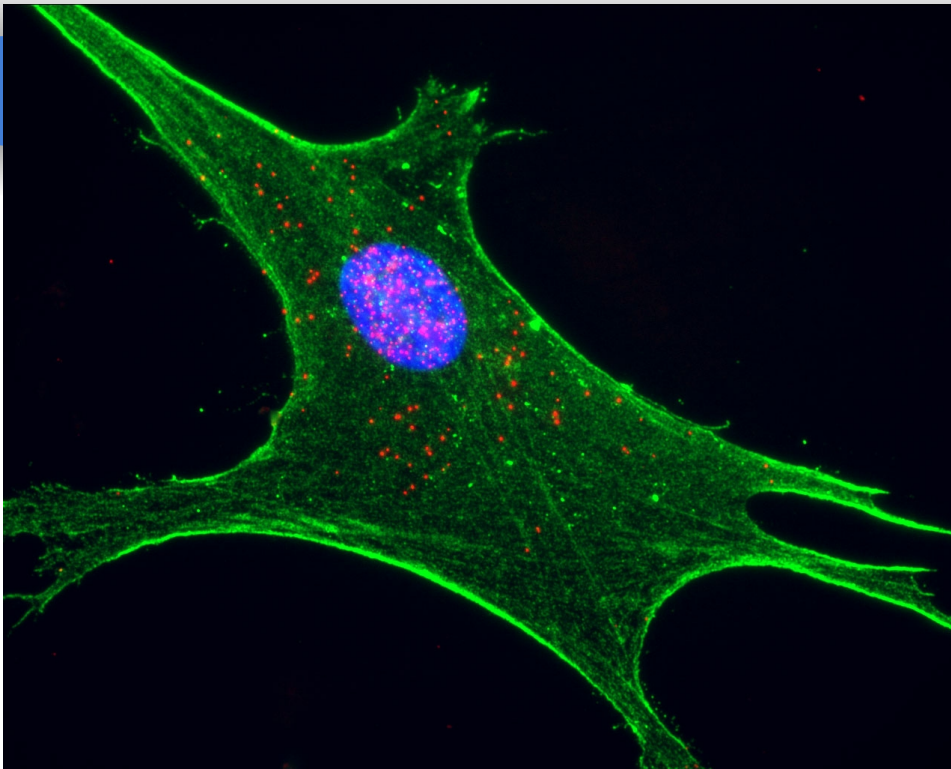
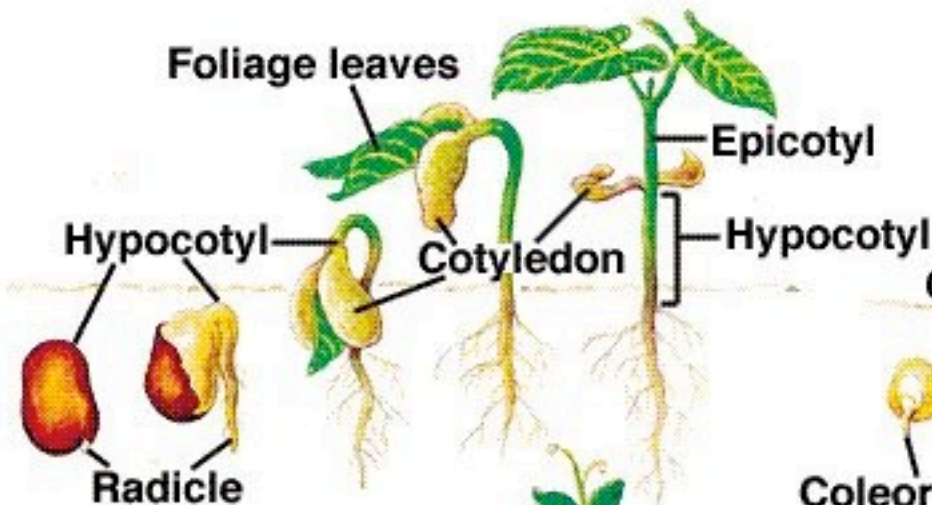


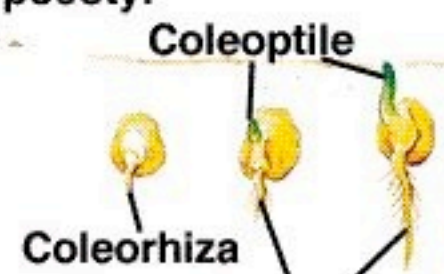
# Control of Cell Function



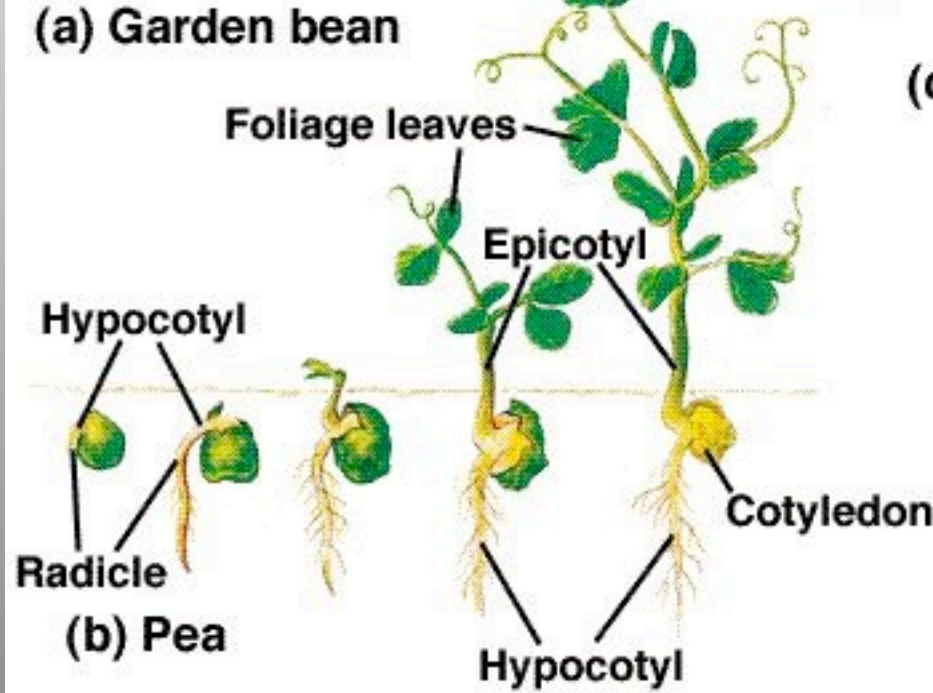
Fluorescent image of  
a cell; nucleus in blue



(a) Garden bean



(c) Corn



(b) Pea

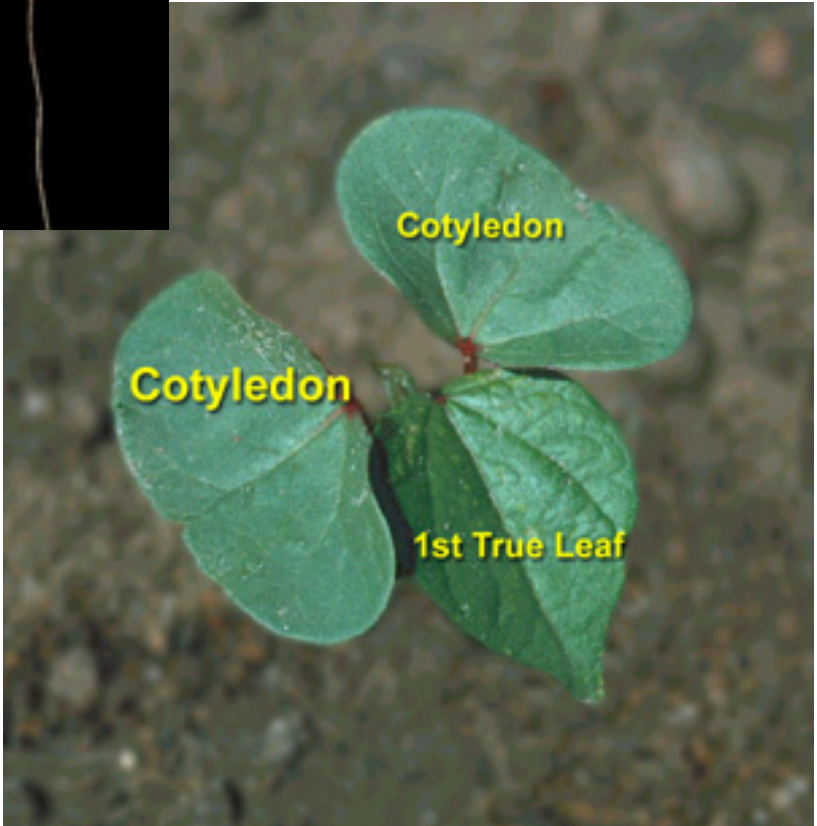
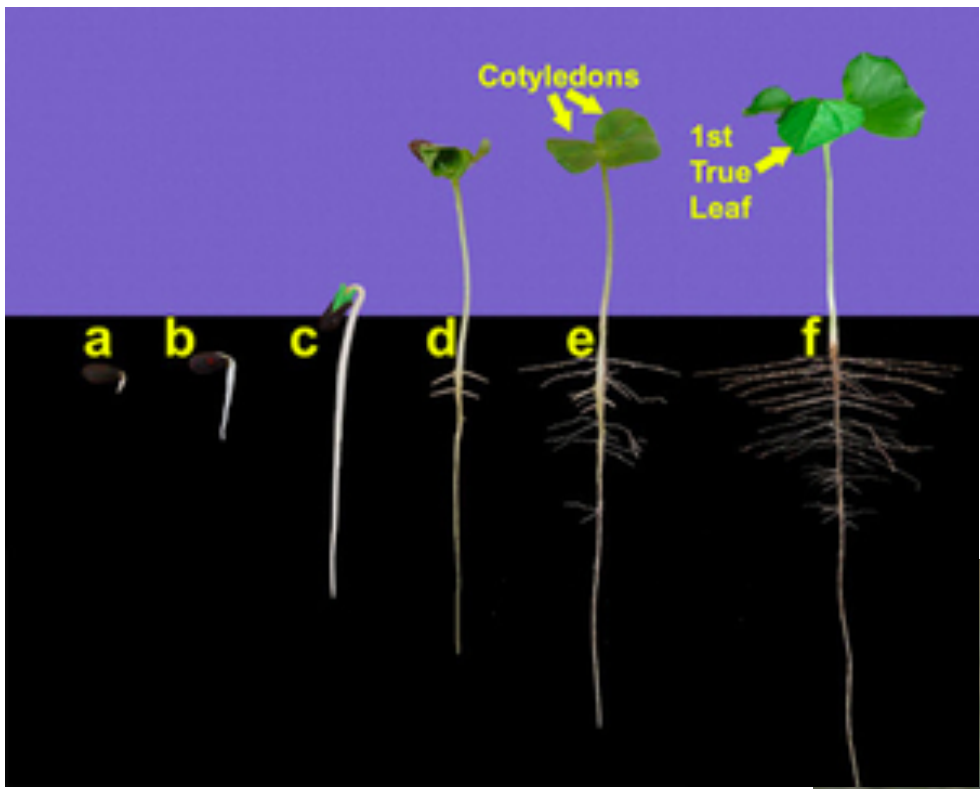
# Seed Germination

# Steps of Germination:

- Imbibition
- Seed swells
- Seed coat ruptures
- Endosperm exposed to oxygen
- Oxygen allows for food stored in endosperm to be converted into energy for growth
- Root grows first
- Stem grows next
- True leaves form
- Cotyledons fall off
- Seedling formed



Radish seed



I. Control of Cell Functions Occurs in the NUCLEOID REGION or NUCLEUS (location of DNA)

A. Chromosomes

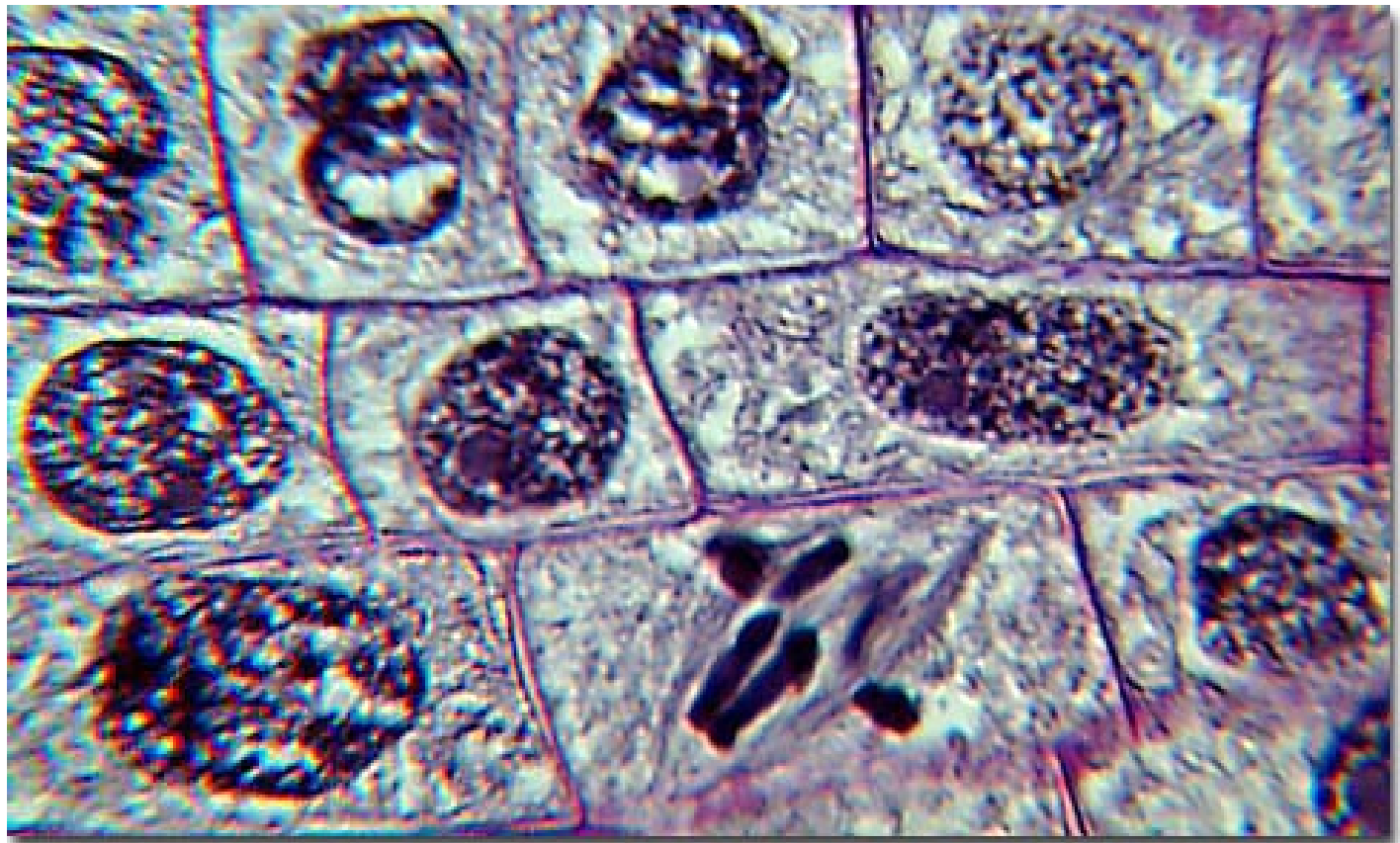
1. Single, circular ones in prokaryotic cells, no packing proteins (naked)

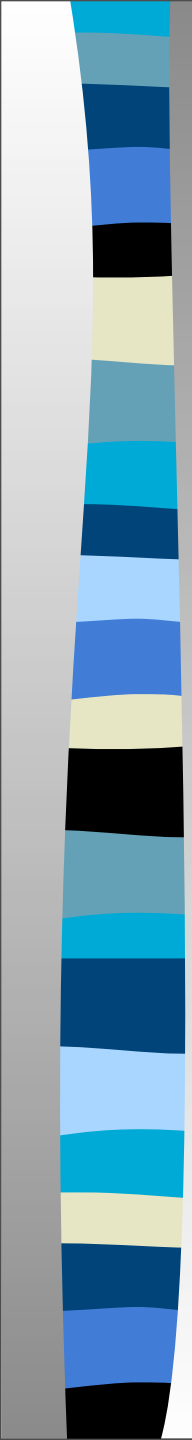
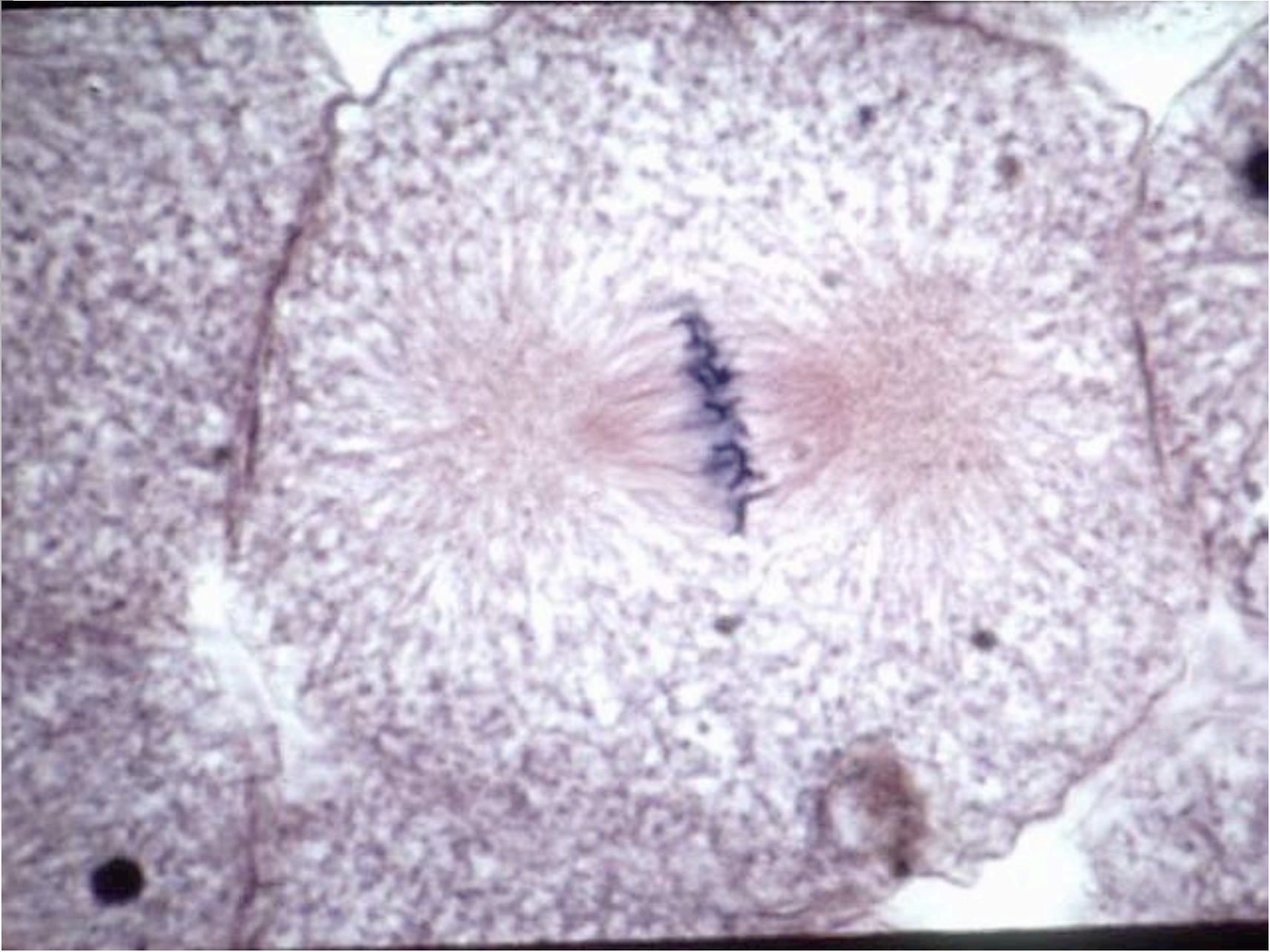


