

# *1. Biology in a nutshell*

# Cells and organisms

## ■ Organisms

### ☐ Unicellular: One single cell (or simpler)

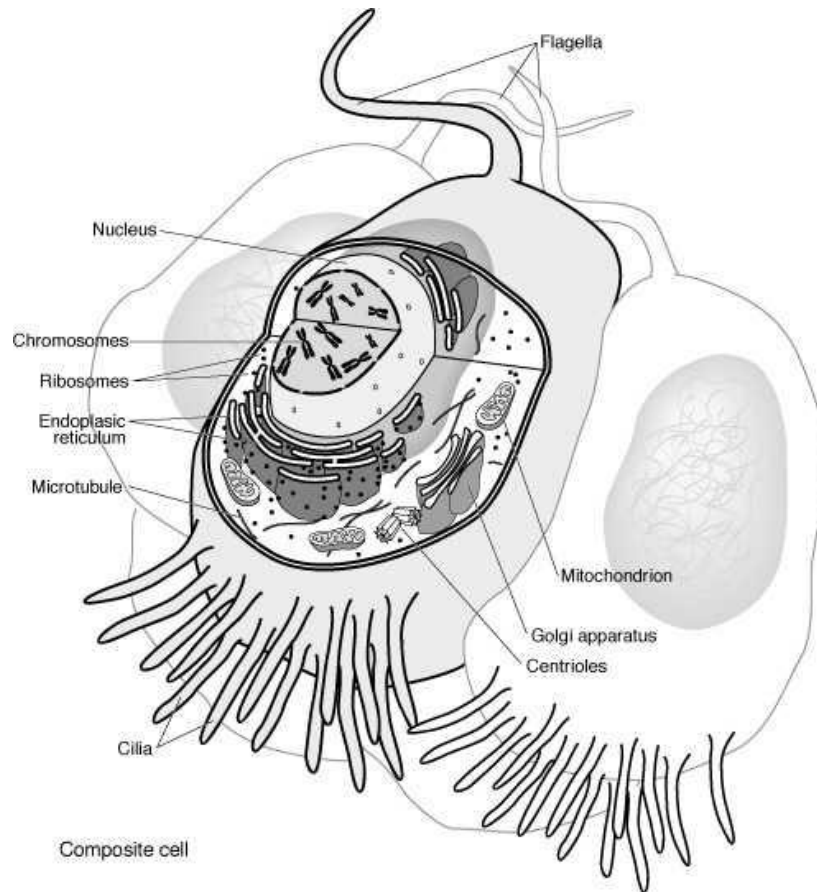
- Archaea
- Prokaryotes: Bacteria, Yeast
- Eukaryote: Protozoos

### ☐ Pluricellular: Eukaryote cells

- Different organisation levels (e.g. tissues)
- Diversity in number, type and syze


### ☐ Viruses are not properly organisms

# Eukaryotes



- “Eu” good,  
“Karyo” nut or kernel →
- Presence of nucleus separated from the cytoplasm by the nuclear envelope.
- DNA: double-stranded, it is organized in chromosomes
- The cell contains other membrane-bound organelles
- Sexual reproduction is common

# Key biomolecules

- The basic components of biological systems are
  - Sugars (carbohydrates)
  - Fats (lipids)
  - Nucleic acids
  - Proteins
- Sugars and lipids have no important role regarding “biological information”   
*Not discussed here*

# Nucleic acids and proteins

- Molecules that contain and transport information
  - DNA (4 different nucleotides)
    - Contains encoded biological information
  - RNA (4 different nucleotides)
    - Carries information from DNA to proteins
  - Proteins (20 different amino acids)
    - Function and structure of living beings

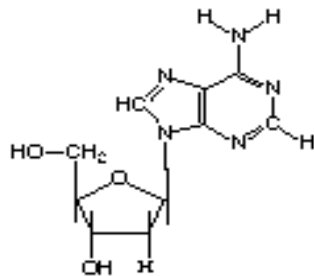
# Nucleotides

- Basic components of nucleic acids
- Consisting of
  - A sugar (Ribose or Deoxyribose)
  - A nitrogen base
  - A phosphate group
- In biological parlance we speak of "bases" instead of nucleotides to describe a string (3000bp = 3000 bp)

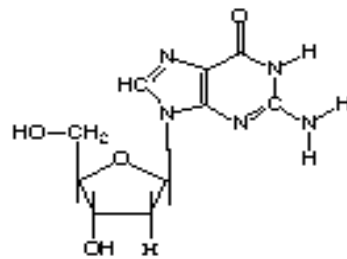
# DNA nucleotides

Deoxyribonucleic Acid (DNA) contains four nucleotide bases.

