



Chapter 5

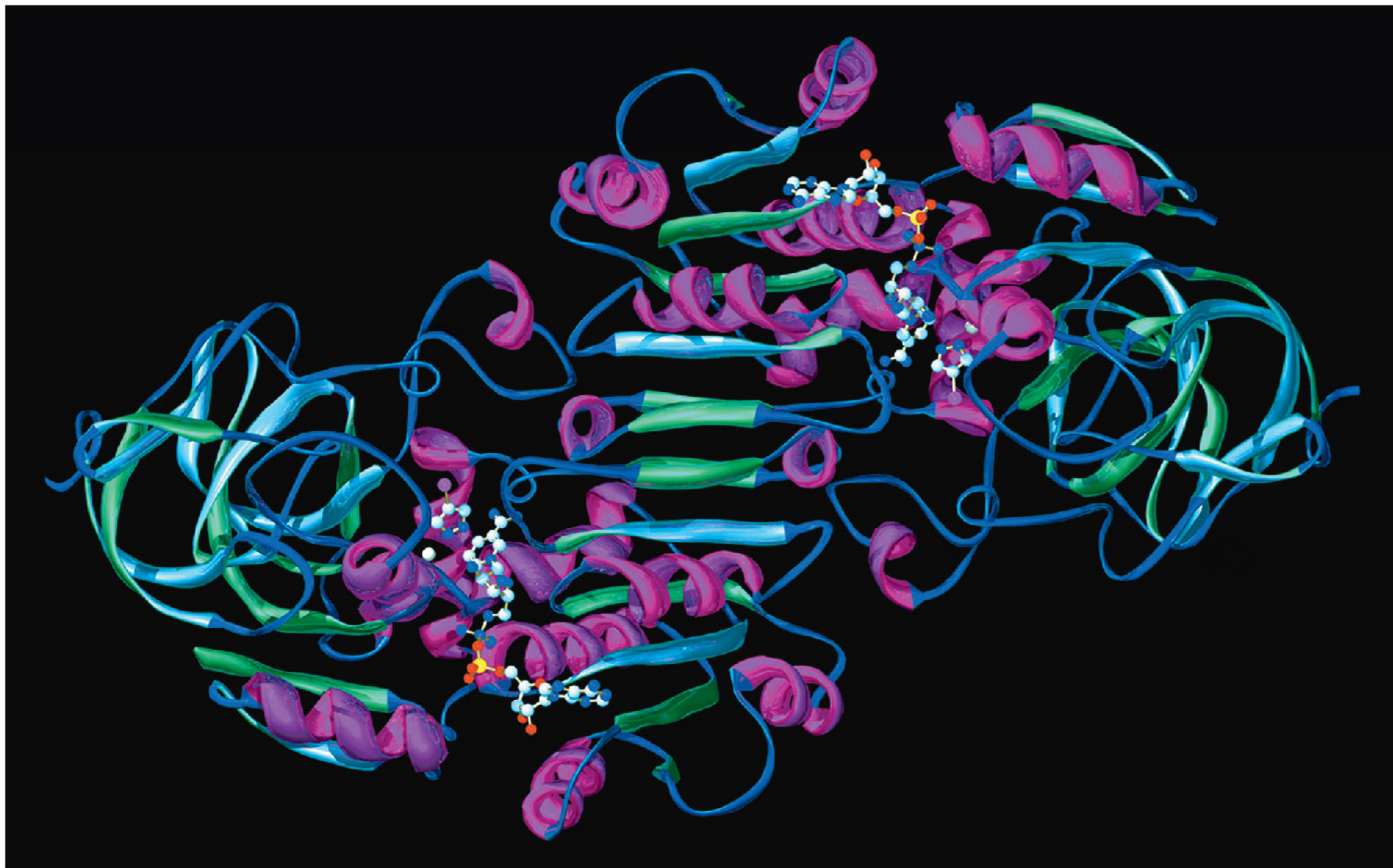
Biological Macromolecules and Lipids

Lecture Presentations by
Nicole Tunbridge and
Kathleen Fitzpatrick

The Molecules of Life

- All living things are made up of four classes of large biological molecules: carbohydrates, lipids, proteins, and nucleic acids
- **Macromolecules** are large molecules and are complex
- Large biological molecules have unique properties that arise from the orderly arrangement of their atoms

Figure 5.1





The scientist in the foreground is using 3-D glasses to help her visualize the structure of the protein displayed on her screen.

Concept 5.1: Macromolecules are polymers, built from monomers

- A **polymer** is a long molecule consisting of many similar building blocks → *not necessarily identical*
- The repeating units that serve as building blocks are called **monomers**
- Carbohydrates, proteins, and nucleic acids are polymers

⊕ lipids are **not** polymers

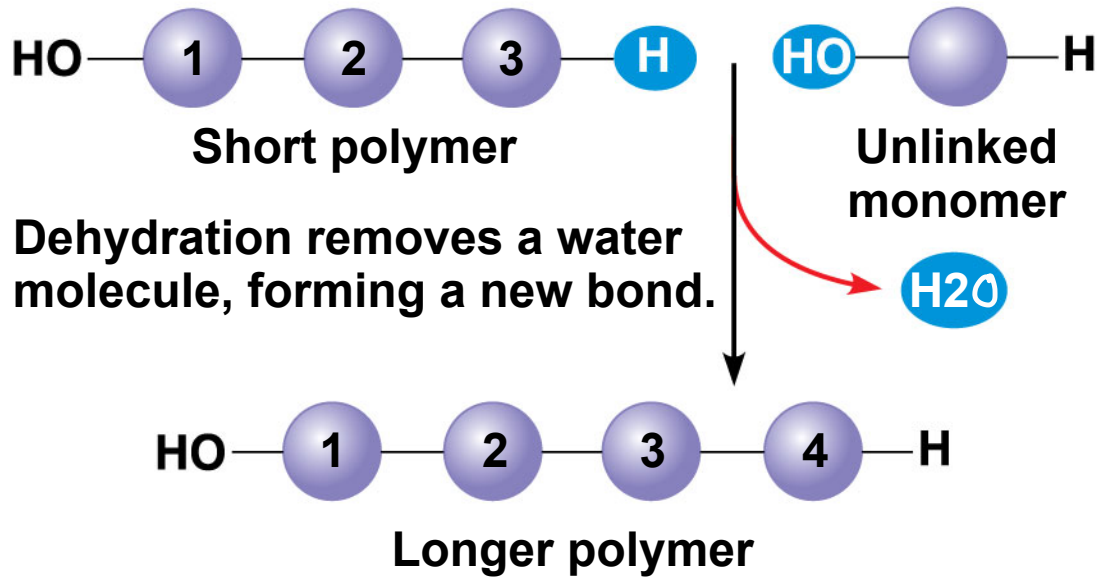
The Synthesis and Breakdown of Polymers

- **Enzymes** are specialized macromolecules that speed up chemical reactions such as those that make or break down polymers
- A **dehydration reaction** occurs when two monomers bond together through the loss of a water molecule
- Polymers are disassembled to monomers by **hydrolysis**, a reaction that is essentially the reverse of the dehydration reaction

