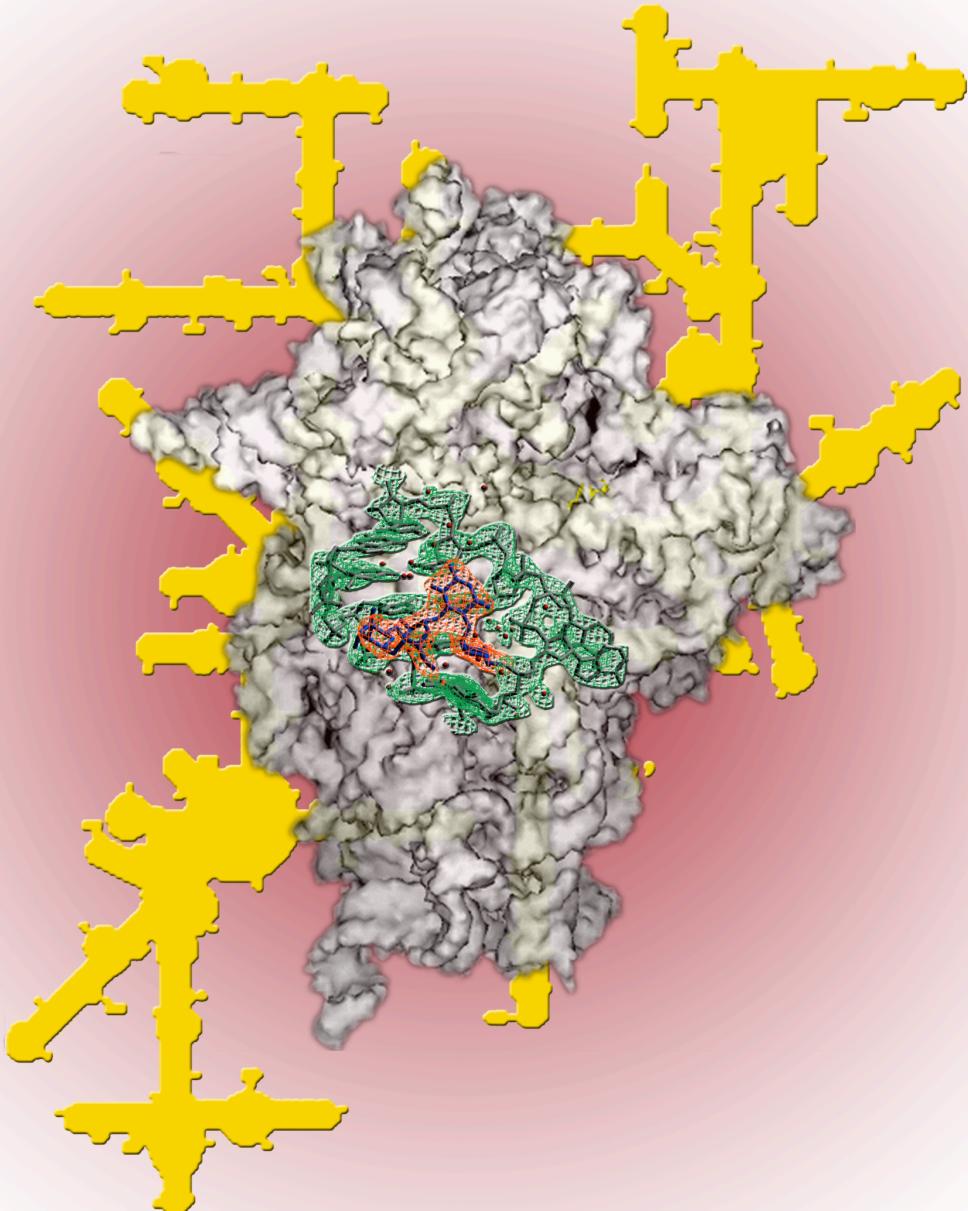


The

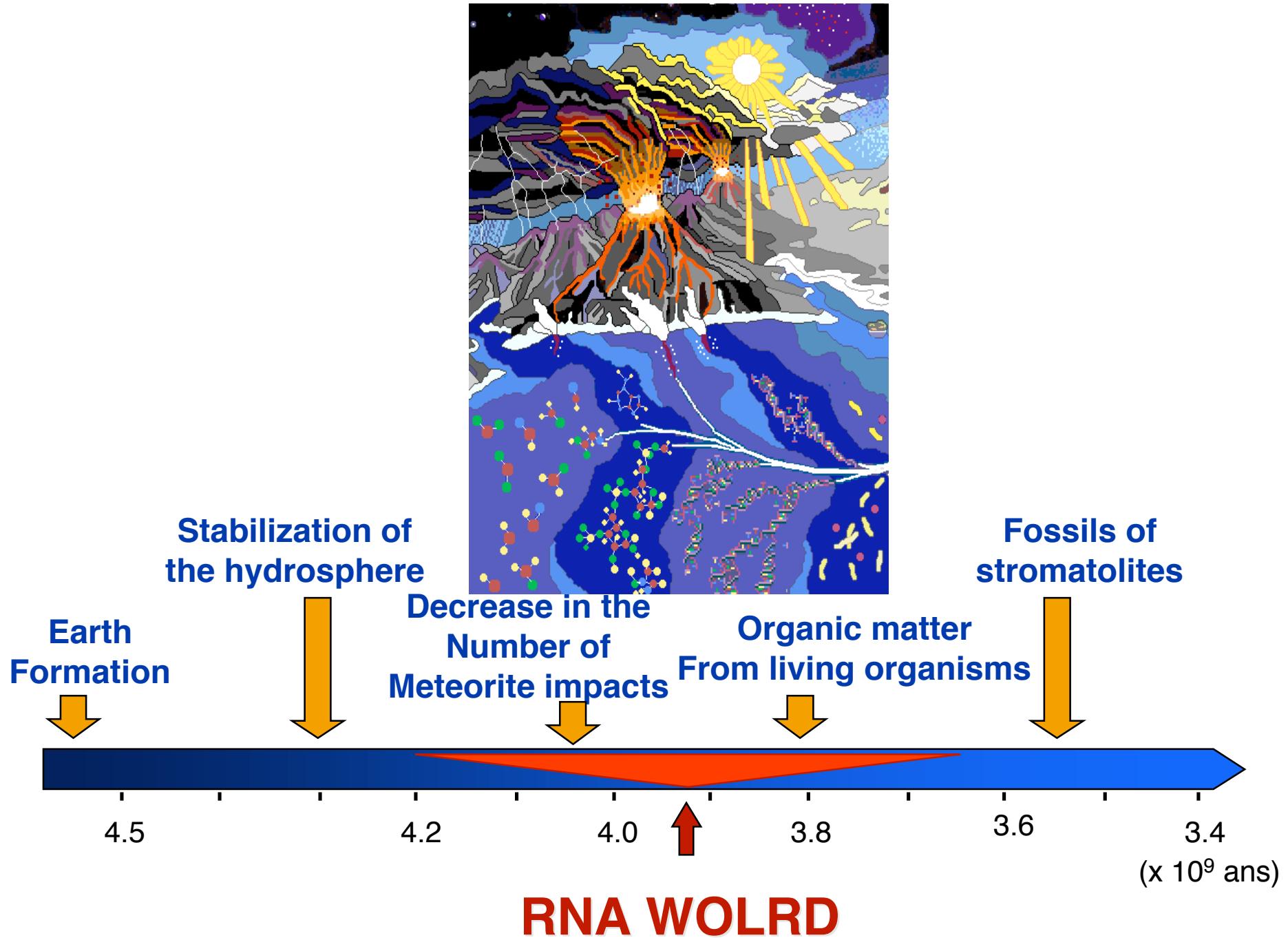
RNA

World

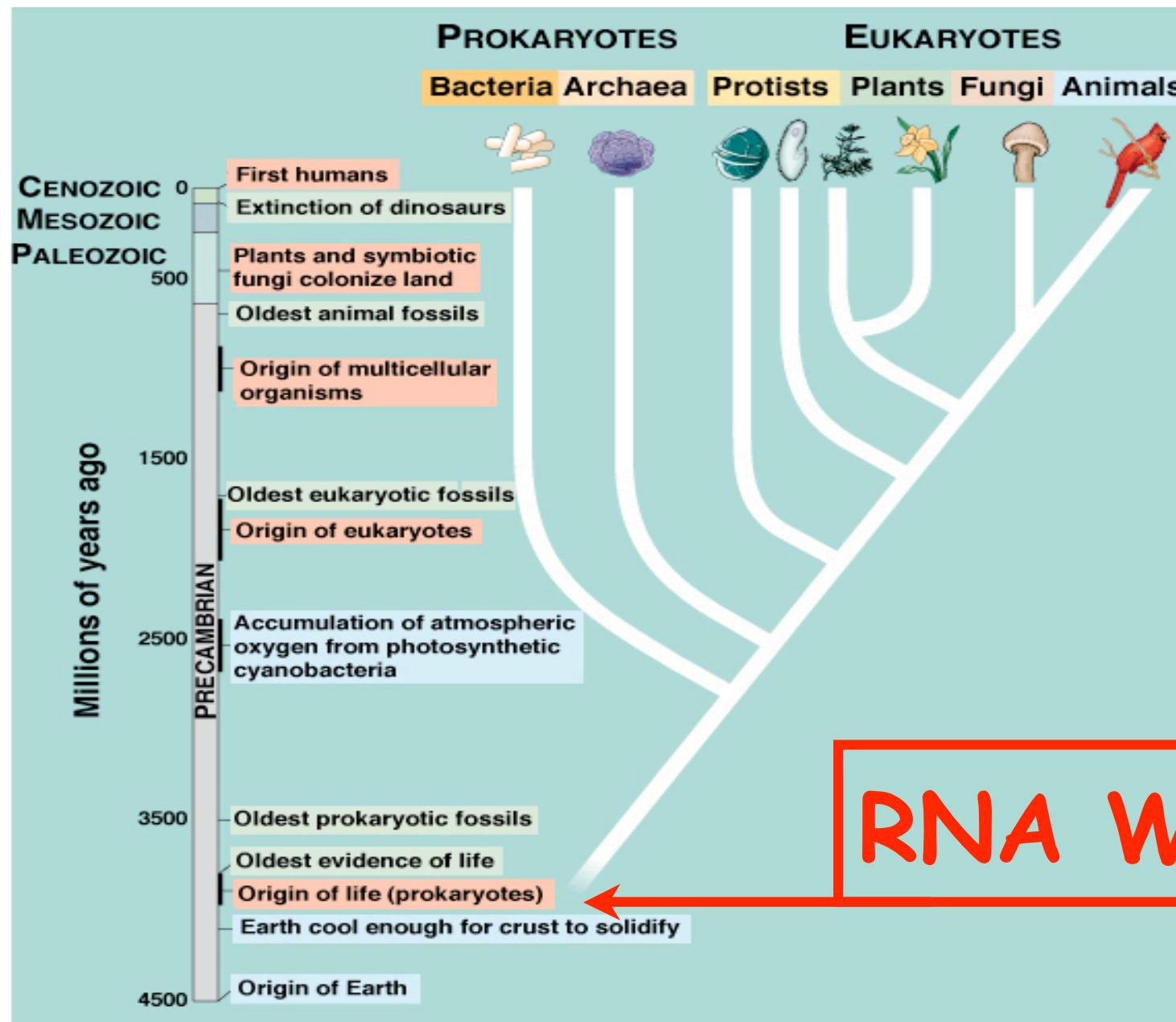
E. Westhof



<http://www-ibmc.u-strasbg.fr/upr9002/westhof/>



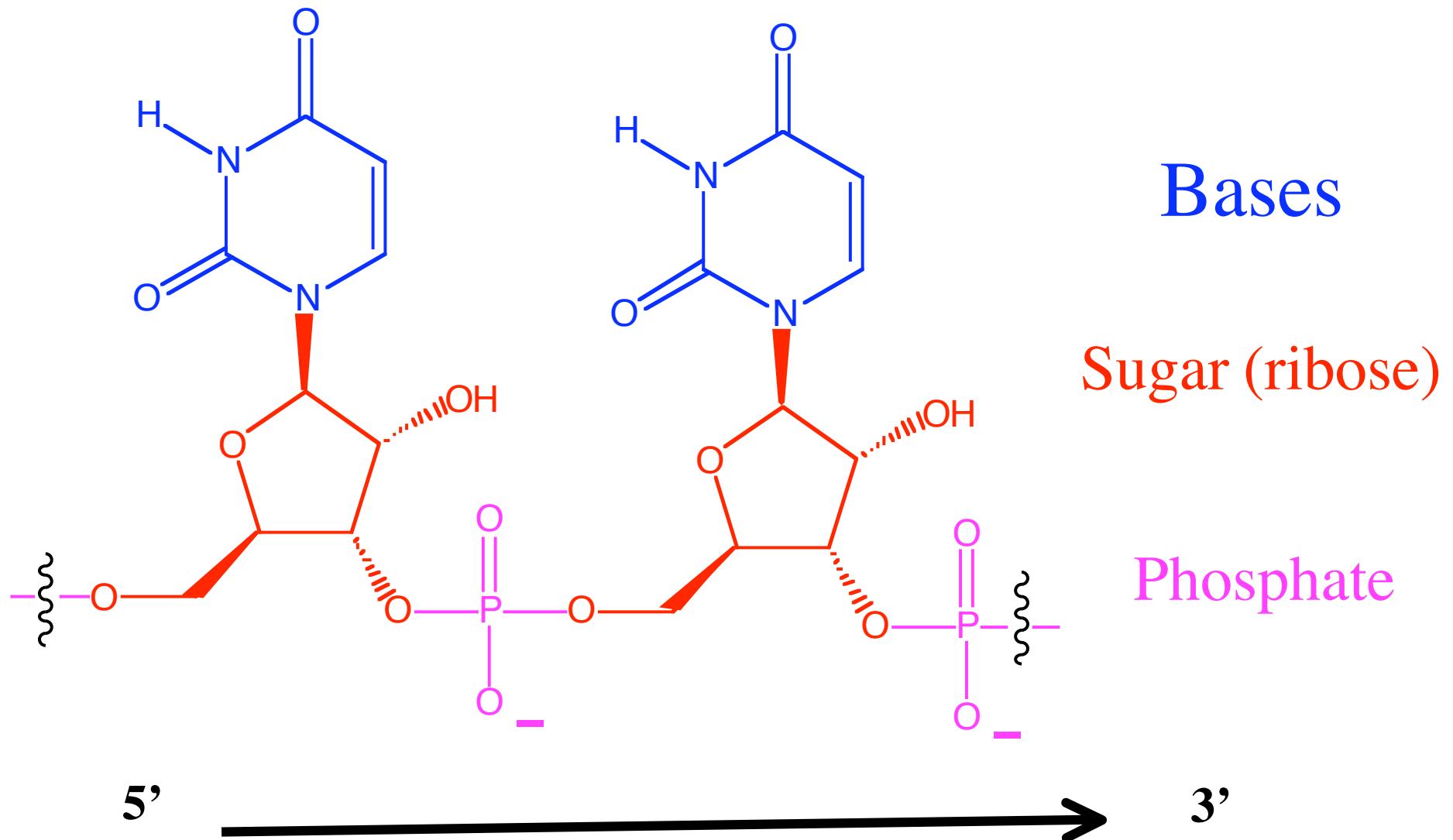
Global overview of all life



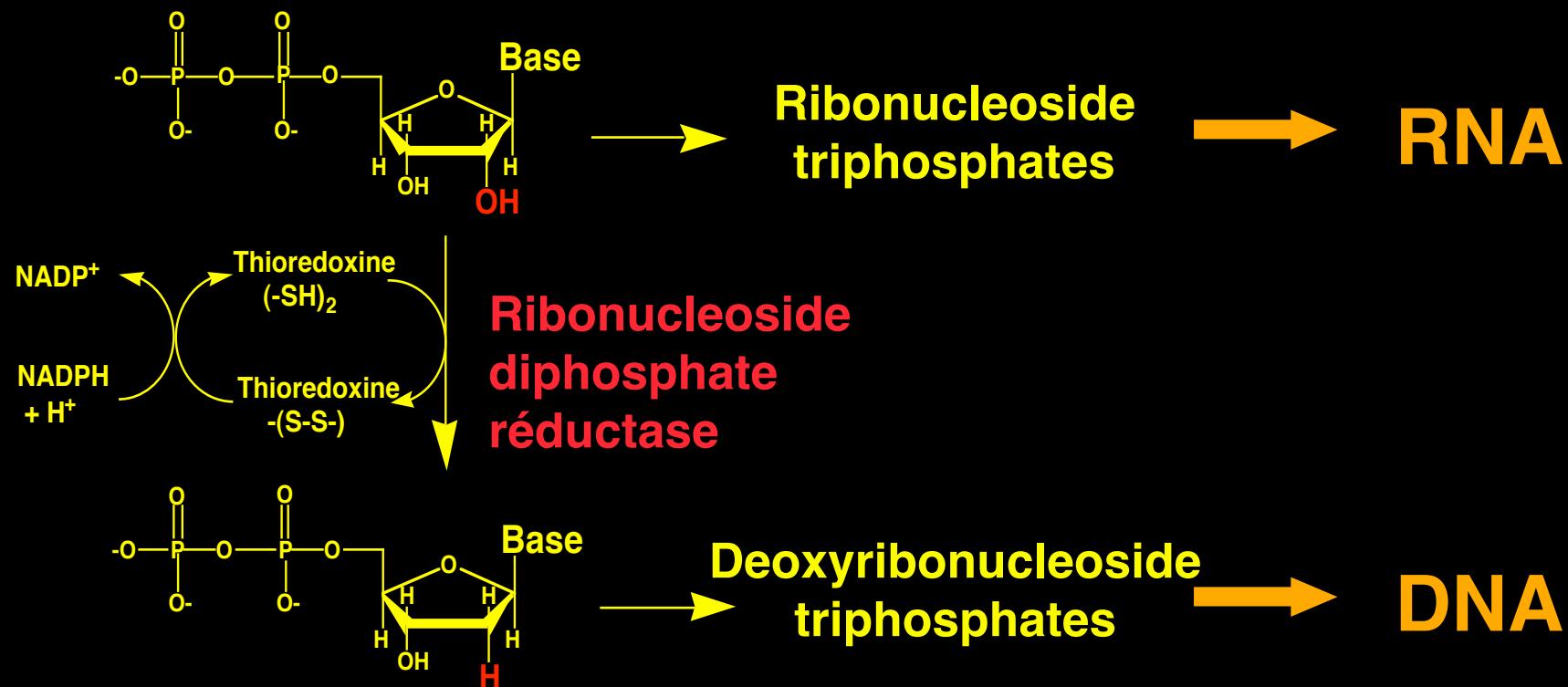
The RNA World

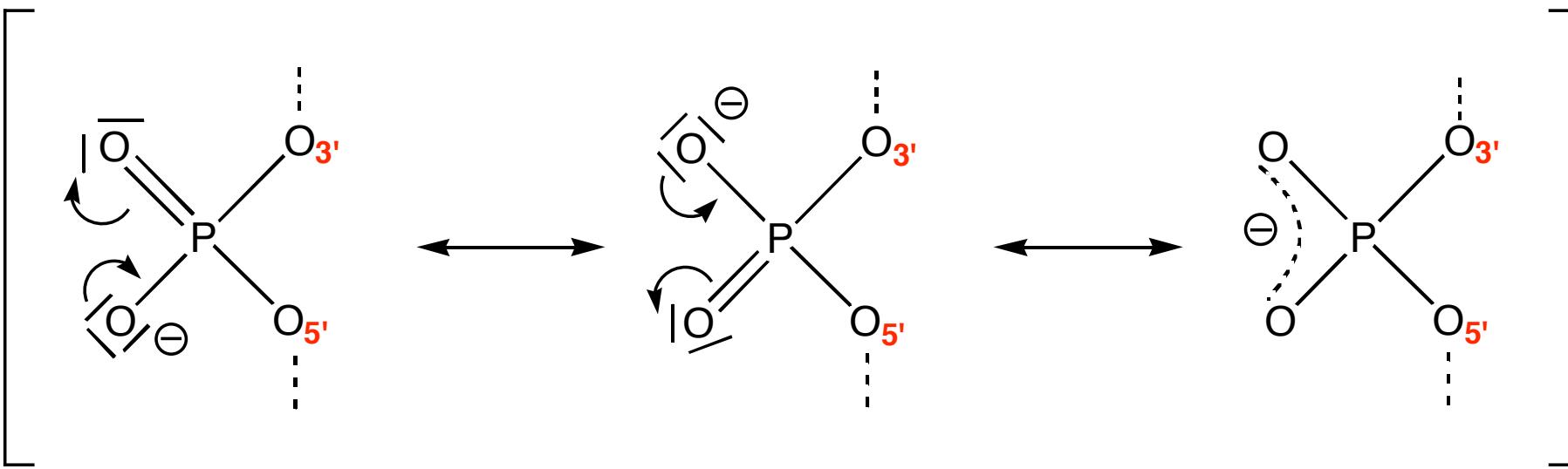
- Origin of life / central dogma paradox
 - DNA needs proteins to replicate
 - Proteins coded for by DNA
- RNA can be code and machinery
 - Selex, aptamers
- RNAs are remnants
 - Ancient and Essential

RNA, Parts

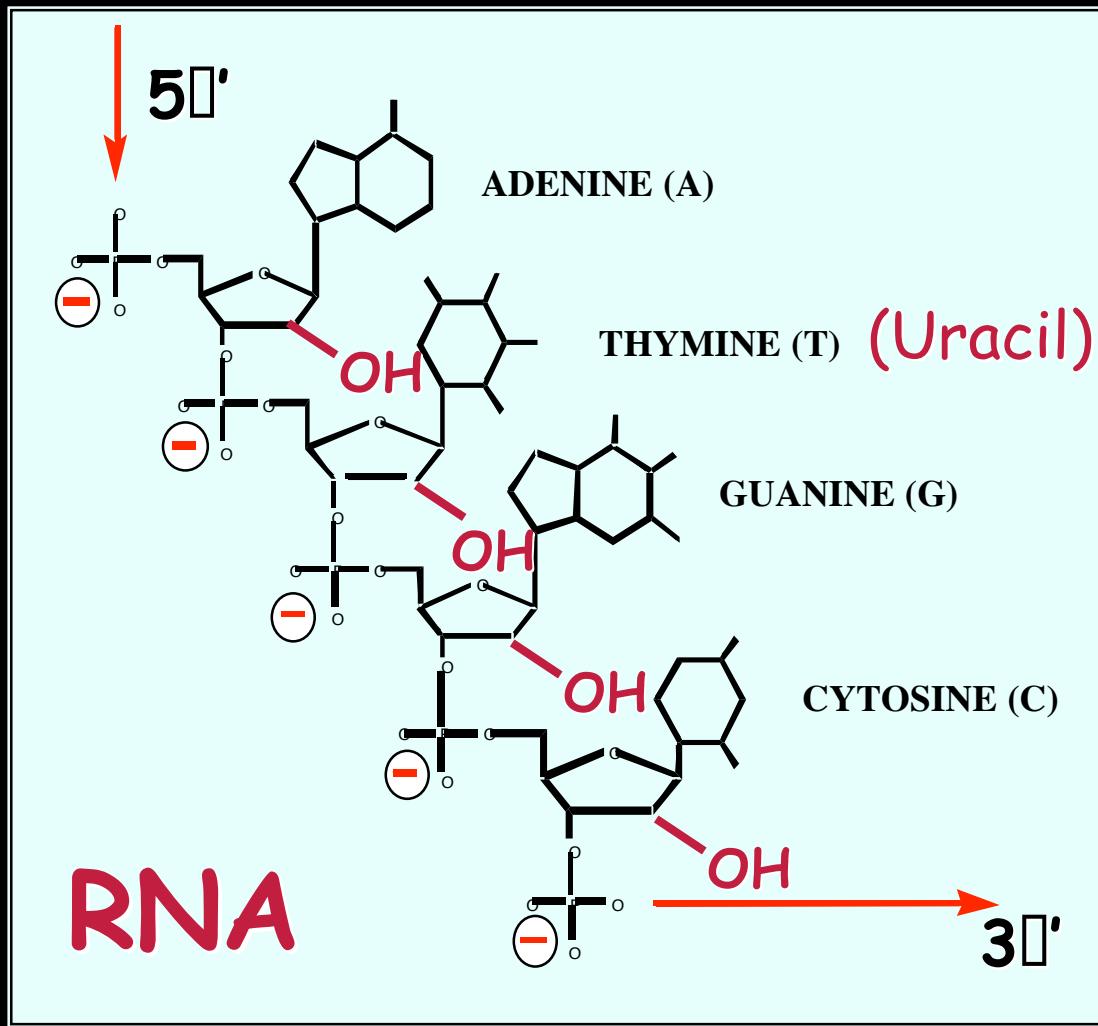


DNA = modified RNA