## The Nucleotides of DNA

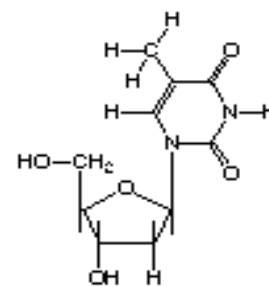


Adenine

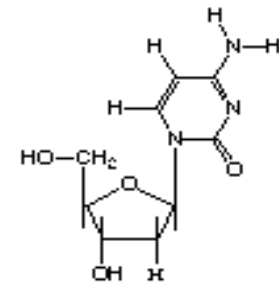


Guanosine

Purines



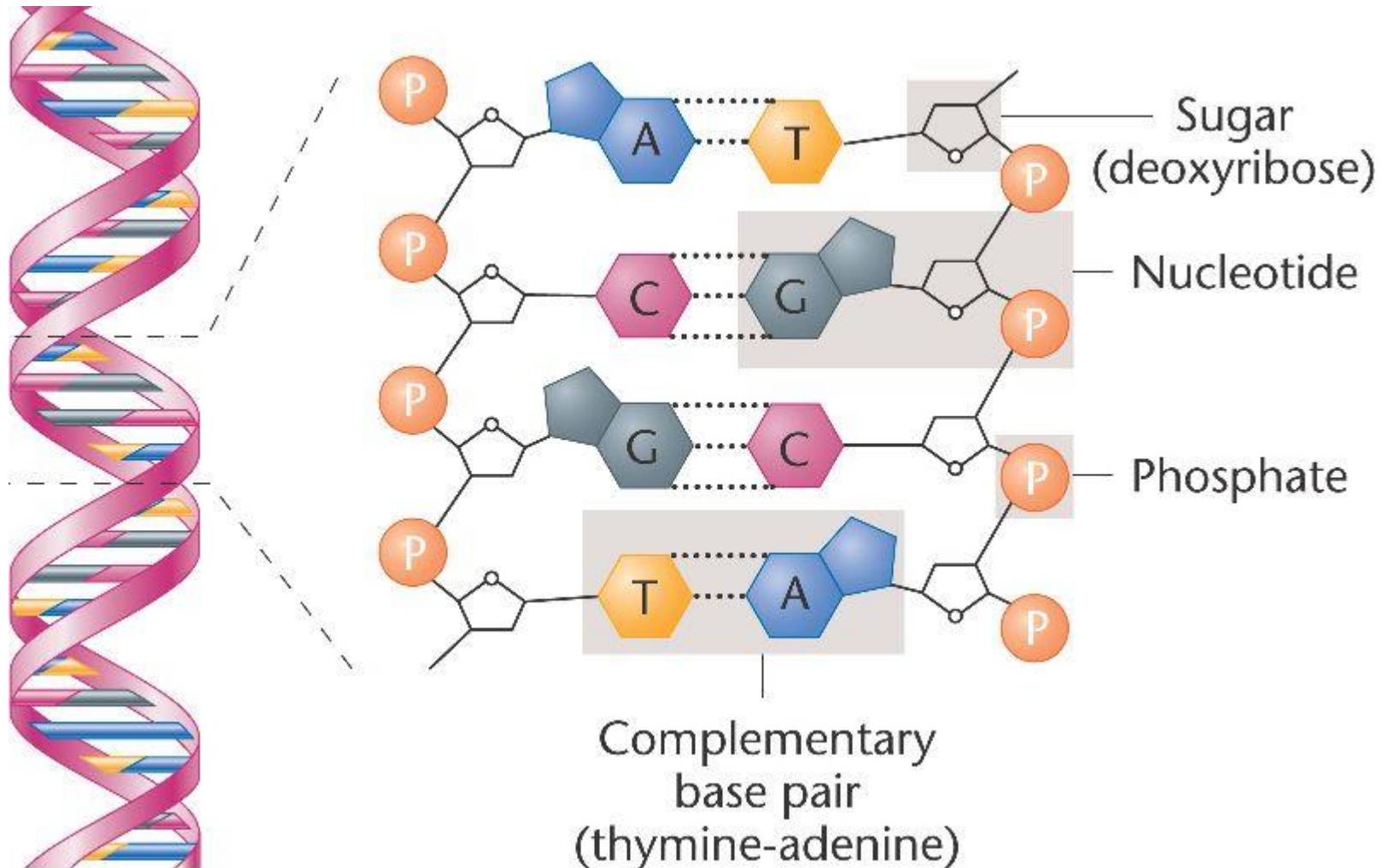
Thymine



Cytosine

Pyrimidines

# The pairing between complementary bases






# The primary structure of DNA

- Sequence of nucleotides
- Forms an unbranched polymer
- Organized in a double-stranded

```
atgaatcgta  ggggtttgaa  cgctggcaat  
acgatgactt  ctcaagcgaa  cattgacgac  
ggcagctgga  aggcggtctc  cgagggcgga
```

# DNA vs RNA

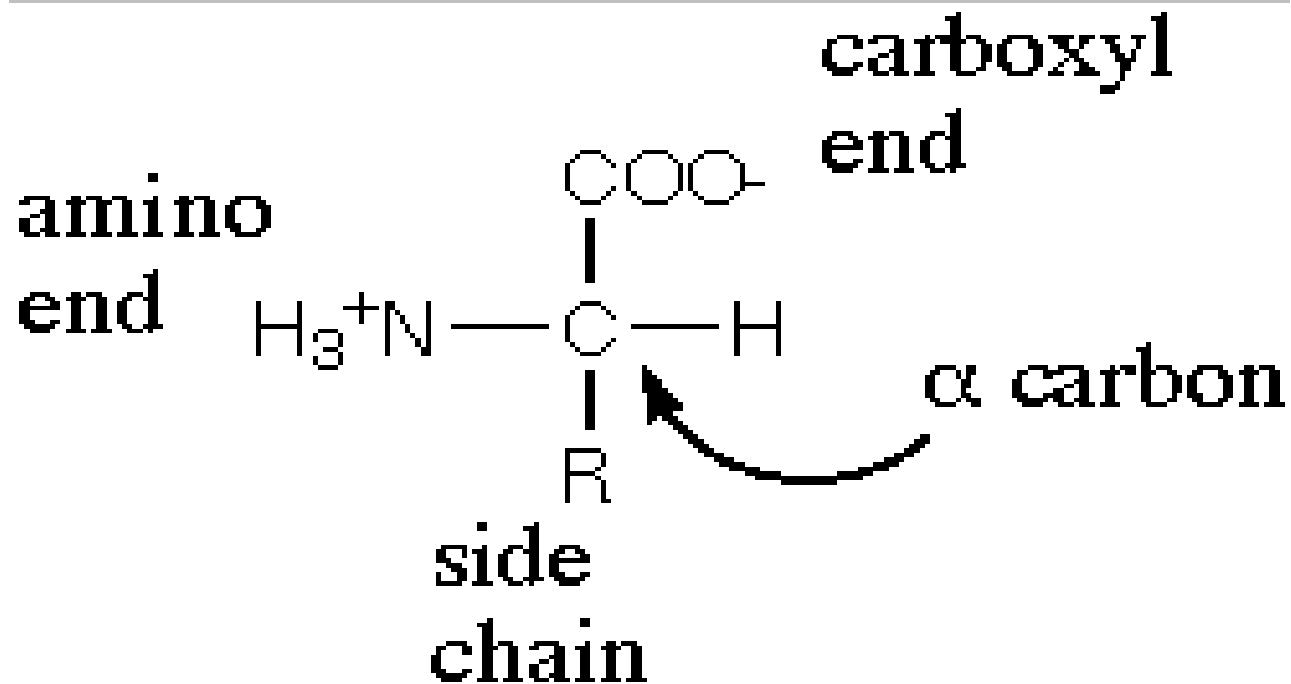
- DNA is organized into a complementary double helix. RNA does not.
- One of the four bases are different
  - DNA  A, C, G, T
  - RNA: A, C, G, U
- Differ from the nucleotide sugar
  - DNA: Deoxyribose
  - ARN: Ribose

