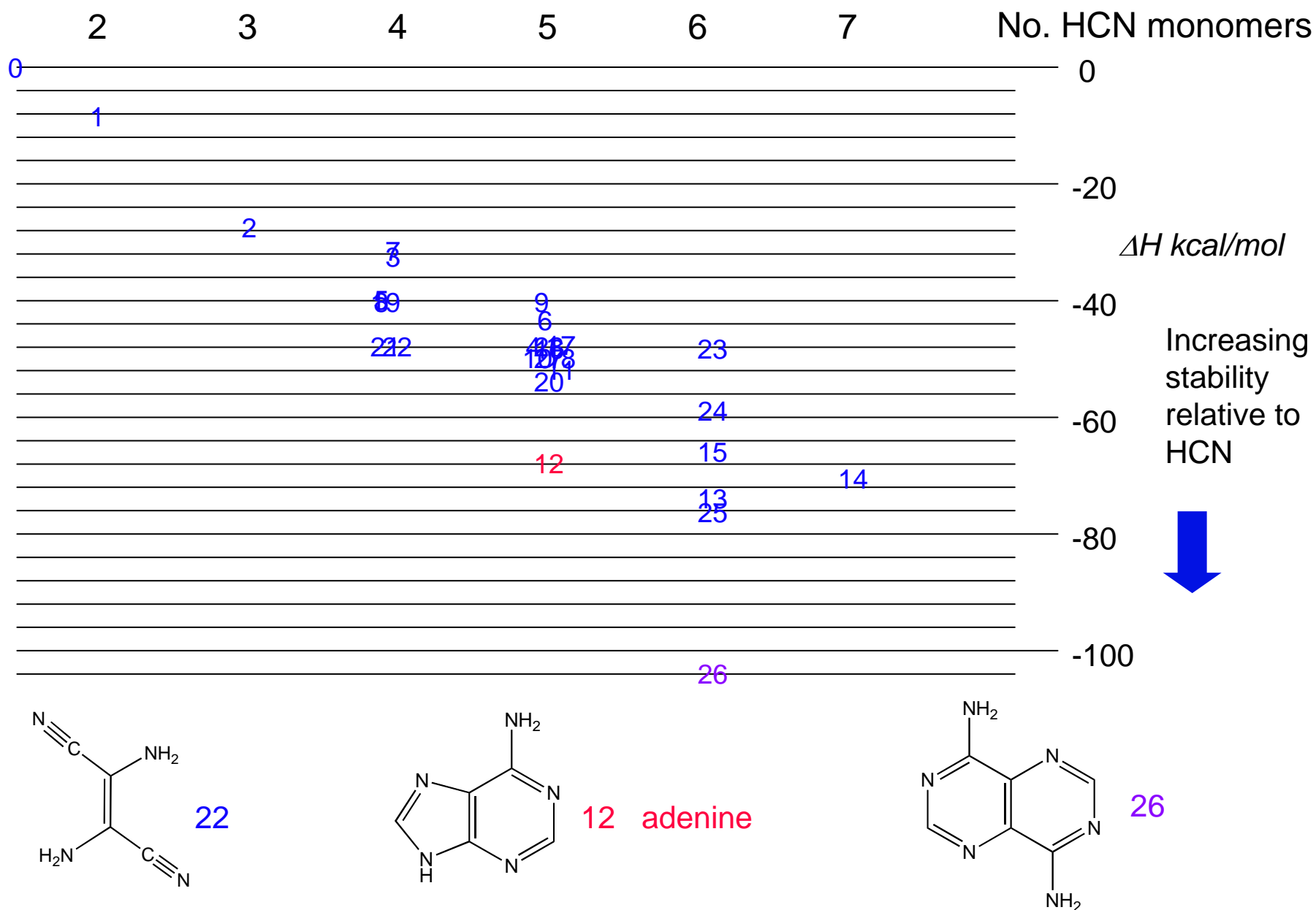


# Portion of hypothetical HCN polymerization landscape

Pathways between compounds in the landscape can be predicted and modeled (a problem not yet adequately studied)



# HCN polymerization landscape

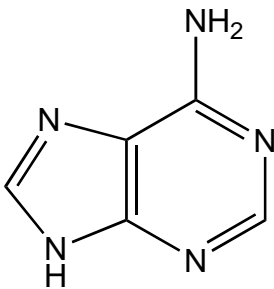


## The path from adenine to adenosine phosphates

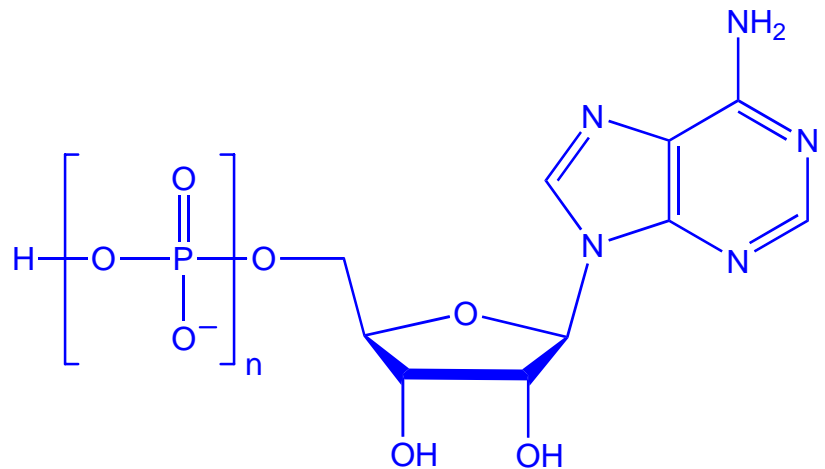
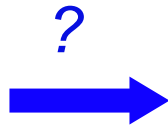
Adenine will be pulled from the HCN domain by interactions with other molecular domains (HCO and polyphosphate). According to *Le Chatelier's* principle other molecules in equilibrium with adenine will be pulled from the landscape as a result. In essence they become fuel for adenine.