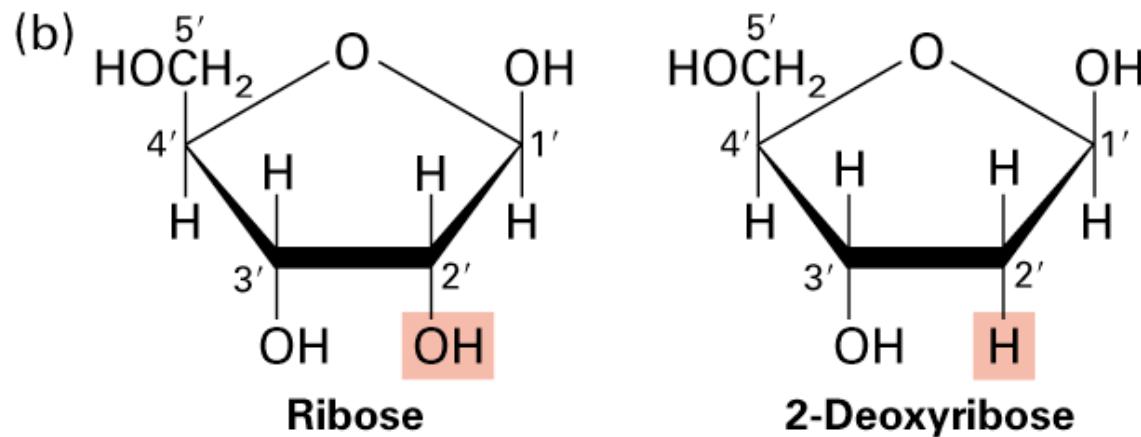


Nucleic acids are negatively charged biopolymers ...

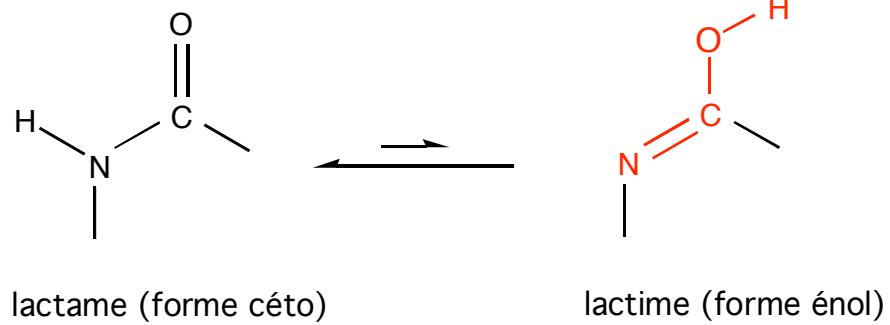
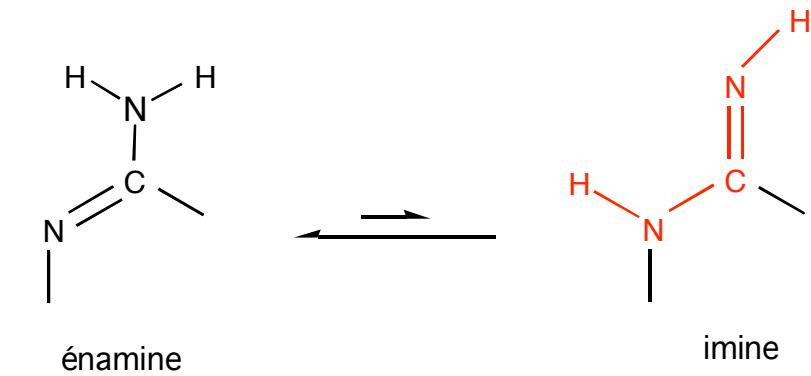


Conformations of RNA

- Primary structure of RNA similar to DNA

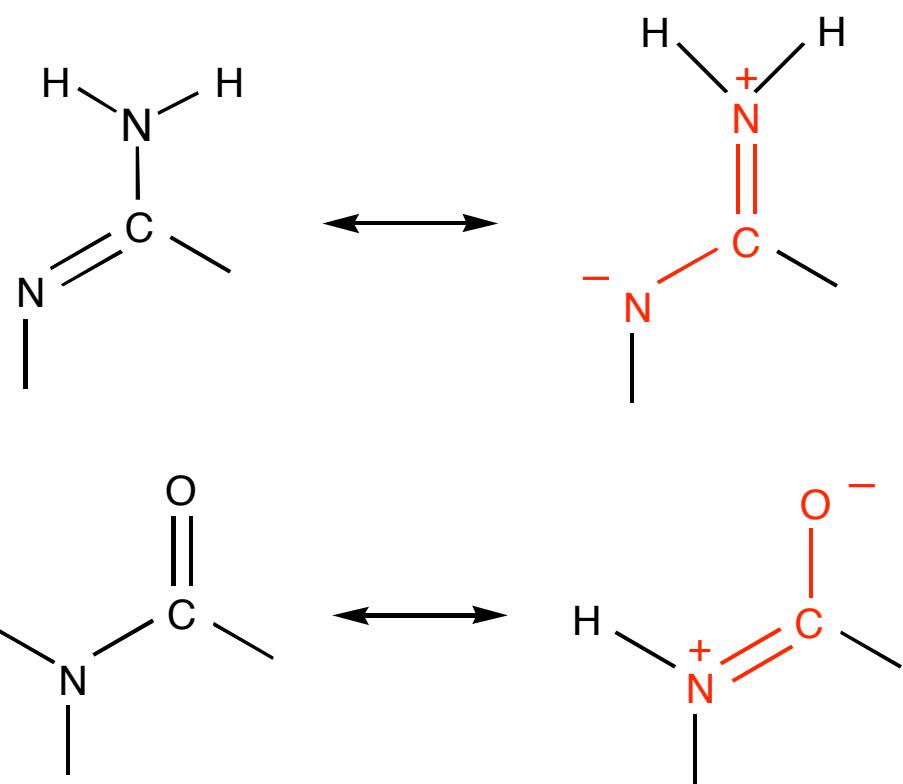


- RNA, like DNA, can be single or double stranded, linear or circular.
- Unlike DNA, RNA can exhibit different foldings
- Different folds permit the RNAs to carry out specific functions in the cell

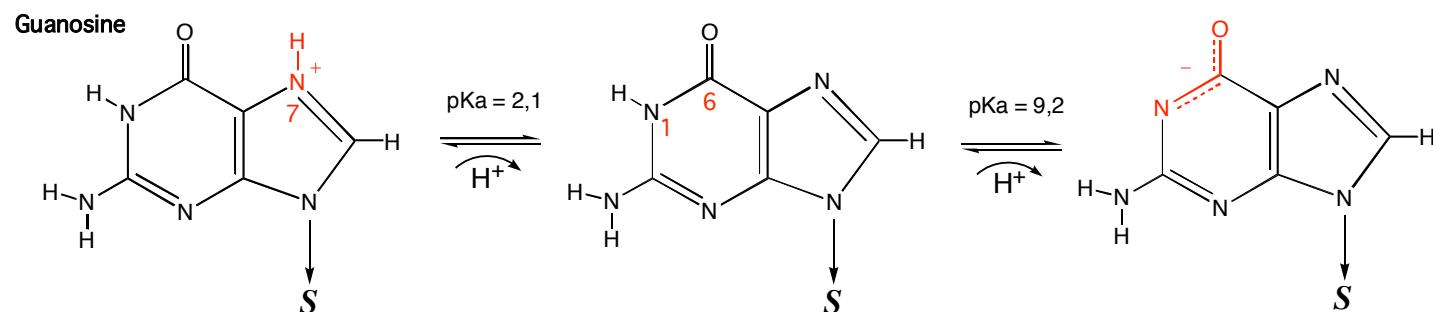
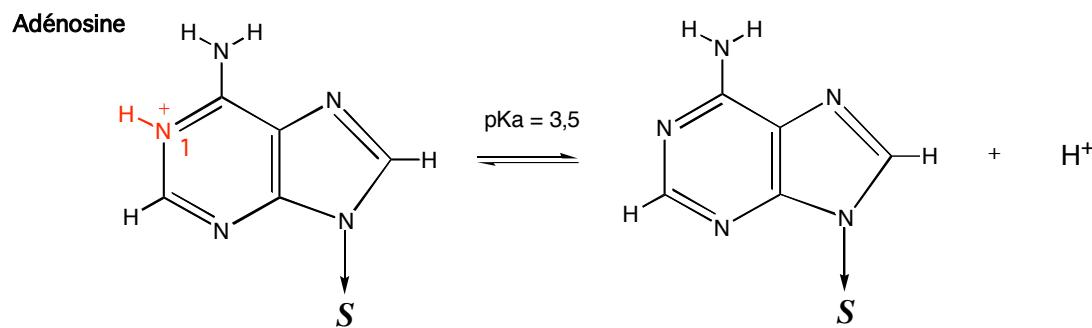
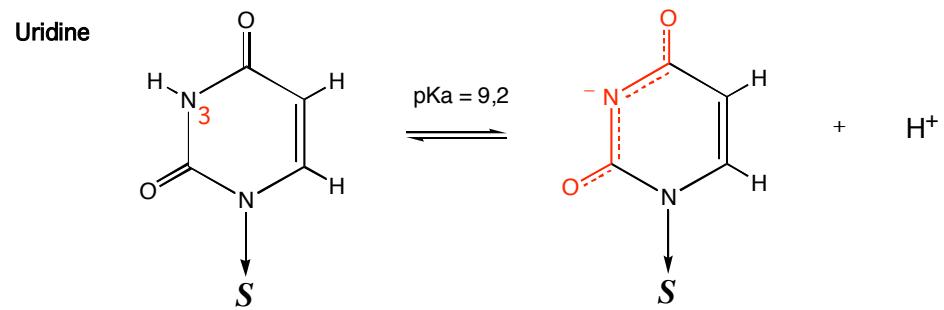
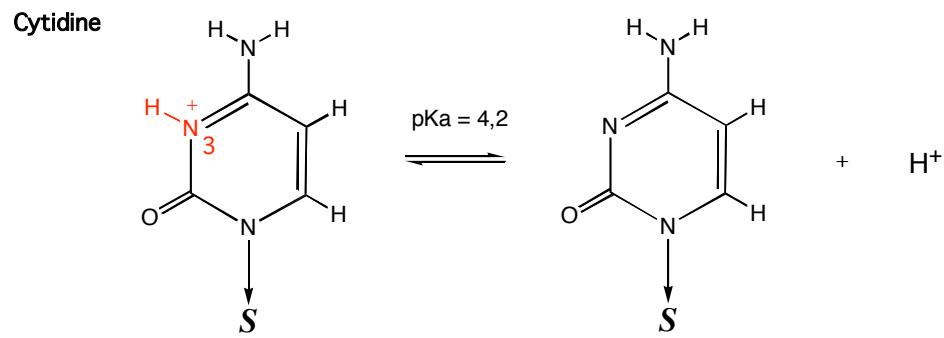