2. Eukaryotic cells: Paired (at least 2 pairs), linear chromosomes.



- Long strands of DNA + packing proteins
- Analogy- thread on a spool
- Can wind up and unwind



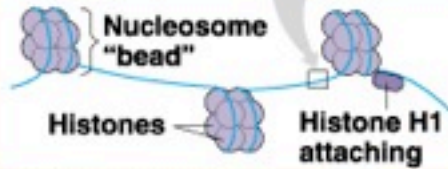






DNA double helix

2 nm

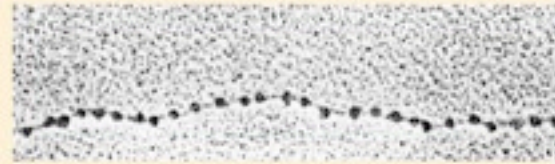


Nucleosome "bead"

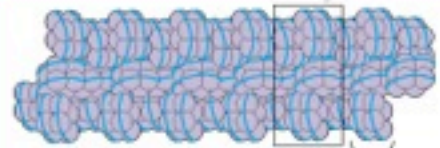
Histones

Histone H1 attaching

11 nm



(a) Nucleosomes ("beads on a string")



Nucleosome

30 nm



(b) 30-nm chromatin fiber



Protein scaffold

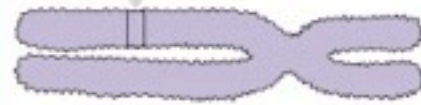
300 nm



(c) Looped domains



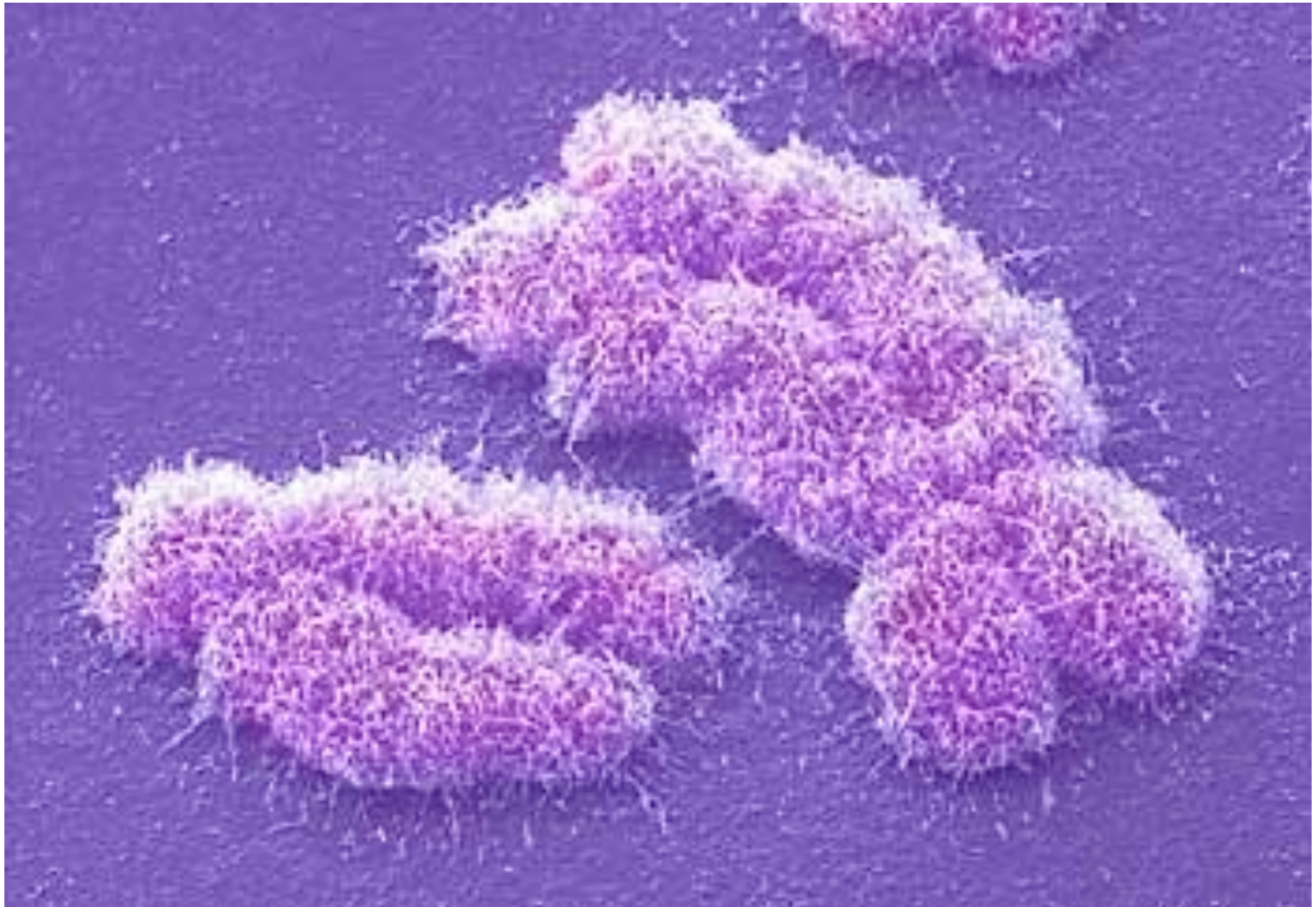
700 nm



1400 nm

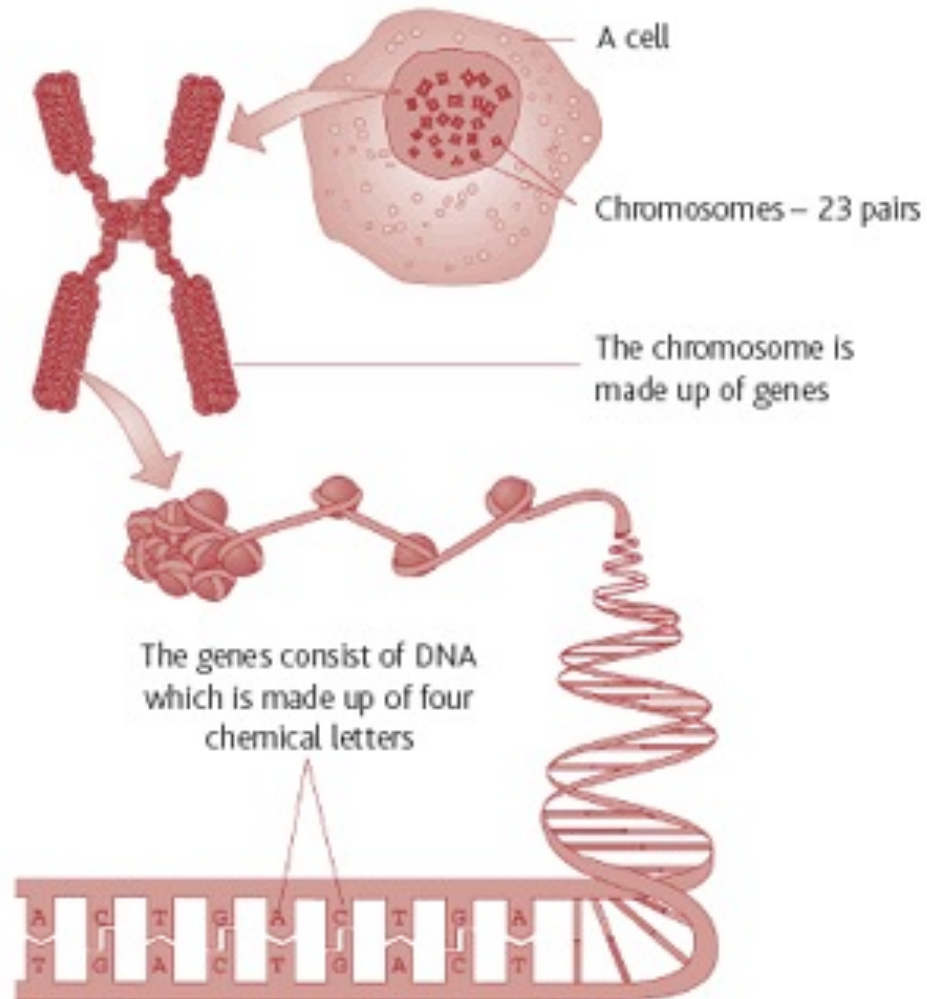


(d) Metaphase chromosome



1400 nm wide = \_\_\_  $\mu\text{m}$  = \_\_\_ mm?

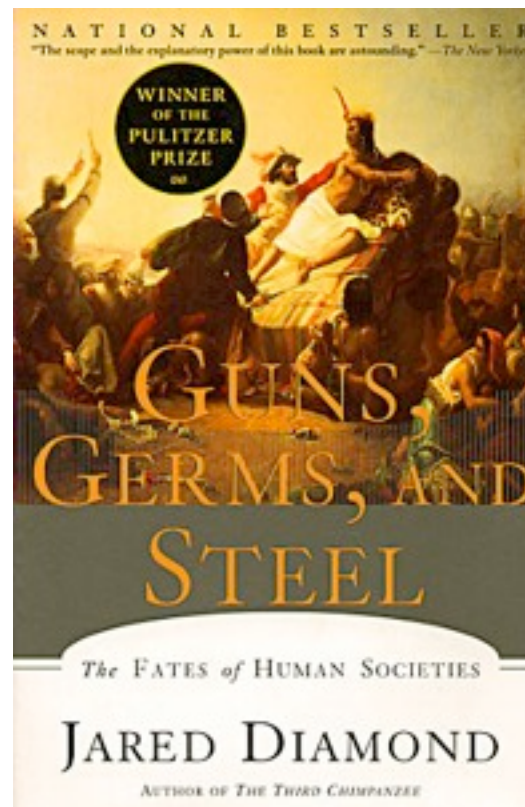
Review: Eukaryotic chromosomes are linear DNA molecules wrapped around proteins.



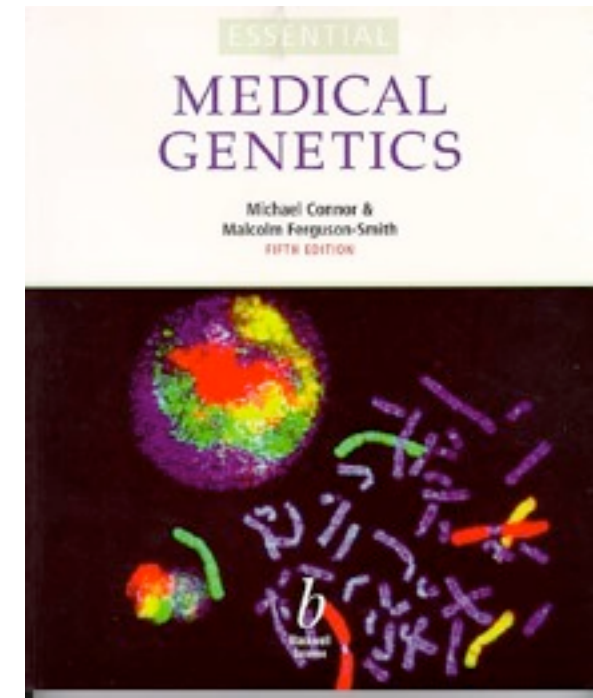
NOTE: A **genome** is the total and unique collection of genetic instructions encoded in the DNA of the chromosomes of each species.

In humans, almost every cell has the complete genome.