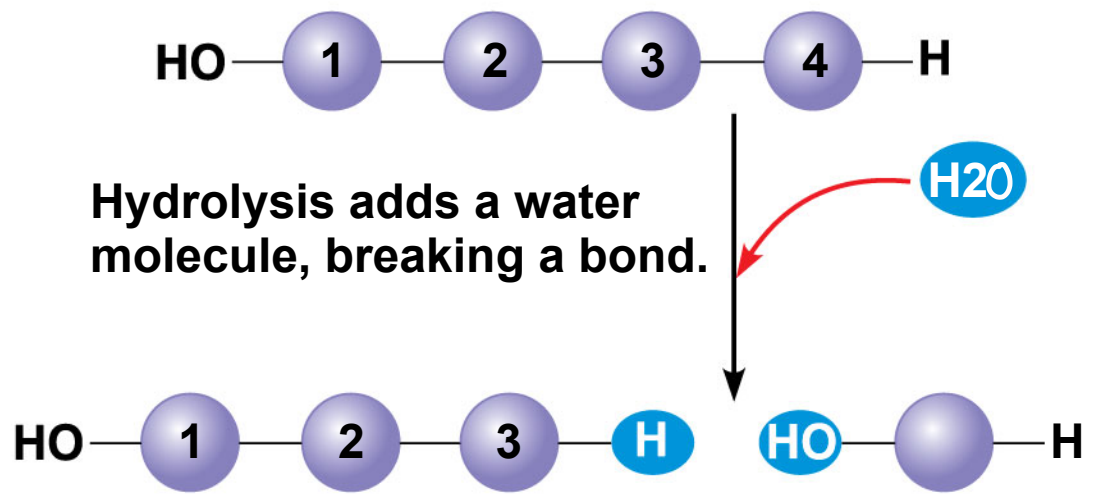
⊗ Dehydration is a type of condensation reactions.

Figure 5.2

(a) Dehydration reaction: synthesizing a polymer



(b) Hydrolysis: breaking down a polymer



The Diversity of Polymers

- A cell has thousands of different macromolecules
- Macromolecules vary among cells of an organism, vary more within a species, and vary even more between species
- A huge variety of polymers can be built from a small set of monomers

n^r = different possibilities

n = no. of different monomers
 r = no. of monomers in the polymer

Concept 5.2: Carbohydrates serve as fuel and building material

- Carbohydrates include sugars and the polymers of sugars
- The simplest carbohydrates are monosaccharides, or simple sugars
- Carbohydrate macromolecules are polysaccharides, polymers composed of many sugar building blocks

Sugars

$(CH_2O)_n$ General Formula

● **Monosaccharides** have molecular formulas that are usually multiples of CH_2O empirical formula

● **Glucose** ($C_6H_{12}O_6$) is the most common monosaccharide

● **Monosaccharides** are classified by

● The location of the carbonyl group (as aldose or ketose)

aldehyde sugar

ketone sugar

● The number of carbons in the carbon skeleton

Ranges from 3 to 7 atoms "C".

Figure 5.3

Aldoses (Aldehyde Sugars)	Ketoses (Ketone Sugars)
Trioses: three-carbon sugars (C₃H₆O₃)	
$ \begin{array}{c} \text{H} \\ \diagdown \\ \text{C}=\text{O} \\ \\ \text{H}-\text{C}-\text{OH} \\ \\ \text{H}-\text{C}-\text{OH} \\ \\ \text{H} \end{array} $ <p style="text-align: center;">Glyceraldehyde</p>	$ \begin{array}{c} \text{H} \\ \\ \text{H}-\text{C}-\text{OH} \\ \\ \text{C}=\text{O} \\ \\ \text{H}-\text{C}-\text{OH} \\ \\ \text{H} \end{array} $ <p style="text-align: center;">Dihydroxyacetone</p>
Pentoses: five-carbon sugars (C₅H₁₀O₅)	
$ \begin{array}{c} \text{H} \\ \diagdown \\ \text{C}=\text{O} \\ \\ \text{H}-\text{C}-\text{OH} \\ \\ \text{H}-\text{C}-\text{OH} \\ \\ \text{H}-\text{C}-\text{OH} \\ \\ \text{H}-\text{C}-\text{OH} \\ \\ \text{H} \end{array} $ <p style="text-align: center;">Ribose</p>	$ \begin{array}{c} \text{H} \\ \\ \text{H}-\text{C}-\text{OH} \\ \\ \text{C}=\text{O} \\ \\ \text{H}-\text{C}-\text{OH} \\ \\ \text{H}-\text{C}-\text{OH} \\ \\ \text{H}-\text{C}-\text{OH} \\ \\ \text{H} \end{array} $ <p style="text-align: center;">Ribulose</p>
Hexoses: six-carbon sugars (C₆H₁₂O₆)	
$ \begin{array}{cc} \begin{array}{c} \text{H} \\ \diagdown \\ \text{C}=\text{O} \\ \\ \text{H}-\text{C}-\text{OH} \\ \\ \text{HO}-\text{C}-\text{H} \\ \\ \text{H}-\text{C}-\text{OH} \\ \\ \text{H}-\text{C}-\text{OH} \\ \\ \text{H} \end{array} & \begin{array}{c} \text{H} \\ \diagdown \\ \text{C}=\text{O} \\ \\ \text{H}-\text{C}-\text{OH} \\ \\ \text{HO}-\text{C}-\text{H} \\ \\ \text{HO}-\text{C}-\text{H} \\ \\ \text{H}-\text{C}-\text{OH} \\ \\ \text{H}-\text{C}-\text{OH} \\ \\ \text{H} \end{array} \end{array} $ <p style="text-align: center;">Glucose Galactose</p>	$ \begin{array}{c} \text{H} \\ \\ \text{H}-\text{C}-\text{OH} \\ \\ \text{C}=\text{O} \\ \\ \text{HO}-\text{C}-\text{H} \\ \\ \text{H}-\text{C}-\text{OH} \\ \\ \text{H}-\text{C}-\text{OH} \\ \\ \text{H}-\text{C}-\text{OH} \\ \\ \text{H} \end{array} $ <p style="text-align: center;">Fructose</p>

Figure 5.3a

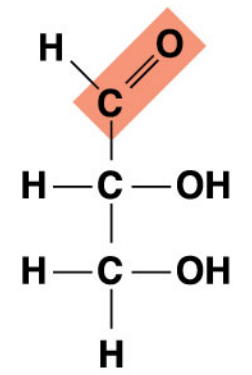
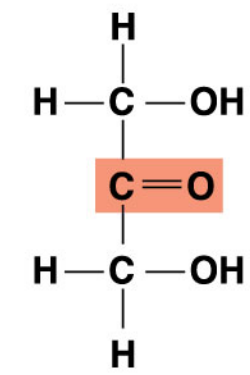
Aldose (Aldehyde Sugar)	Ketose (Ketone Sugar)
Trioses: three-carbon sugars (C₃H₆O₃)	
 <p>The structure shows a vertical chain of three carbon atoms. The top carbon is part of an aldehyde group, with a hydrogen atom (H) to its left and a double-bonded oxygen atom (O) to its right. This top carbon and its double bond are highlighted with a red diamond. The middle carbon is bonded to a hydrogen atom (H) on the left and a hydroxyl group (OH) on the right. The bottom carbon is bonded to a hydrogen atom (H) on the left and a hydroxyl group (OH) on the right.</p>	 <p>The structure shows a vertical chain of three carbon atoms. The top carbon is bonded to a hydrogen atom (H) above it, a hydrogen atom (H) to its left, and a hydroxyl group (OH) to its right. The middle carbon is part of a ketone group, with a double-bonded oxygen atom (O) to its right. This middle carbon and its double bond are highlighted with a red rectangle. The bottom carbon is bonded to a hydrogen atom (H) to its left and a hydroxyl group (OH) to its right.</p>
Glyceraldehyde	Dihydroxyacetone