Charge delocalization

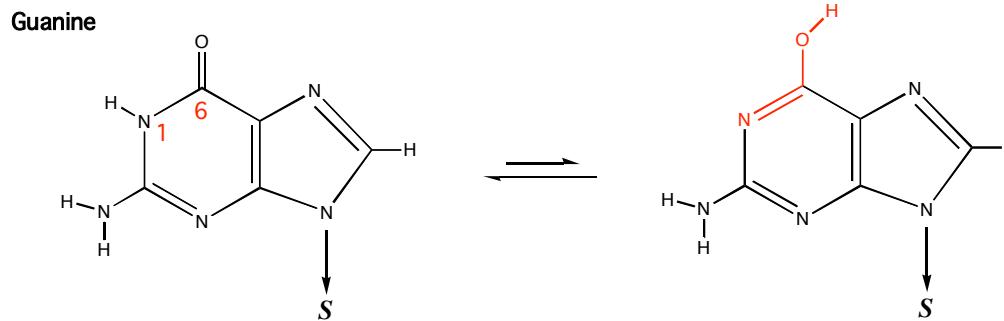
Tautomeric forms



Protonation possibilities

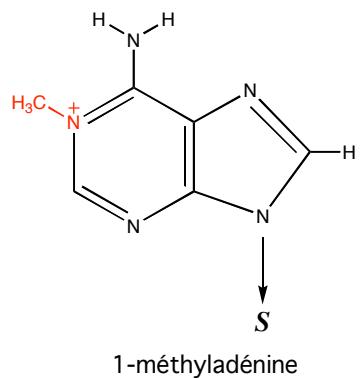


**Always
seen**

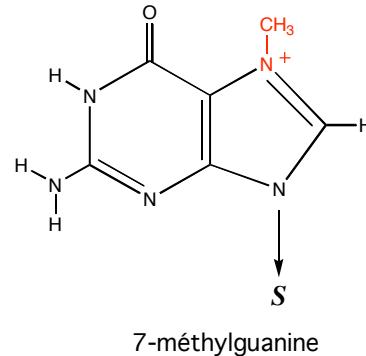


**Never
seen**

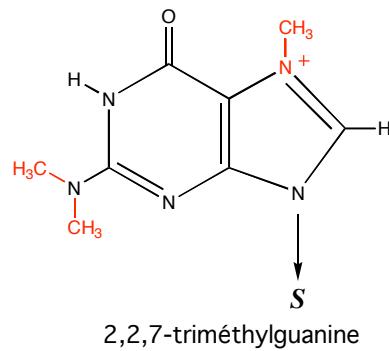
Modified bases have different



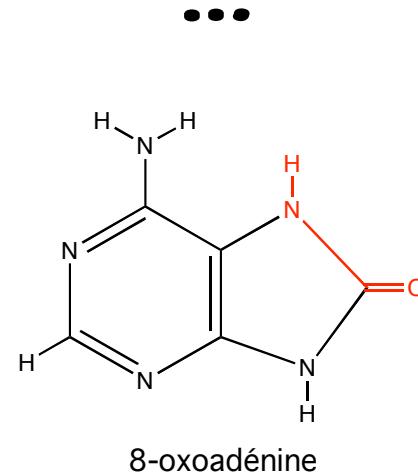
1-méthyladénine



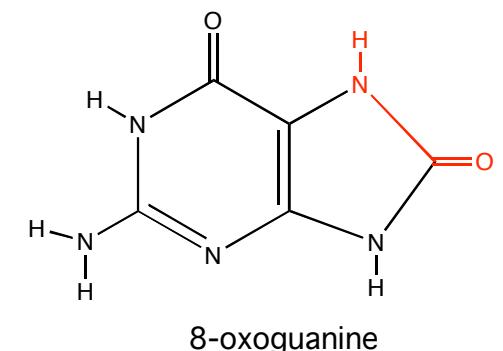
7-méthylguanine



2,2,7-triméthylguanine



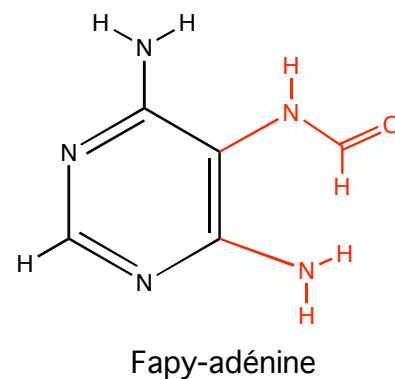
8-oxoadénine



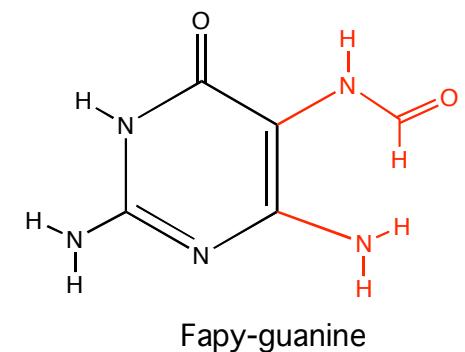
8-oxoguanine

...

electronic properties

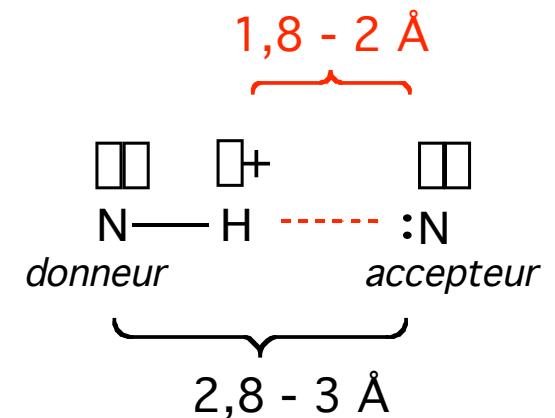
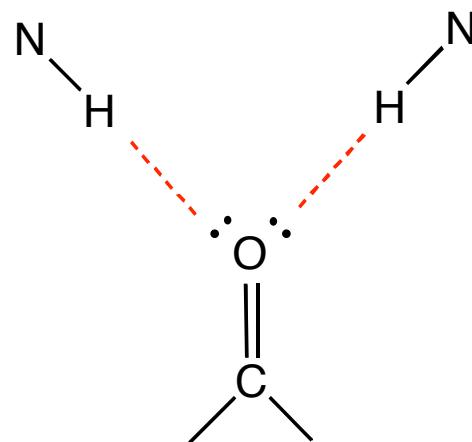


Fapy-adénine

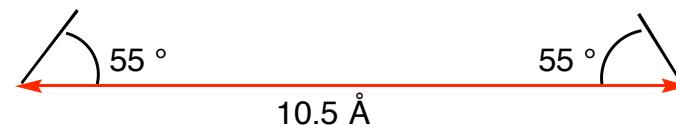
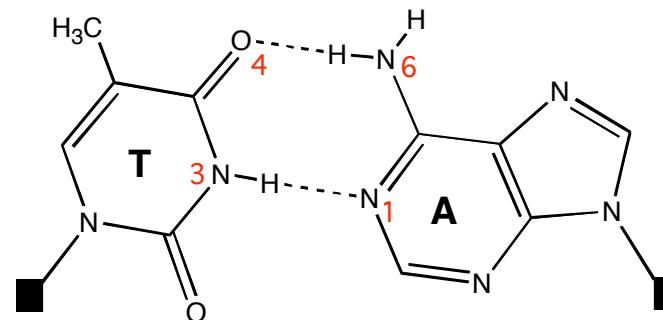
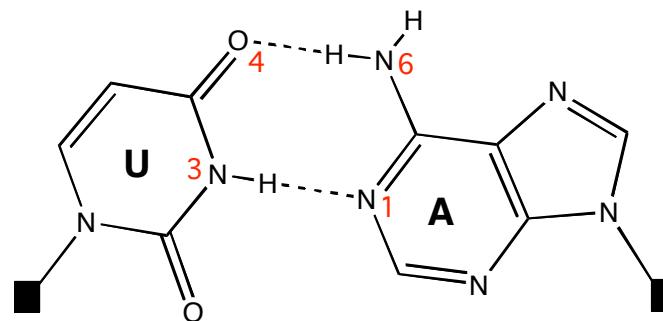
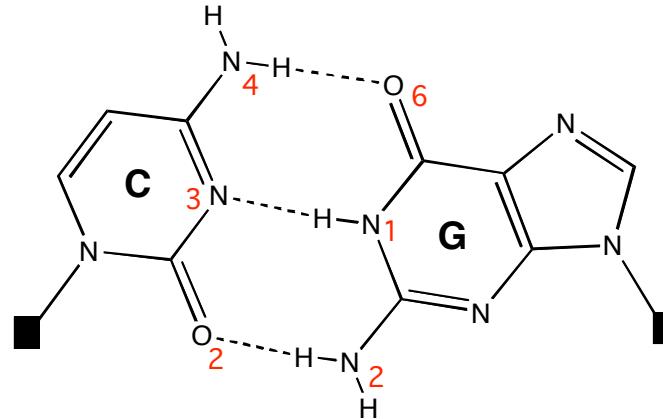


Fapy-guanine

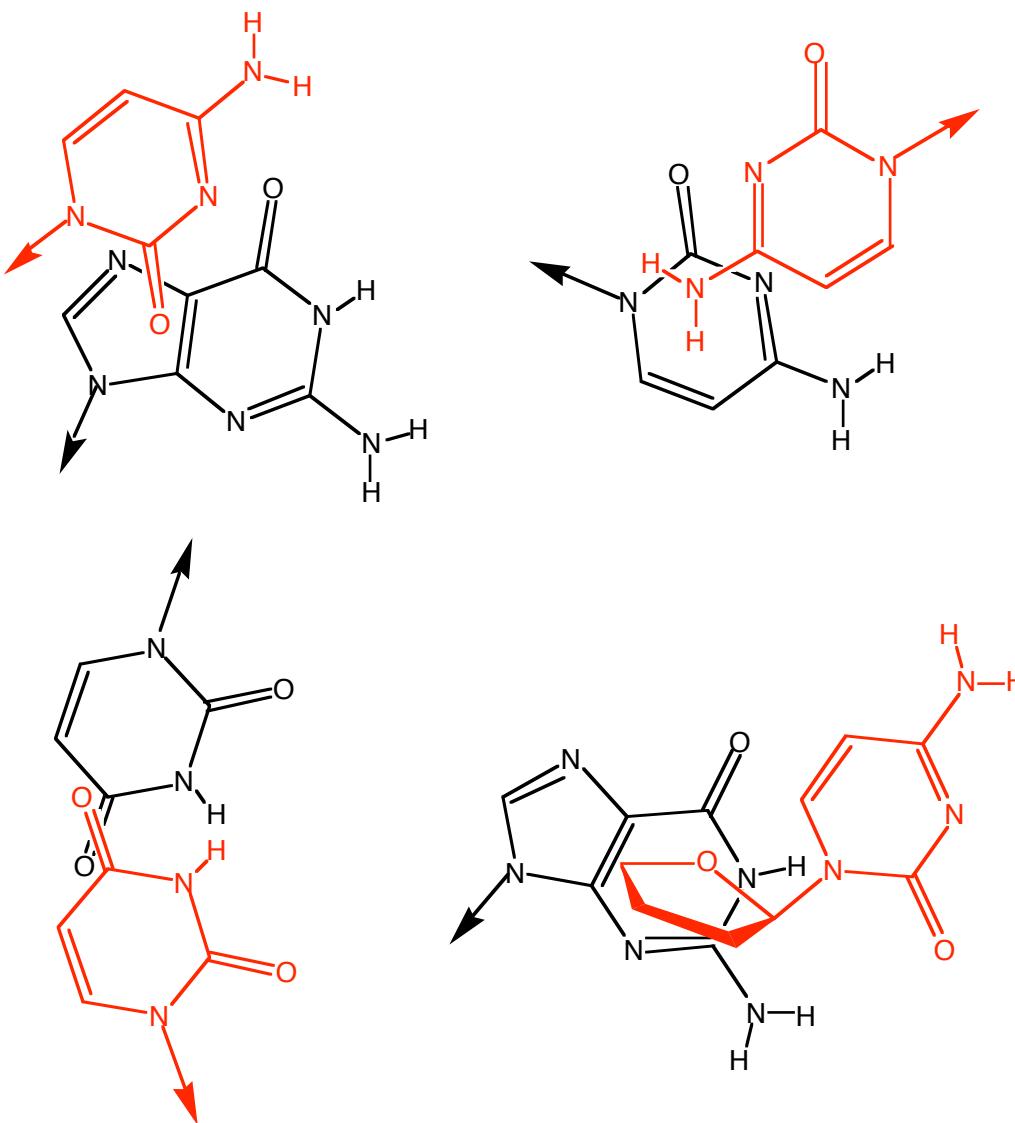
H-bond characteristics



Horizontal
Interactions
Base pairing.
In helices
Complementary
Watson-Crick



Vertical interactions : stacking



Stacking forces

- Driving Force : hydrophobic effect.
 - Not very specific
- Partition in
- very polar regions (phosphates) &
less polar ones (exocyclic groups of bases)

History of the many roles of RNA

Central dogma

The flow of genetic information

transcription

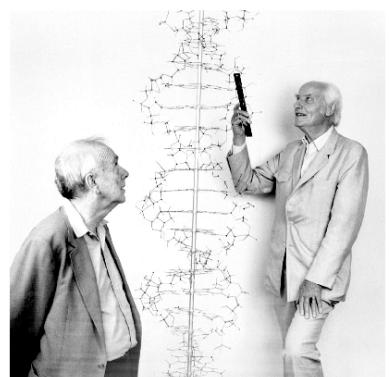
translation



DNA notoriety

NCBE | DNA50
Double helix 1953–2003

"We have discovered the secret of life"



I'm Watson, I'm Crick,
Let us show you our trick.
We've solved the riddle
seed of life sprang from.
We believe we're a stew
Of molecular goo
With a period of thirty-four Angstroms.

E.S. Anderson, et al. 1953

celebrating life
DNA: 50 years 1953-2003

- introduction
- DNA ephemera
- DNA models
- Read all about it
- Timeline
- DNA Easter eggs
- DNA dance
- DNA drama
- The DNA cocktail
- Practical protocols
- Links to other sites

Rosalind Franklin: Dark Lady of DNA

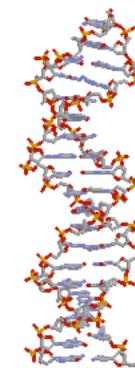
Book Sheds New Light on a Scientific Landmark



Rosalind Franklin
Photo: courtesy
HarperCollins

Oct. 6, 2002 -- Early next year, scientific institutions in the United States and Great Britain will mark the 50th anniversary of one of the greatest discoveries in science. In April of 1953, James Watson, Francis Crick and Maurice Wilkins identified the substance of life -- the structure of DNA.

They later shared a Nobel Prize. Their discovery depended heavily on the work of a woman, chemist Rosalind Franklin, whose research was used without her knowledge or permission. Watson's memoir of the discovery dismisses Franklin as frumpy, hostile and unimaginative. A later work by a friend casts Franklin as a feminist icon, cheated of recognition.

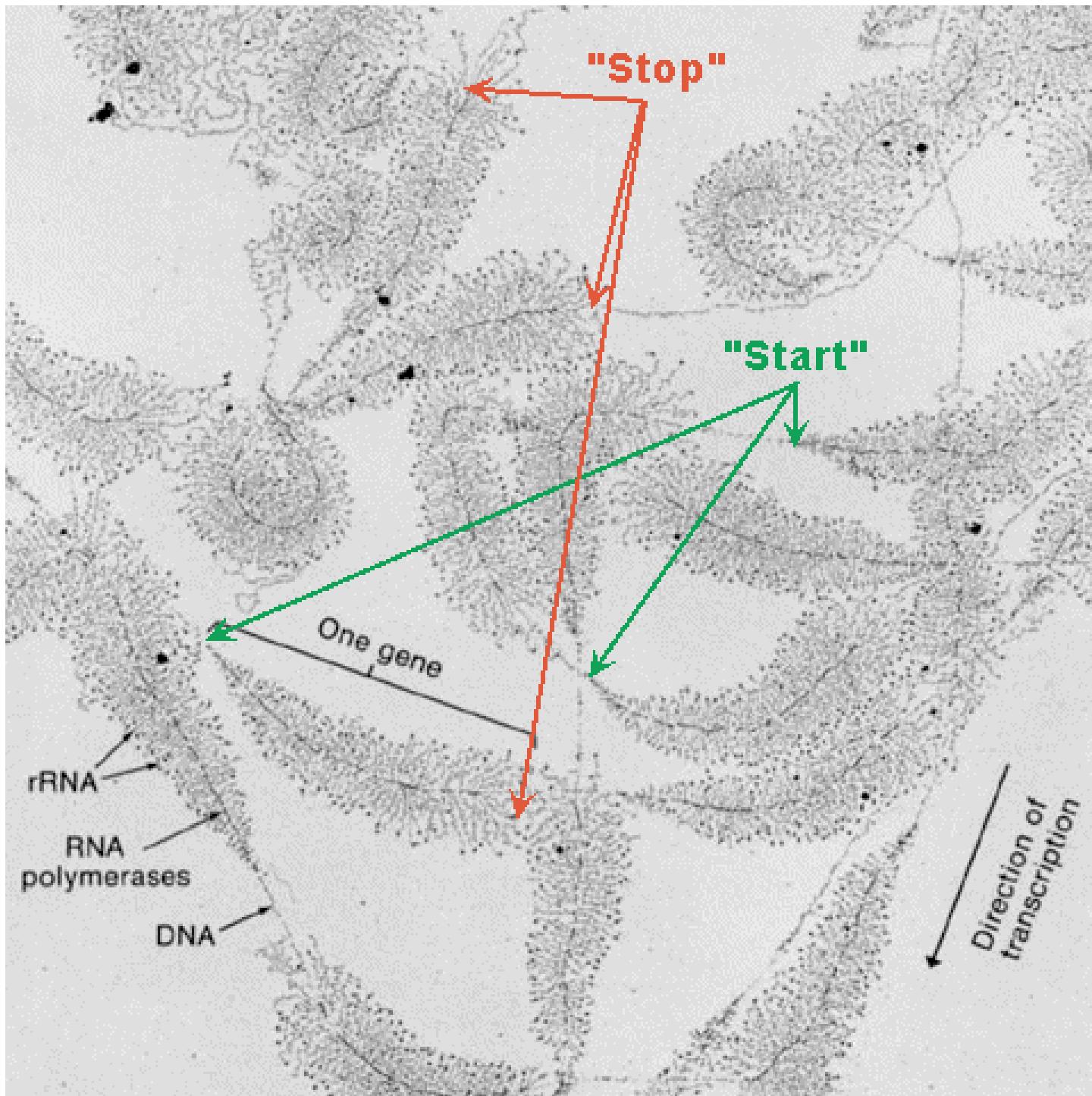


Short history of RNA

- Late 1800's - 2nd kind of nucleic acid not in the nucleus (rRNA)
- 1920's - sugar for DNA vs RNA
- 1958 - tRNA (Hoagland)
- 1960's - mRNA

