

Introduction to molecular biology



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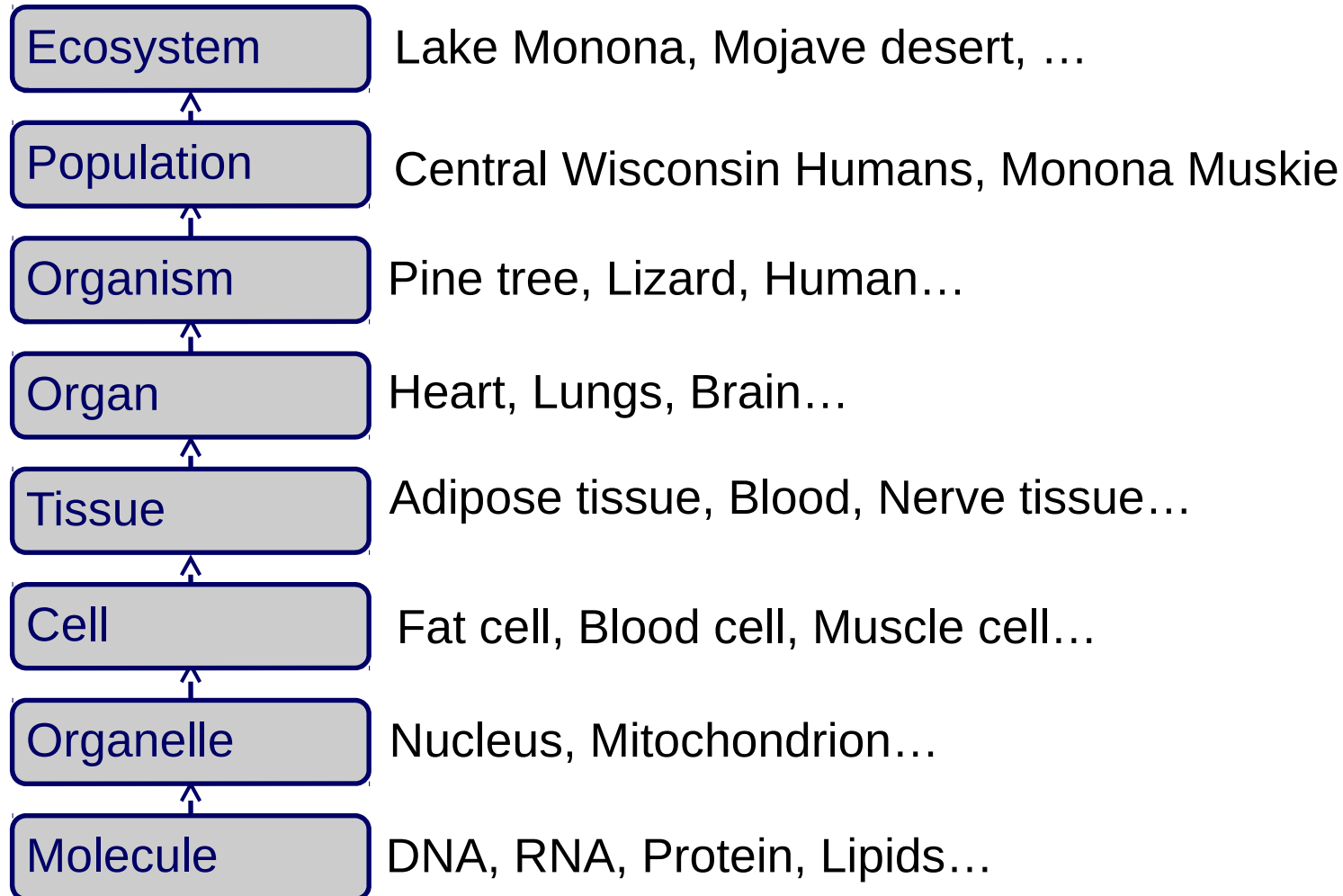
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Inspired by: Mark Craven (craven@biostat.wisc.edu), BMI/CS 576

www.biostat.wisc.edu/bmi576/

Biological hierarchy



DNA

- “blueprint” for an organism
- composed of small molecules called *nucleotides*
- four different nucleotides distinguished by the four *bases*: adenine (A), cytosine (C), guanine (G) and thymine (T)
- is a *polymer* : large molecule consisting of similar units (nucleotides in this case)
- nucleotides form a strand of DNA. It can be thought of as a string composed of the four letters: A, C, G, T

**ctgctggaccgggtgctaggaccctgactgcccggggccgggggtgcg
gggcccgctgag...**

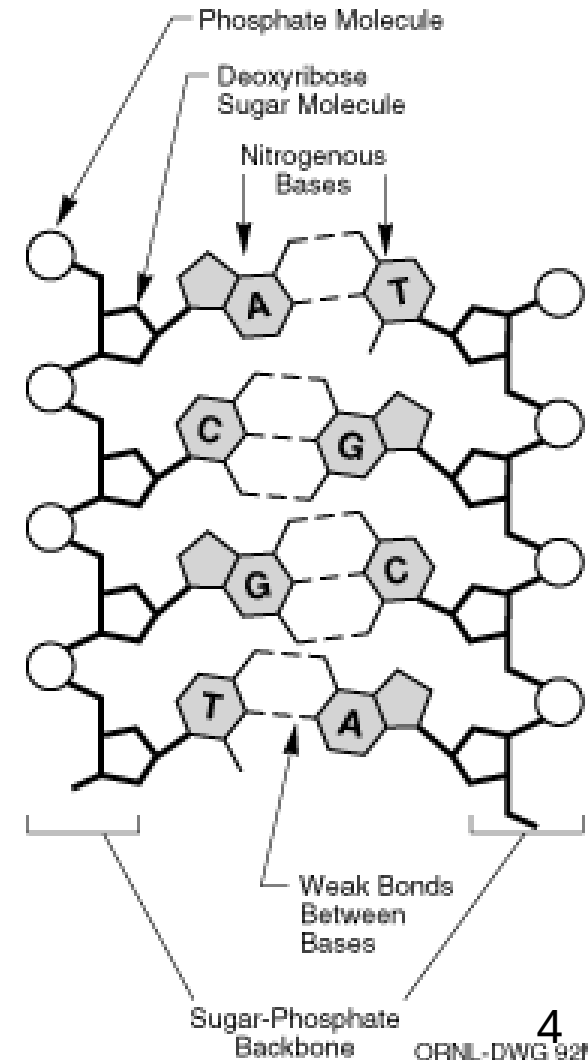
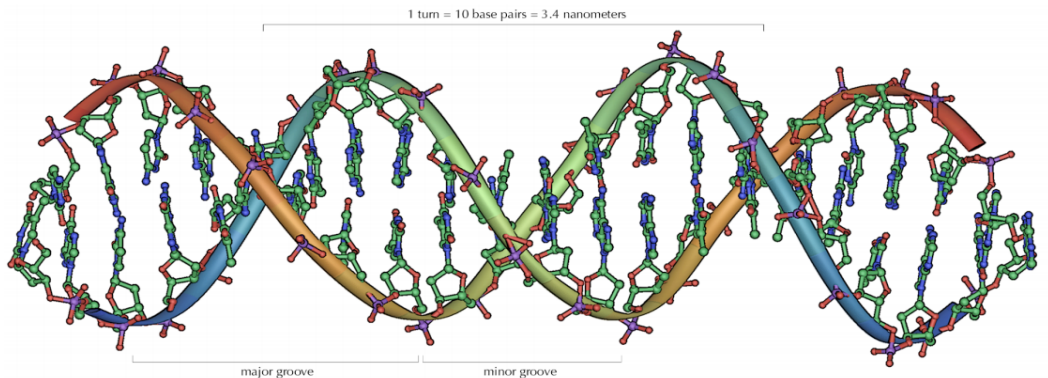
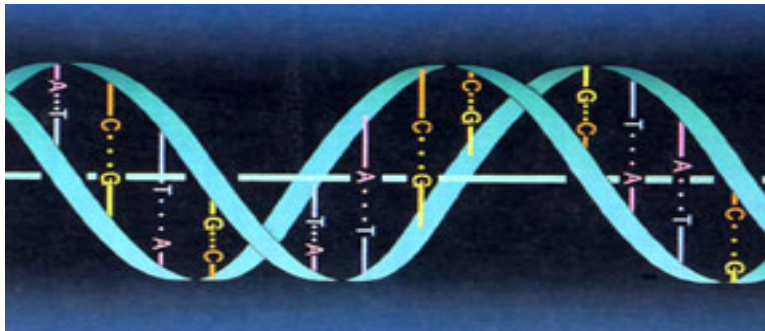
- DNA molecules usually consist of two strands arranged in the double helix