# Proteins

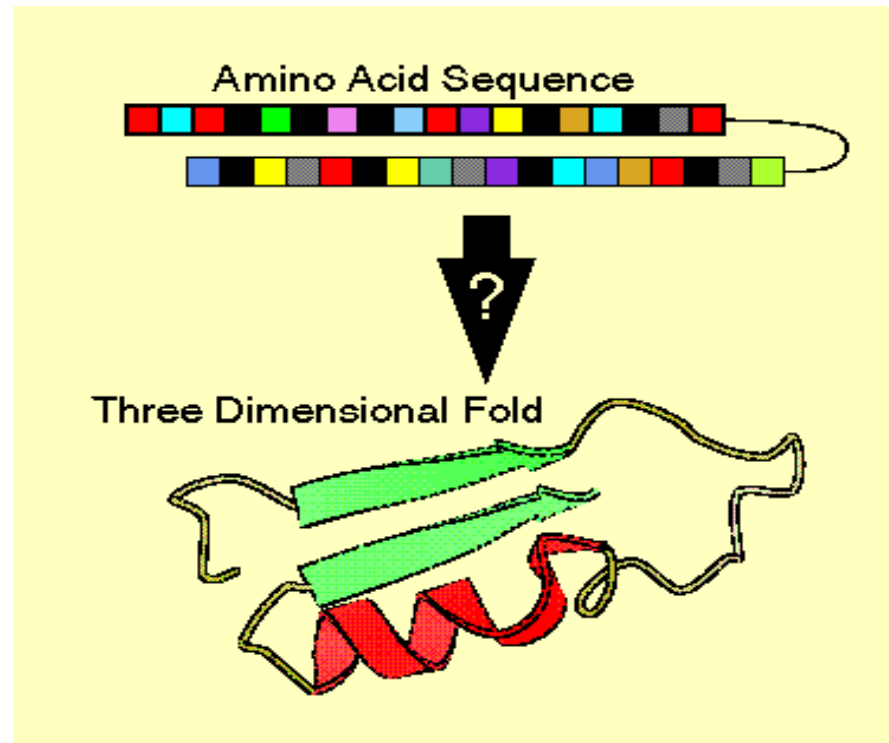
- Amino acid sequence
- Forms an unbranched polymer
- There are 20 different amino acids (AA)
- The key function of proteins is in its three dimensional structure

# Amino acids

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# Proteins “fold” into conformational structure



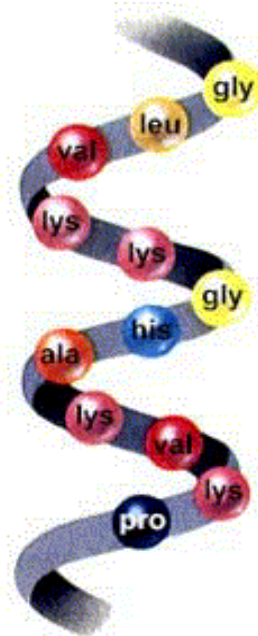
# Protein's structure

Proteins fold into lowest to highest complexity level:

- Primary structure: amino acid sequence
- Secondary structure: regularly repeating local structures stabilized by hydrogen bonds.
- Tertiary structure: fold into 3-dimensional structures.
- Quaternary structure: structure formed by several protein molecules (protein complex).



primary structure  
(amino acid sequence)



secondary structure  
( $\alpha$ -helix)

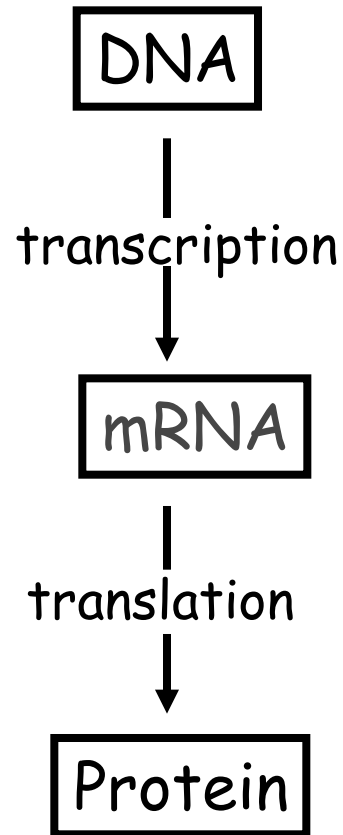
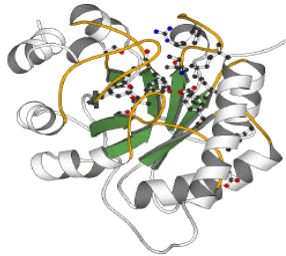
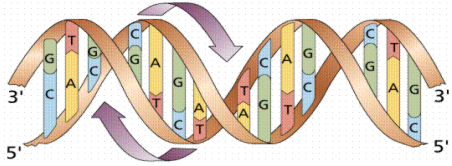


tertiary structure  
(folded individual peptide)



quaternary structure  
(aggregation of two or more peptides)

