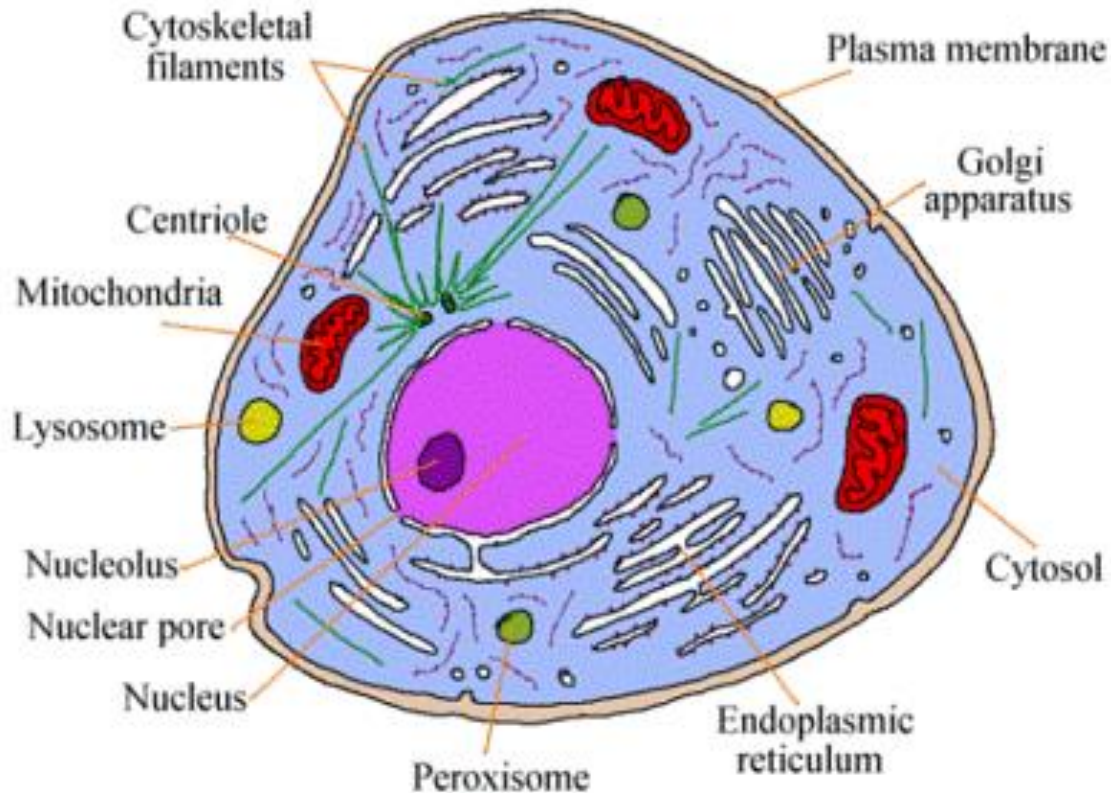
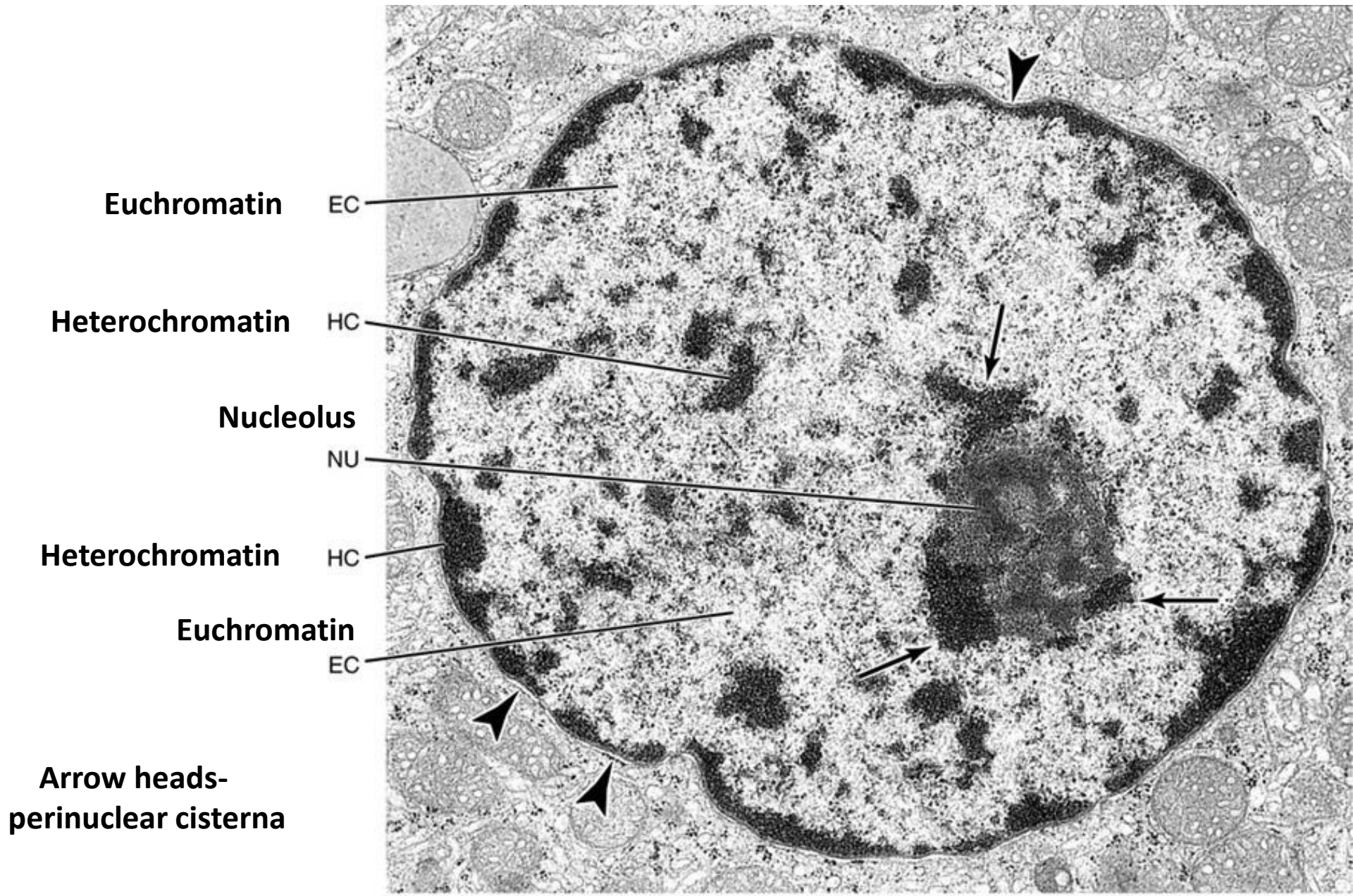


NUCLEUS - the largest organelle of the cell

The main function of the cell nucleus is to control of gene expression and mediate the replication of DNA during the cell cycle.

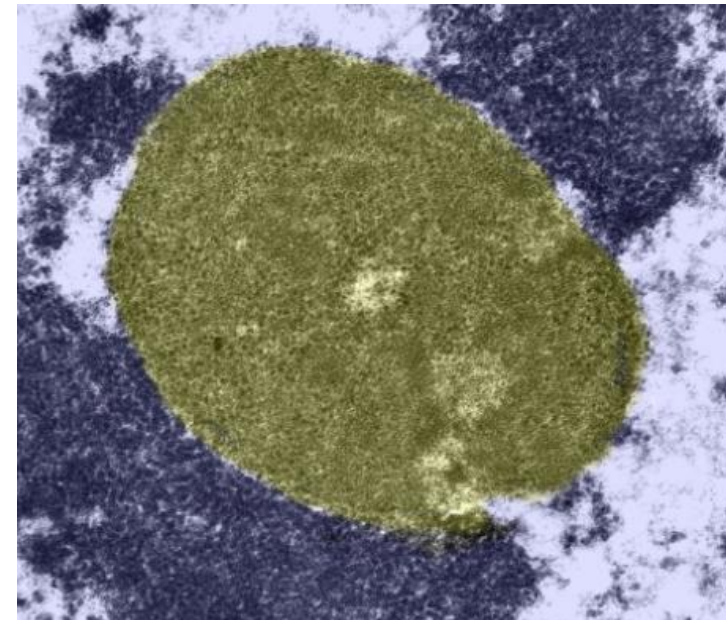
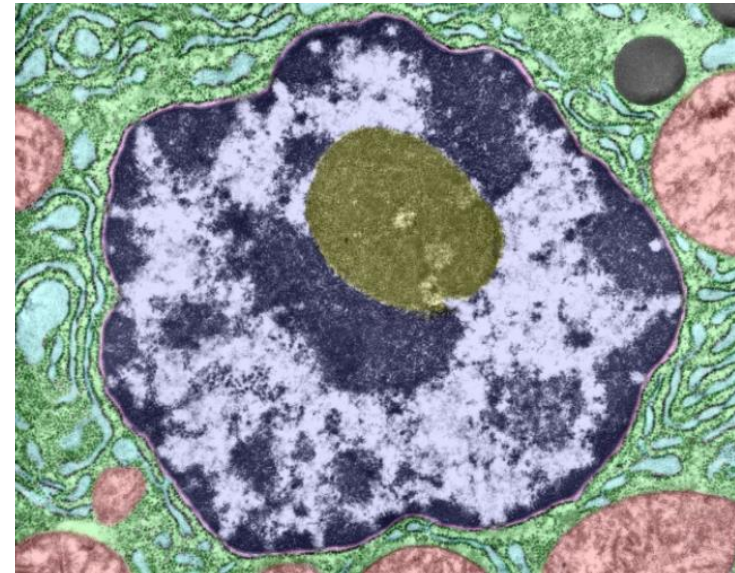


Cell nucleus in electron microscope



Nucleolus

- There may be 1 to 5 nucleoli in the nucleus
- Nucleolus contains genes necessary for ribosome production
- these genes come from chromosome 13, 14, 15, 21 and 22 (NOR)

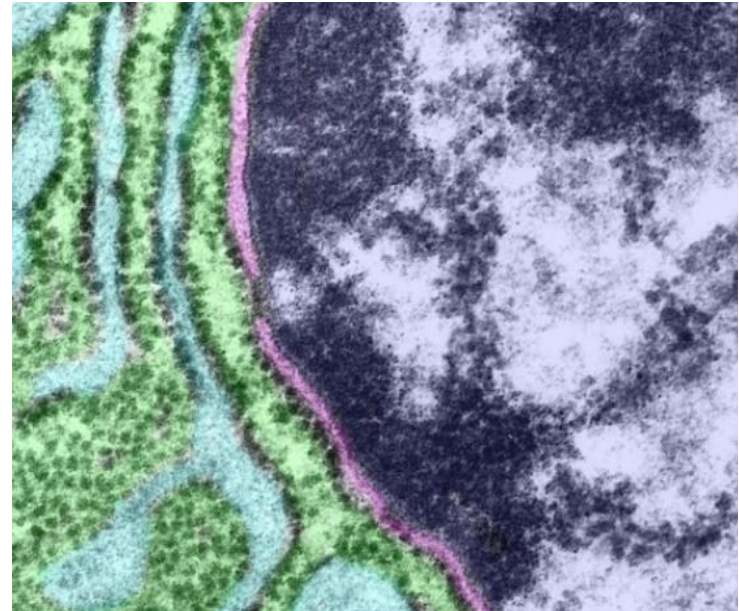
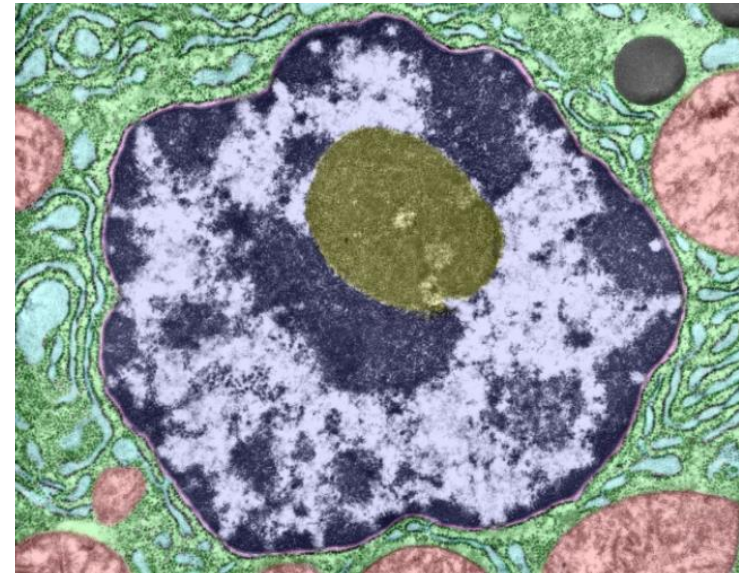


Nuclear envelope

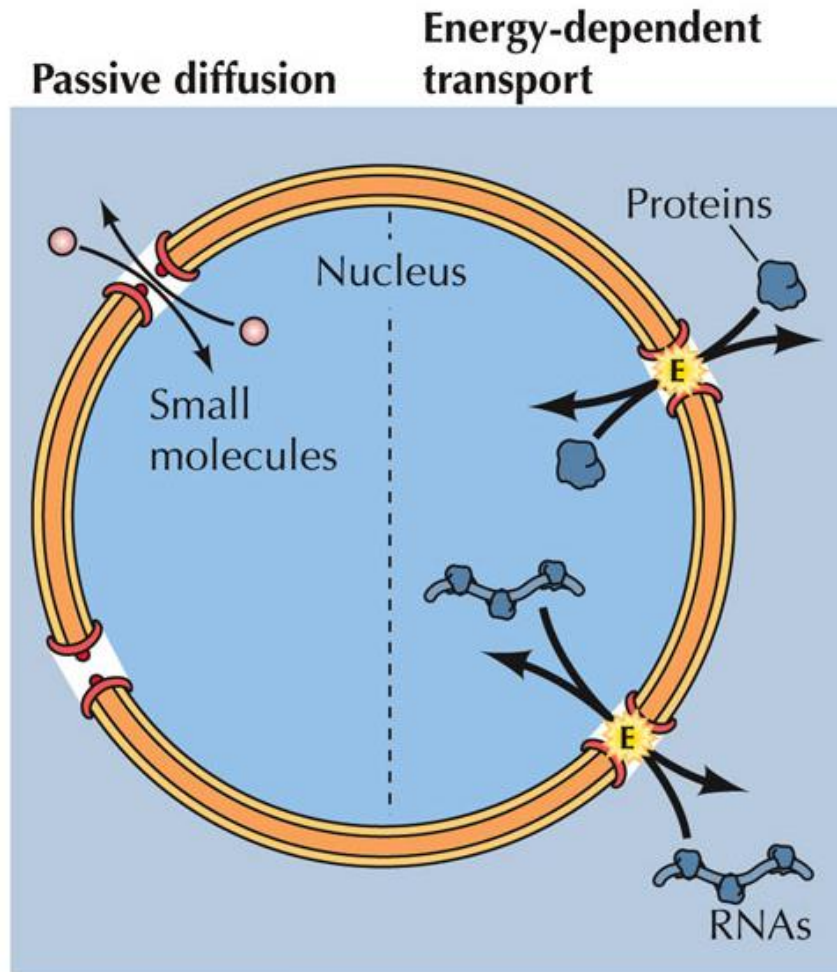
Outer nuclear membrane

Inner nuclear membrane:

Separated from nuclear material by the **nuclear lamina**

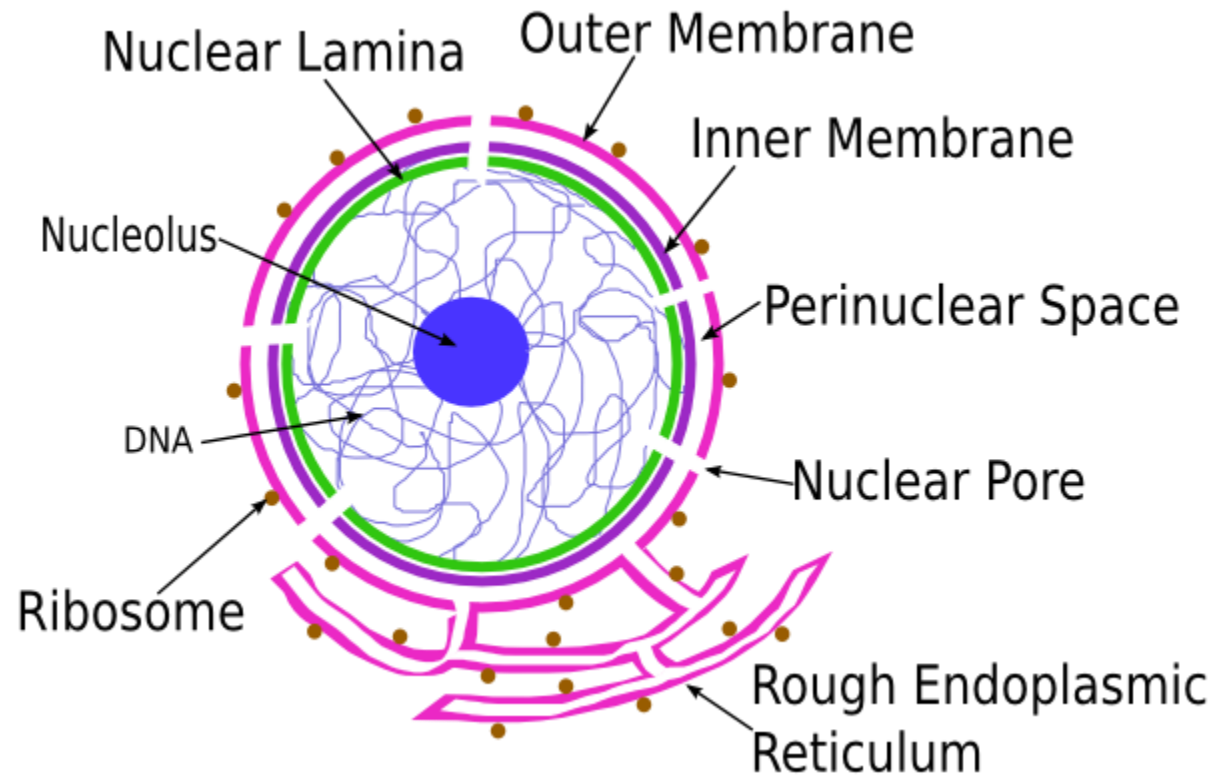


Nuclear pore



THE CELL, Fourth Edition, Figure 9.6 © 2006 ASM Press and Sinauer Associates, Inc.

Laminopathies are a group of genetic disorders caused by mutations in genes encoding proteins of the nuclear lamina.