Watson-Crick base pairs

- in double-stranded DNA

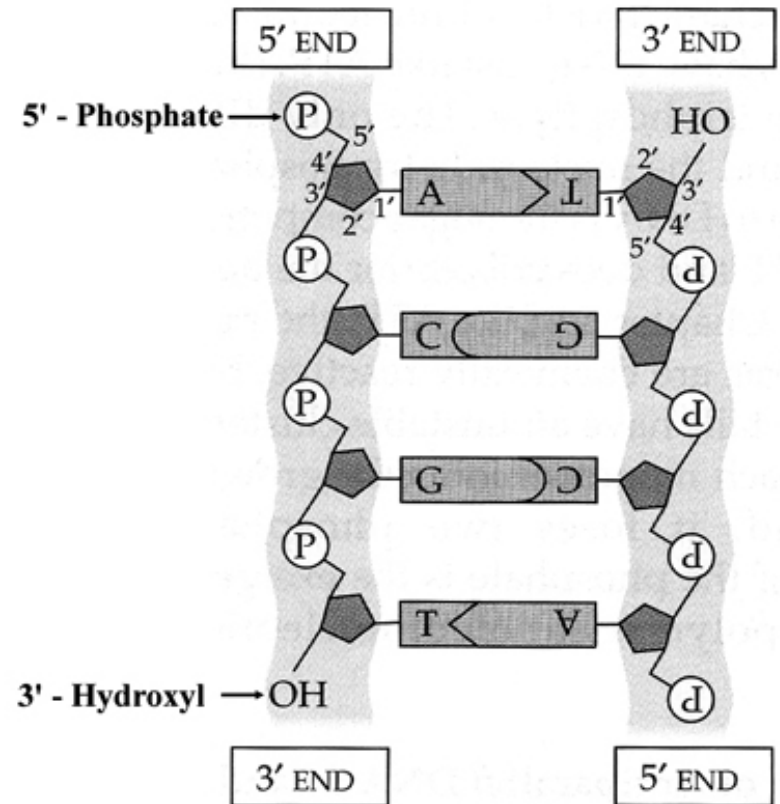
A always bonds to T

C always bonds to G



The double helix

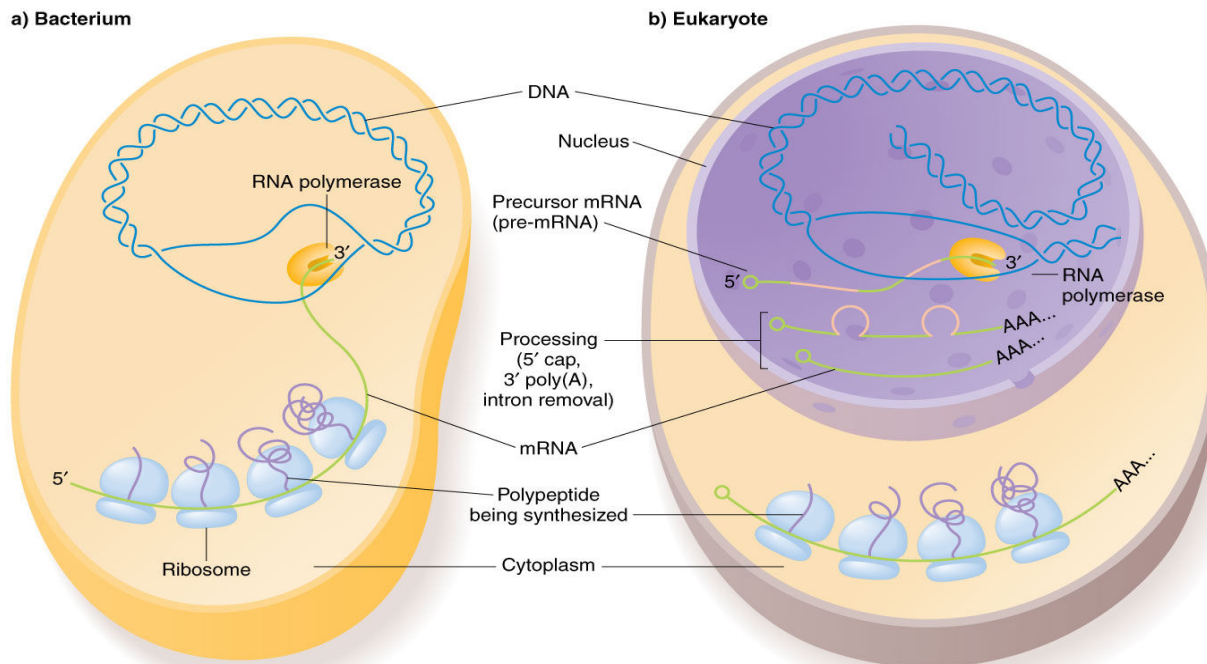
- each strand of DNA has a “direction”
- at one end, the terminal carbon atom in the backbone is the 5' carbon atom of the terminal sugar
- at the other end, the terminal carbon atom is the 3' carbon atom of the terminal sugar
- therefore we can talk about the 5' and the 3' ends of a DNA strand
- in a double helix, the strands are *antiparallel* (arrows drawn from the 5' end to the 3' end go in opposite directions)



http://hshgp.genome.washington.edu/teacher_resources/modules-view.htm

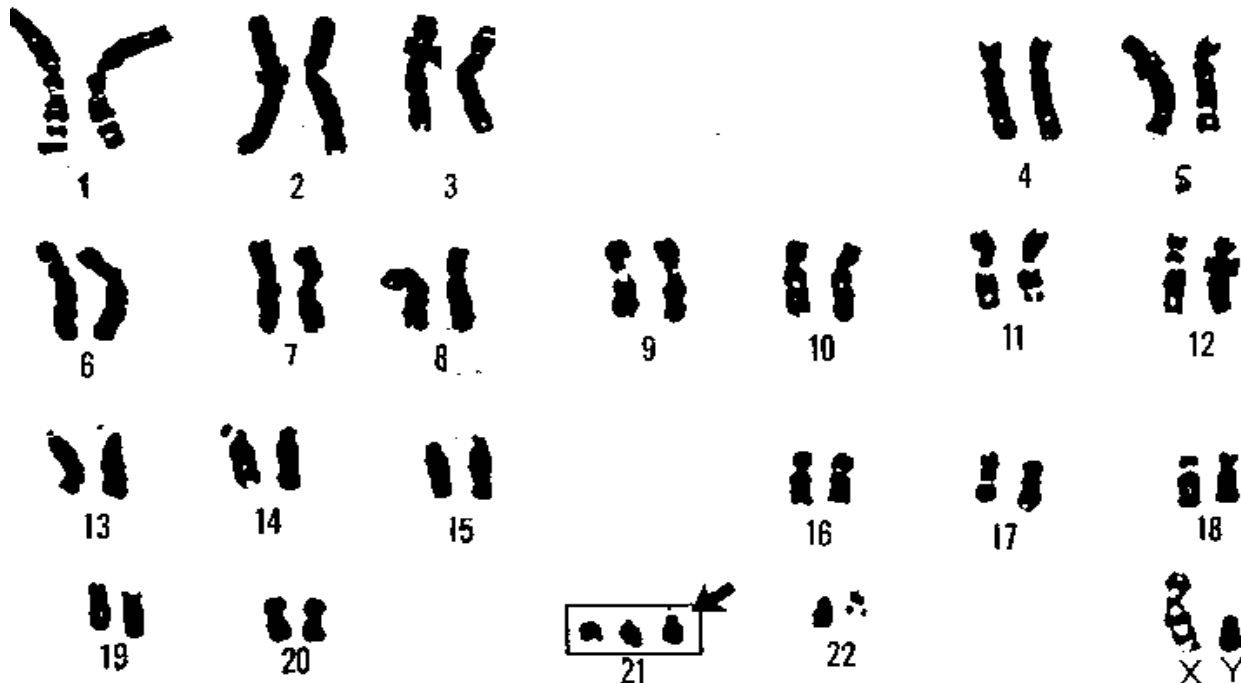
Chromosomes

- DNA is packaged into individual *chromosomes* (along with proteins)
- *prokaryotes* (single-celled organisms lacking nuclei) typically have a single circular chromosome
- *eukaryotes* (organisms with nuclei) have a species-specific number of linear chromosomes



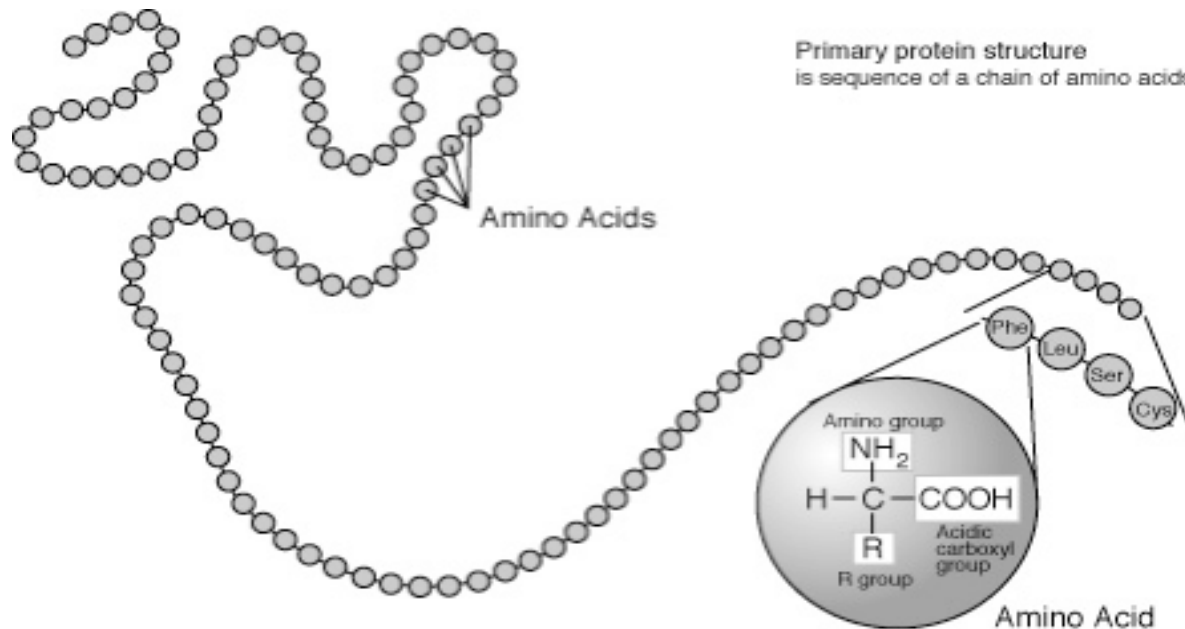
Genome

- the term *genome* refers to the complete complement of DNA for a given species
- the human genome consists of 46 chromosomes
- every cell (except sex cells and mature red blood cells) contains the complete genome of an organism



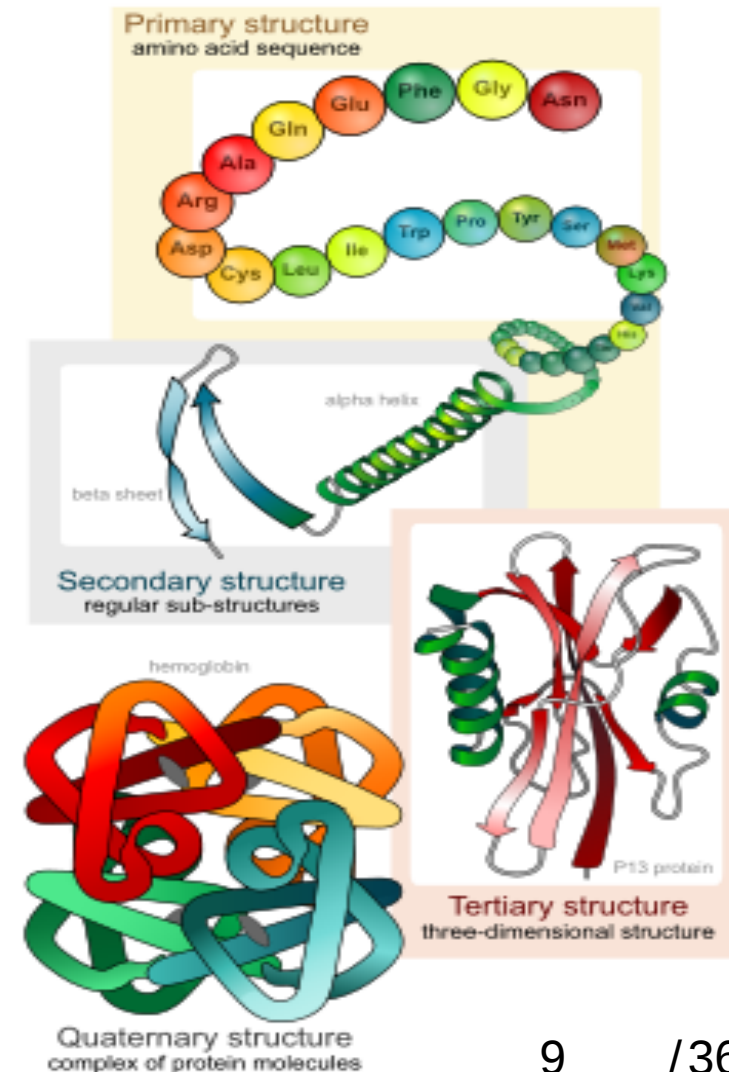
Proteins

- proteins are molecules composed of one or more *polypeptides*
- a polypeptide is a polymer composed of *amino acids*
- cells build their proteins from 20 different amino acids
- a polypeptide can be thought of as a string composed from a 20-character alphabet