Figure 5.3b

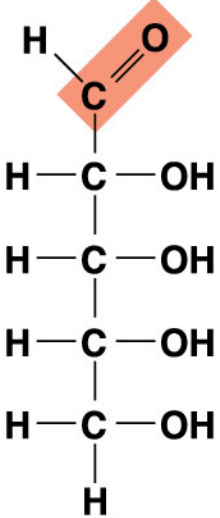
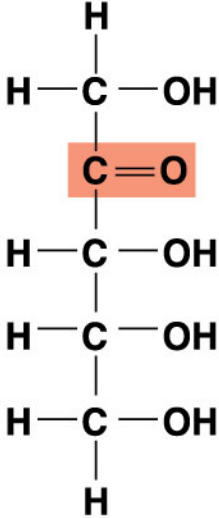
Aldose (Aldehyde Sugar)	Ketose (Ketone Sugar)
Pentoses: five-carbon sugars (C₅H₁₀O₅)	
 <p style="text-align: center;">Ribose</p>	 <p style="text-align: center;">Ribulose</p>

Figure 5.3c

	Aldose (Aldehyde Sugar)	Ketose (Ketone Sugar)
Hexoses: six-carbon sugars (C₆H₁₂O₆)		
1		
2	H — C — OH	H — C — OH
3	HO — C — H	HO — C — H
4		
	H — C — OH	H — C — OH
	H — C — OH	H — C — OH
	H	H
	Glucos	Galactos
	e	e
		H — C — OH
		HO — C — H
		H — C — OH
		H — C — OH
		H — C — OH
		H
		Fructos
		e

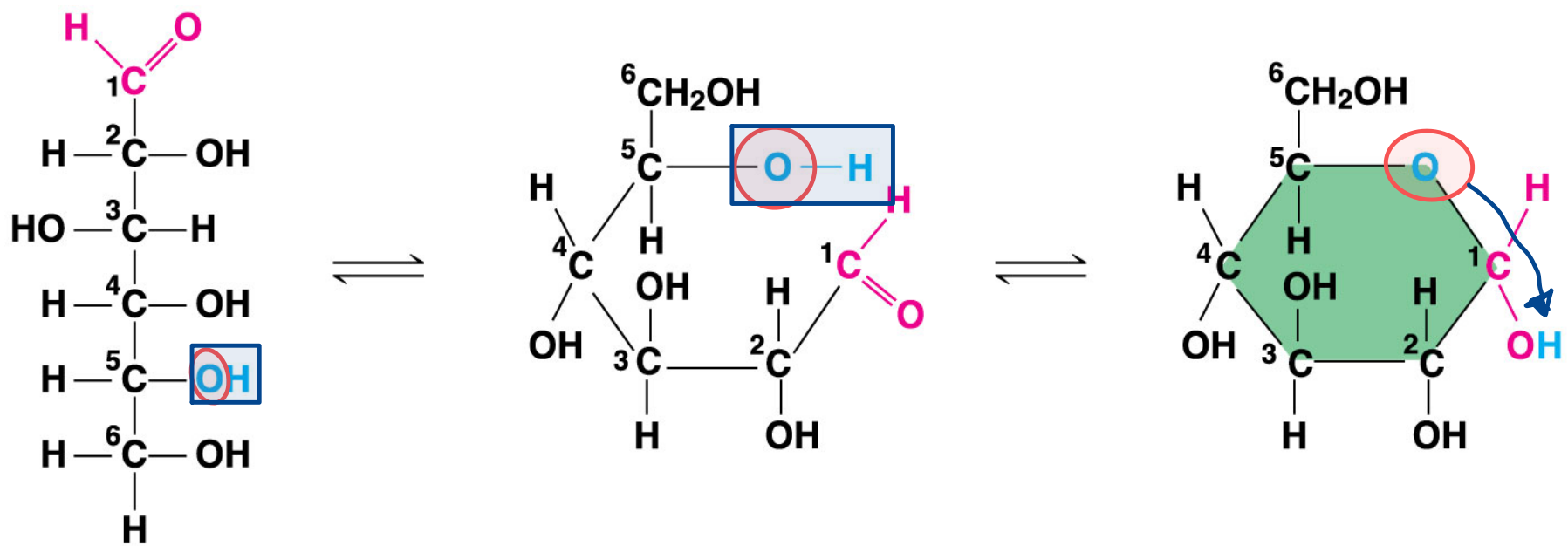
1
2
3
4

Spatial arrangement around C4

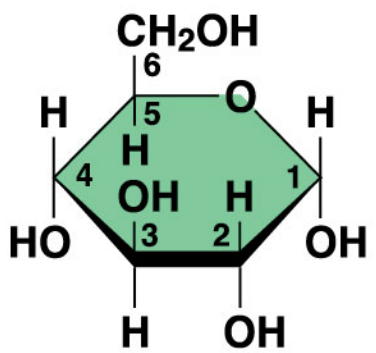
- Though often drawn as linear skeletons, in aqueous solutions many sugars form rings ~~⊗~~
- Monosaccharides serve as a major fuel for cells and as raw material for building molecules ^①
_②

~~⊗~~ Rings are more stable than linear structures
in aqueous solutions.

Figure 5.4



(a) Linear and ring forms




(b) Abbreviated ring structure

*α-glucose
(later)*

Empty angles represent carbon atoms.

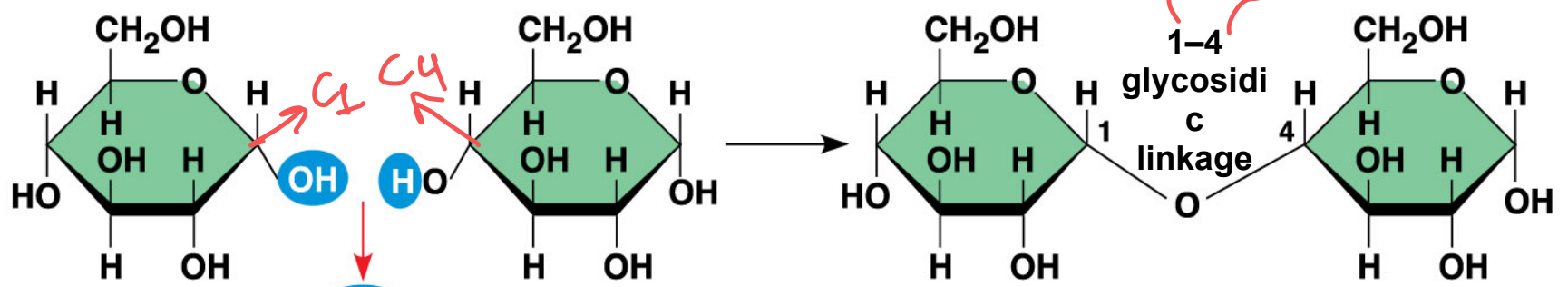
- A **disaccharide** is formed when a **dehydration reaction** joins two monosaccharides
- This **covalent bond** is called a **glycosidic linkage**