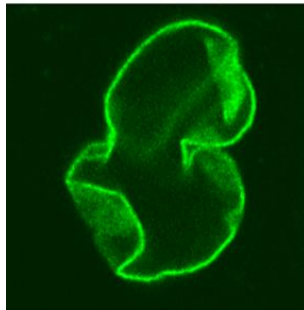
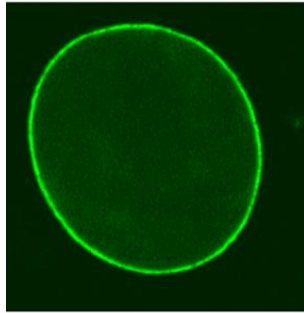


Emery-Dreifuss muscular dystrophy - A muscle wasting disease

Progeria - Premature aging

Restrictive dermopathy - A disease associated with extremely tight skin and other severe neonatal abnormalities

Hutchinson Gilford Progeria Syndrome



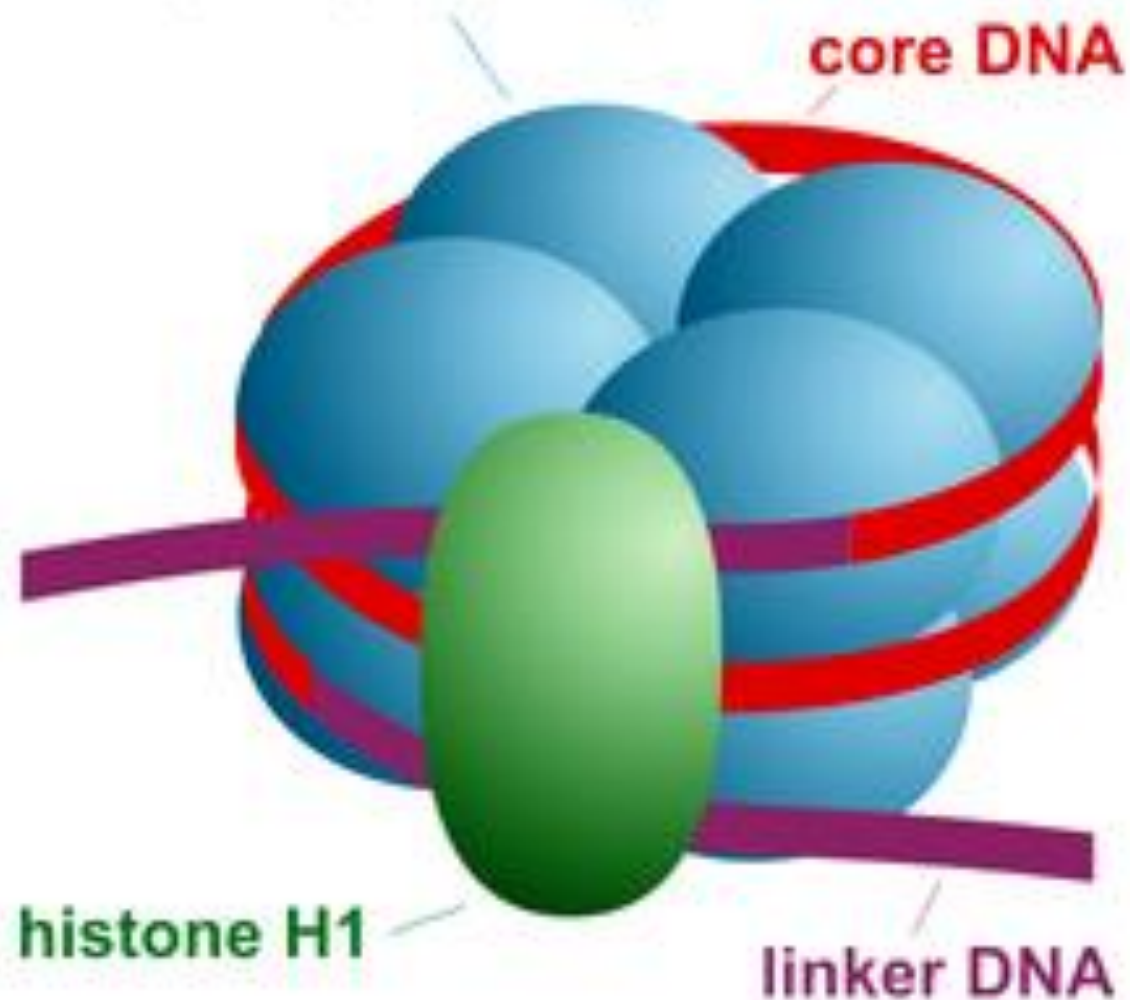
normal fibroblast and
of fibroblast of patient
with progeria.



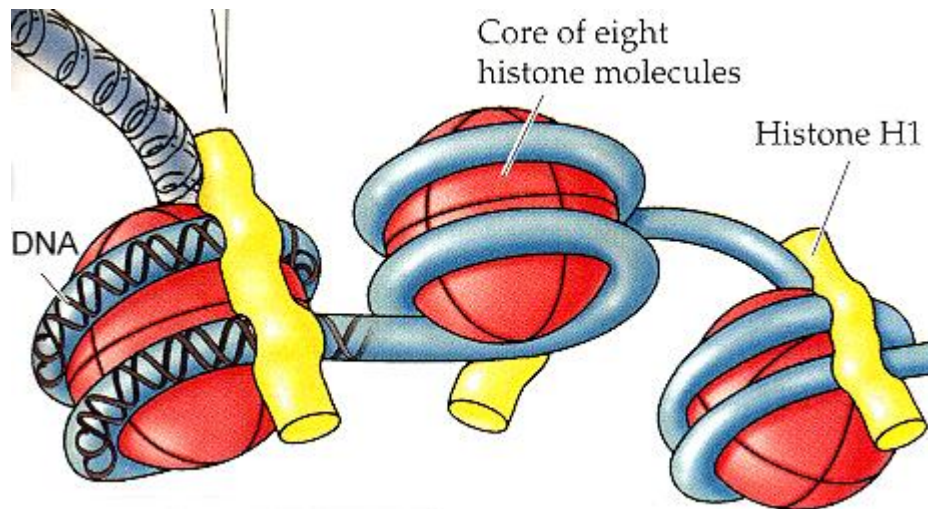
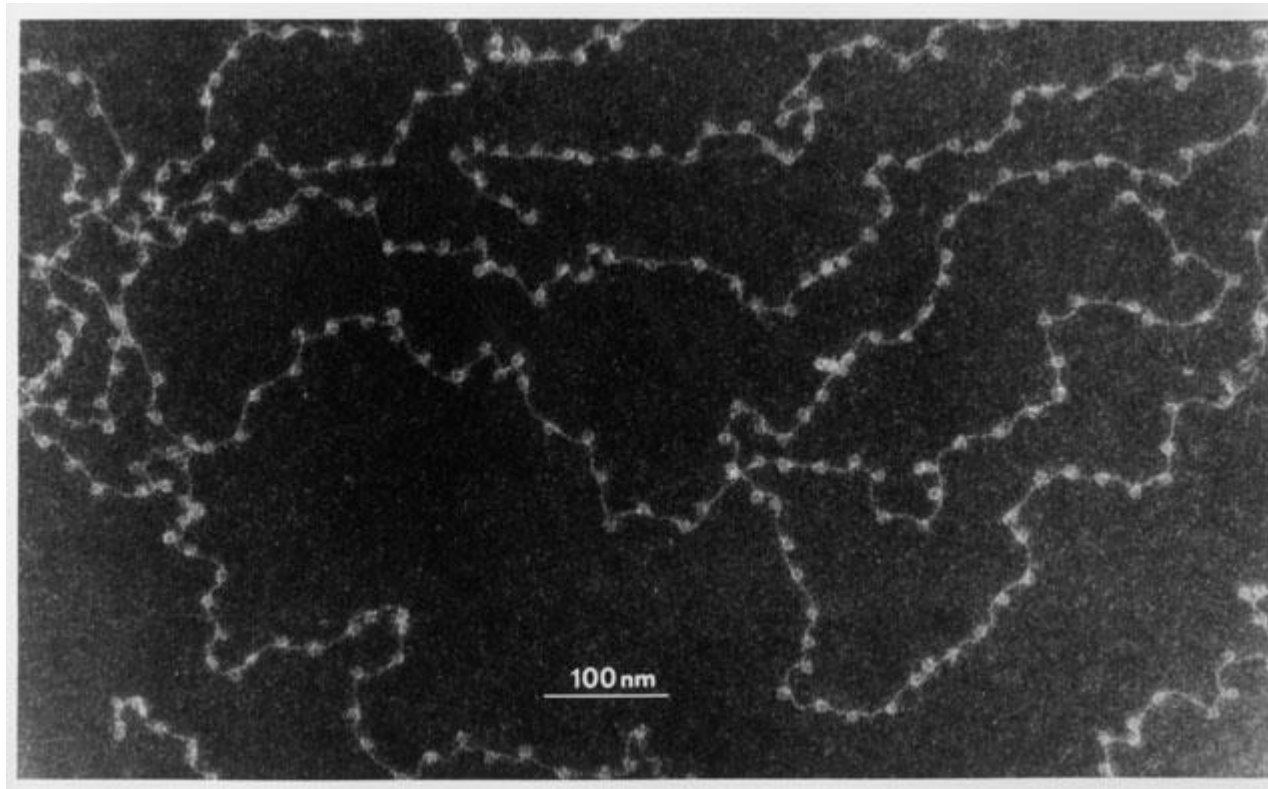
Extremely rare genetic disease, symptoms resembling aging are manifested during their first few months after birth. The average life expectancy for a child with progeria is about 13 years.

Symptoms: Scleroderma-like skin condition, limited growth and full-body alopecia (hair loss), atherosclerosis, kidney damage, loss of eyesight, and cardiovascular problems. People diagnosed with this disorder usually have small, fragile bodies, like those of elderly people.

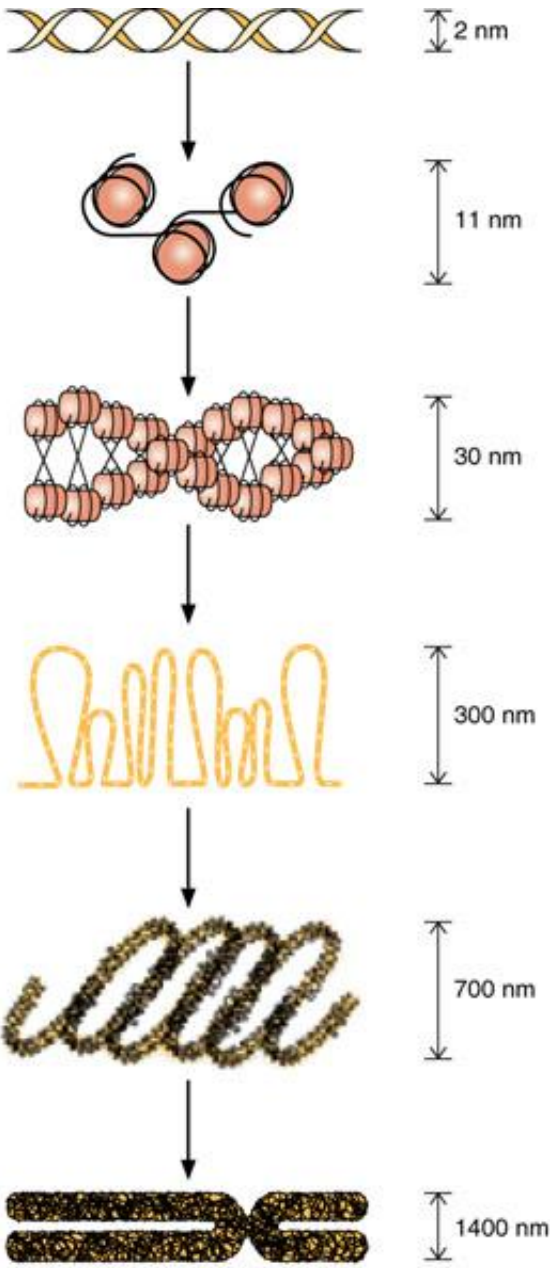
octamer of core histones:
H2A, H2B, H3, H4 (each one $\times 2$)



"beads on a string"
structure (euchromatin).



The order of chromatin packing in the metaphase chromosome



2-nm DNA double helix

„Beads on a string”

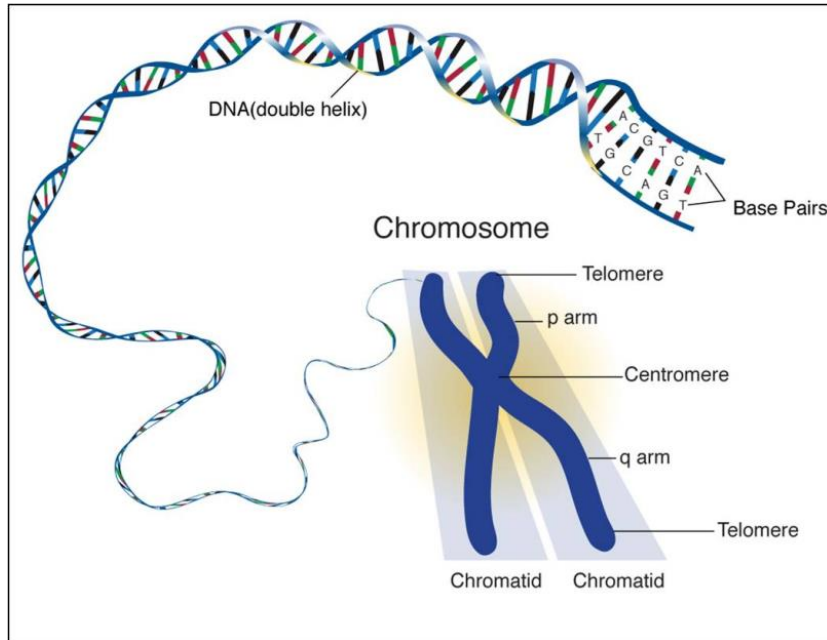
Association of DNA with octamer of histones;
formation of 11nm

And with H1 - 30 nm fibers of chromatin

Further condensation,
formation of 300 and
700 nm filaments

Maximal packing of
DNA in metaphase
chromosome

CHROMOSOMES - are maximally condensed in metaphase.

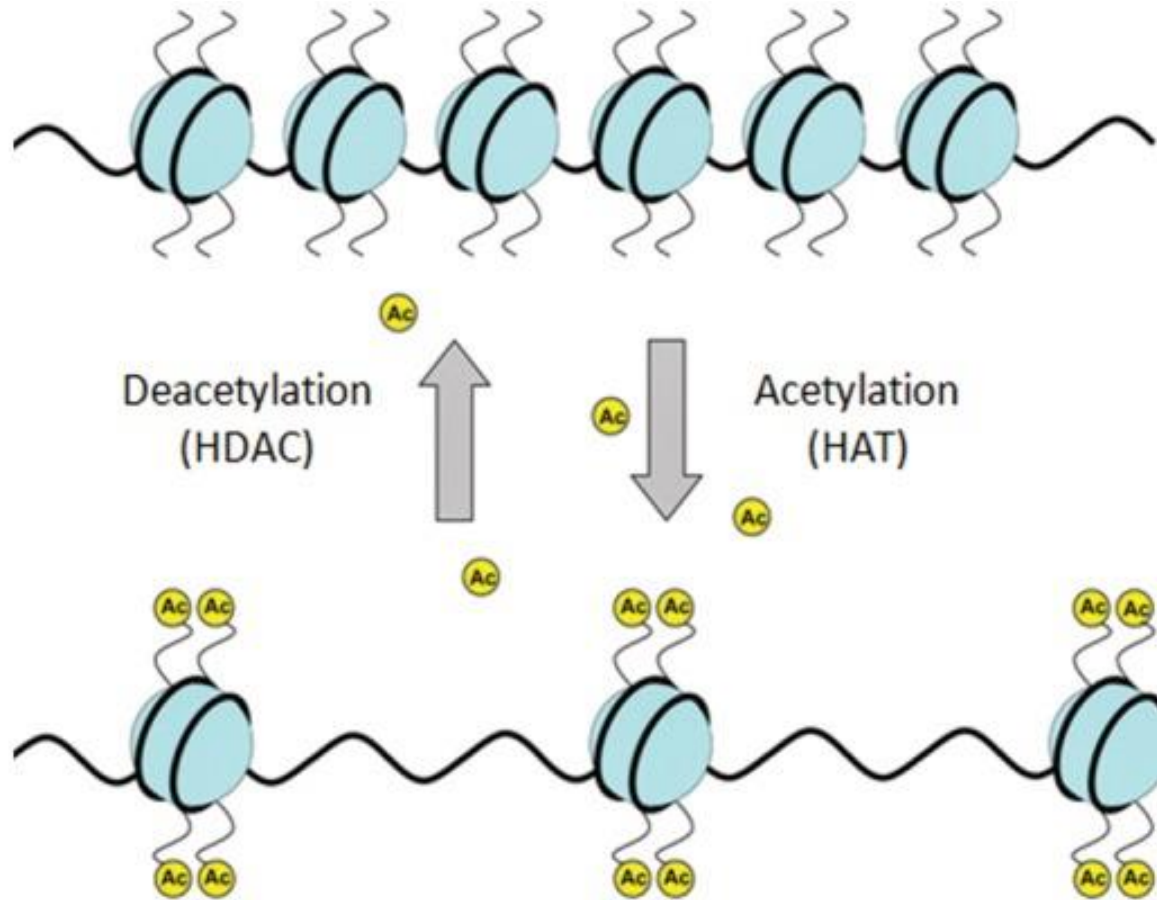


Human genome consists of 23 homologous pairs of chromosomes (maternal and paternal) – 22 autosomes and 1 pair of sex chromosomes.