Protein structure

- **Primary structure**
amino acid linear sequence of the polypeptide chain
- **Secondary structure**
highly regular local sub-structures (alpha helix, beta strand or beta sheets)
- **Tertiary structure**
three-dimensional structure of a protein molecule (given by hydrophilic and hydrophobic properties of molecules)
- **Quaternary structure**
three-dimensional structure of a multi-subunit protein and how the subunits fit together
- **Protein structure**



The alphabet of proteins

Alanine	Ala	A
Arginine	Arg	R
Aspartic Acid	Asp	D
Asparagine	Asn	N
Cysteine	Cys	C
Glutamic Acid	Glu	E
Glutamine	Gln	Q
Glycine	Gly	G
Histidine	His	H
Isoleucine	Ile	I
Leucine	Leu	L
Lysine	Lys	K
Methionine	Met	M
Phenylalanine	Phe	F
Proline	Pro	P
Serine	Ser	S
Threonine	Thr	T
Tryptophan	Trp	W
Tyrosine	Tyr	Y
Valine	Val	V

Protein functions

- structural support
- storage of amino acids
- transport of other substances
- coordination of an organism's activities
- response of cell to chemical stimuli
- movement
- protection against disease
- selective acceleration of chemical reactions
- **ATP synthase - powering your cells**

DNA variations - mutations

ATG ACC CAG CAG CCA ATG AAA
Met Thr Gln Gln Pro Met Lys

Normal sequence
(Reading frame marked with spaces)

▼
ATG **CCC** CAG CAG CCA ATG AAA
Met **Pro** Gln Gln Pro Met Lys

Missense substitution
(threonine replaced by proline)

▼
ATG ACC **TAG** CAG CCA ATG AAA
Met Thr **STOP** - - - -

Nonsense substitution
(altered DNA sequence prematurely signals the cell to stop building a protein)

▼
ATG **ACA** CAG CAG CCA ATG AAA
Met Thr Gln Gln Pro Met Lys

Silent substitution
(threonine encoded by different codon)

▼▼▼
ATG --- CAG CAG CCA ATG AAA
Met - Gln Gln Pro Met Lys

Deletion without shift of a reading frame
(deletes one amino acid)

▼
ATG **-CCC** AGC AGC CAA TGA AA
Met **Pro Ser Ser Gln STOP** -

Deletion with shift of a reading frame
(causes changes in following amino acid sequence and causes premature termination)

▼▼▼ ▼▼▼ ▼▼▼
ATG ACC CAG CAG **CAG CAG CAG** CCA ATG AAA
Met Thr Gln Gln **Gln Gln Gln** Pro Met Lys

Expansion of a trinucleotide repeats

DNA variations - polymorphism

▼
ATGCC**C**AGCAGCCAAT
▼
ATGCC**T**AGCAGCCAAT

Single nucleotide polymorphism

Chromosome A

Chromosome B

Three possible genotypes

▼ ▼ ▼ ▼ ▼ ▼ ▼
ATGCC**CCACACACACACAC**AGAAA
▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼
ATGCC**CCACACACACACACACACACAC**AGAAA
▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼
ATGCC**CCACACACACACACACACACACACACACACAC**AGAAA

Short tandem repeat

Allele A

Allele B

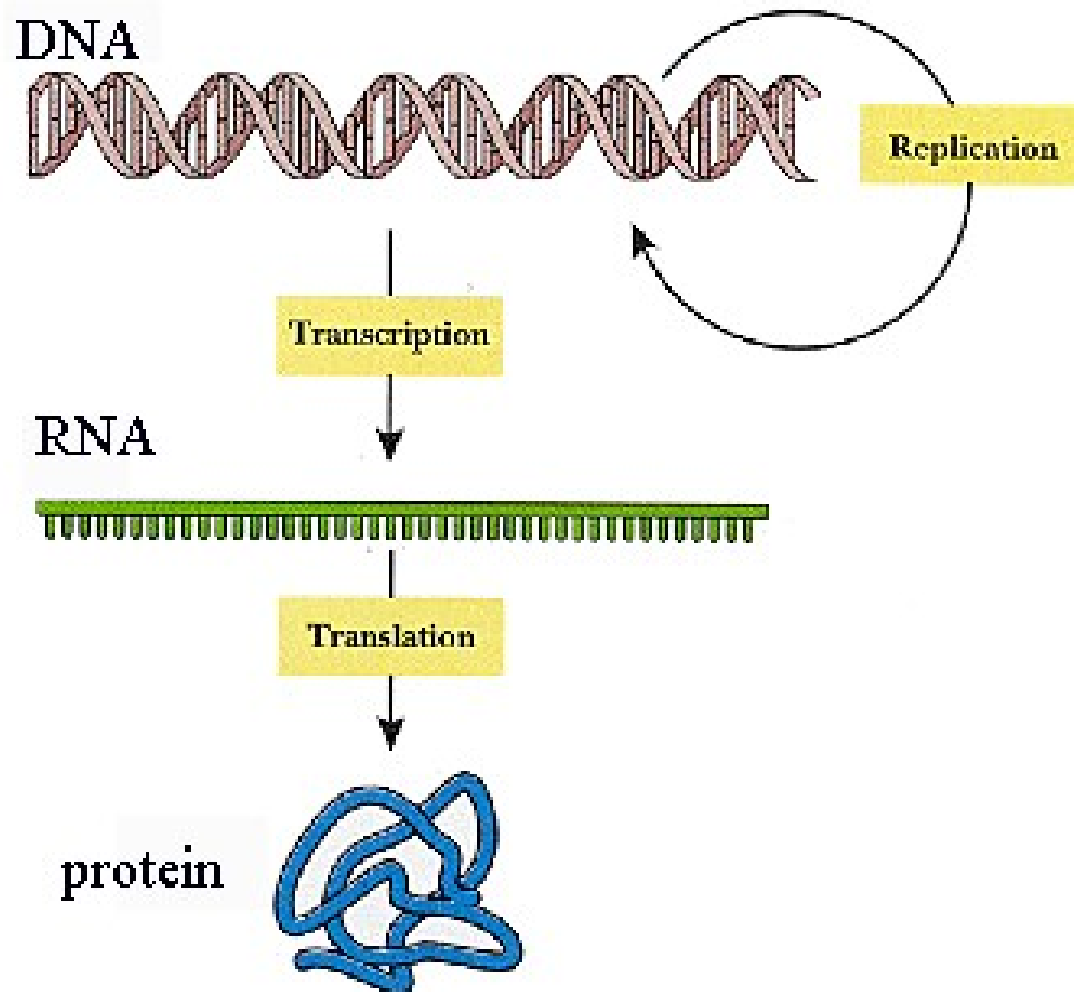
Allele C

Many possible genotypes



How to construct a protein from a DNA?

The central dogma



DNA content

- Coding DNA

genes (gene is a sequence of DNA bases that carries the information required for constructing a particular protein – encodes a protein)

the human genome comprises ~ 25,000 protein-coding genes

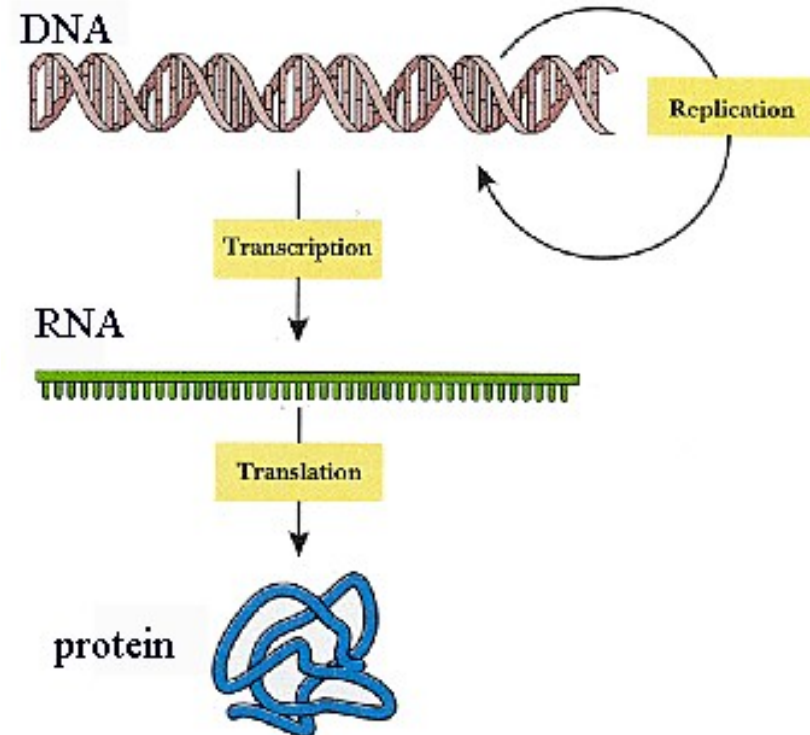
- Non-coding DNA

transcribed into functional non-coding RNA molecules, give rise to RNA transcripts

- *ribosomal RNA* (rRNA), which includes major constituents of ribosomes
- *transfer RNAs* (tRNAs), which carry amino acids to ribosomes
- *micro RNAs* (miRNAs), which play an important regulatory role in various plants and animals
- etc.