# The central dogma of molecular biology



CCTGAGCCAAC TATTGATGAA



CCUGAGCCAACUAUUGAUGAA



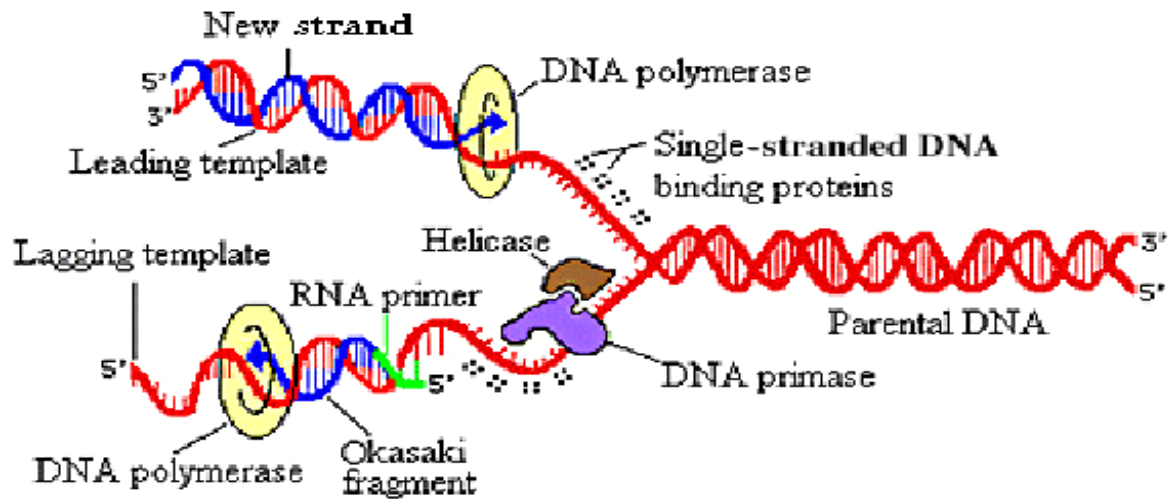
PEPTIDE



# From DNA to proteins

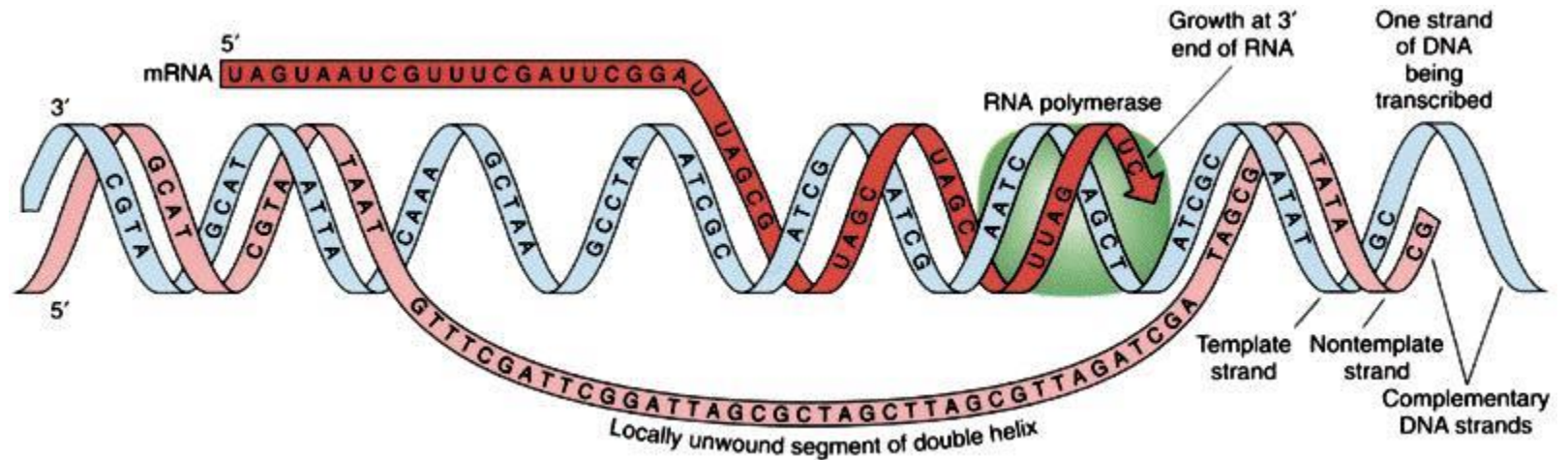
- DNA is replicated in a complex process involving many enzymes: replication
- DNA is copied in a string of complementary messenger RNA (mRNA): transcription transcription
- In eukaryotic cells, the mRNA is processed splicing eliminating coding fragments (“splicing”) and migrates from the nucleus to the cytoplasm.
- The mRNA carries coded information to ribosomes (ribosomal RNA) that "read" and perform protein synthesis: translation

# 1. Replication

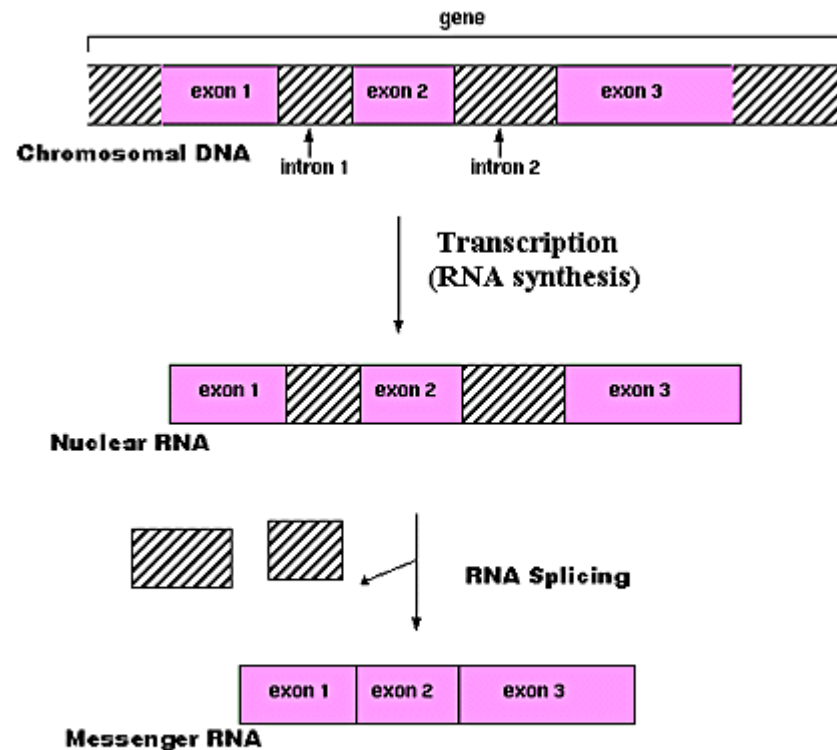


- Animations (1), (2), (3)