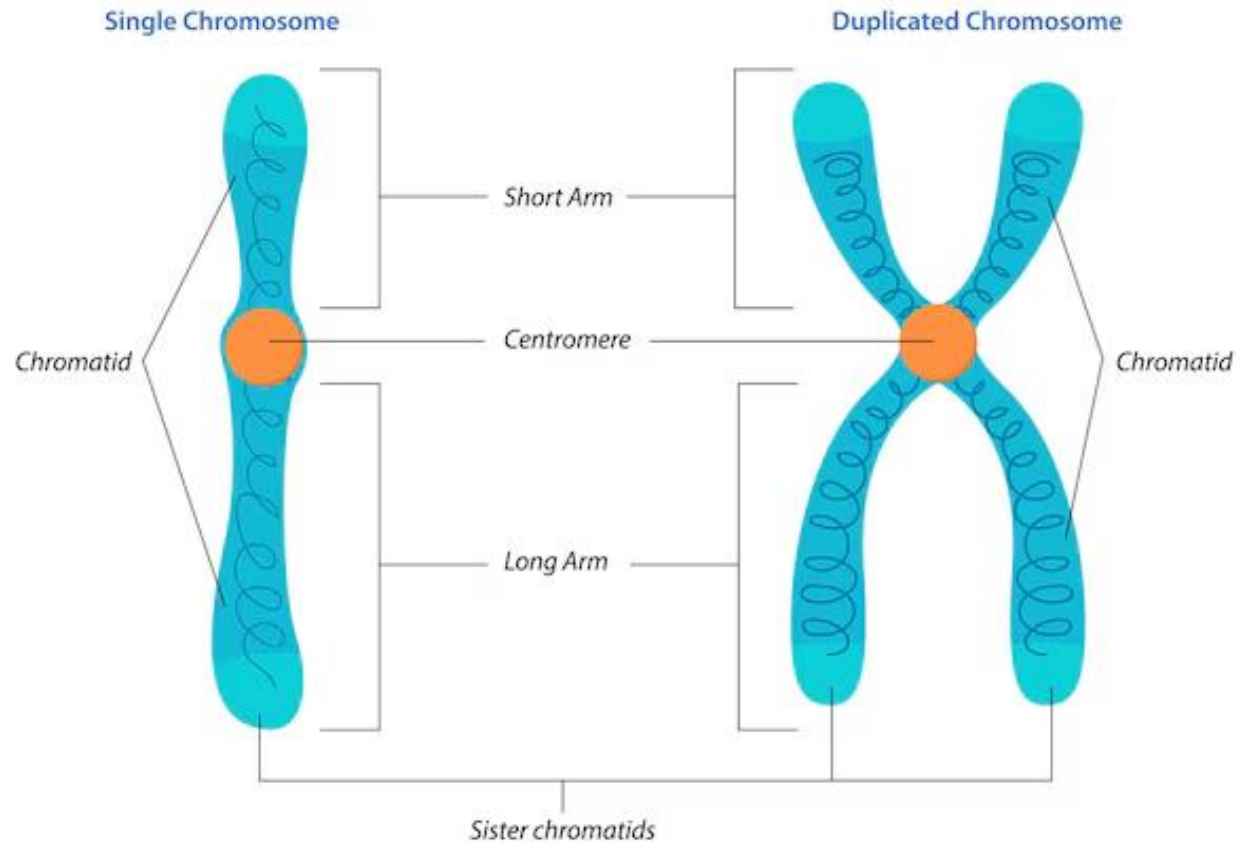
Cells containing full set of chromosomes are diploid – **2n**. They have 23 pairs of chromosomes. These cells have also 2 copies of each gene – **2c**.

Gametes are haploid **1n**. They have only 1 copy of each gene – **1c**.

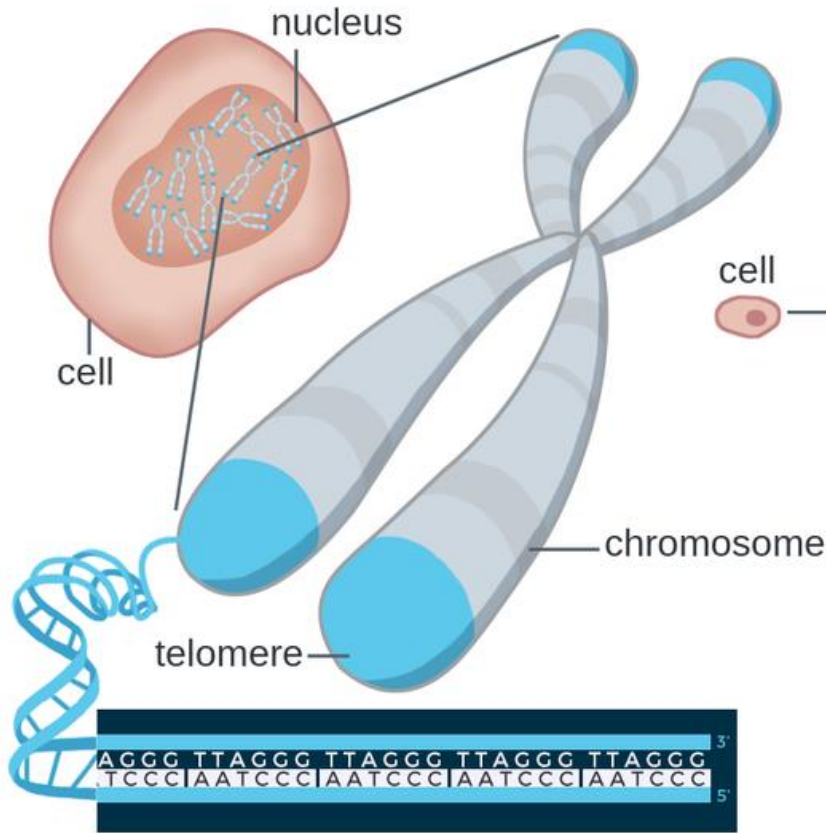
Histone modification – ACETYLATION and DEACETYLATION



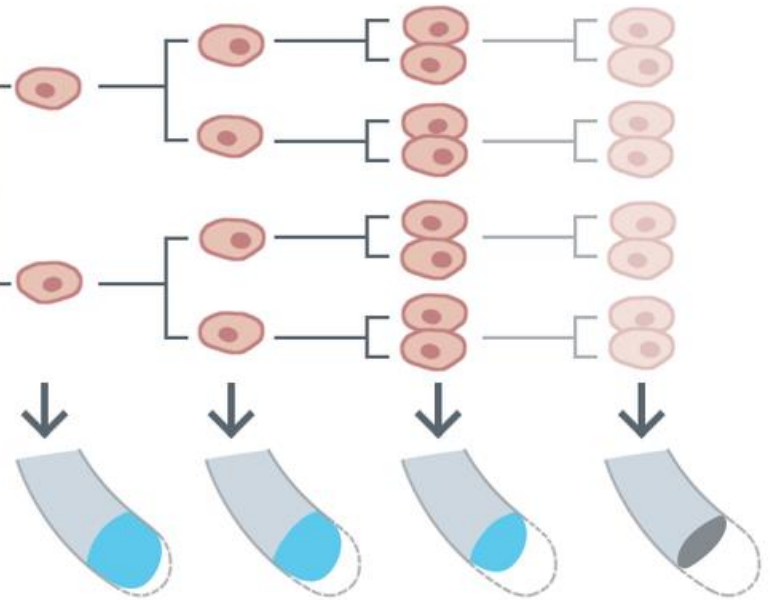
Chromosome structure



Telomeres



As the cell divides over time (healthy cell)..



...telomeres shorten, eventually signaling the cell to stop dividing (senescence).

Telomerase

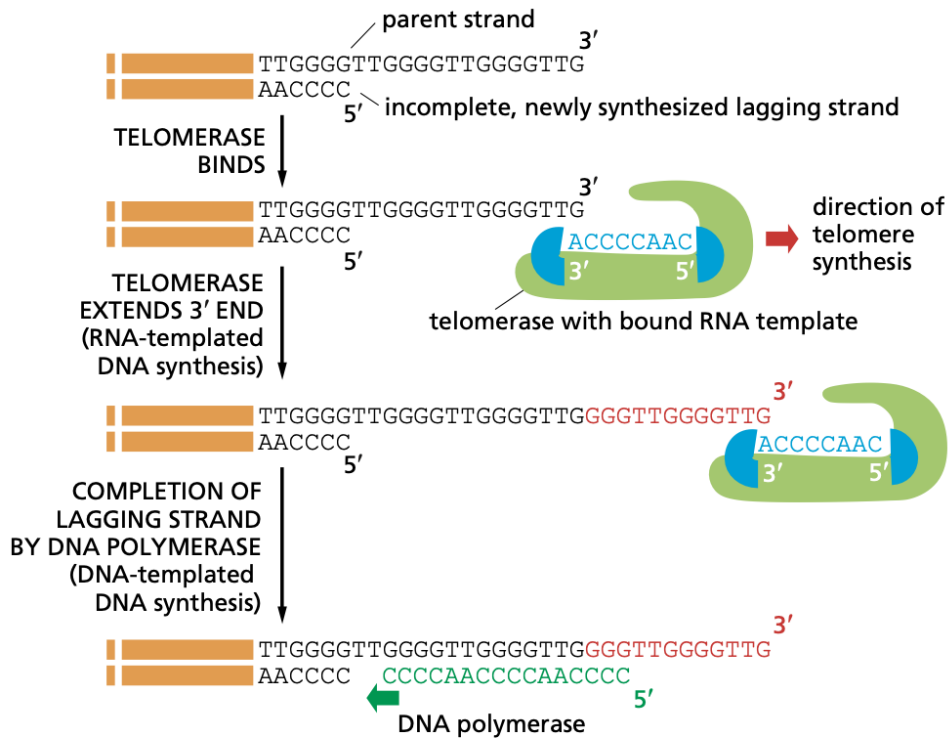
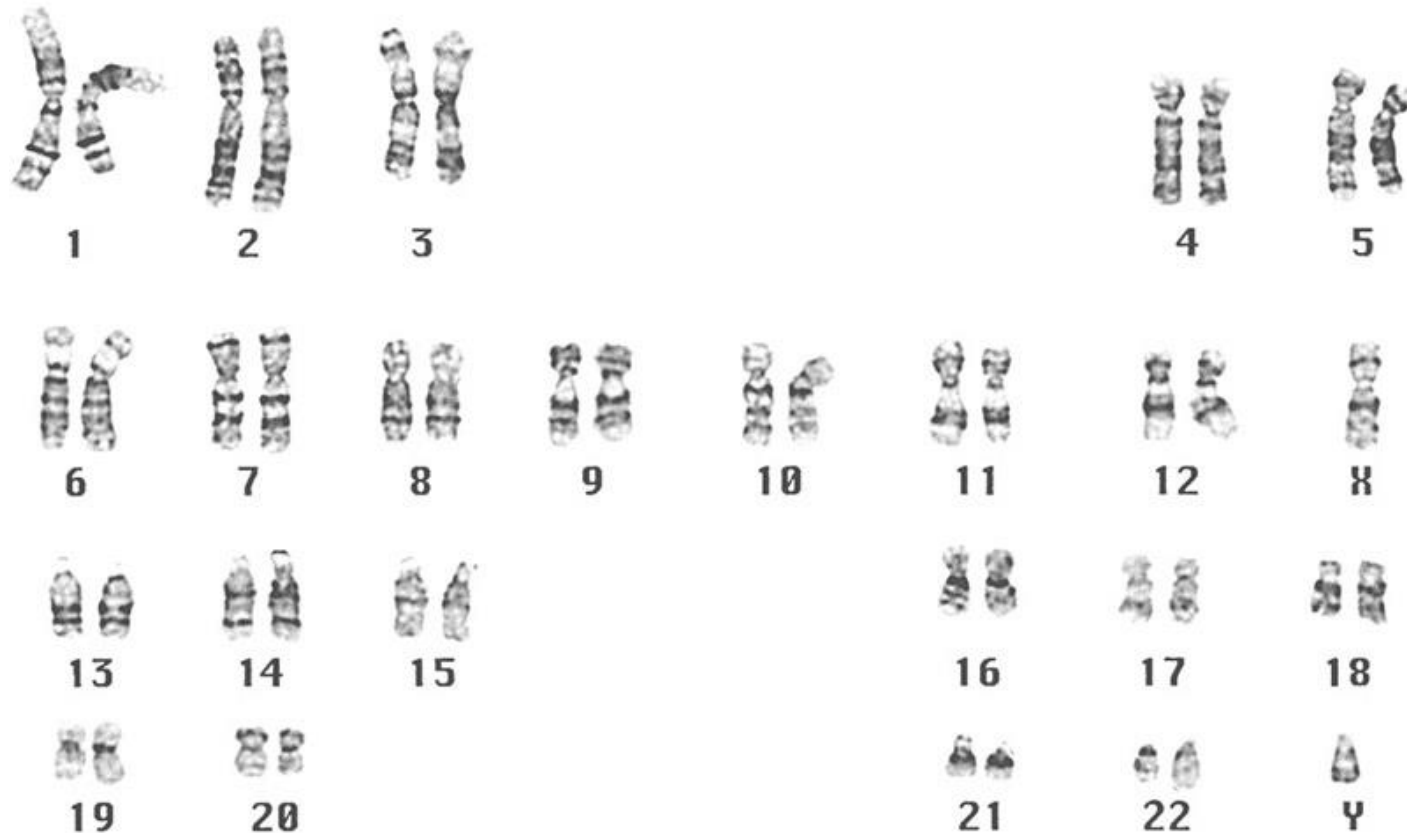


Figure 5–34 Telomere replication. Shown here are the reactions that synthesize the repeating sequences that form the ends of the chromosomes (telomeres) of diverse eukaryotic organisms. The 3' end of the parental DNA strand is extended by RNA-templated DNA synthesis; this allows the incomplete daughter DNA strand that is paired with it to be extended in its 5' direction. This incomplete, lagging strand is presumed to be completed by DNA polymerase α , which carries a DNA primase as one of its subunits (**Movie 5.6**). The telomere sequence illustrated is that of the ciliate *Tetrahymena*, in which these reactions were first discovered.

Human karyotype - A karyotype is the number and appearance of chromosomes in the nucleus of a cell.

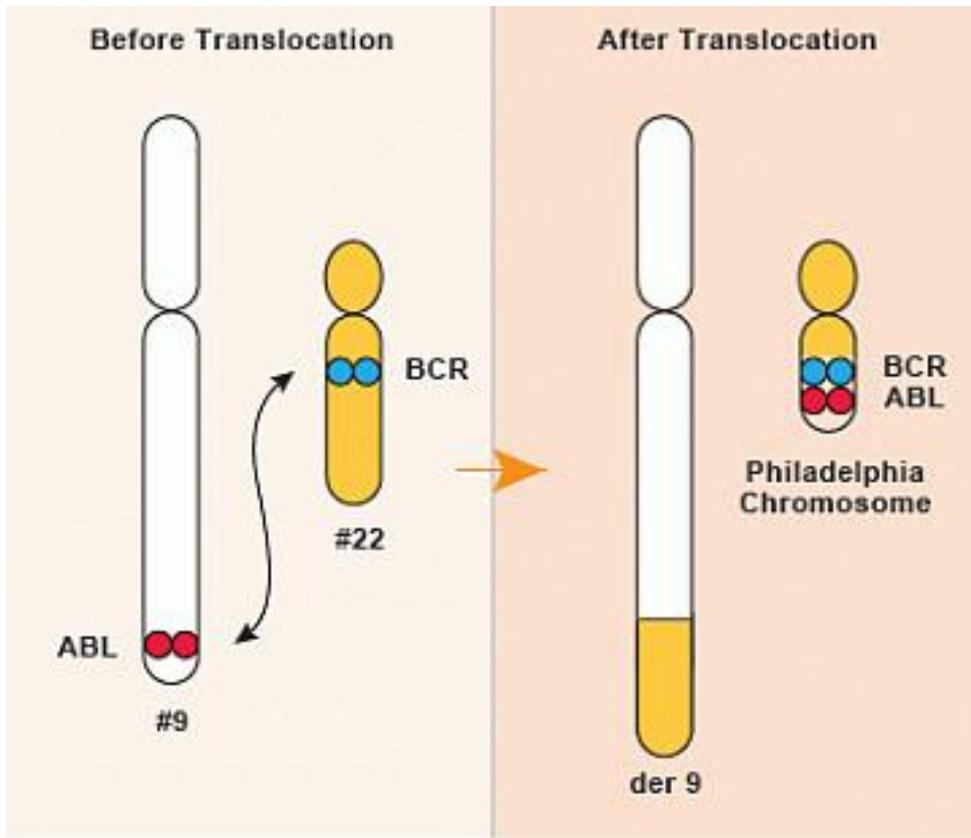
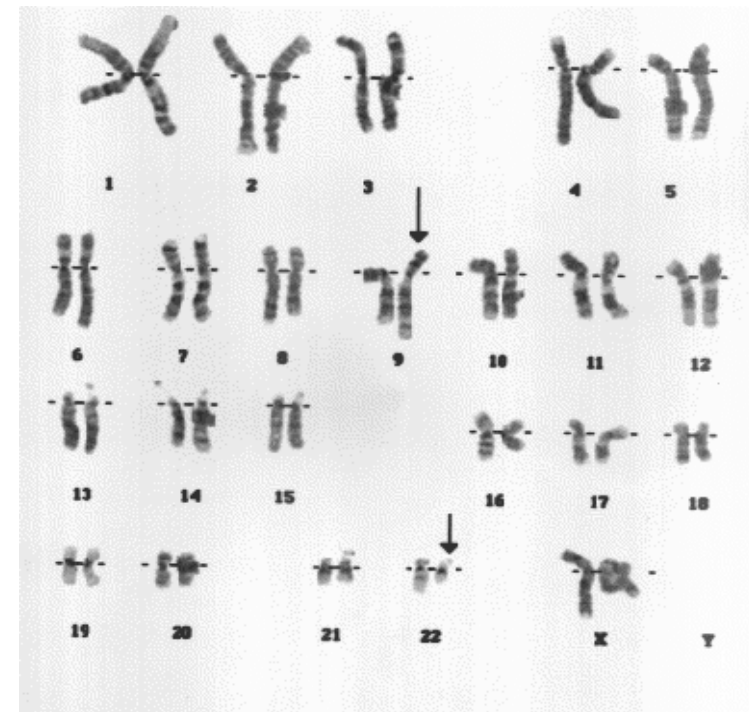
Karyotype is used to study chromosomal aberrations



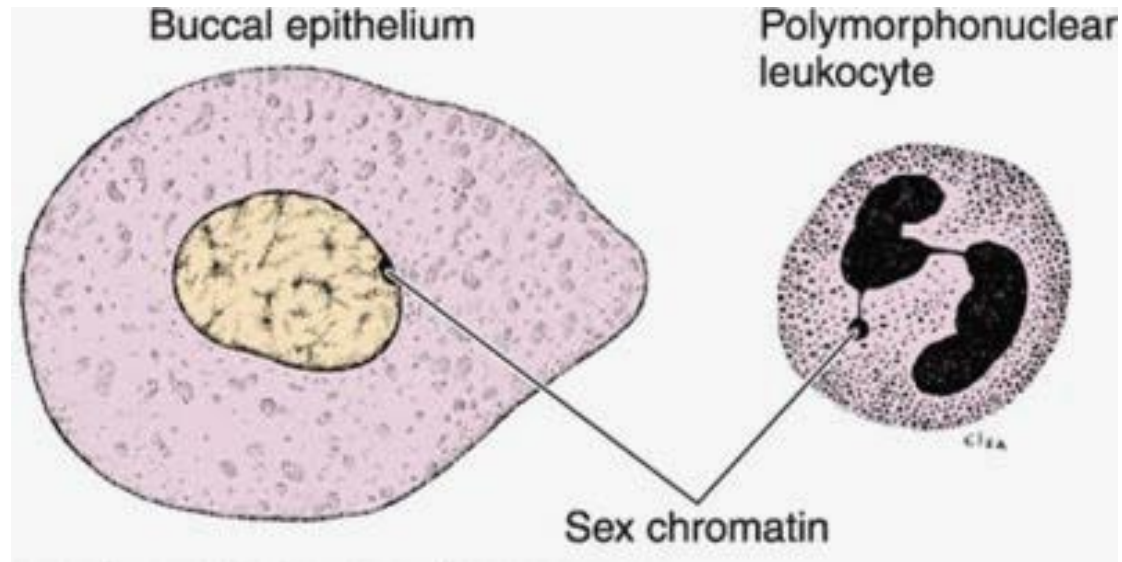
Banding is obtained after special staining, e.g. with Giemsa stain (adenine-thymine-rich regions of chromosomes).