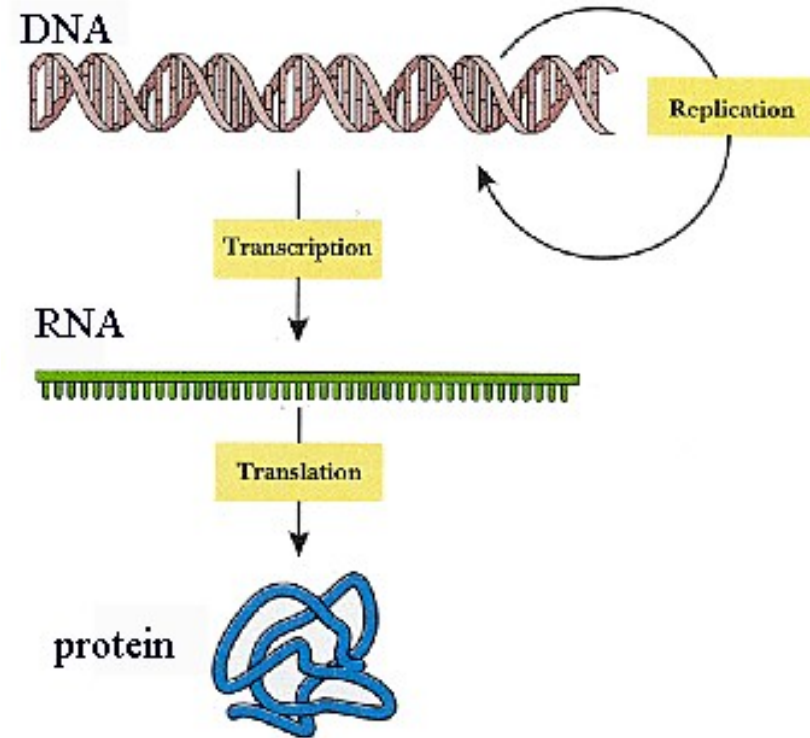
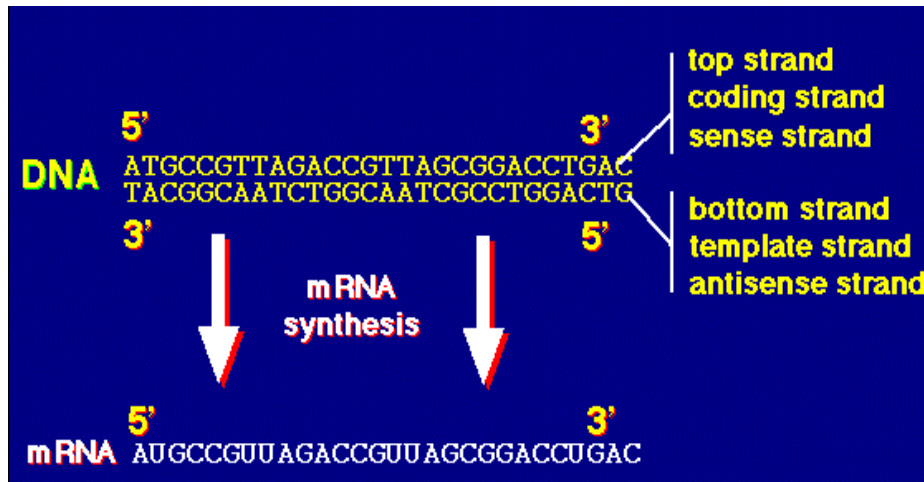
- Junk DNA

no known biological function

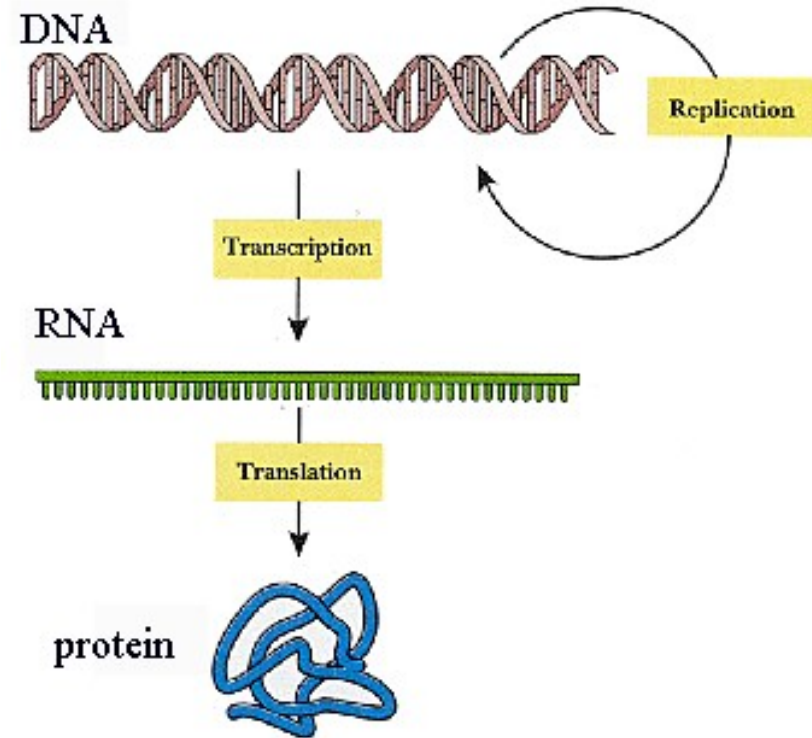
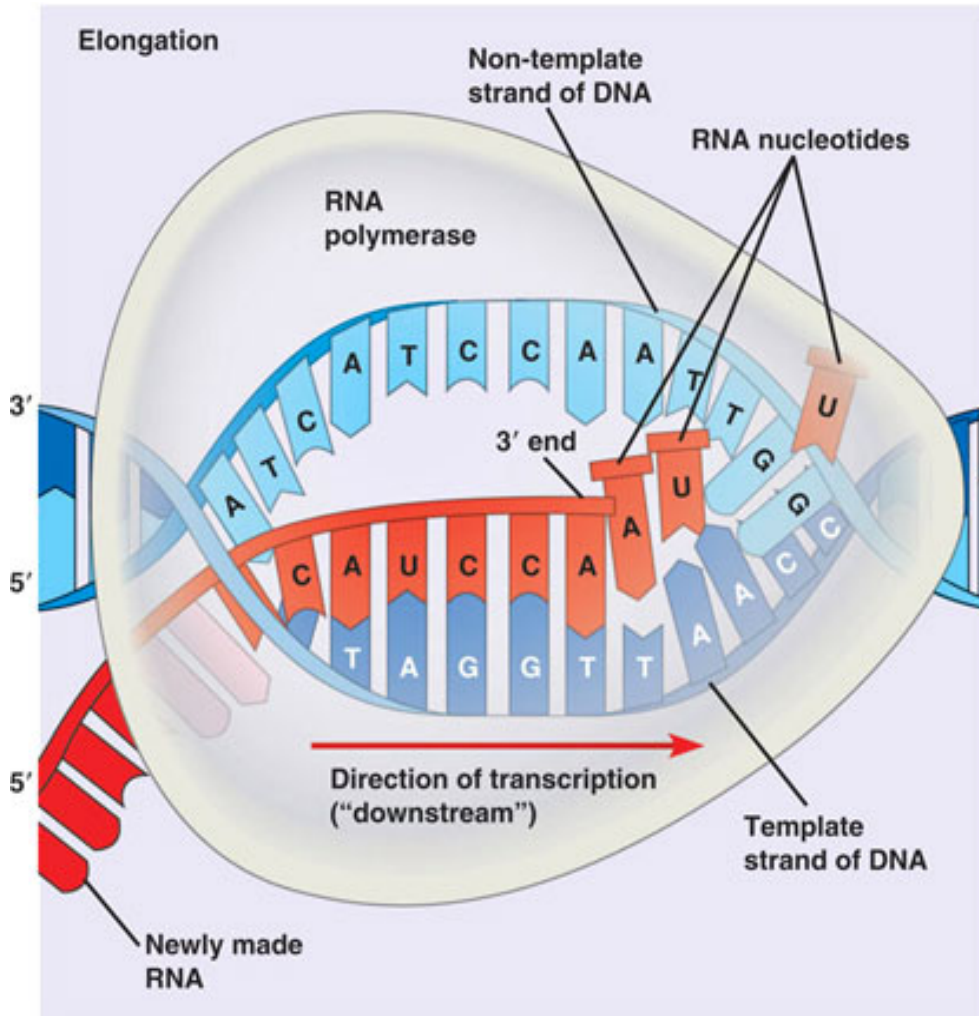


Transcription

- Transcription is the process of creating a complementary RNA copy of a DNA subsequence
- *RNA polymerase* is the enzyme that builds an RNA strand from a gene
- RNA that is transcribed from a gene is called *messenger RNA* (mRNA)

