

Molecular and Cellular Biology (MCB)

BB 101

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Class 6: learning objectives

- Inherited DNA leads to specific traits
 - Coat color, white eyes, phenylketonuria
- DNA: packaging, accessing the stored message
 - Book shelves, storage compactors, chromosomes
- DNA to protein – basic principles
 - Transcription and translation
- The genetic code and mutations

How is information stored in DNA?

Figure 16.22



DNA (chromosome)



Specific sequence of nucleotides along the strands of DNA

part of Figure 17.4

Replication of DNA - copying of information

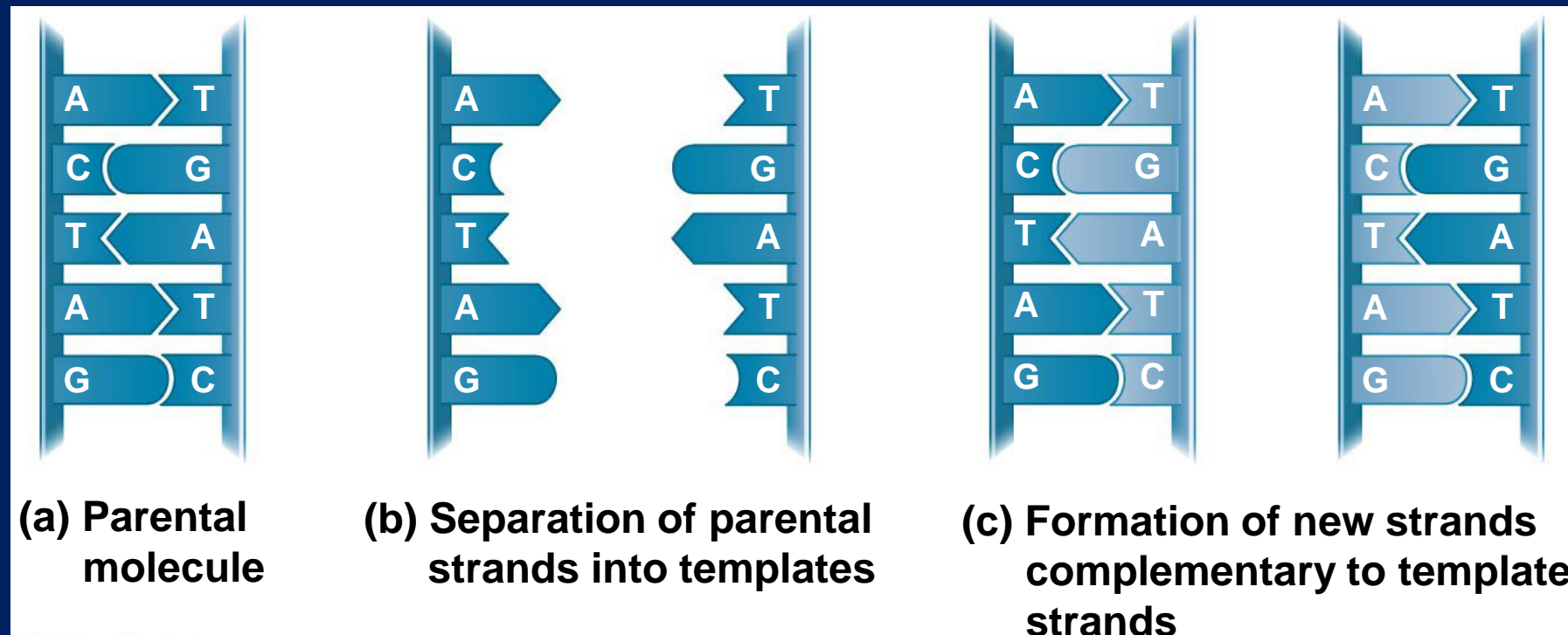
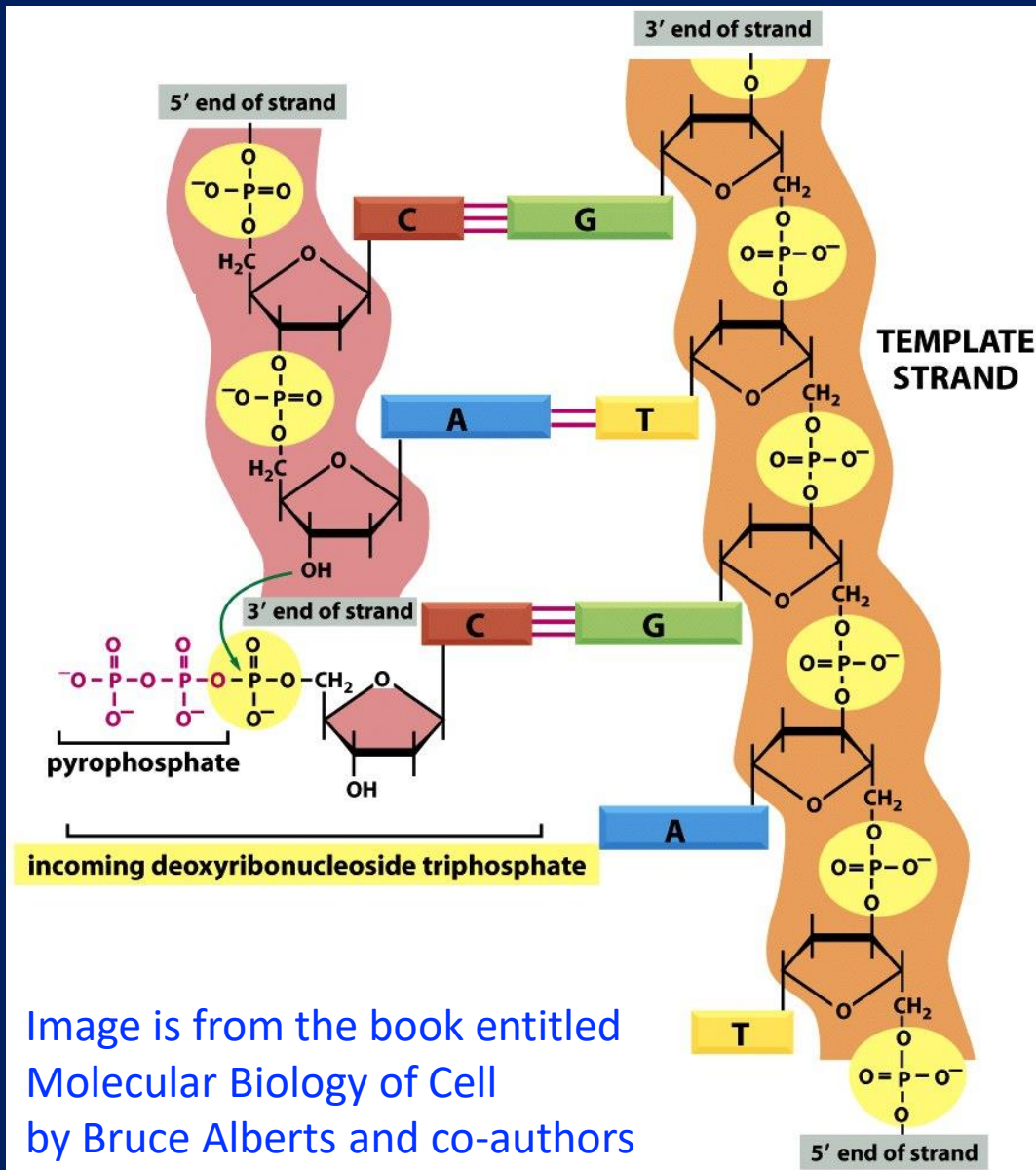


Figure 16.9

Will there be (or can there be) errors during replication?
If yes, what are the consequences?

Directional synthesis



DNA synthesis proceeds in 5' to 3' direction

DNA polymerases are enzymes which catalyze DNA synthesis

DNA polymerases have error checking and correction activity also



http://www.school-clipart.com/school_clipart_images/clip_art_illustration_of_a_short_pencil_with_an_eraser_0515-1007-2718-0950.html

Mutations

MUTATION

Change in “information”
i.e., change in DNA sequence

Can be natural or induced

Error by DNA polymerase
External causes (e.g., UV light)

MUTATION

Can be silent i.e., no APPARENT
effect on the organism

Wild type and mutants

WILD TYPE

The reference
(genetic) state
of an organism

“trait normally found
in a population”

Mutation



MUTANT

Organism that has
undergone genetic change
with respect to the wild
type

- Wild type and mutant are two DISTINCT organisms
- Wild type and mutant are WITH RESPECT TO A TRAIT or GENE
- A mutant can have one or more genetic changes with reference to the wild type

Wild type and mutants



[ADP Opens Global 'Center of Excellence' in Bucharest - New Jersey Business Magazine \(njbmagazine.com\)](#)



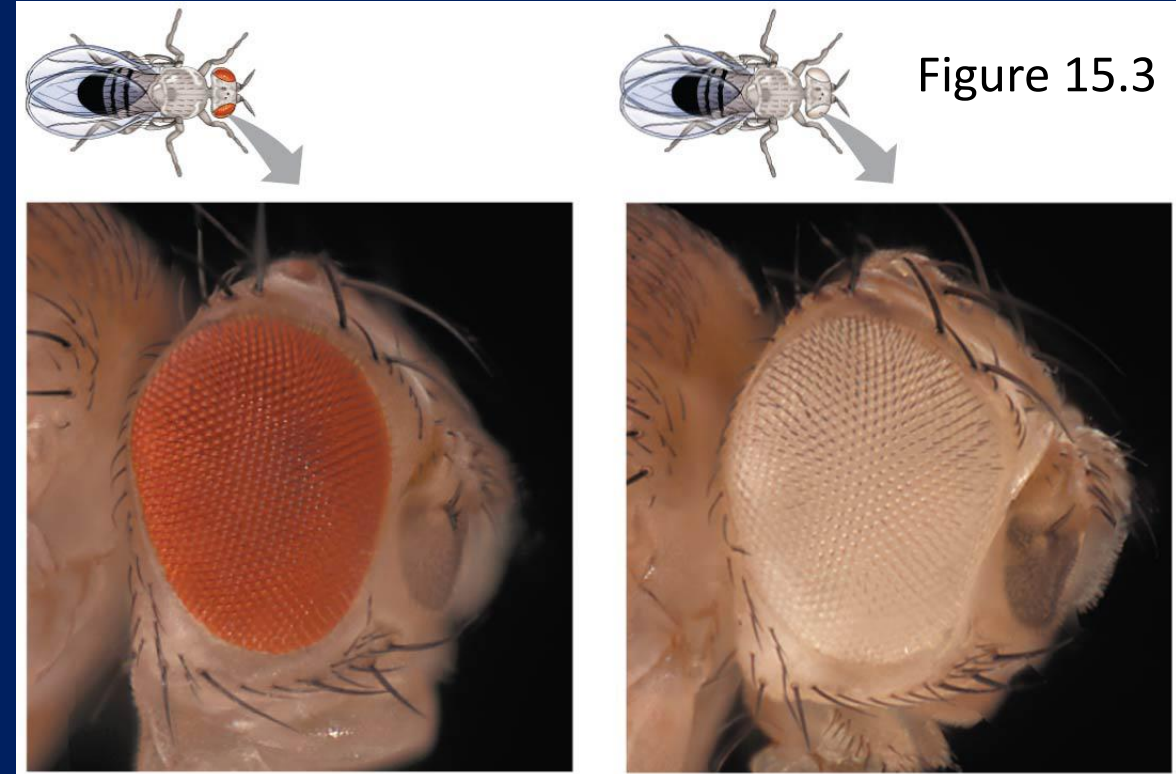
[Who Are The New Mutants In X-Men: Days Of Future Past? | Movies | Empire \(empireonline.com\)](#)

Wild type and mutants



Figure 17.1

Biology. A global approach



Fruit fly with
white eyes

Fruit fly: *Drosophila melanogaster*

Germ cells (gametes) and somatic cells

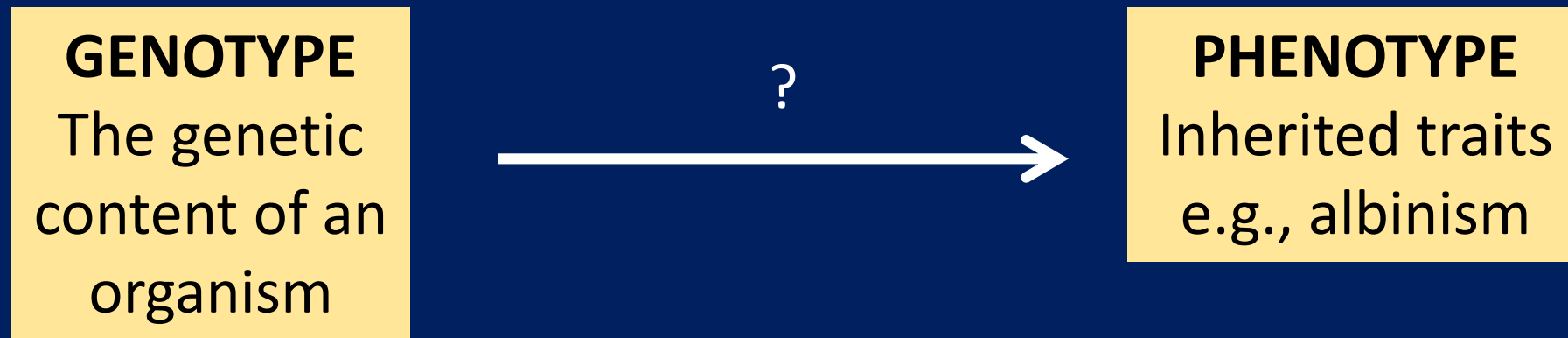
In multi-cellular organisms, egg and sperm are called germ cells (or gametes)

All other cells are called somatic cells

Only mutations in germ cells are transmitted

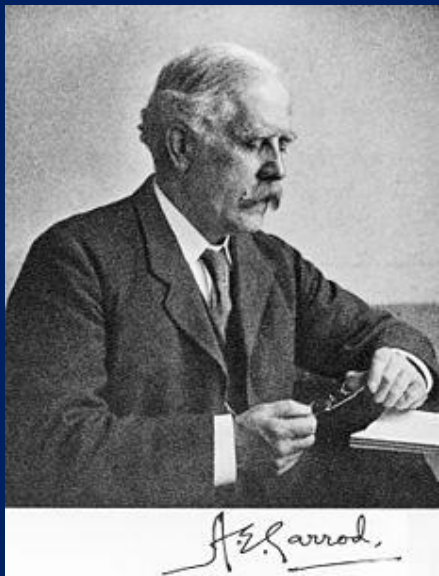
Genetic makeup and observed traits

Inherited DNA leads to specific traits

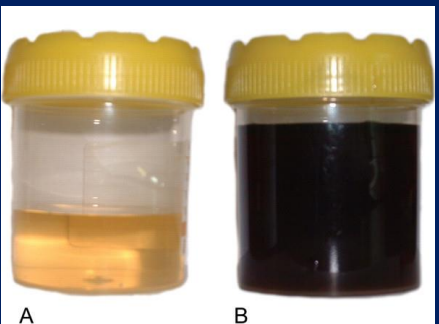


How does the information flow?

Genes dictate phenotype through enzymes



- Urine becomes black because it contains the chemical alkapton
- Most people have an enzyme that converts (i.e., metabolizes) alkapton
- People who do not inherit this enzyme suffer from the disease alkaptonuria
- Garrod named this as an “inborn error of metabolism”



He was ahead of his time!



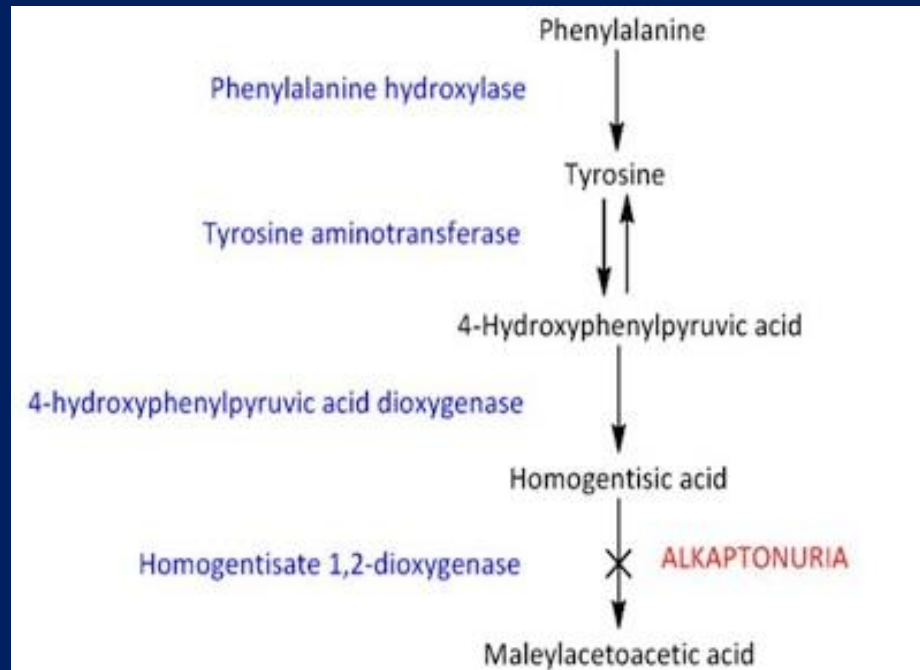
Archibald Garrod (1857-1936) English Physician

Image of Garrod taken from Wikipedia

Images of urine and pigmented hands taken from <https://lrodgers93.wordpress.com/>

Metabolic pathways

Gregor Mendel's principles of heredity apply to humans as well as peas



Genes dictate production of specific enzymes

Cells synthesize or degrade organic molecules via metabolic pathways

Each chemical reaction in a pathway is catalyzed by a specific enzyme

<http://www.rarediseasesindia.org/aku>

Neither the chemical reactions nor the enzymes that catalyze were known

Relationship between genes and enzymes

Which genes are responsible for the
coat color in animals
white eyes of *Drosophila*
Black-urine in alkaptonuria
...