### *Le Chatelier's principle*

*If a chemical system at equilibrium experiences a change in concentration, temperature, volume, or total pressure; the equilibrium will shift in order to minimize that change.*



Adenine

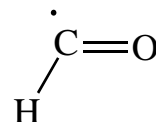


AMP, ADP, ATP, Ap<sub>n</sub>A

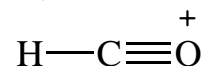
# Interstellar CHO species

COH	formyl radical
HCO <sup>+</sup>	formyl ion
HOC <sup>+</sup>	hydroxymethylidynium
HOCO	hydroxyformyl radical
HOCO <sup>+</sup>	protonated carbon dioxide
H <sub>2</sub> CO	formaldehyde
HCOOH	formic acid
CH <sub>3</sub> O	methoxy radical
H <sub>2</sub> COH <sup>+</sup>	protonated formaldehyde
CH <sub>3</sub> OH	methanol
H <sub>2</sub> CCO	ketene
CH <sub>3</sub> CHO	acetaldehyde
CH <sub>2</sub> CHOH	vinyl alcohol
c-C <sub>2</sub> H <sub>4</sub> O	ethylene oxide
HCOOCH <sub>3</sub>	methyl formate
CH <sub>3</sub> COOH	acetic acid
CH <sub>2</sub> OHCHO	glycolaldehyde
CH <sub>3</sub> CH <sub>2</sub> OH	ethanol
CH <sub>3</sub> OCH <sub>3</sub>	dimethyl ether
HC <sub>2</sub> CHO	propynal
(H <sub>2</sub> CO) <sub>3</sub>	trioxane
CH <sub>3</sub> COCH <sub>3</sub>	acetone

