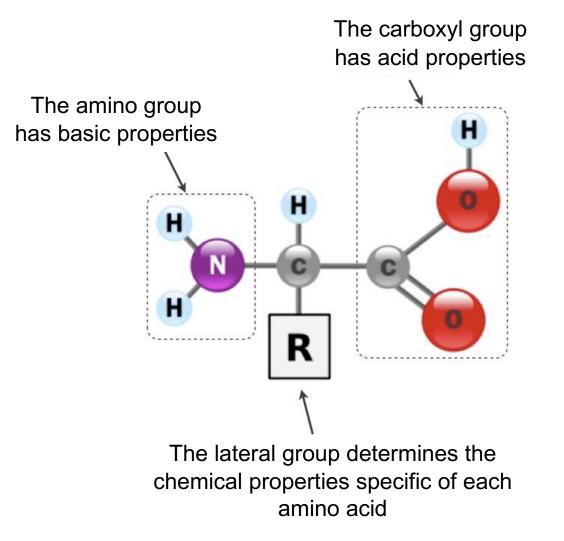
Astrobiology Lecture 4

Metabolic & genetic macromolecules

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Biological amino acids Constituents of proteins



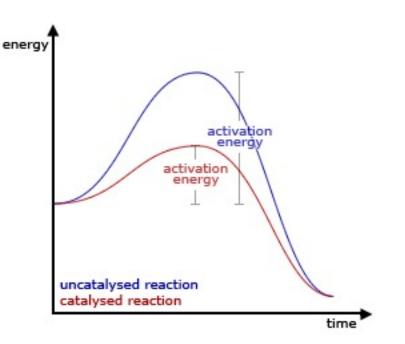
Biological macromolecules Proteins

Proteins are polymers of amino acids

Short chains of amino acids are called <u>peptydes</u>
Long, unbranched peptyde chains are called <u>polypeptides</u>
Proteins are formed by one or more chains of polypeptides
Molecular masses of proteins vary between ~10³ e ~10⁶ atomic mass units
They contribute to about half the mass of the cell

Proteins play fundamental functions in living organisms

Mostly <u>structural</u> and <u>enzymatic</u> (i.e., catalytic) functions



Terrestrial biological amino acids

Proteins use only 20 types of amino acids

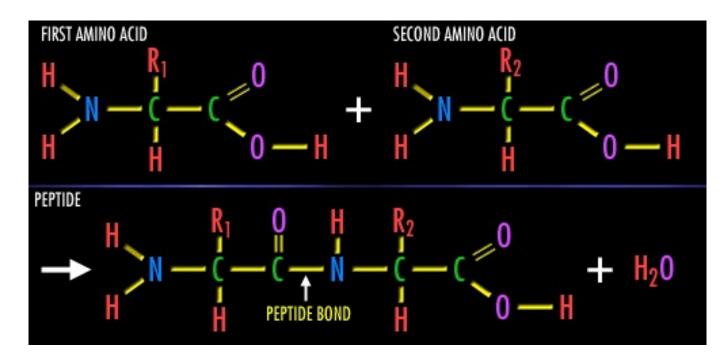
Organic chemistry allows for the existence of <u>thousands</u> of amino acids

Apparently, terrestrial life has "<u>chosen</u>" a <u>short list</u> of amino acids, sufficiently representative of the different types of chemical properties that are required to build up the variety of proteins necessary to living organisms