What do genes do?

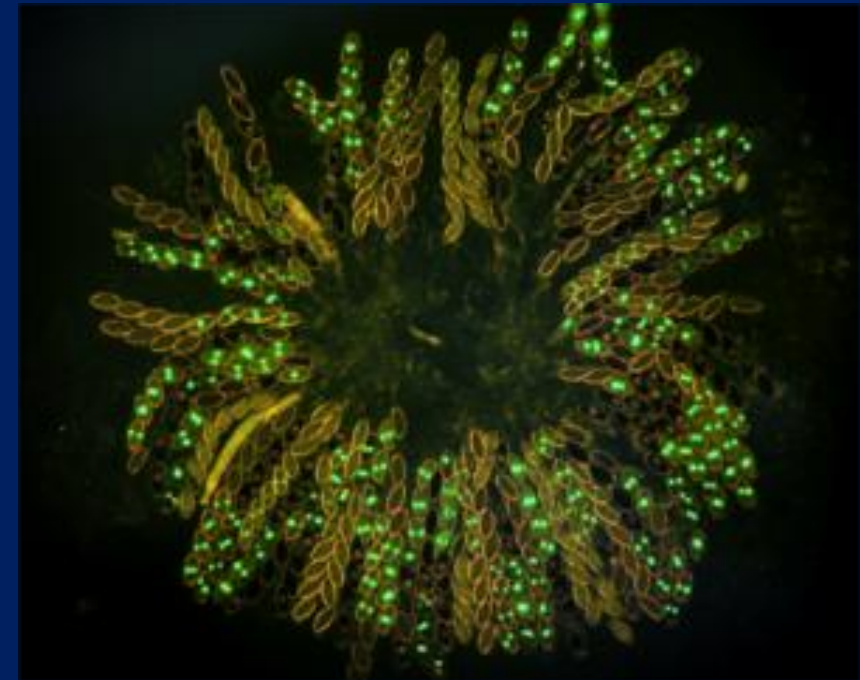
Relationship between genes and enzymes



Figure 1 | **George Beadle (left) and Edward Tatum (right) receiving their Nobel Prizes.** Image courtesy of the Karolinska Institute, Stockholm, Sweden.

Taken from the article published in the journal
Nature Review Genetics (2004) Vol 5, p949

Worked at the Stanford University



It is a bread mold

It is a fungus

Its scientific name is *Neurospora crassa*

Mold is a type of fungus

Relationship between genes and enzymes

Which genes are responsible for the coat color, white eyes, etc.?
What do genes do?

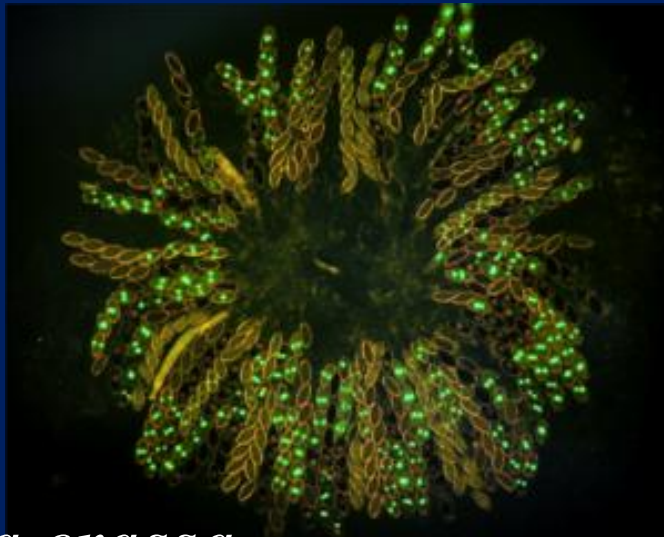
Beadle and Tatum's approach:

Instead of asking, "which genes are responsible for observed phenotype"
Seek (or look for) genes that control already known reactions

Wild type *Neurospora* can grow on minimal media

Minimal medium consists of
inorganic salts,
glucose and
biotin (a vitamin)
in agar, a support medium

Complete medium consists of
minimal media,
all 20 amino acids and
a few other nutrients
in agar, a support medium



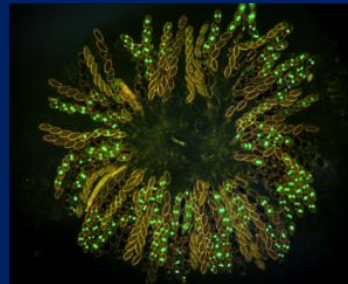
Wild type *Neurospora* can grow even
in minimal medium because it can
biosynthesize all the other molecules
required for growth

Neurospora crassa

Finding out what genes do

- George Beadle and Edward Tatum exposed bread mold to X-rays
- X-ray were known to cause mutations (change in the genetic make up)
- By this process, they isolated mutants that were unable to survive on minimal media

Relationship between genes and enzymes



*Neurospora
crassa*

Wild type *Neurospora*

X-ray irradiation

Mutant *Neurospora*

Complete
medium

Minimal
medium

Growth

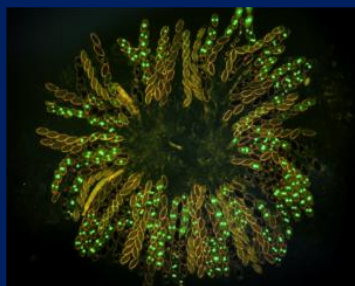
No growth

Exposure to x-rays results in mutants which do NOT grow even in complete medium. Such mutants are discarded

Exposure to x-rays alters the genetic make up (i.e., causes mutations)

Isolate ONLY such mutants which grow in complete medium but not in the minimal medium

Relationship between genes and enzymes



*Neurospora
crassa*

Wild type *Neurospora*

X-ray irradiation

Mutant *Neurospora*

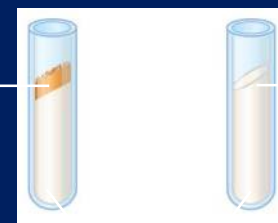
Complete
medium

Minimal
medium

Growth

No growth

Figure 17.2



Growth:
Wild-type
cells growing
and dividing

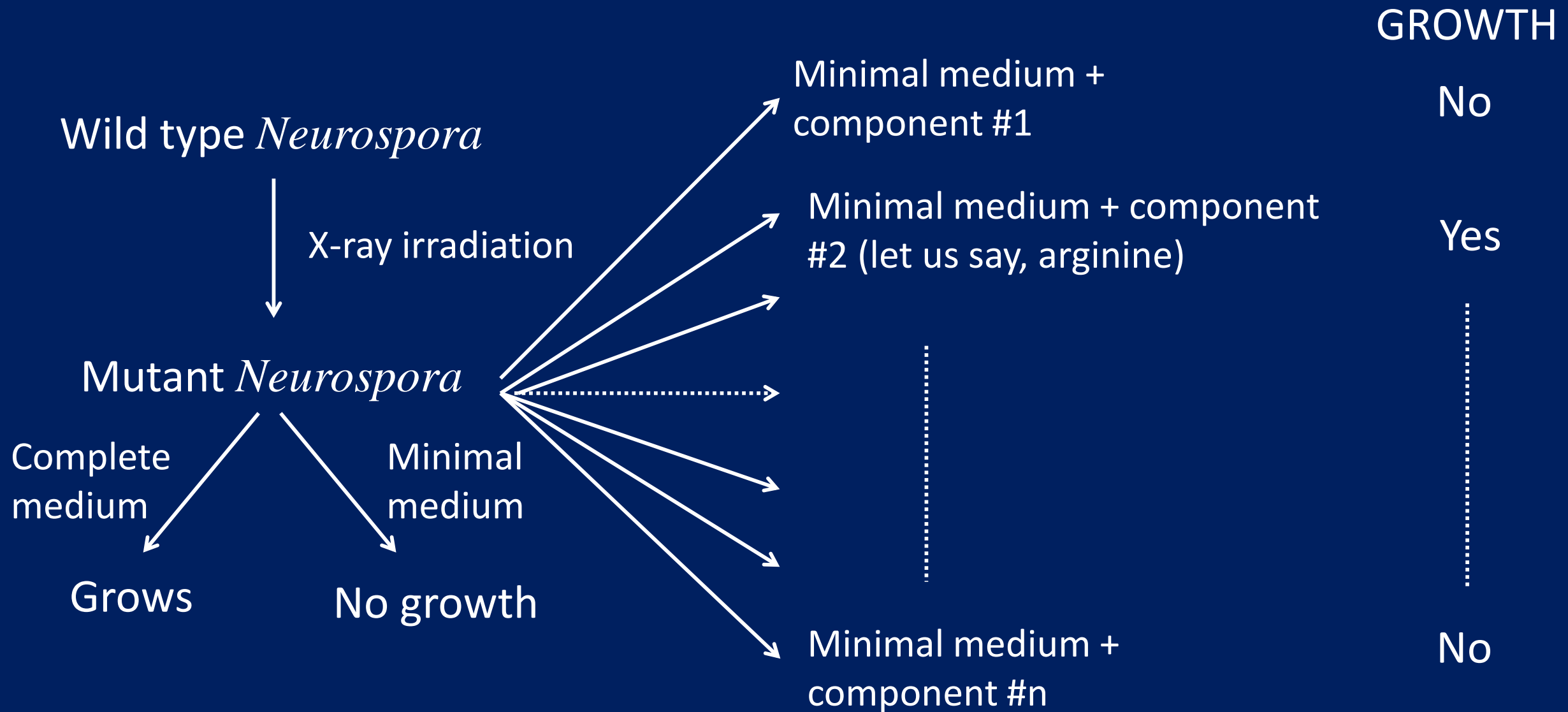
No growth:
Mutant cells
cannot grow
and divide

Control:
Minimal
medium

INFERENCE

Mutant is not able to synthesize
one or more essential components

Relationship between genes and enzymes



Mutant is NOT able to biosynthesize component #2 (e.g., arginine)

Finding out what genes do



Adrian M. Srb

Adrian Srb



Norman Horowitz

Norman Horowitz

Photographs are from the biographical memoirs available at the National Academy of Sciences (www.nasonline.org)

Used Beadle and Tatum's method to investigate the biochemical pathway for arginine biosynthesis

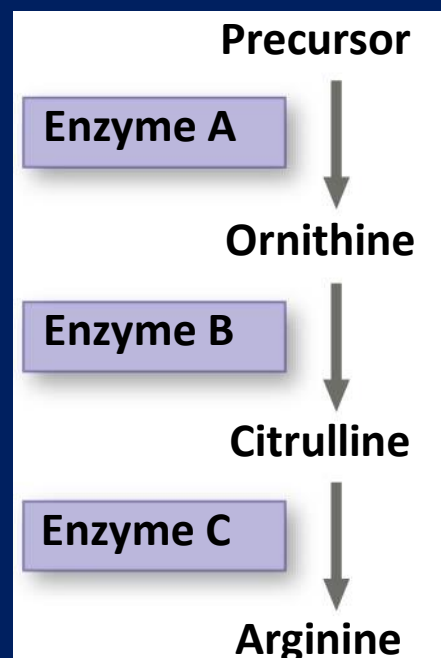
Arginine is one of the amino acids that is found in natural proteins

Worked at the Stanford University

A (part of a) metabolic pathway

Experiments of Adrian Srb and Norman Horowitz

















Figure 17.2



Pathway for arginine biosynthesis described in the mammalian liver

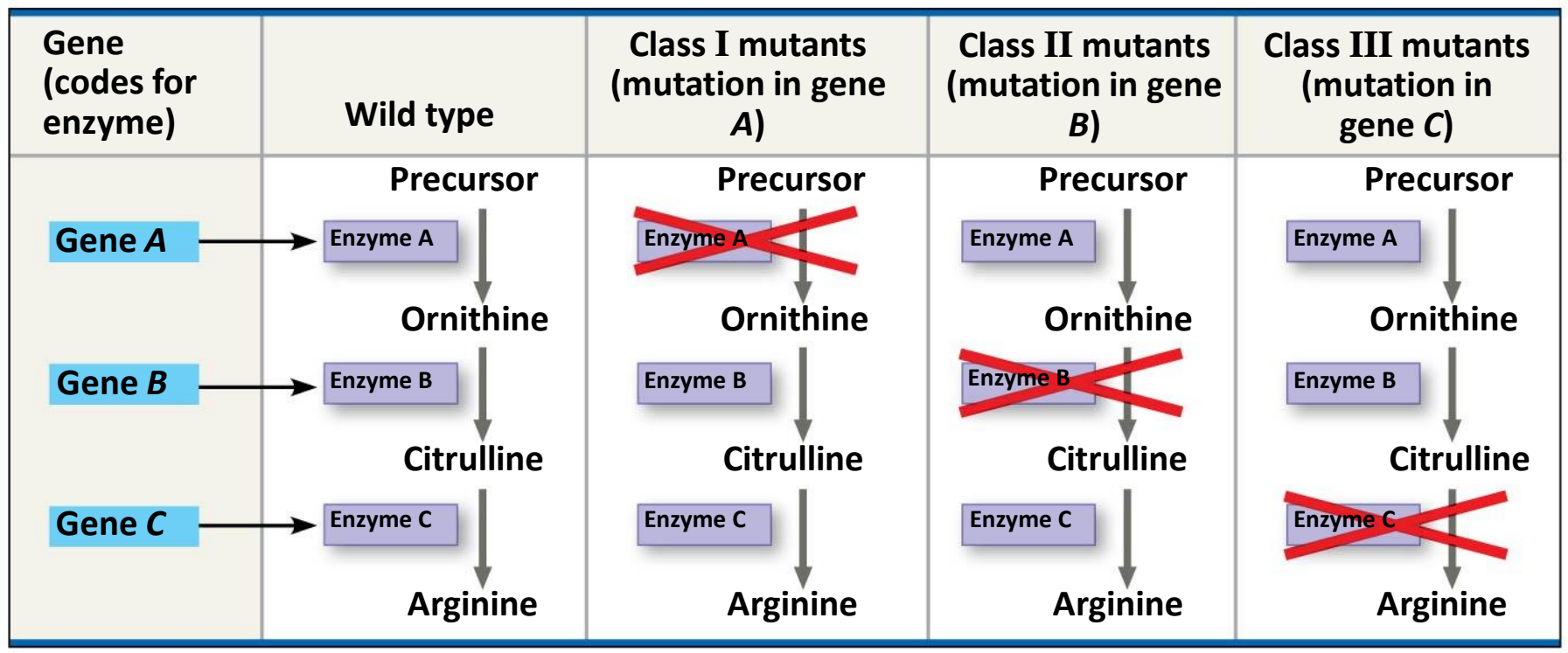
Arginine is one 20 amino acids that make up proteins in ALL organisms

Experimental set up

Results Table		Classes of <i>Neurospora crassa</i>				Figure 17.2
Condition		Wild type	Class I mutants	Class II mutants	Class III mutants	
	Minimal medium (MM) (control)					
	MM + ornithine					
	MM + citrulline					
	MM + arginine (control)					
	Summary of results	Can grow with or without any supplements	Can grow on ornithine, citrulline, or arginine	Can grow only on citrulline or arginine	Require arginine to grow	

Interpretation

Figure 17.2



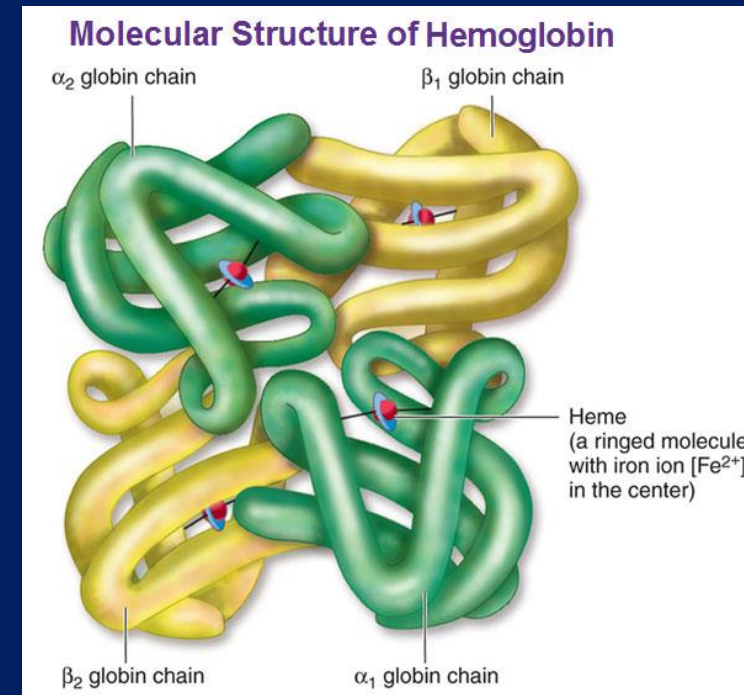
Information content of DNA and traits

- DNA inherited by an organism leads to specific traits by dictating the synthesis of proteins
- Proteins are the links between genotype and phenotype



One gene – one polypeptide

- Some proteins are not enzymes... one gene—one protein
- Many proteins have several polypeptides, each with its own gene
- Beadle and Tatum's hypothesis: one gene—one polypeptide



<http://antranik.org/blood-components-hemoglobin-typerh-factor-agglutination/>

Chromosomes and genes

- Observable traits can be inherited
 - Gregor Mendel's studies with pea plants
 - Flower color (white or purple)
 - Seed shape (round or wrinkled)
 - Pod color (yellow or green)
 - Thoman Morgan's studies on fruit fly
- Inherited traits are determined by genes
- Genes are present in chromosomes

Mendel's : Law of segregation

- Mendel crossed true-breeding purple flowered plants and white flowered plants.
- F1 hybrids were allowed to self-pollinate or were cross-pollinated with other F1 hybrids
- Both purple-flowered and white-flowered plants appeared in the F2 generation, in a ratio of approximately 3:1.
- The “heritable factor” for the recessive trait (white flowers) had not been destroyed, deleted, or “blended” in the F1 generation but was merely masked by the presence of the factor for purple flowers, which is the dominant trait.