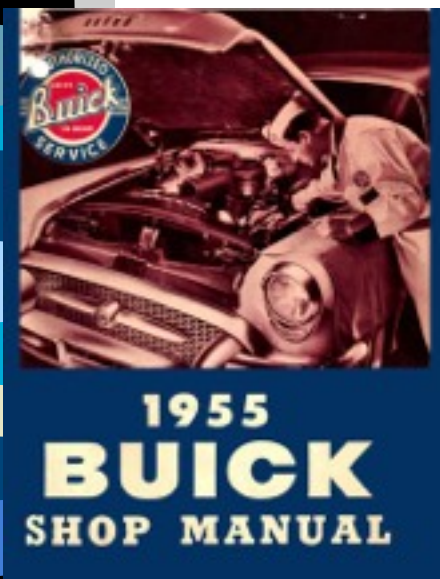
*A history book*



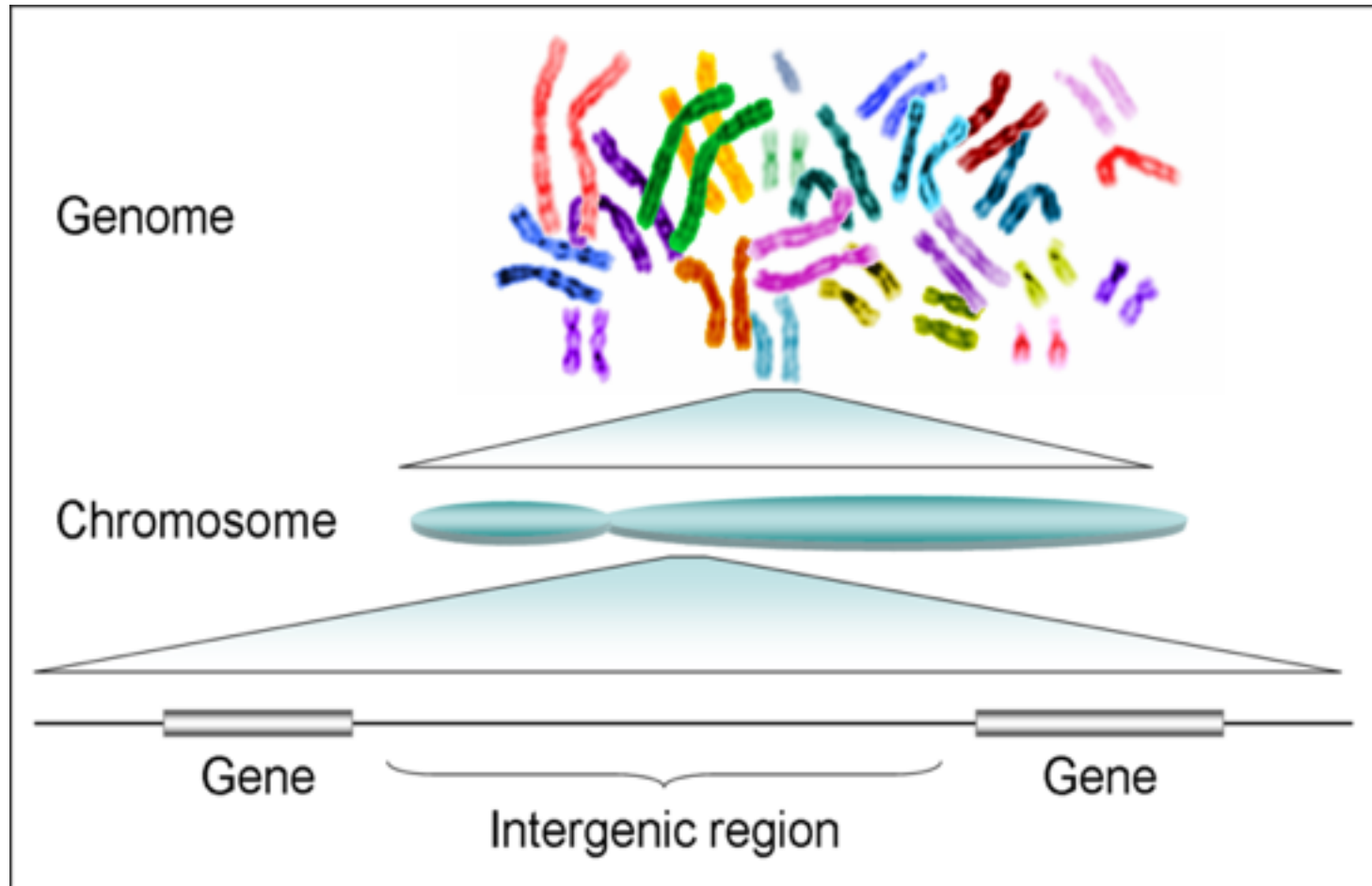
*A medical textbook*

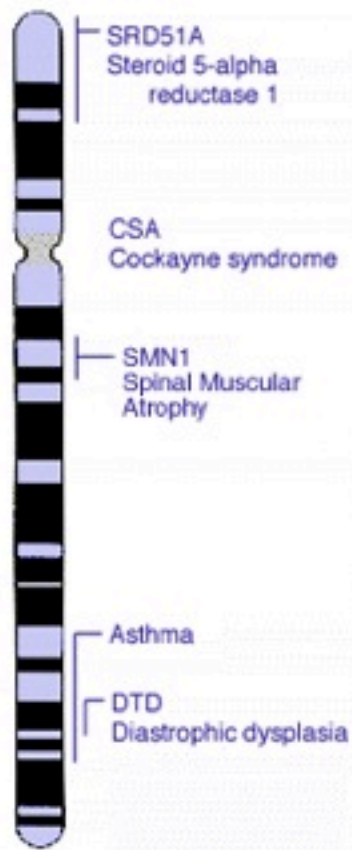


*A shop manual*

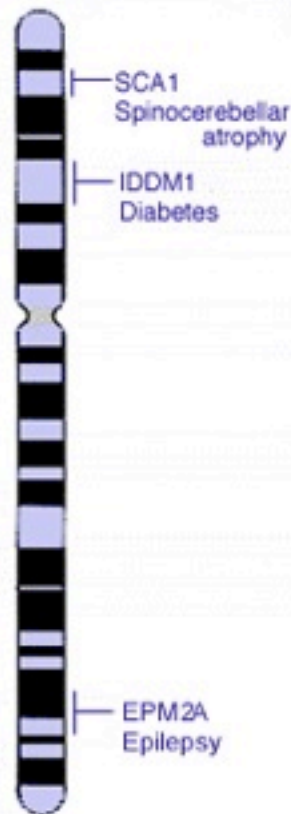


## B. DNA strands have CODING and NON-CODING regions

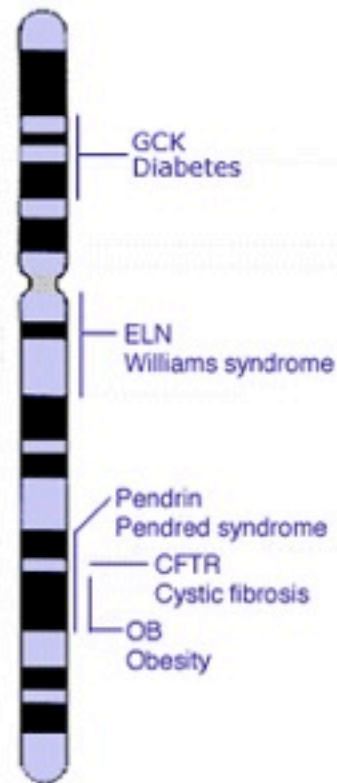




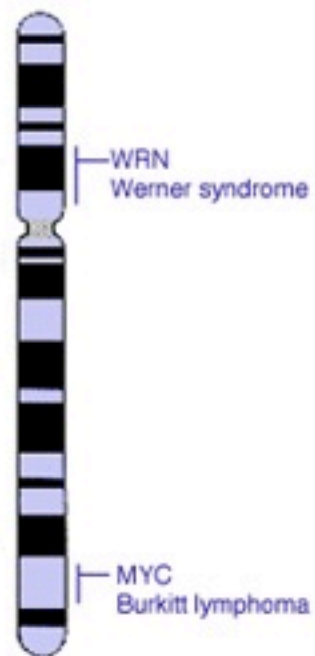
**Chromosome 5**



**Chromosome 6**






**Chromosome 7**

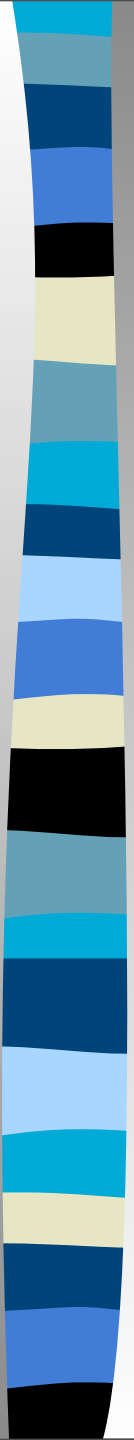


**Chromosome 8**

**Key**

-  centromere
-  rDNA
-  noncentromeric heterochromatin

# 1. Coding regions





# 1. Coding regions

- Make up <10% of DNA





# 1. Coding regions

- Make up <10% of DNA
- Discreet sections are called **genes**



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- Genes code for a cell's proteins → proteins determine an organism's traits and behaviors



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- Humans have an estimated 500-25,000 genes (number keeps falling)

# 1. Coding regions

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## 2. Non-coding Regions



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- Make up >90% of DNA
- Some are repetitive sequences- may play a role in cell longevity and/or chromosome organization



## 2. Non-coding Regions

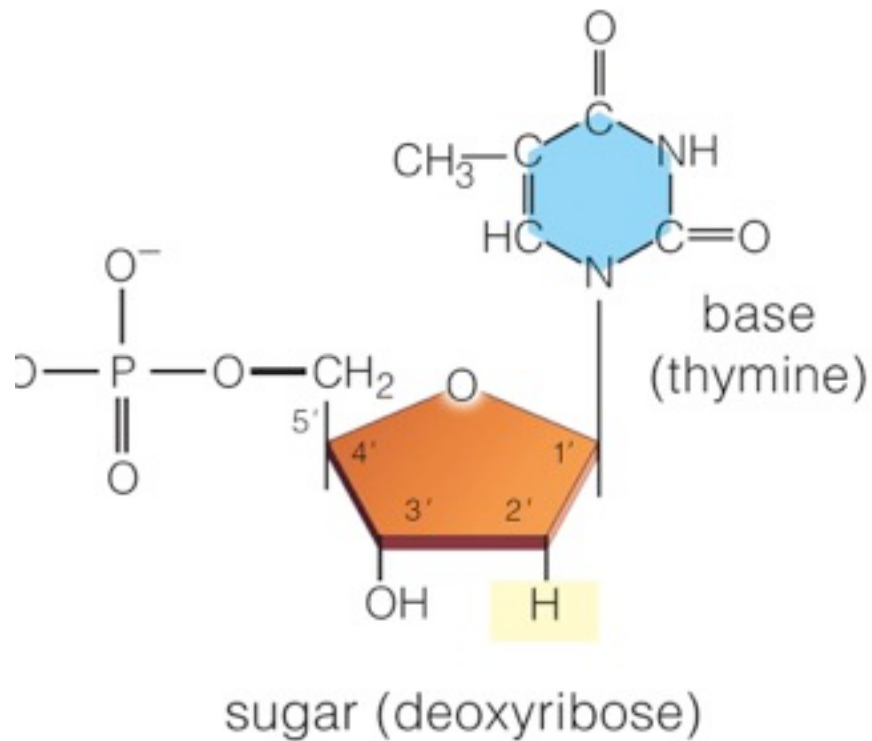
- Make up >90% of DNA
- Some are repetitive sequences- may play a role in cell longevity and/or chromosome organization
- The rest is known as 'junk' or 'nonsense' DNA- may be old viruses, old unused genes ????????



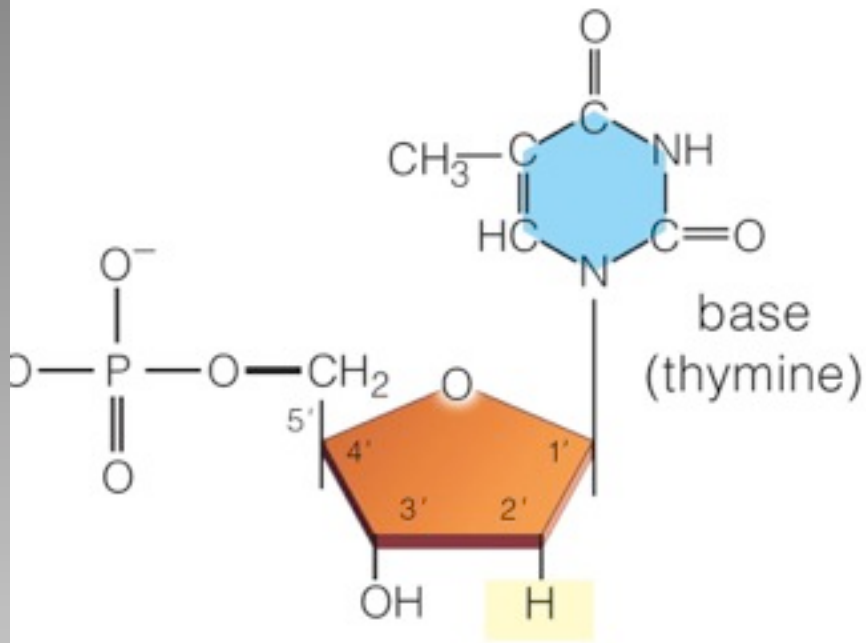
## II. DNA Molecules

### A. Structure

#### 1. Nucleotides

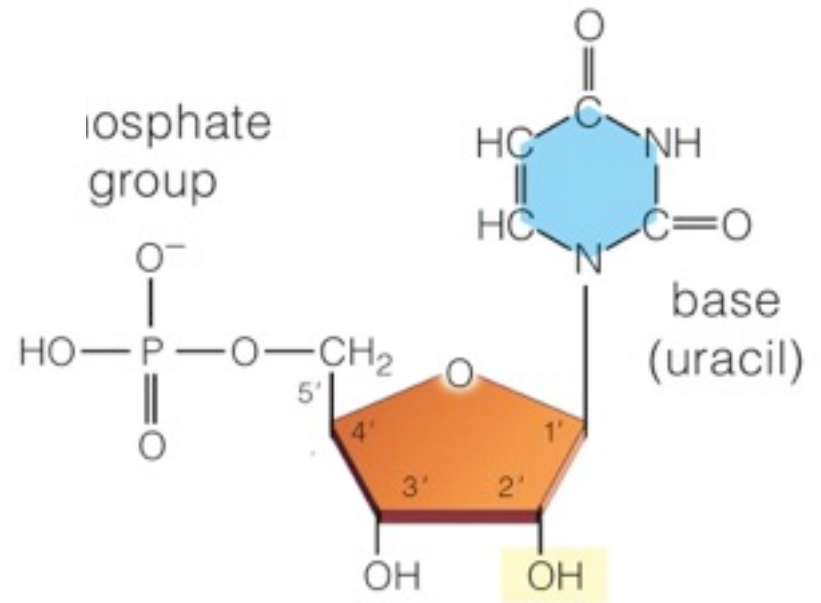


- Phosphate group
- Deoxyribose sugar
- Nitrogen base

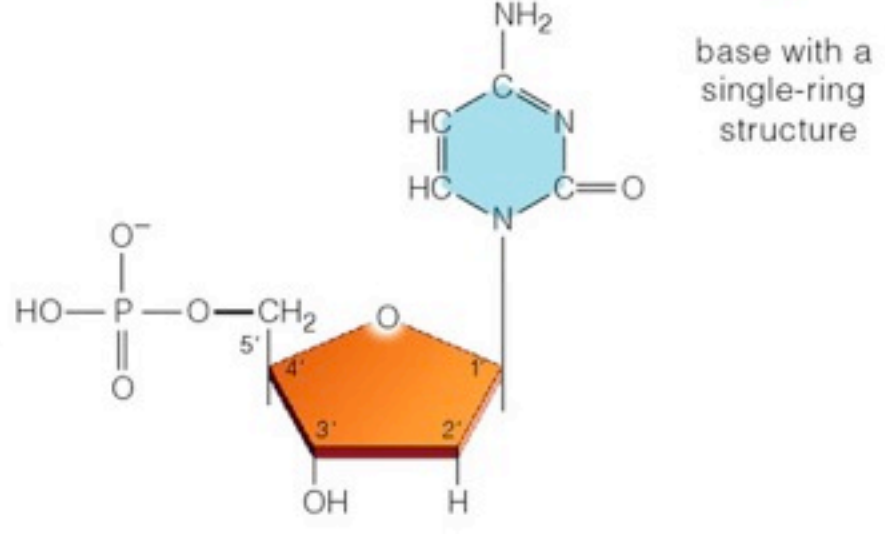
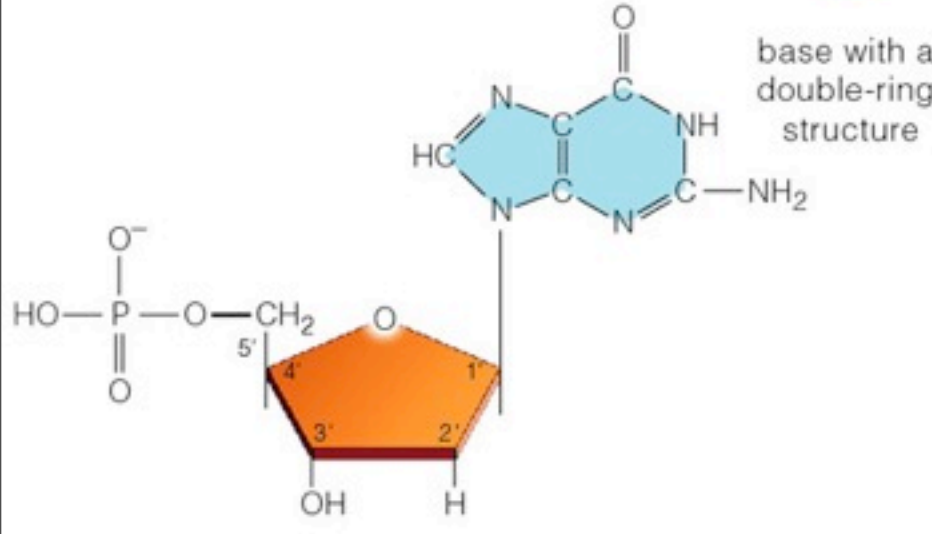
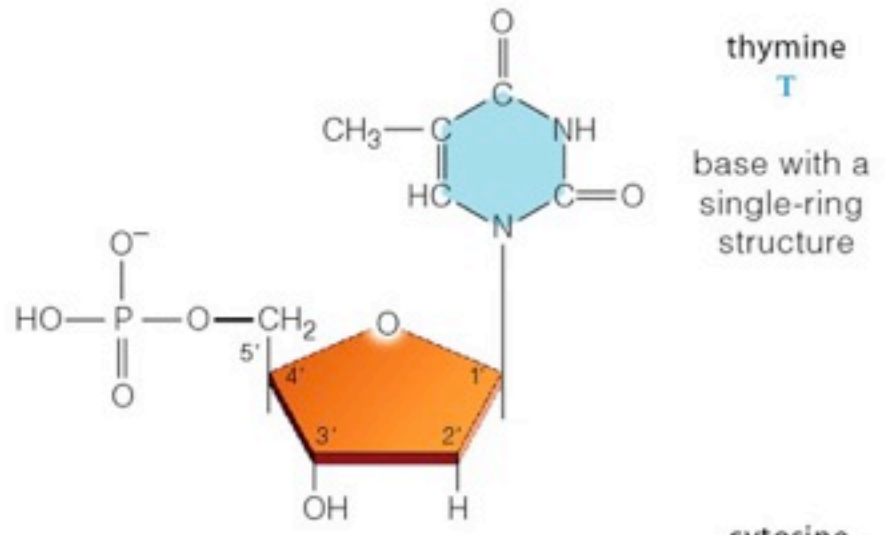
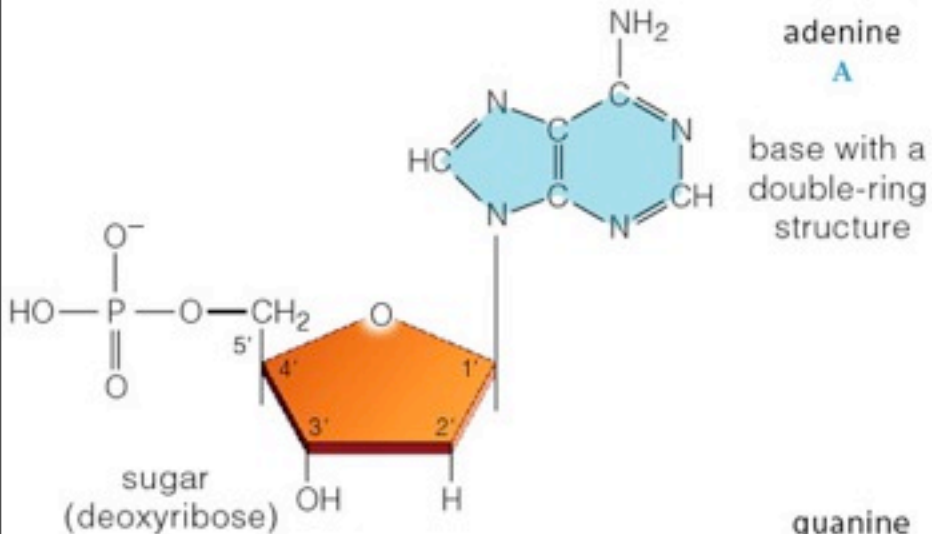