Relative Amount of
RNA in E. coli

rRNA	80%
tRNA	15%
mRNA	5%



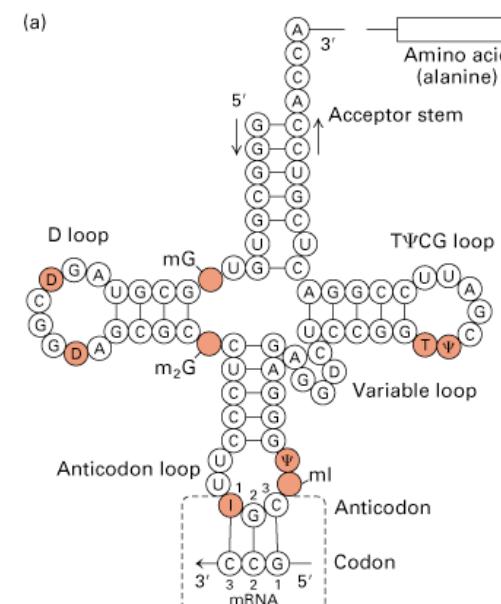
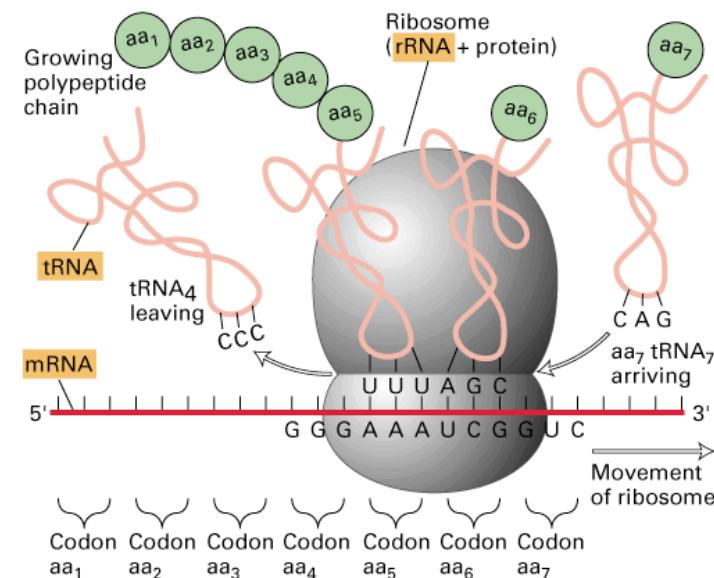
RNA

- “Three” different types of RNA

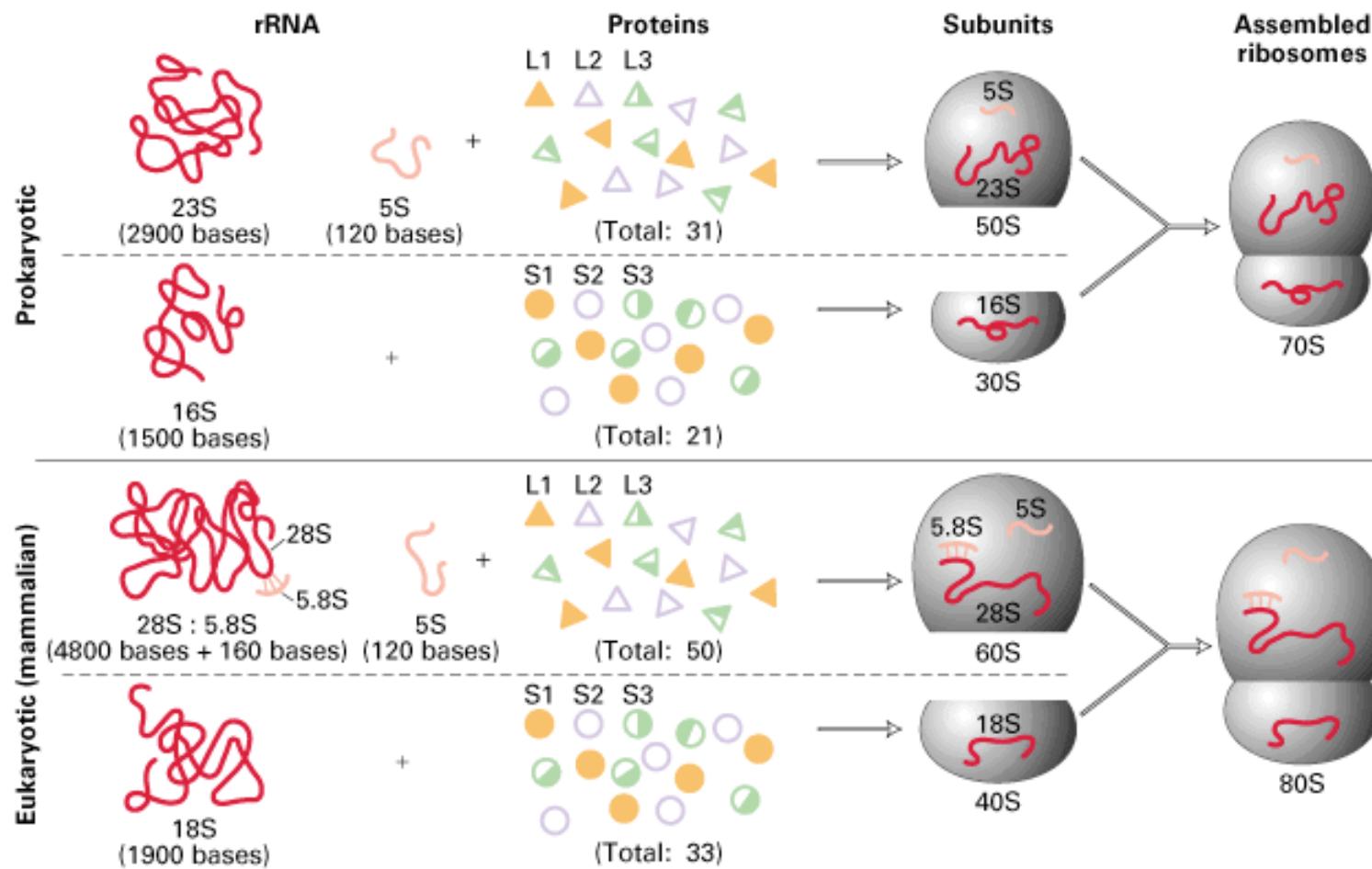
- mRNA - messenger RNA, specifies order of amino acids during protein synthesis

- tRNA - transfer RNA, during translation mRNA information is interpreted by tRNA

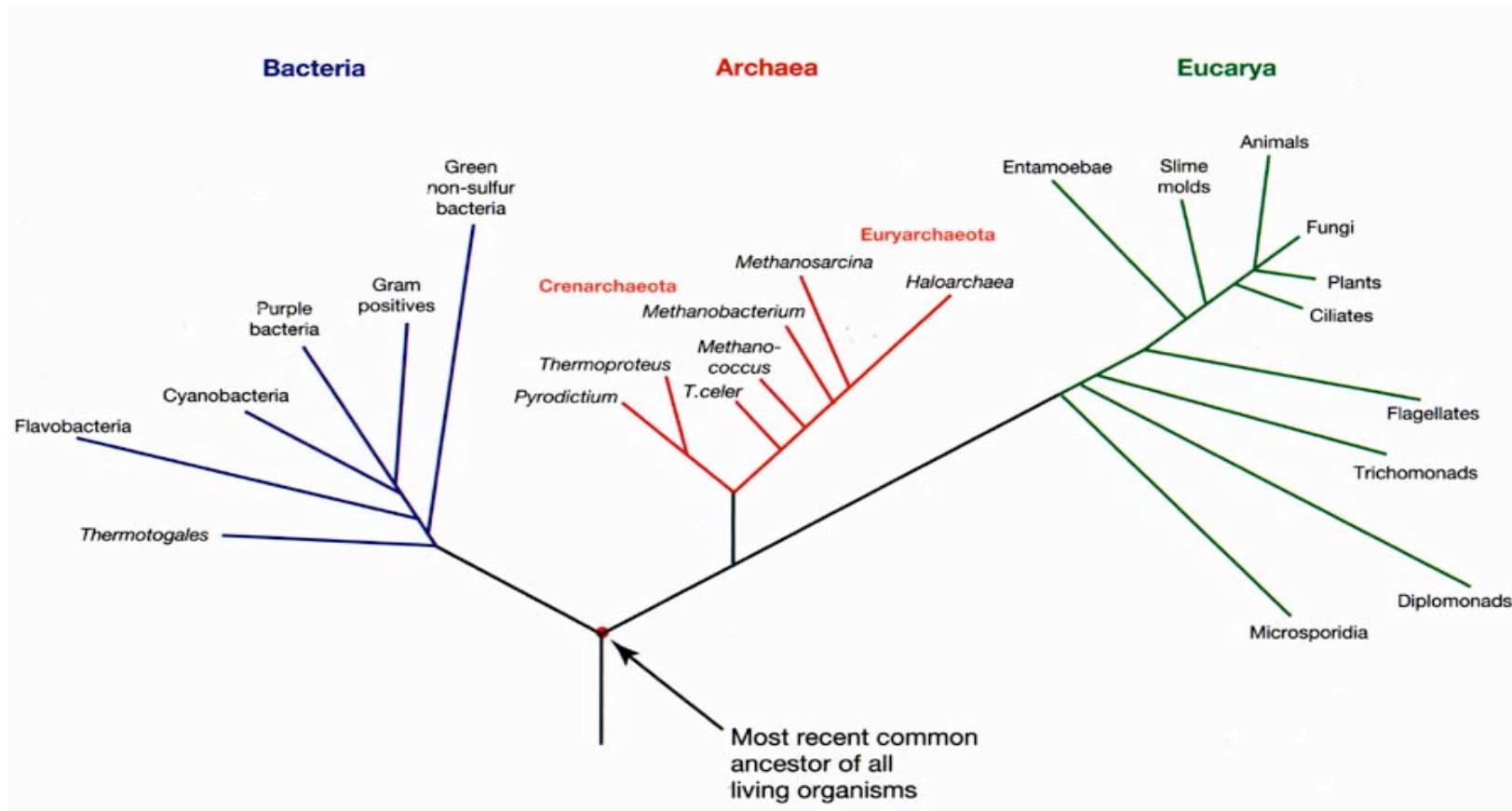
- rRNA – ribosomal RNA, combined with proteins aids tRNA in translation



Ribosomal RNA (rRNA)

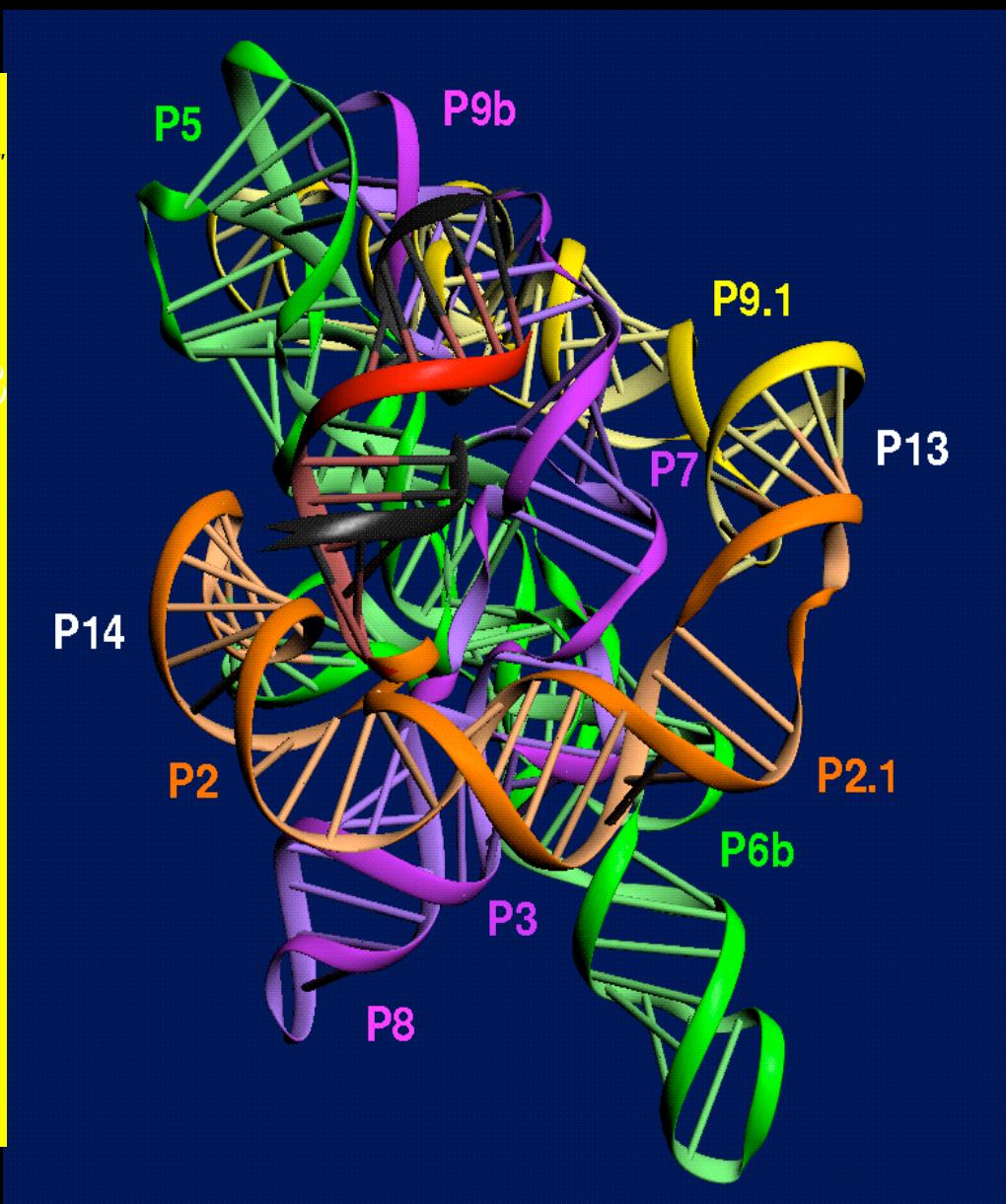
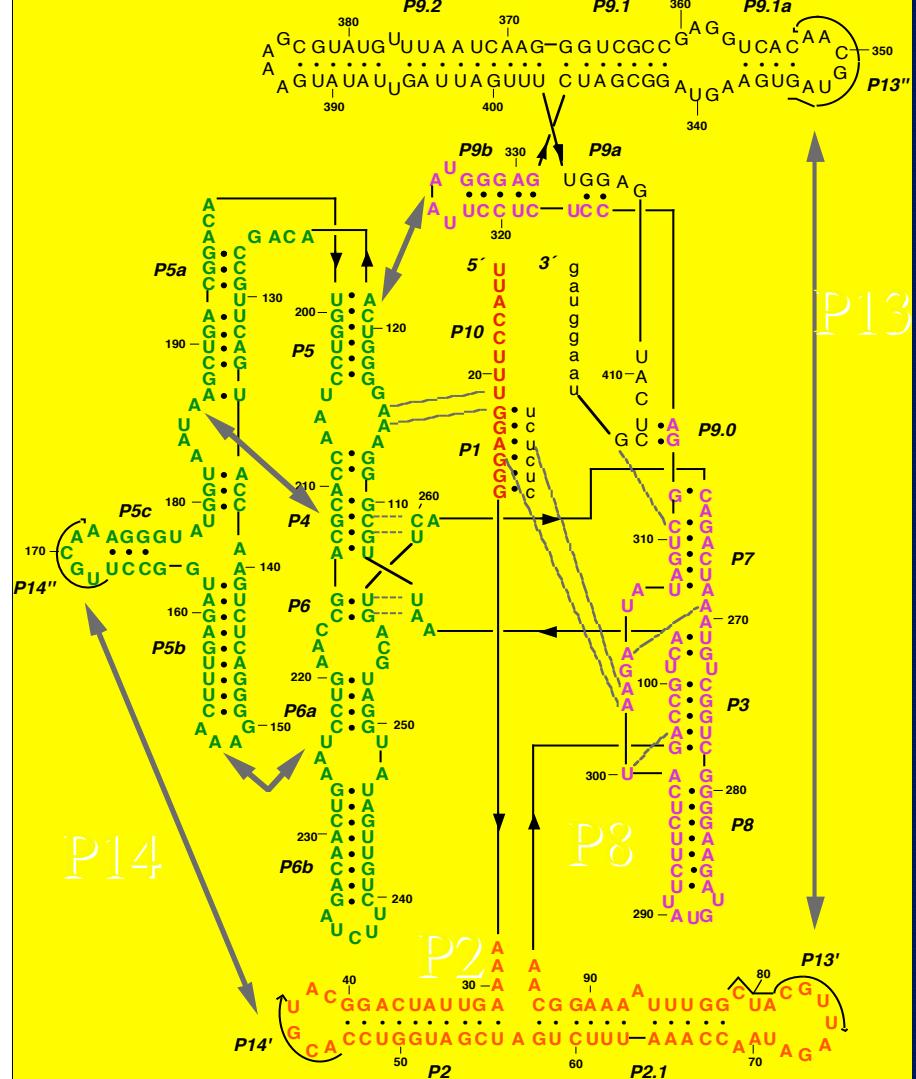