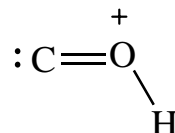
Formyl radical



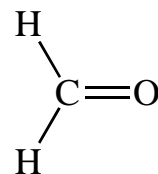
Formyl ion



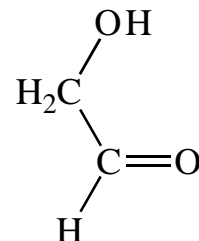
Hydroxymethylidynium



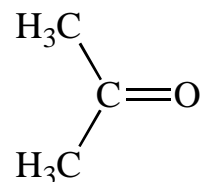
Formaldehyde



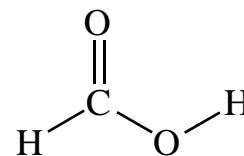
Glycolaldehyde



Acetone

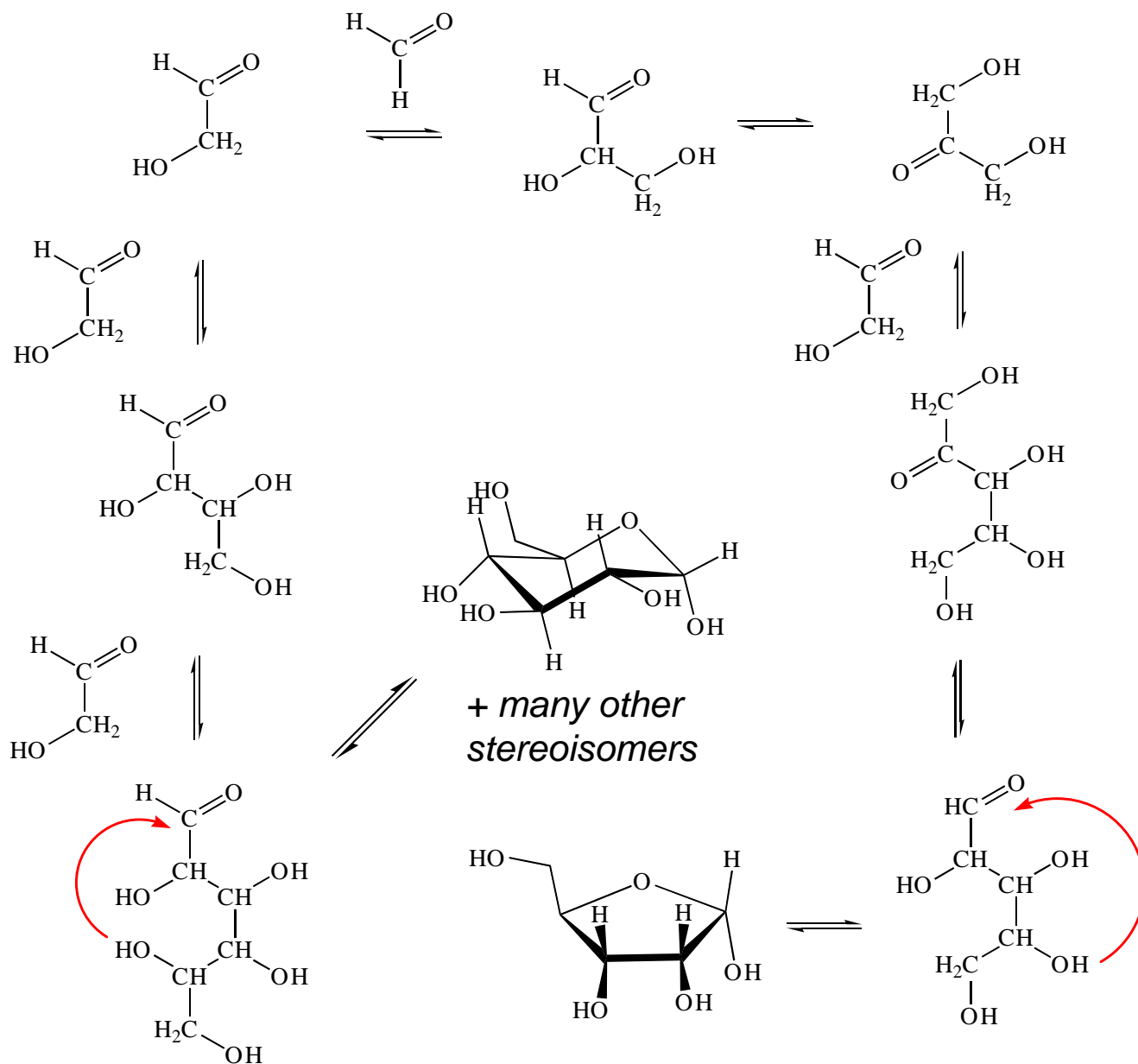


Formic acid

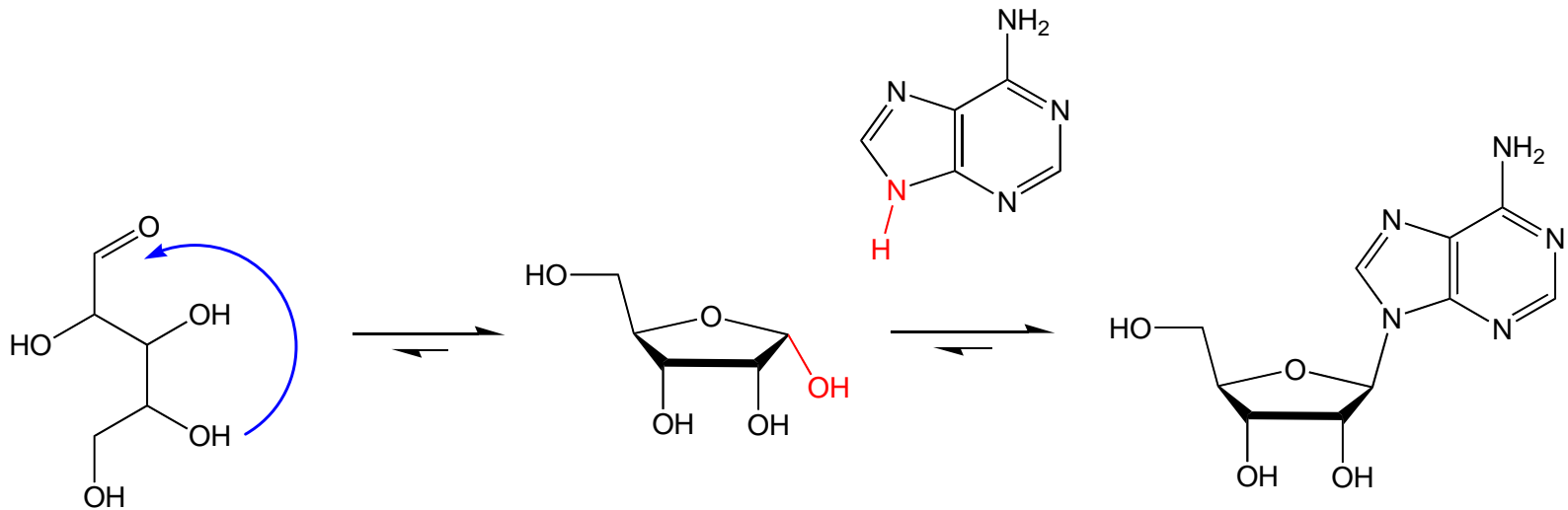


From **Constants for molecules of astrophysical interest in the gas phase: photodissociation, microwave and infrared spectra**. J. Crovisier Observatoire de Paris, Section de Meudon F-92195 Meudon, France  
<http://www.lesia.obspm.fr/~crovisier/basemole/>

# Reaction pathways to monosaccharides

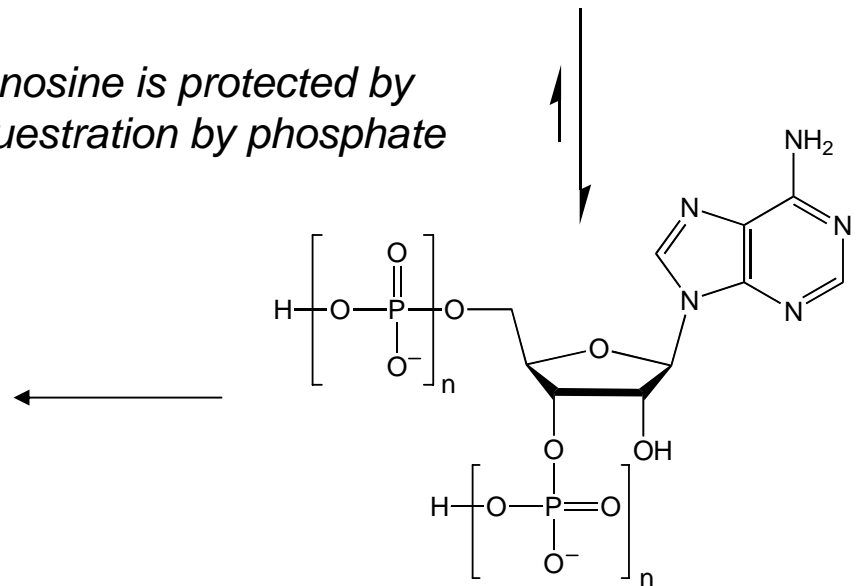


# Path to an adenosine centric world



*Adenosine is protected by sequestration by phosphate*

*An adenosine centric world evolves as networks with other molecules are established; eventually leading to the RNA world*