Syndrom	Karyotyp	Symptome
Patau		
Down		
Edwards		
Klinefelter		
Turner		

Philadelphia chromosome - is a specific chromosomal abnormality that is associated with chronic myelogenous leukemia (acute lymphoblastic leukemia and acute myelogenous leukemia).

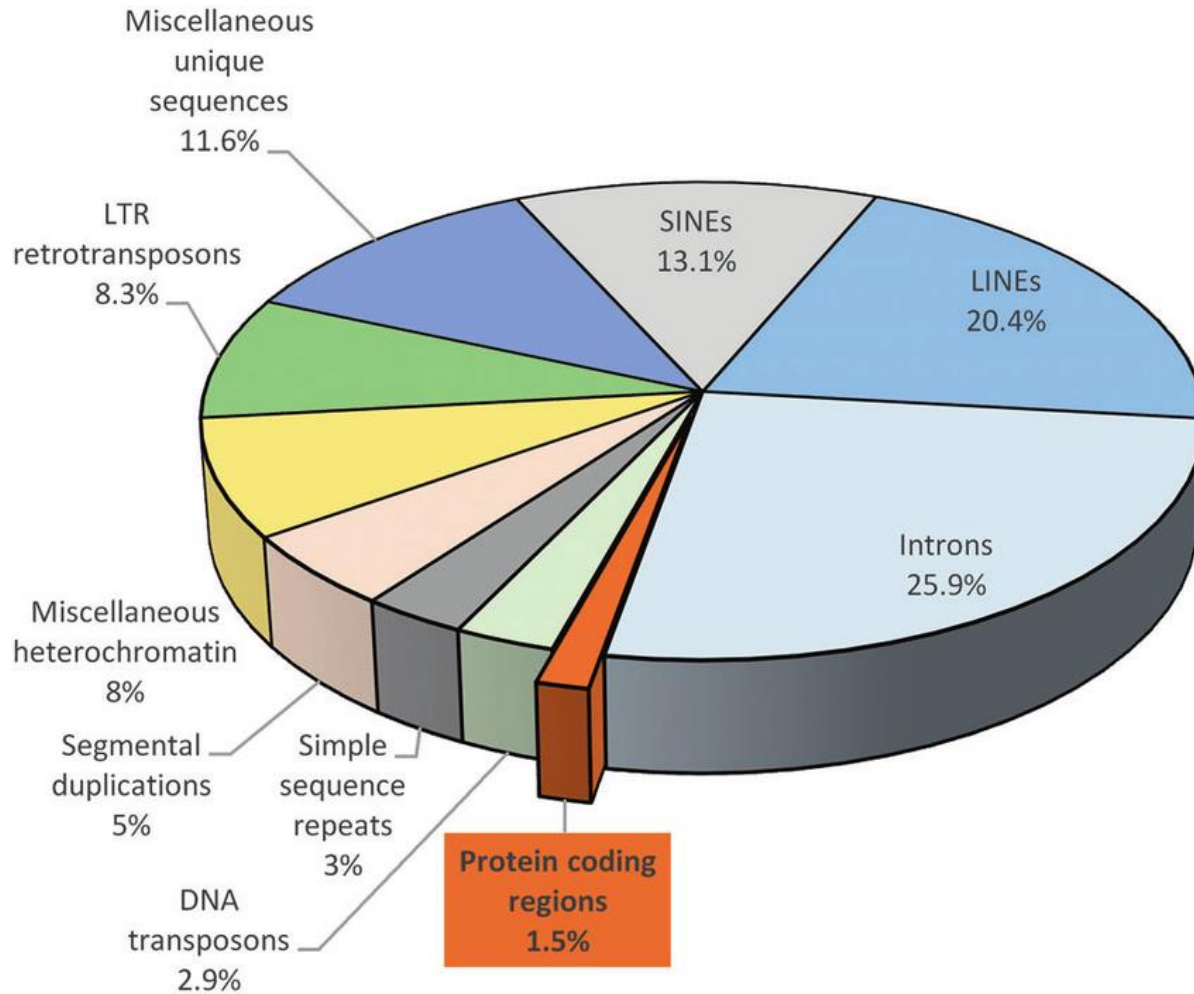


Sex chromatin in female epithelium (dense granule) and leukocyte (drumstick shape)

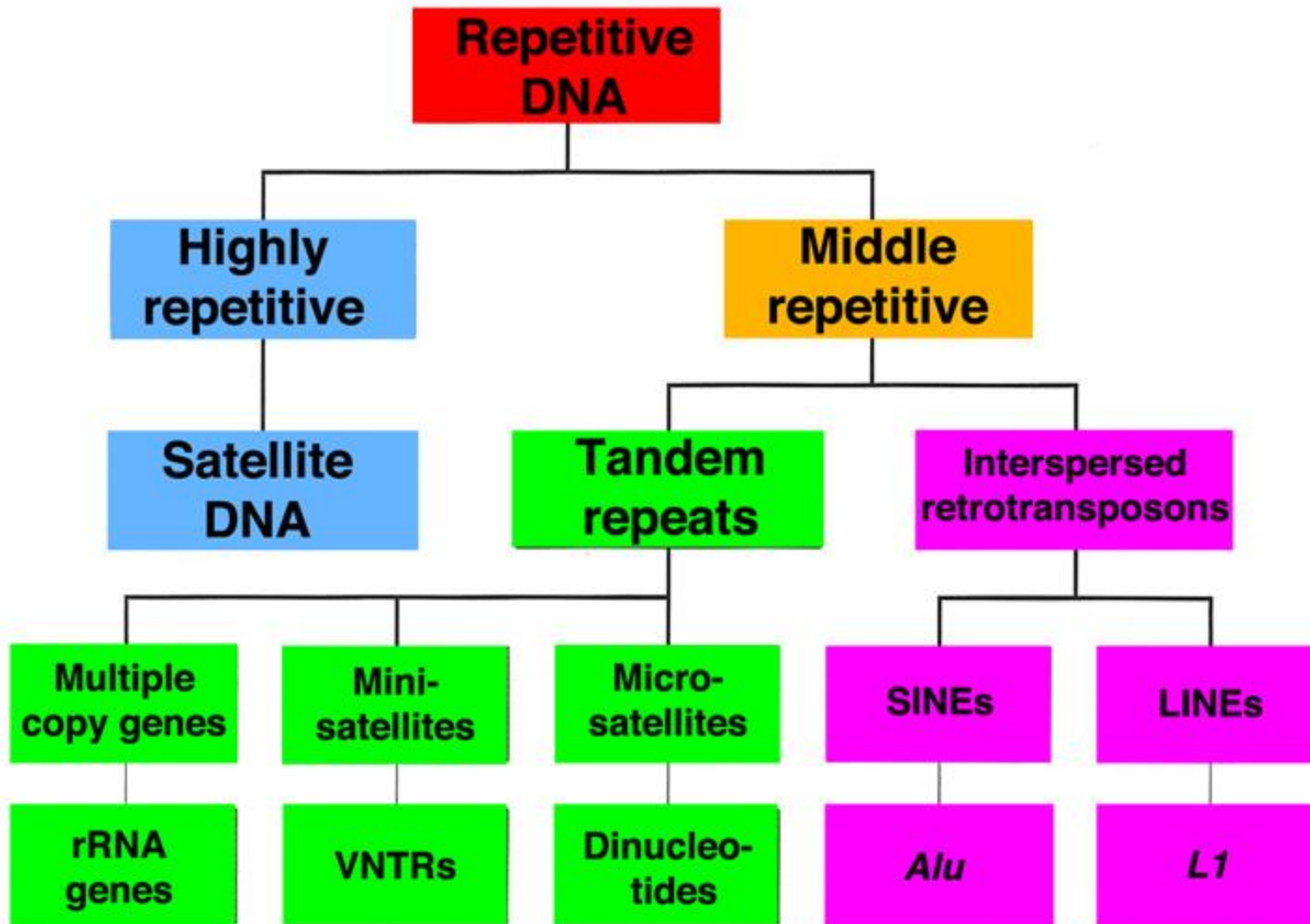


Only one of the two X chromosomes in female somatic cells is transcriptionally active. The inactive X chromosome remains inactive throughout the life of that individual.