Phylogeny of Life using SSU rRNA



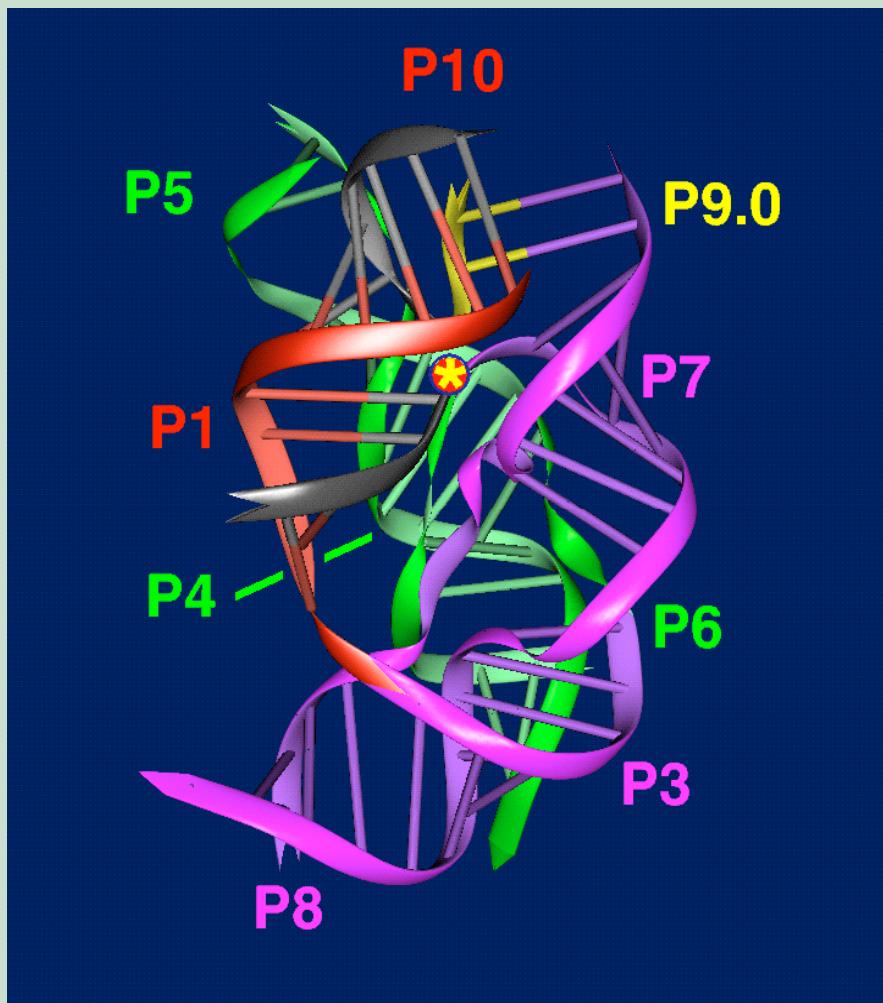
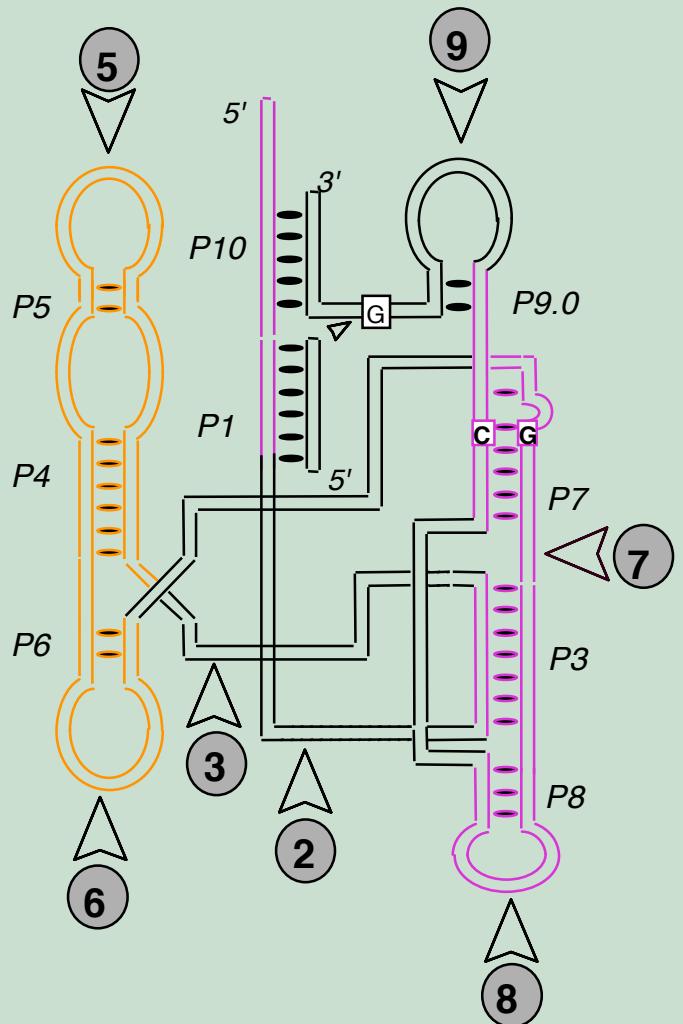
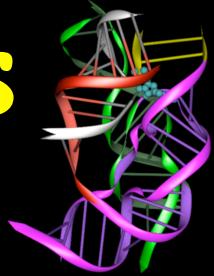
Discovery of catalytic RNA

- Early 1980's :
- Tom Cech & Sidney Altman
- Self-splicing group I introns and
- Ribonuclease P

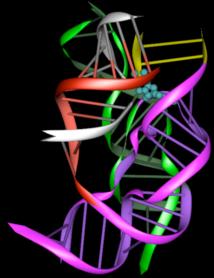


Group I self-splicing introns

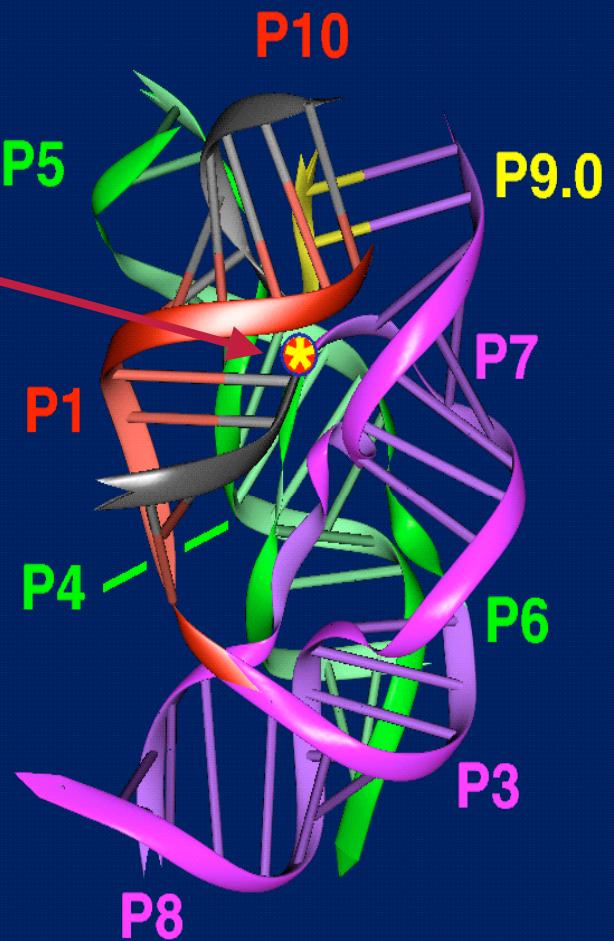
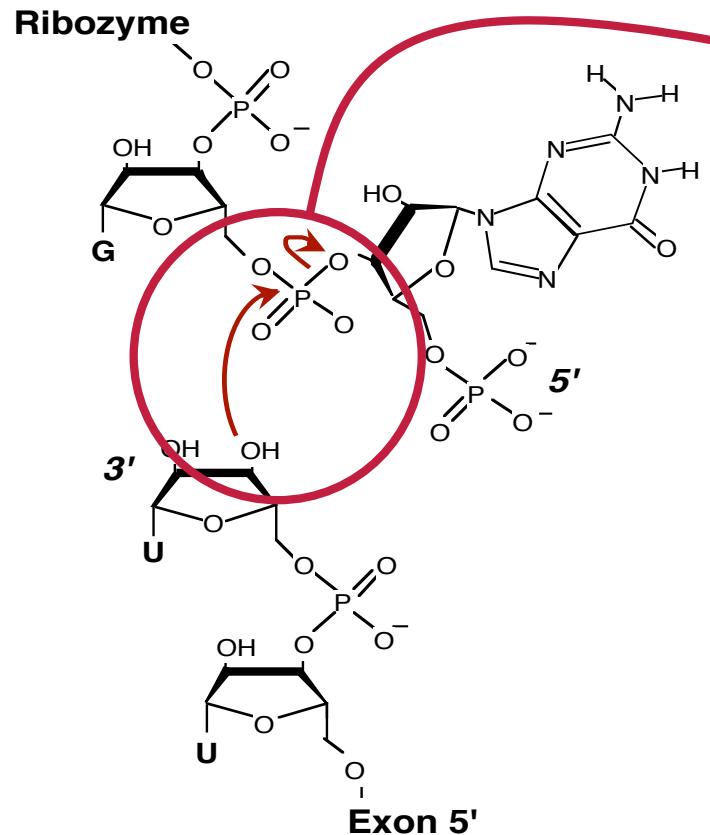
Core structure



Catalytic Core of Group I introns



Transesterification



Second discovery of RNA

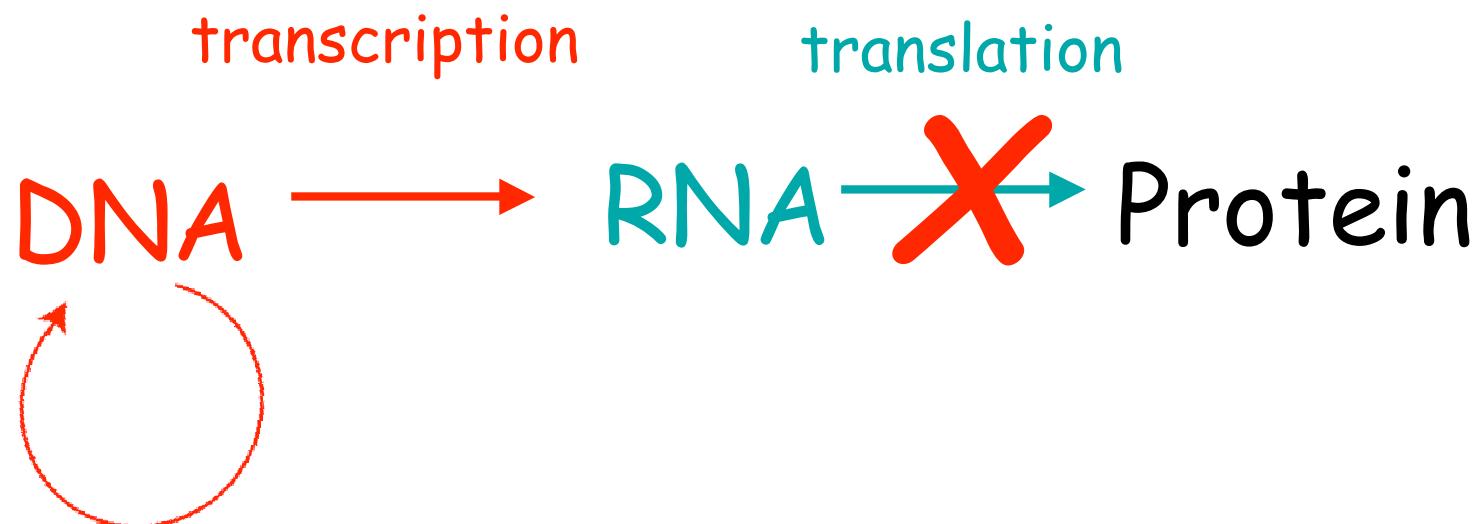
- **ncRNAs** - functional RNA molecules rather than proteins; RNA other than mRNA (ex.XIST)
- **RNAi** - RNA interference
- **siRNA**- active molecules in RNA interference; degrades mRNA (act where they originate)
- **miRNAs** - tiny 21–24-nucleotide RNAs; probably acting as translational regulators of protein-coding mRNAs (regulate elsewhere)

Other Roles of RNA

- **stRNA** - Small temporal RNA; (ex. lin-4 and let-7 in *Caenorhabditis elegans* : development
- **snRNA** - Small nuclear RNA; includes spliceosomal RNAs
- **snoRNA** - Small nucleolar RNA; most known snoRNAs are involved in rRNA modification
- **RNA world** - RNA as catalyst

Central dogma

The flow of genetic information



ncRNA genes

- Genomic dark matter
 - Ignored by gene prediction methods
 - Not in EnsEMBL
 - Computational complexity
- ~10% of human gene count?

RNA interference

Short interfering RNAs (siRNAs)

A class of double-stranded RNAs of 21–22 nucleotides in length, generated from dsRNAs. siRNAs silence genes by promoting the cleavage of mRNAs with exactly complementary sequences, or recruiting inhibitory proteins to, or directing the modification of, DNAs with exactly complementary sequences.

MicroRNAs (miRNAs)

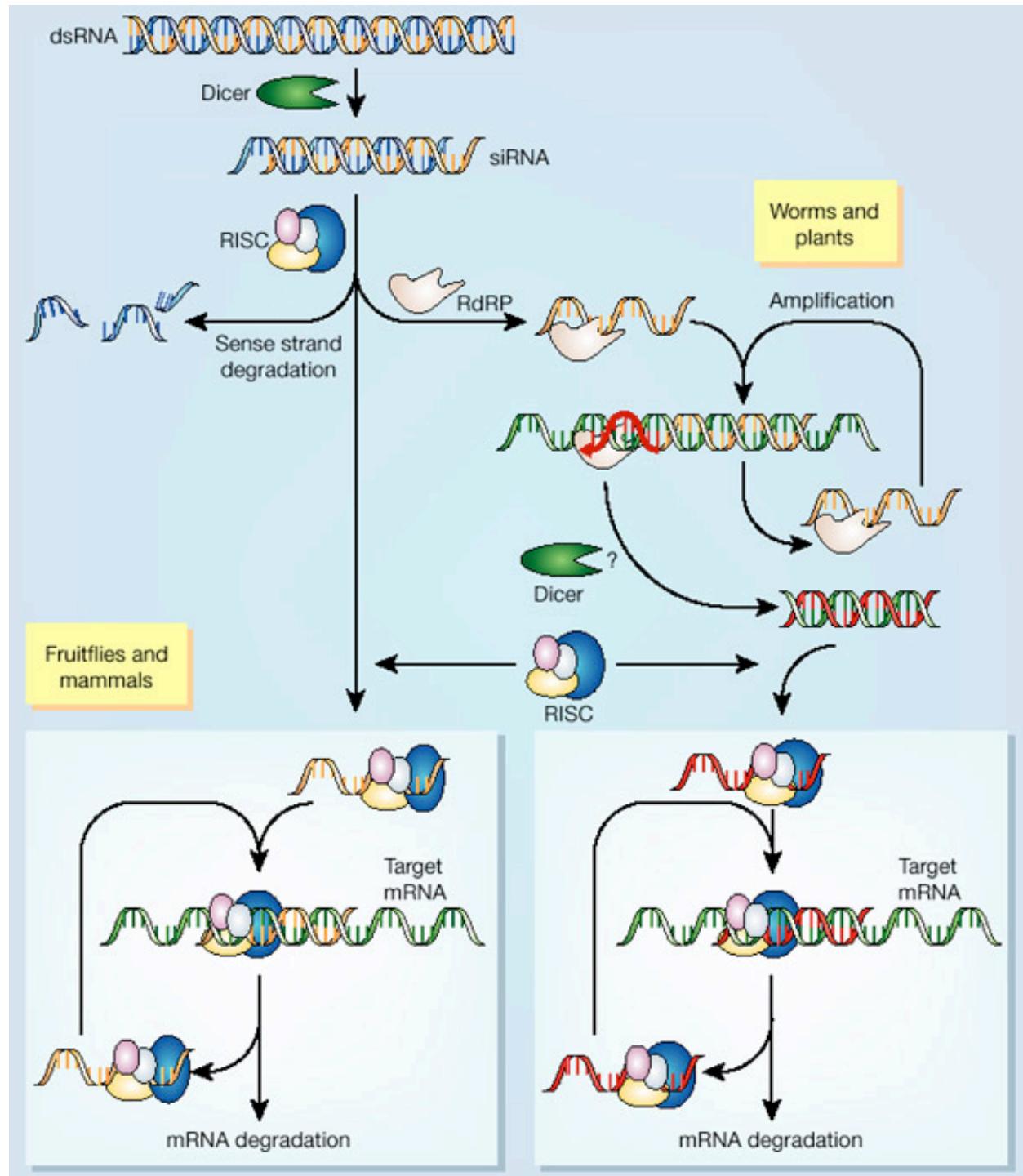
A class of 19–25-nucleotide, single-stranded RNAs that are encoded in the genomes of most multicellular organisms studied. Some are evolutionarily conserved and are developmentally regulated. They silence certain cellular genes at the stage of protein synthesis.