Table 7.2 The Twenty Amino AcidsFound in Living Organisms

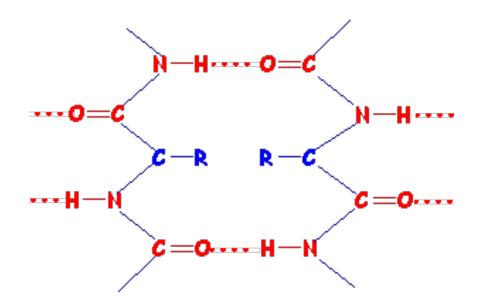
 Amino Acid*	Chemical Formula	Number of Atoms
L-Alanine	$C_3H_7O_2N$	13
L-Arginine	$C_{6}H_{15}O_{2}N_{4}$	27
L-Asparagine	$C_4H_8O_3N_2$	17
L-Aspartic Acid	$C_4H_6O_4N$	15
L-Cysteine	$C_3H_7O_2NS$	14
L-Glutamic Acid	$C_5H_8O_4N$	18
L-Glutamine	$C_5H_{10}O_3N_2$	20
Glycine	$C_2H_5O_2N$	10
L-Histidine	$C_6H_9O_2N_3$	20
L-Isoleucine	$C_{6}H_{13}O_{2}N$	22
L-Leucine	$C_{6}H_{13}O_{2}N$	22
L-Lysine	$C_6H_{15}O_2N_2$	25
L-Methionine	$C_5H_{11}O_2NS$	20
L-Phenylalanine	$C_9H_{11}O_2N$	23
L-Proline	$C_5H_9O_2N$	17
L-Serine	$C_3H_7O_3N$	14
L-Threonine	$C_4H_9O_3N$	17
L-Tryptophan	$C_{11}H_{12}O_2N_2$	27
L-Tyrosine	$C_9H_{11}O_3N$	24
L-Valine	$C_5H_{11}O_2N$	19

From amino acids to polypeptides

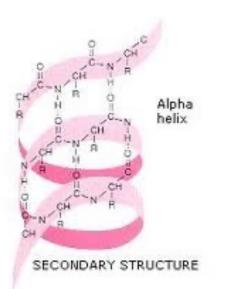


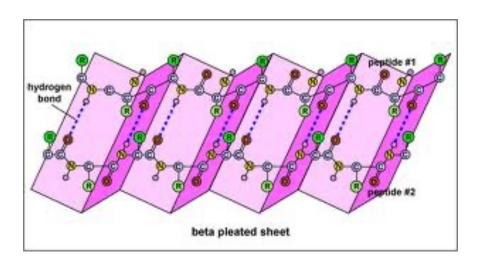
Amino acids are bound to each other with peptide bonds
The carboxyl end ties to the amino end of the next molecule
A sequence OC-NH is formed (peptide bond)
A water molecule is released each time a peptide bond is created
A water molecule is required to break a peptide bond

From polypeptides to 3D proteins Importance of hydrogen bonds as intramolecular forces



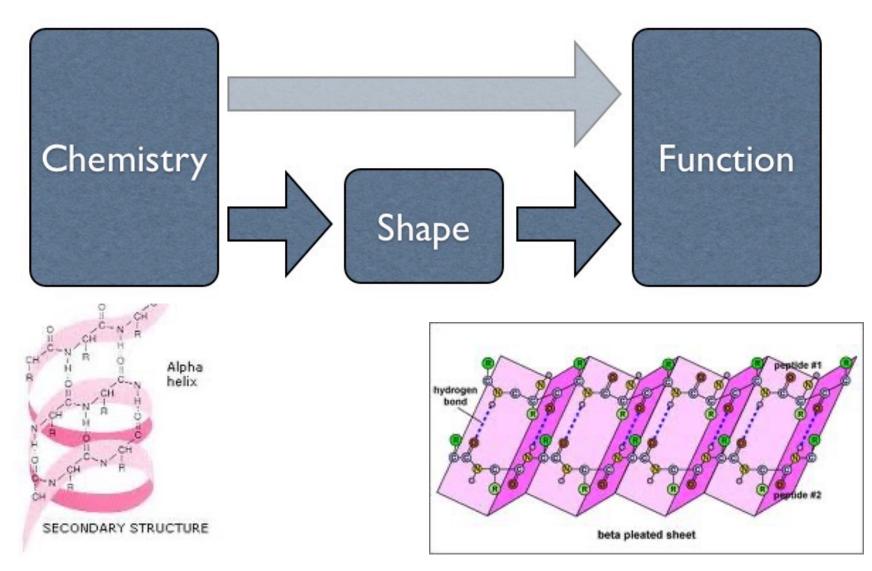
Spontaneous formation of geometrical configurations with lowest potential energy