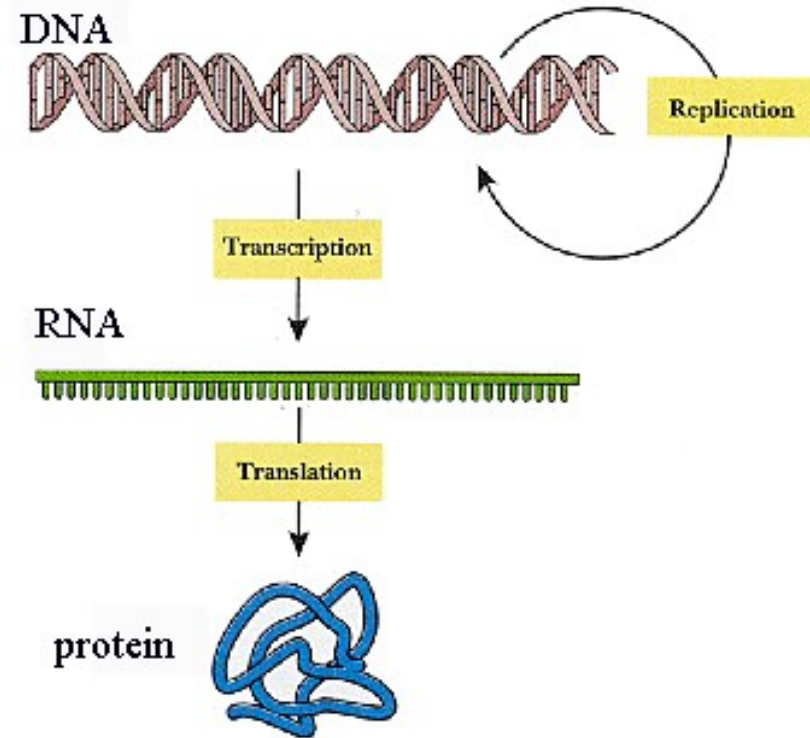


Transcription



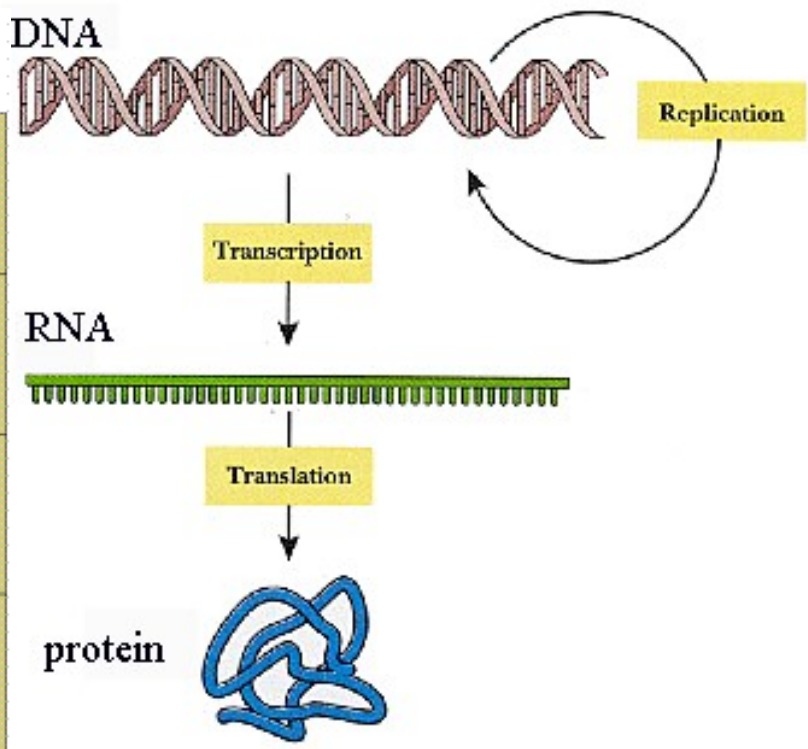
RNA

- RNA is like DNA except:
 - backbone is a little different
 - often single stranded
 - the base uracil (U) is used in place of thymine (T)
 - a strand of RNA can be thought of as a string composed of the four letters: A, C, G, U
- triplets of bases encoding amino acids are called *codons*
- the grouping of codons is called the *reading frame*

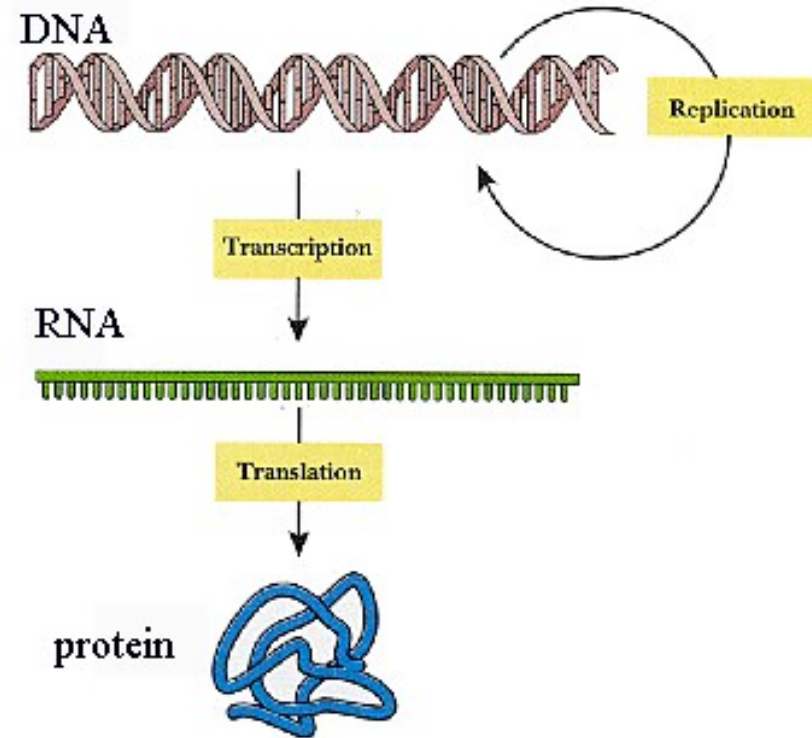
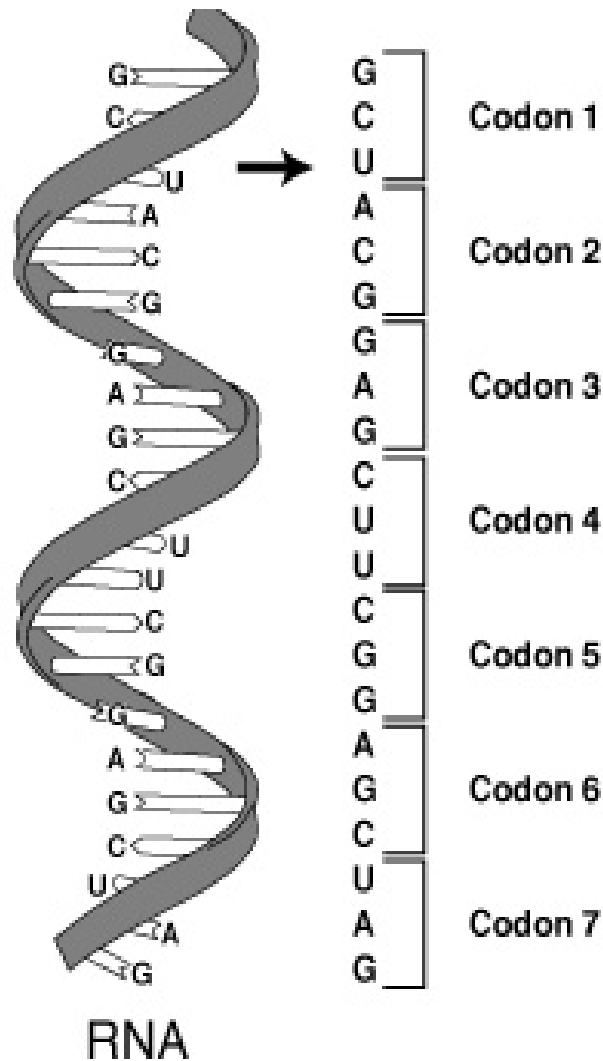


Codons

		Second letter				
		U	C	A	G	
First letter	U	UUU	UCU UCC UCA UCG Serine	UAU	UGU UGC Cysteine	U
		UUC		UAC		Tyrosine
		UUA UUG		UAA UAG	Stop codon Stop codon	UGA UGG
	C	CUU	CCU CCC CCA CCG Proline	CAU	CGU CGC CGA CGG Arginine	U
CUC		CAC		Histidine		C
CUA CUG		CAA CAG		Glutamine	U C A G	
A	AUU	ACU ACC ACA ACG Threonine	AAU	AGU AGC Serine	U	
	AUC		AAC		Asparagine	C
	AUA		AAA AAG	Lysine	AGA AGG	A G
	AUG	GAU GAC	Aspartic acid	GGU GGC GGA GGG Glycine	U C A G	
G	GUU	GCU GCC GCA GCG Alanine	GAA GAG	Glu Glu Glutamic acid	U	
	GUC		Valine		G	
	GUA GUG		G			

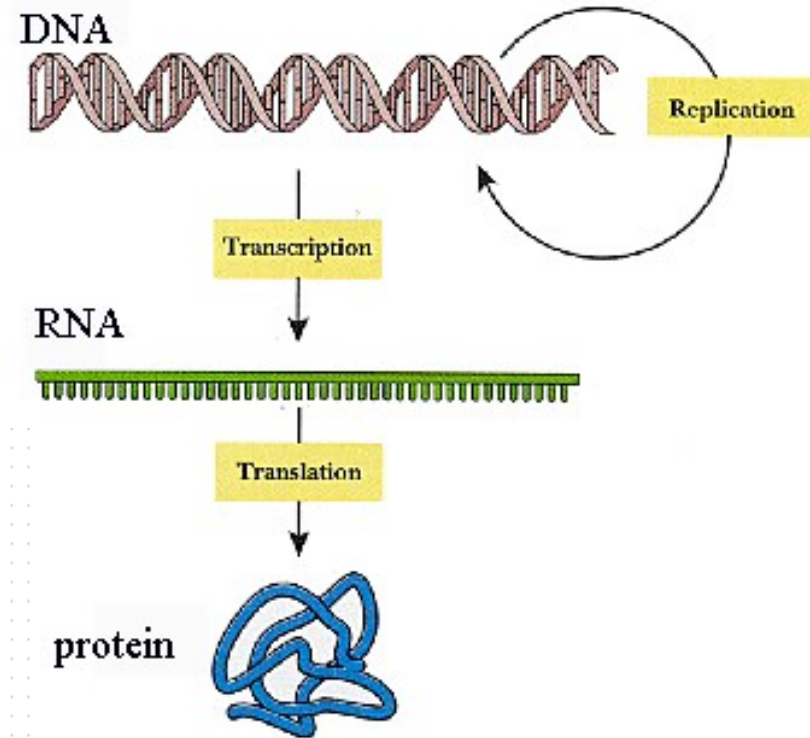
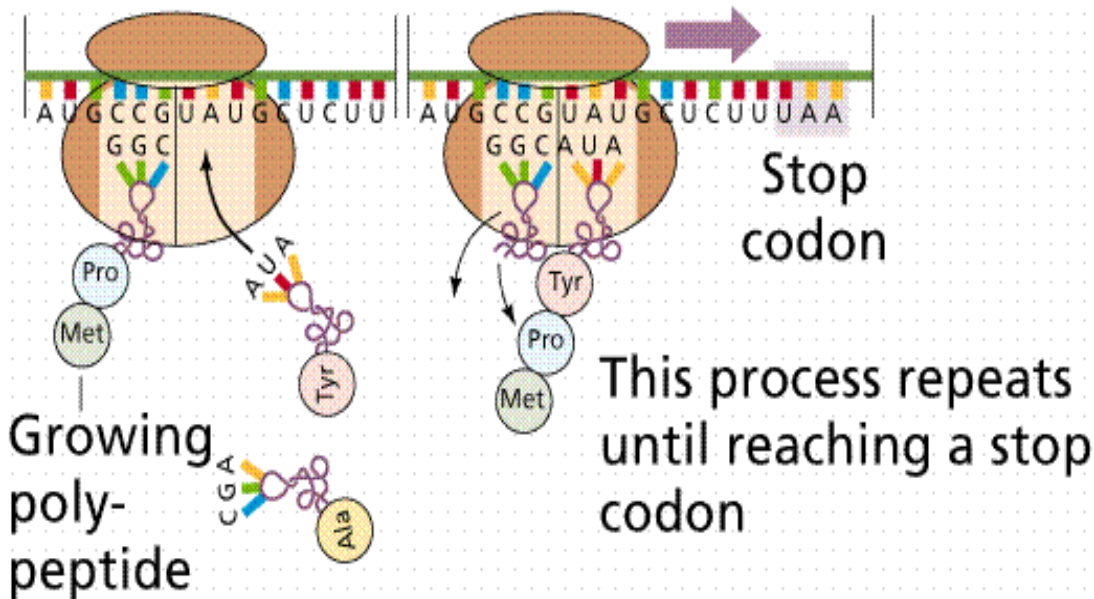