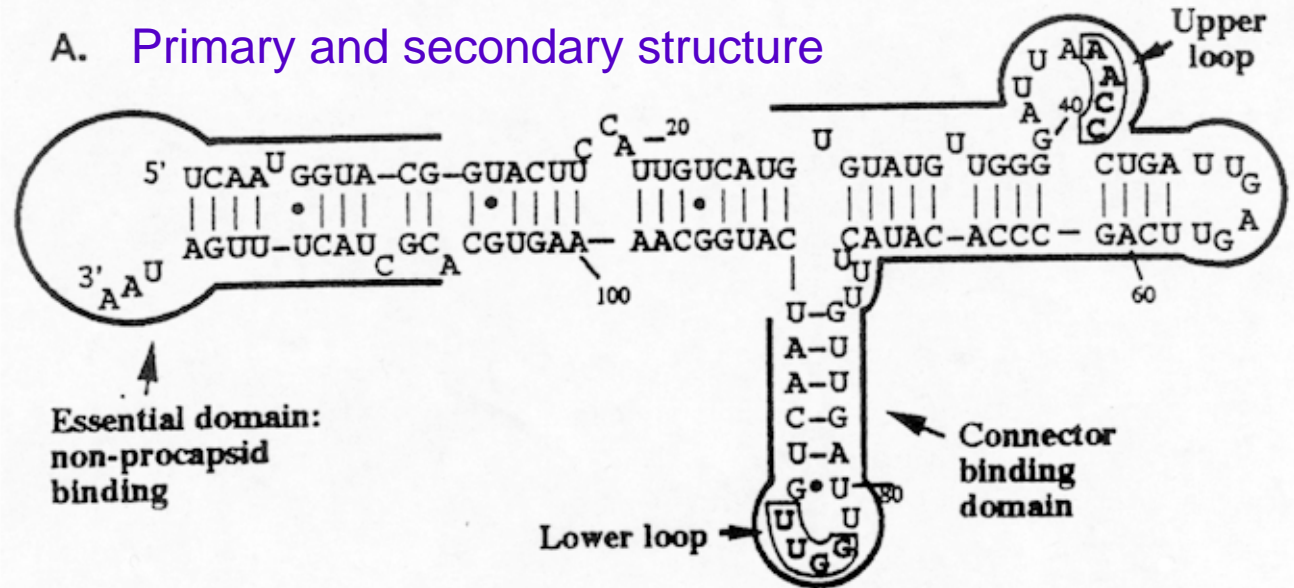


# The path from adenine continues to unfold

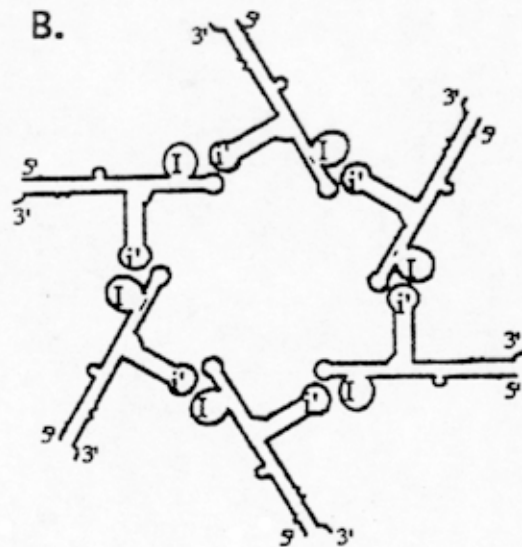
1. Adenine captures monosaccharides
2. Adenosine recruits phosphate and other small molecules in the environment
3. Coenzymes evolve to aid adenosine survival
4. The RNA world is established
5. RNA and peptides co-evolve
6. The translation system evolves
7. DNA is established and co-evolves with RNA and proteins
8. Higher order organisms evolve