Proteins The importance of 3D structures

The shape of proteins determines, in large part, their function



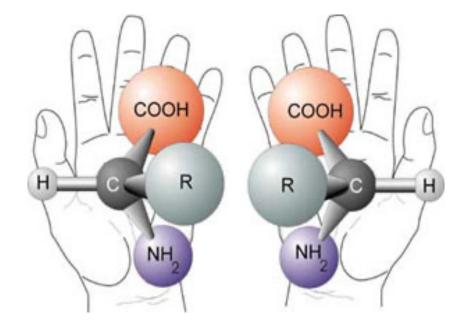
Chirality

Molecules with same chemical formula but different structure are called *isomers*

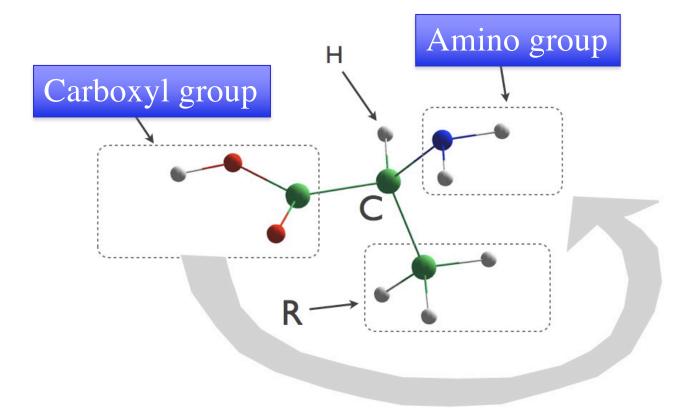
<u>Chiral</u> molecules are isomers with a center of symmetry ("stereocenter") They cannot be superimposed to their mirror image The two mirror images of a chiral molecule are called <u>enantiomers</u>

Amino acids are chiral

The carbon atom at the center of the amino acid is the stereocenter



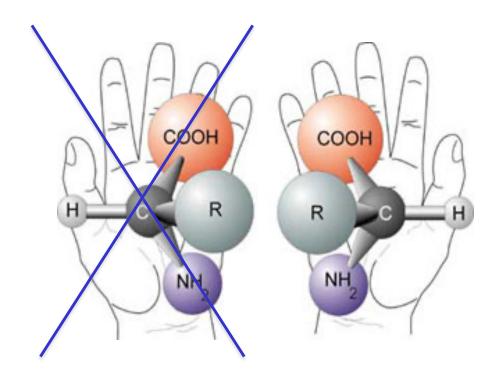
The two enantiomers of amino acids are called L and D according to the "CORN" convention