## 2. Transcription

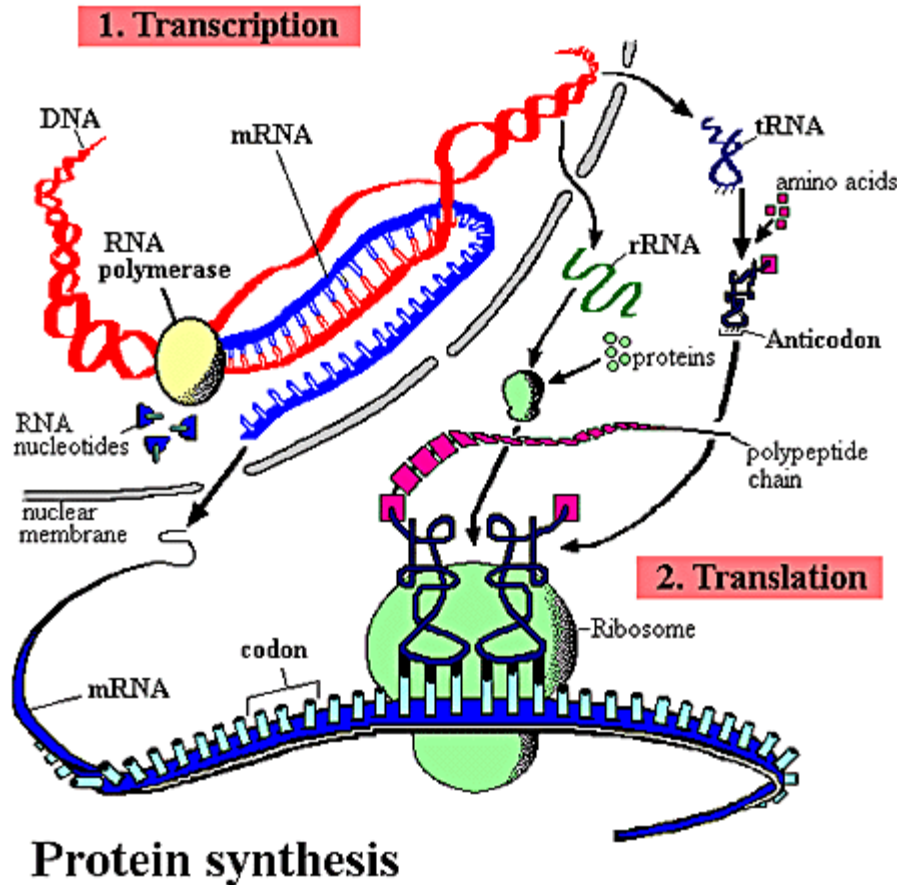


### 3. RNA processing or splicing



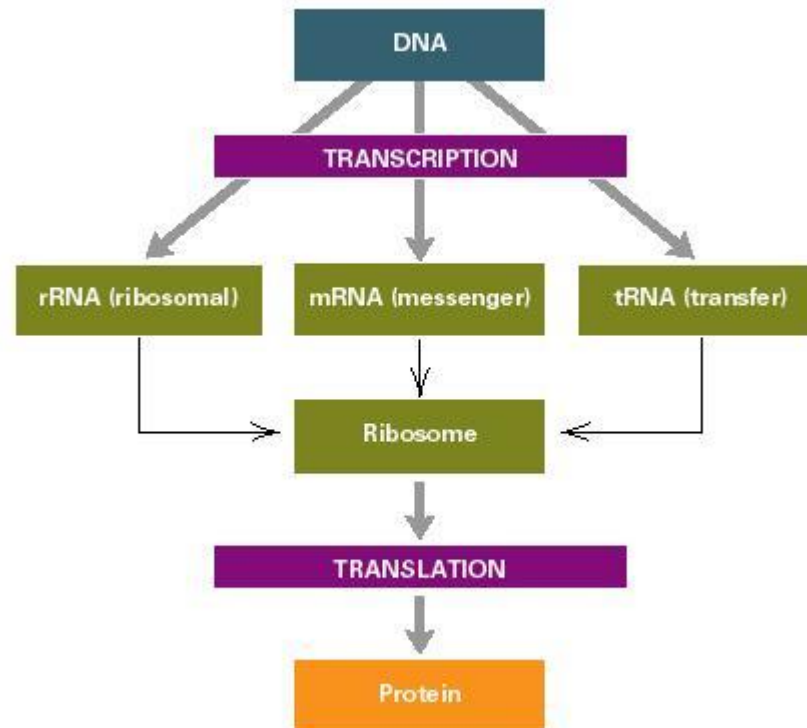
RNA synthesis and processing

# 4. Protein synthesis

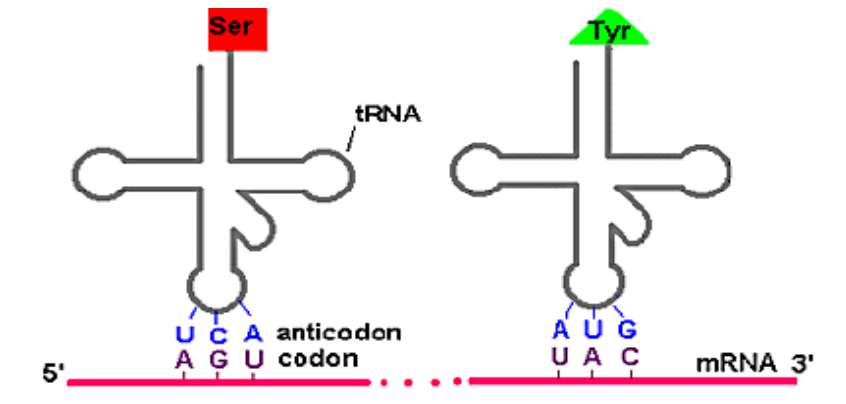


- The mRNA goes to the cytoplasm where it binds to ribosomes.
- Codon: mRNA information unit.
- The tRNA brings the complementary AA tRNA.
- The AA are bound to the protein to complete the sequence.
- Animations (1), (2)