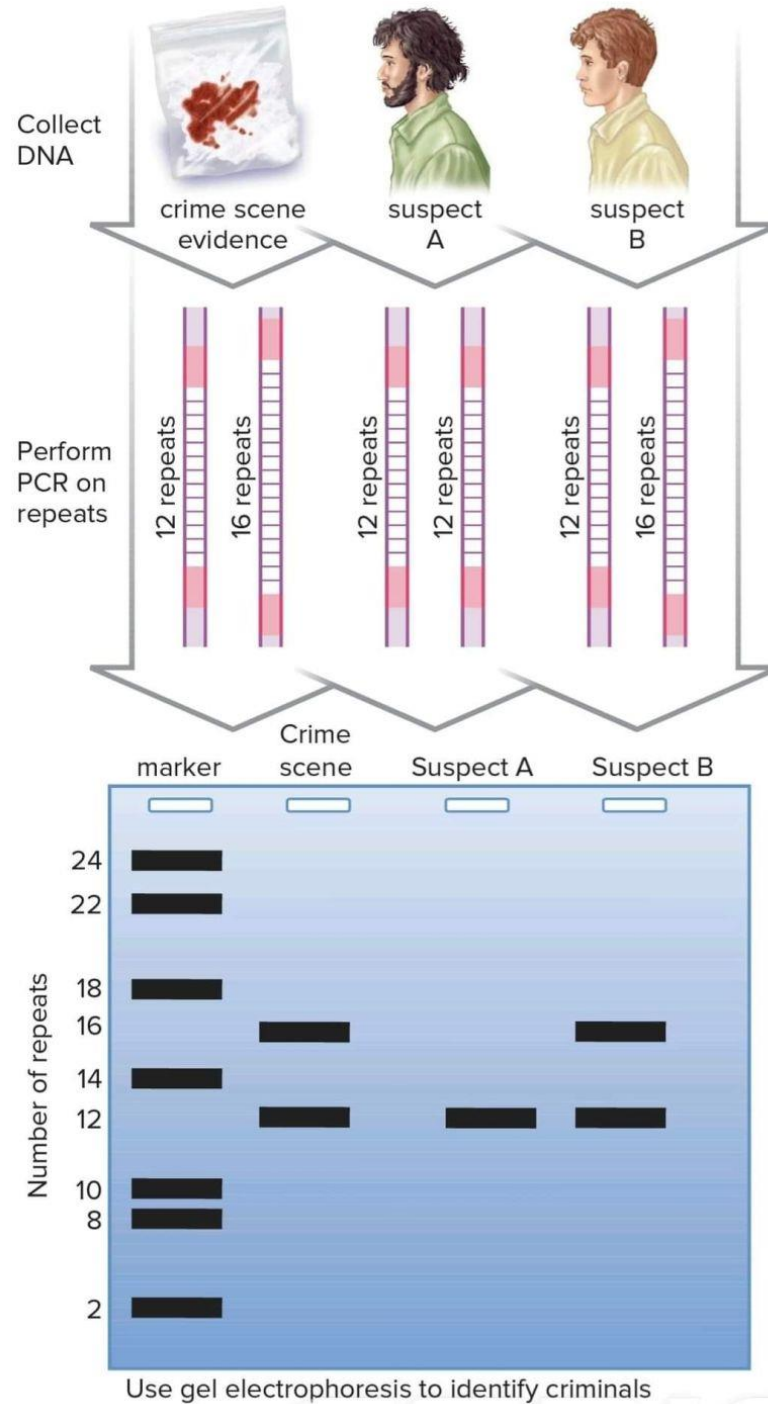
Organisation of human genome



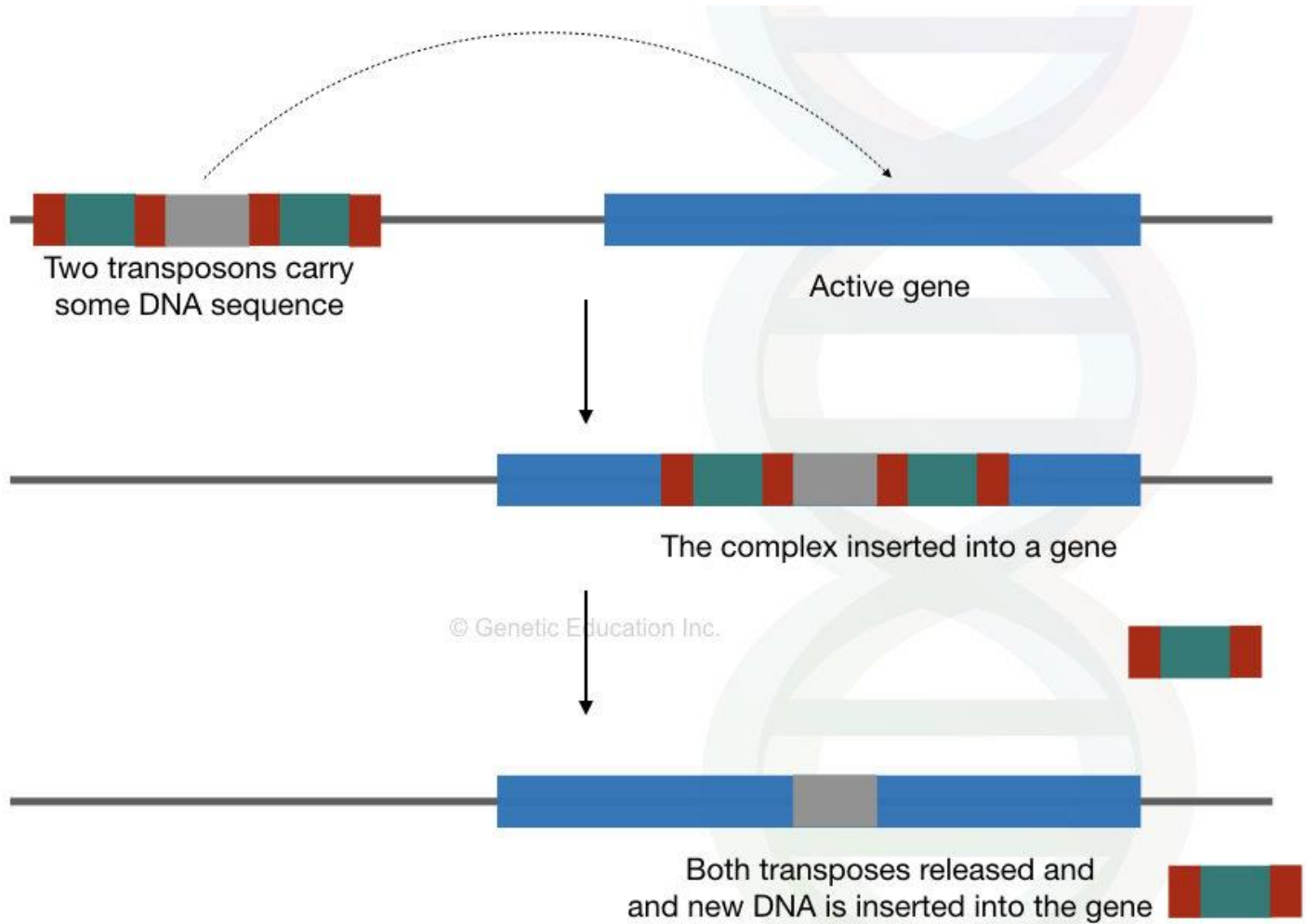
Classification of human repetitive DNA



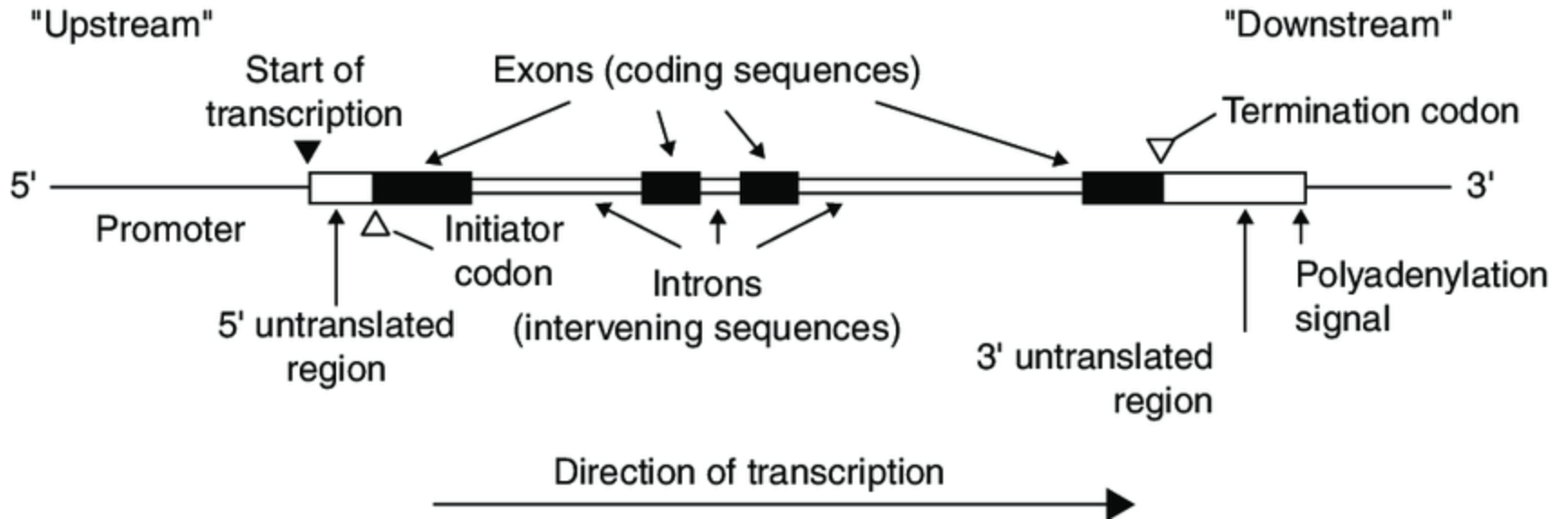
DNA fingerprint



Transposones



Human gene structure



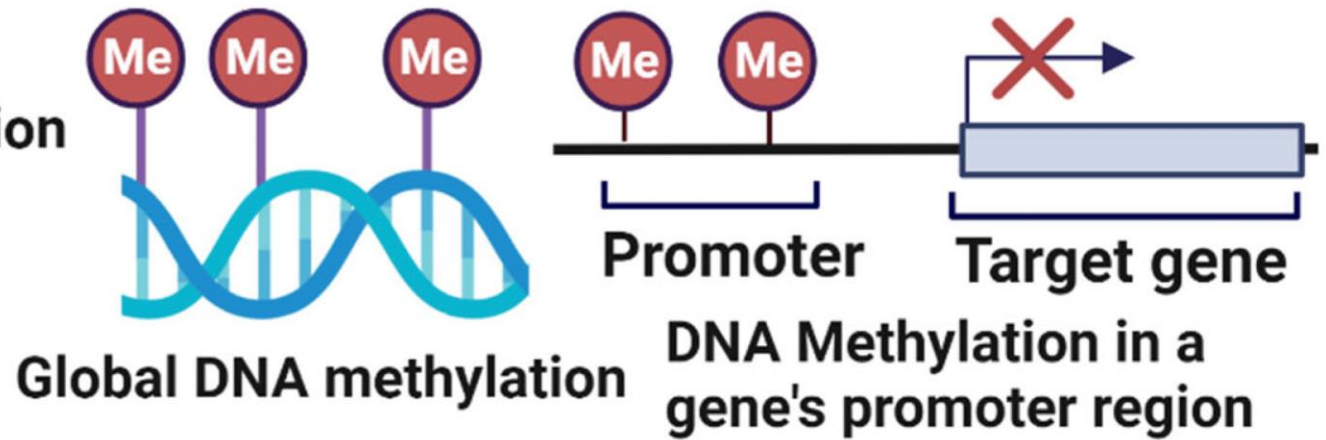
Epigenetic alterations to the genome

a) DNA methylation

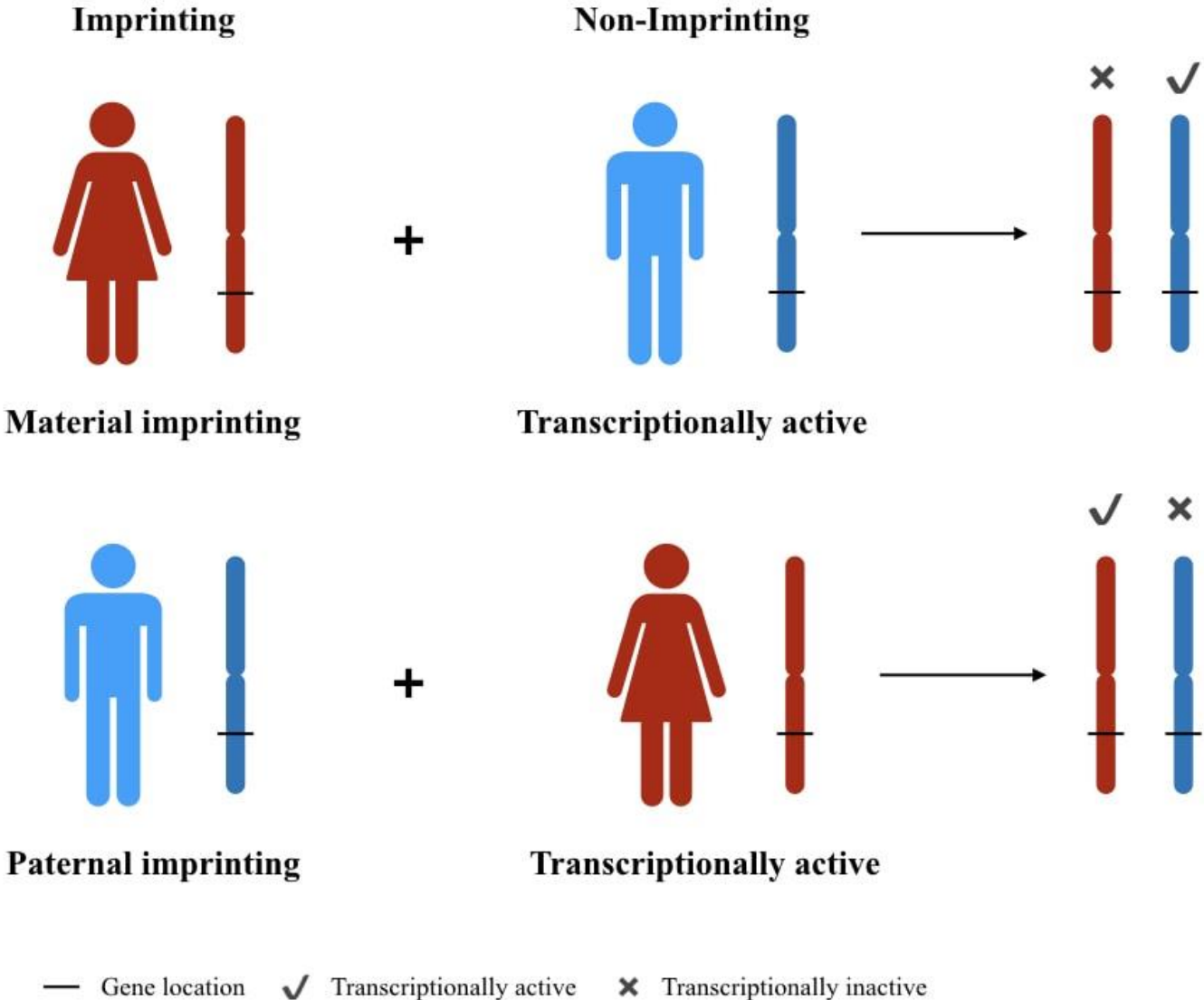
DNA methylation

enzymes:

- DNMTs
- TETs



Genomic imprinting



Prader-Willi/Angelman syndrome

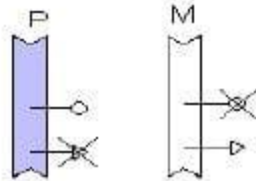
Both produced by same genetic mutation

*PWS: deletion of paternal copy or UPD of chromosome 15q11-q13

*Language, motor and developmental delays, excessive weight gain

*AS: deletion of maternal copy, maternal mutation of UBE3A gene or UPD of chromosome 15q11-q13

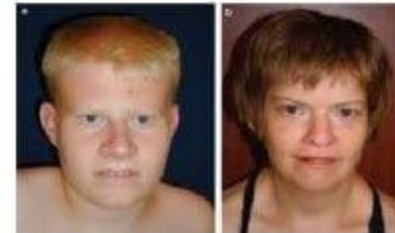
*Severe mental retardation, happy demeanor, non-verbal



Normal



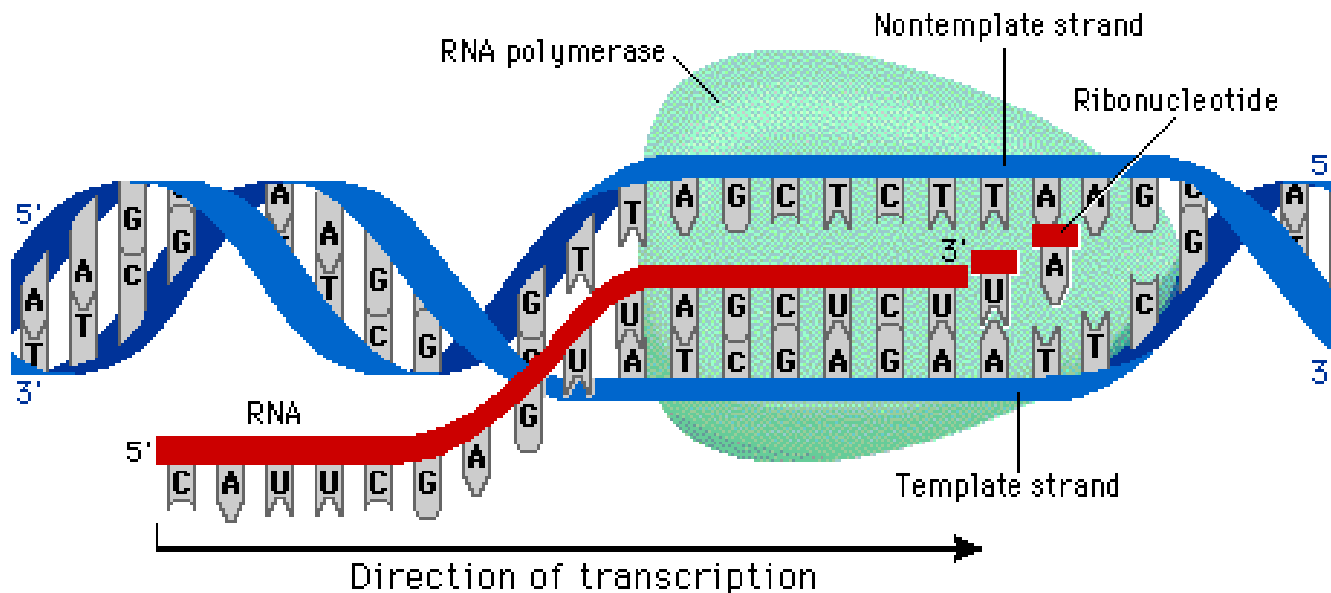
Angelman



Prader-Willi

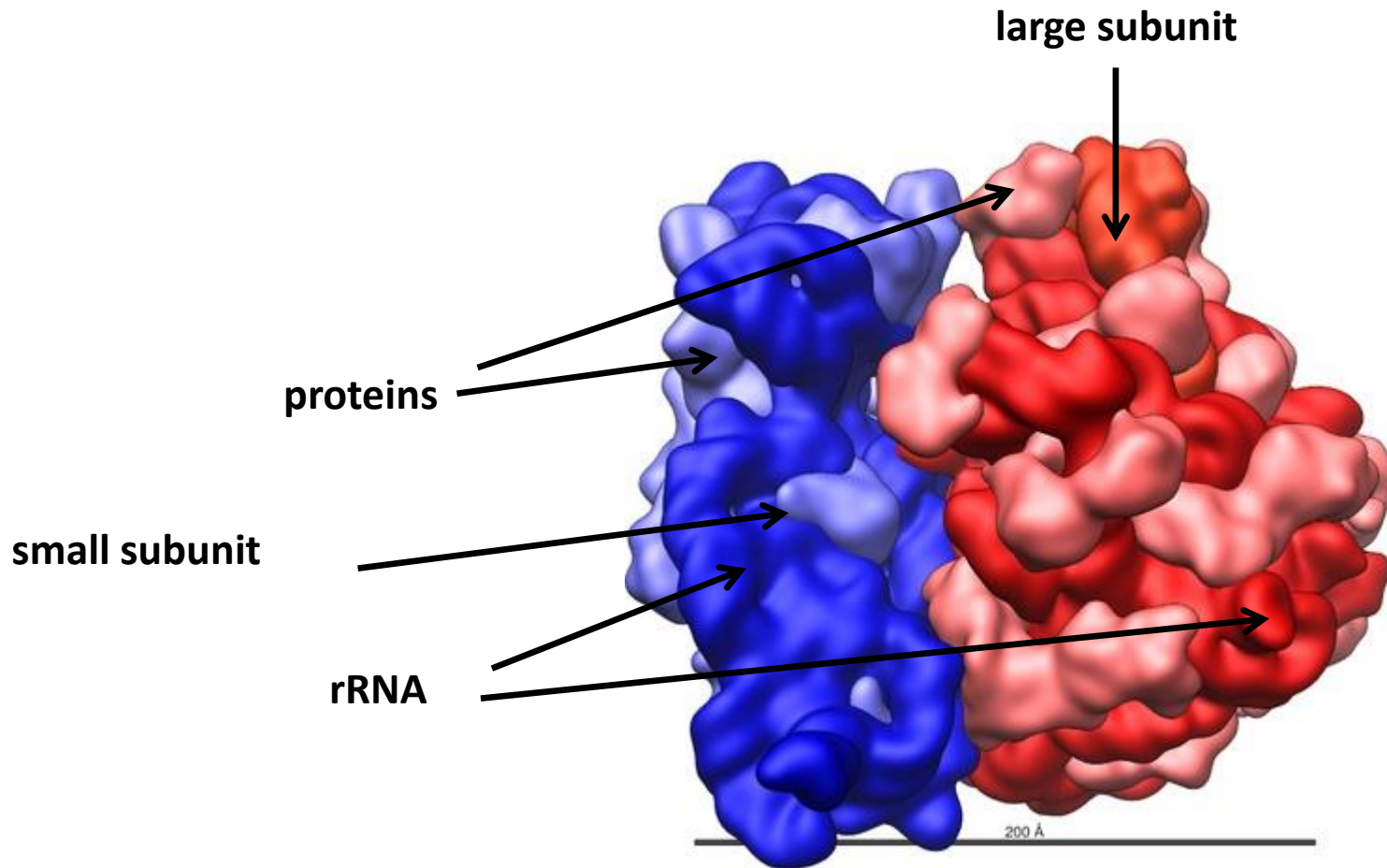
TRANSCRIPTION

- DNA sequence is read by an RNA polymerase, which produces a complementary, antiparallel RNA strand. New-formed RNA contains uracil (U) instead of thymine (T) presented in DNA.



Ribosomal RNA (rRNA)

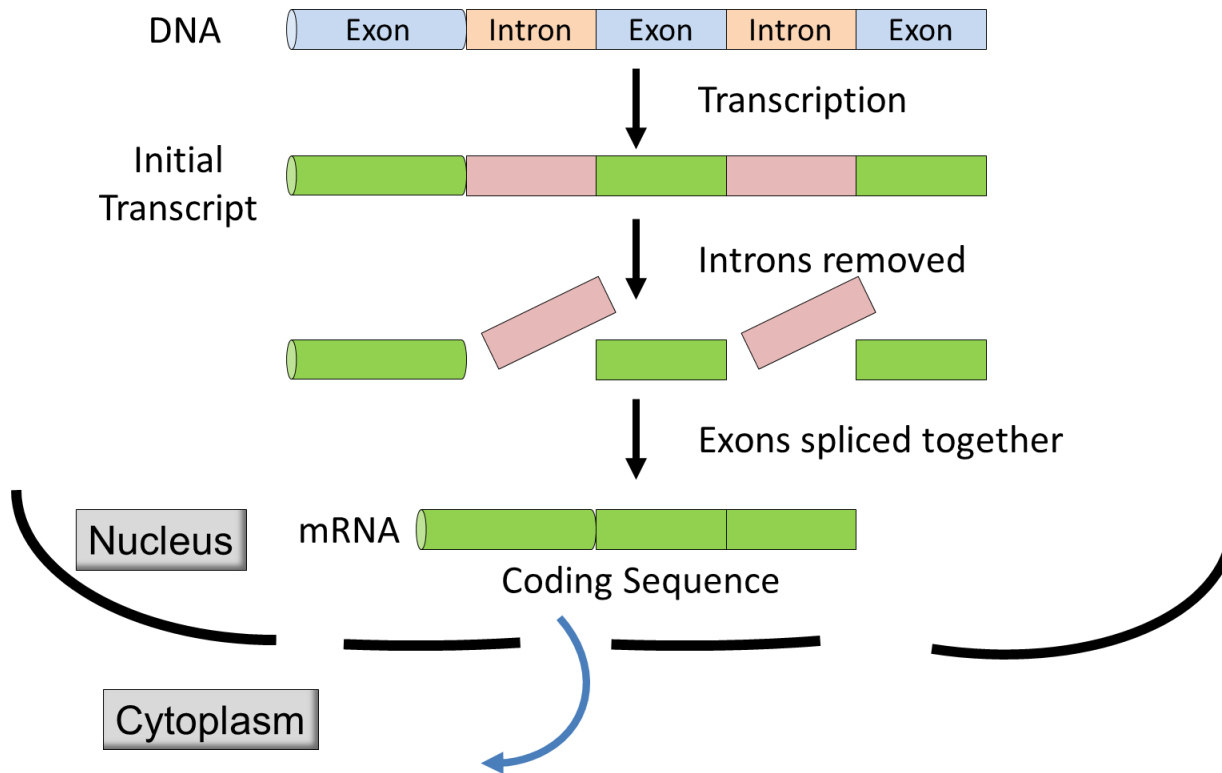
- rRNA is produced **nucleolus** by RNA polymerase I.



Messenger RNA (mRNA)

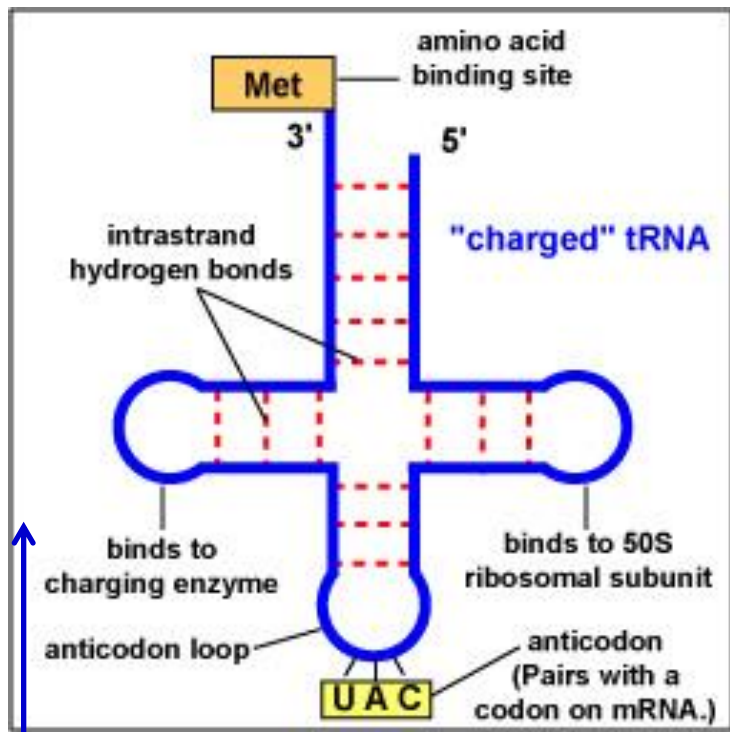
Produced from DNA by RNA polymerase II

- It is complementary copy of one gene
- contains coding segments - exons and noncoding segments – introns. Introns must be removed, exons must be spliced together.
- After processing mRNA is transported into cytoplasm. .