# RNA can assemble into higher order structures

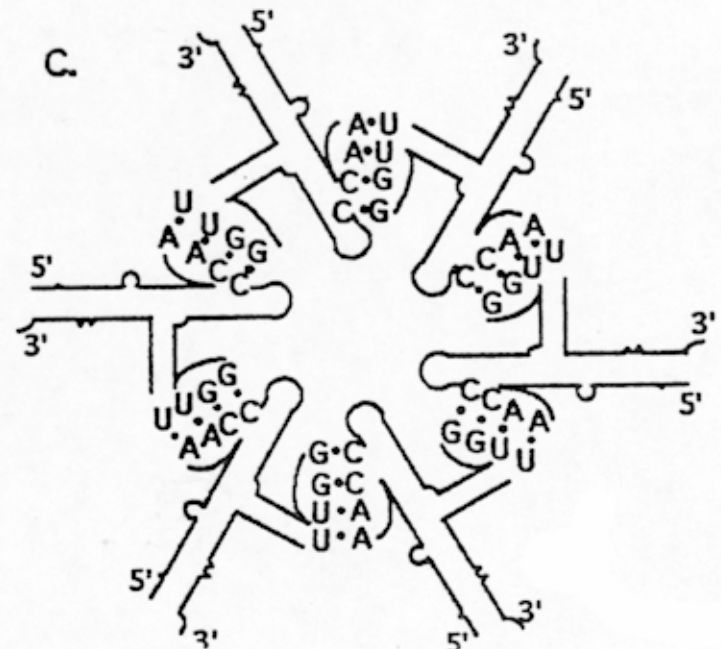
## A. Primary and secondary structure



## B.



## C.



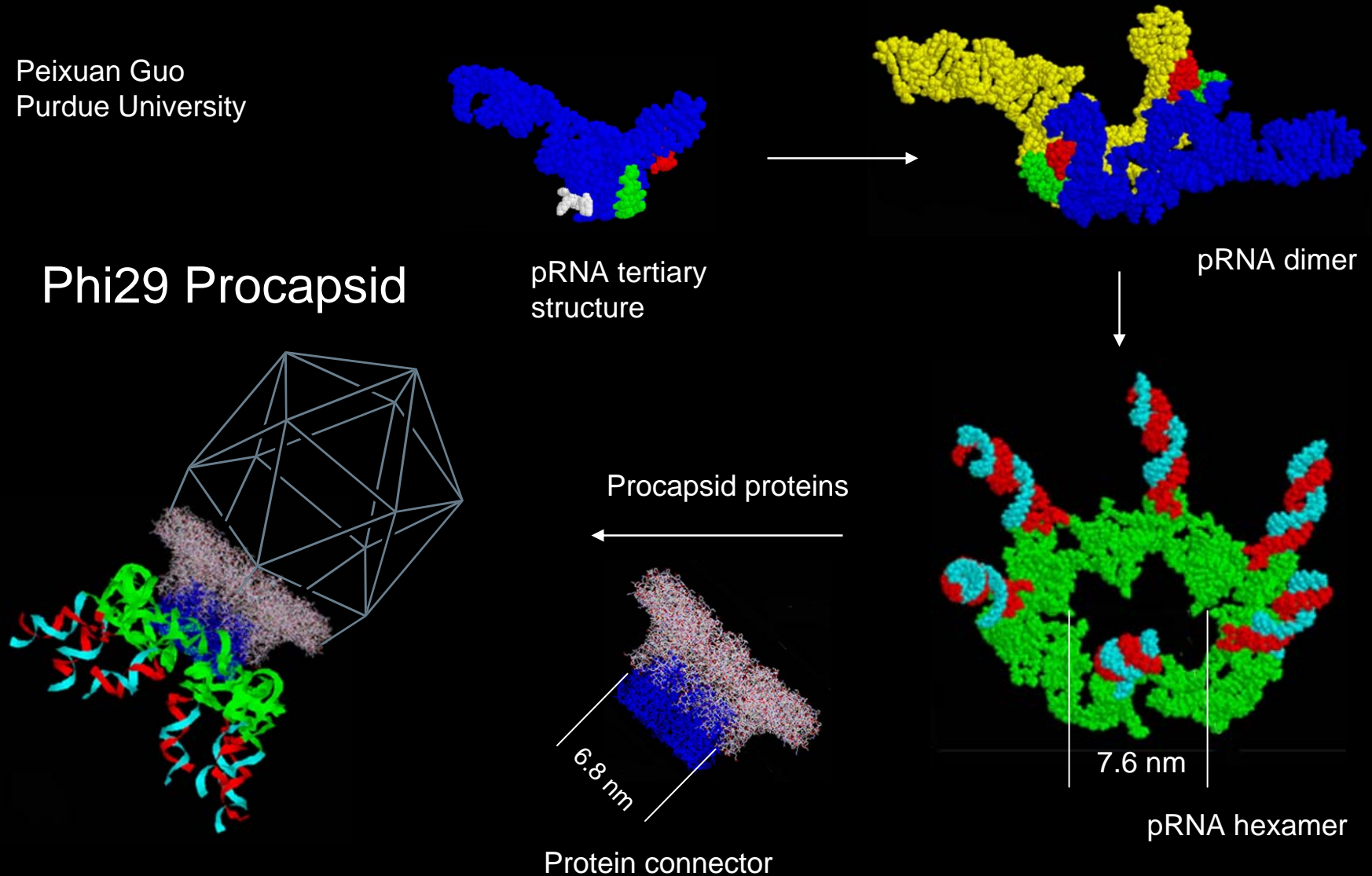
pRNA hexamer  
from the  
bacteriophage  
phi29

Peixuan Guo  
Purdue University



# pRNA hexamer assembles into a molecular motor for packaging DNA into the phi29 procapsid

Peixuan Guo  
Purdue University



# The path from adenine continues to unfold - 2

1. Adenine captures monosaccharides
2. Adenosine recruits phosphate and other small molecules in the environment
3. Coenzymes evolve to aid adenosine survival
4. The RNA world is established
5. RNA and peptides co-evolve
6. The translation system evolves
7. DNA is established and co-evolves with RNA and proteins
8. Higher order organisms evolve
9. Humans are recruited to develop adenine derived materials

# Why nucleic acids are important materials for nanotechnology?

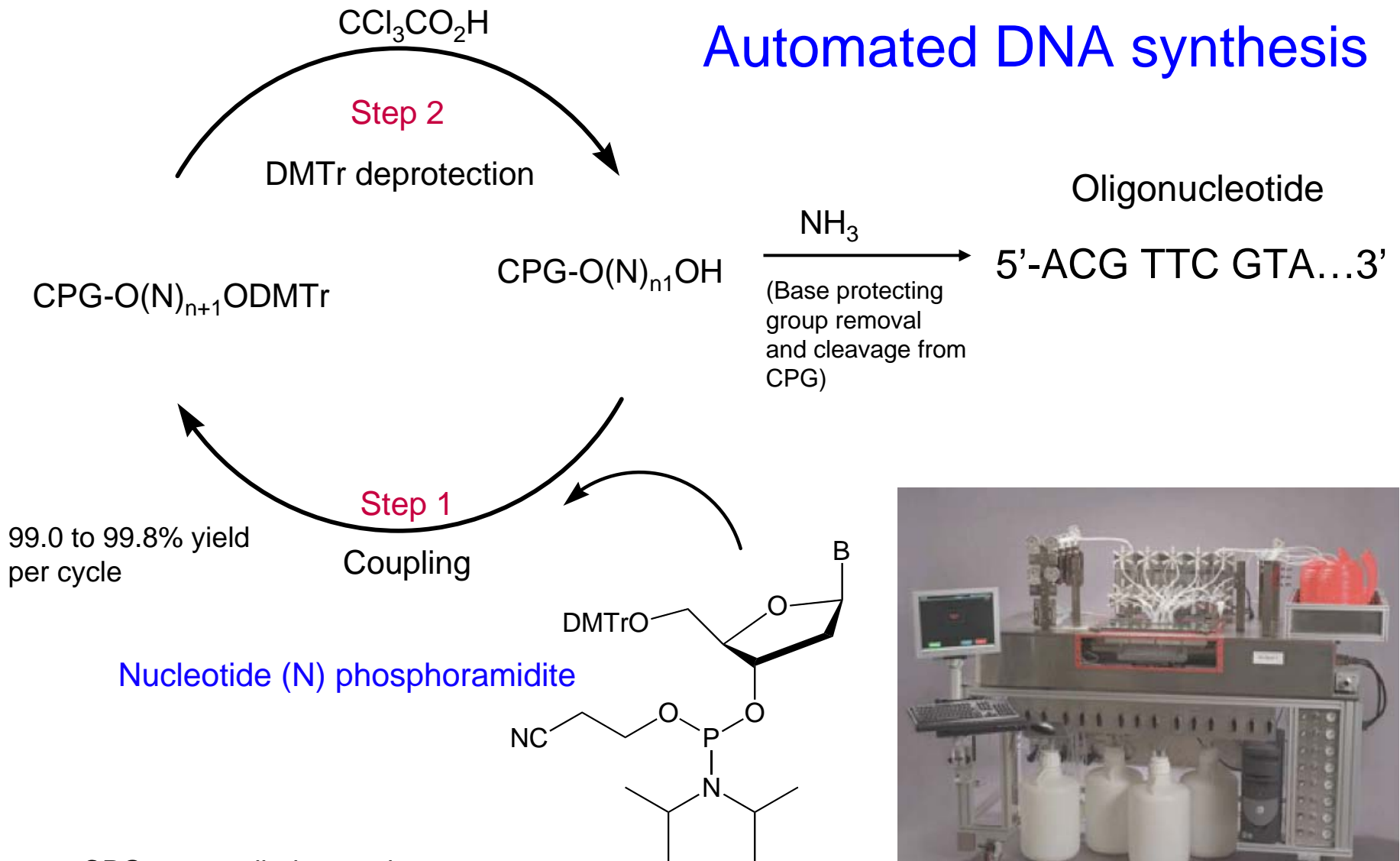
1. High sensitivity identification of any organism by nucleic acid sequence will be most efficiently accomplished with nanoscale devices.
2. High-throughput sequencing of the human genome for genetic diseases will only be economical if done with nanoscale technologies.
3. Nanoscale devices carrying nucleic acid components will be needed for nucleic acid based drugs and gene therapies.
4. Synthetic modified nucleic acids constitute unique materials for self-assembly of multi-functional nanoscale devices.
5. Nucleic acids can be used as components in nano-scale computing strategies.