Tiny non-coding RNAs (tncRNAs)

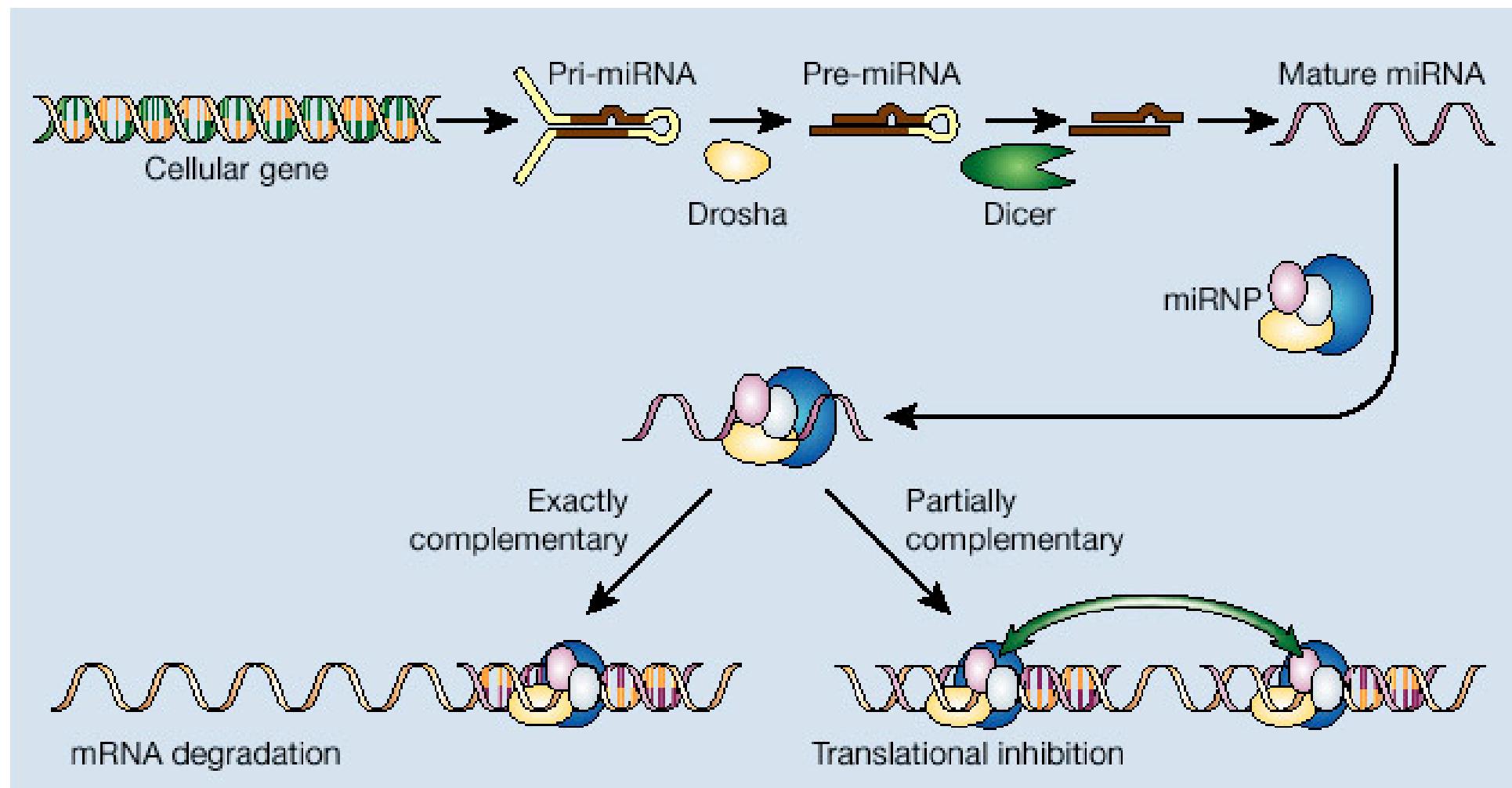
A newly discovered class of short, 20–22-nucleotide RNAs that are encoded in the genome of *C. elegans*. They are not evolutionarily conserved, but some are developmentally regulated. Their function is still unknown.

Small modulatory RNA (smRNA)

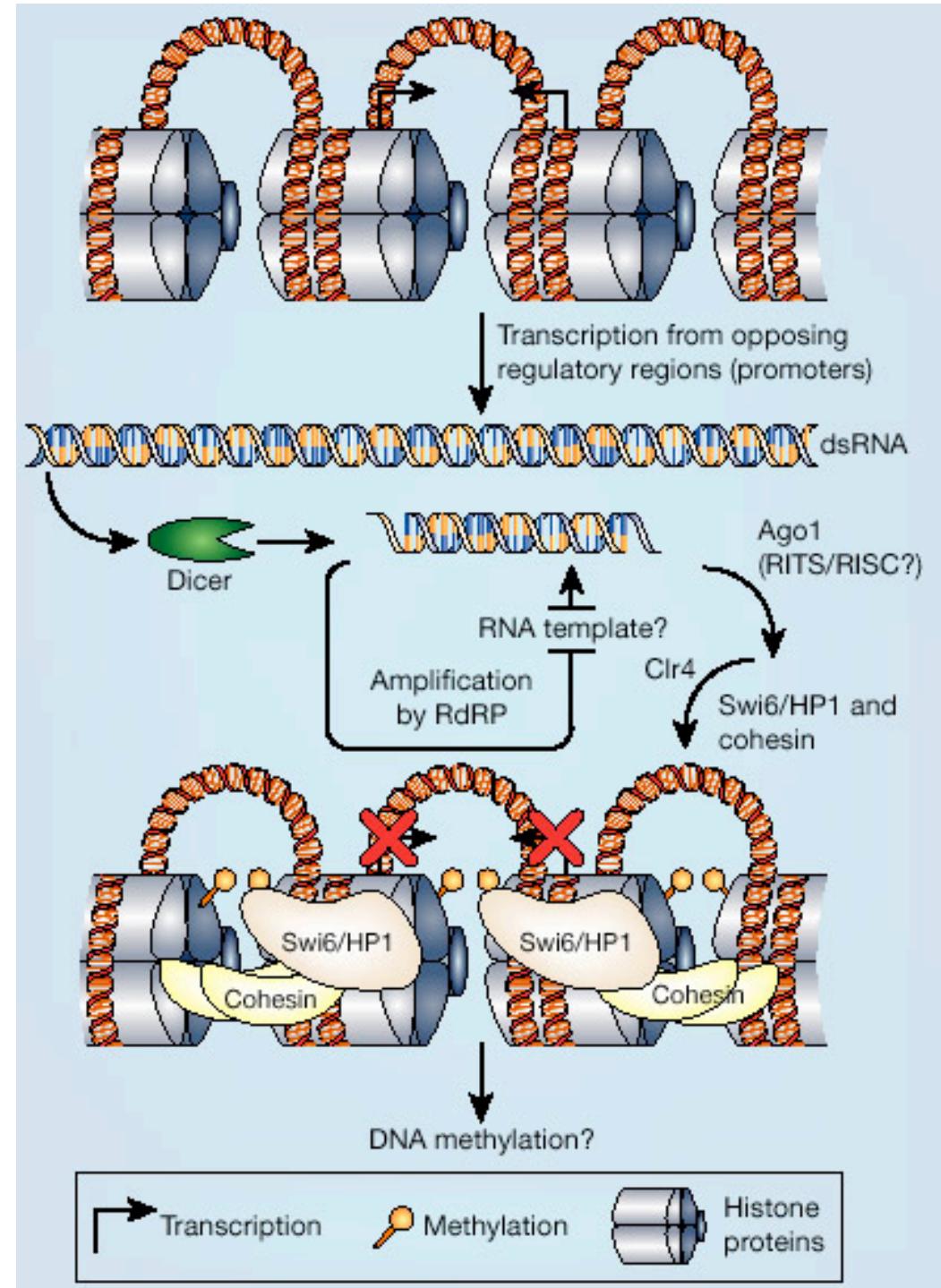
A short, dsRNA, identified earlier this year in mice, that allows the expression of neuron-specific genes only in adult neurons.



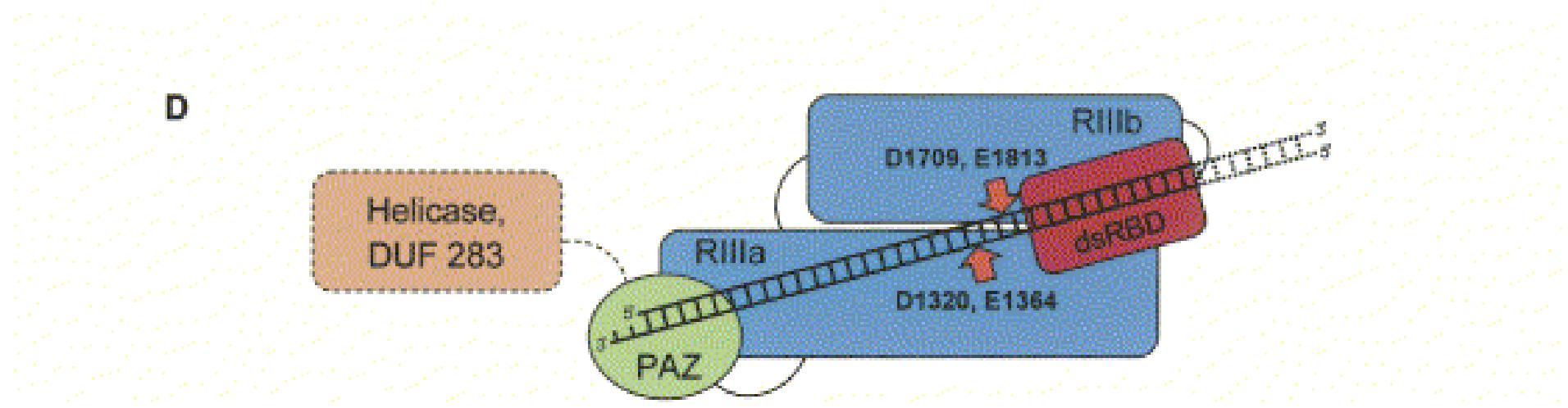
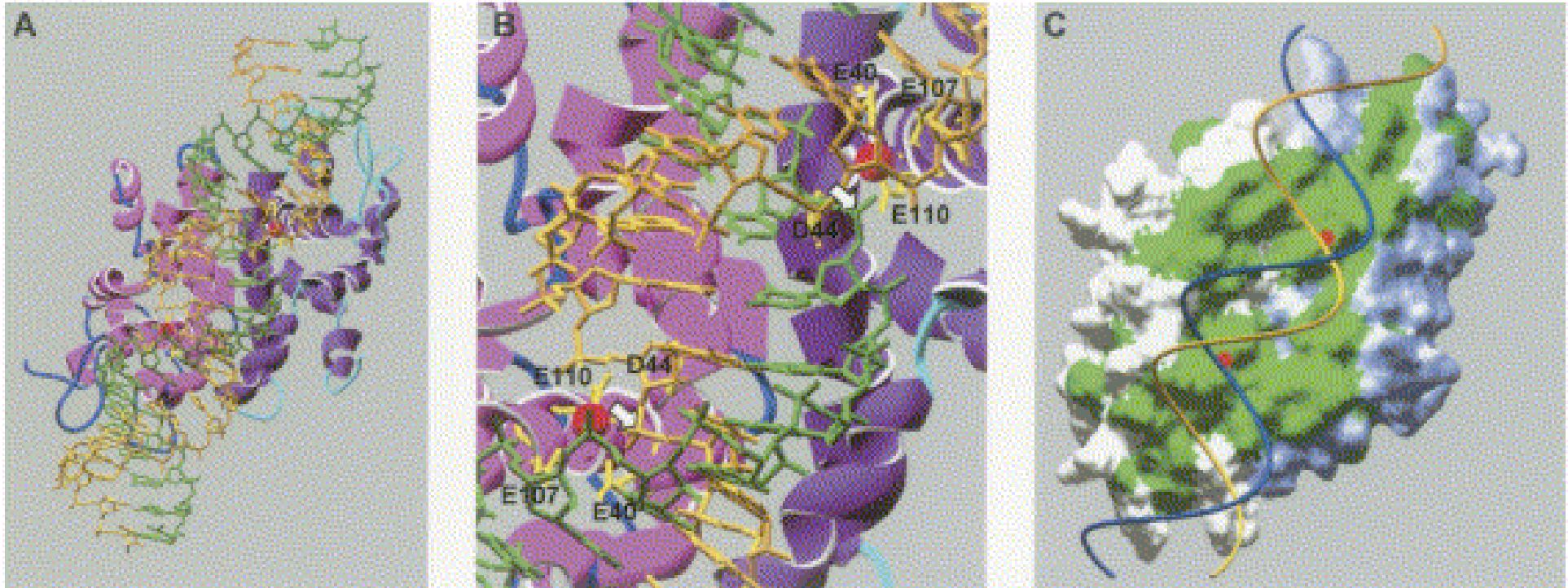
Intricate maturation pathways



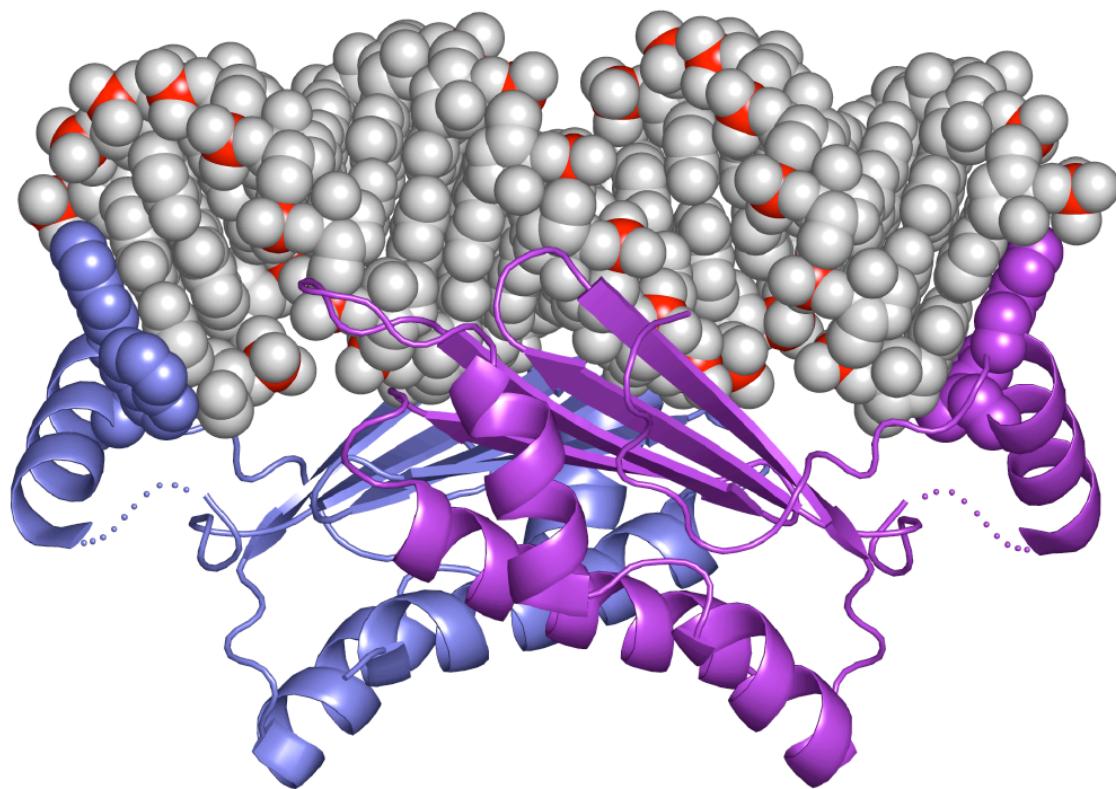
Implicated in chromatine remodelling



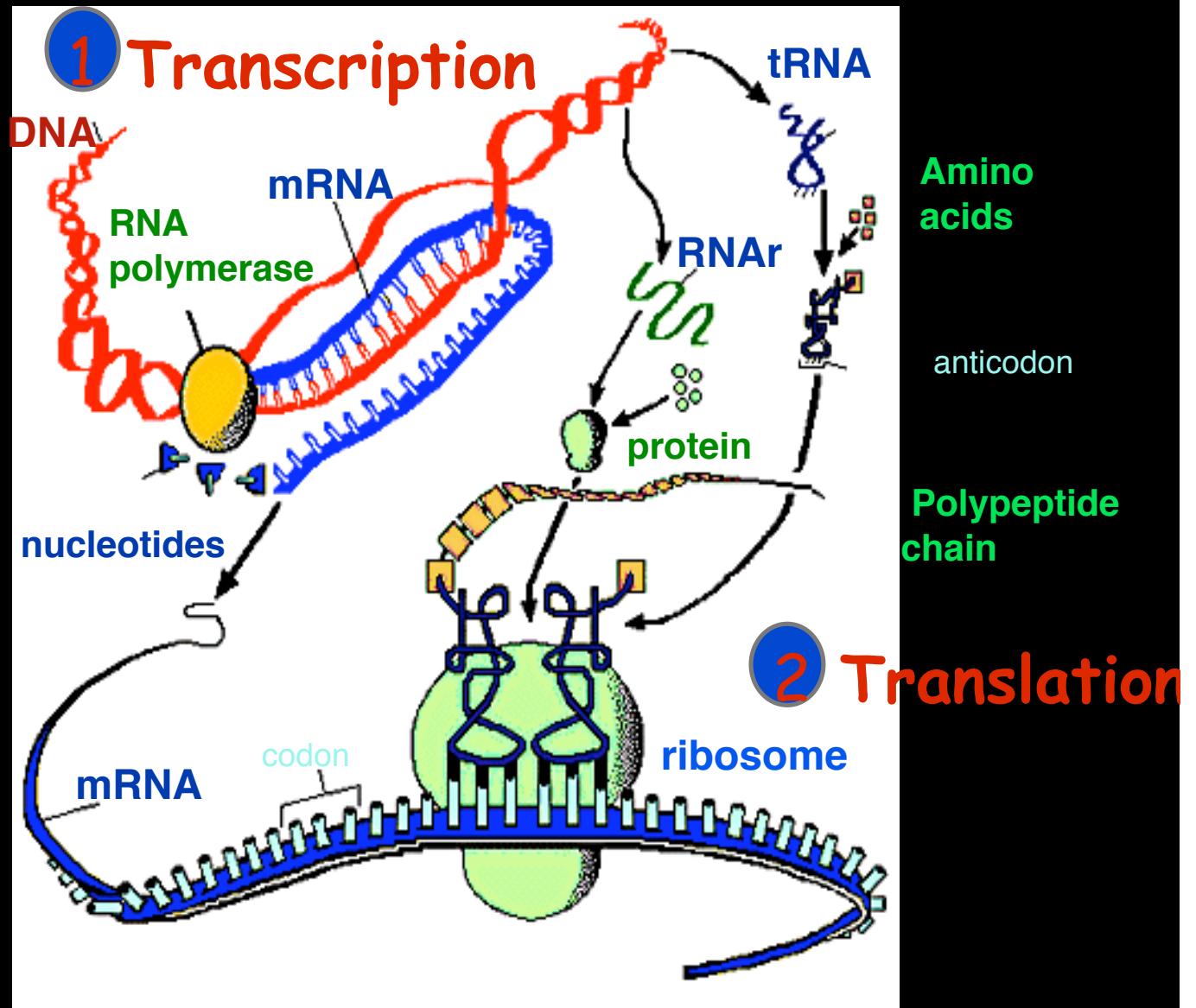
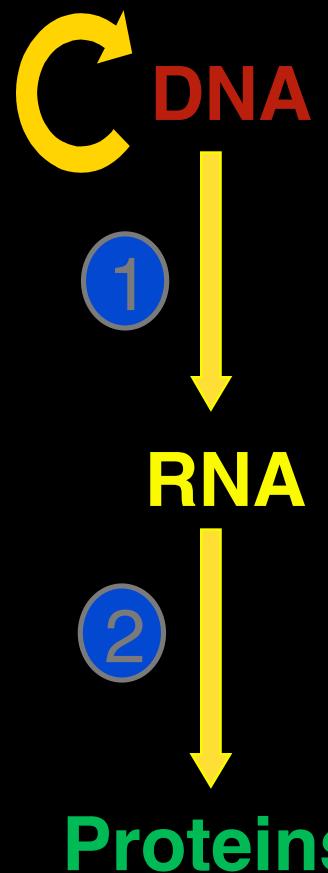
Cleavage part of DICER