Transfer RNA (tRNA)

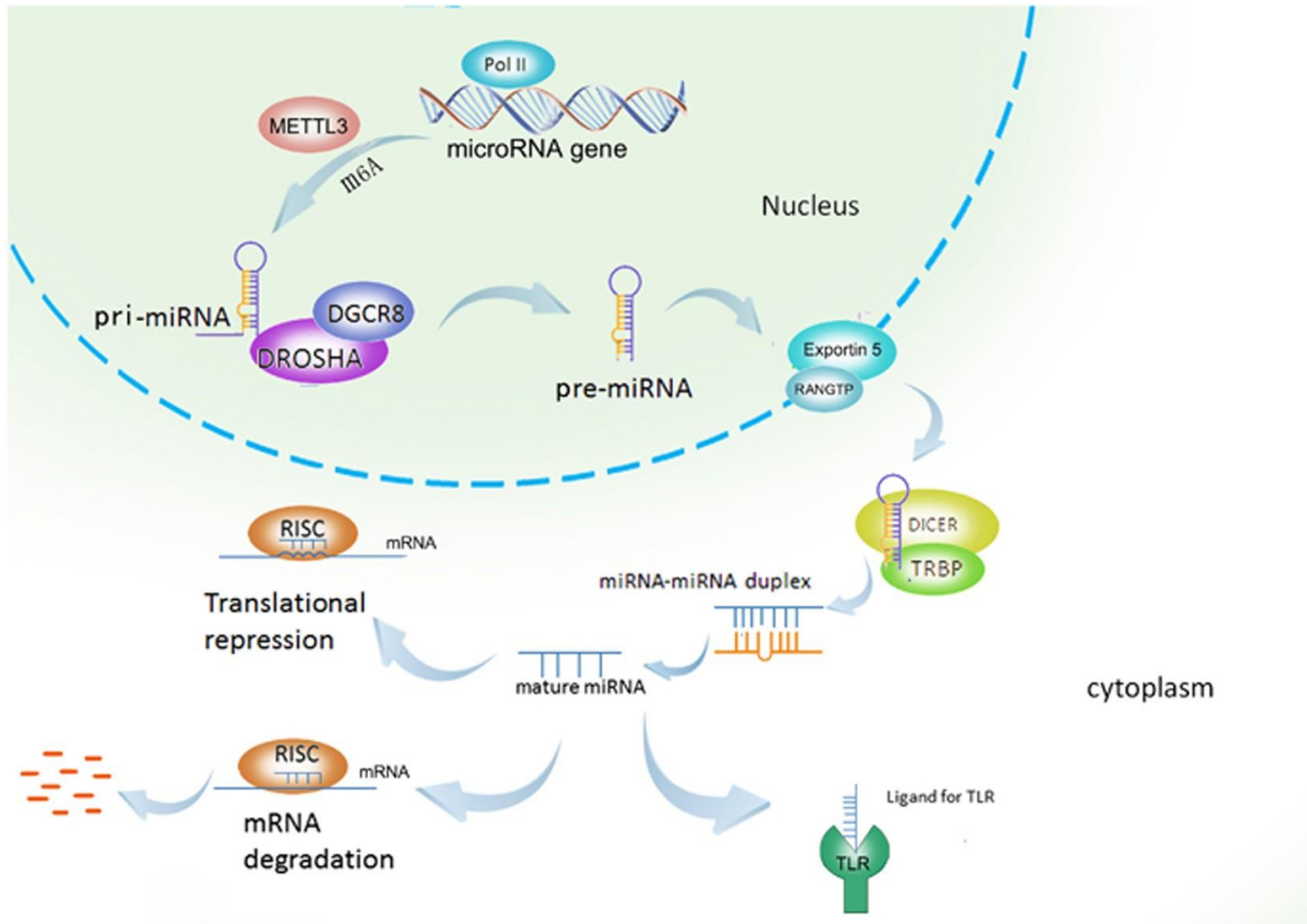
Produced from DNA by RNA polymerase III



aminoacyl tRNA synthetase

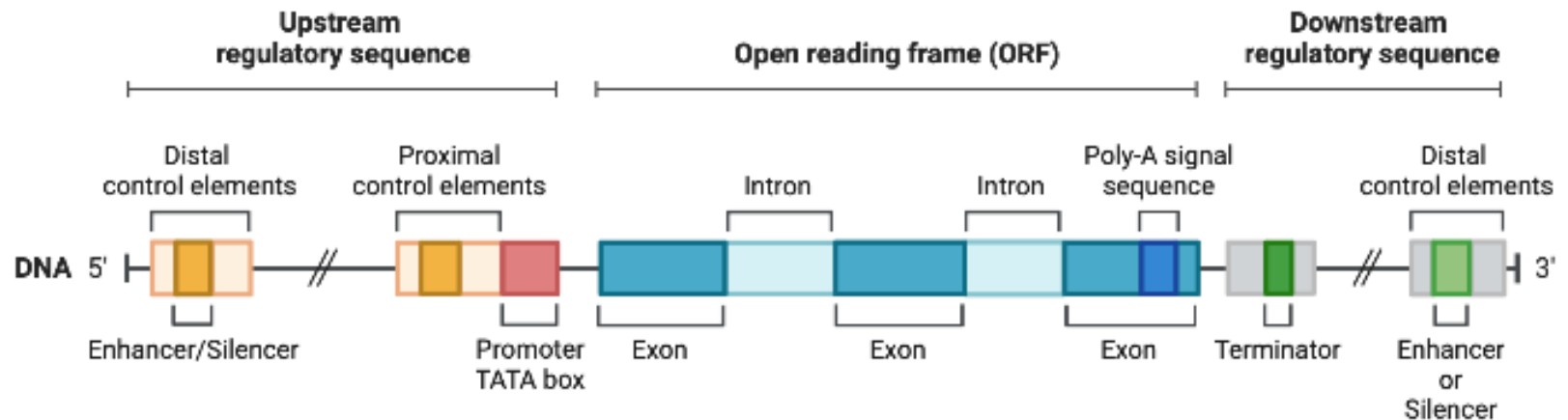
tRNA transfers activated amino acids to the ribosome-mRNA complex. Here amino acids are incorporated into forming protein.

micro RNA (miRNA)



Gene structure

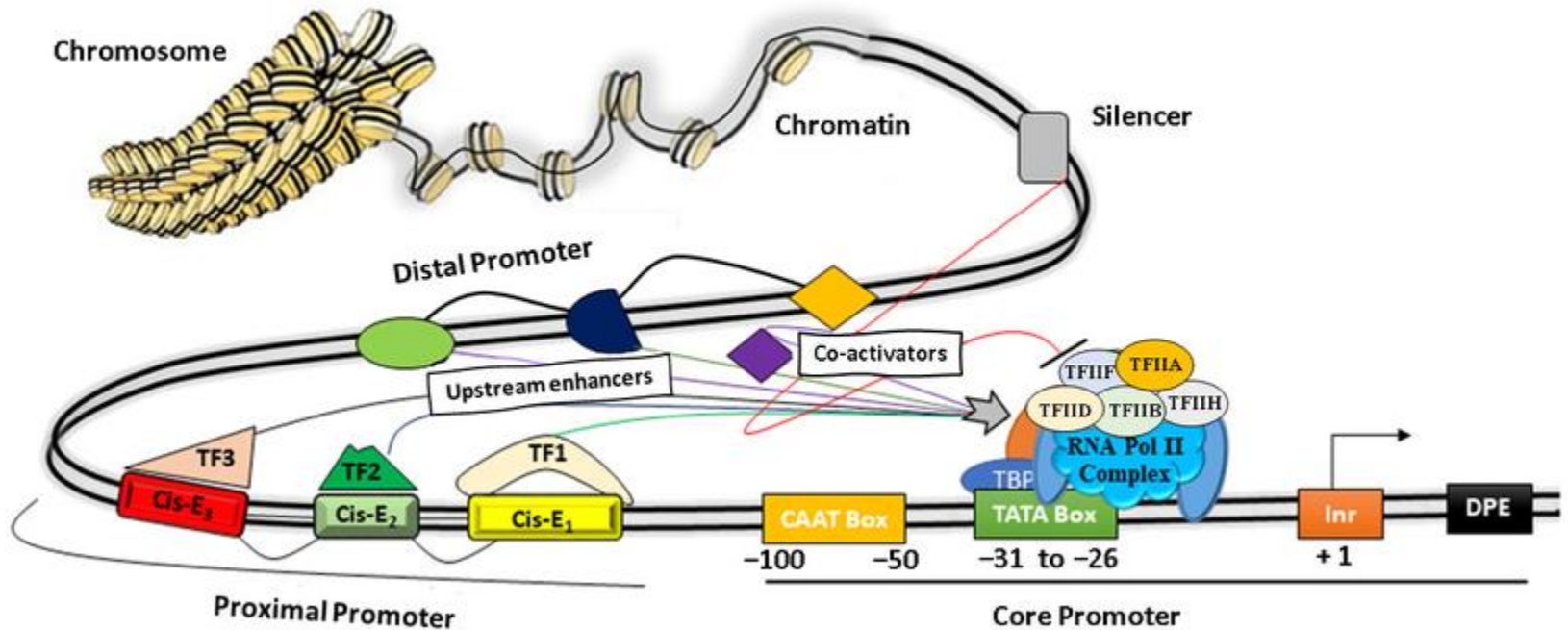
Eukaryotic Gene Structure



TYPES OF RNA POLYMERASE

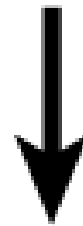
- **RNA polymerase I** - is located in the nucleolus and synthesizes 28S, 18S, and 5.8S rRNAs.
- **RNA polymerase II** - is located in the nucleoplasm and synthesizes hnRNA/mRNA and some snRNA.
- **RNA polymerase III** - is located in the nucleoplasm and synthesizes tRNA, some snRNA, and 5S rRNA

Basal promoter and enhancer



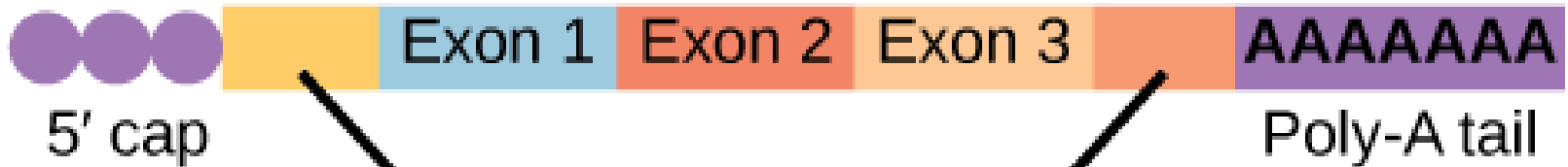
RNA processing

Primary RNA transcript



RNA processing

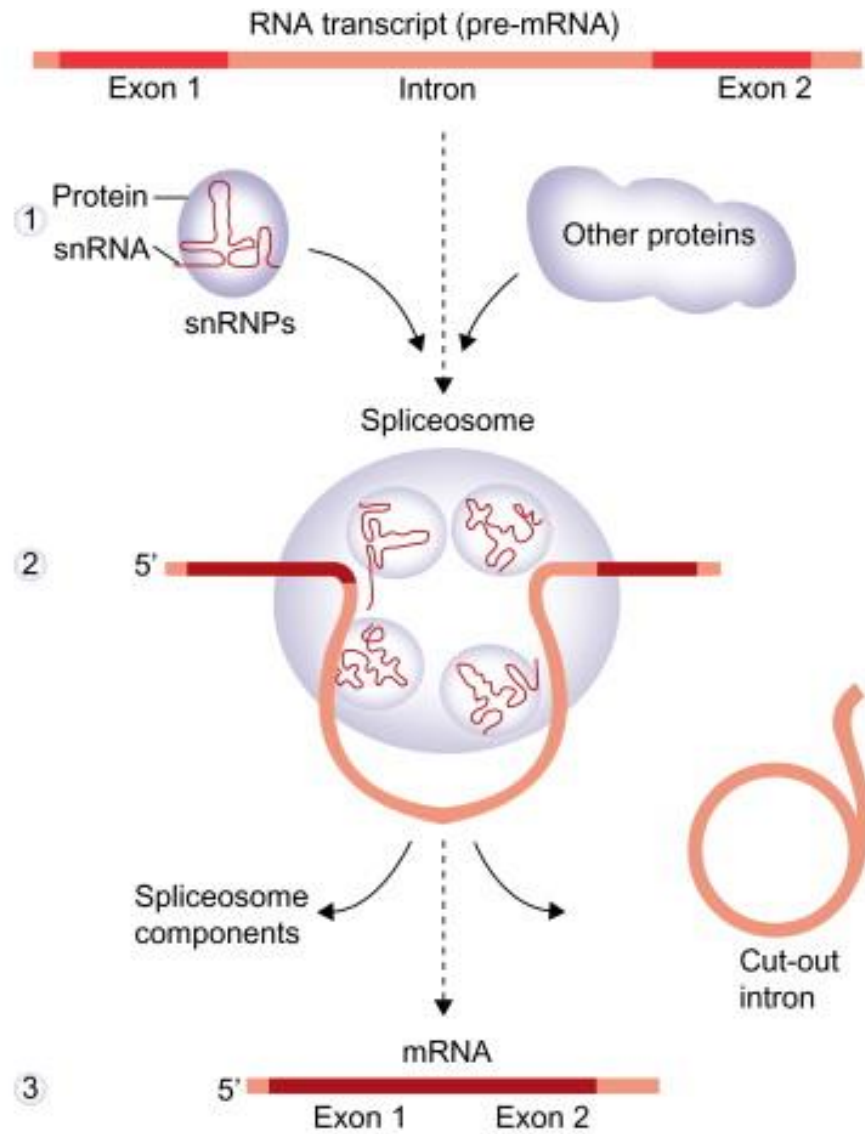
Spliced RNA



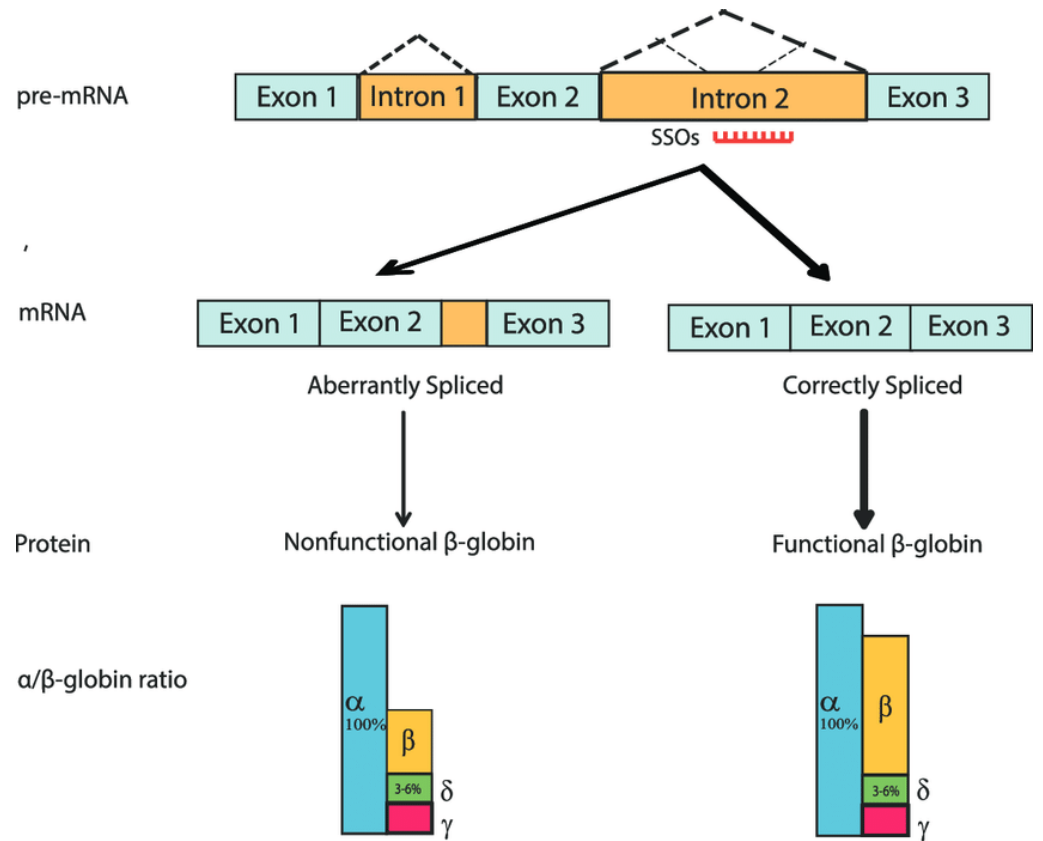
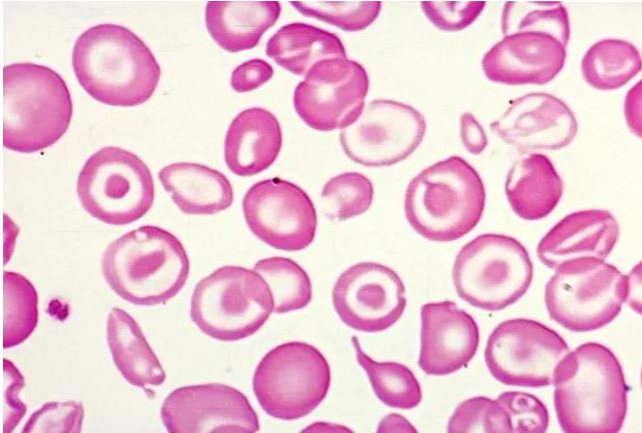
5' untranslated region

3' untranslated region

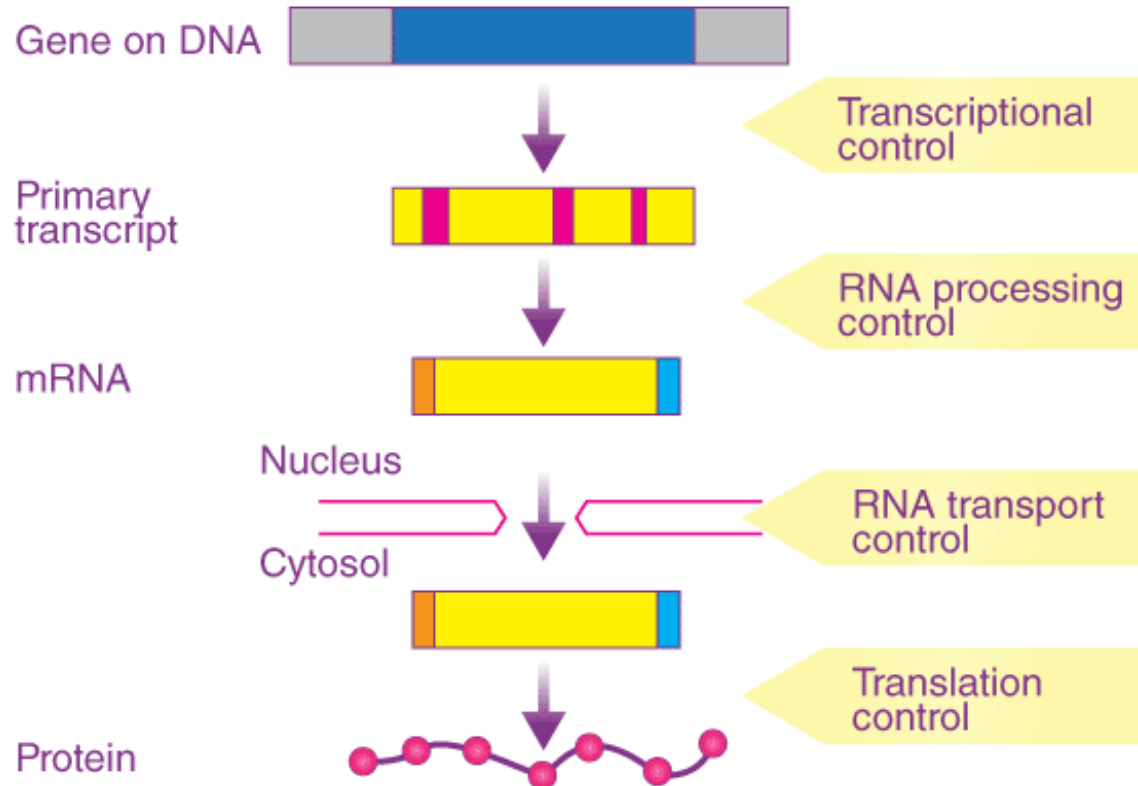
Spliceosome



Thalassemias



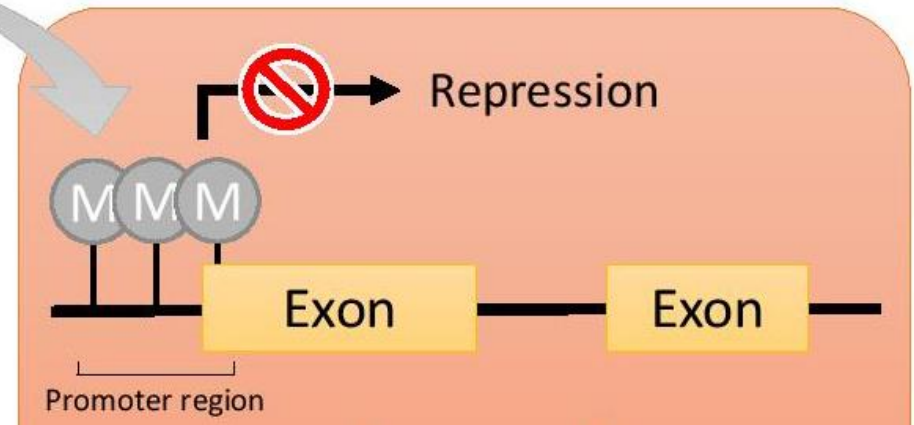
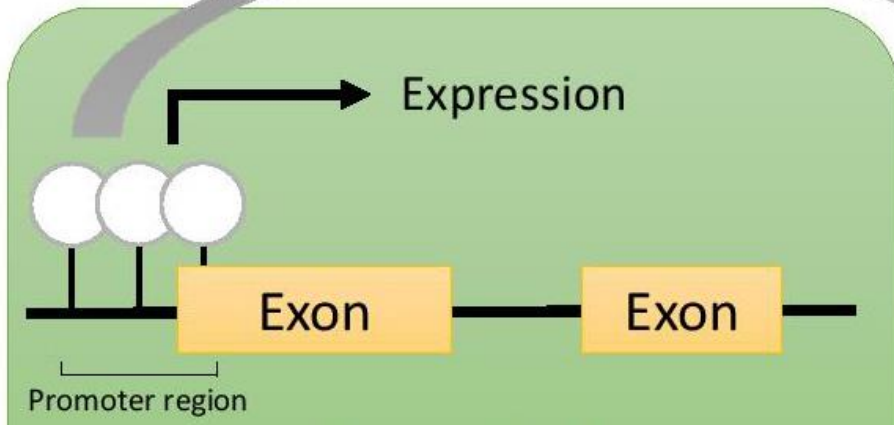
Gene expression regulation