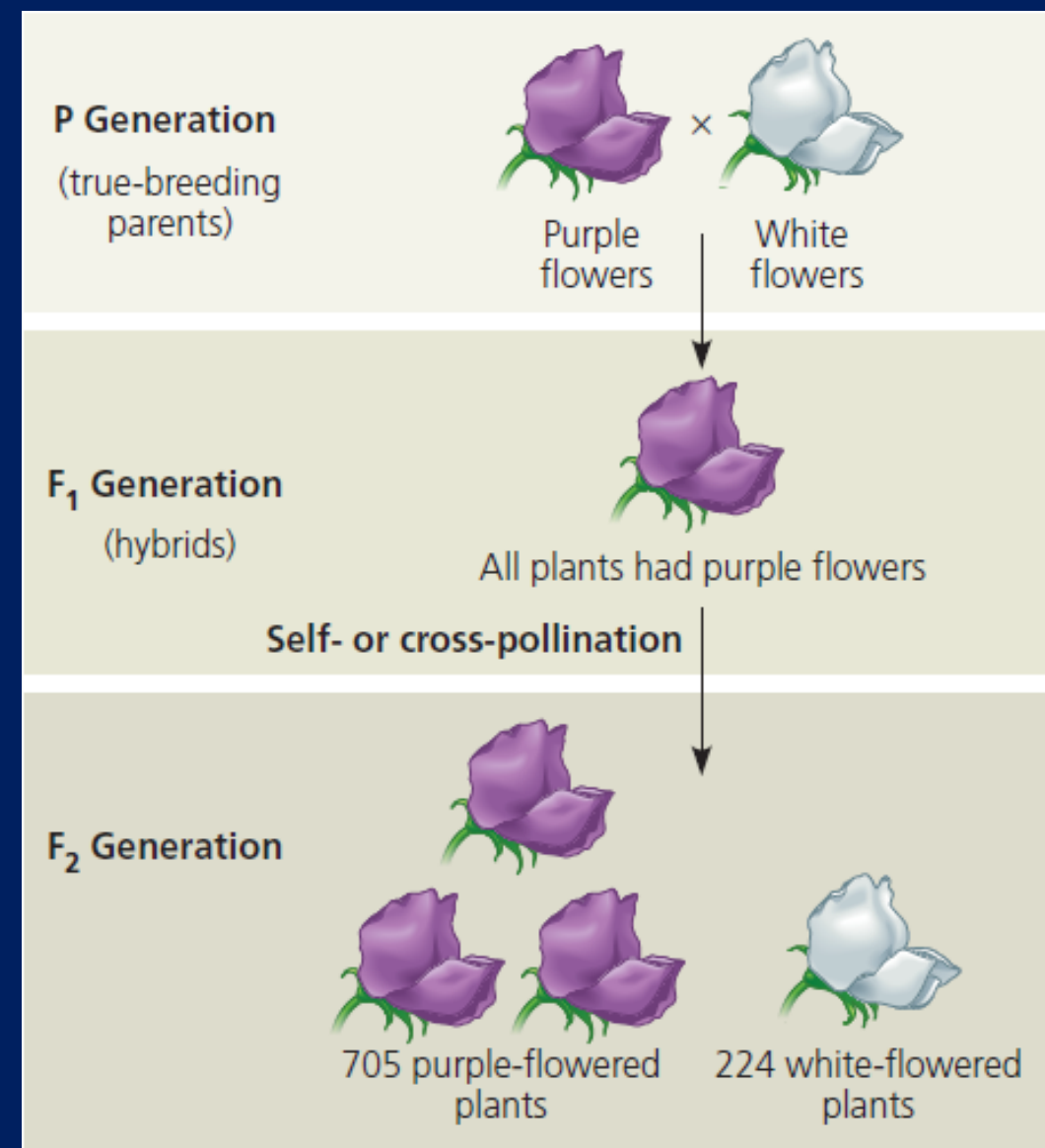


Figure 14.3

Mendel's Law of segregation

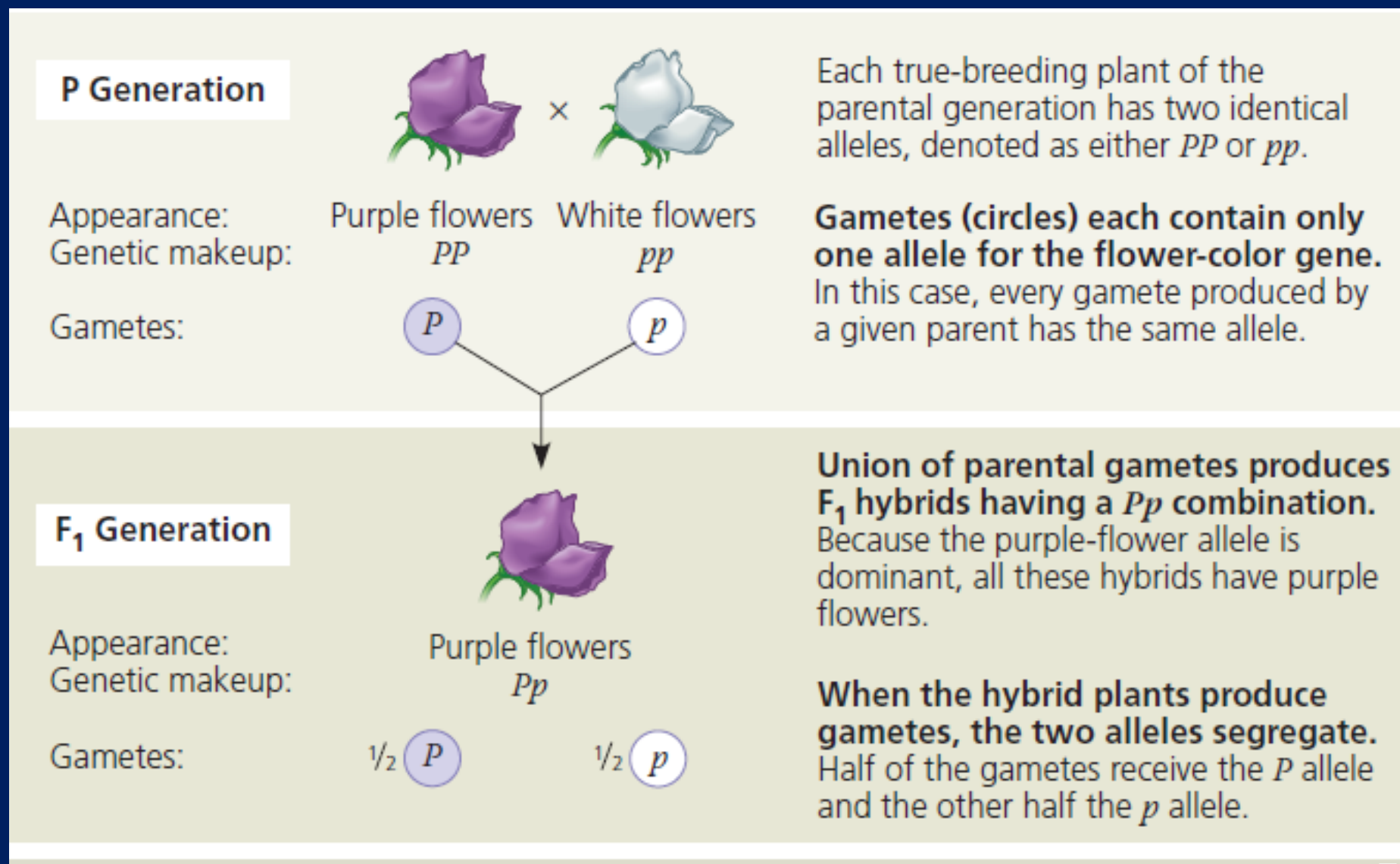
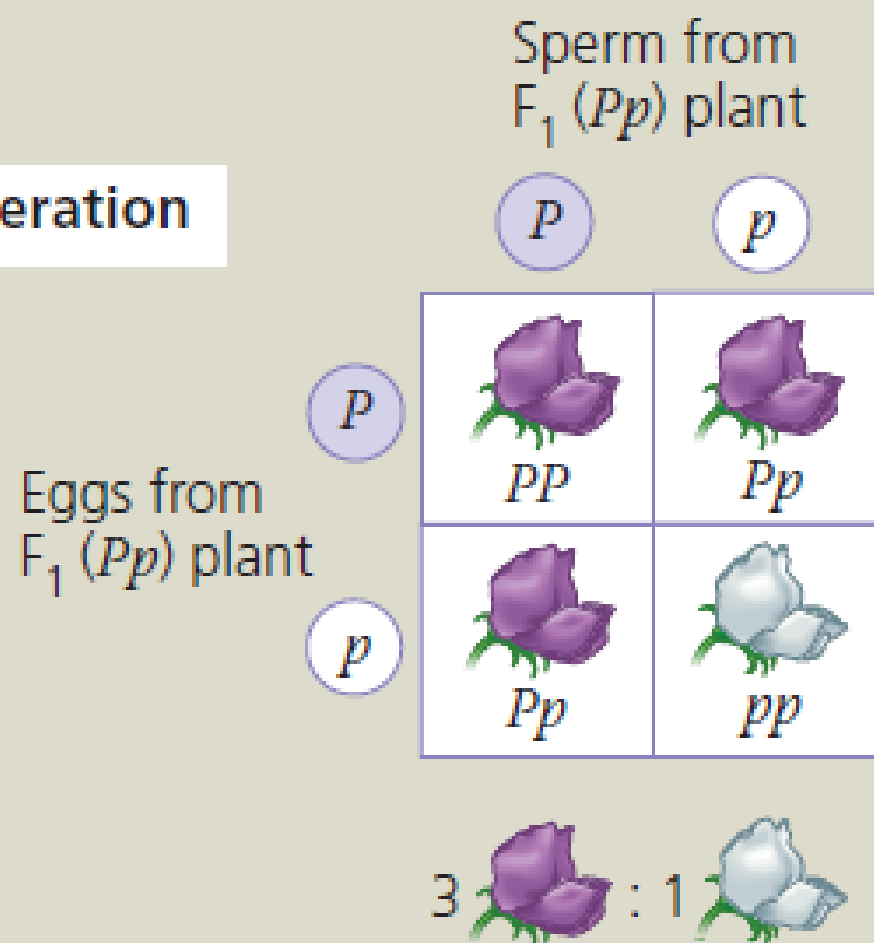


Figure 14.5

Mendel's Law of segregation

F₂ Generation



This box, a Punnett square, shows all possible combinations of alleles in offspring that result from an F₁ × F₁ (*Pp* × *Pp*) cross. Each square represents an equally probable product of fertilization. For example, the bottom left box shows the genetic combination resulting from a *p* egg fertilized by a *P* sperm.

Random combination of the gametes results in the 3:1 ratio that Mendel observed in the F₂ generation.