Hydroxyl - Hydroxyl
linkage

Figure 5.5

(a) Dehydration reaction in the **synthesis of maltose**

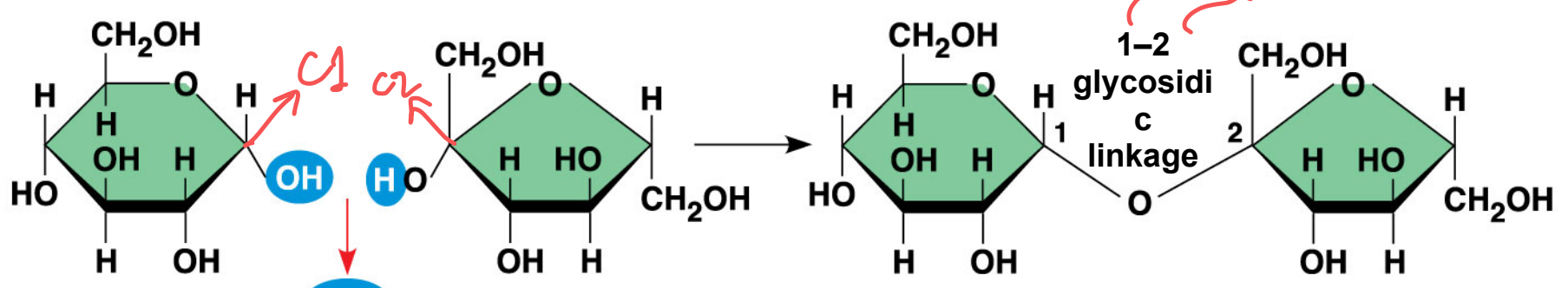


Glucose

Glucose

Maltose

(b) Dehydration reaction in the **synthesis of sucrose**



Glucose

Fructose

Sucrose

⊗ sucrose: used in plants when transporting carbohydrates to non-photosynthetic organs — roots ...

Polysaccharides

- **Polysaccharides**, the **polymers of sugars**, have **storage** and **structural** roles
- The **architecture and function** of a **polysaccharide** are **determined by its sugar monomers** and the **positions of its glycosidic linkages**

⊕ additional Disaccharide example:

Lactose (milk sugar): glucose + galactose

Storage Polysaccharides

- **Starch**, a **storage** polysaccharide **of plants**, consists of **glucose monomers**
- Plants store surplus starch as **granules** within **chloroplasts and other plastids**
- **The simplest form of starch is amylose**
(unbranched)

excess

Small particles

Figure 5.6

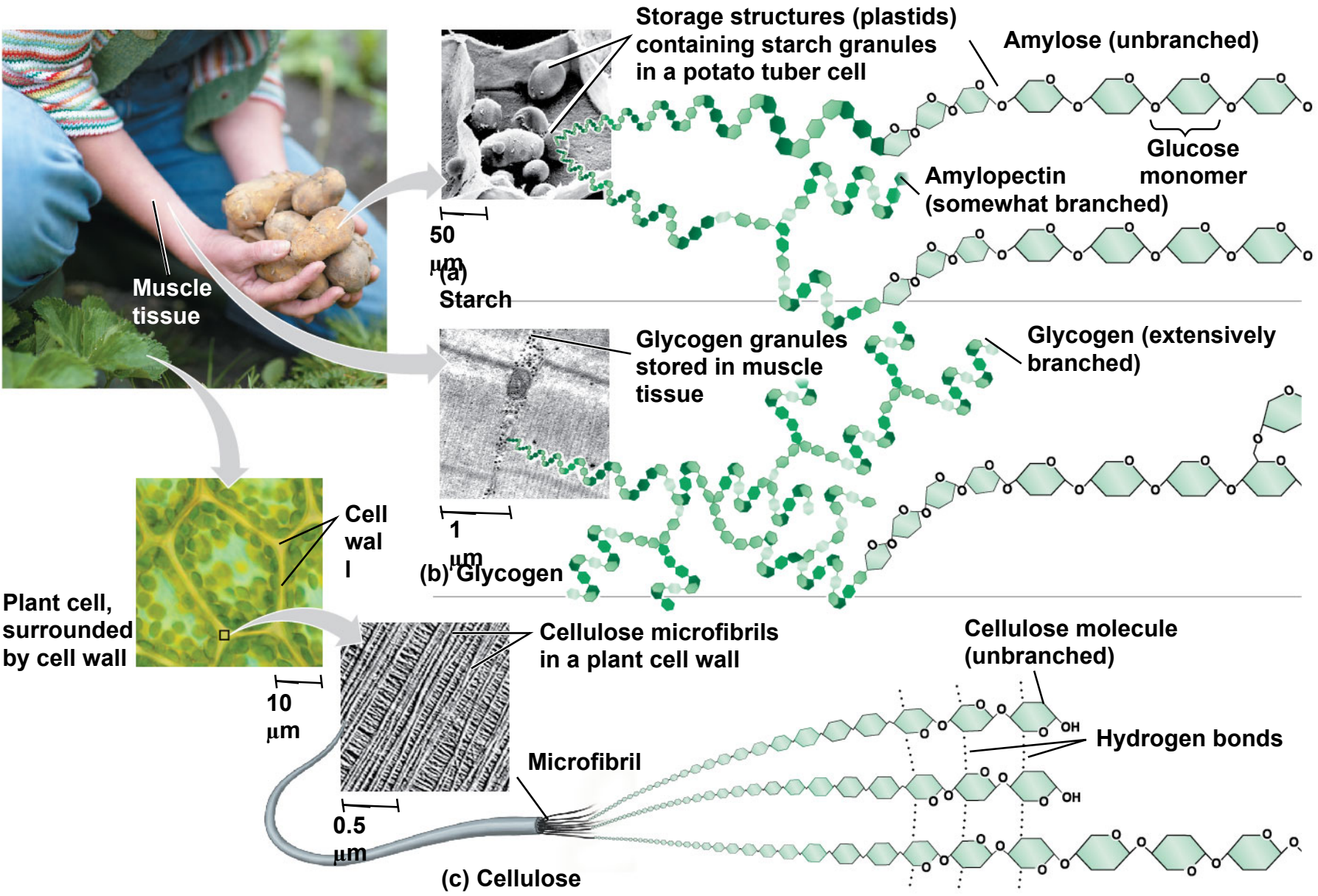
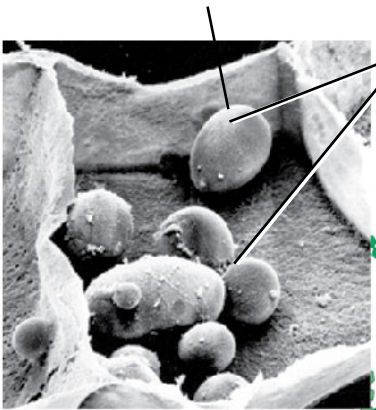
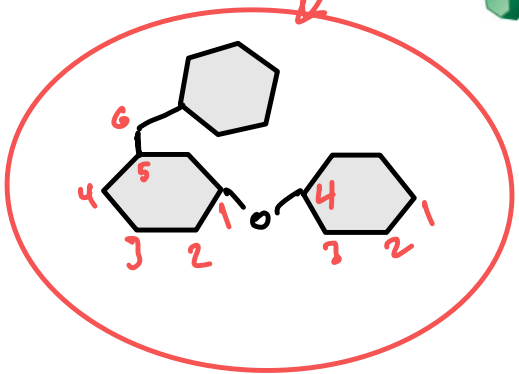
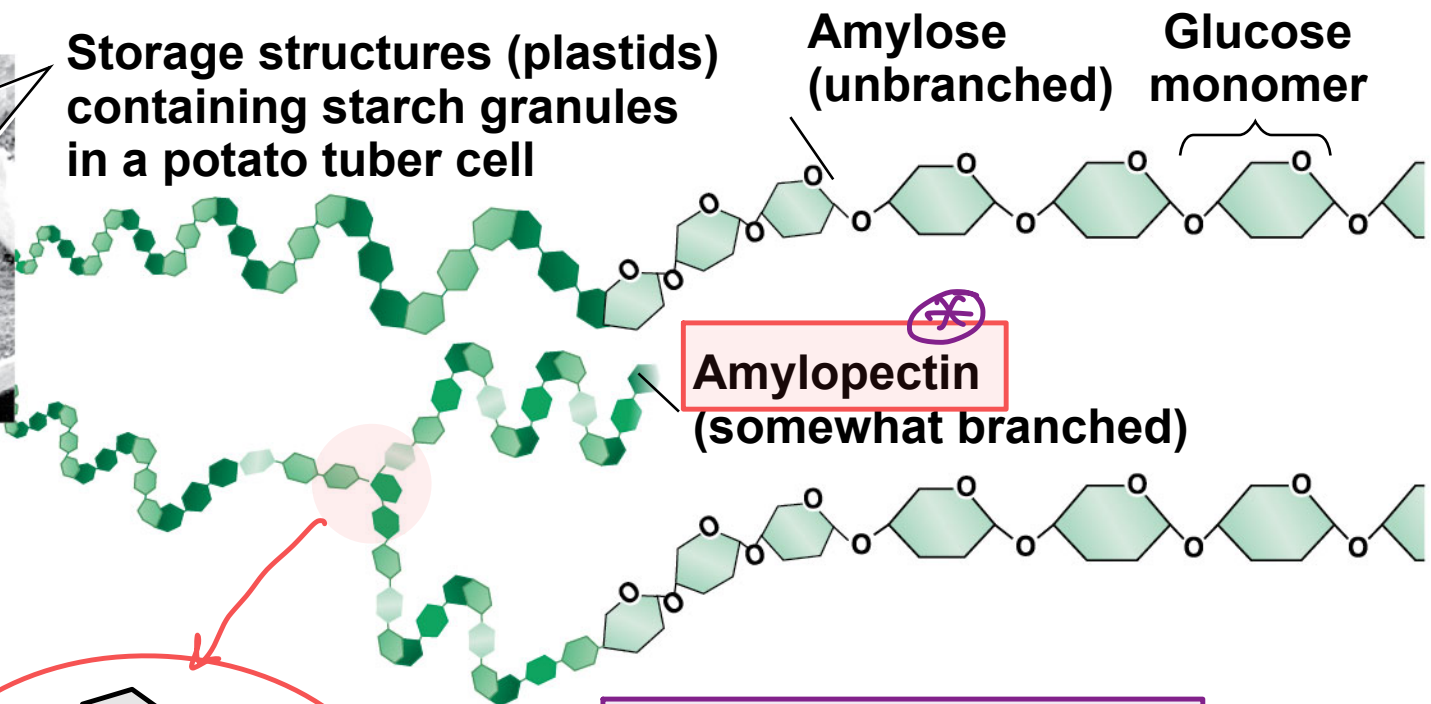