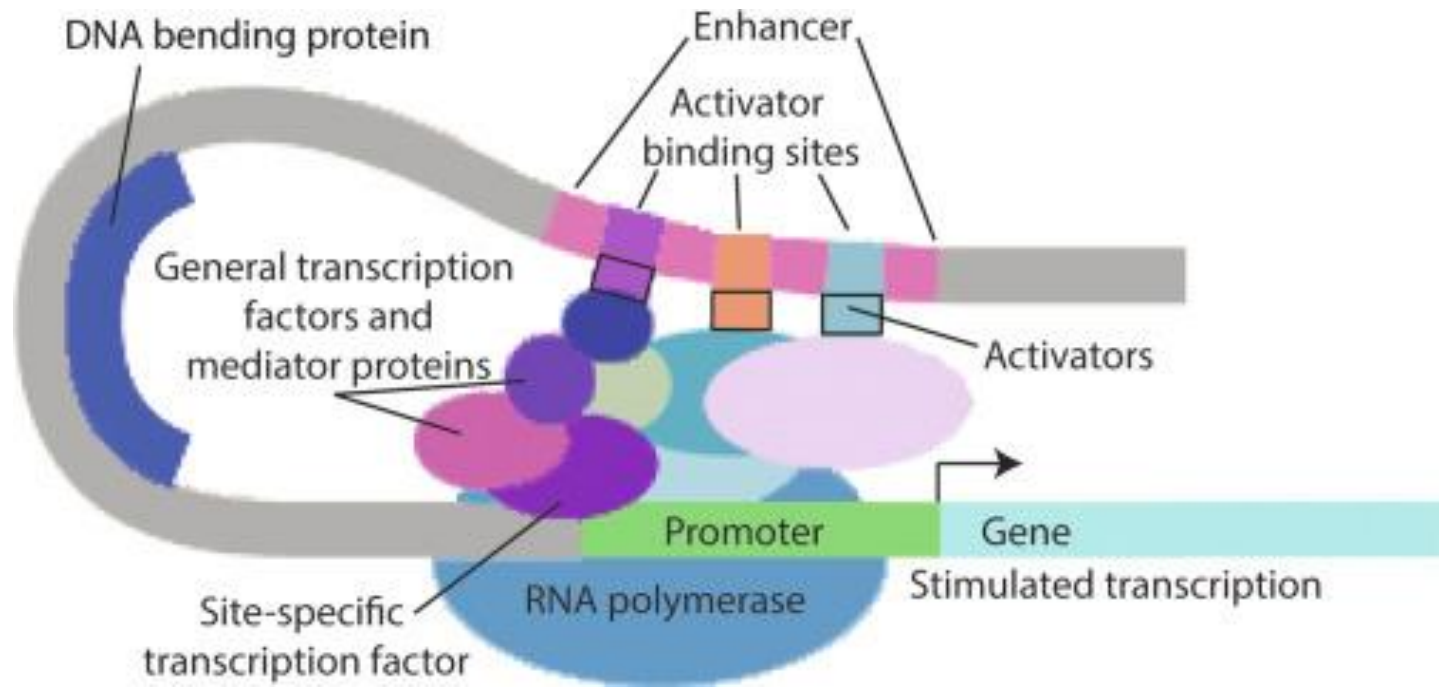
DNA methylation

DNA methylation


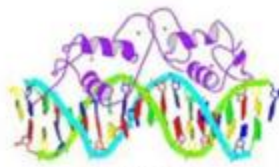

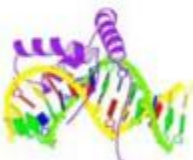





- Unmethylated
- M Methylated



Basal and specific transcription factors



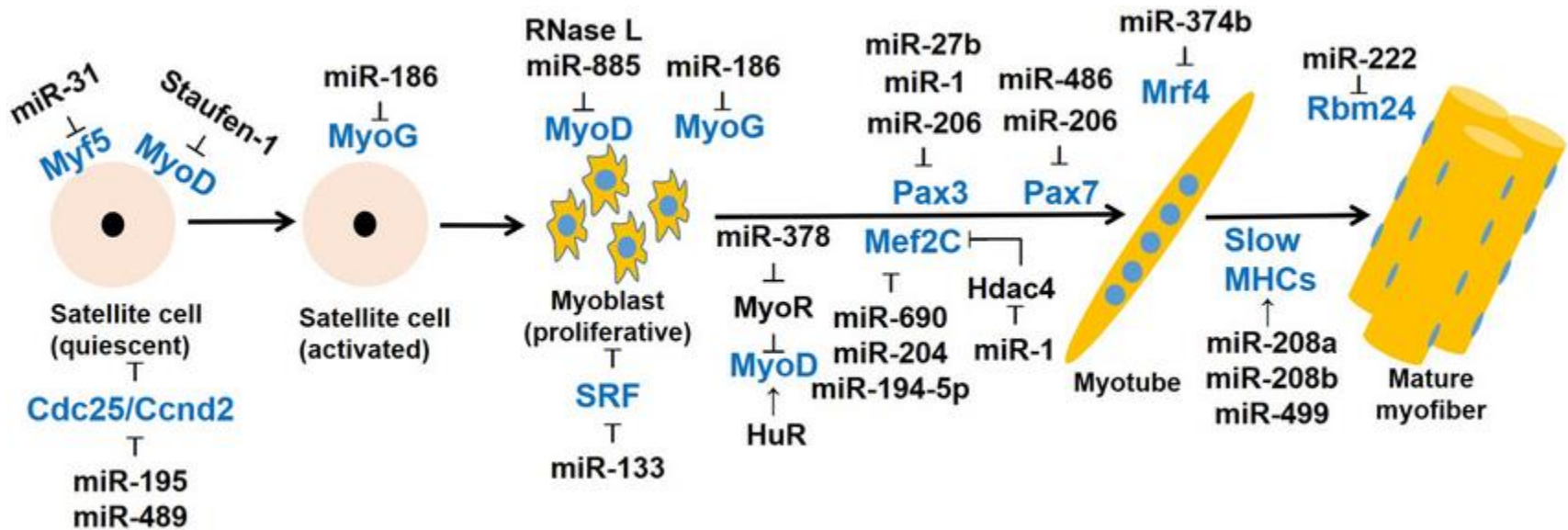
Specific transcription factors

<p>1. Basic domains</p>  <p>CREB (1DH3)</p>	<p>2. Zinc-coordinating DNA-binding domains</p>  <p>GR (1R4R)</p>	<p>3. Helix-turn-helix domains</p>  <p>Pax-6 (6PAX)</p>
<p>4. Other all-α-helical DNA-binding domains</p>  <p>SRY (1J46)</p>	<p>5. α-Helices exposed by β-structures</p>  <p>MEF2A (1C7U)</p>	<p>6. Immunoglobulin fold</p>  <p>NF-kappaB p50 (1SVC)</p>
<p>7. β-Hairpin exposed by an α β-scaffold</p>  <p>SMAD3 (1MHD)</p>	<p>8. β-Sheet binding to DNA</p>  <p>TBP (1CDW)</p>	<p>9. β-Barrel DNA-binding domains</p>  <p>YB-1 (1H95)</p>

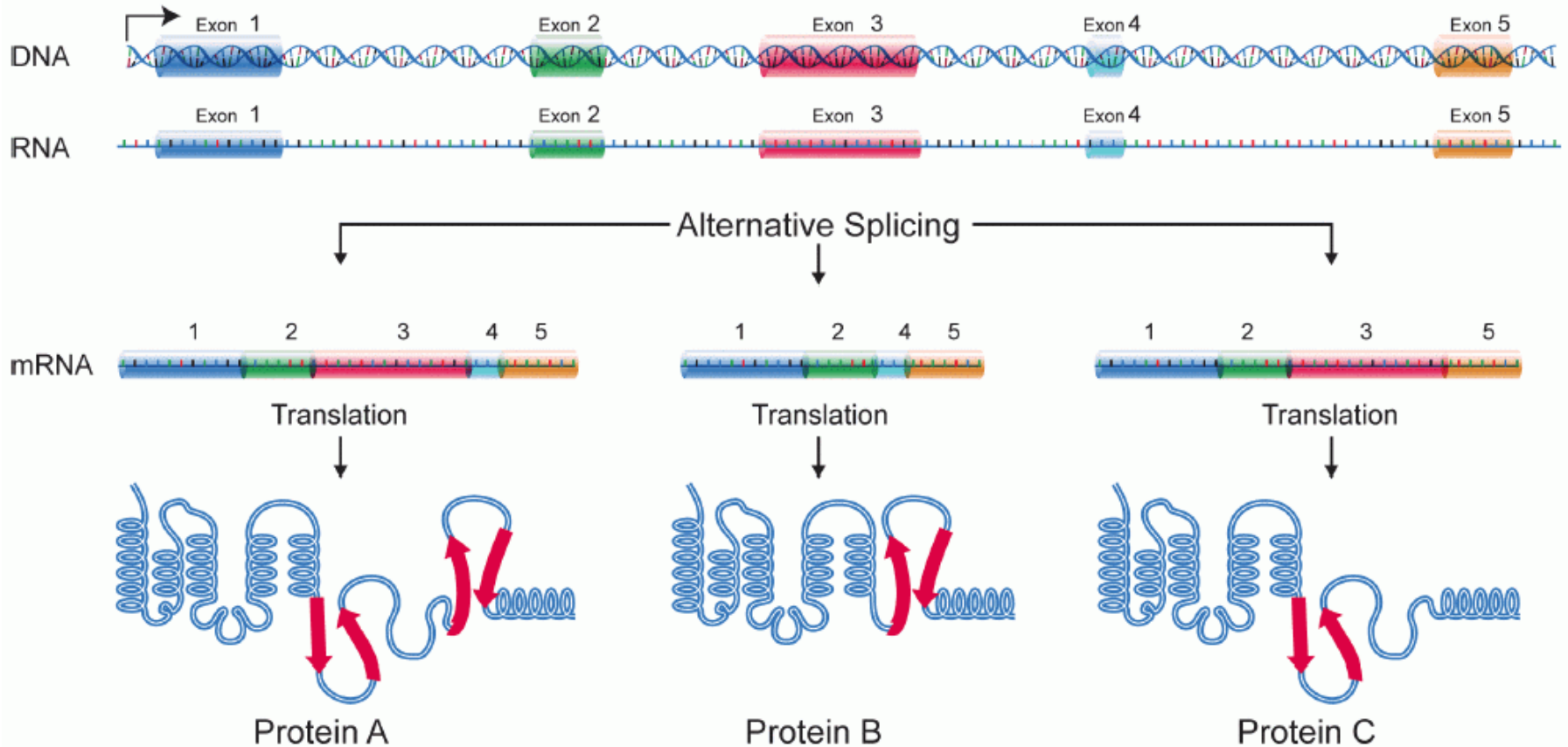
Specific transcription factors

GENE TRANSCRIPTION	
Transcription Factor	Sequence Recognized
Myc and Max	CACGTG
Fos and Jun	TGACTCA
TR (thyroid hormone receptor)	GTGTCAAAGGTCA
MyoD	CAACTGAC
RAR (retinoic acid receptor)	ACGTCATGACCT

Specific transcription factors myoD



Alternative RNA splicing



Alternative RNA splicing - lamins

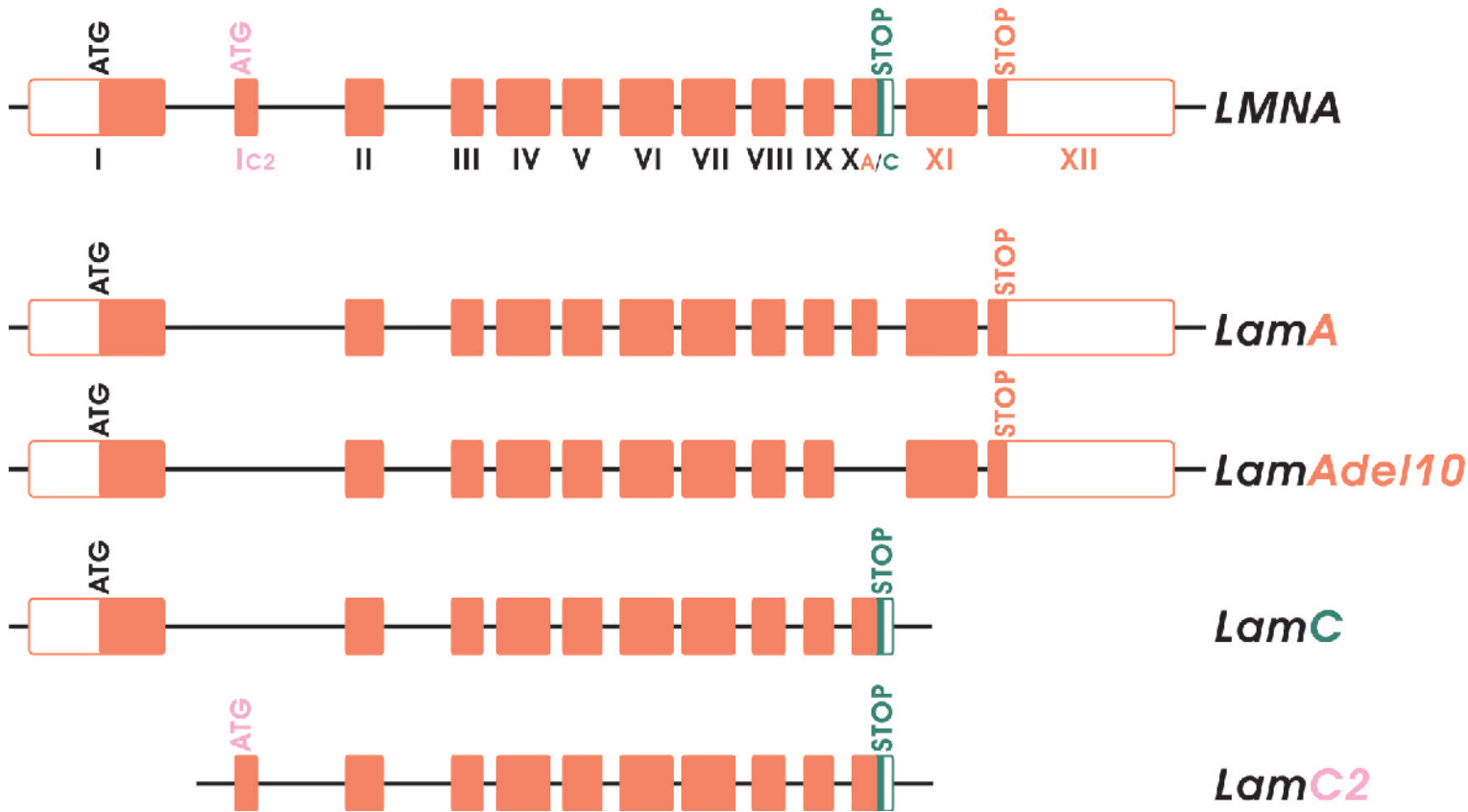
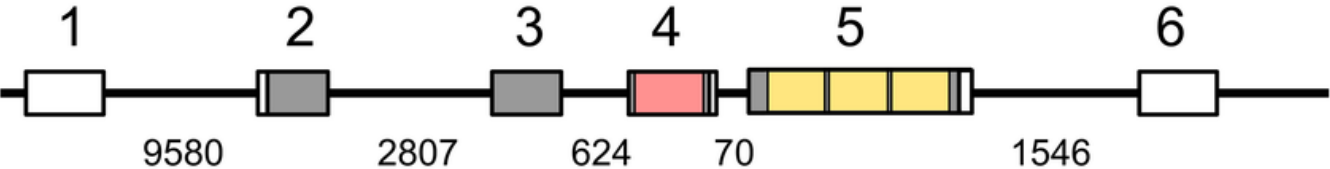


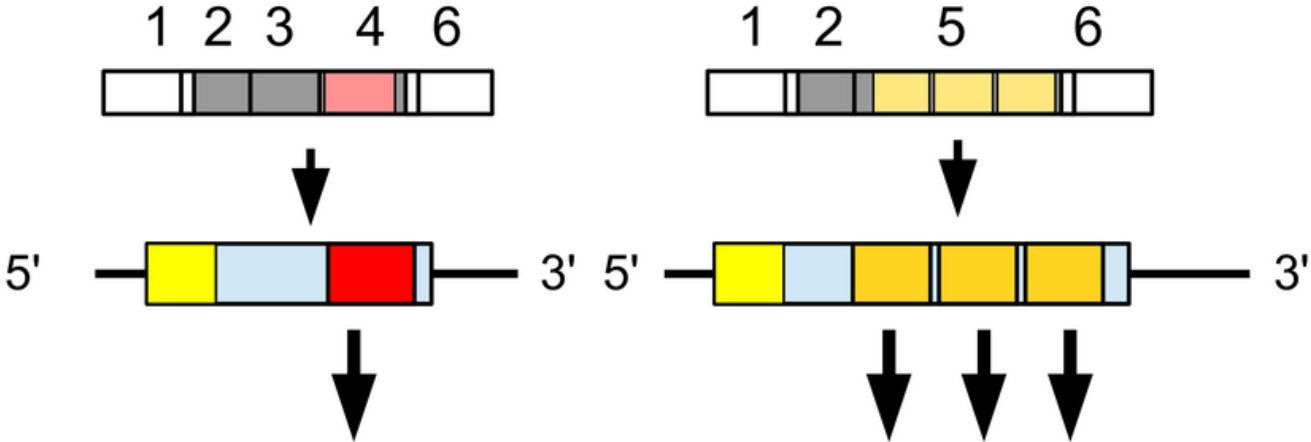
Fig. 1. *LMNA* and A-type lamins. The *LMNA* gene has 12 exons and contains two polyadenylation signals, which are located in exon 10 and 12, and are utilized by lamin C/C₂ and lamin A/A Δ 10 respectively. As a consequence of alternative splicing, the first exon in lamin C₂ is different from other A-type lamins, and lamin A Δ 10 shares identical exons with lamin A but lacks exon 10.