Figure 14.5

Mendel's Law of Independent Assortment

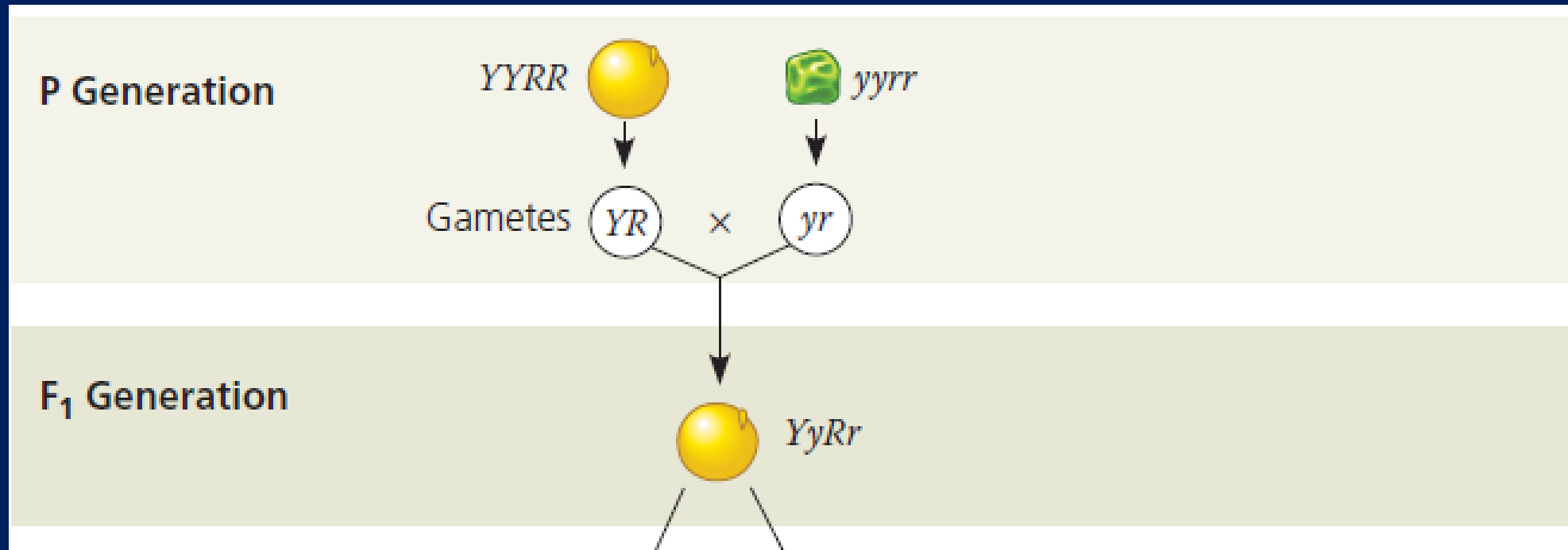


Figure 14.8

Mendel's Law of Independent Assortment

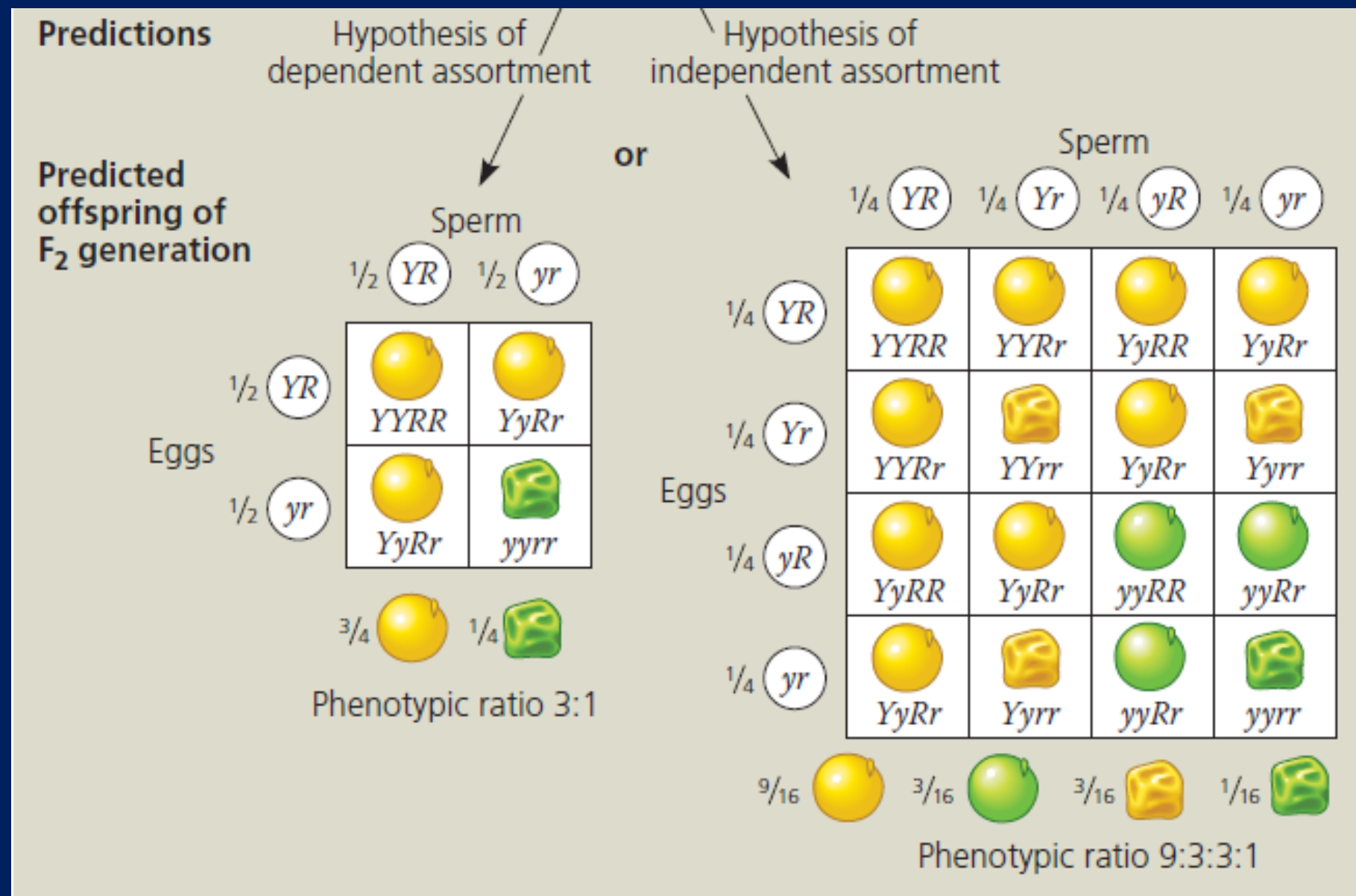


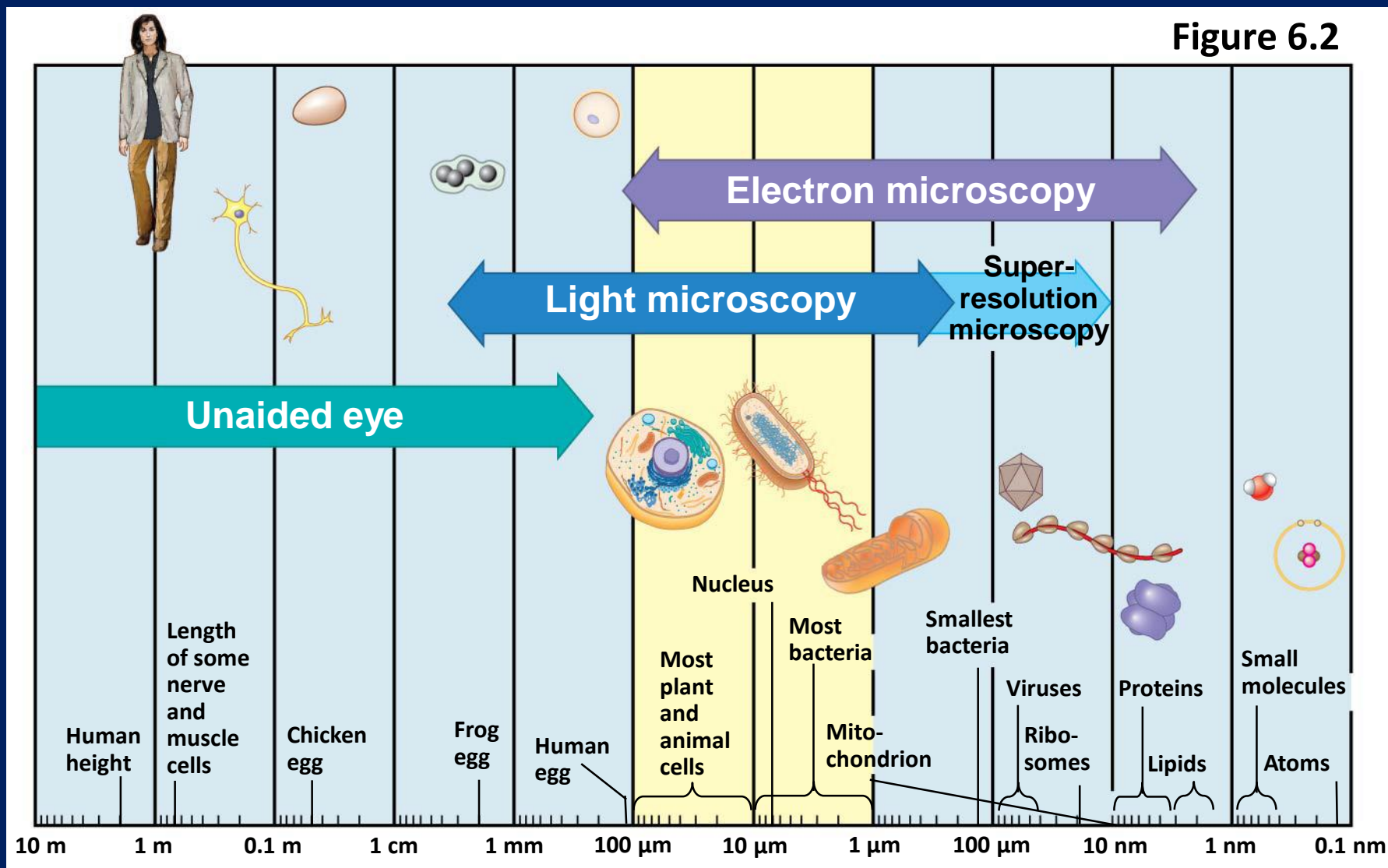
Figure 14.8

Class 4: learning objectives

- Inherited DNA leads to specific traits
 - Coat color, white eyes, phenylketonuria
- DNA: packaging, accessing the stored message
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Length comparison: DNA and cell

Figure 6.2



Logarithmic scale

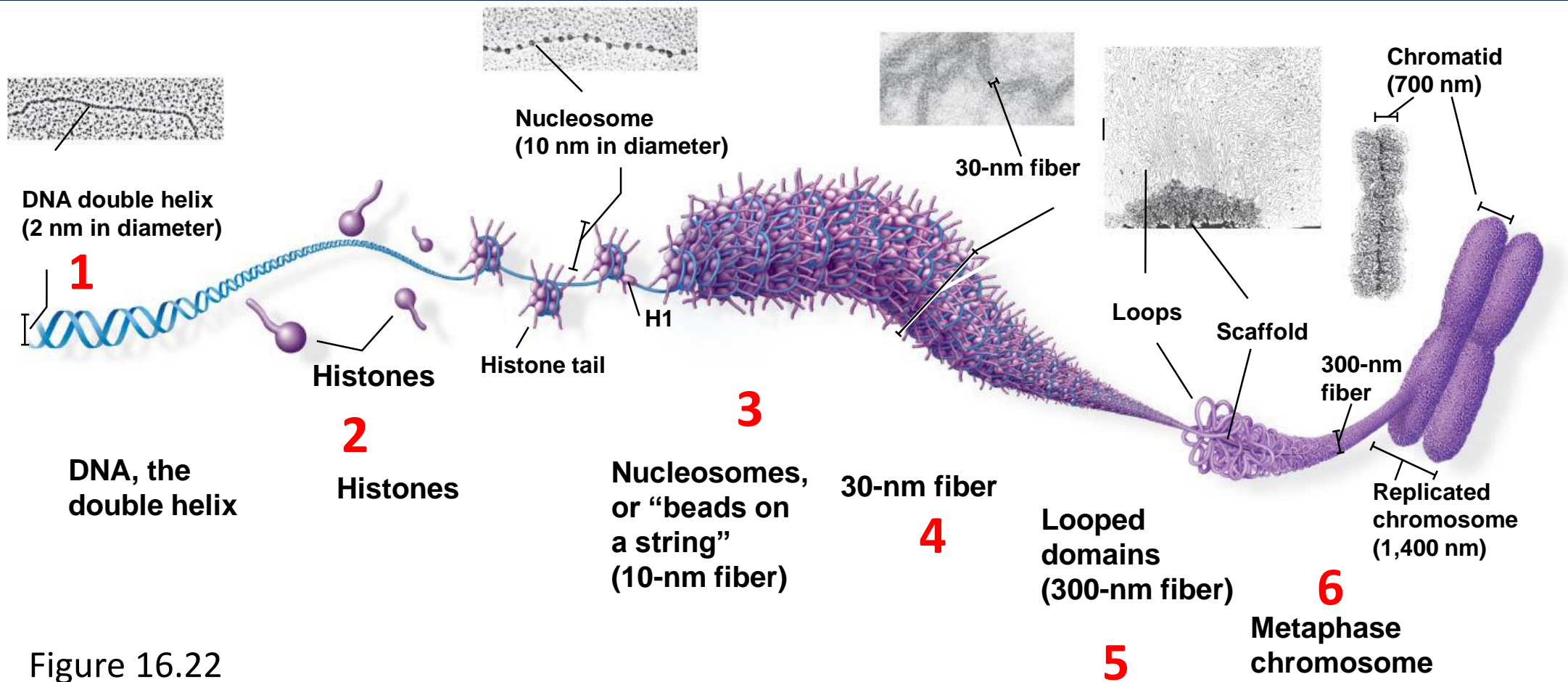
Length of DNA
Watson-Crick model

10 base pairs: 3.4 nm

Number of bases in
human chromosome
#1 is approximately
250 million base pairs

Length:
 $250 \times 10^6 \times 0.34 \text{ nm}$
85 mm

Packaging of DNA



1. Ribbon model of DNA. Each ribbon represents a DNA strand

Packaging of DNA

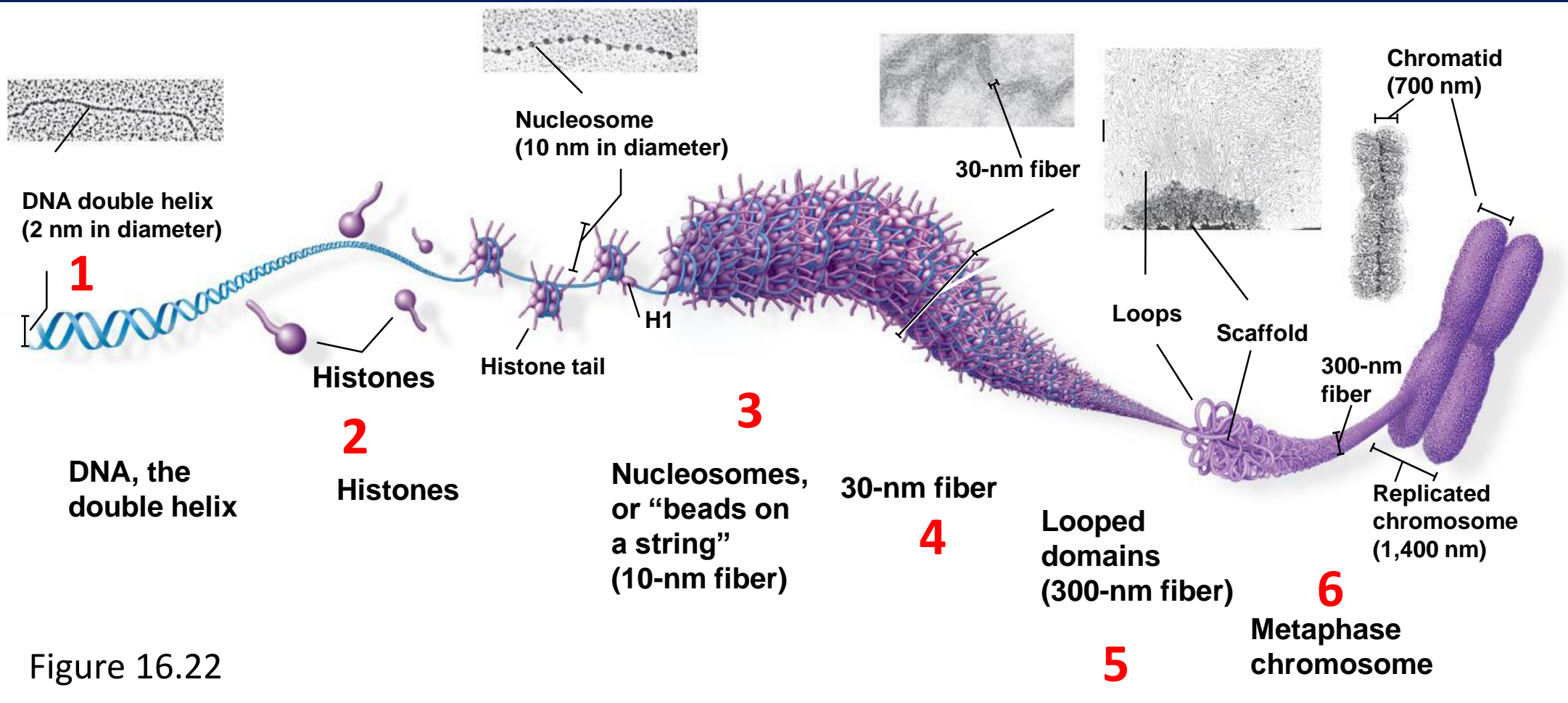
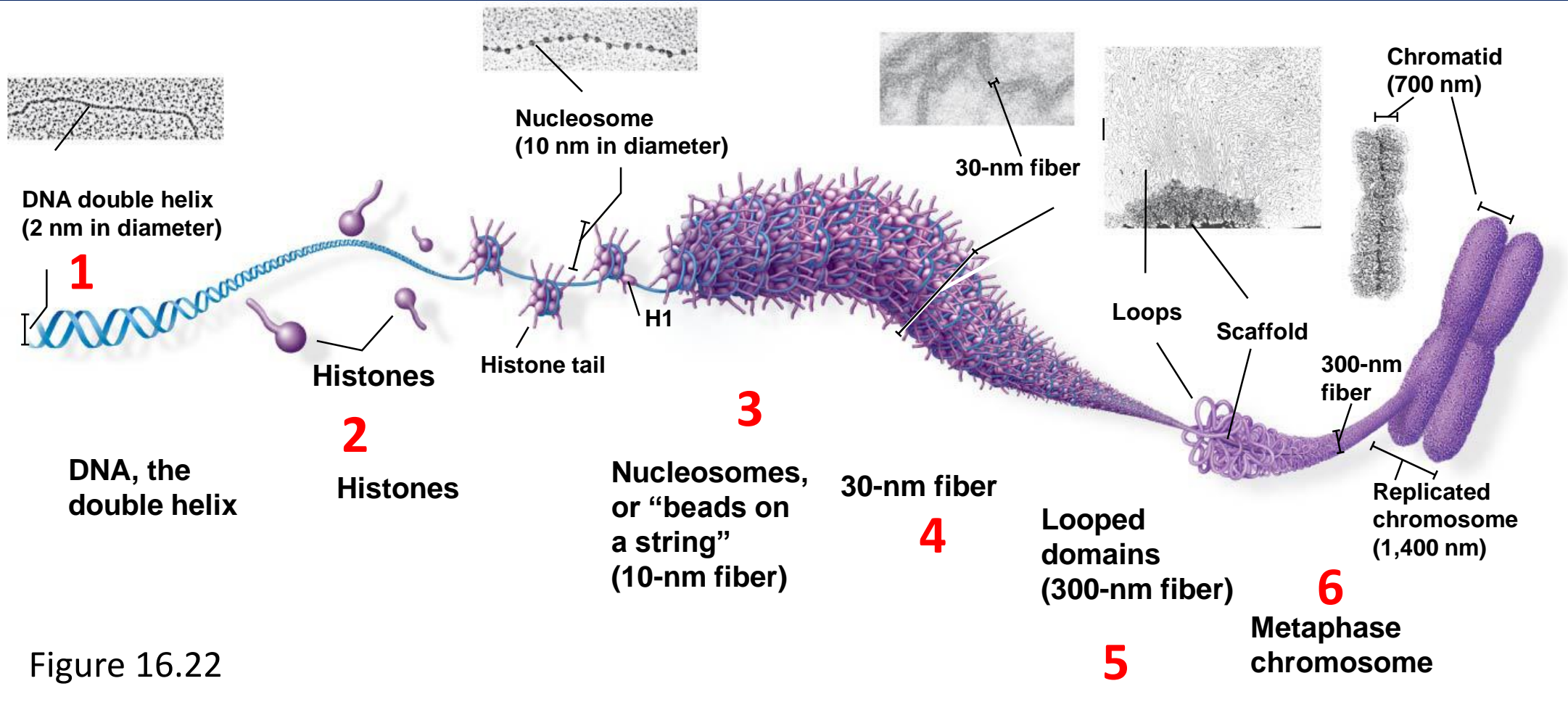