← DNA nucleotide

RNA nucleotide →

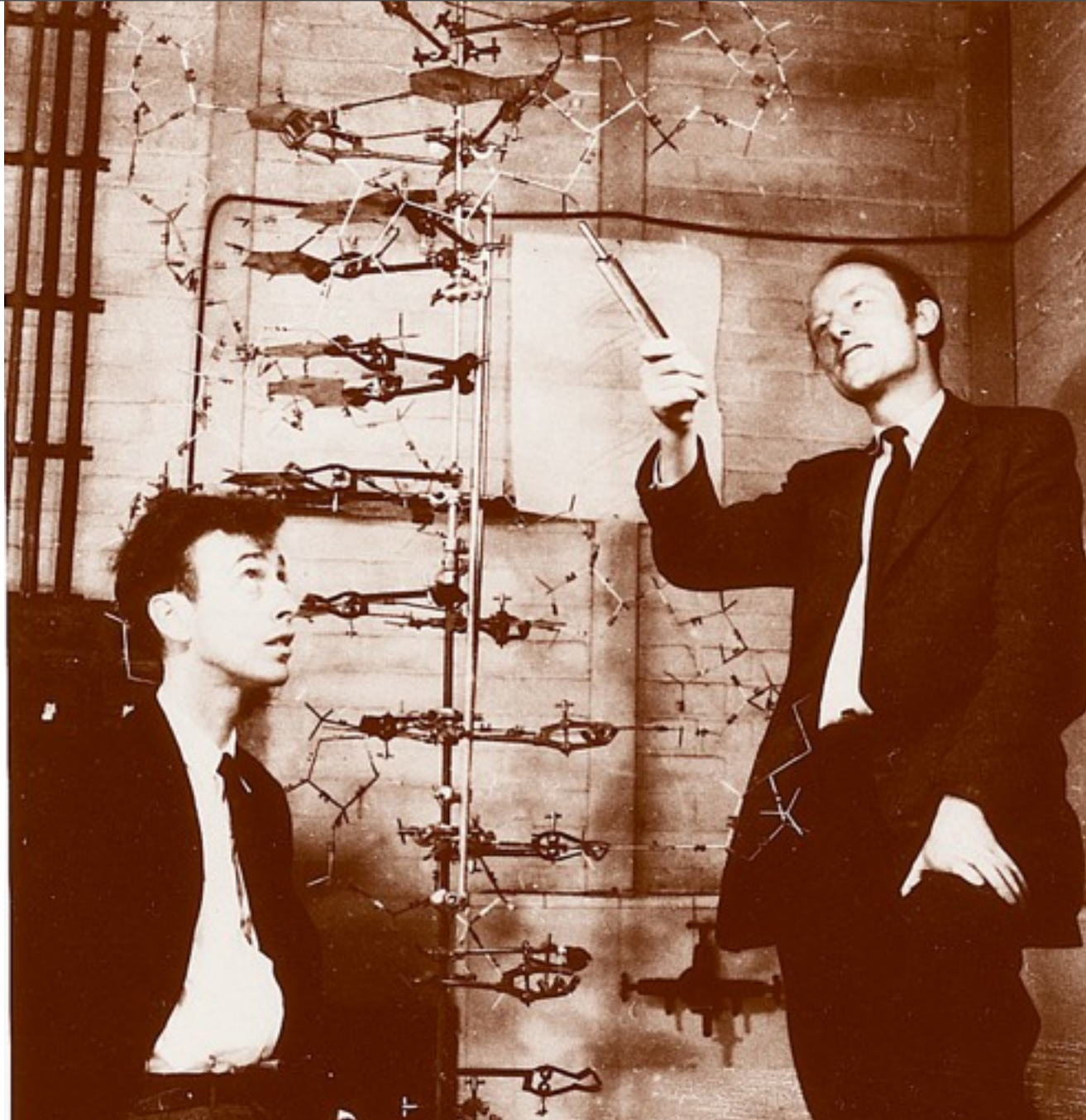


a



Double-ring bases  
A, G

Single-ring bases  
T, C

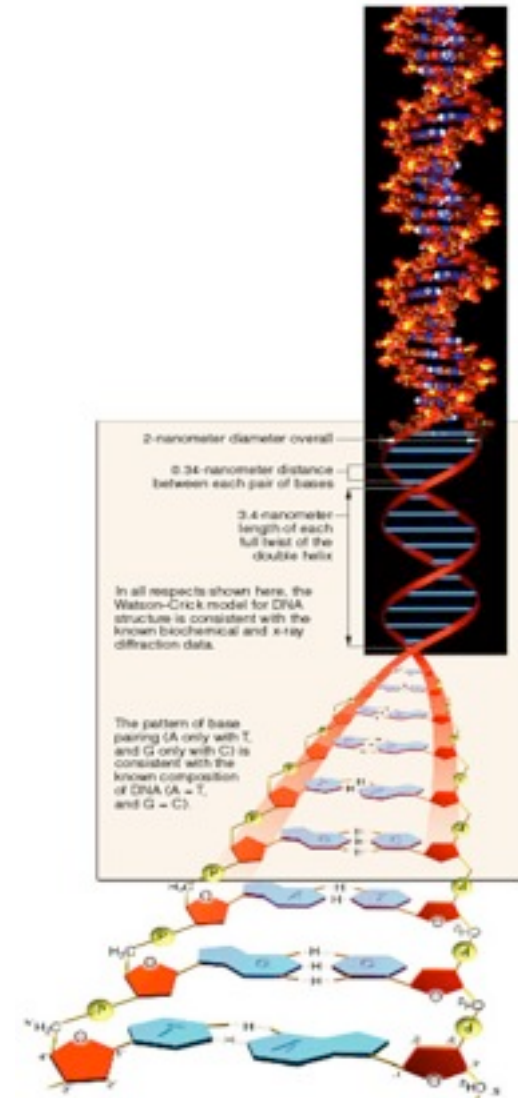


Watson

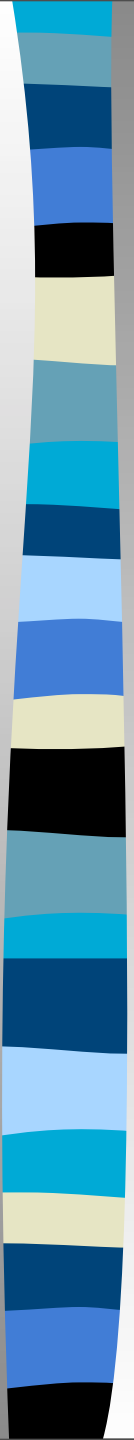
Crick

## 2. Watson-Crick Model

- Double helix
- Sugar-phosphate backbone
- C-G with 3 H bonds
- T-A with 2 H bonds
- Antiparallel



# B. Central Dogma





## B. Central Dogma

■ 1st            1 gene → 1 enzyme



## B. Central Dogma

- 1st            1 gene  $\rightarrow$  1 enzyme
- Next           1 gene  $\rightarrow$  1 protein





## B. Central Dogma

- 1st            1 gene  $\rightarrow$  1 enzyme
- Next           1 gene  $\rightarrow$  1 protein
- Recent        1 gene  $\rightarrow$  1 polypeptide



## B. Central Dogma

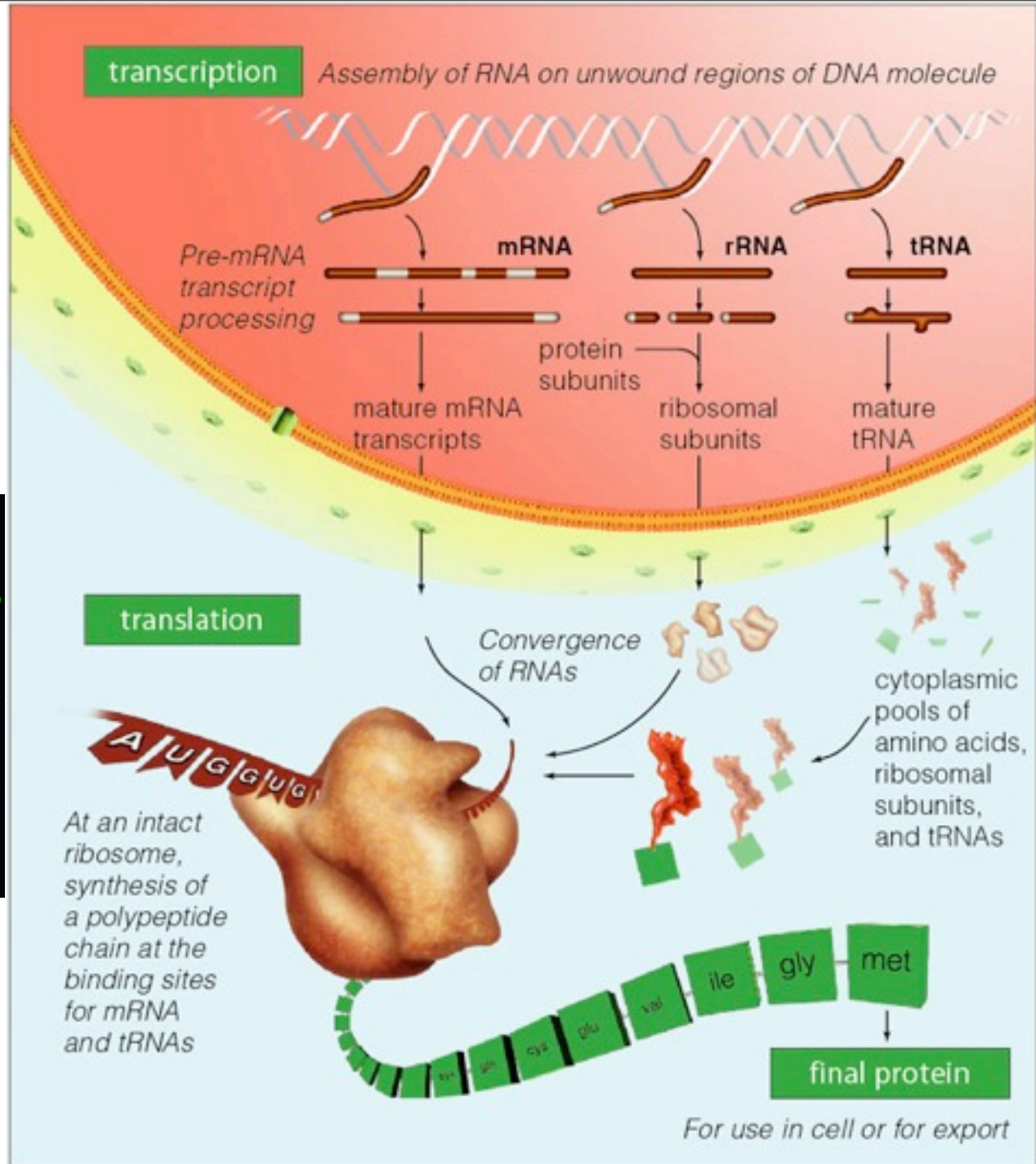
- 1st            1 gene → 1 enzyme
- Next           1 gene → 1 protein
- Recent        1 gene → 1 polypeptide
- Now            1 gene → 1-10 polypeptides



## B. Central Dogma

- 1st            1 gene → 1 enzyme
- Next           1 gene → 1 protein
- Recent        1 gene → 1 polypeptide
- Now            1 gene → 1-10 polypeptides
- Humans have 25,000 genes or less and make about 100,000 different proteins. How?

# OVERVIEW





# -RNA

- A. Solves 2 problems for DNA
  1. Getting information from nucleus to RER or cytoplasm.
  2. Translating from the language of nucleotides (4 bases) to the language of proteins (20 amino acids).



# -RNA

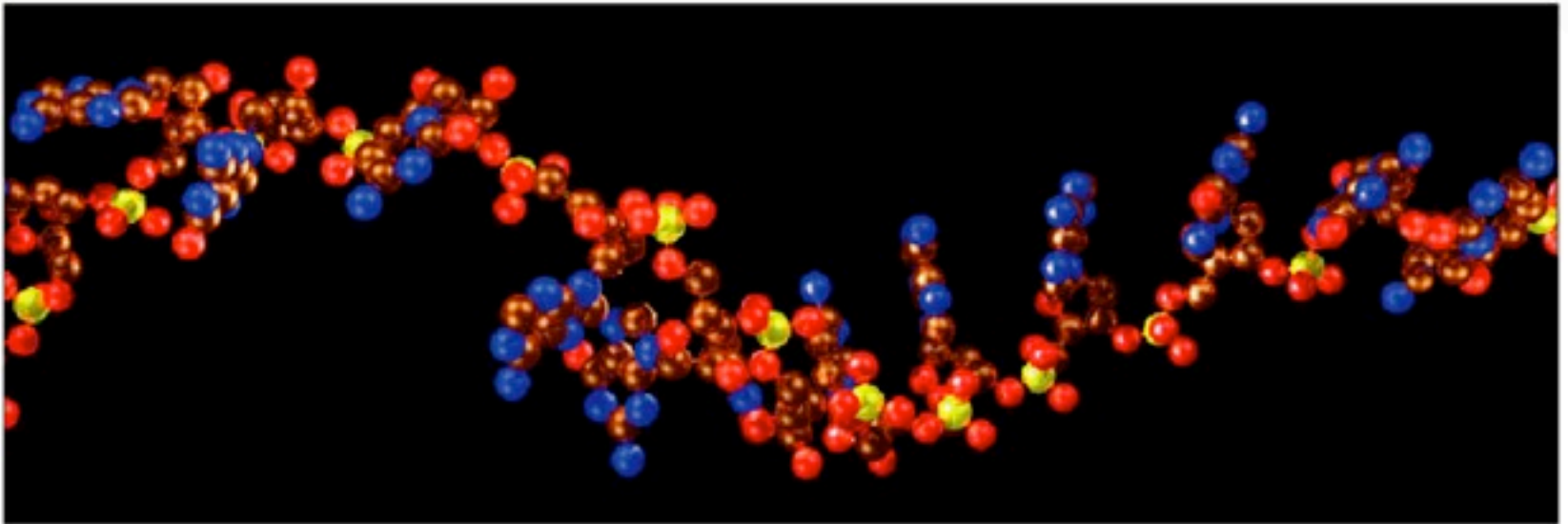
A. Solves 2 problems for DNA

1. Getting information from nucleus to RER or cytoplasm.
2. Translating from the language of nucleotides (4 bases) to the language of proteins (20 amino acids).

B. DNA vs. RNA

DNA	RNA
Deoxyribose- more stable	Ribose- less stable
Double-stranded	Mostly single-stranded
Nucleus or nucleoid region	Nucleus and cytoplasm
Thymine	Uracil- single ring

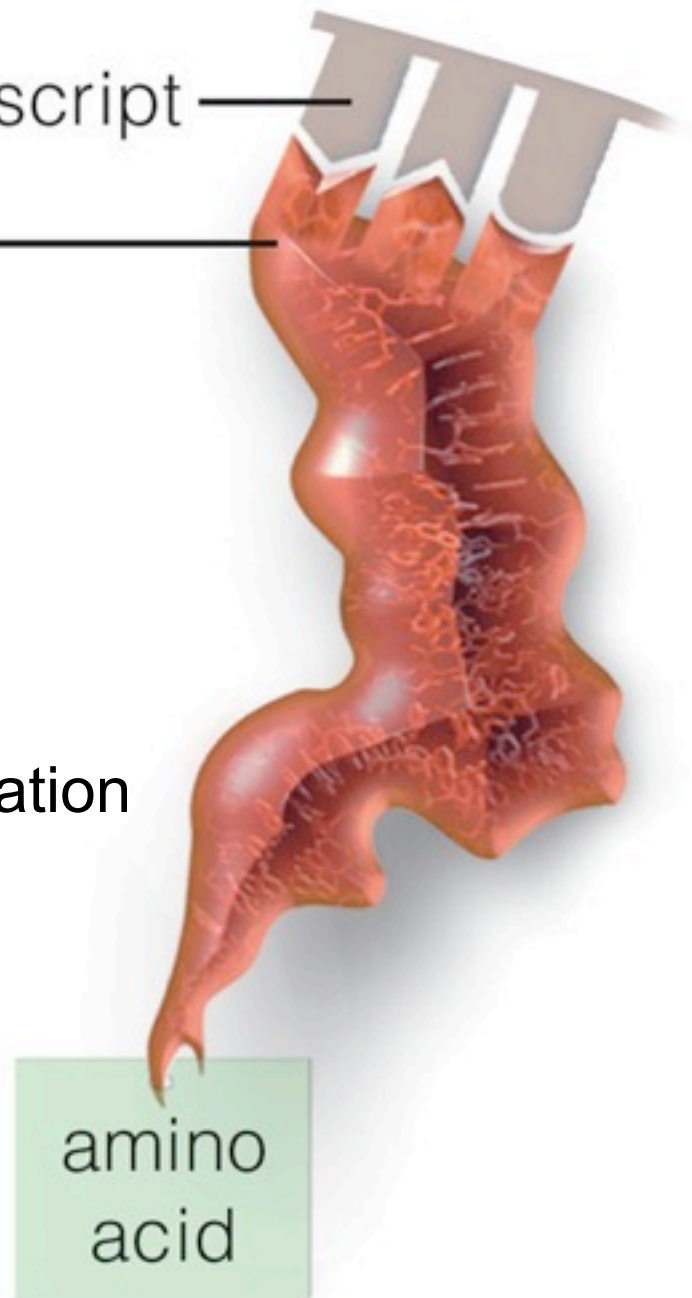
- C. Types and Shapes and Functions
1. mRNA = messenger RNA  
contains the code for amino acid  
sequences in polypeptides



codon in mRNA transcript

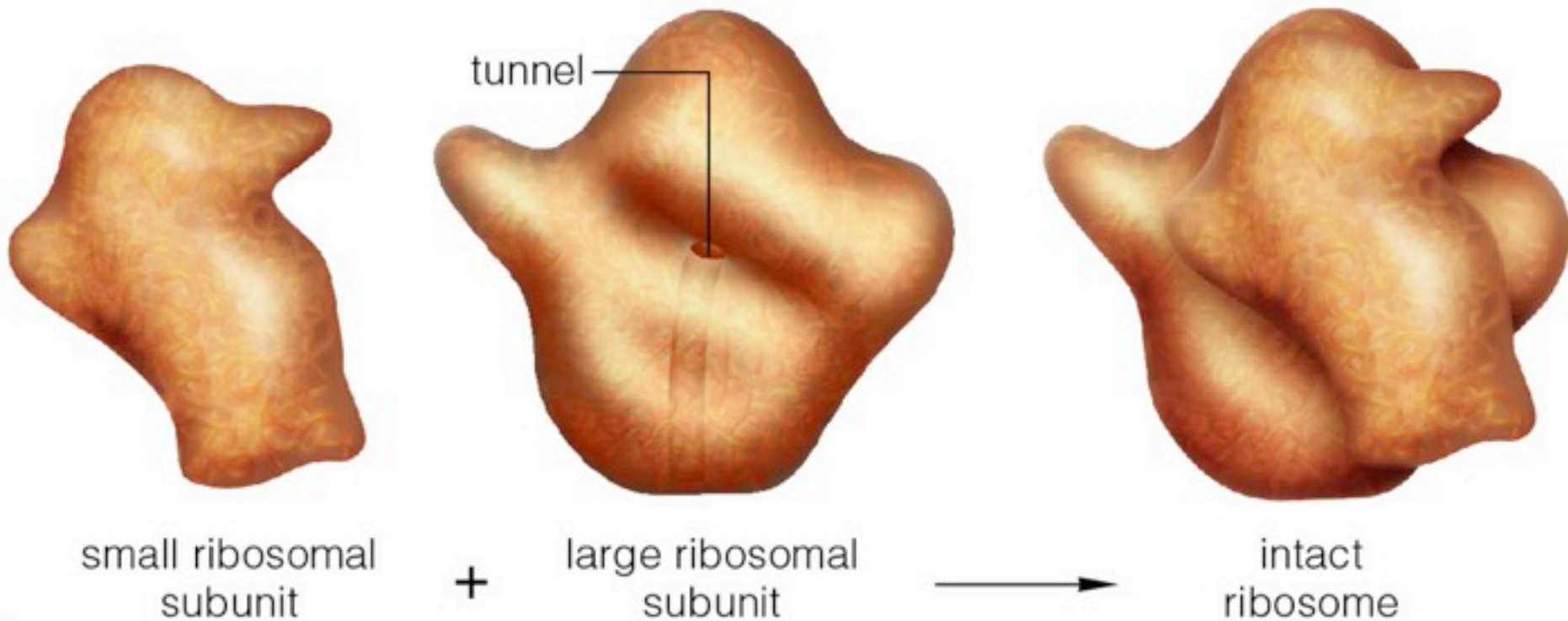
anticodon in tRNA

- tRNA= transfer RNA**  
brings an amino acid to  
a ribosome for incorporation  
into a polypeptide