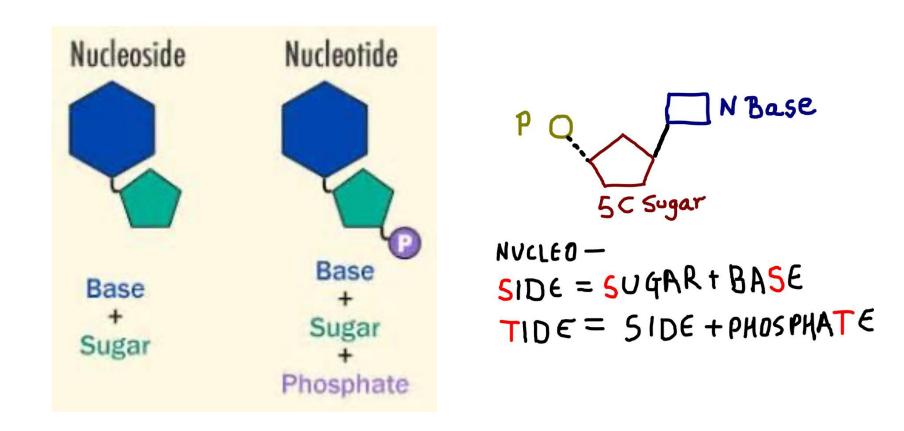
Example of L-type amino acid: L-alanine

Homochirality of biomolecules

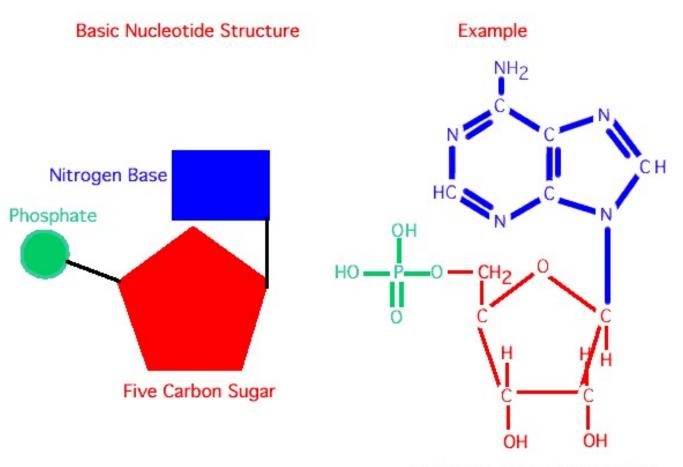
Protein amino acids are <u>homochiral</u>: they only show the L-type enantiomer



Nucleosides and nucleotides Constituents of the DNA and RNA

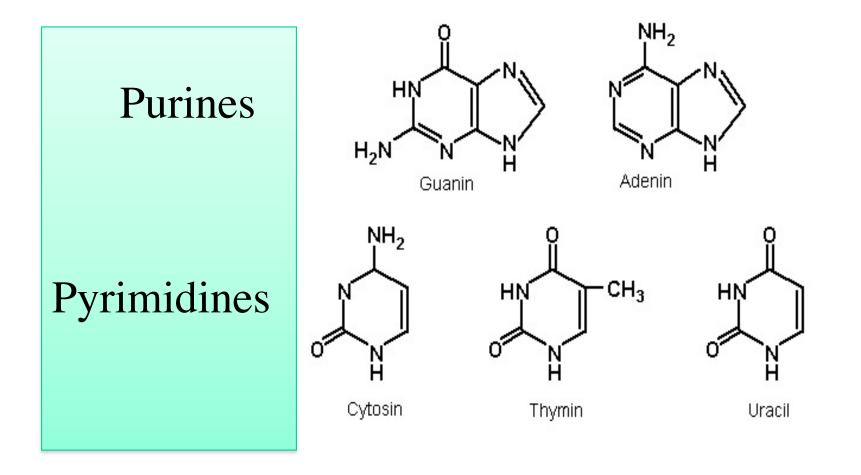


Nucleotides Constituents of the DNA and RNA



Adenosine 5' phosphoric acid

Nitrogen bases aromatic rings with N substitutions



Biological macromolecules Nucleic acids

Nucleic acids are polymers of <u>nucleotides</u>

Depending on the type of organism, they may contain $\sim 10^6 - 10^8$ nucleotides

Nucleic acids store and use the genetic information There are two types: DNA and RNA DNA preserves the information RNA, which comes in different forms, uses the information for driving metabolic/replication processes

Nucleic acids: RNA

RNA has a single strand of nucleotides

The backbone of the strand is made up of a sequence of phosphate groups and <u>ribose</u> sugars

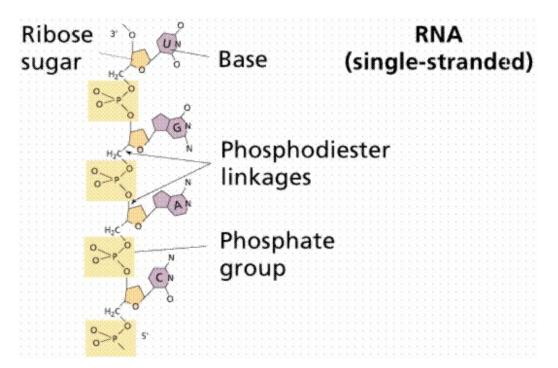
Has 4 types of nucleobases

Purines

Adenine, Guanine Pyrimidines Cytosine, Uracyl

RNA drives the synthesis of proteins

The order of the nitrogen bases on the backbone determines the sequence in which amino acids are assembled



Nucleic acids: DNA

DNA has two strands that form a double helix structure

The backbone of each strand is made up of a sequence of phosphate groups and <u>deoxyribose</u> sugars