Figure 5.6a



Storage structures (plastids) containing starch granules in a potato tuber cell

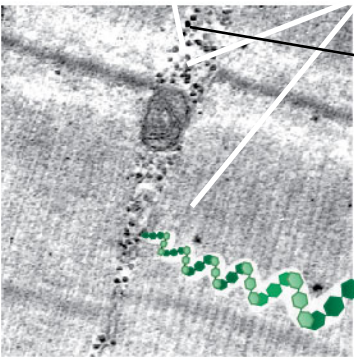
50 μm
(a)
Starch



1-6 linkage

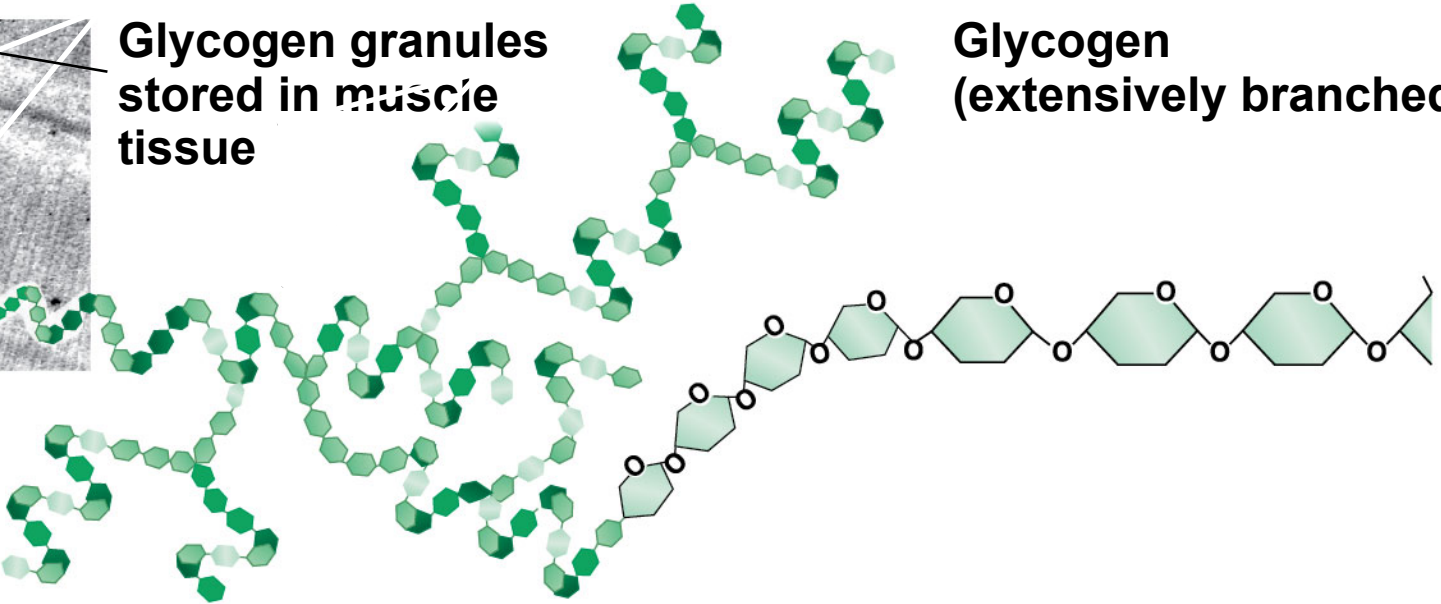
in amylopectin branches.