Figure 16.22

Chromatin = protein + DNA complex

2. First level of packing of DNA in the chromatin

Packaging of DNA



Bead = nucleosome; string – linker DNA

3. An aggregate of histones = resembles a ball of wool
 DNA is wound twice around histones = nucleosome

Packaging of DNA

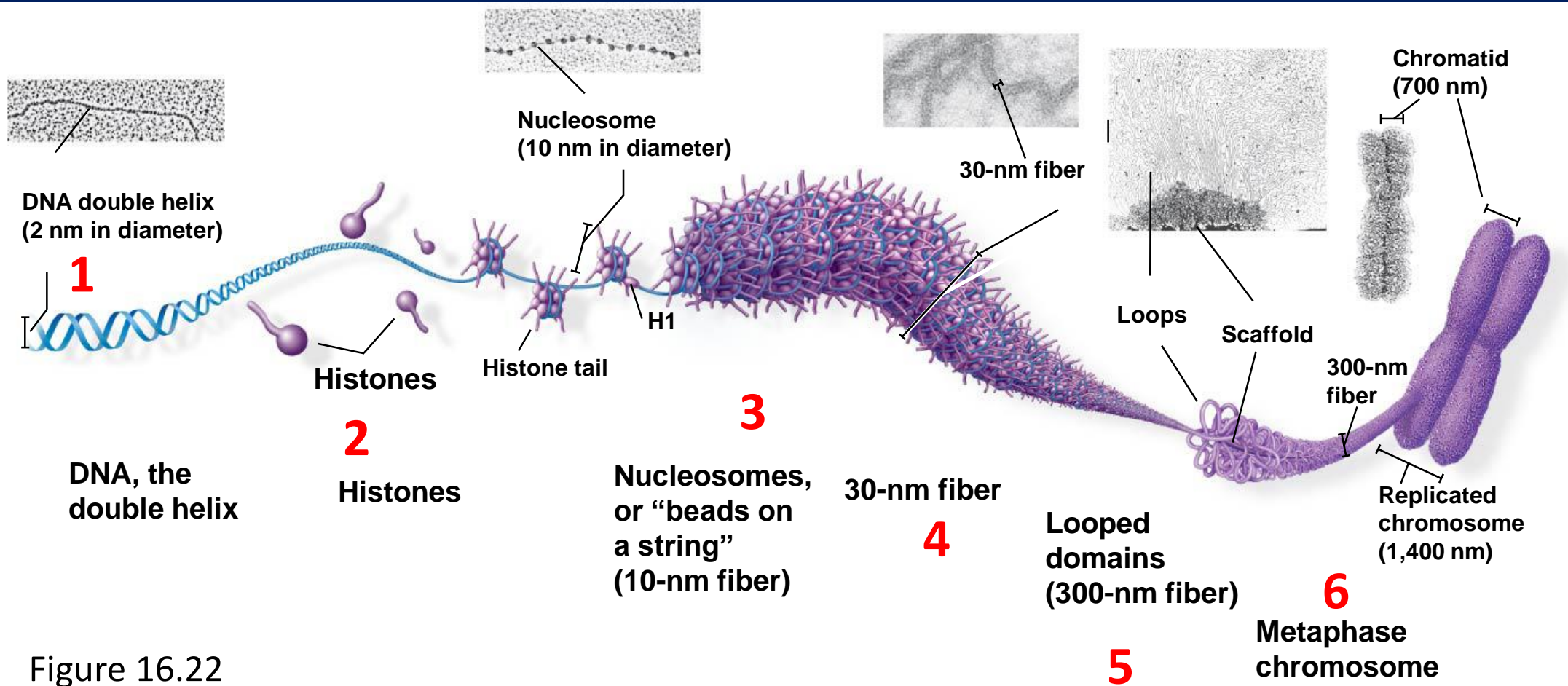


Figure 16.22

4. Second level of packing. Involves interactions of histone tails

Packaging of DNA

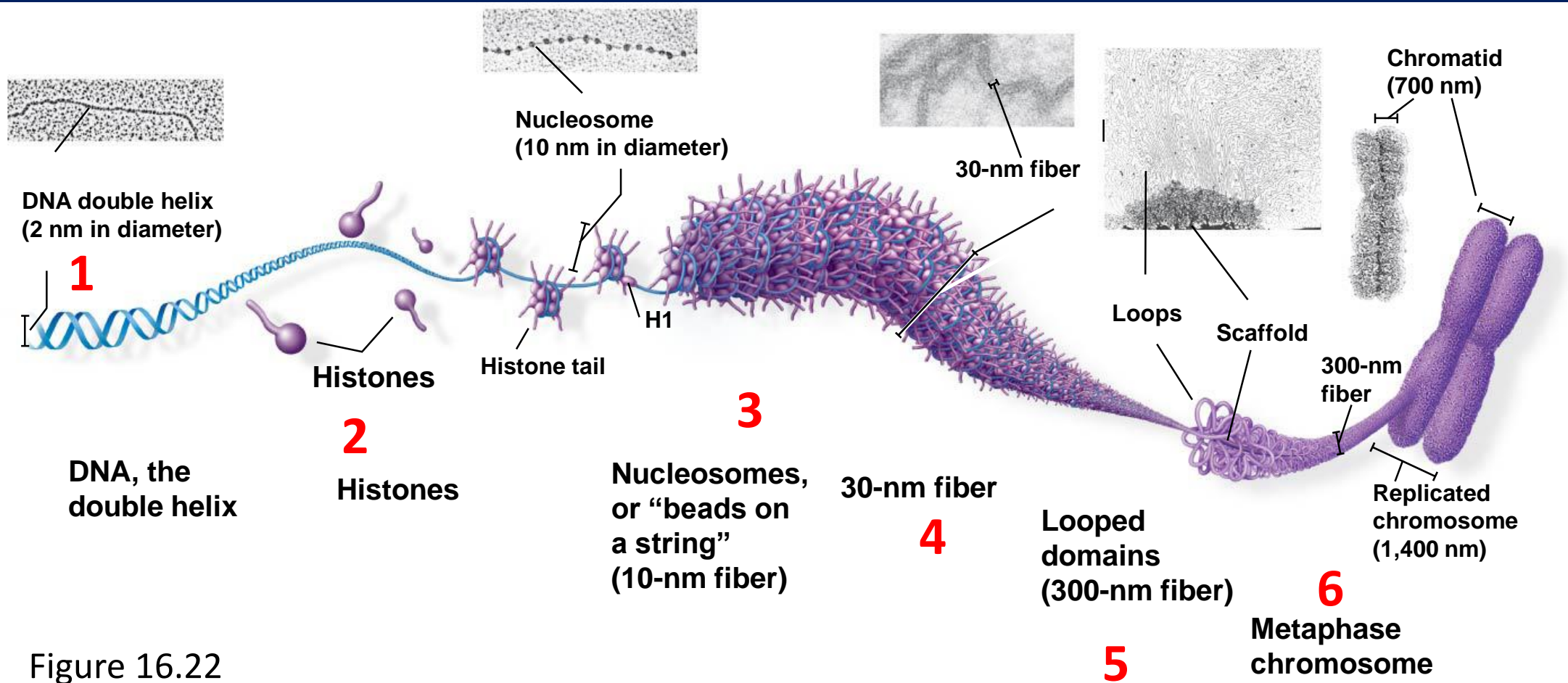
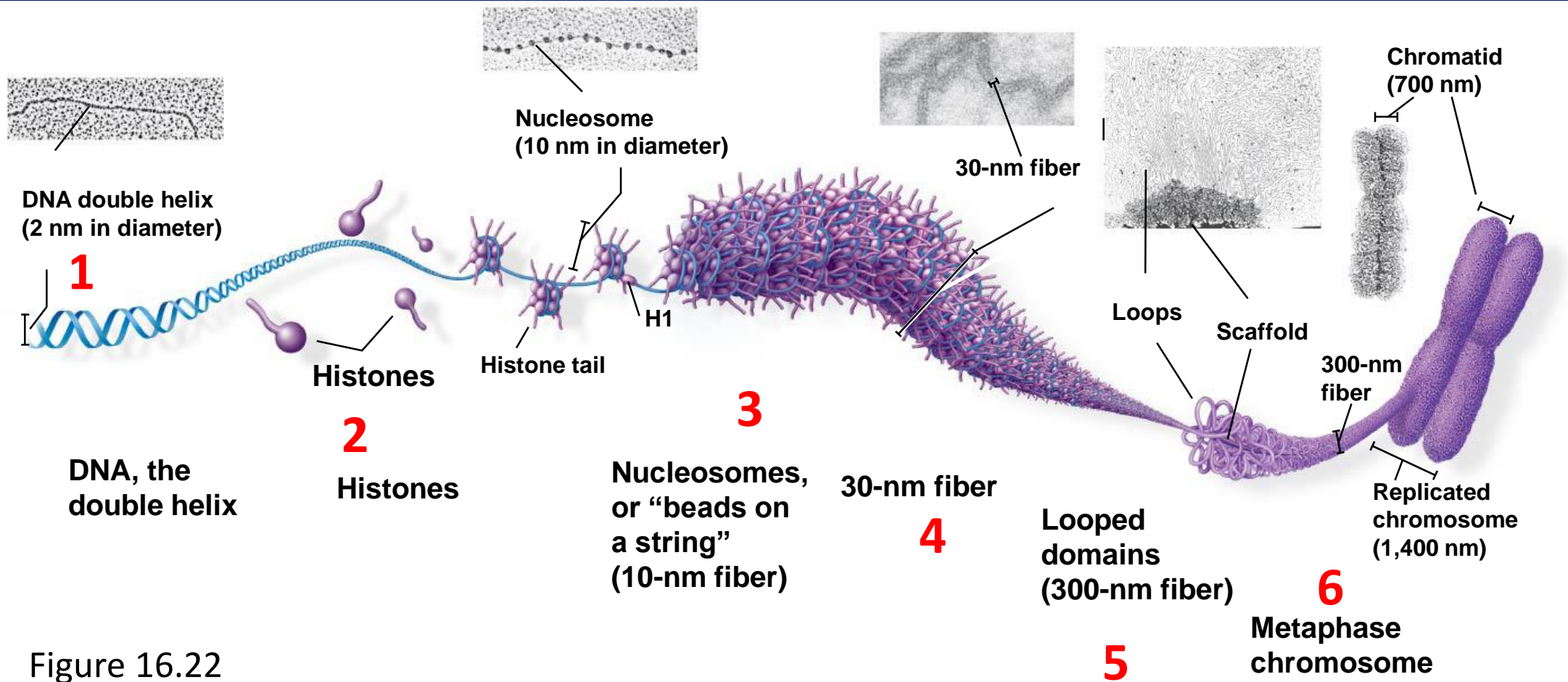


Figure 16.22

5. Formation of loops by 30-nm fibers

Packaging of DNA



6. Coiling and folding of 300-nm fibers leading to further compaction

Accessing books in a library



Books arranged in shelves



Shelves spaced out to facilitate access

Photo courtesy:
Sri Manju Naika
Central Library
IIT Bombay

Requires a lot of space

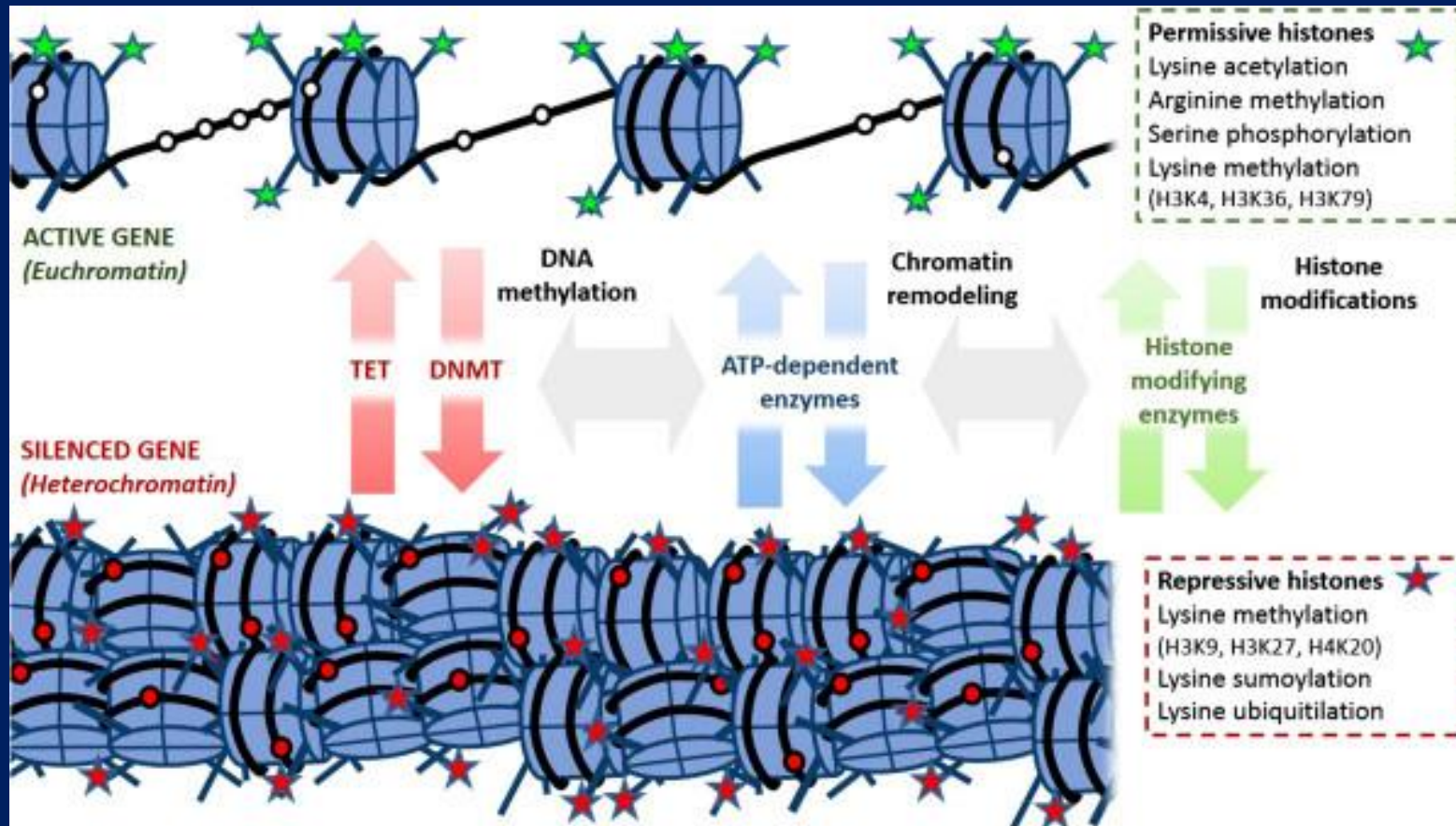
Not all shelves are accessed simultaneously

Sub-optimal space utilization

Packed and accessible forms

Euchromatin: dispersed, less compact, accessible form

Heterochromatin: compact form, not possible to access genes



This figure is taken from a research article published in the year 2014

Name of the journal: Allergy Asthma and Clinical Immunology

Volume 10; article # 27.