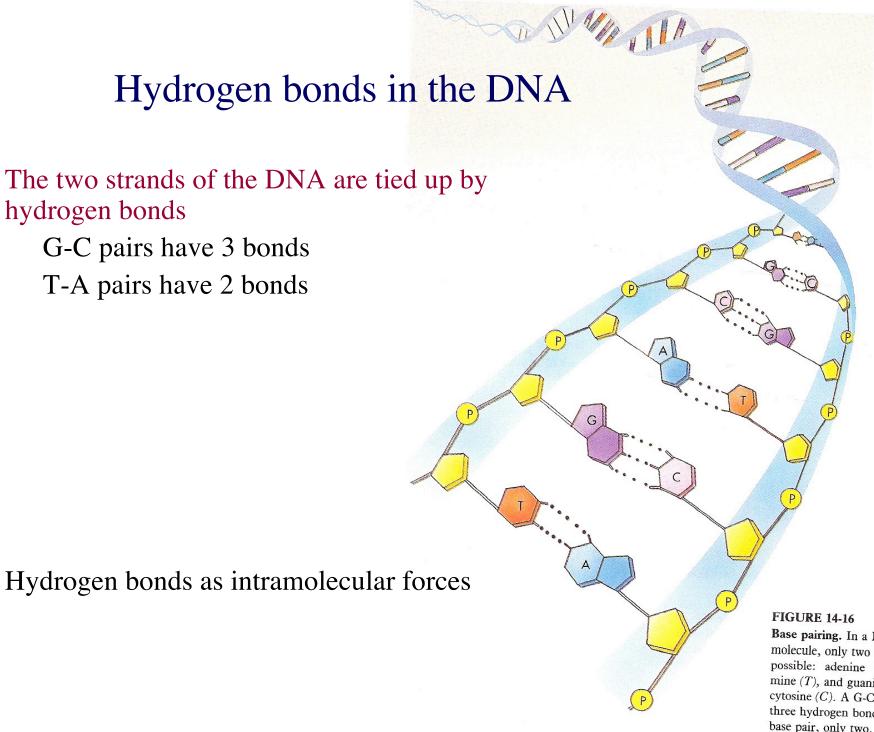
DNA has 4 types of nucleobases

- 2 purins
 - Adenine e Guanine
- 2 pyrimidins
 - Cytosine e Thymine

Thymine replaces Uracyl, which is instead used in the RNA

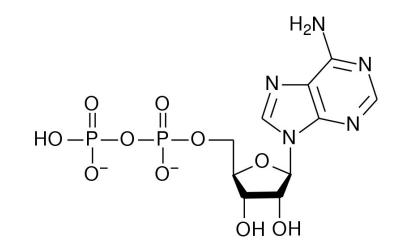
The complementarity of purines and pyrimidines is essential for pairing the two strands

FIGURE 14-16 Base pairing. In a DNA molecule, only two base 1 possible: adenine (A) x mine (T), and guanine (Gcytosine (C). A G-C base three hydrogen bonds; an base pair, only two.



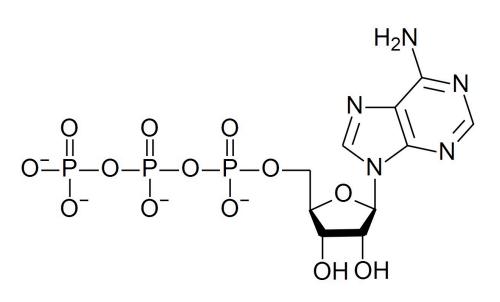
Base pairing. In a DNA molecule, only two base 1 possible: adenine (A) v mine (T), and guanine (Gcytosine (C). A G-C base three hydrogen bonds; an

ADP & ATP Energy exchange molecules



ADP

Adenosine diphosphate



ATP

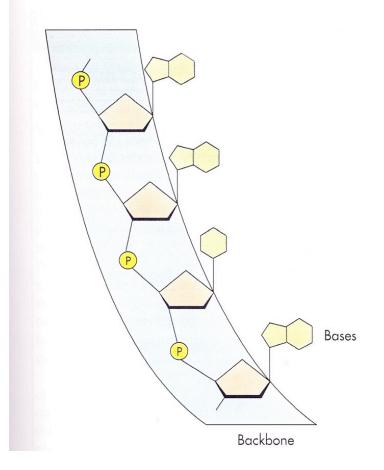
Adenosine triphosphate

Genetic information

The <u>order</u> of the nucleobases attached to the backbone of the nucleic acids determines the genetic information, which is therefore stored in <u>digital form</u> The <u>order of the nucleobases</u> is not constrained by chemical laws We believe this is the result of natural selection at early stages of life evolution