# Notice: 1 DNA vs. 3 RNA's !



# The genetic code



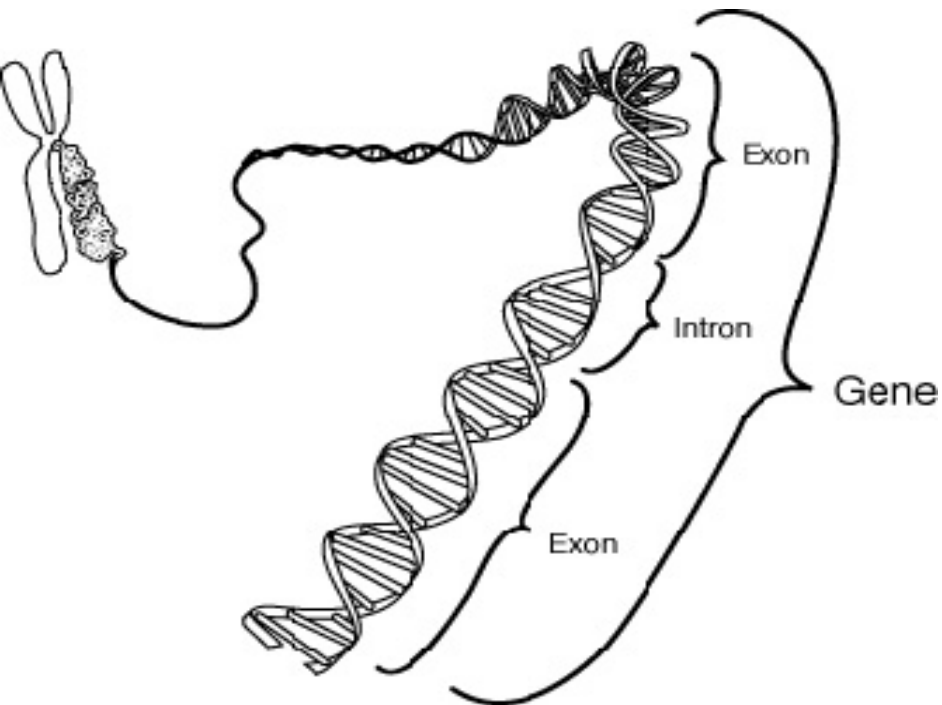
		2nd base in codon				3rd base in codon
		U	C	A	G	
1st base in codon	U	Phe Phe Leu Leu	Ser Ser Ser Ser	Tyr Tyr STOP STOP	Cys Cys STOP Trp	U C A G
	C	Leu Leu Leu Leu	Pro Pro Pro Pro	His His Gln Gln	Arg Arg Arg Arg	U C A G
	A	Ile Ile Ile Met	Thr Thr Thr Thr	Asn Asn Lys Lys	Ser Ser Arg Arg	U C A G
	G	Val Val Val Val	Ala Ala Ala Ala	Asp Asp Glu Glu	Gly Gly Gly Gly	U C A G

# What is a gene? (pre ENCODE)

- A gene used to be defined as ...
  - The functional and physical unit of heredity,
  - transmitted from one generation to their offsprings,
  - consisting of DNA fragments,
  - with the information needed for the synthesis of a specific protein (most of them...)



# Gene components



	DNA	Transcription	mRNA	tRNA	Amino Acid	Polypeptide chain
G	Exon	A C G C G A T A G	T G C G C U A U C G A C U G C A C A U G A C C U A	A C G	◆	◆ * ●
				C G C	*	
				G A U		
E	Intron	O T G A C G T T G T A C T G A A	C A C U G C A C C A U G A C C U A	G O U G A C G U G U A C U G A A		
N	Exon	T T G T A C T G A A	A C C A U G A C C U A	A C C	▲	▲ ● ■
				C A U	●	
				G A C	■	



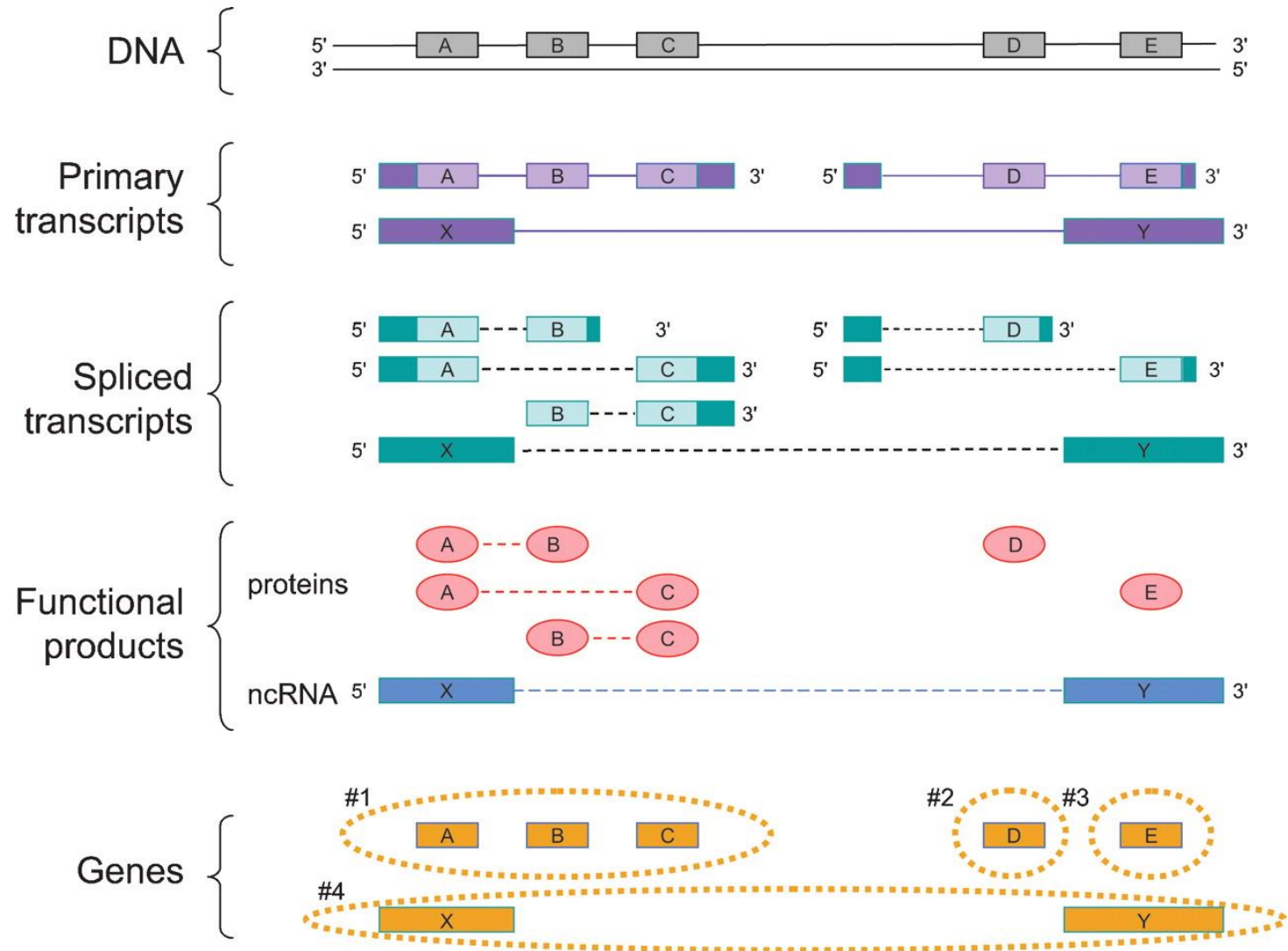
# The ENCODE project

In September 2003 The National Human Genome Research Institute (NHGRI) launched a public research consortium named ENCODE, the Encyclopedia Of DNA Elements