*A dominate theme in 21st century technology will be the merging of biology and material science.*

# Advantages of Synthetic Nucleic Acids as Engineering Materials

- Automated high-throughput synthesis is fast and inexpensive
- Chemically stable and biologically compatible
- Sequence-specific self-assembly into stable duplex and higher order structures
- Probes and functional groups can be attached at many different sites
- Structure can be extensively modified yet retain self-assembly properties
- Sequences can be attached to virtually any material
- Electrical and optical properties can be tuned

# Automated DNA synthesis

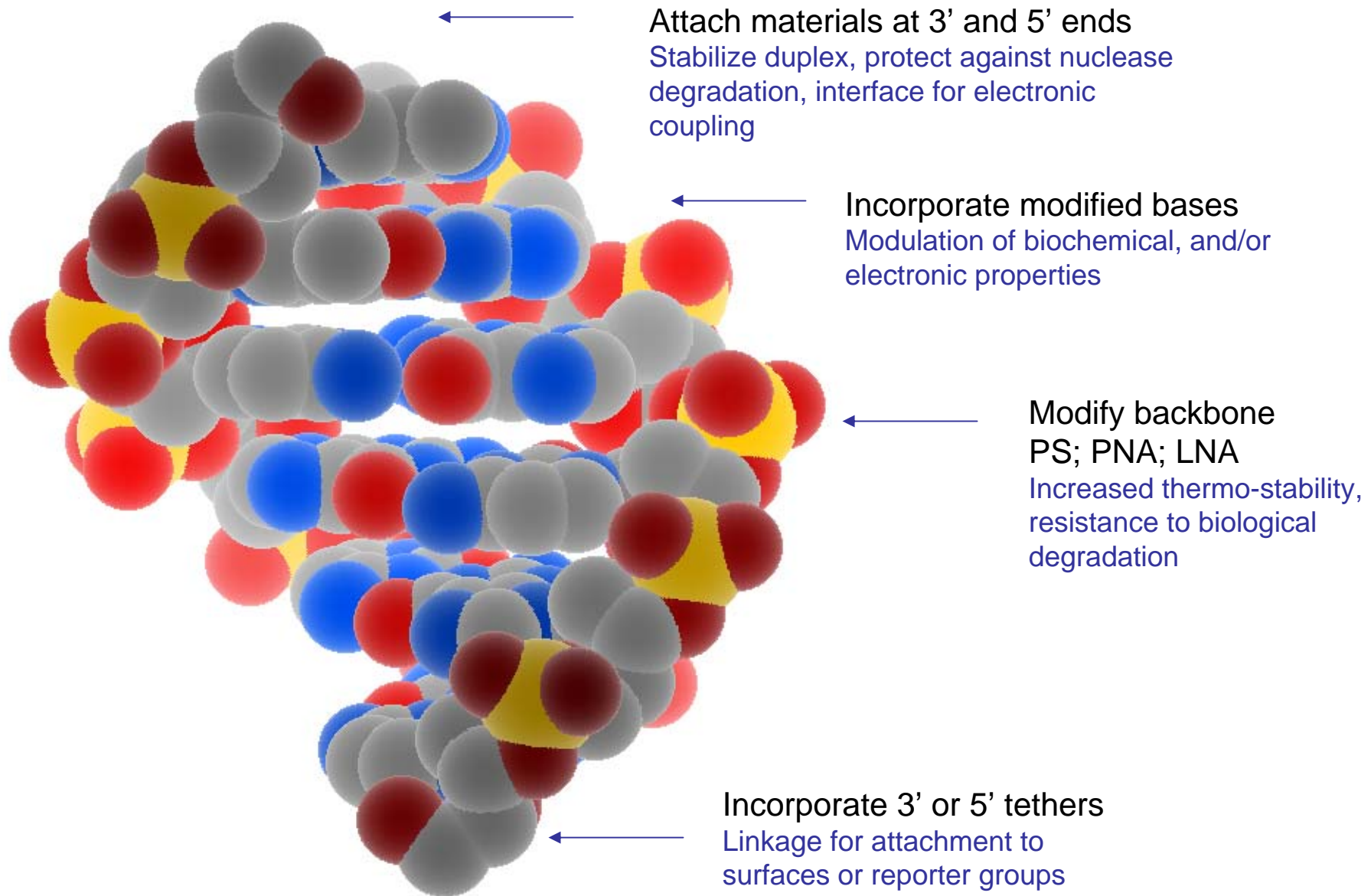


384 oligonucleotide parallel synthesizer

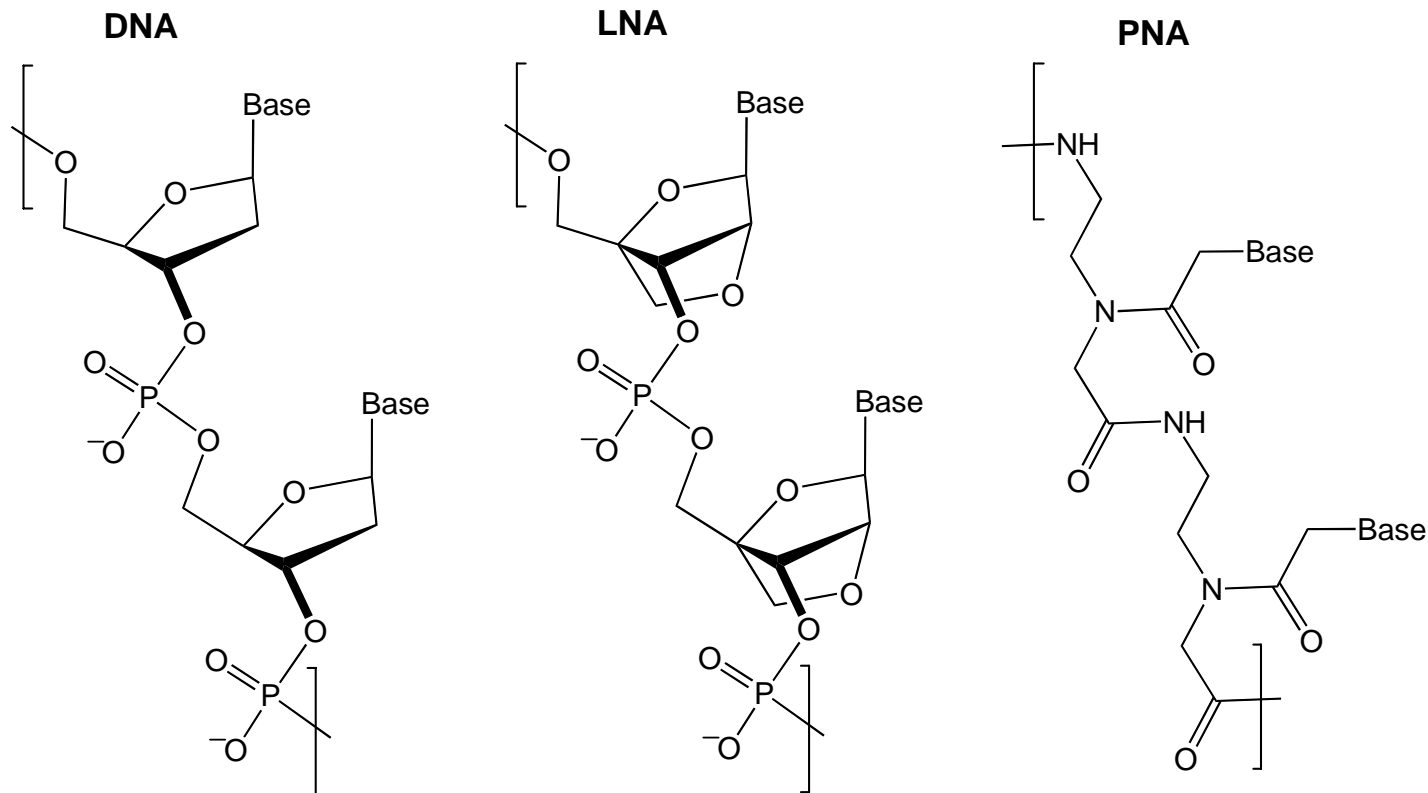
CPG = controlled pore glass

DMTr = dimethoxytrityl

B = protected adenine, cytosine, guanine, thymine, or modified base

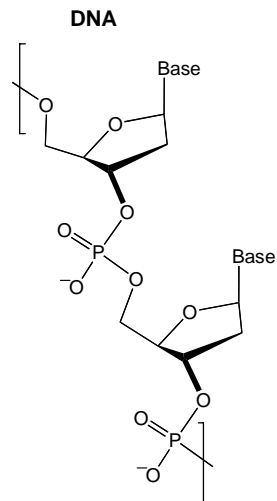
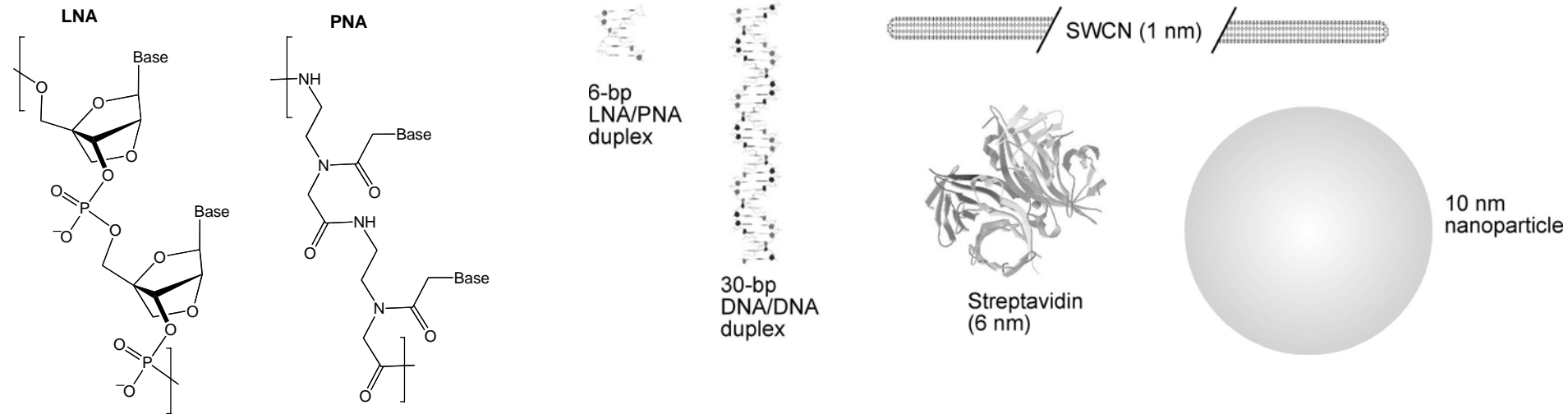