Codons and reading frames



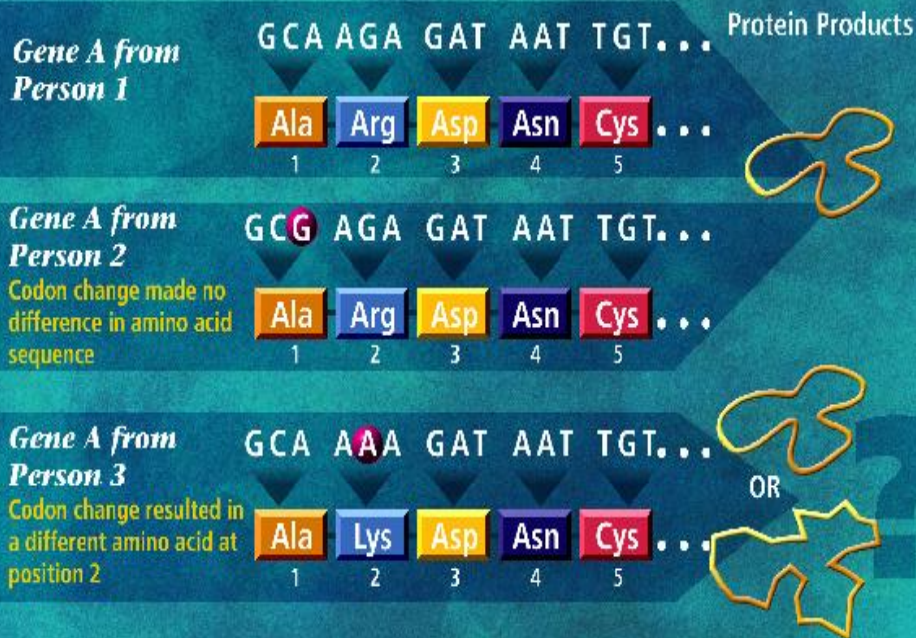
Translation

- *ribosomes* are the machines that synthesize proteins from mRNA
- translation begins with the *start codon* and ends with the *stop codon*

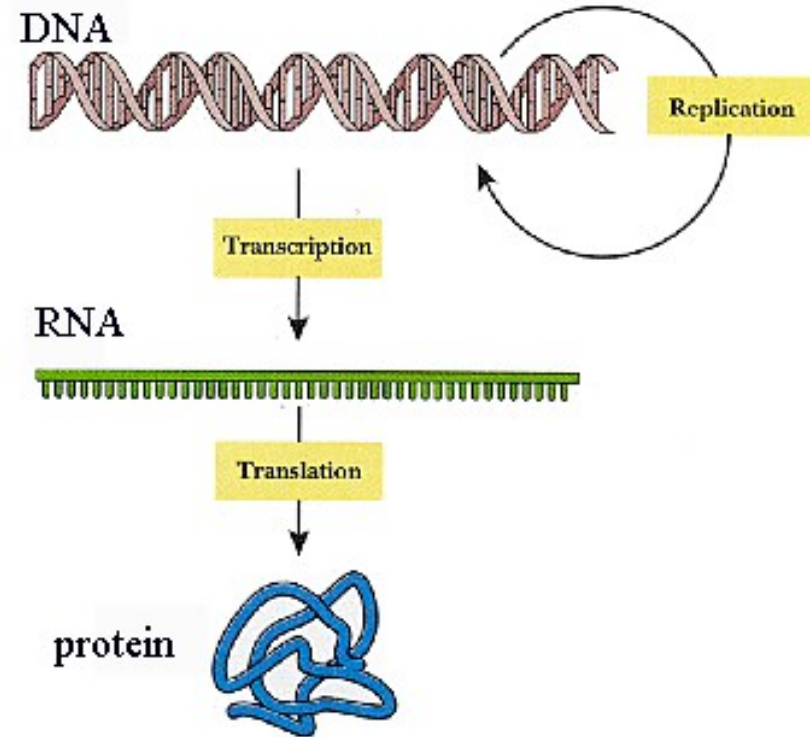


Translation

DNA Sequence Variation in a Gene Can Change the Protein Produced by the Genetic Code

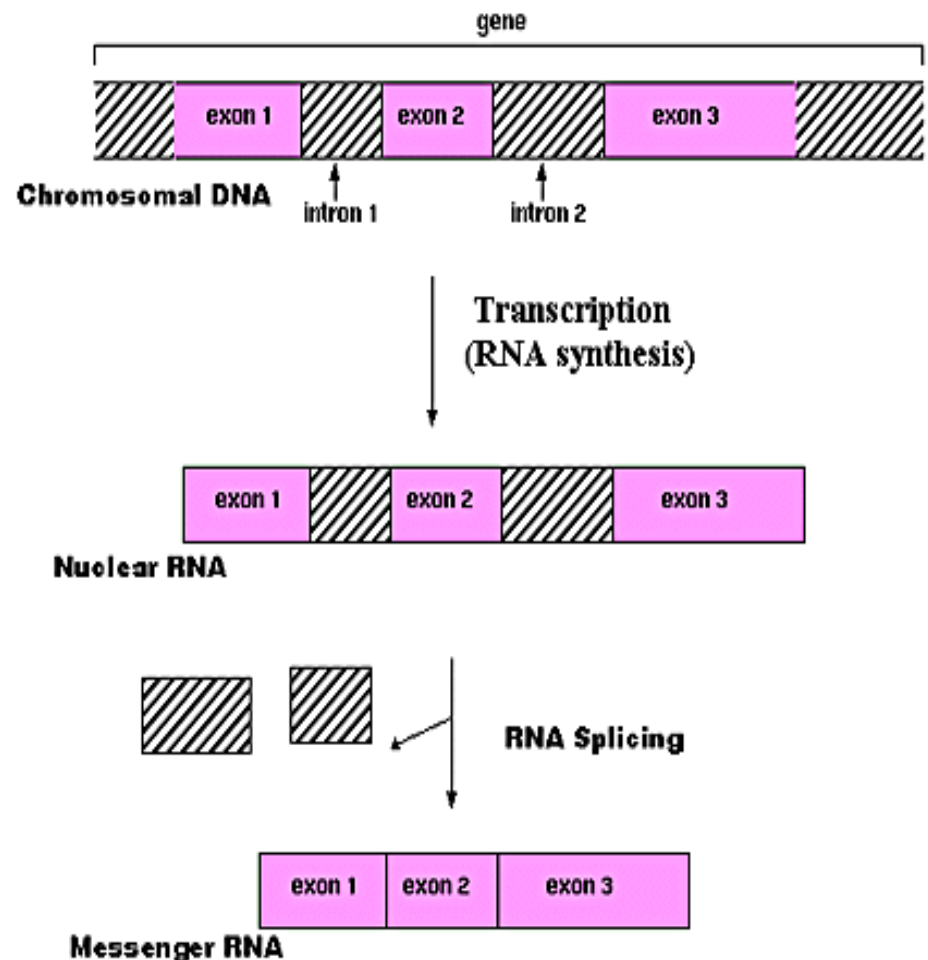


Y-GA 99-649

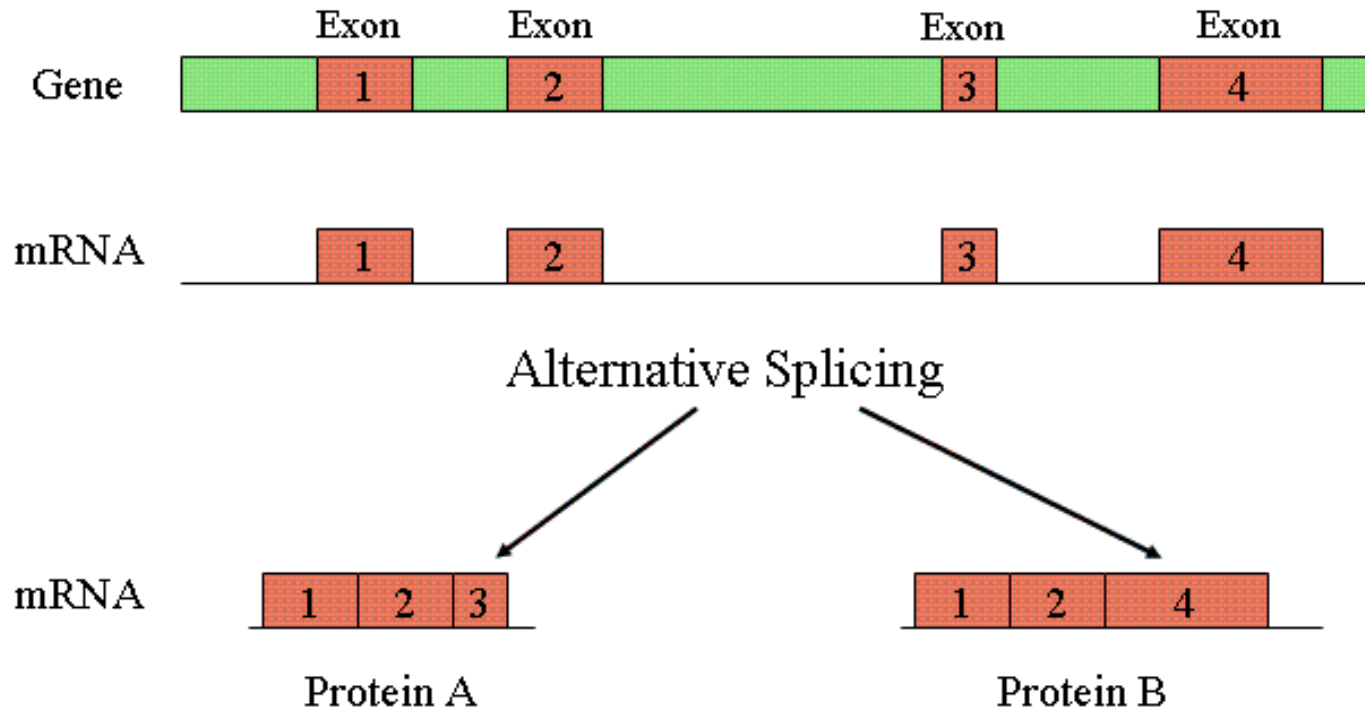


RNA processing in eukaryotes

- *eukaryotes* are organisms that have enclosed nuclei in their cells
- in many eukaryotes, genes/mRNAs consist of alternating *exon/intron* segments
- exons are the coding parts
- introns are spliced out before translation