Its goal was *“to carry out a project to identify all functional elements in the human genome sequence”*

The findings of the ENCODE project have led to reconsider fundamental ideas such as the definition of gene

# What is a gene (post-ENCODE)



# Gene expression

- Genes can be “turned on” or “off”
- When a gene is turned on we say it is *expressing* or *being expressed*
  - *The central dogma implies that when a gene is expressed it is, at least transcribed*
  - *That is: expression  $\rightarrow$  mRNA*
- Each cell expresses only a fraction of its genes
- Remaining genes are repressed ("off")

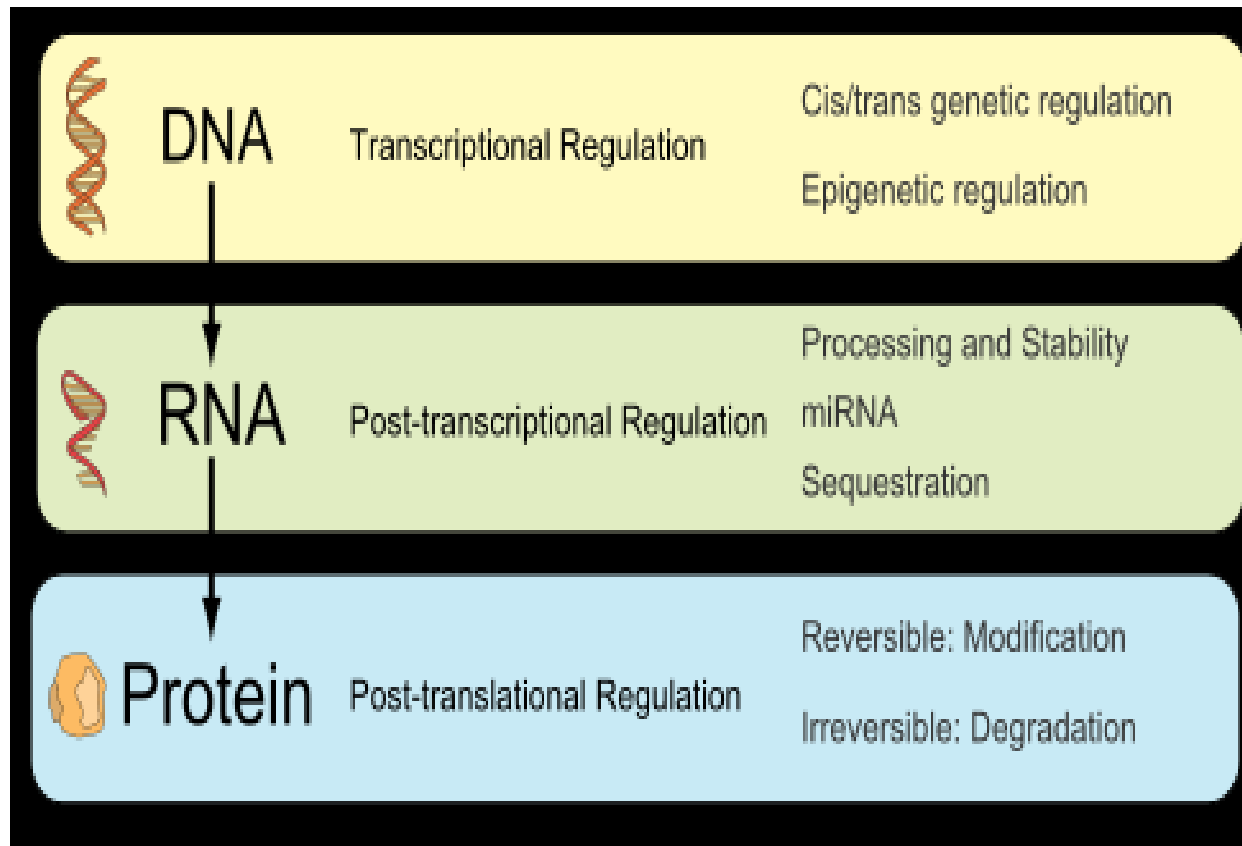
# Regulation of gene expression

- Each cell expresses (or active or "on") only a fraction of their genes.
- Remaining genes are repressed ("off").
- The process consisting of activating some genes and inhibiting others is globally known as *gene regulation*.
- Gene regulation determines:
  - The appearance and different function of different cells types
  - The ability of some cells to react quickly to environmental changes

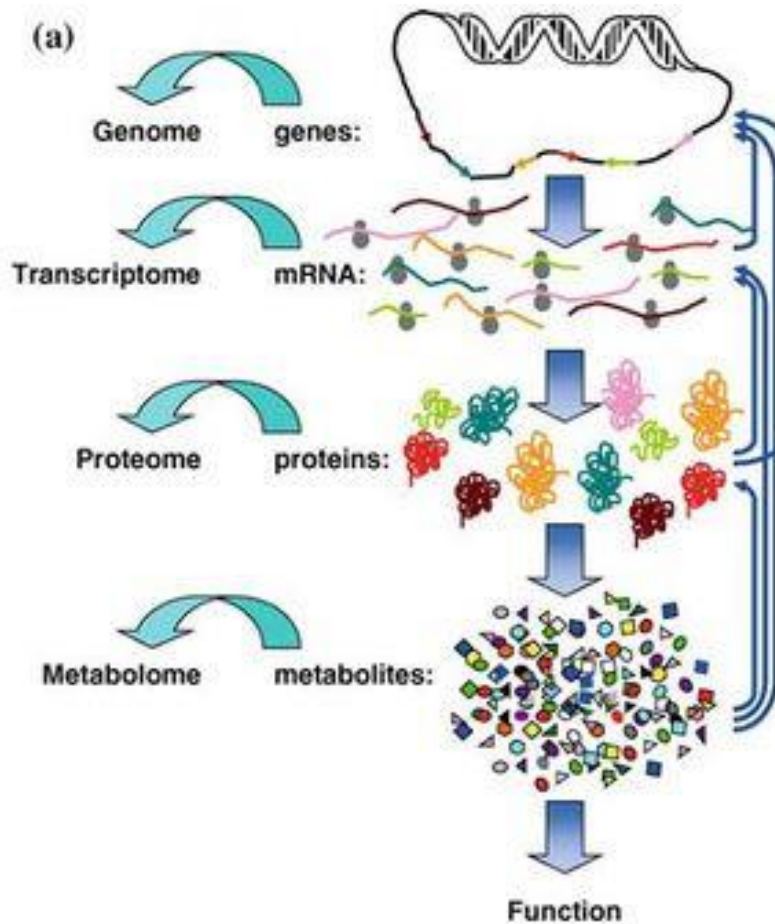
# How are genes regulated?