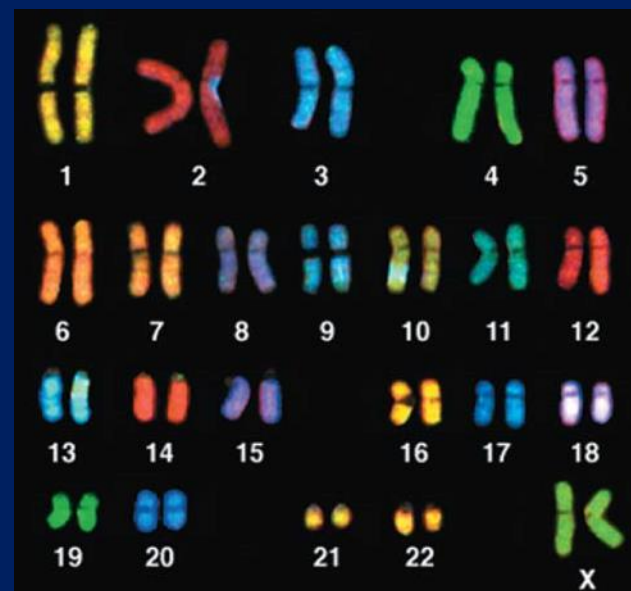
How are human chromosomes arranged?

Figure 16.23

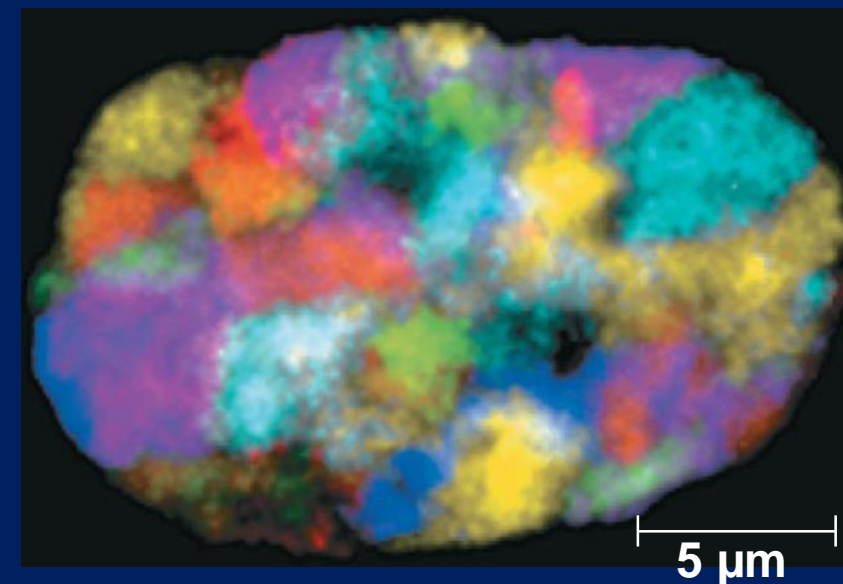
Chromosomes spread out



Chromosomes arranged into a karyotype



Tagging each chromosome such that it can be seen in a different color



Arrangement in a cell that is not undergoing cell division

Use as and when required



Encyclopedia comes in several volumes – all volumes are probably of same size

DNA of an organism also comes as ‘volumes’ called chromosomes

Chromosomes are of different sizes

Each chromosome (except XX or XY) come in two copies with ‘slightly’ different sequence

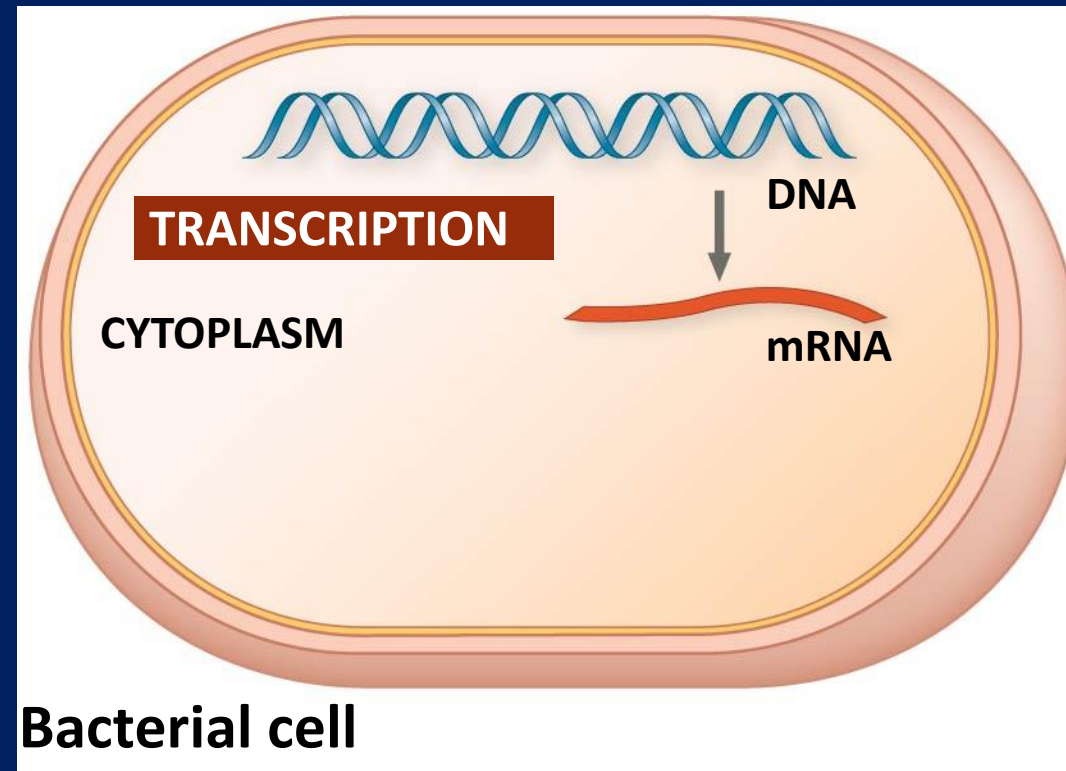
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Transcription in prokaryotes

Figure 17.3

Transcription:
synthesis of RNA
using information
in DNA

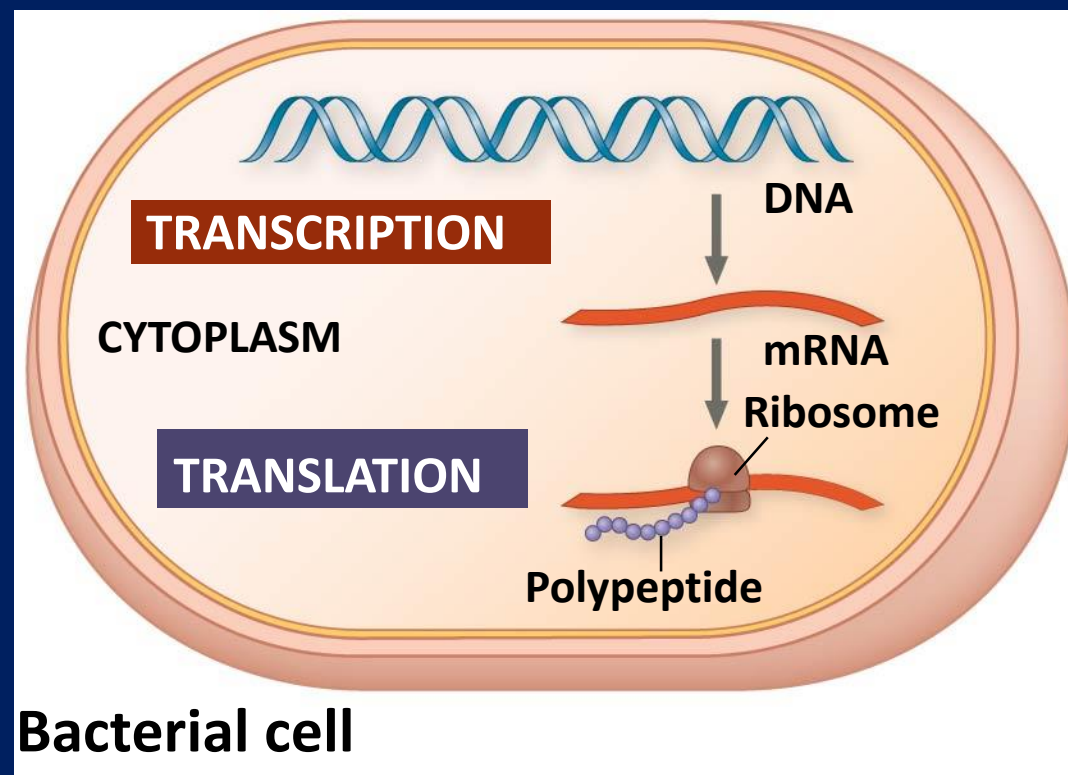


RNA: bridge
between genes
and proteins
that they
encode

Transcription produces messenger RNA (mRNA)

Translation in prokaryotes

Figure 17.3



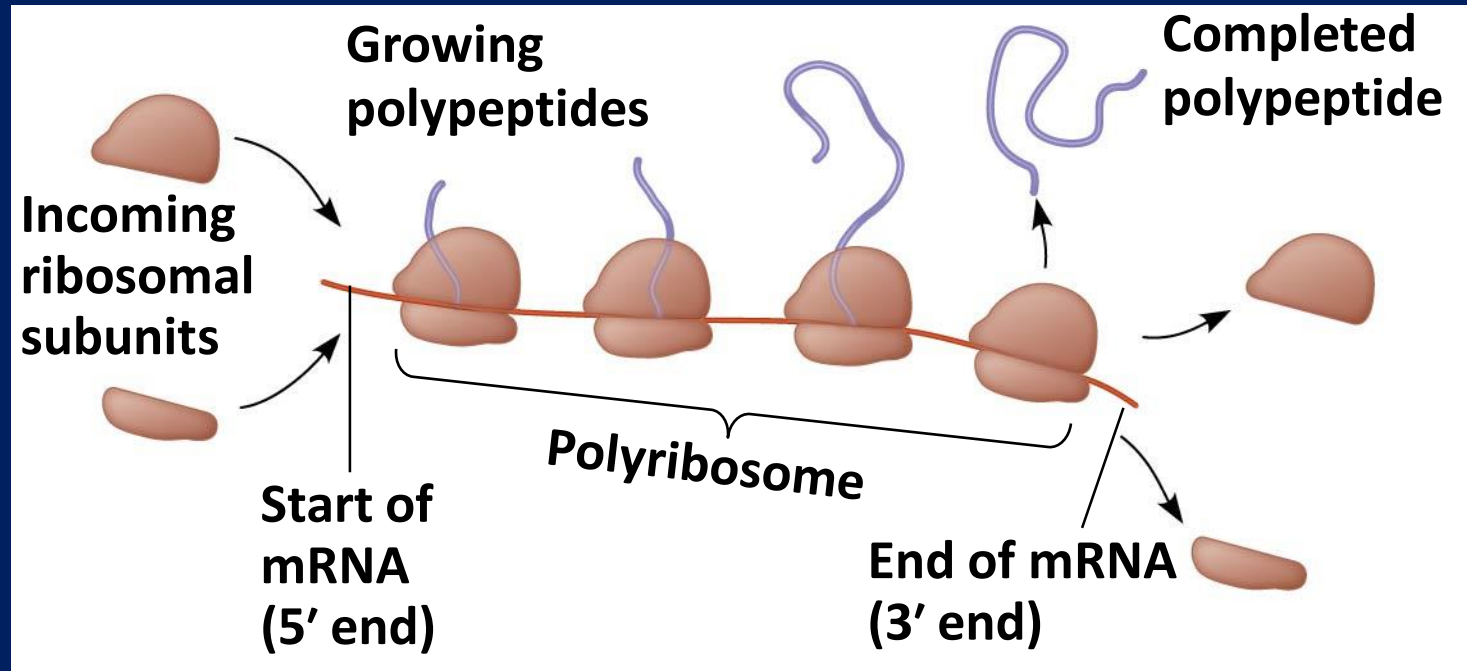
Ribosomes are the sites of translation

Translation: synthesis of a polypeptide, using information in the mRNA

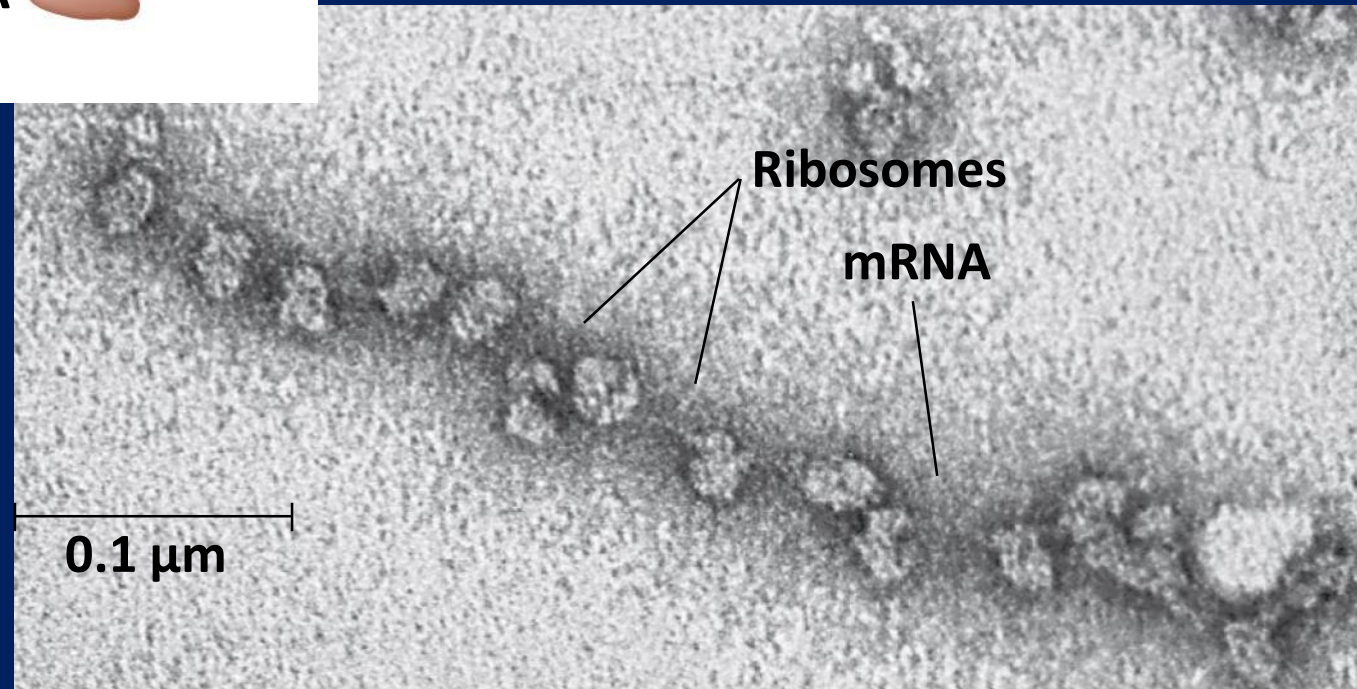
Translation can be concurrent with transcription