Figure 5.6b



**Glycogen granules
stored in muscle
tissue**

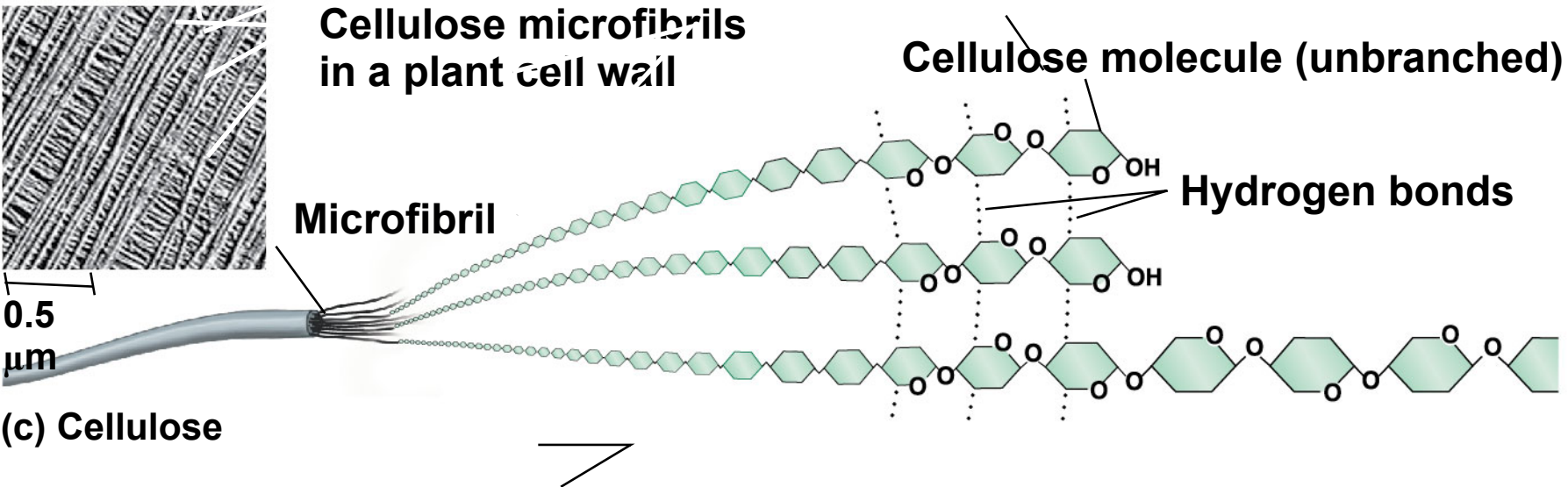
**Glycogen
(extensively branched)**



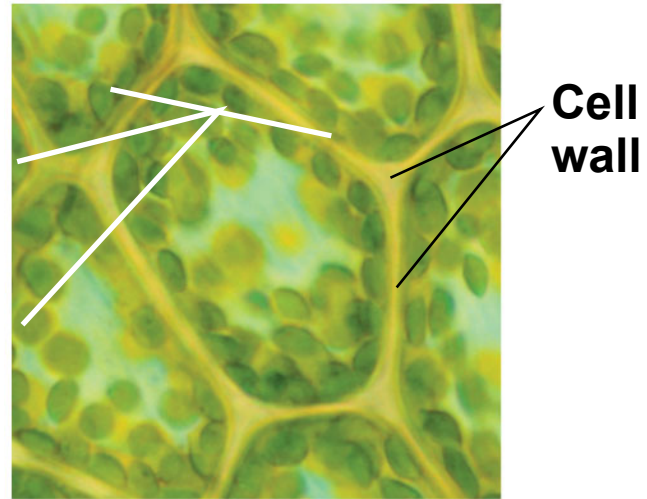
1
μm

(b) Glycogen

Figure 5.6c



(c) Cellulose



**Plant cell,
surrounded
by cell wall**

**10
µm**

Not long sustaining (unlike starch)

- **Glycogen** is a **storage** polysaccharide **in animals**
- Glycogen is stored mainly in **liver** and **muscle** cells
- **Hydrolysis** of glycogen in these cells **releases glucose** when the demand for sugar increases

Structural Polysaccharides

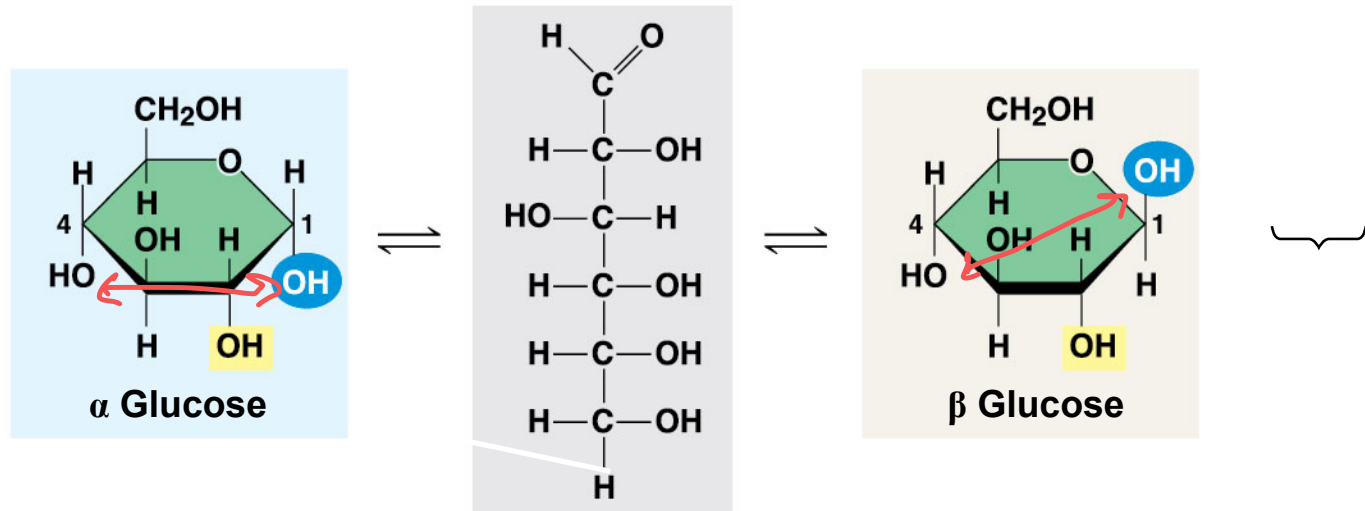
Annually $\rightarrow 10^{14}$ kg

the most abundant organic compound on Earth.

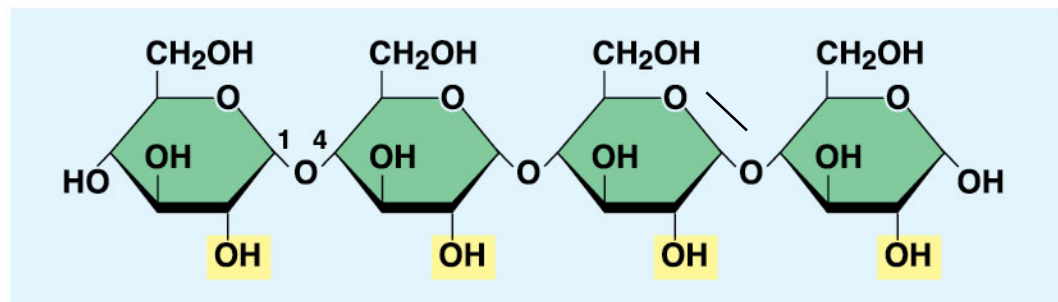
- The polysaccharide **cellulose** is a major component of the tough wall of plant cells
- Like starch, cellulose is a polymer of glucose, but the glycosidic linkages differ } \rightarrow important
- The difference is based on two ring forms for glucose: alpha (α) and beta (β)