Digital information is more stable than analogic information

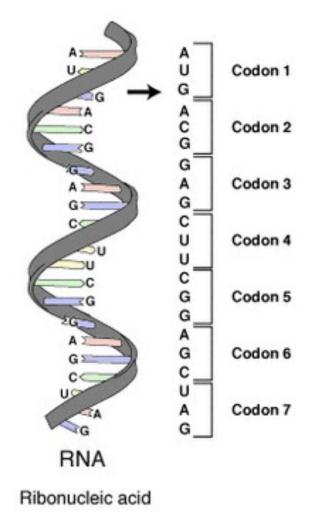
Terrestrial life has started to use digital information more than 3.5 billon years ago



The genetic information and genetic code

The <u>digital information</u> is coded <u>in triplets of</u> <u>nucleobases</u> called <u>codons</u>

Each codon uses 3 of the 4 nucleobases and can express $4^3=64$ possible combinations, equivalent to 6 bits of information (64=2⁶)



Each codon uniquely identifies a single amino acid Some aminoacids are coded by more than one codon (example of unplanned evolution) Some codons are used as a "stop" signal of the sequence

Correspondenc e between RNA codons and amino acids

> A=Adenine G=Guanine C=Citosine U=Uracyl

			0000				
		U	С	А	G		
	U	UUU UUC UUA UUA UUG	UCU UCC UCA UCG	UAU UAC Tyr UAA Stop UAG Stop	UGU UGC UGA Stop UGG Trp	U C A G	Third letter
First letter	С	CUU CUC CUA CUG	CCU CCC CCA CCG	CAU CAC CAA CAA CAG Gln	CGU CGC CGA CGG	U C A G	
Firs	A	AUU AUC AUA AUG Met	ACU ACC ACA ACG	AAU AAC AAA AAG Lys	AGU AGC AGA AGG Arg	U C A G	
	G	GUU GUC GUA GUG	GCU GCC GCA GCG	GAU GAC GAA GAA GAG	GGU GGC GGA GGG	U C A G	

Second letter

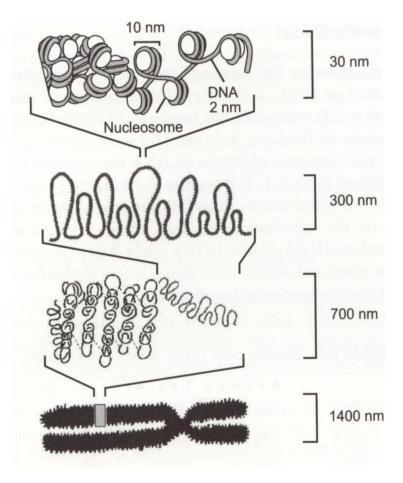
Genes

From the structural point of view, a gene is a <u>sequence of nucleobases</u> along a strand of a nucleic acid