Chemists have created many other nucleic acid-like polymers that self-assemble

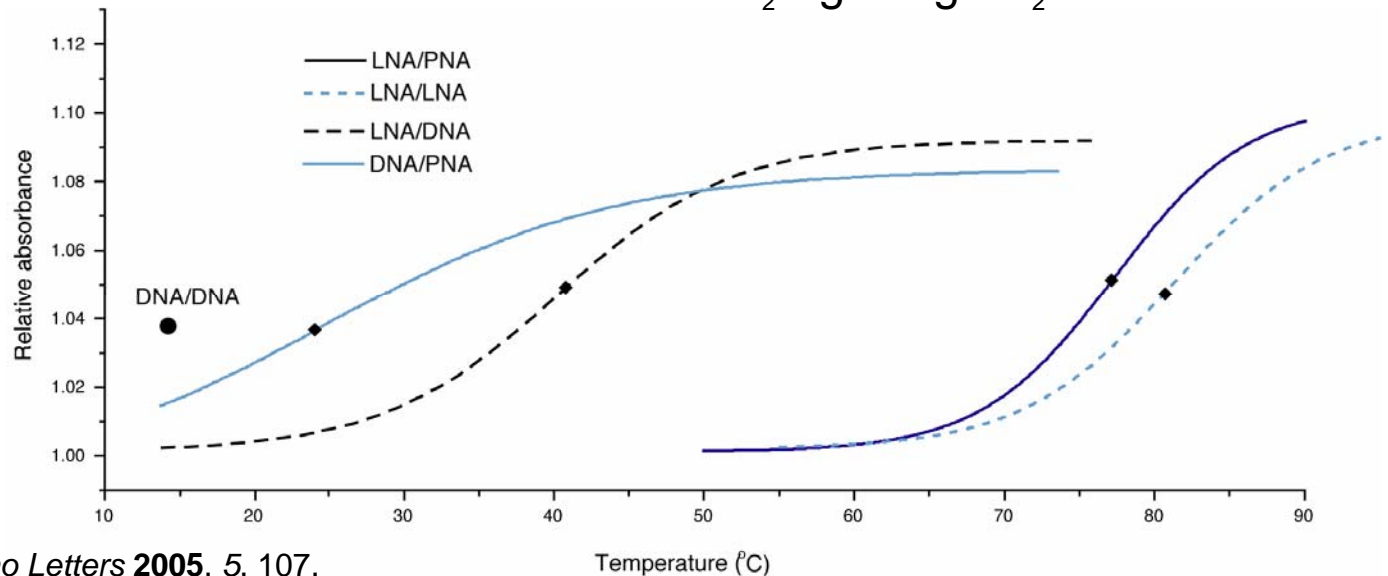


*Locked nucleic acids (LNA) and peptide nucleic acids (PNA) form duplex structures that are considerably more stable than DNA and RNA*

# Alternative nucleic acid analogs for programmable assembly: Hybridization of LNA to PNA



5'-CGATGC-3'/N<sub>2</sub>H-gcatcg-CO<sub>2</sub>H





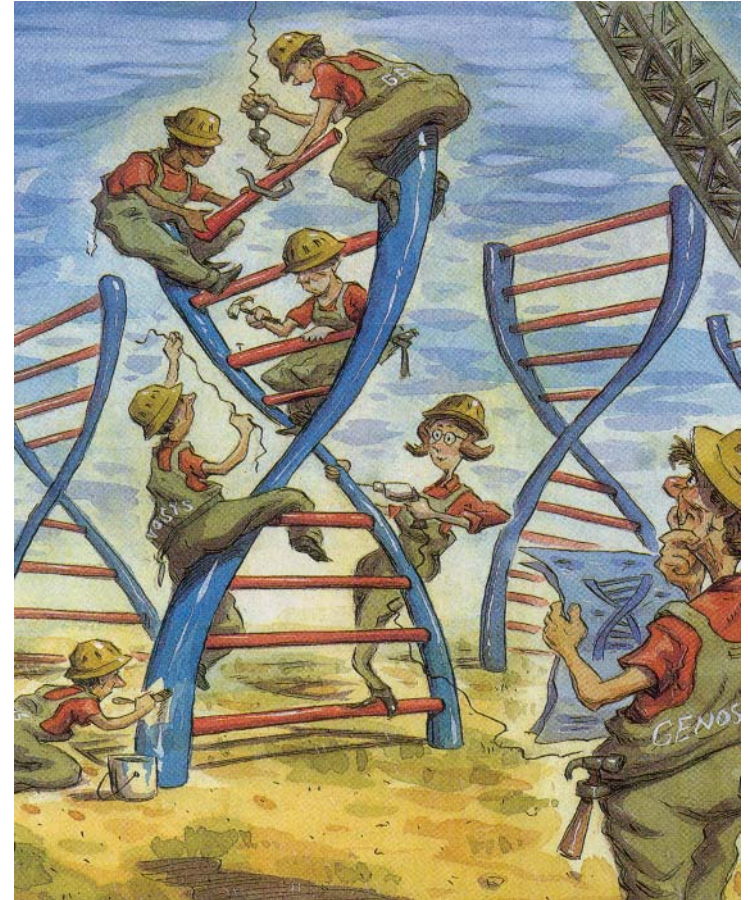
# Re-engineering nucleic acids and nucleic acid components: future opportunities for scientists and engineers

## Biologicals

- Tools for investigations of enzyme mechanism
- Tools for structural studies on protein-DNA complexes
- Components of nucleic acid based assay systems
- Diagnostic probes
- Tools for drug target validation
- Therapeutics
  - Antisense oligonucleotides
  - Decoy oligonucleotides
  - Modified oligonucleotides for gene correction
  - Sequence specific prodrug activators

## Materials

- Self-assembling scaffolds for nanoscale devices
- Assemblers for reaction discovery
- Computing



# Questions & Answers