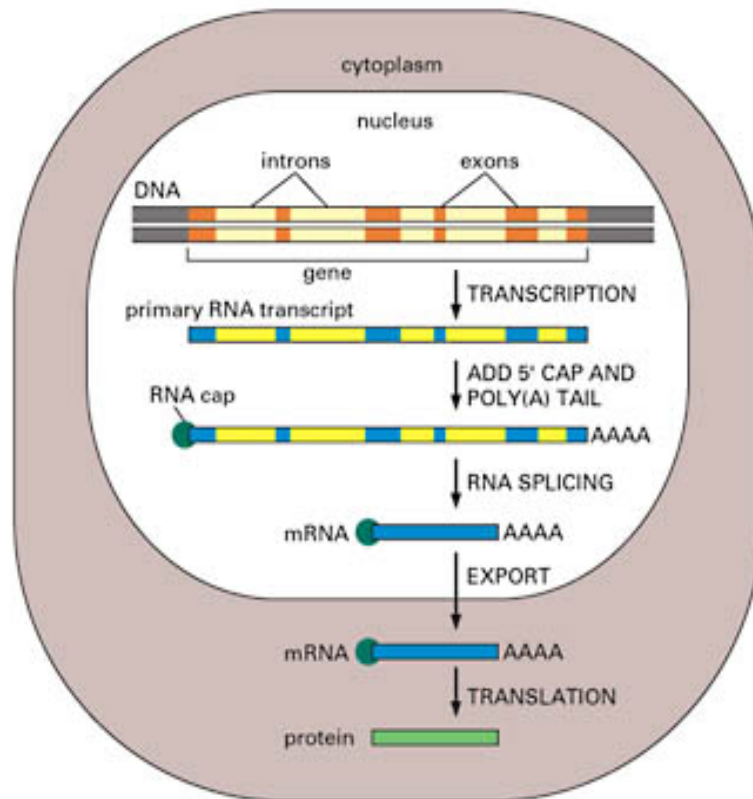


Alternative splicing

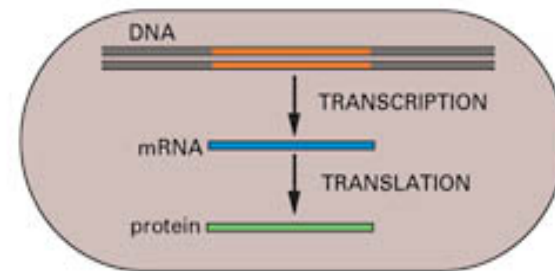


Protein synthesis in eukaryotes vs. prokaryotes

(A) EUCARYOTES



(B) PROCARYOTES



From DNA to protein

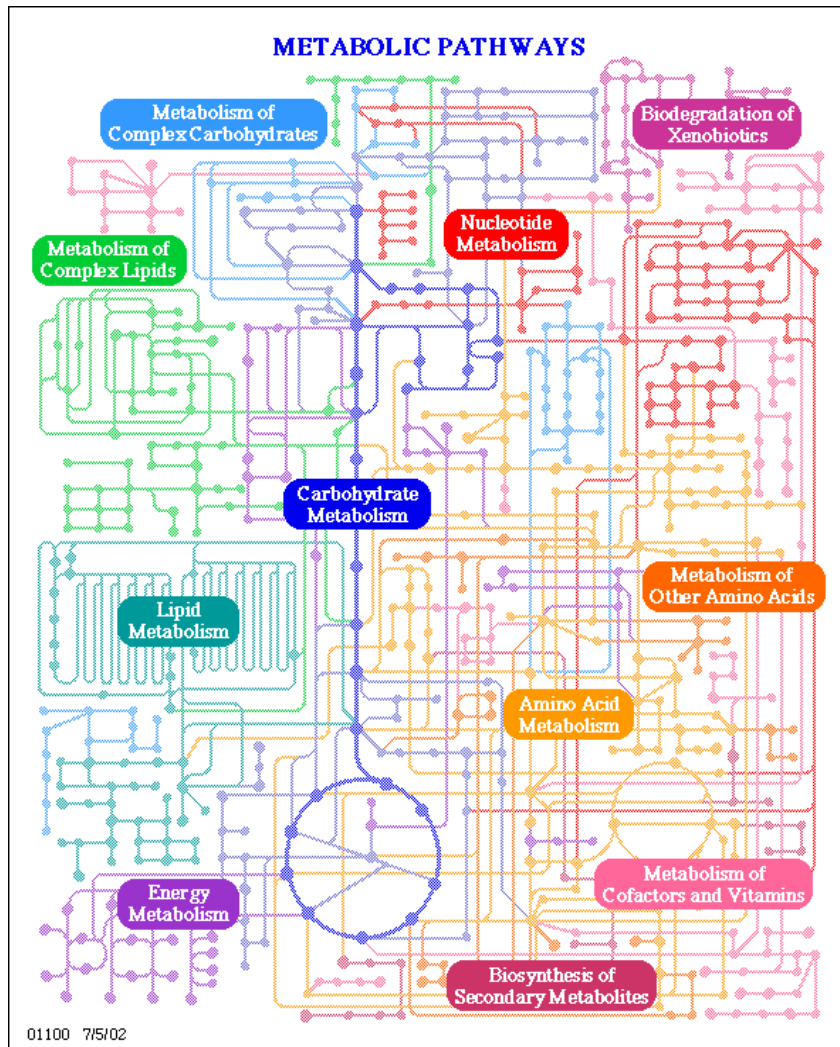


All cells in an organism have the same genomic data. How is possible to create different kind of cells and tissue?

The dynamics of cells

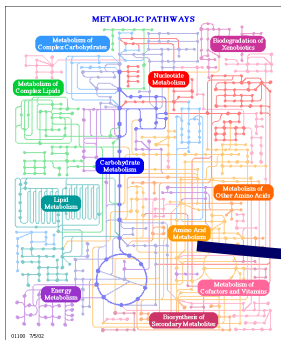
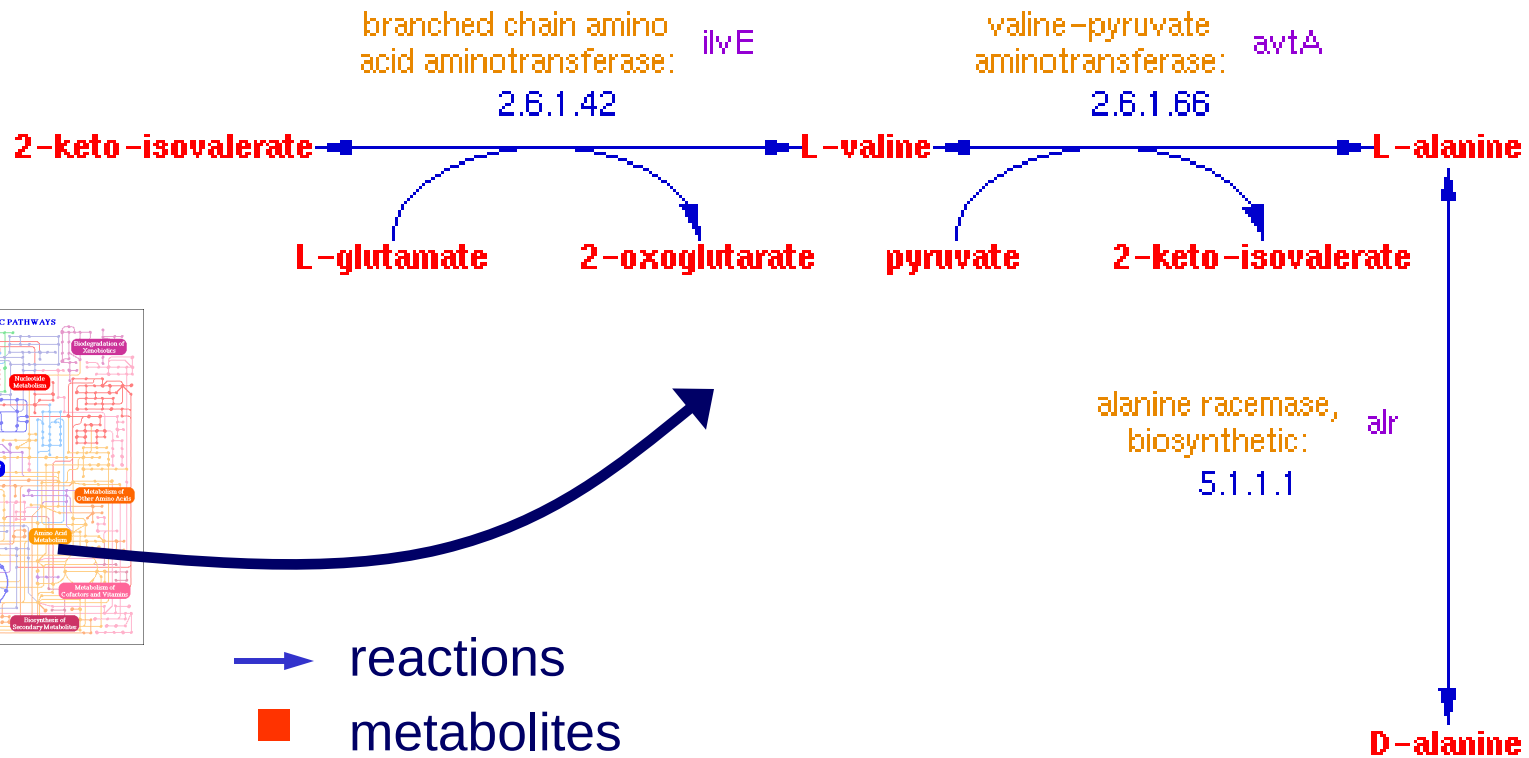
- all cells in an organism have the same genomic data, but the genes expressed in each vary according to cell type, time, and environmental factors
- there are networks of interactions among various biochemical entities in a cell (DNA, RNA, protein, small molecules) that carry out processes such as
 - metabolism
 - intra-cellular and inter-cellular signaling
 - regulation of gene expression

E. coli metabolic pathway map



- circles represent metabolites (small molecules)
- lines represent reactions

The metabolic pathway for synthesizing the amino acid alanine



- reactions
- metabolites
- enzymes (proteins that catalyze reactions)
- genes encoding the enzymes