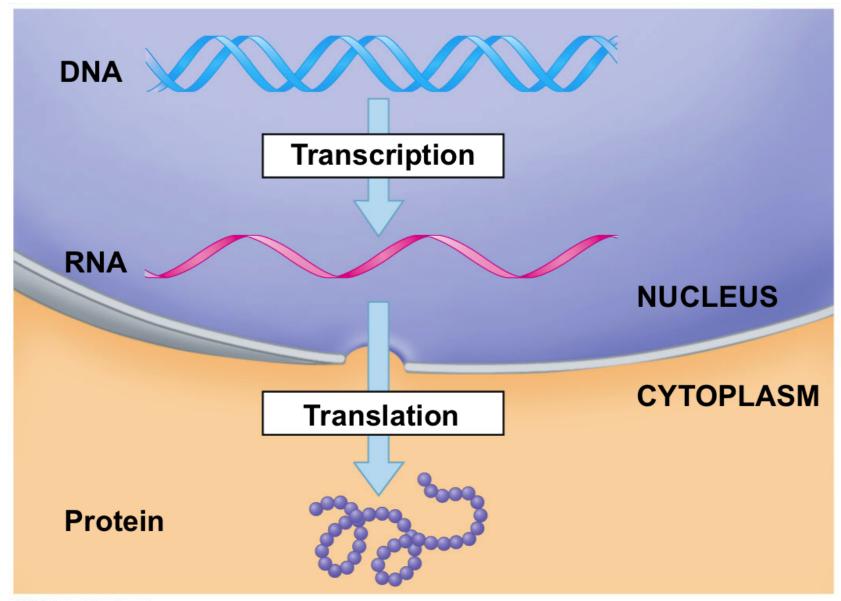
From the point of view of the information content, a gene is a <u>sequence of instructions</u> with a specific function

As an example, a sequence that specifies how to build up a specific amino acid

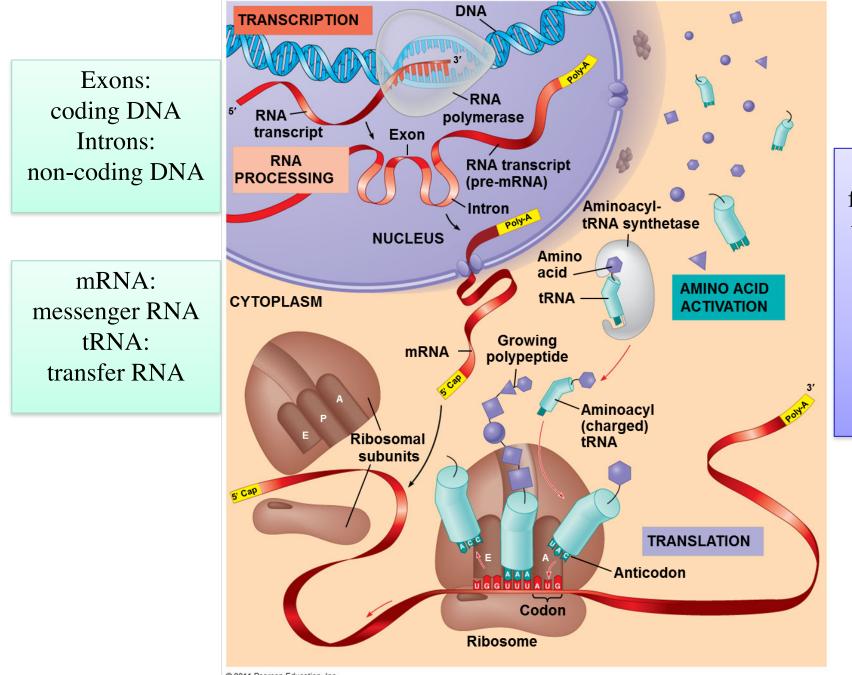
In complex organisms, the number of genes is extremely high and this is why DNA needs to be stored in very compact structures, such as chromosomes



The "central dogma"



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A closer look to the molecular machinery

The functioning of the molecular machinery requires the presence of nuclei acids <u>and proteins</u> (enzymes) Appendix: the DNA as a digital storage device

- The density of information (MB/unit volume) stored in the DNA can be calculated as follows
- The radius of the helix is $r_h \sim 1$ nm, while the perpendicular distance between adjacent nucleobasis $d_n \sim 0.34$ nm
- The volume occupied by a codon (3 nucleobasis) is therefore $V_{\text{codon}} \cong 3 d_n \pi r_h^2 \cong 3.2 \text{ nm}^3$
- Each codon has 6 bits of information (64 combinations=2⁶), corresponding to 0.75 bytes
- The density of information is therefore $0.75 \text{ B}/(3.2 \text{ nm}^3) = 0.23 \text{ B/nm}^3 = 2.3 \times 10^5 \text{ TB/mm}^3$
- This is the maximum density of information that can be obtained by compactified DNA strands
- Exercise: check that this density of information is largest than that of present-day storage devices by several orders of magnitude