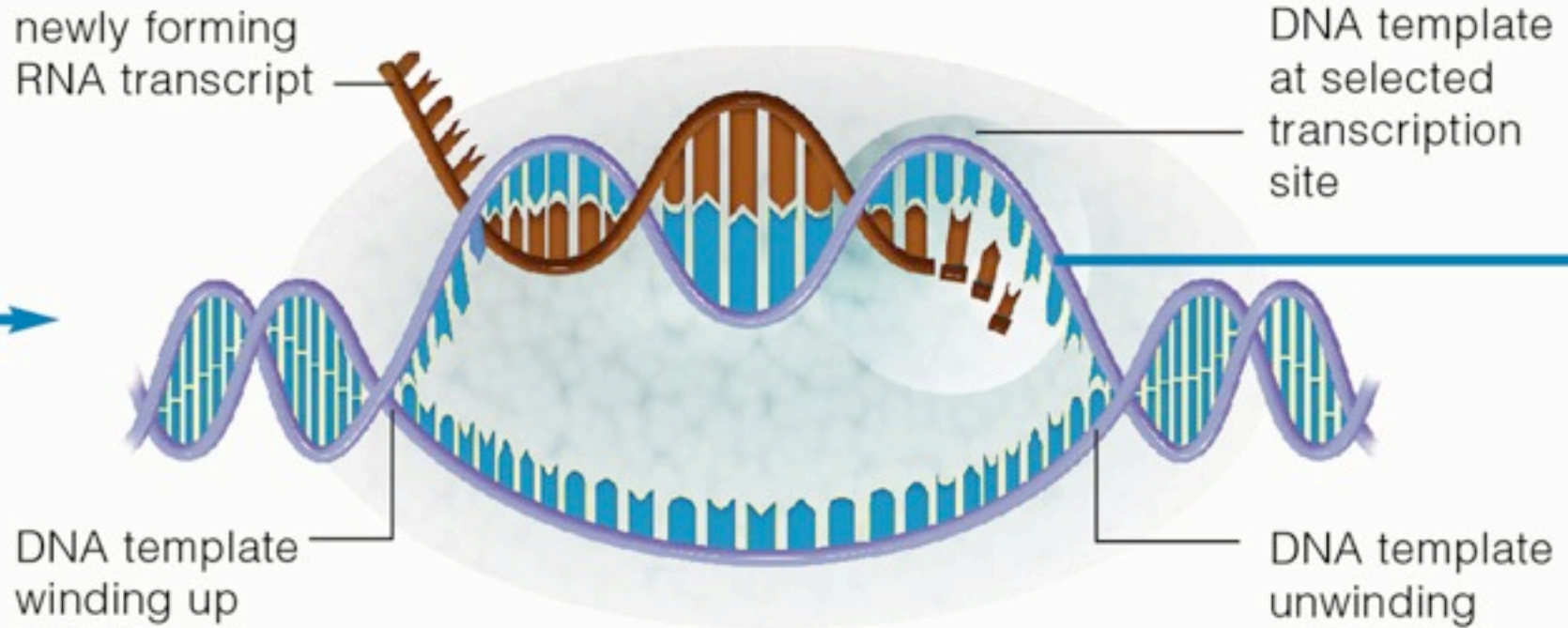




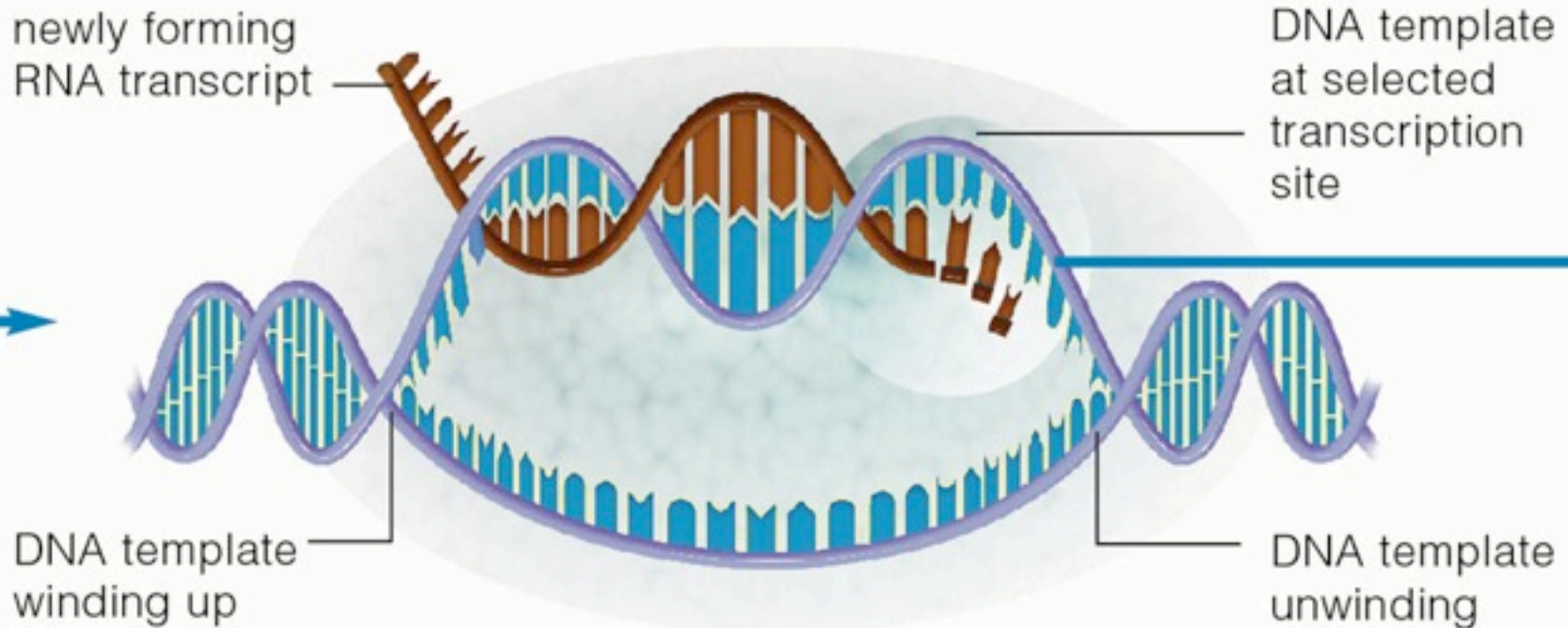
3. rRNA = ribosomal RNA-  
joins with other proteins to make 2 ribosomal subunits- one small and one large, these come together to make a ribosome which assembles amino acids into polypeptides



## IV. Transcription (RNA Synthesis)

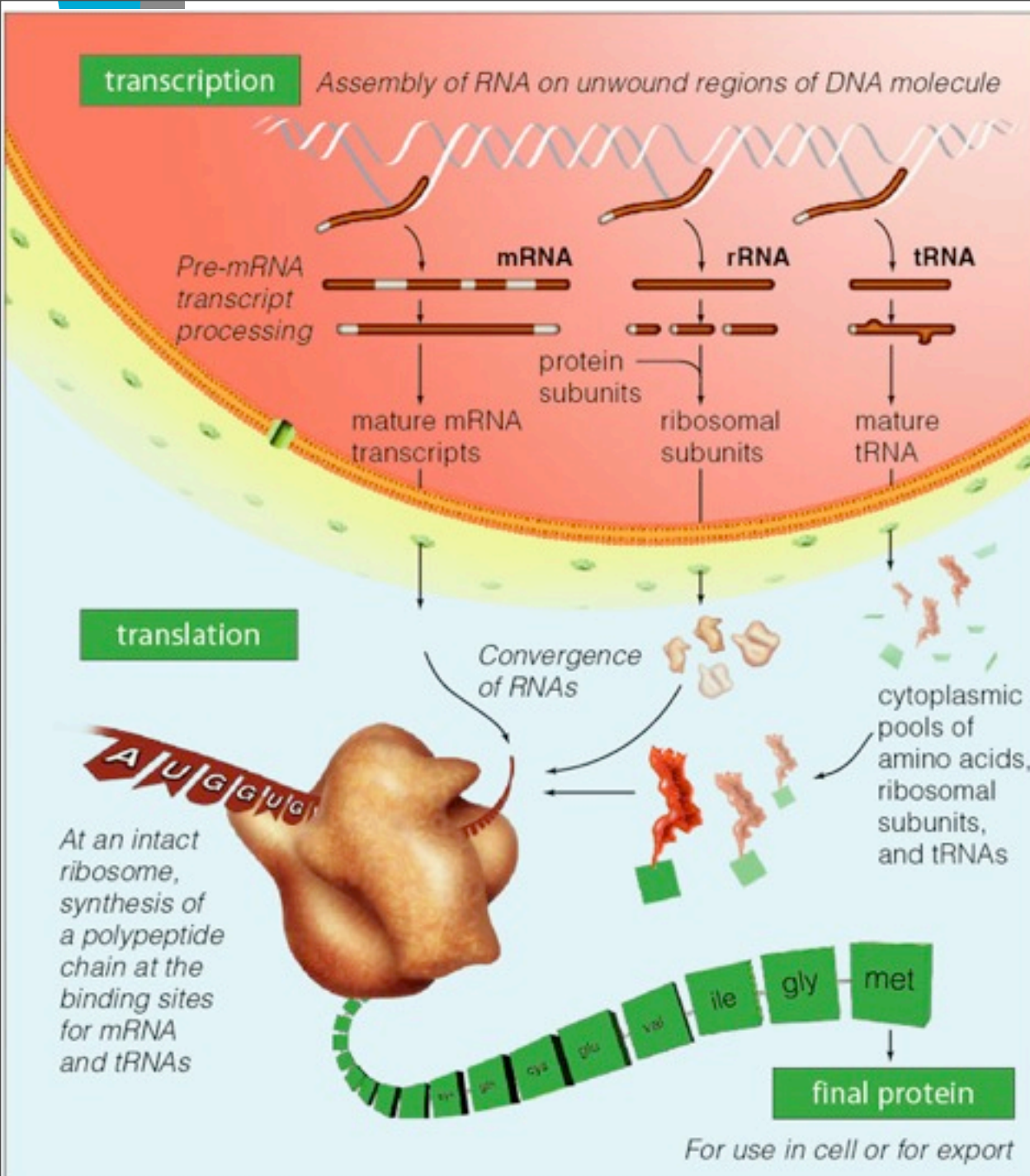


## IV. Transcription (RNA Synthesis)



© 2006 Brooks/Cole - Thomson

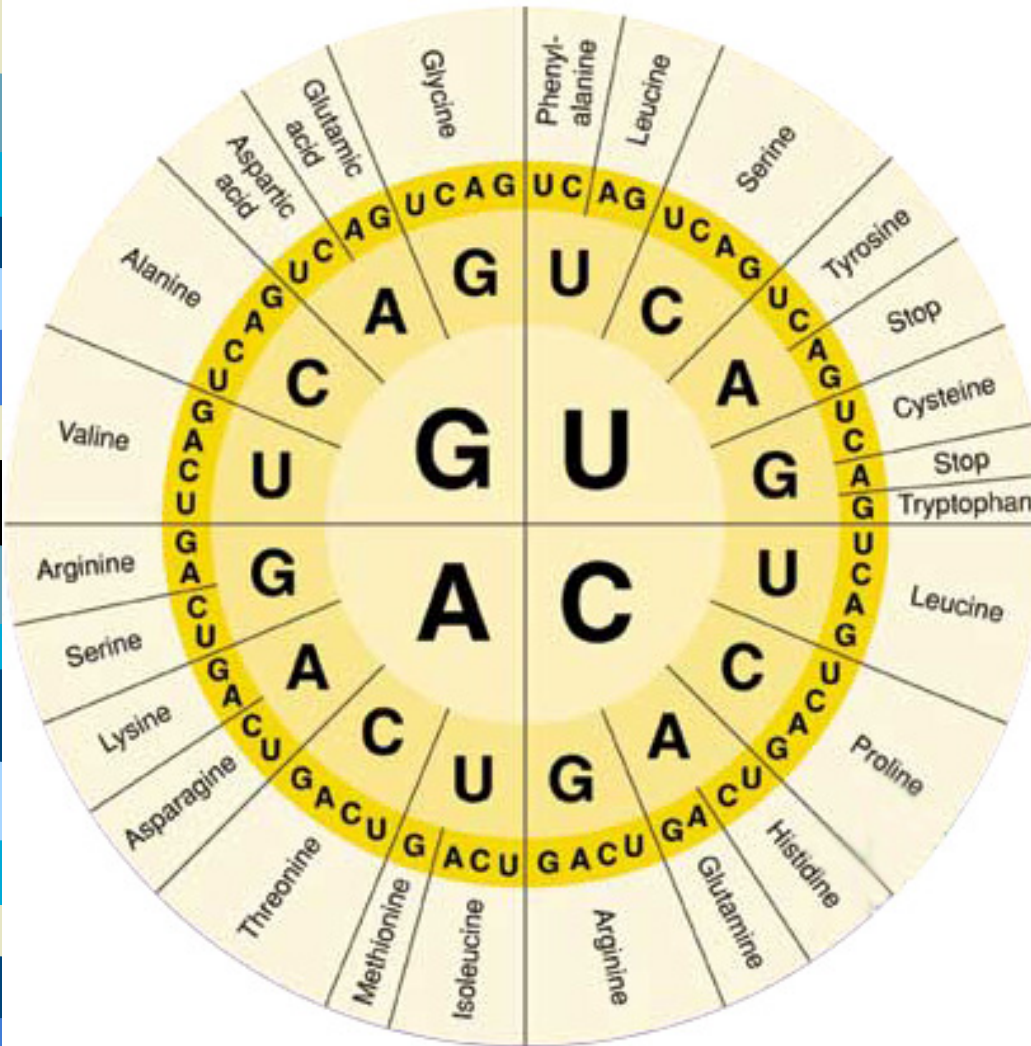
- Uses the enzyme RNA polymerase
- RNA made in a 5' → 3' direction
- Many copies can be made at once
- RNA will be processed before used



## Processing:

- 5' 'cap' added
- Poly A tail added to mRNA
- Cutting, looping, and folding
- Transported to cytoplasm (\*RER)

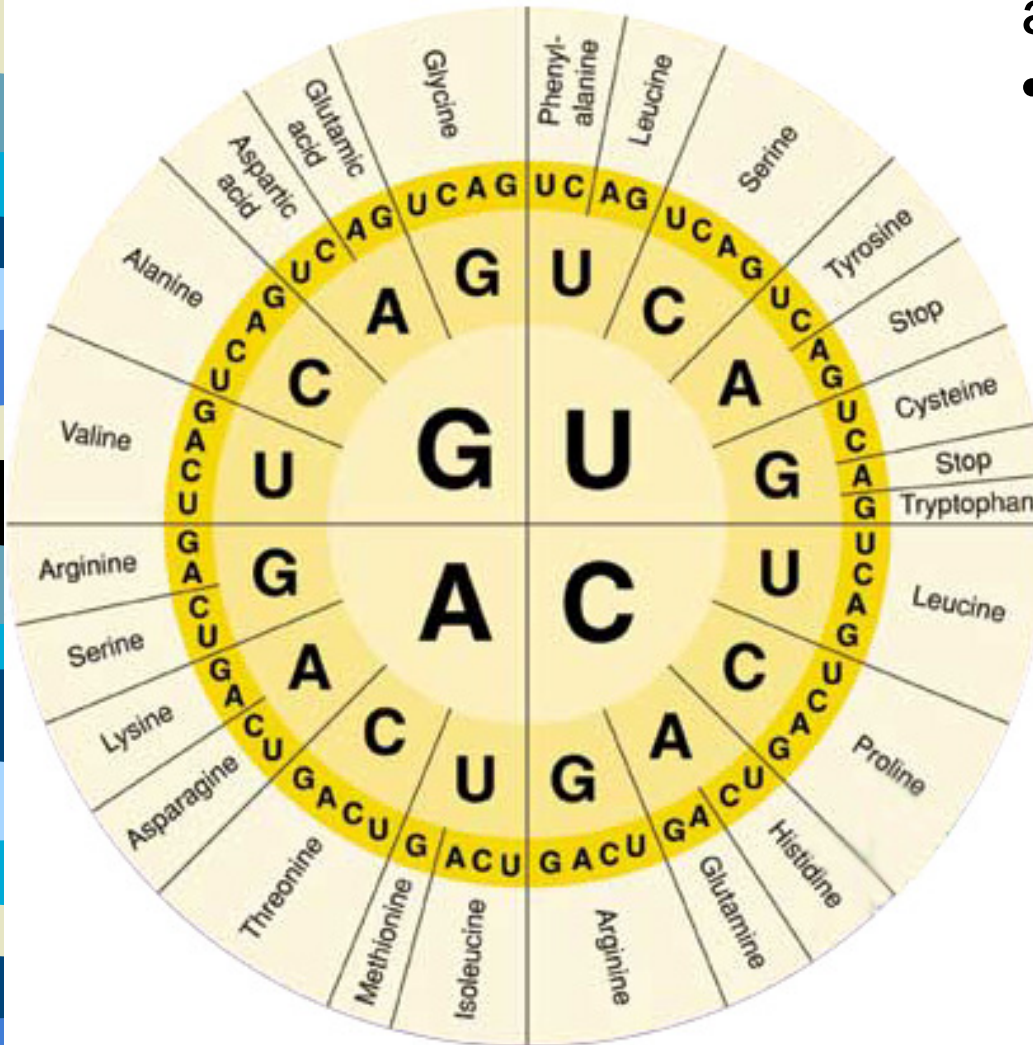
V. Translation (Protein synthesis)  
A. Genetic Code:



# V. Translation (Protein synthesis)

## A. Genetic Code:

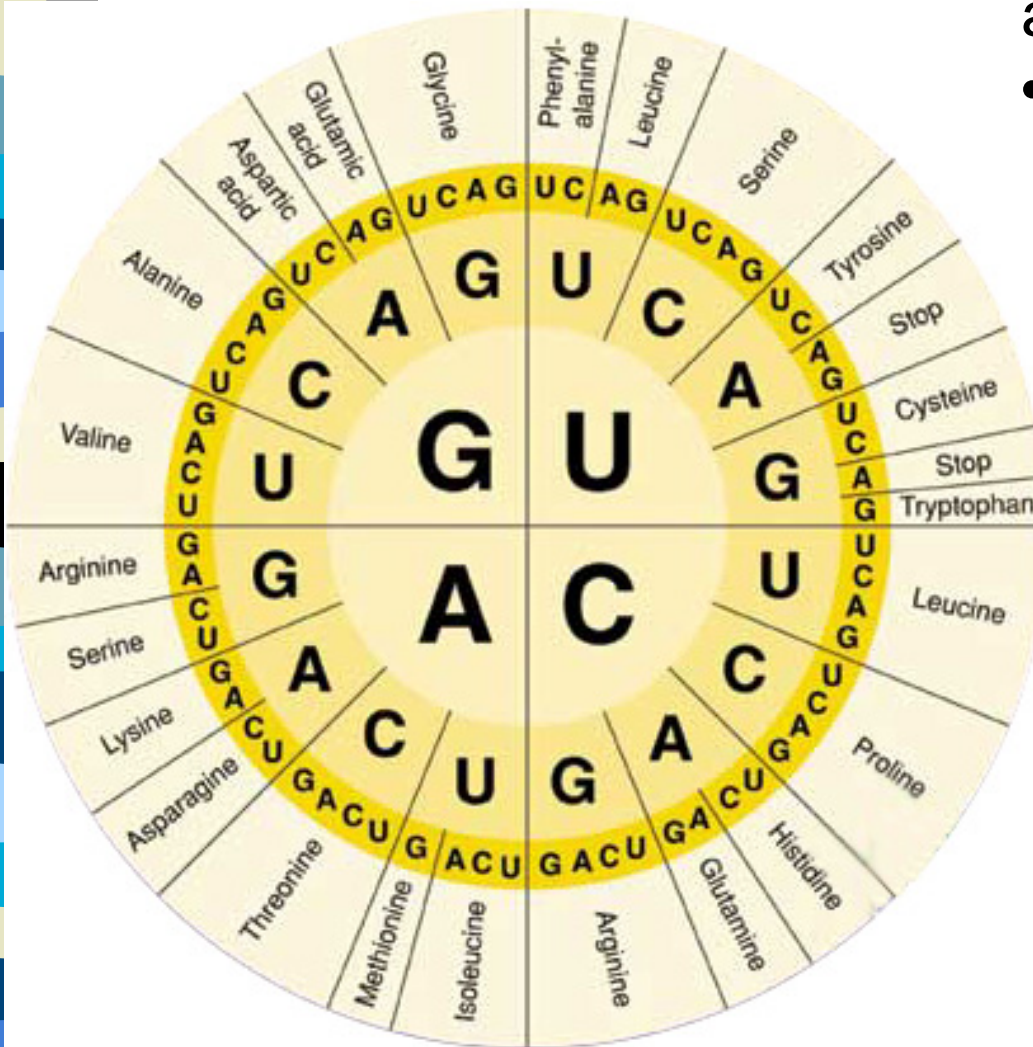
- 3 bases (codon) = 1 amino acid
- Start and stop signals