Alternative RNA splicing - calcitonin



mRNA1 : Exons 1-4,6

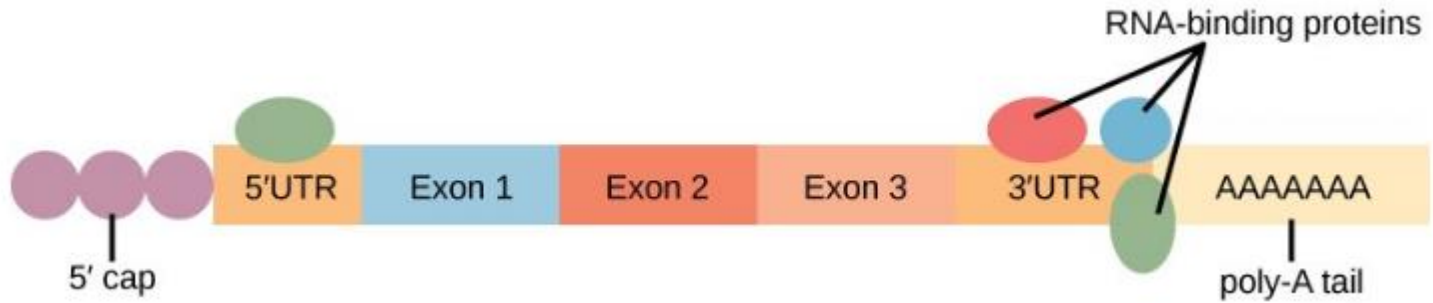
mRNA2 : Exons 1-2,5-6



Calcitonin-A

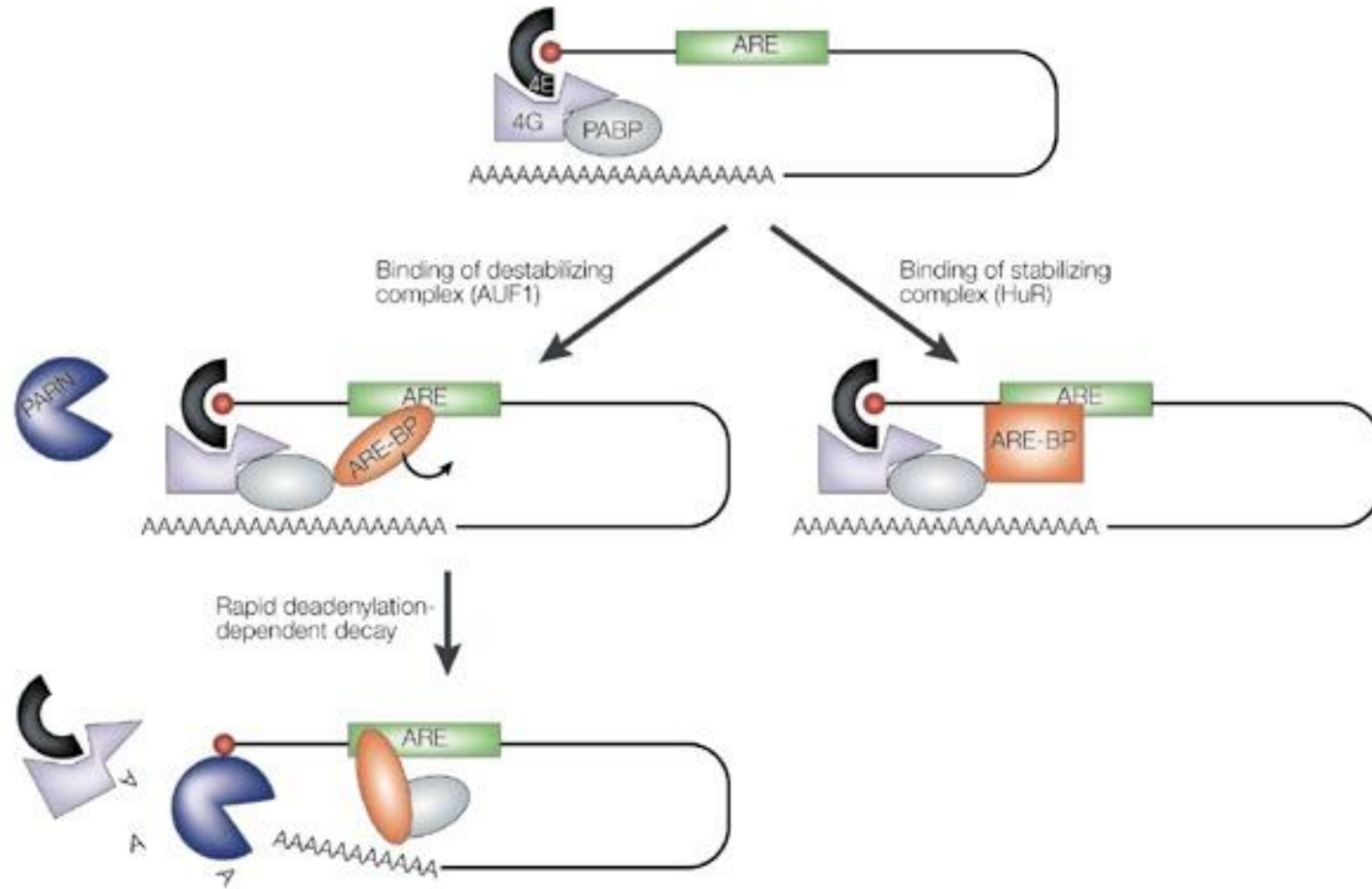
Calcitonin-B1, B2 and B3

Stability control

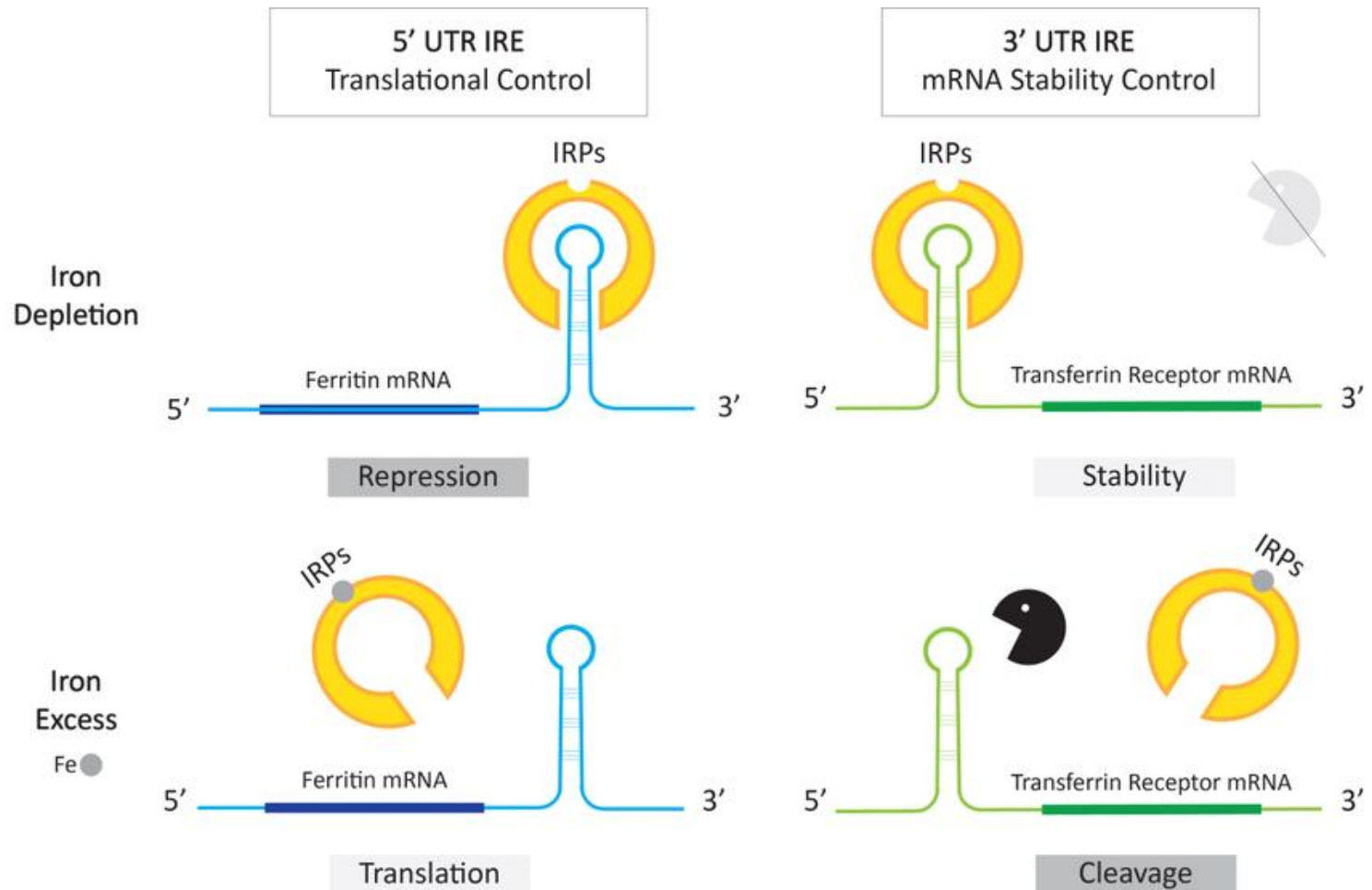


Unstable mRNA – regulatory proteins
Stable mRNA – house keeping proteins

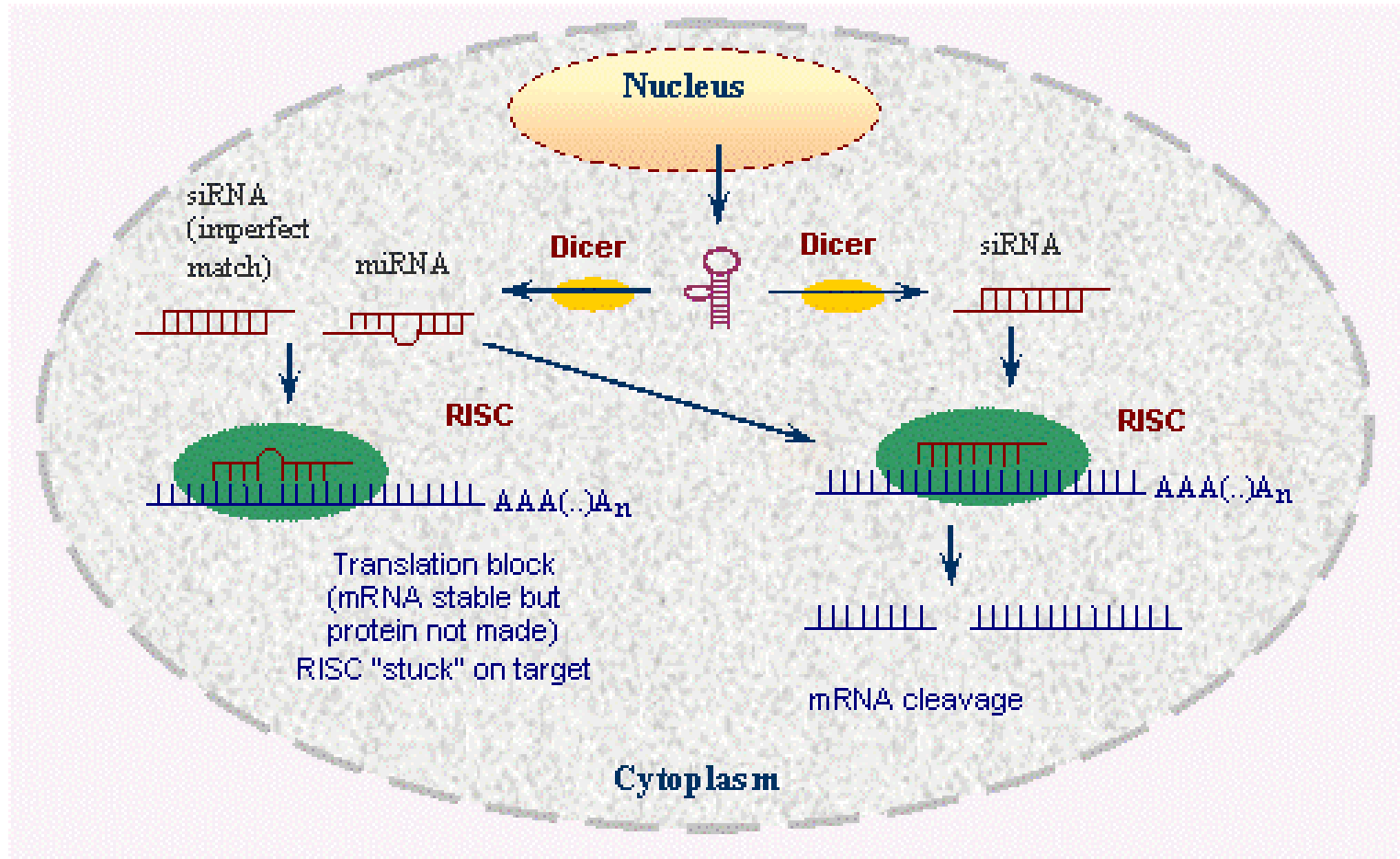
AUUA repeats on 3' end



3' and 5' control – Iron Response Elements IRE



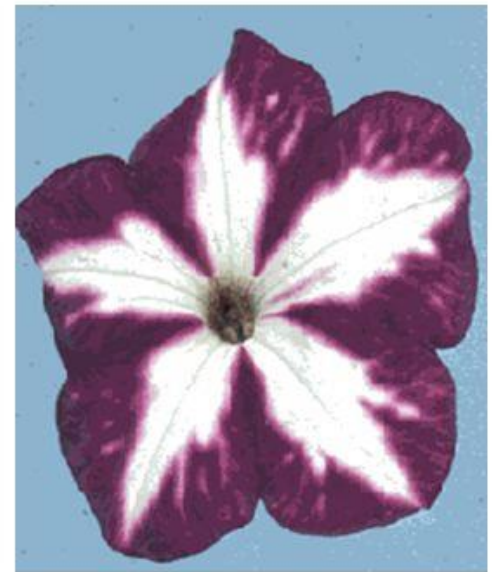
Regulatory RNA



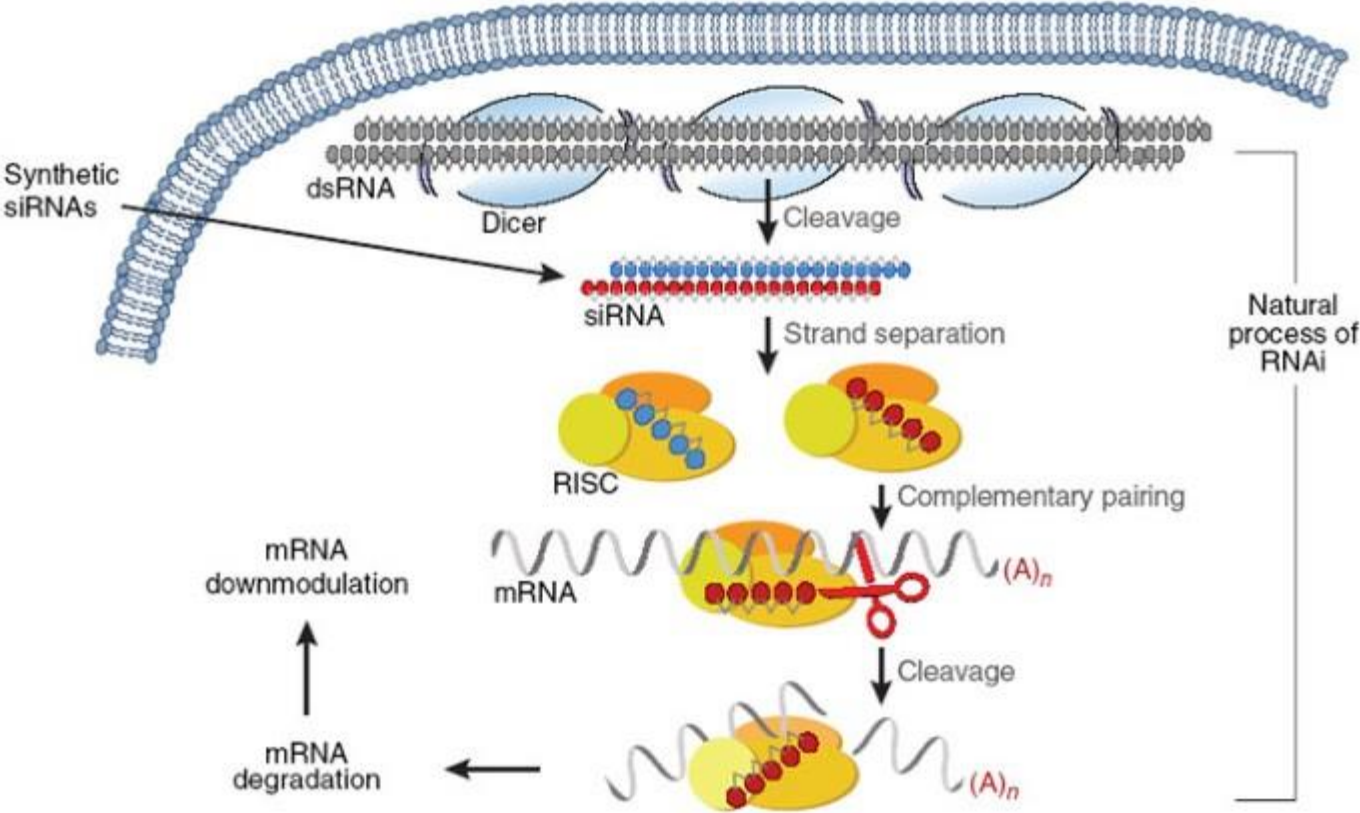
Phenomena first observed in petunia

Attempted to overexpress chalcone synthase (anthocyanin pigment gene) in petunia. (trying to darken flower color)

Caused the loss of pigment.



Antisense nucleotides as a therapeutic agent



Post translational regulation - proteasome