Operon

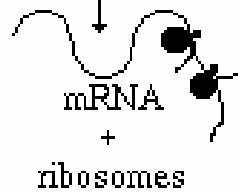
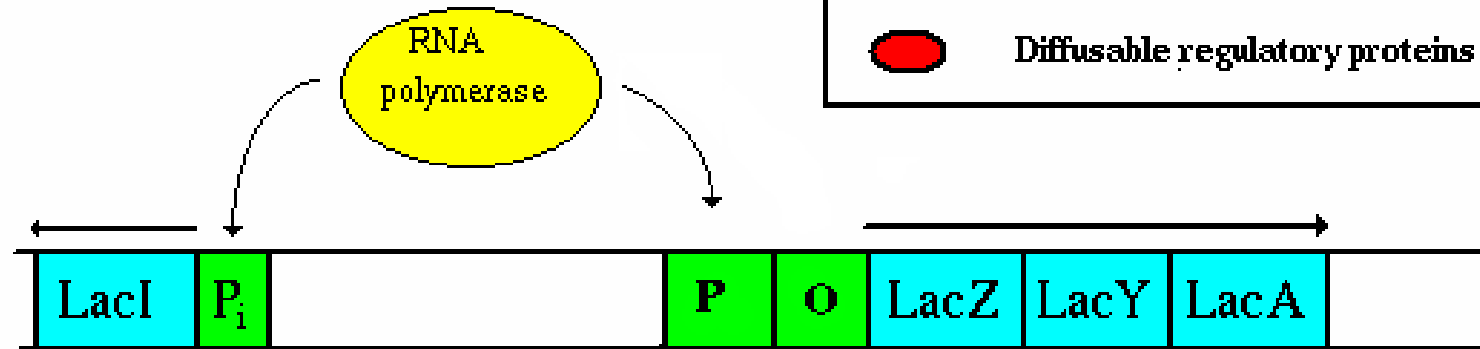
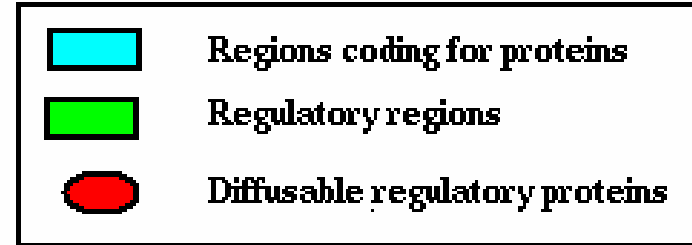
- Operon is a part of the DNA molecule of the prokaryotic organism that contains two or more genes, but operates as a single transcriptional unit.
- The genes in an operon are transcribed together (and often regulated together).

Gene expression

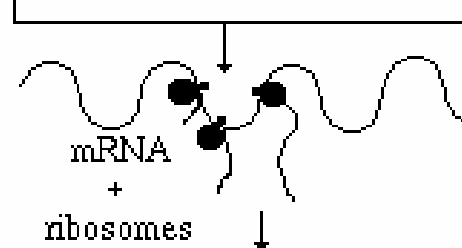
Lac operon

Gene regulation example: the lac operon

THE LAC OPERON

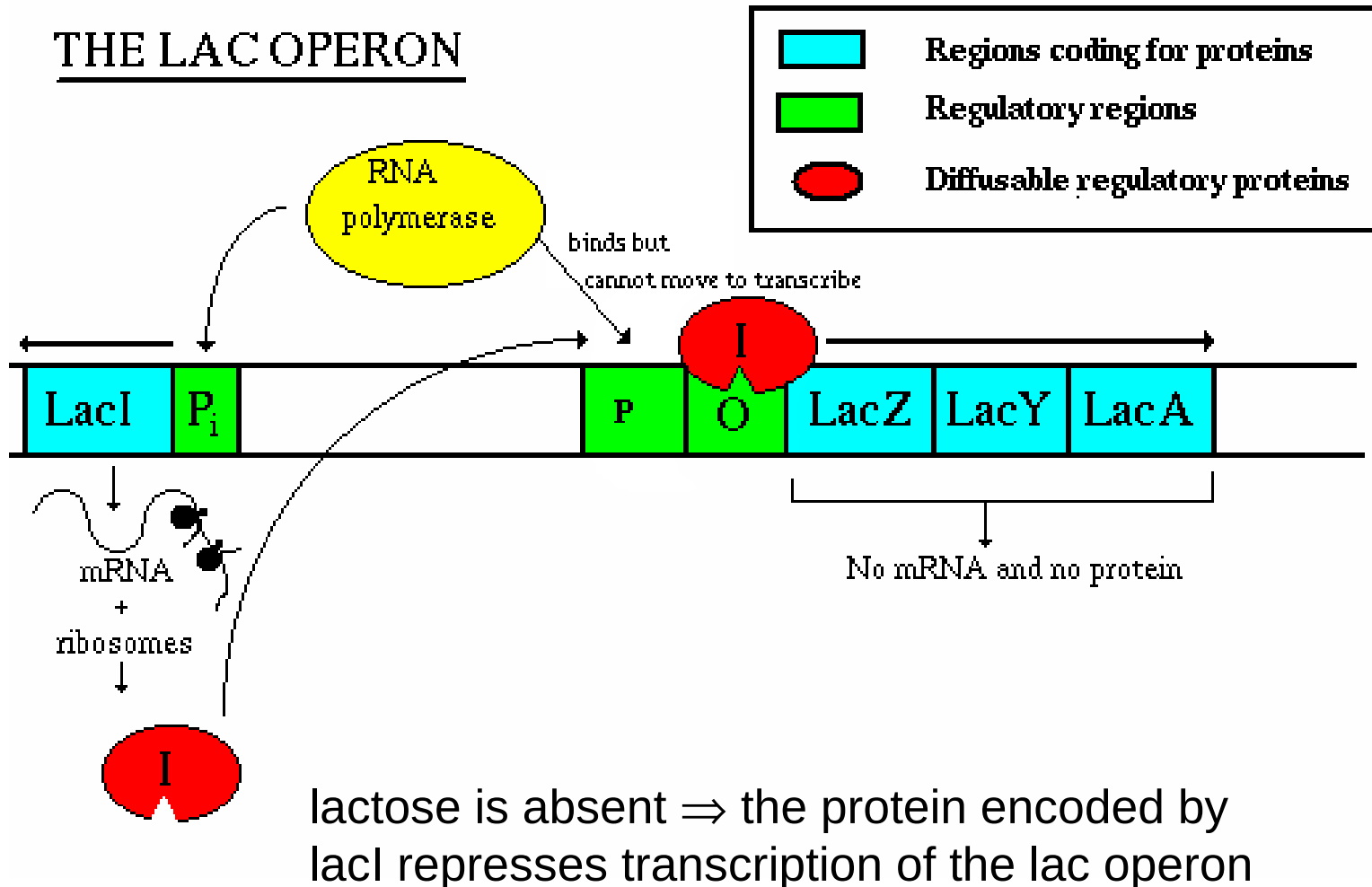


this protein regulates the transcription of LacZ, LacY, LacA

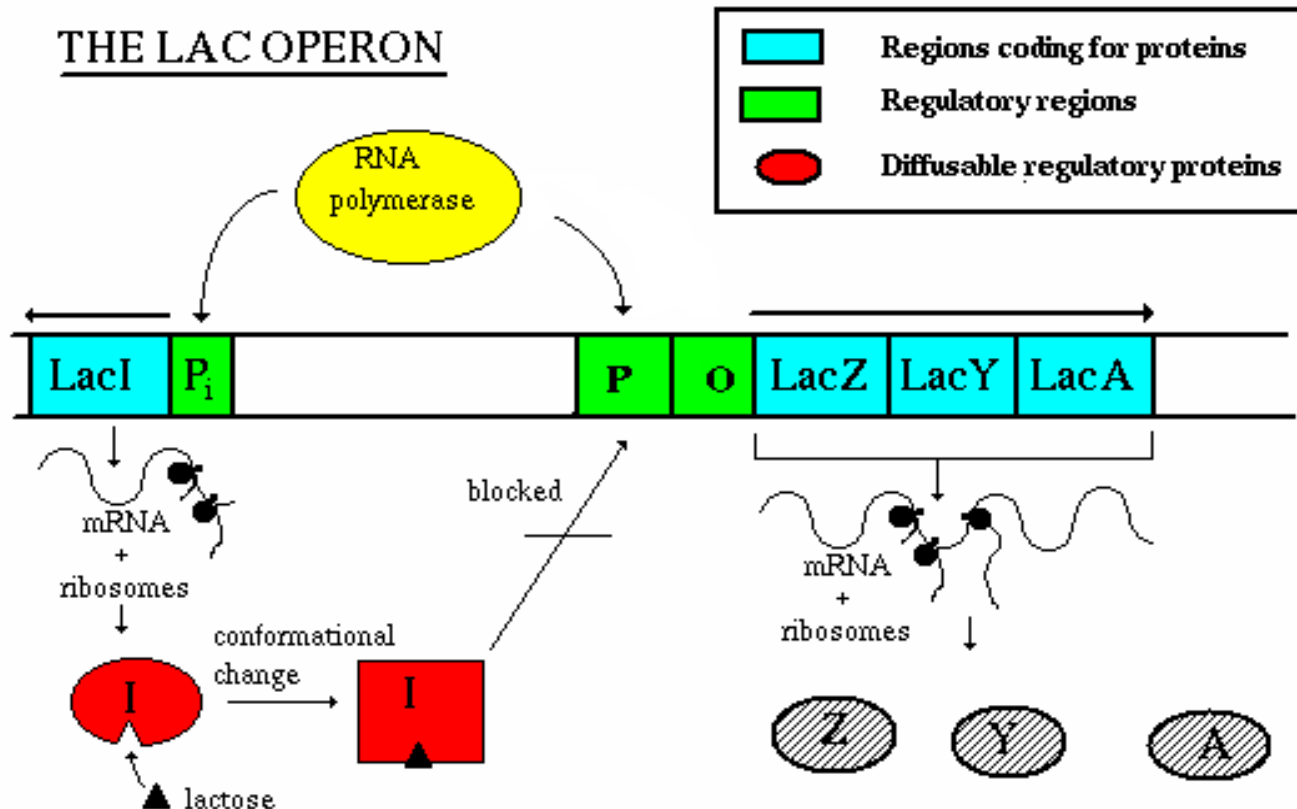


these proteins metabolize lactose

Gene regulation example: the lac operon



Gene regulation example: the lac operon



lactose is present \Rightarrow it binds to the protein encoded by lacI changing its shape; in this state, the protein doesn't bind upstream of the lac operon; therefore the lac operon can be transcribed

Gene regulation example: the lac operon

- this example provides a simple illustration of how a cell can regulate (turn on/off) certain genes in response to the state of its environment
- an *operon* is a sequence of genes transcribed as a unit
- the lac operon is involved in metabolizing lactose
- it is “turned on” when lactose is present in the cell
- the lac operon is regulated at the transcription level

- the depiction here is incomplete; for example, the level of glucose in the cell also influences transcription of the lac operon



Questions