## V. Translation (Protein synthesis)

### A. Genetic Code:

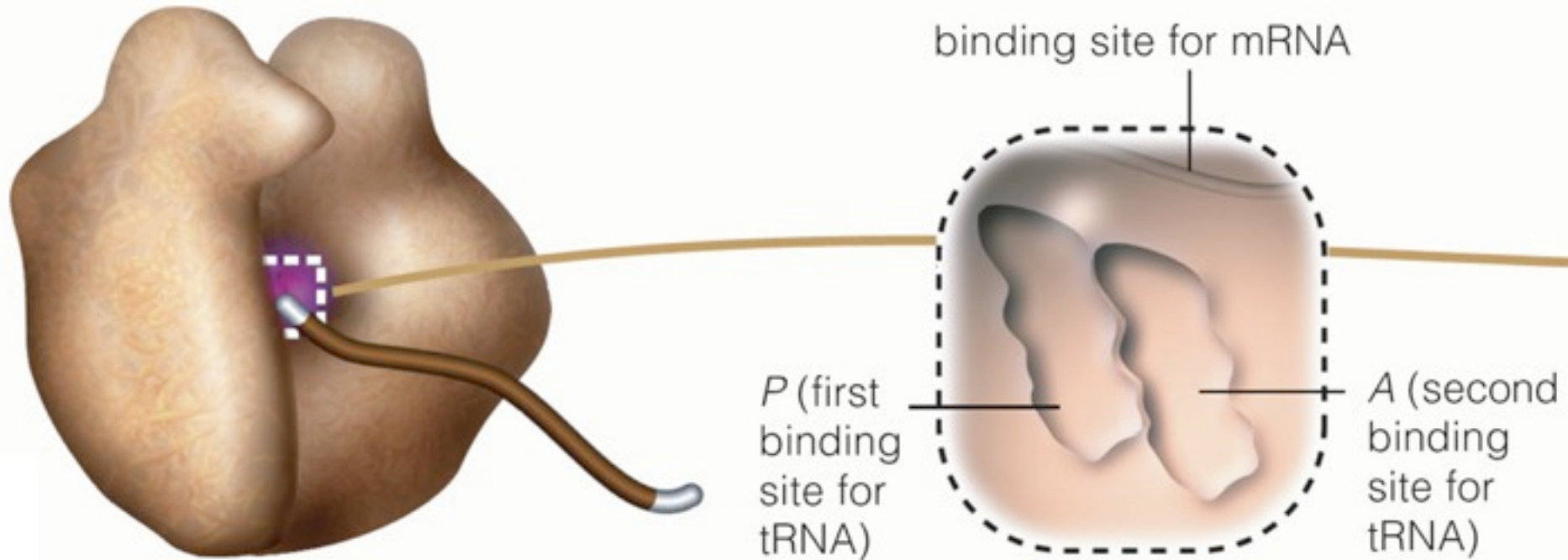
- 3 bases (codon) = 1 amino acid
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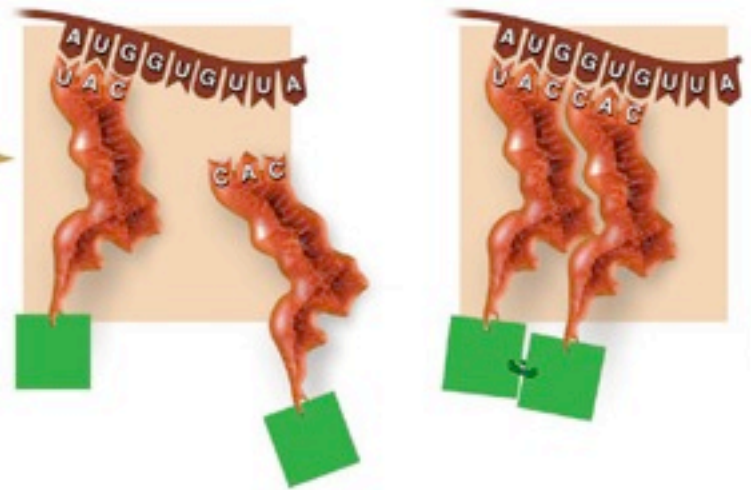
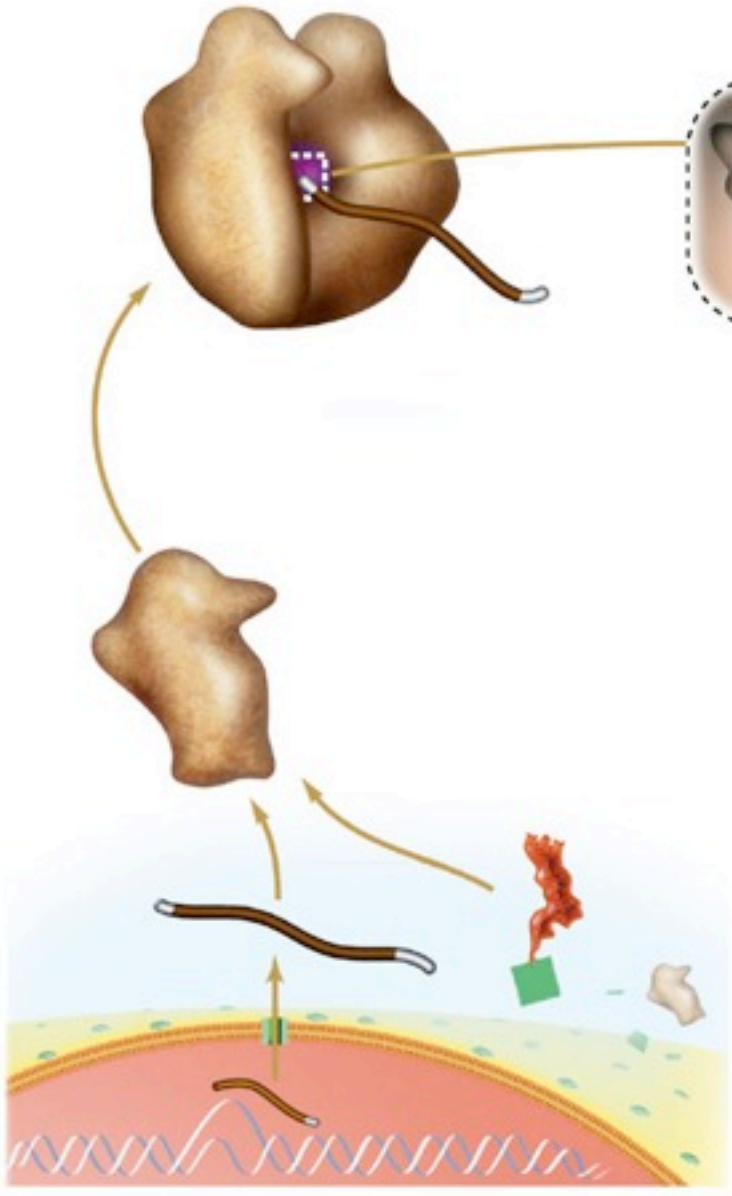
### B. Components

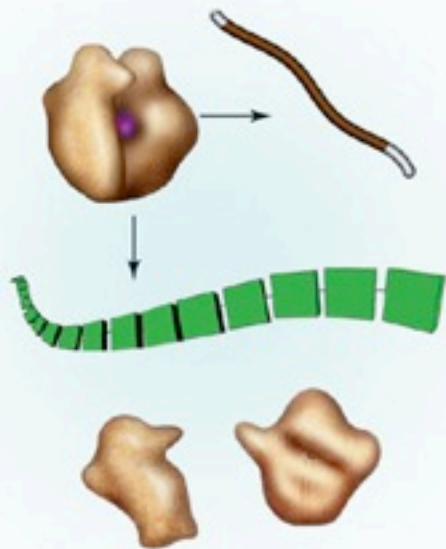
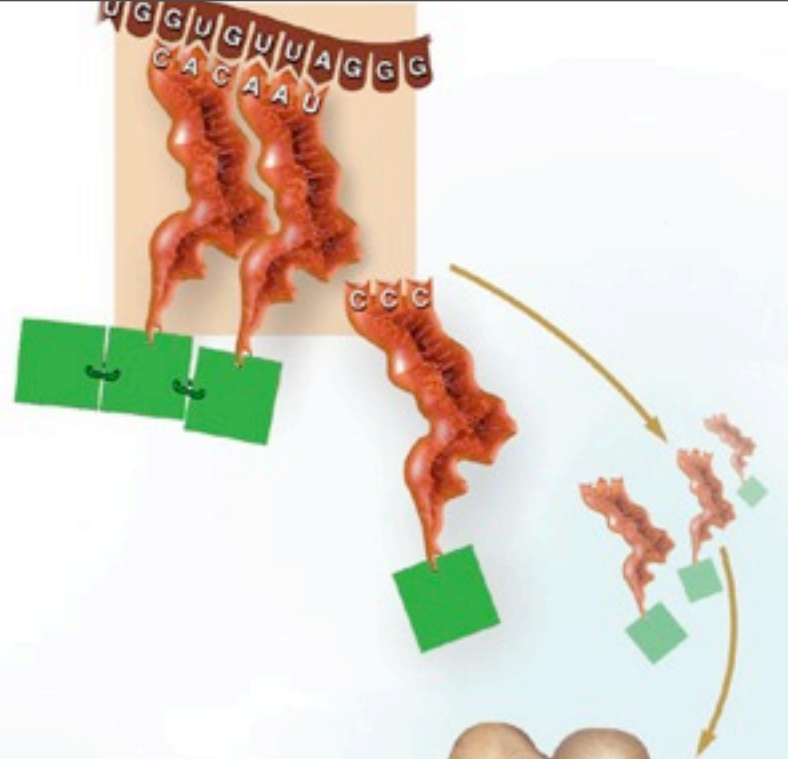
- mRNA
- Ribosomes
- tRNAs
- Pool of amino acids

## C. Steps



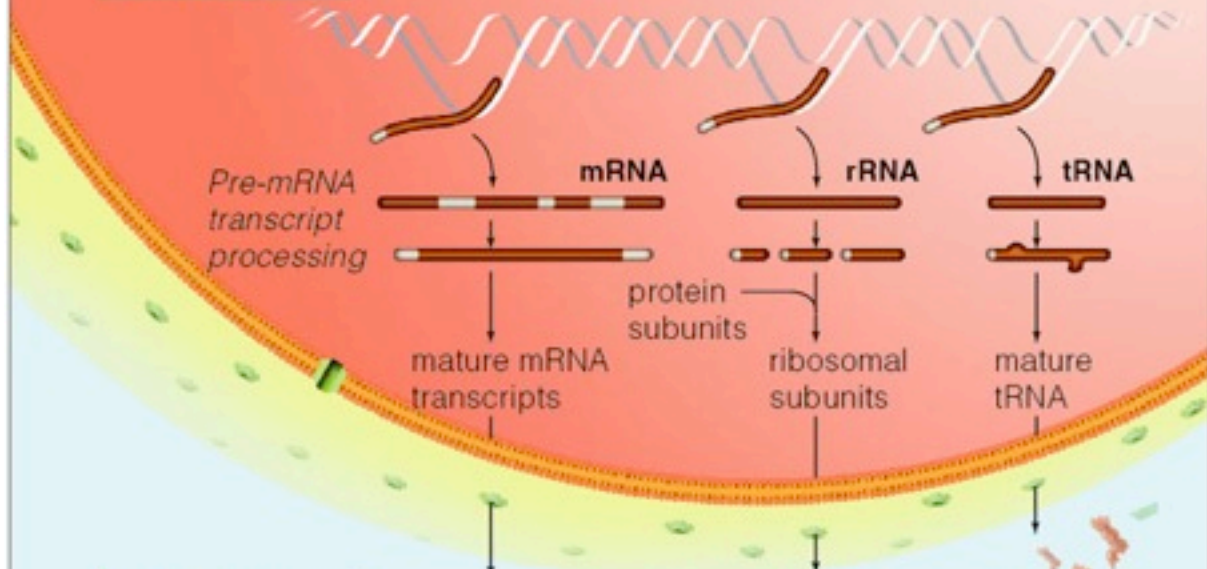




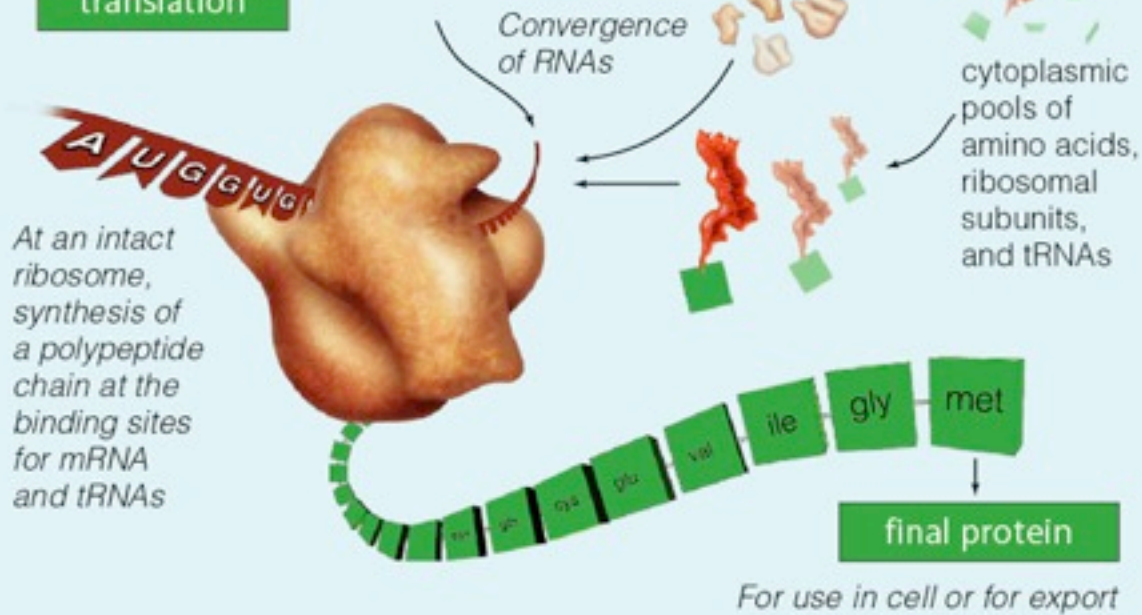


**transcription**

Assembly of RNA on unwound regions of DNA molecule



**translation**



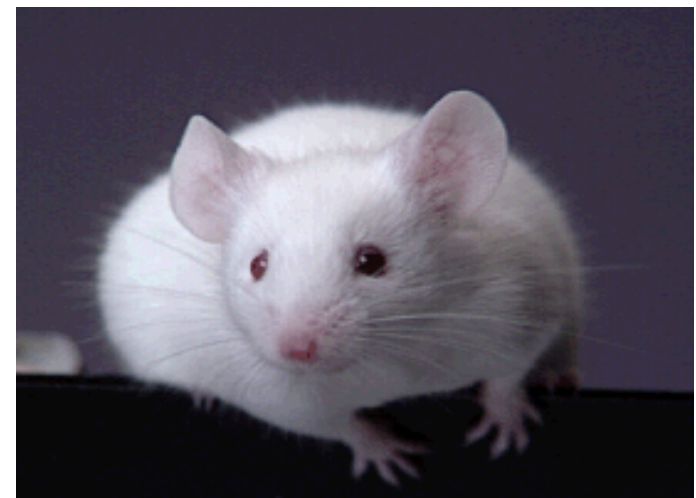
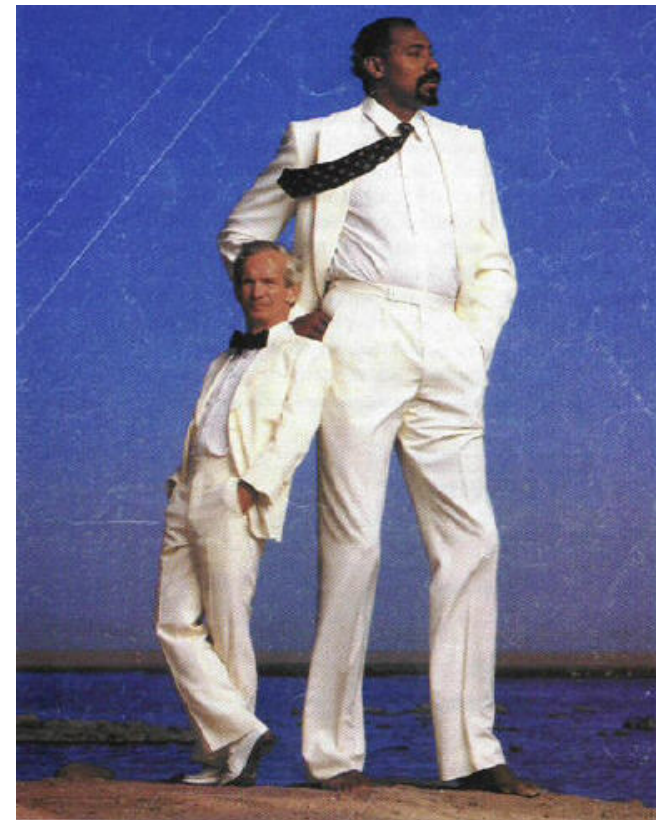


Questions for thought-

Why do we look and act different from each other?  
(99.9% of our DNA is the same)

Why do we look and act different from other organisms?  
(90% of our DNA is the same as in mice.)

What **CAUSES** these differences?