Measuring device for 21 nt



ONE Role of RNA in cells



Versatility of RNA functions

- Not only messenger RNAs (alternative splicing)
- Number & variety of non-coding RNAs : ribosomal RNAs, snRNAs, snoRNAs, and numerous regulator RNAs..
- Cofactor RNAs : telomerase,
- to be discovered

Properties of RNA molecules

Assemble in double-starnded helices like DNA

→ Carry GENETIC INFORMATION like DNA

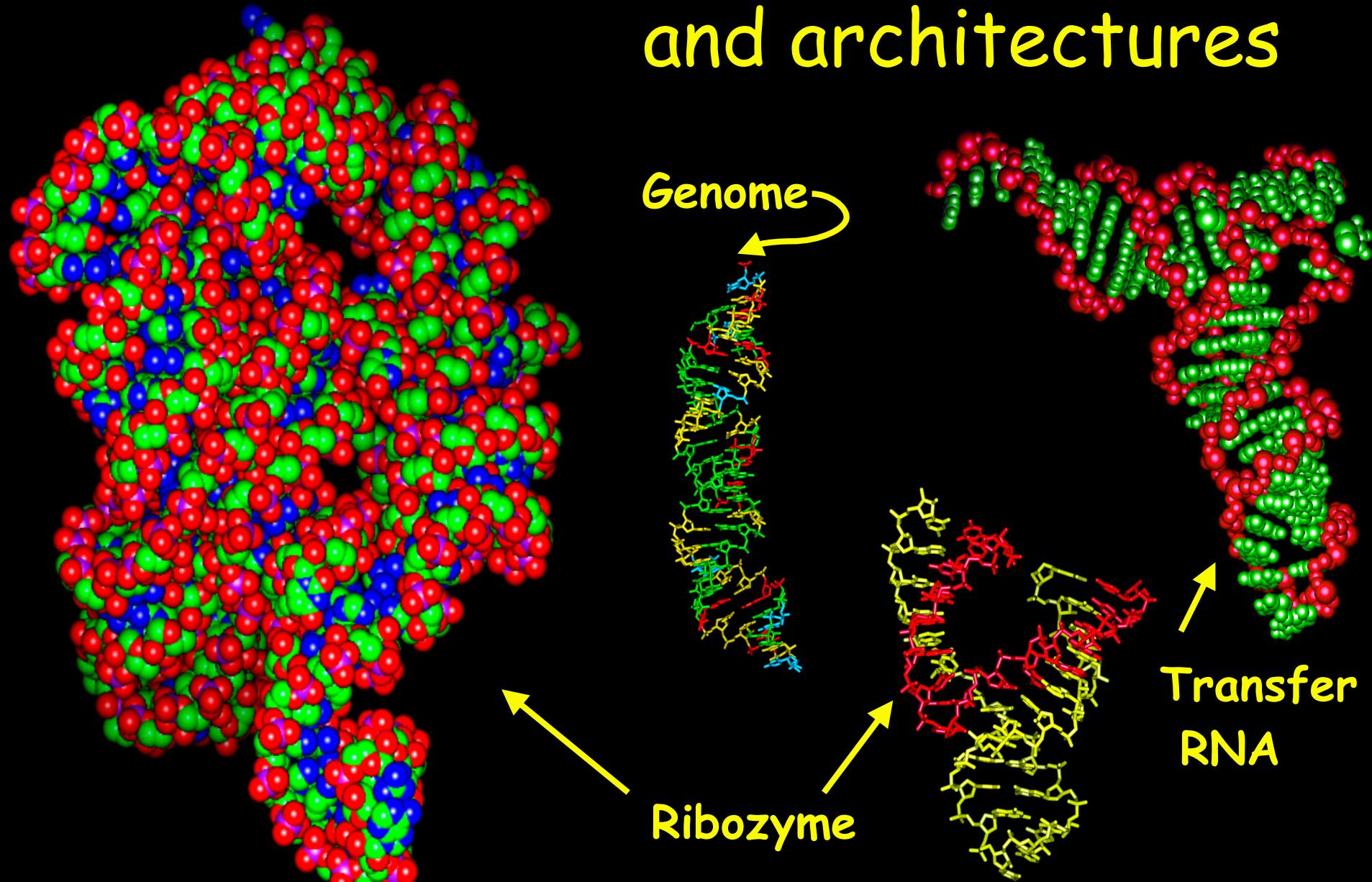
Fold in complex tertiary architectures like proteins

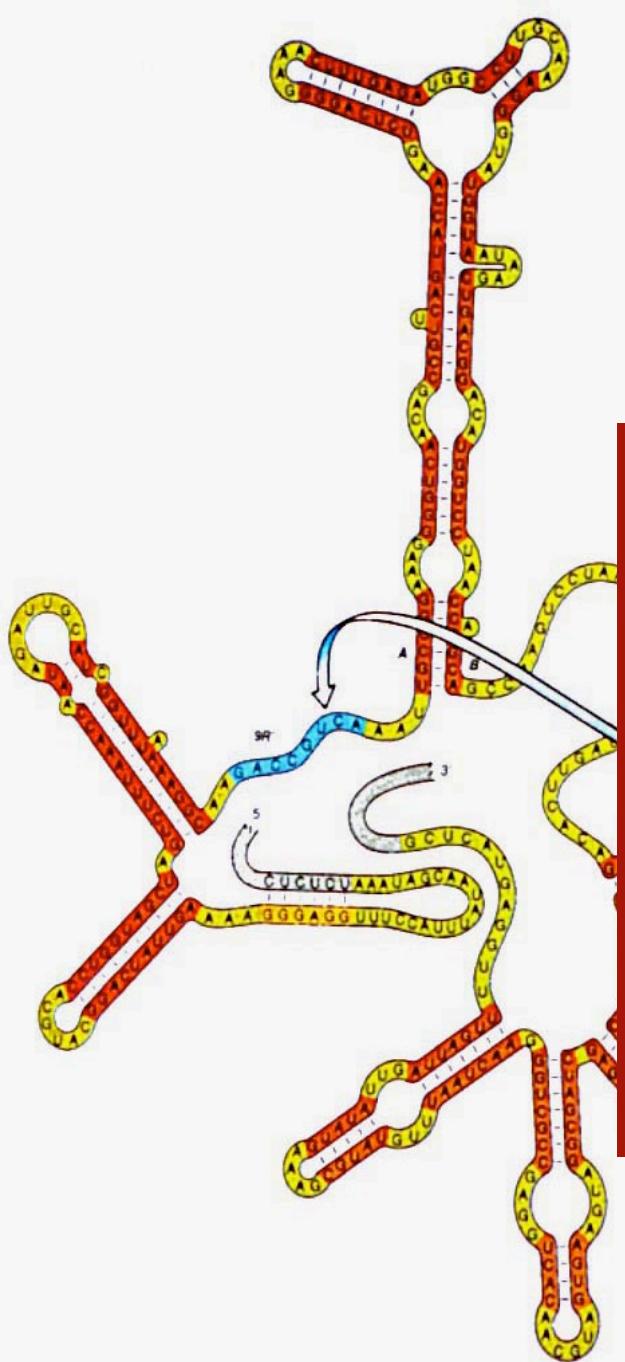
→ Perform CHEMICAL CATALYSIS like proteins

Catalytic RNAs

- Ribonuclease P
- Self-splicing introns
- Hepatitis delta virus
- The ribosome and peptide bond formation
- The spliceosome (not fully proven yet)
- and ...

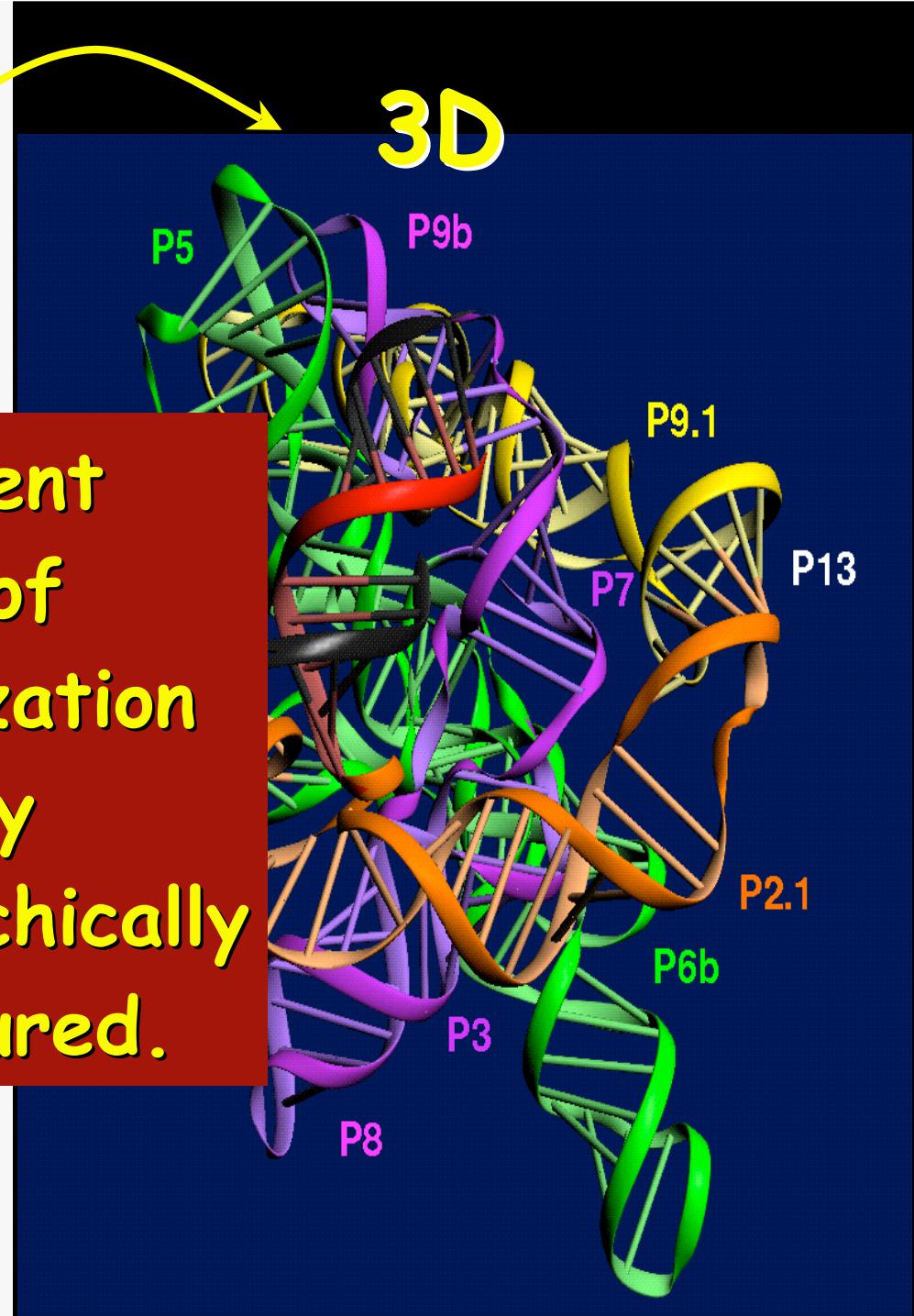
Great diversity of RNA functions and architectures