Poly-ribosomes

Figure 17.21

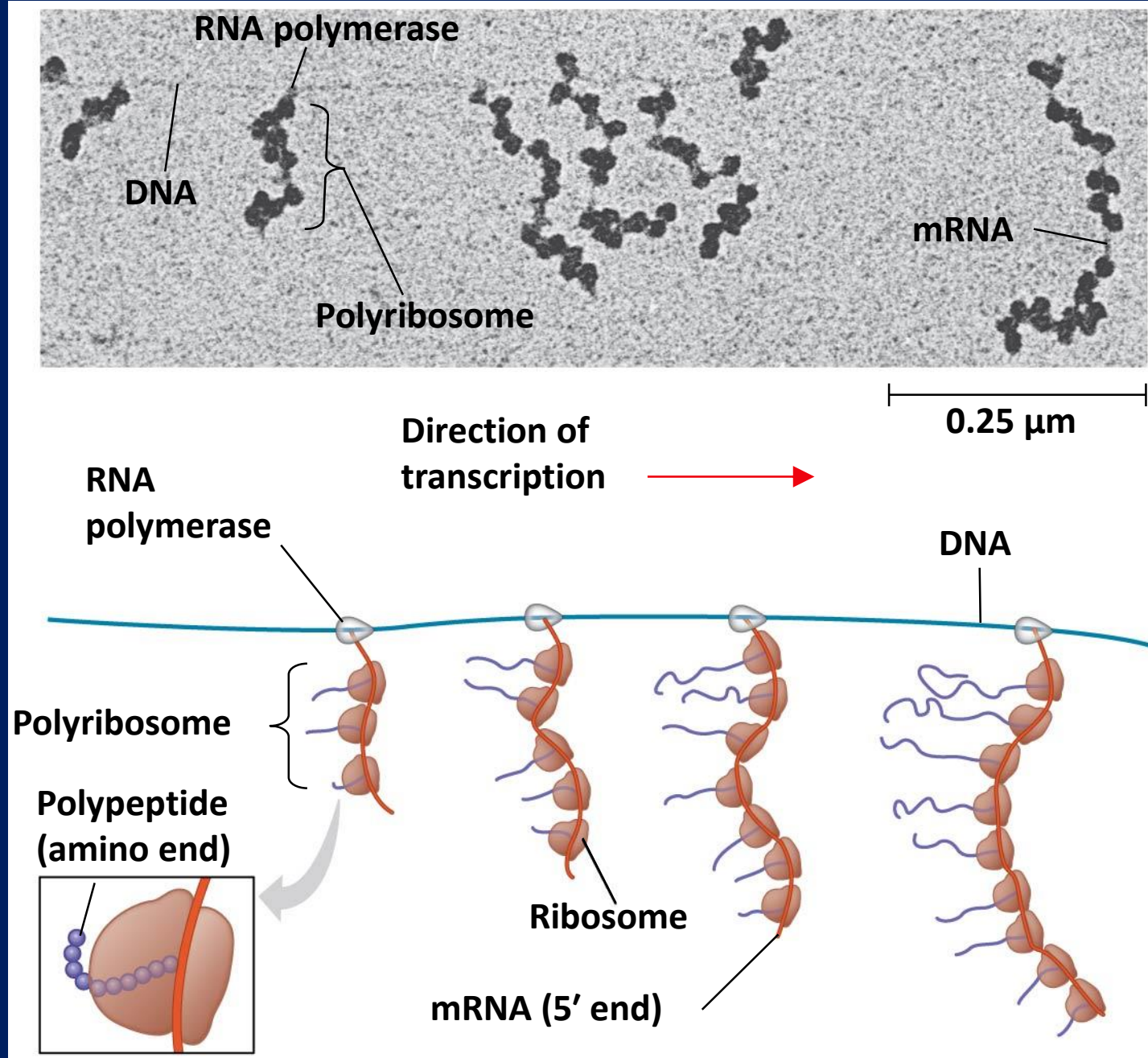


Several copies of a polypeptide chain are made simultaneously



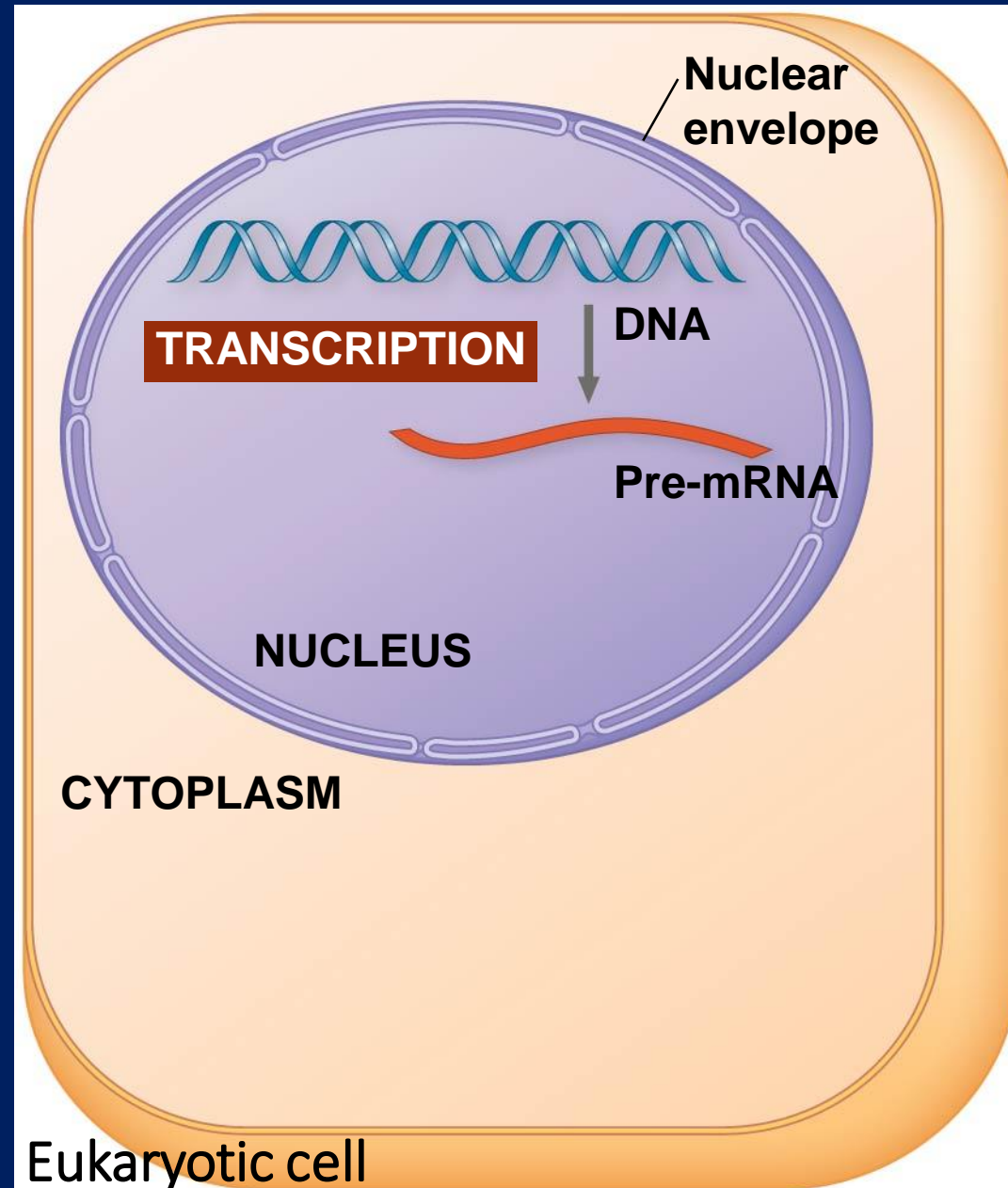
Coupling transcription and translation

Figure 17.25



Transcription in eukaryotes

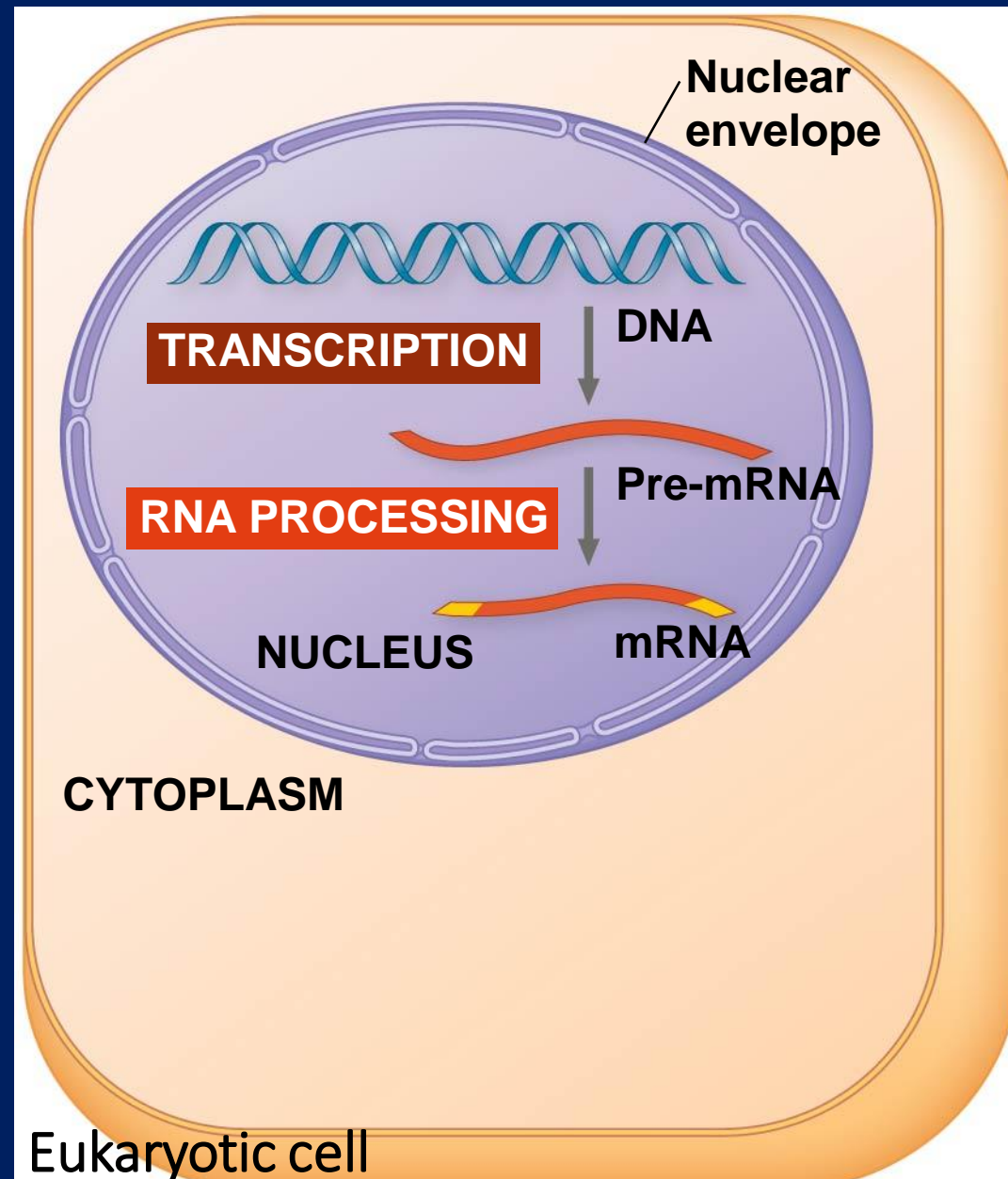
Figure 17.3



mRNA undergoes processing

Figure 17.3

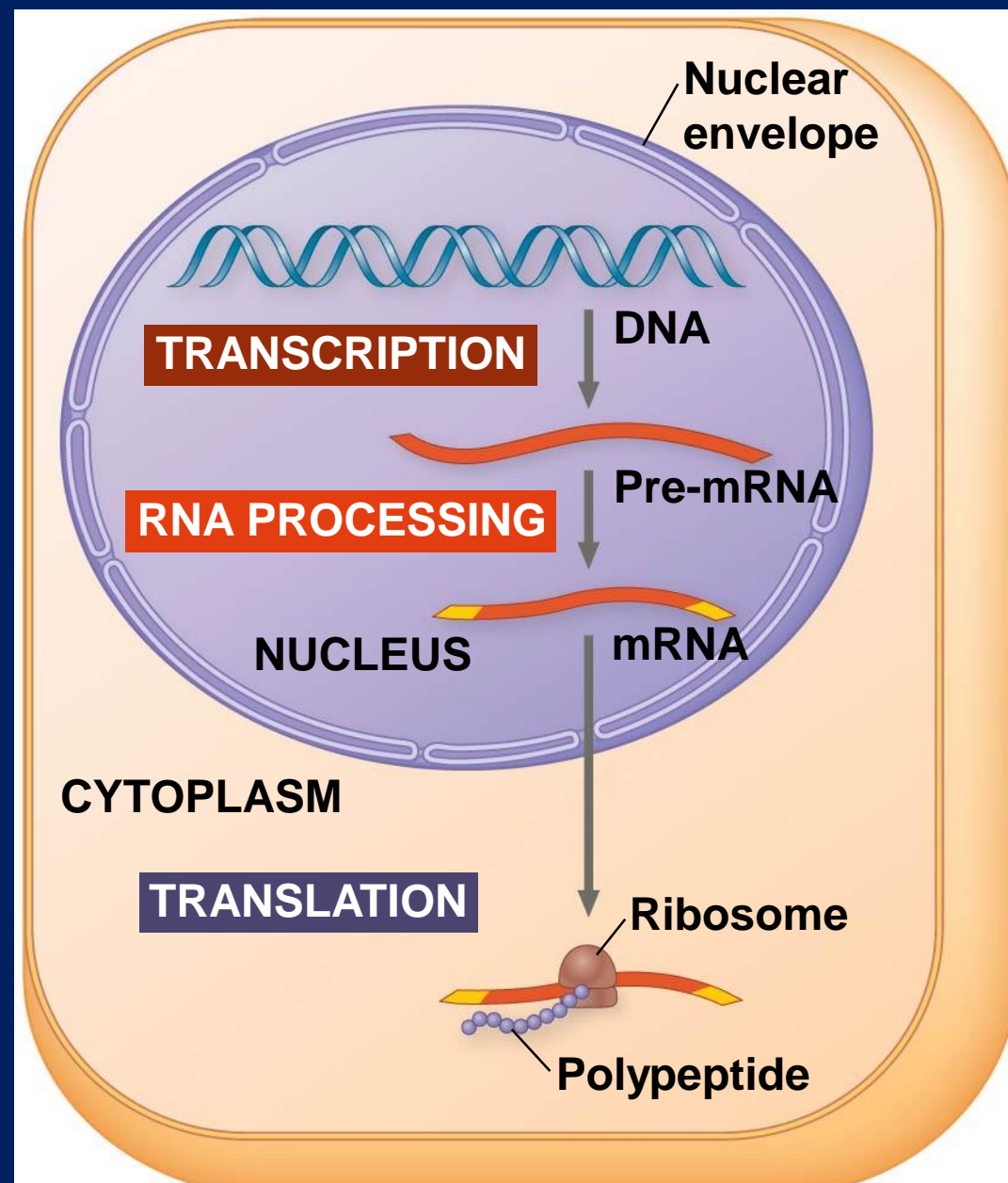
Primary transcript:
initial mRNA
transcript prior to
processing