## VI. Mutations-

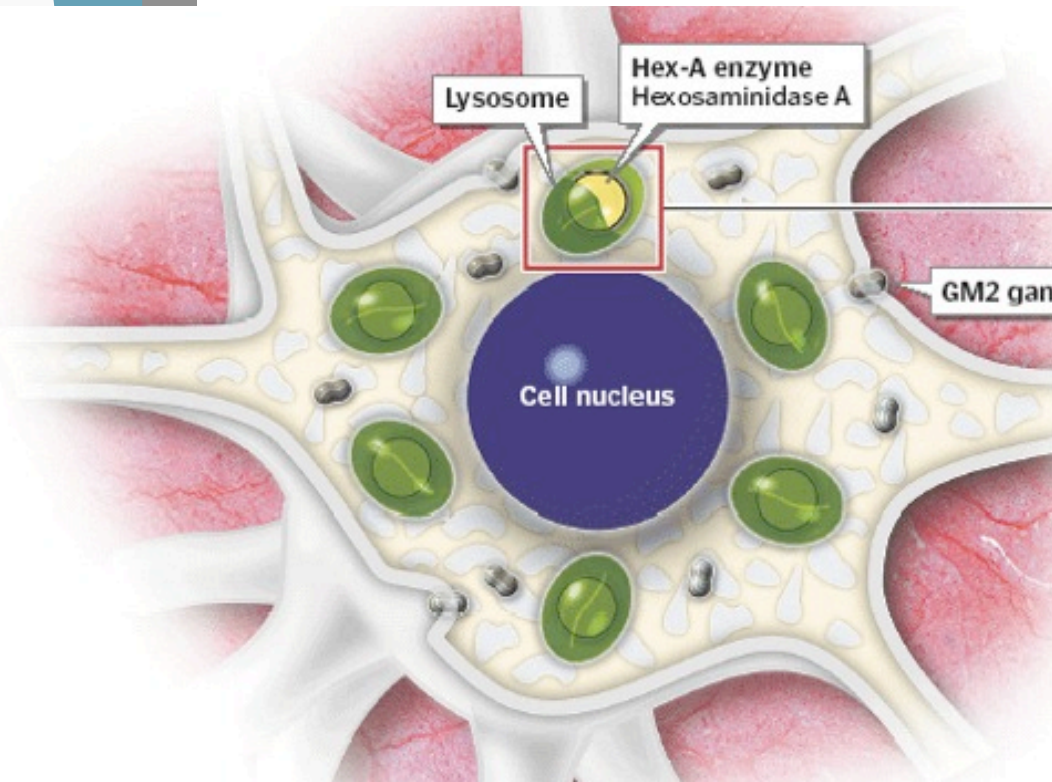
### A. Types of Gene or Point Mutations-

#### 1. Insertion-

extra bases are inserted into the genetic code

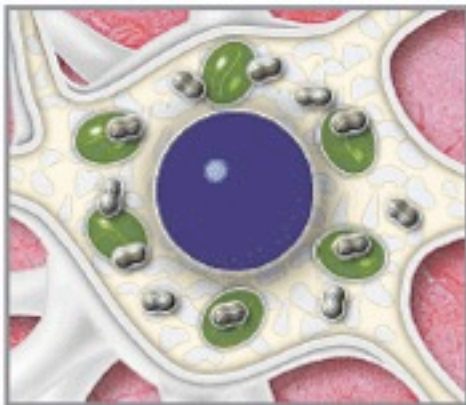
ex. Tay-Sachs Disorder



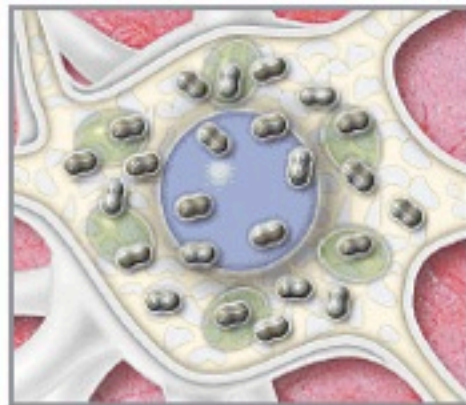


- 4 base insertion in the Hex A gene causes Tay-Sachs
- The gene normally codes for an enzyme in lysosomes that breaks down the food that brain cells eat- GM2- a lipid
- Leads to slow brain death by 2 years of age

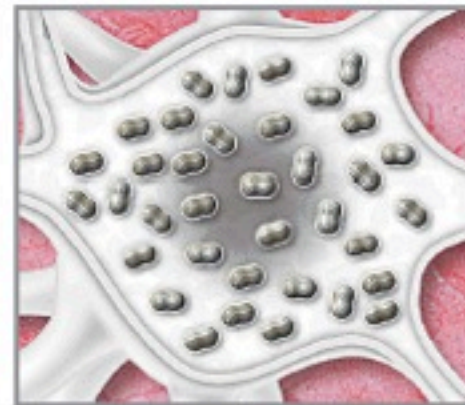
If Hex-A enzyme is not present ...



... GM2 accumulates ...



... and in time chokes off the cells.



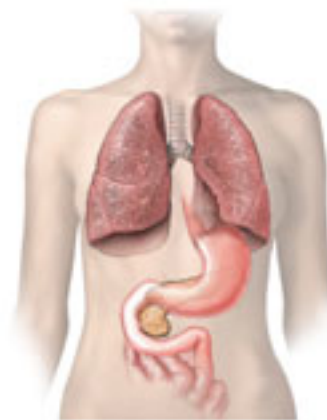
## 2. Deletion- extra bases are deleted from the genetic code

ex. Cystic fibrosis

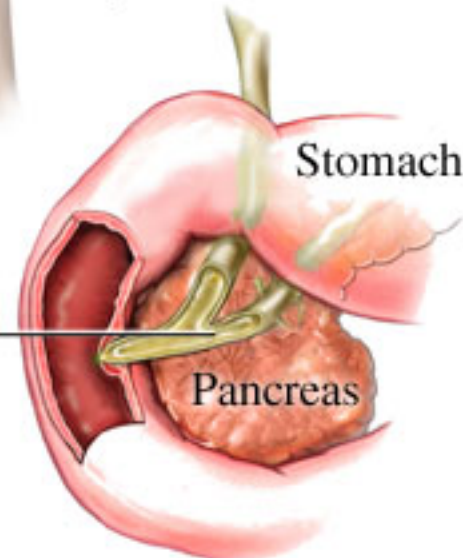
- 3 base deletion in gene for a Cl<sup>-</sup> transport protein
- The protein normally has 1480 amino acids
- 1/20 Caucasians are carriers



Mucus blocks  
air sacs (alveoli)  
in the lungs



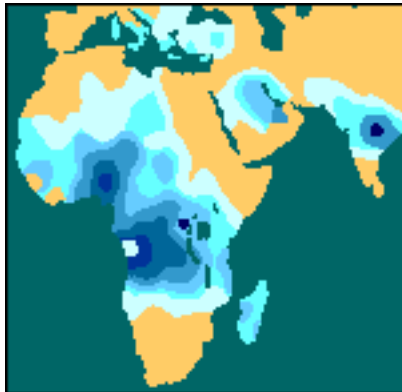
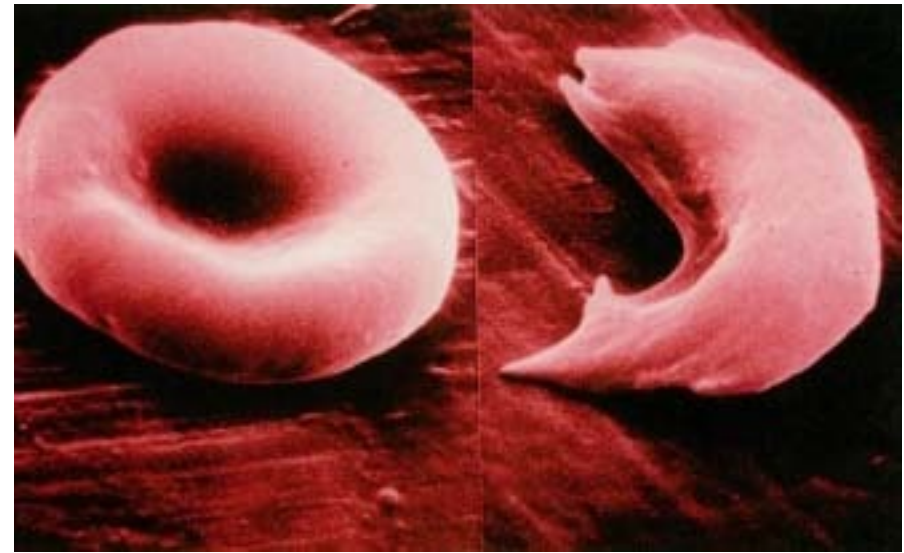
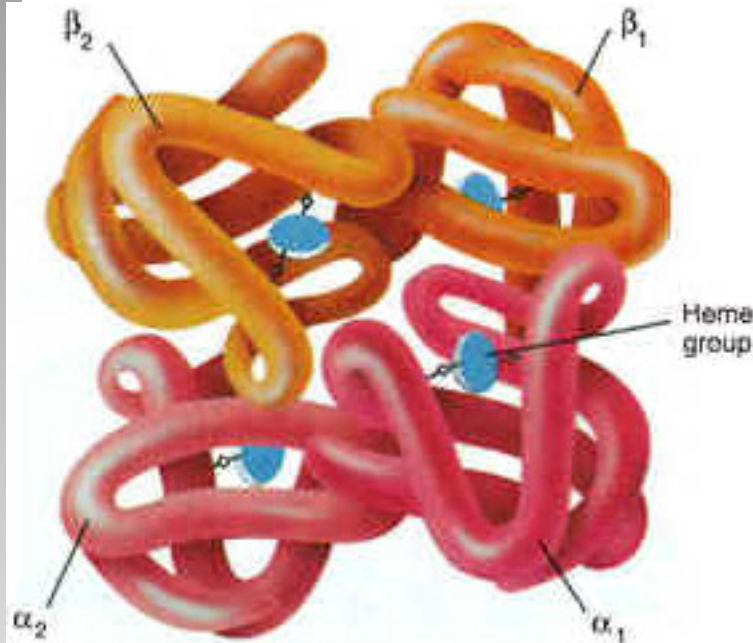
Mucus blocks  
pancreatic ducts



Pancreatic  
duct

Pancreas

3. Substitution-a base or many bases are swapped in the DNA code  
ex. Sickle-cell disease

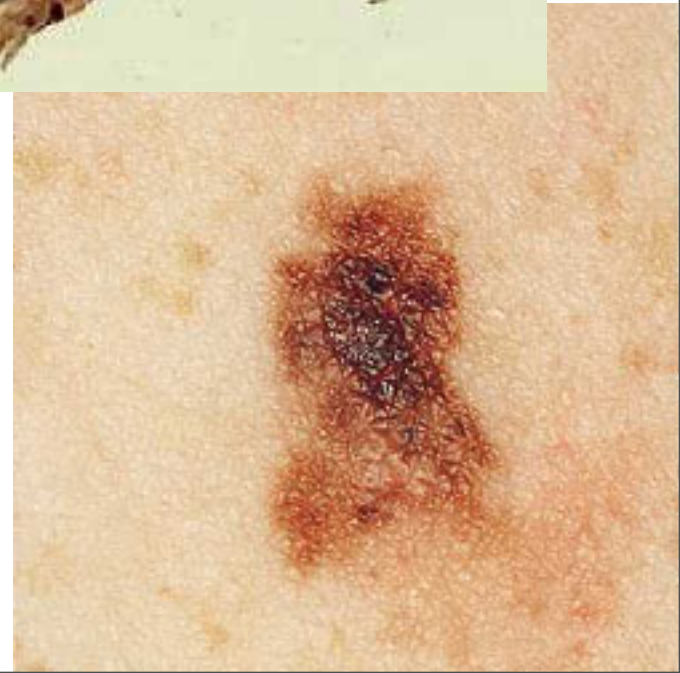
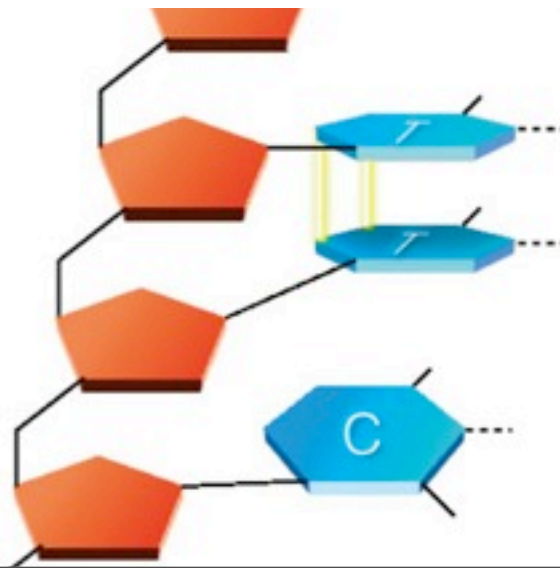


- 6th codon CTC  $\rightarrow$  CAC in beta-globin gene
- 1/12 African Americans are carriers



## B. Causes

1. Radiation- nuclear, UV, x-ray
2. Viruses and other microorganisms
3. Environmental poisons- formaldehyde, nicotine, asbestos
4. Alcohol and diet?
5. Random mistakes



# C. Effects

1. Harmful
2. Neutral
3. Beneficial



# VII. Viruses

NON-CELLULAR



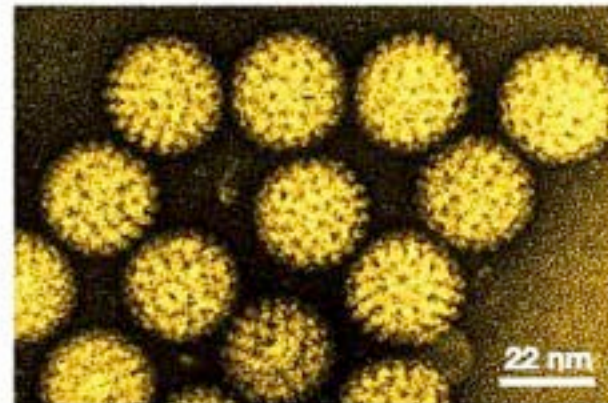
Ebola virus



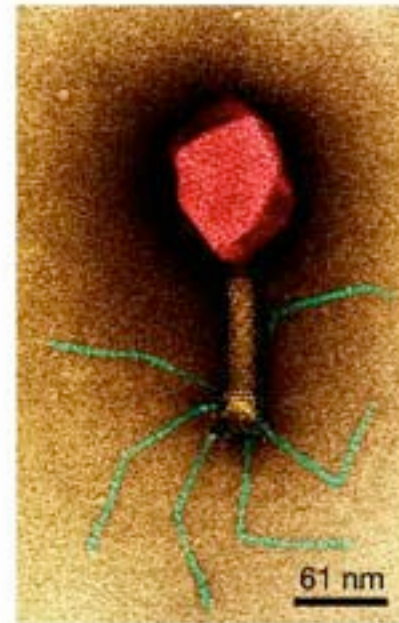
tobacco mosaic virus



adenovirus



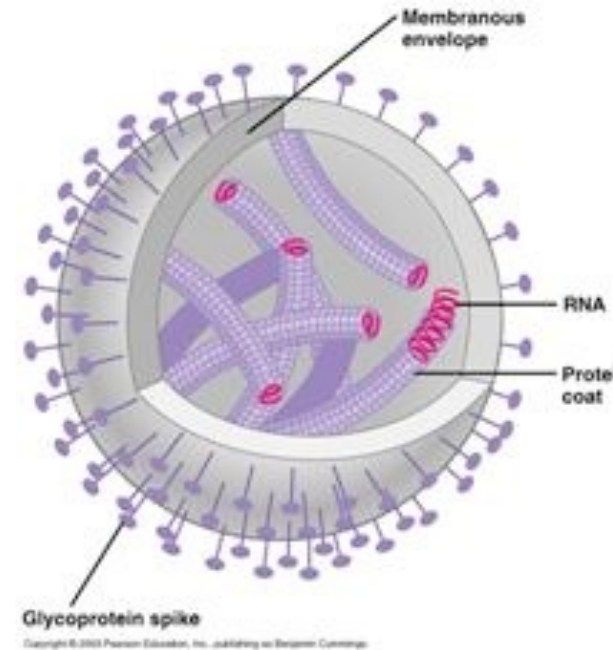
influenza virus



bacteriophage T4

## A. Characteristics

1. Protein capsule and nucleic acid (DNA or RNA) core
2. Obligate parasite- is a parasite that cannot complete its life cycle without exploiting a suitable host.