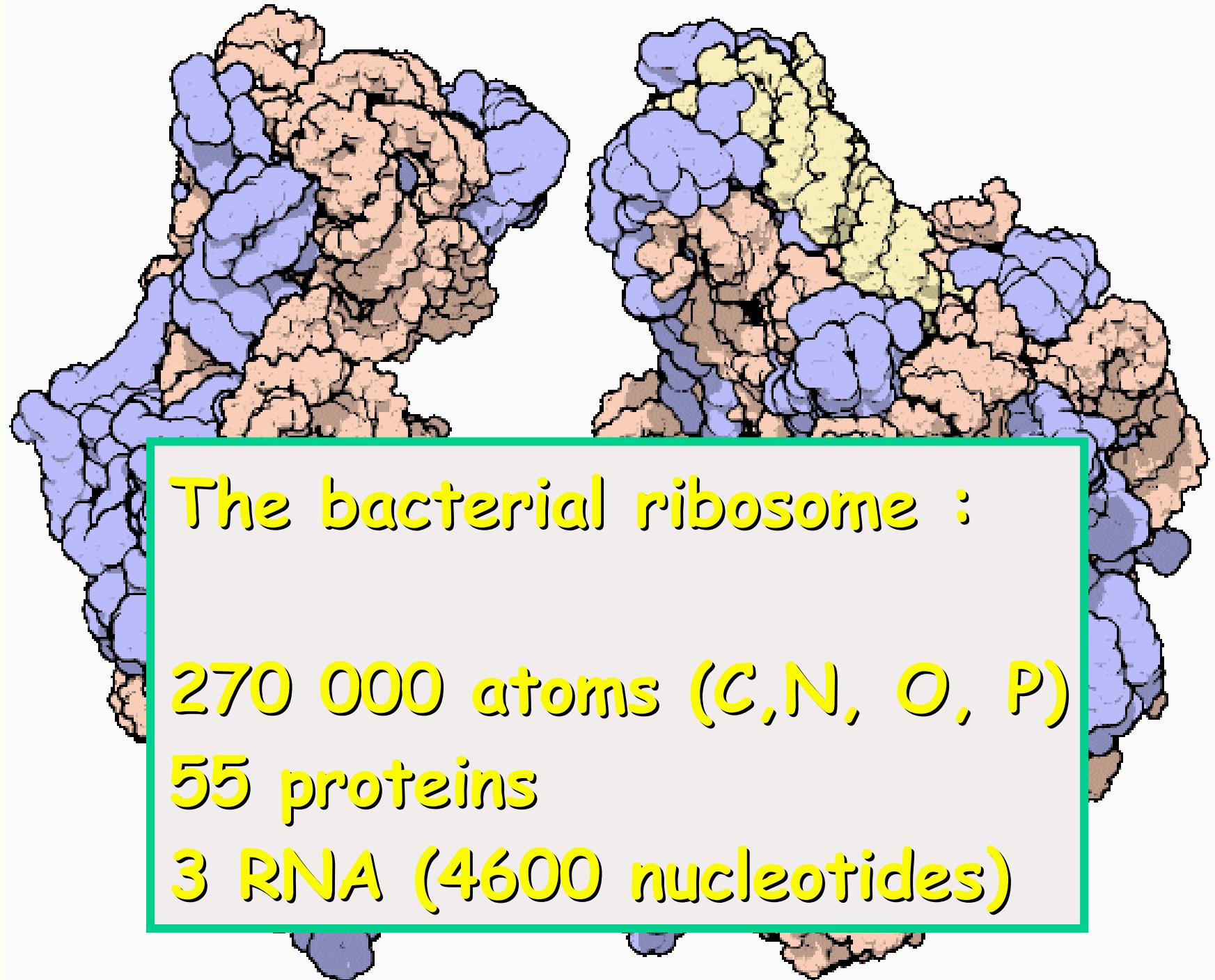


2D

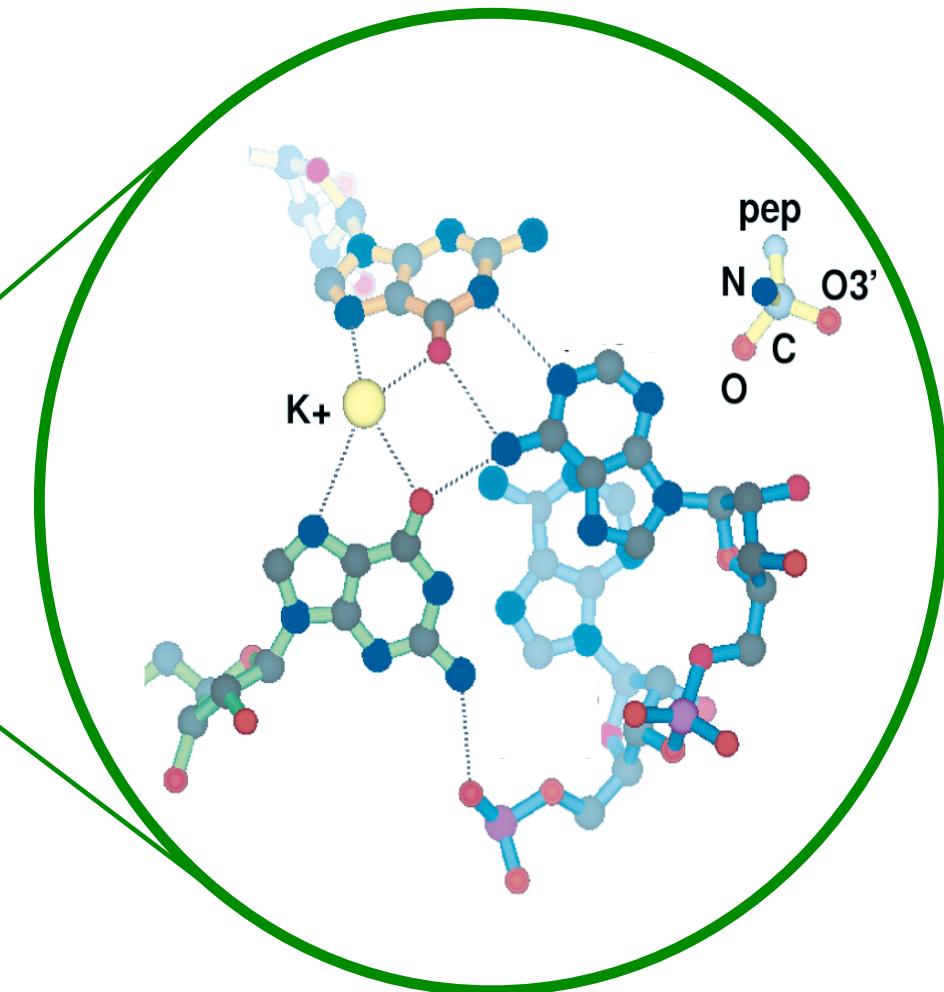
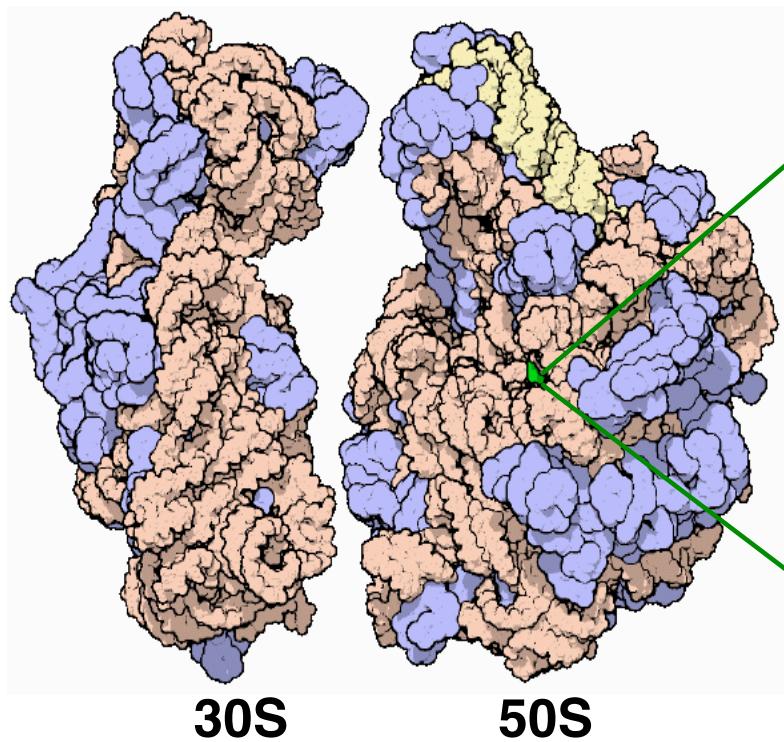
Different
levels of
organization
possibly
hierarchically
structured.



3D



Ribosome active site

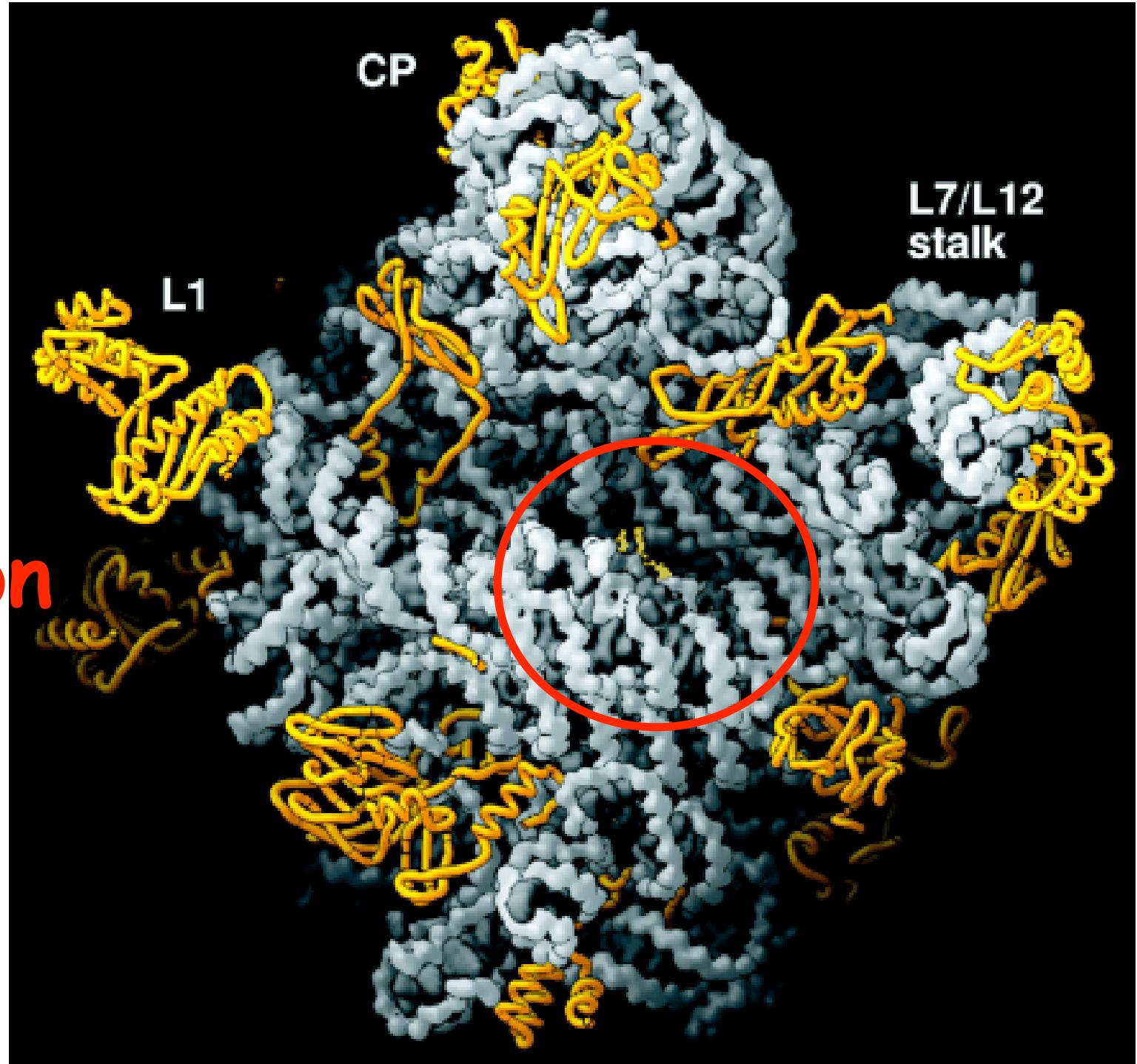


Nissen et al. (2000) Science, 289

H. F. Noller, T. A. Steitz, P. B. Moore, A. Yonath, V.
Ramakrishnan

Luc Jaeger ©

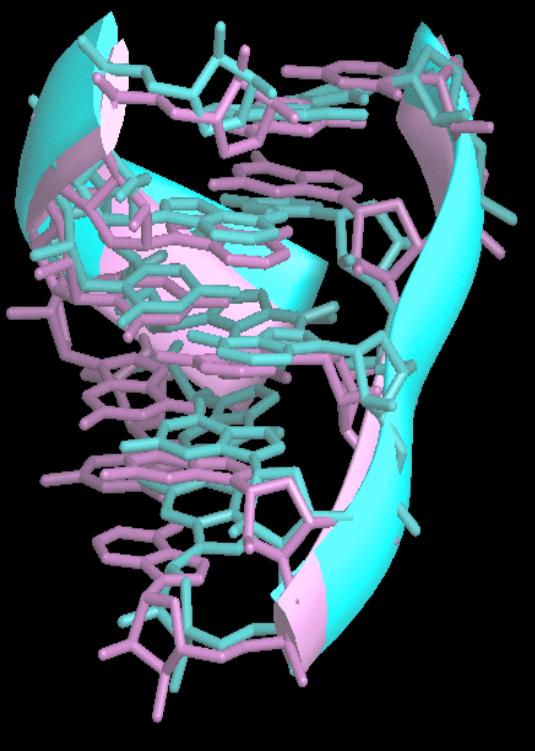
Only
RNA
in
the
reaction
site



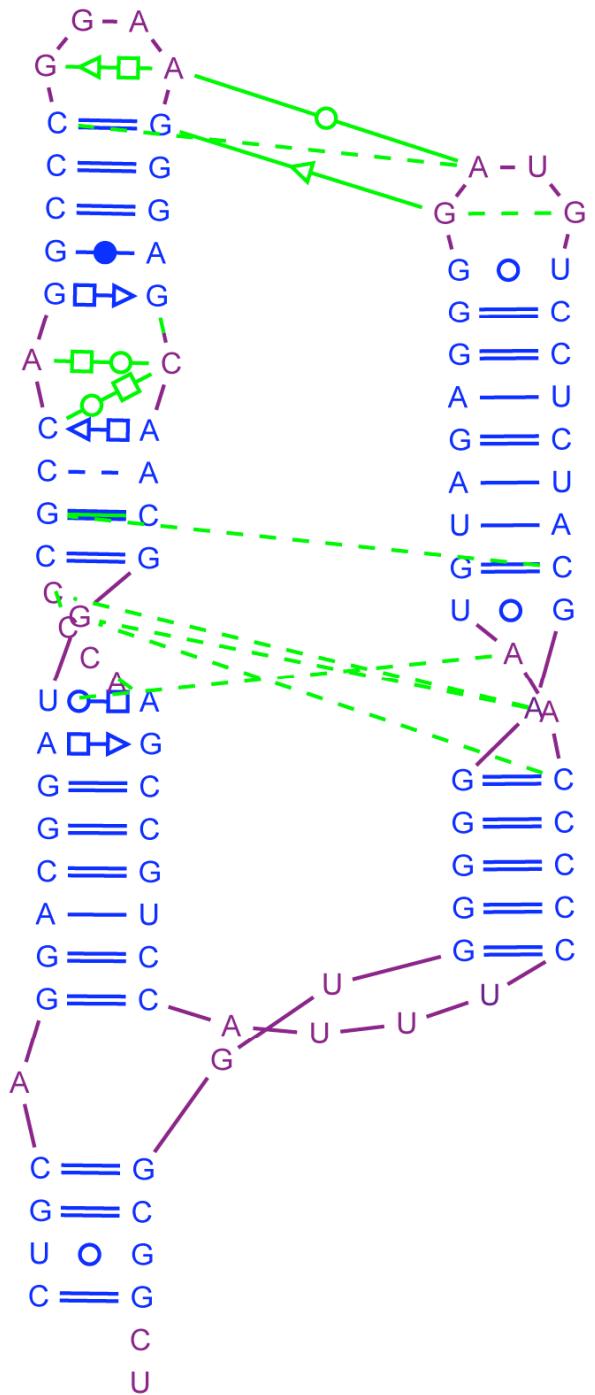
Comparisons of 3D structures



Comparisons of sequences



((((UUAGG	GGA	GUUUUA	UCC	AGCGU	CAG-C
GCCG	UUAGG	GGA	GUUUCA	UCC	AGCGA	UGG-C
GUUG	UAGG	GGA	GUCUCA	UCC	AGCA	CAA-C
GCUG	GAGG	GAA	GC AA	UUC	AGCA	CAG-C
ACUU	CAGU	GGA	GC AA	UCC	AGCA	GAGAU
ACUU	CAGU	GGA	GC AA	UCC	AGCA	GAGAU
GAUG	GAGG	UUG	G AAA	CAA	UGCA	CAU-C
GGGC	CAGG	GGU	G AAA	ACC	AGCA	GCC-A
GGCC	UAGG	UCG	G AAA	CGG	AGCA	GGU-C
GGCC	CAGG	UCG	G AAA	CGG	AGCA	GGU-C
GGCC	CAGG	UCG	G AAA	CGG	AGCA	GGU-C
GGCC	CAGG	UCG	G AAA	CGG	AGCA	GGU-C
GGCC	CAGG	UCG	G AAA	CGG	AGCA	GGU-C



Biological sequence analysis

Protein easy
RNA hard

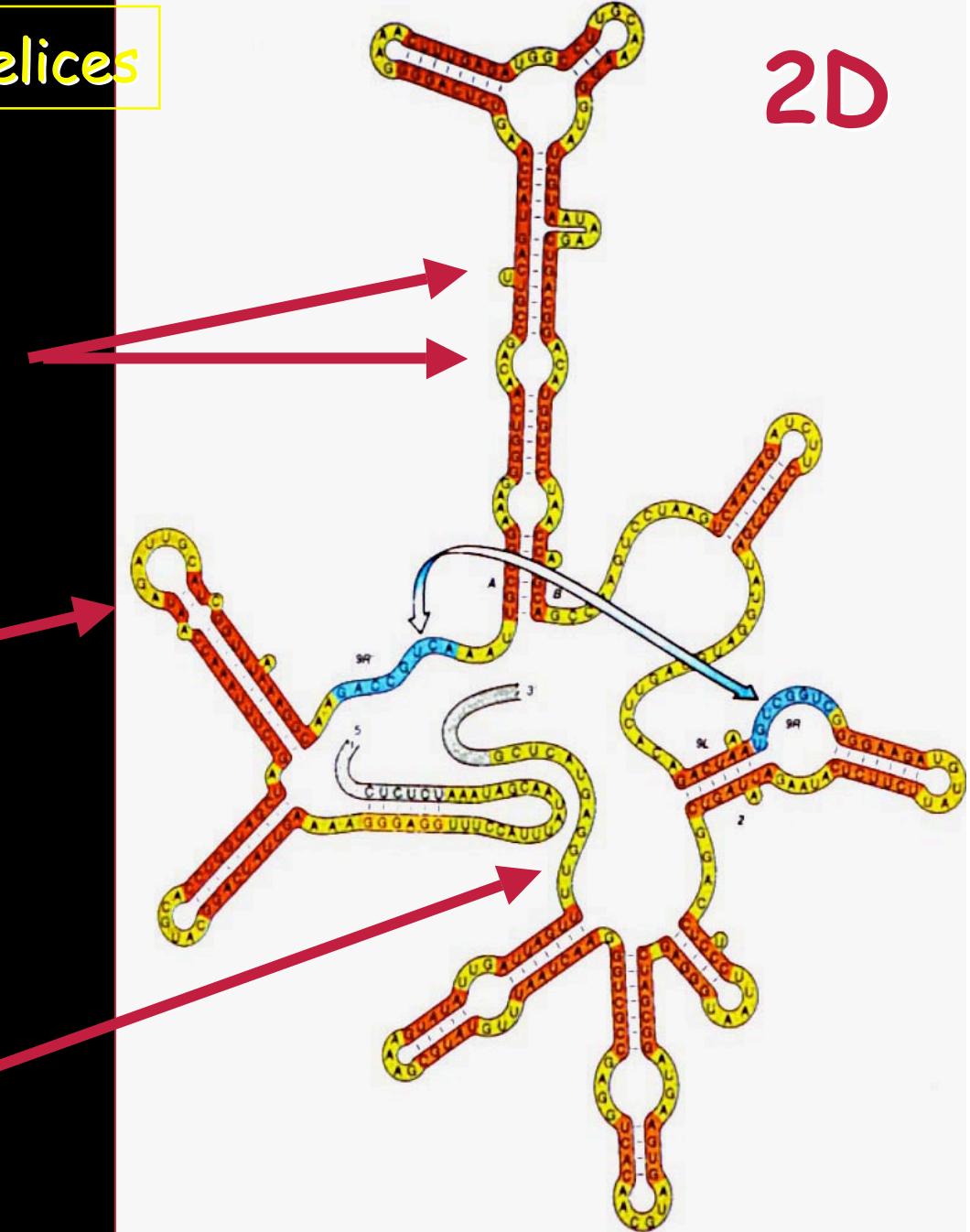
Watson-Crick base paired helices

2D

Internal loops (symmetric,
Asymmetric, bulge)

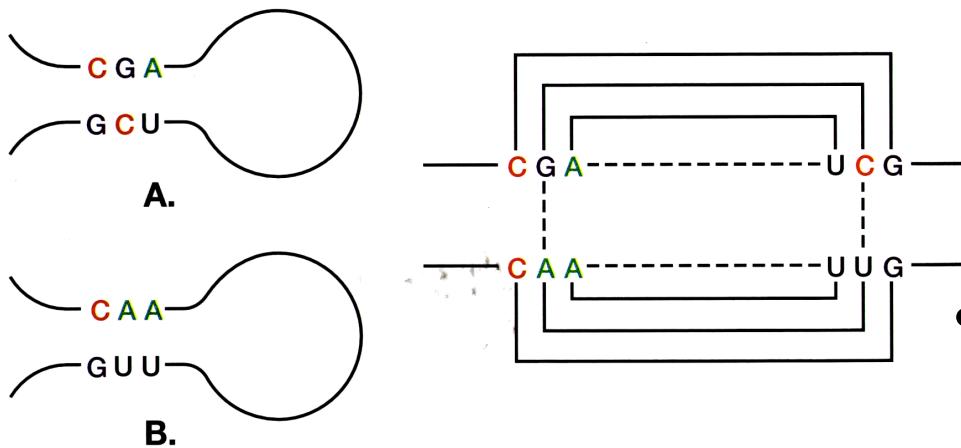
Hairpin loops

Single-strands junctions



RNA alignments

- RNA sequences are aligned/compared differently because sequence variation in RNA maintain base-pairing patterns
 - Thus an alignment will exhibit **covariation** at interacting basepairs



- RNA specifying genes will have conserved regions reflecting common ancestry

Main building block :
the RNA double helix
held together by
Watson-Crick pairs