mRNA is modified
through RNA
processing to yield
the finished mRNA

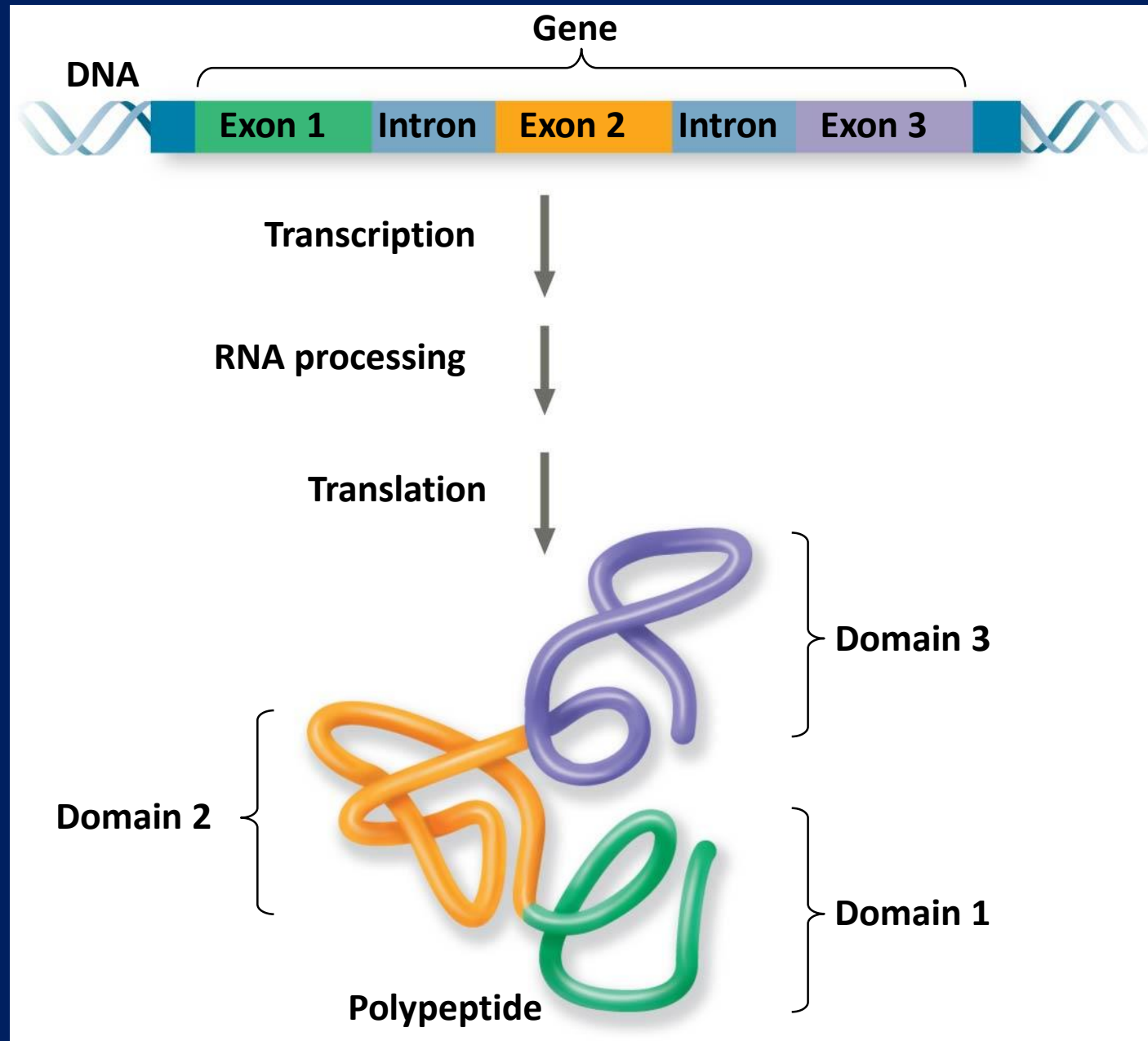
Translation in eukaryotes

Figure 17.3



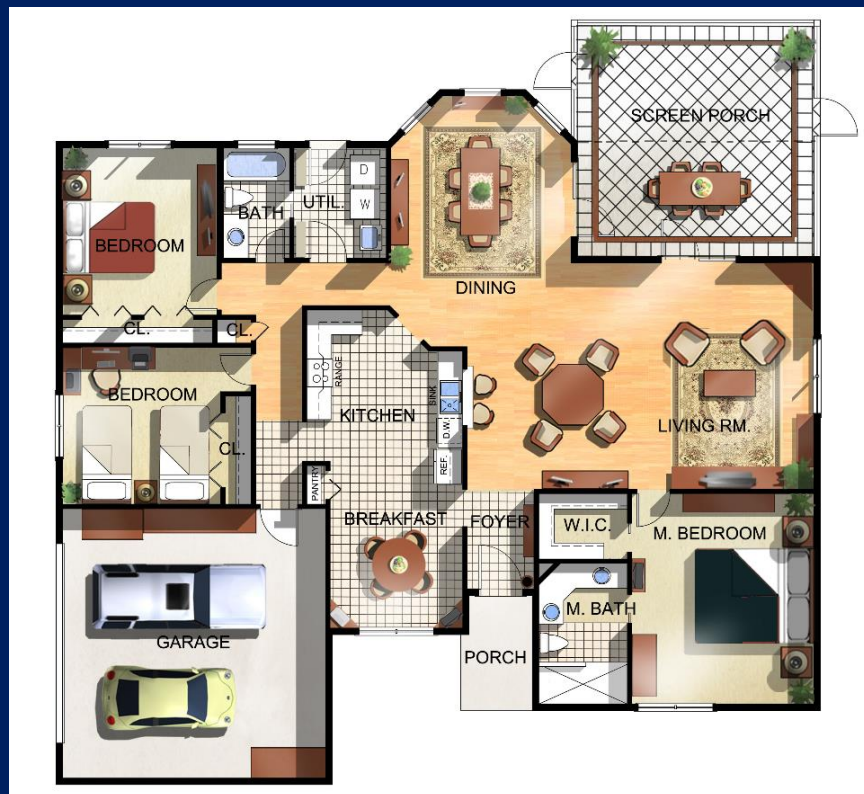
Eukaryotes: nuclear envelope separates transcription from translation

RNA processing



Compartmentalization

Floor plan of an affluent home



http://zenlibs.com/a_floorplan-of-a-house/

A 1-room tenement

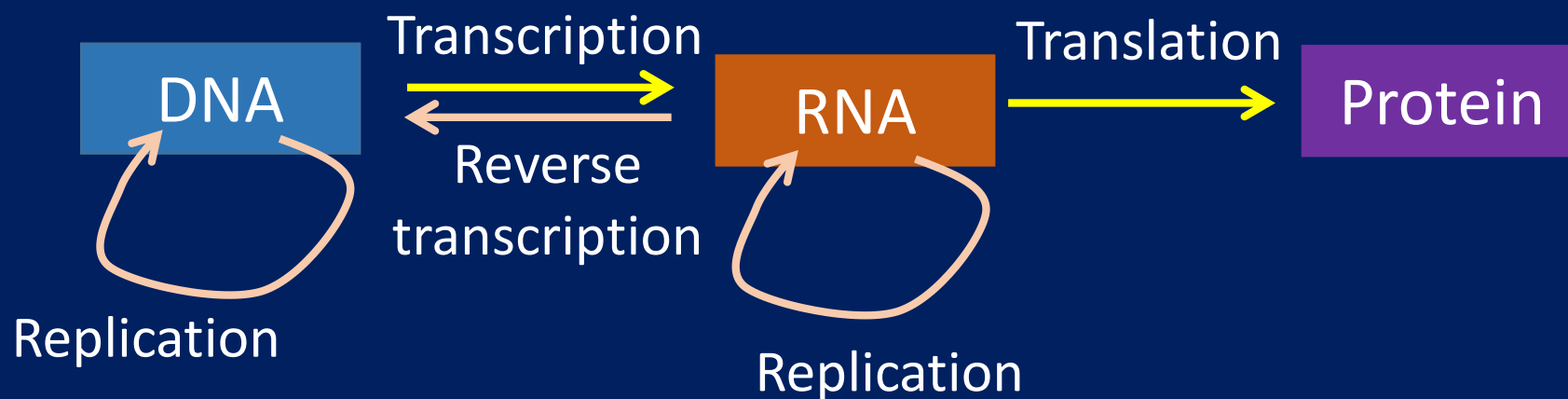


www.studenthandouts.com/01-Web-Pages/01-Picture-Pages/10.07-Industrial-Revolution/1-Riis-Family-Living-in-One-Room-New-York-City-Slum-1890.htm

Tenement: a room that by itself is a residence

Central dogma

Central dogma is the concept that cells are governed by a cellular chain of command



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DNA and protein alphabets

How are instructions for assembling amino acids into proteins encoded into DNA?

The alphabet of English: 26 letters

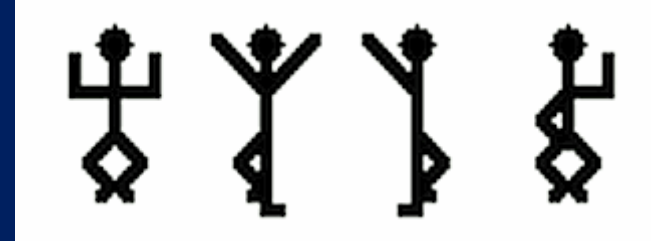
The alphabet of Hindi (or Assamese or ...): __ letters

The alphabet of DNA: 4 letters (A, C, G, T)

The alphabet of protein: 20 letters (names intentionally omitted here)

Decoding the genetic code

Sherlock Holmes in The dancing men



How can one possibly use 4 dancing men to send messages in a script that contains 20 letters?

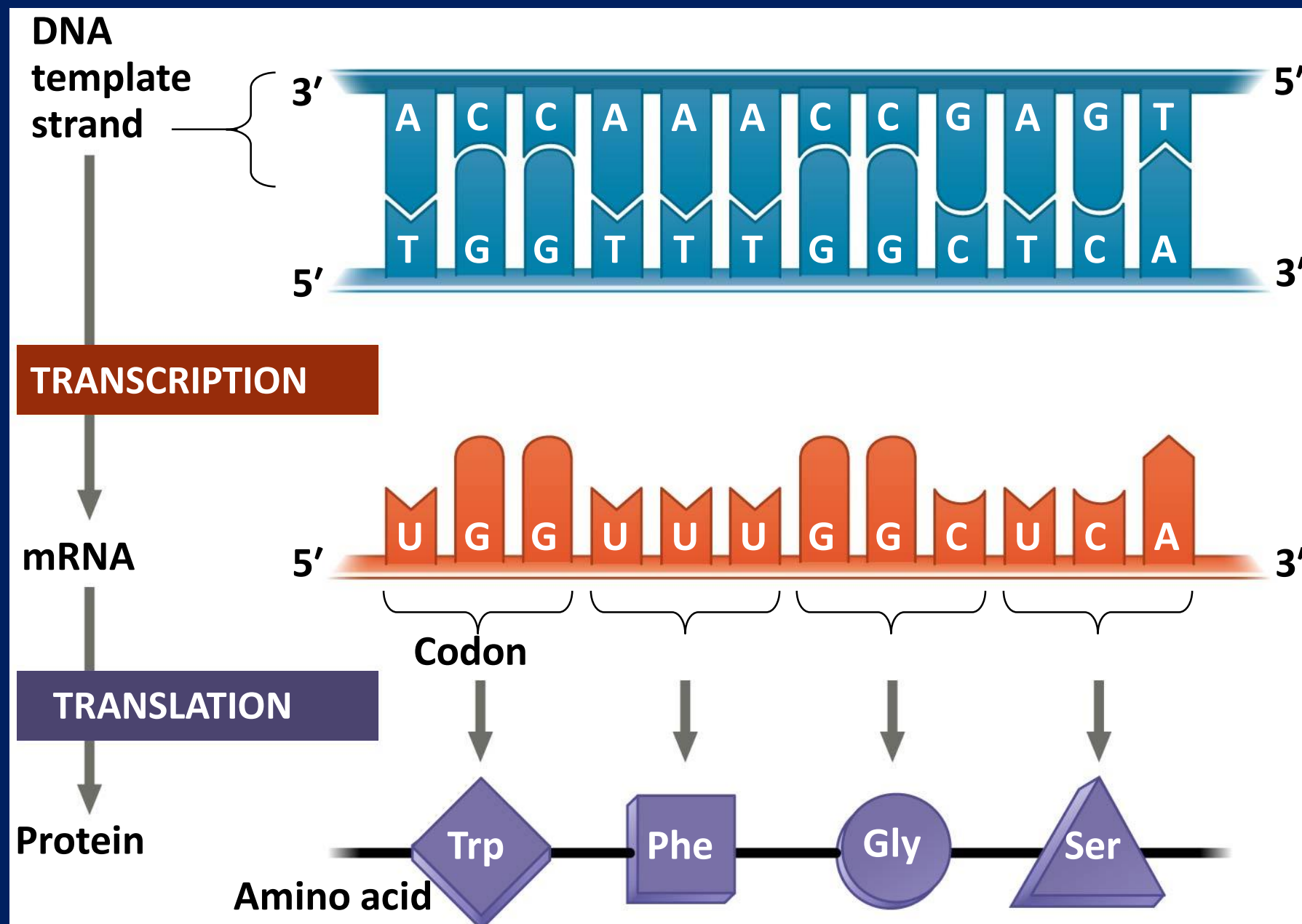
$$4^1 = 4$$

$$4^2 = 16$$

$$4^3 = 64$$

Non-overlapping triplet code

Figure 17.4



Is there an equivalent of full stop?

Template strand

3' -T A C T T C A A A C C G A T T-5'

5' -A T G A A G T T T G G C T A A-3'

mRNA

5' -A U G A A G U U U G G C U A A-3'

Protein

Met

Lys

Phe

Gly

Stop

The “full stop” or stop codons: UAA, UAG and UGA

The triplet genetic code: problem of plenty?

- Number of types of bases in DNA 4
- Number of types of amino acids in proteins 20 + at least 1 full stop
- Number of possible triplet codons 64

What about the additional $44 - 1 = 43$ triplet codons?

The degenerate genetic code

Refer to
figure 17.25

Codon	Amino acid
UUU	Phenyl alanine
UUC	Phenyl alanine
UUA	Leucine
UUG	Leucine
GUU	Valine
GUC	Valine
GUA	Valine
GUG	Valine

Substitution mutation. 1. Silent mutation

Refer to
figure 17.25

Codon	Amino acid
UUU	Phenyl alanine
UUC	Phenyl alanine
UUA	Leucine
UUG	Leucine
GUU	Valine
GUC	Valine
GUA	Valine
GUG	Valine

Mutating UUU to UUC

Same amino acid
Mutation is “silent”

Substitution mutation. 2. Missense mutation

Refer to
figure 17.25

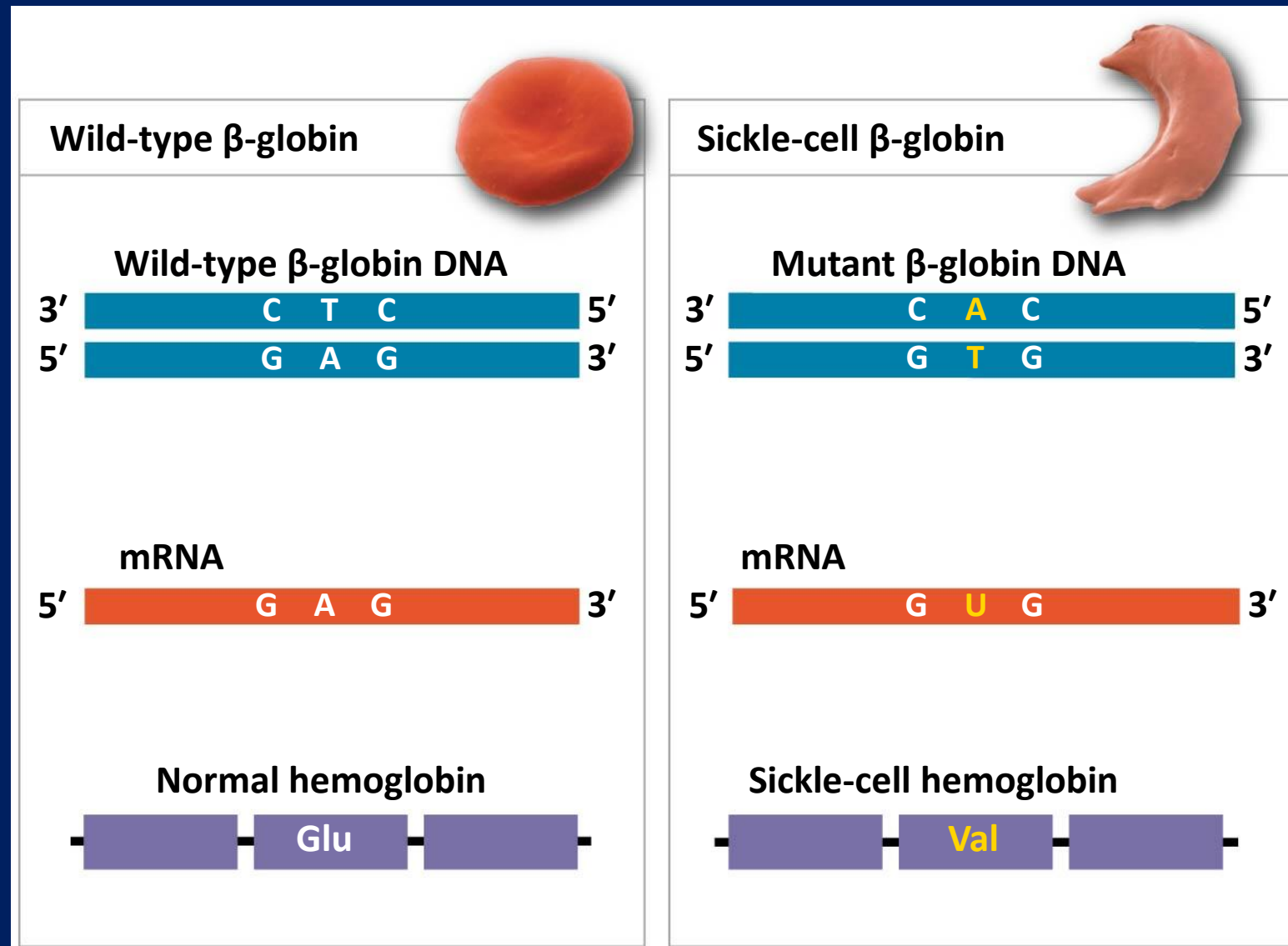
Codon	Amino acid
UUU	Phenyl alanine
UUC	Phenyl alanine
UUA	Leucine
UUG	Leucine
GUU	Valine
GUC	Valine
GUA	Valine
GUG	Valine

Mutating UUU to GUU

Valine in place of phenyl alanine
Mutation is “mis-sense”

Sickle cell anaemia

Figure 17.25



Substitution mutation. 3. Nonsense mutation

Refer to
figure 17.25

Codon	Amino acid
UUU	Phenyl alanine
UUC	Phenyl alanine
UUA	Leucine
UUG	Leucine
GUU	Valine
GUC	Valine
GUA	Valine
GUG	Valine

Mutating UUA to UAA

UAA is a STOP codon

Protein synthesis stops

Mutation is termed “non-sense”

Insertion or deletion: frameshift mutation

Refer to
figure 17.25

Template strand

3' -T A C T T C A A A C C G A T T-5'
5' -A T G A A G T T T G G C T A A-3'

mRNA

5' -A U G A A G U U U G G C U A A-3'

Template strand

3' -T A C G T T C A A A C C G A T T-5'
5' -A T G C A A G T T T G G C T A A-3'

mRNA

5' -A U G A A G U U U G G C U A A-3'
5' -A U G C A A G U U U G G C U A A-3'

Universality of the genetic code

Figure 17.6



(a) Tobacco plant expressing a firefly gene



(b) Pig expressing a jellyfish gene