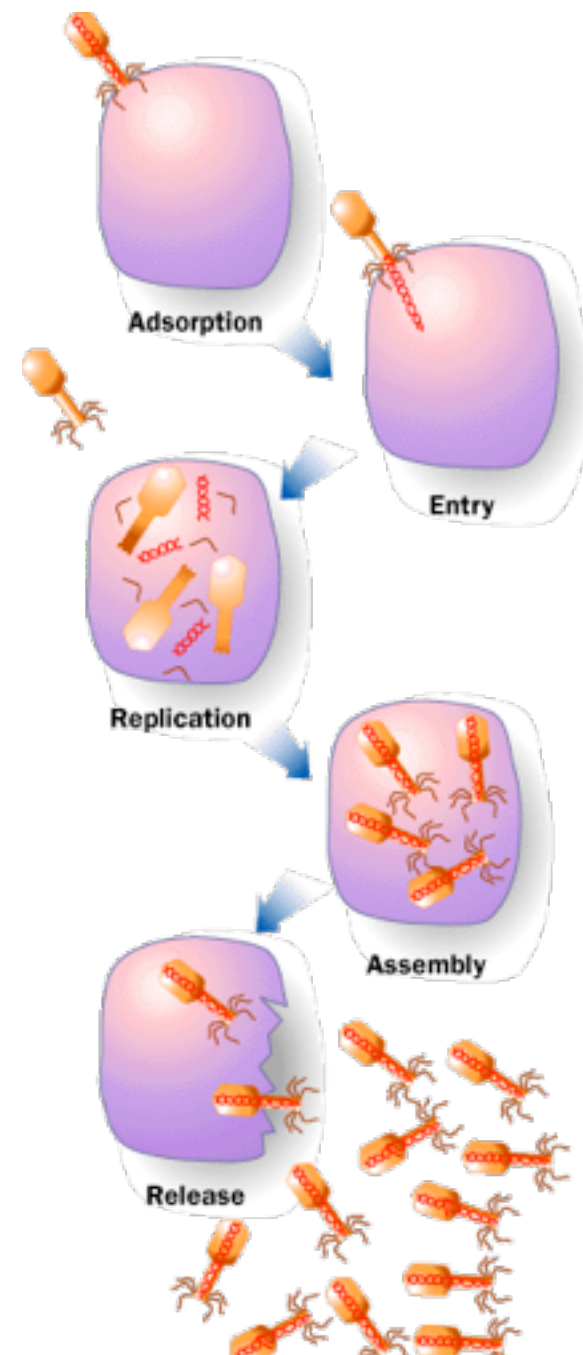


## B. Actions in Host Cells

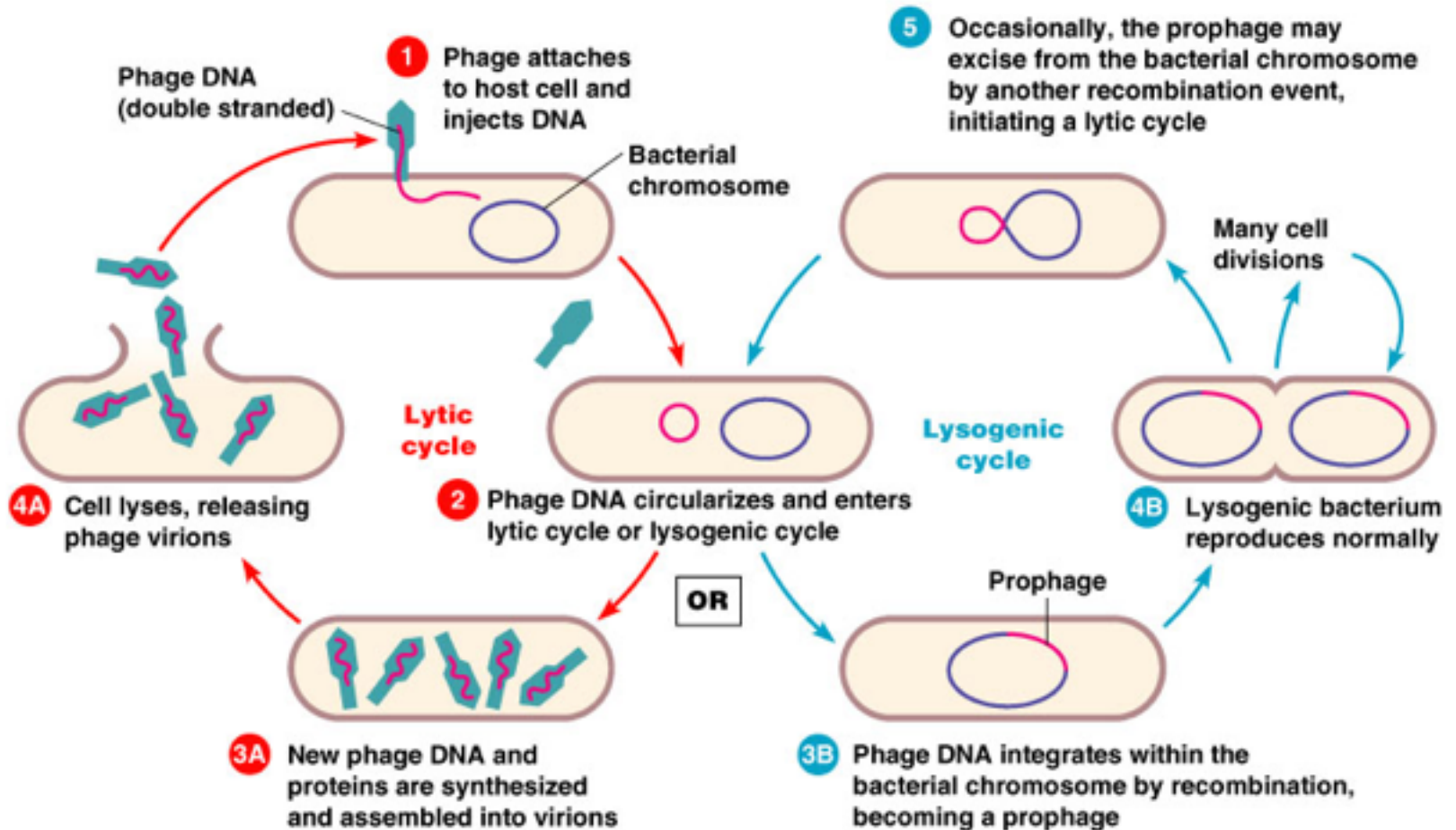
1. Lytic Cycle-
2. Lysogenic Cycle-

*get ready to draw 😊*

# Lytic Pathway:

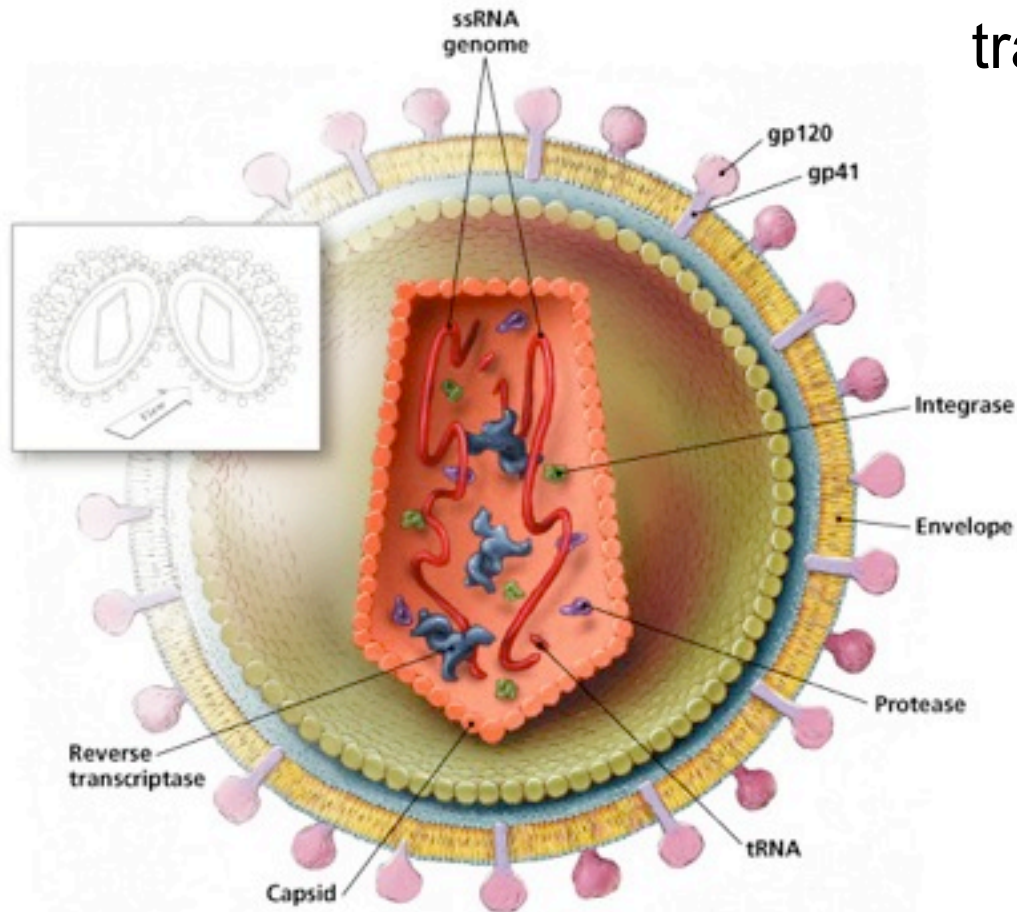


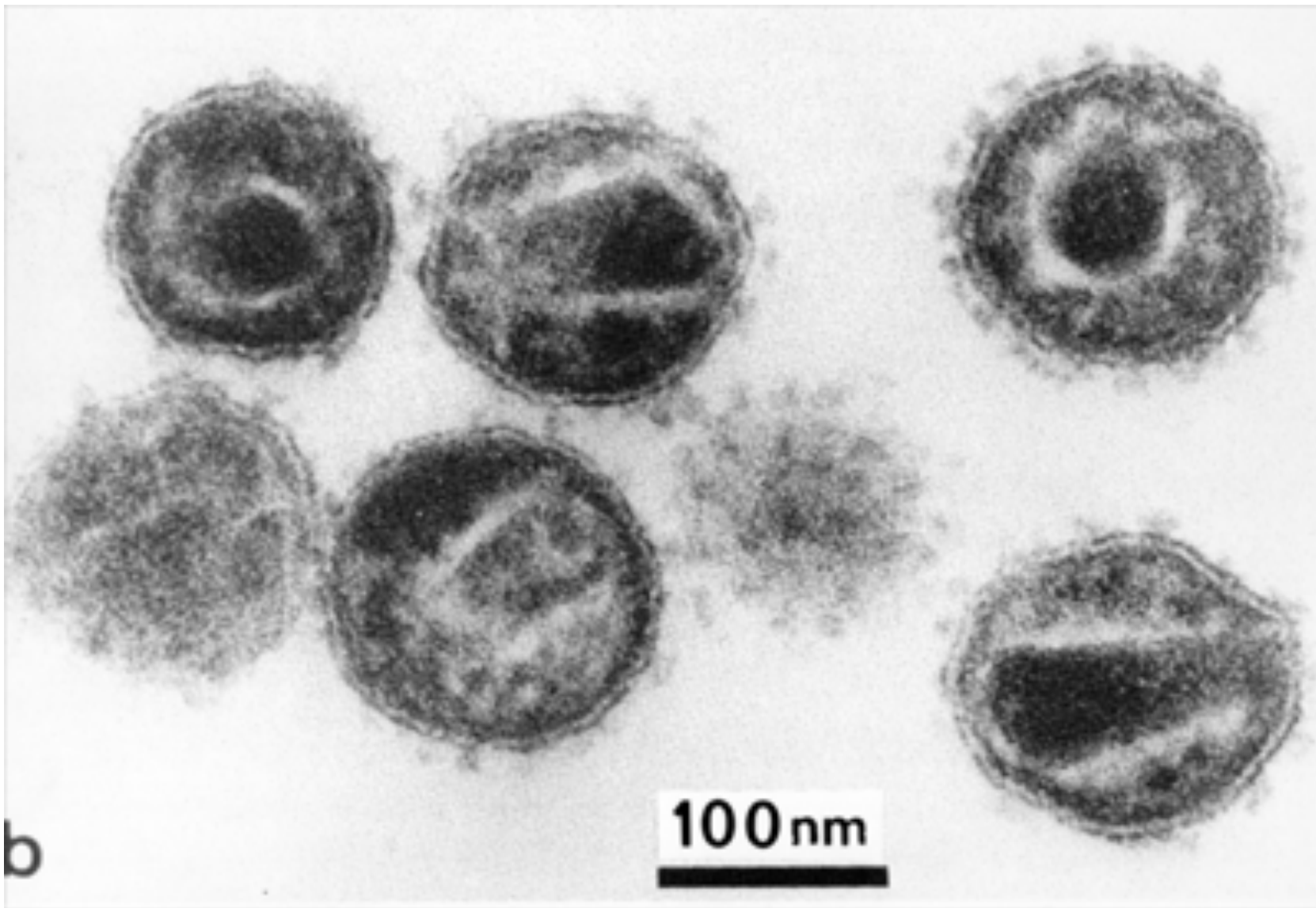
# Lytic and Lysogenic Pathway



# C. Special Case- HIV- retrovirus

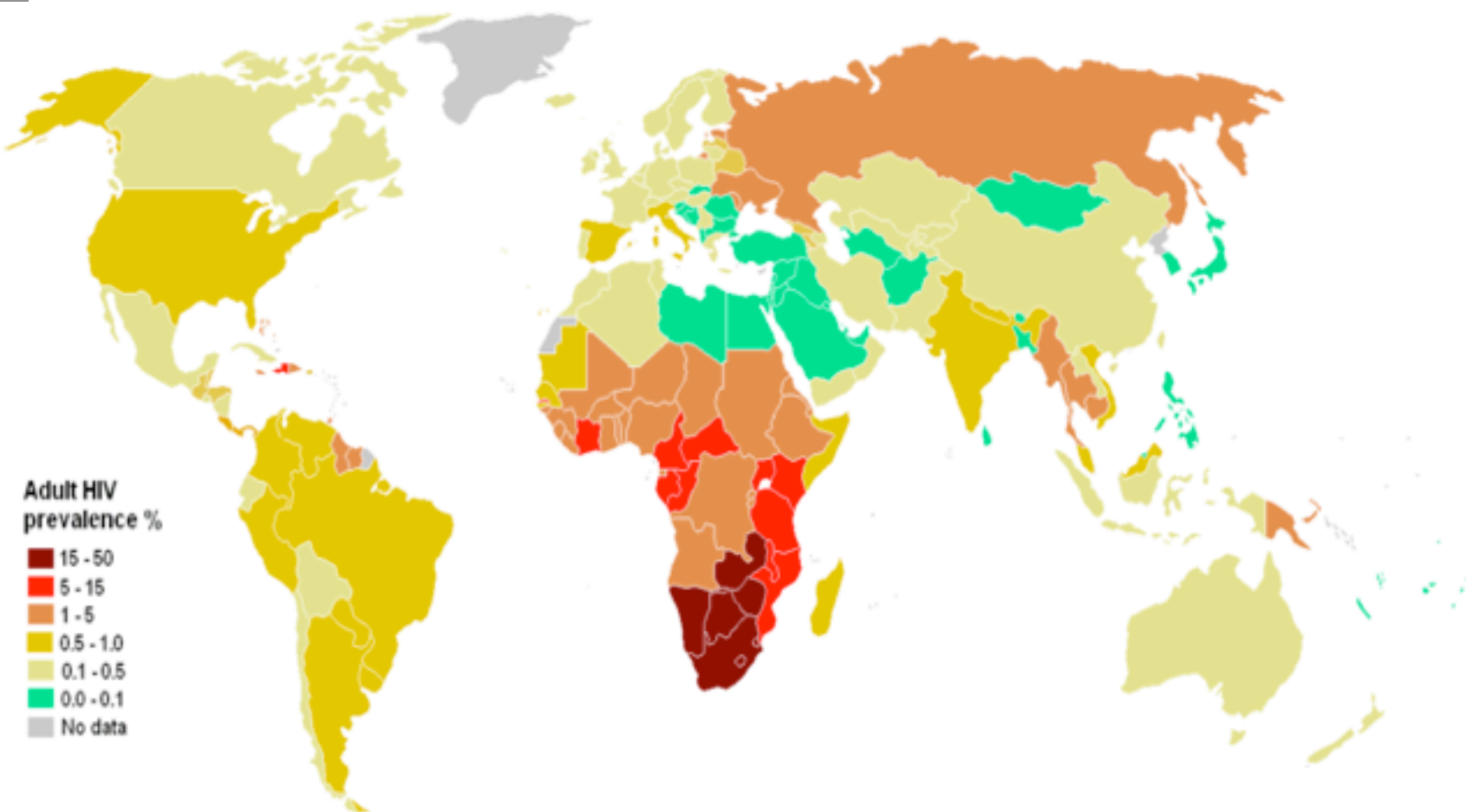
Features: Reverse transcriptase





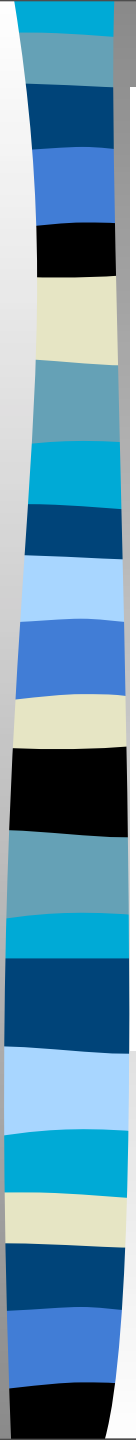


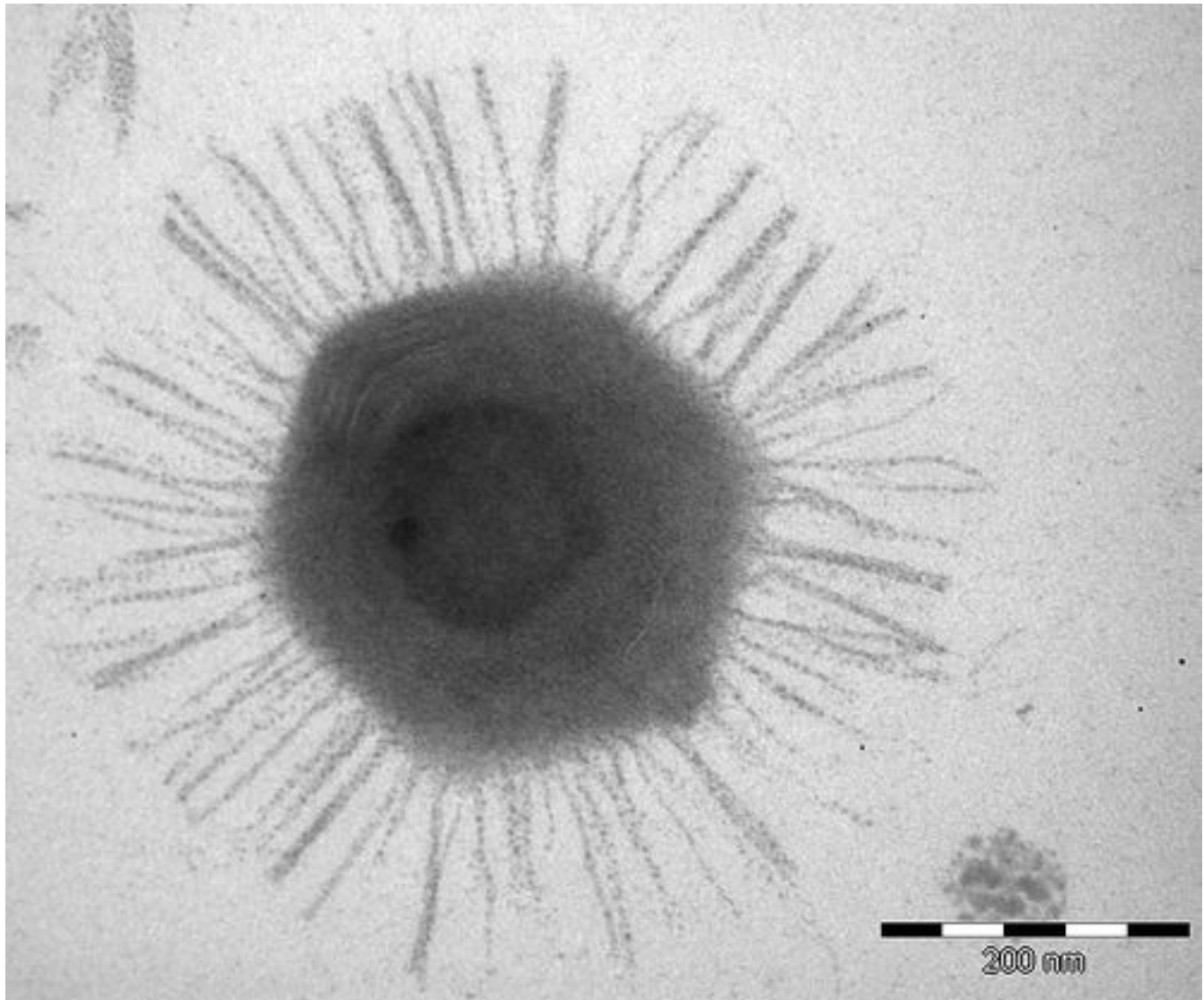
# 6. Epidemiology-



Adult HIV prevalence %

- 15 - 50
- 5 - 15
- 1 - 5
- 0.5 - 1.0
- 0.1 - 0.5
- 0.0 - 0.1
- No data





Mimivirus.....