⊗ The linear structure of glucose does not differ

Figure 5.7

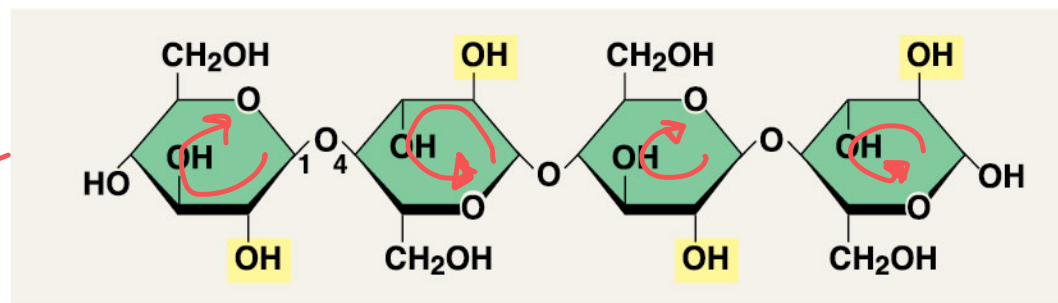


(a) α and β glucose ring structures



(b) Starch: 1-4 linkage of α glucose monomers

Corresponding



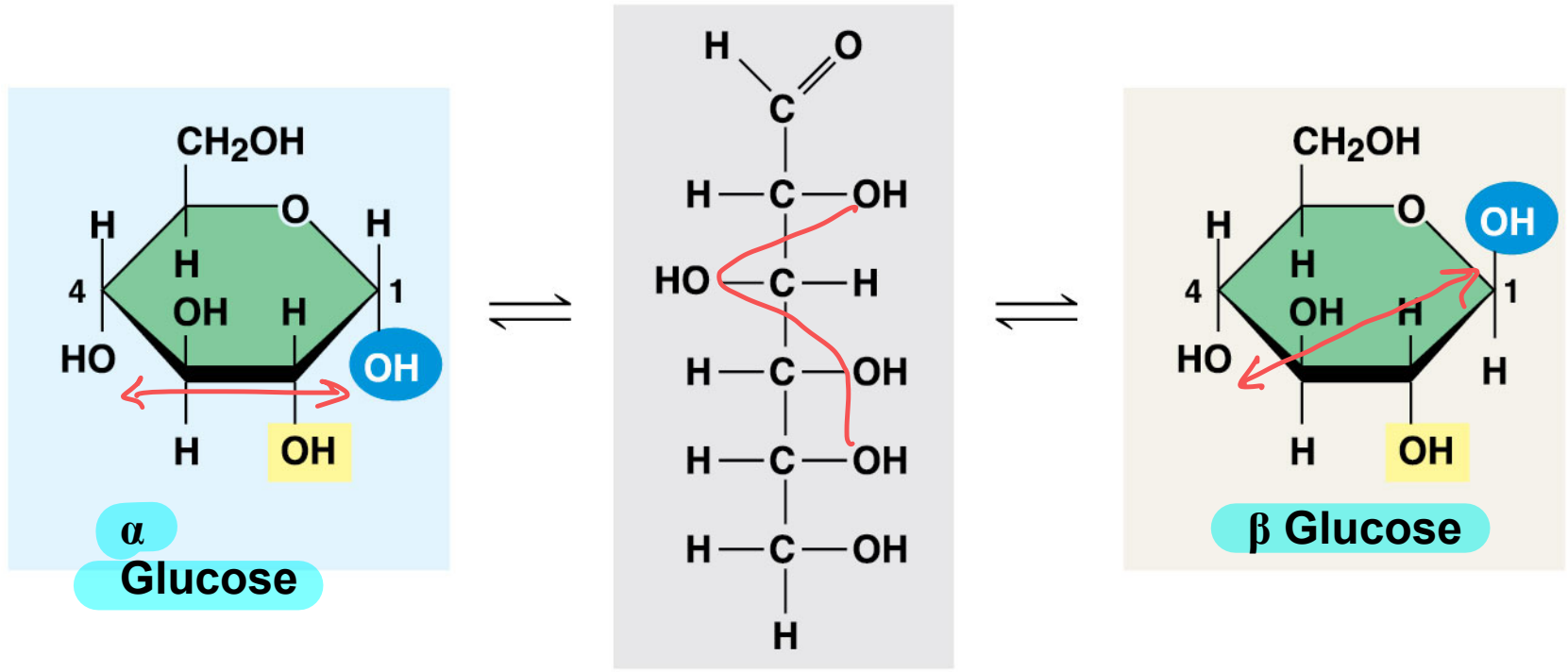
(c) Cellulose: 1-4 linkage of β glucose monomers

Alternating



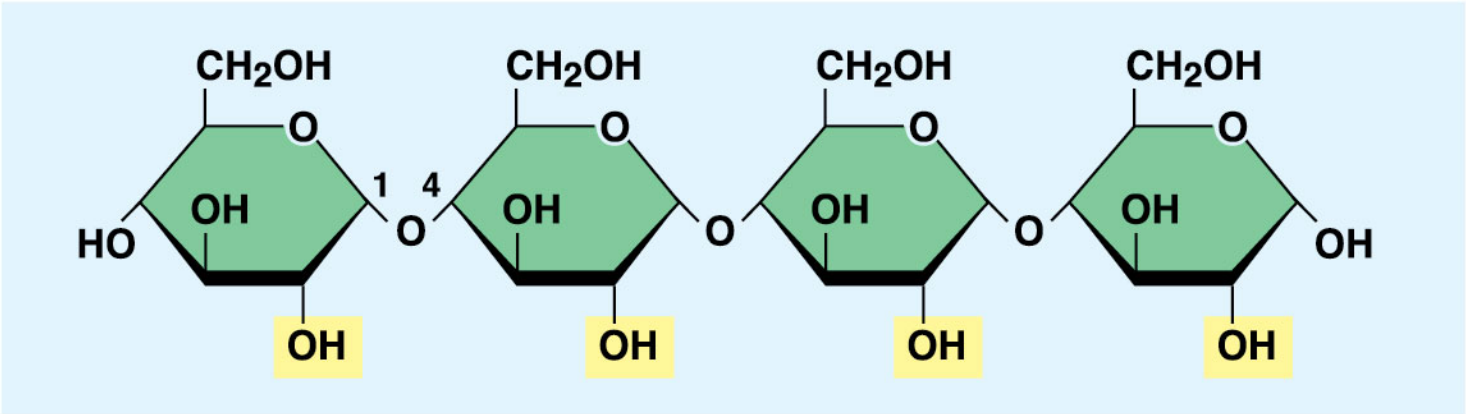
arrows are for numbering of Carbon atoms