- Gene regulation can occur at any point in the process of expression but often occurs during transcription.
- Environmental signs or other cells activate proteins called *transcription factor*.
- They bind to the *regulatory regions* of genes, increasing or decreasing the level of transcription → They control the amount of gene product produced by the gene in every moment.

# Forms of gene expression regulation



# Put it altogether: *The OMICS cascade*



## The “Omics” Cascade

*What can happen*

GENOME

*What appears to be happening*

TRANSCRIPTOME

*What makes it happen*

PROTEOME

*What has happened and is happening*

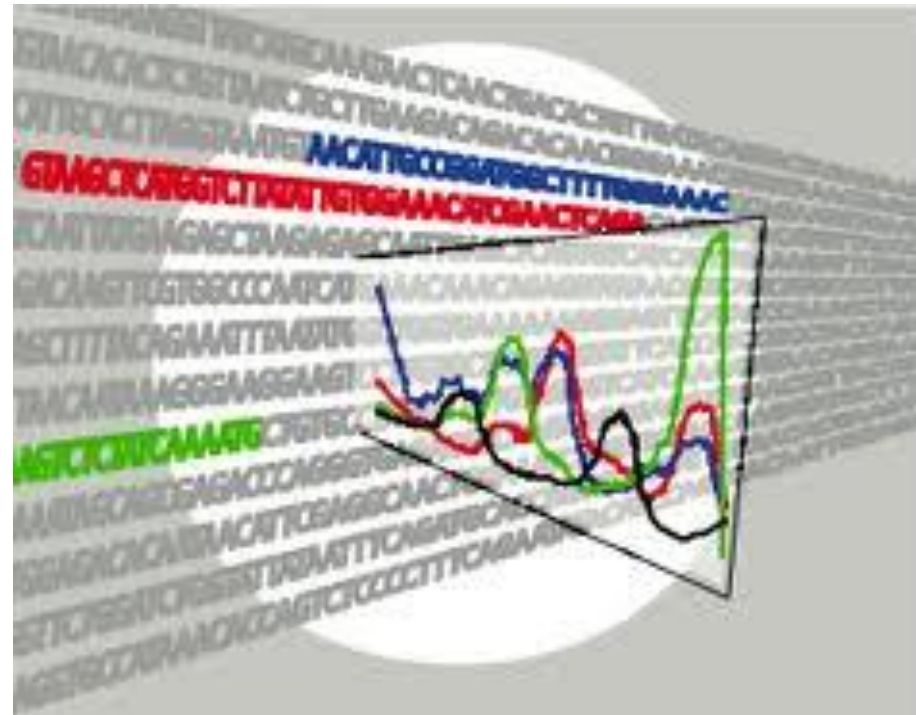
METABOLOME

PHENOTYPE



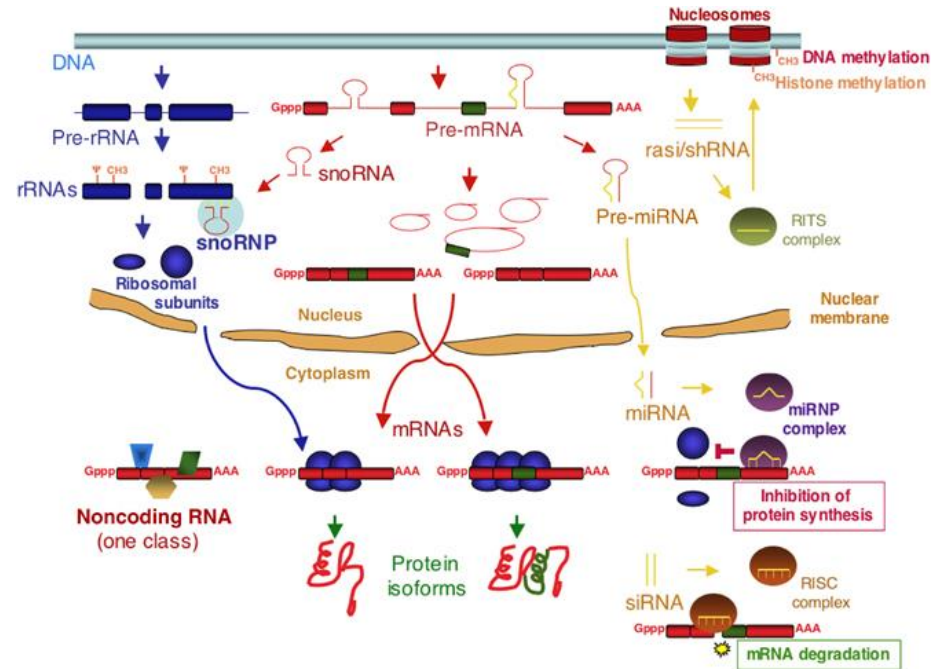
# Genomics

- A genome is an organism's complete set of DNA, including all of its genes
- Genomics relies on *sequencing technologies* to study genomes and intragenomic phenomena



# Transcriptomics

- The transcriptome is the set of all RNA molecules, in one or a population of cells.
- **Transcriptomics**, examines expression levels of mRNAs in a given cell population, often using high-throughput techniques such as
  - RTqPCR
  - microarrays or
  - NGS.



# 'omics' are high throughput

- Most 'omic' approaches generate huge quantities of data.
- The management, storage, analysis and interpretation of these high throughput datasets cannot be conceived without all type of computing and quantitative resources
- ***Bioinformatics and computational biology are a must for omics sciences***