⊗ Both α & β share the same linear configuration

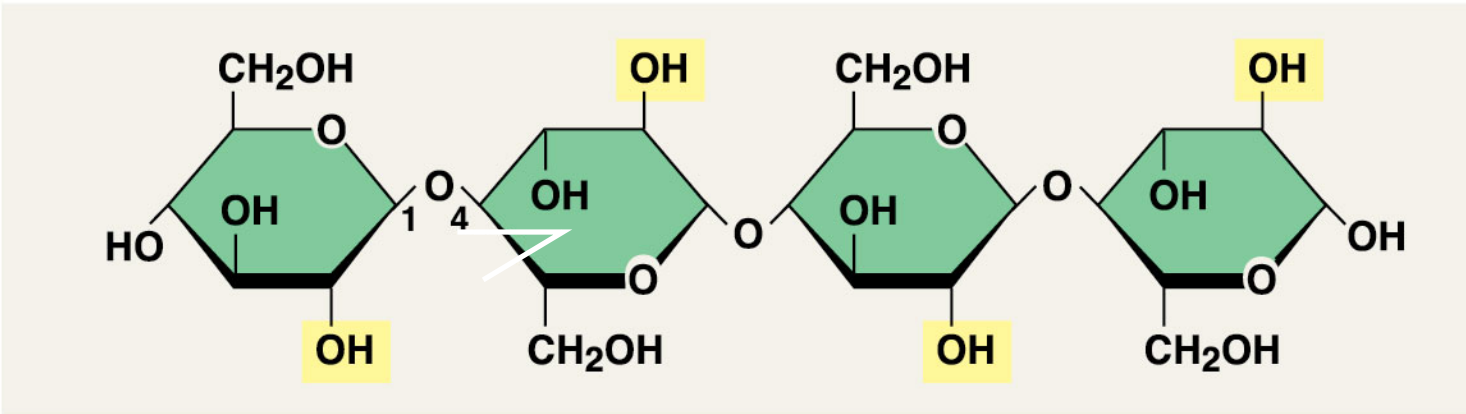


(a) α and β glucose ring structures

Figure 5.7b



(b) Starch: 1–4 linkage of α glucose monomers



(c) Cellulose: 1–4 linkage of β glucose monomers

Starch (α configuration) is largely helical

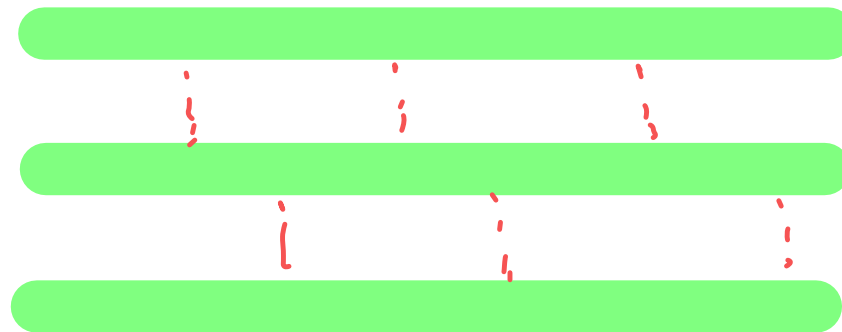
Cellulose molecules (β configuration) are straight and unbranched

Some hydroxyl groups on the monomers of cellulose can hydrogen-bond with hydroxyls of parallel cellulose molecules

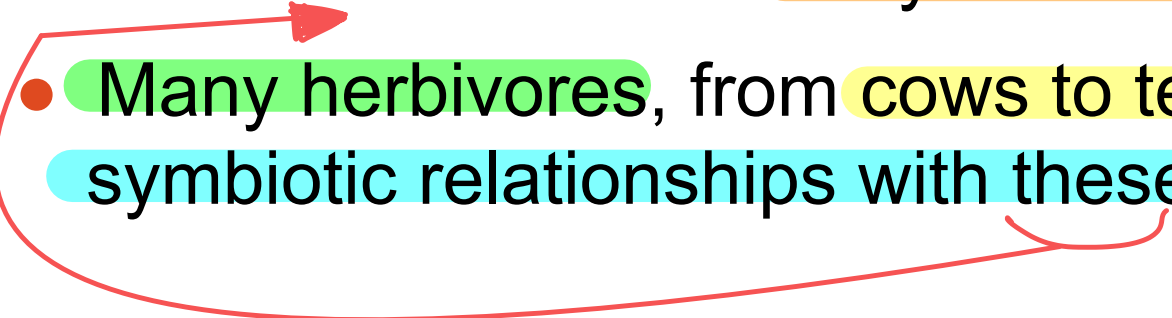
(:·)



H-Bonding



Micro
fibrils

- Enzymes that digest starch by hydrolyzing α linkages can't hydrolyze β linkages in cellulose
 - The cellulose in human food passes through the digestive tract as "insoluble fiber"
 - Some microbes use enzymes to digest cellulose
 - Many herbivores, from cows to termites, have symbiotic relationships with these microbes
- 

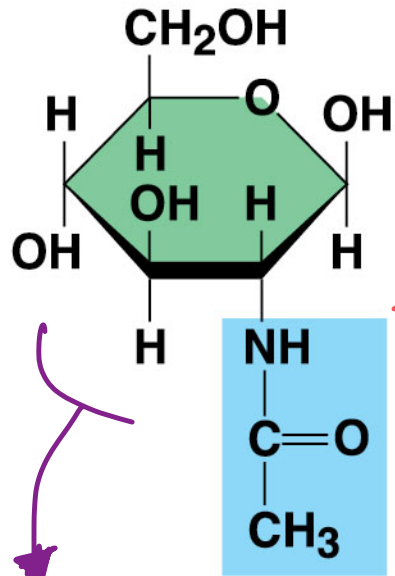
(Kaitin)

like cellulose

- **Chitin**, another **structural** polysaccharide, is found in the **exoskeleton of arthropods**
- Chitin also **provides structural support for the cell walls of many fungi**

Figure 5.8

⊕ Chitin is decomposed using a different enzyme than cellulase ⇒ **Chitinase**





The structure of the chitin monomer


Nitrogen-containing group.

Similar to glucose (β)
Because Chitin is like cellulose

◀ Chitin, embedded in proteins, forms the exoskeleton of arthropods.

Concept 5.3: Lipids are a diverse group of hydrophobic molecules

- **Lipids** are the one class of large biological molecules that does not include true polymers 
- The unifying feature of lipids is that they mix poorly, if at all, with water
- Lipids consist mostly of hydrocarbon regions
- The most biologically important lipids are fats, phospholipids, and steroids 

 That is why lipids are excluded from the title of the chapter unlike other macromolecules

- Carbohydrates
- Proteins
- Nucleic acids.

Fats

- **Fats** are constructed from two types of smaller molecules: glycerol and fatty acids.
- Glycerol is a three-carbon alcohol with a hydroxyl group attached to each carbon
- A **fatty acid** consists of a carboxyl group attached to a long carbon skeleton

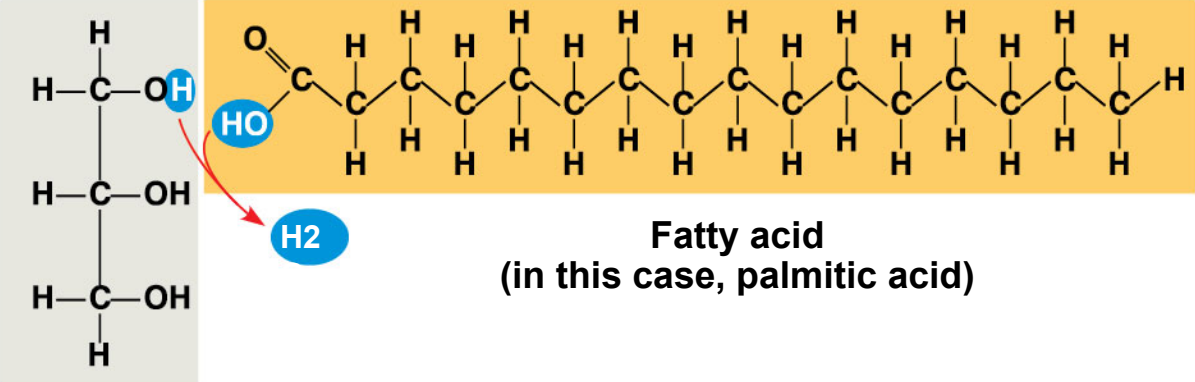
usually

16 or 18

carbon atoms.

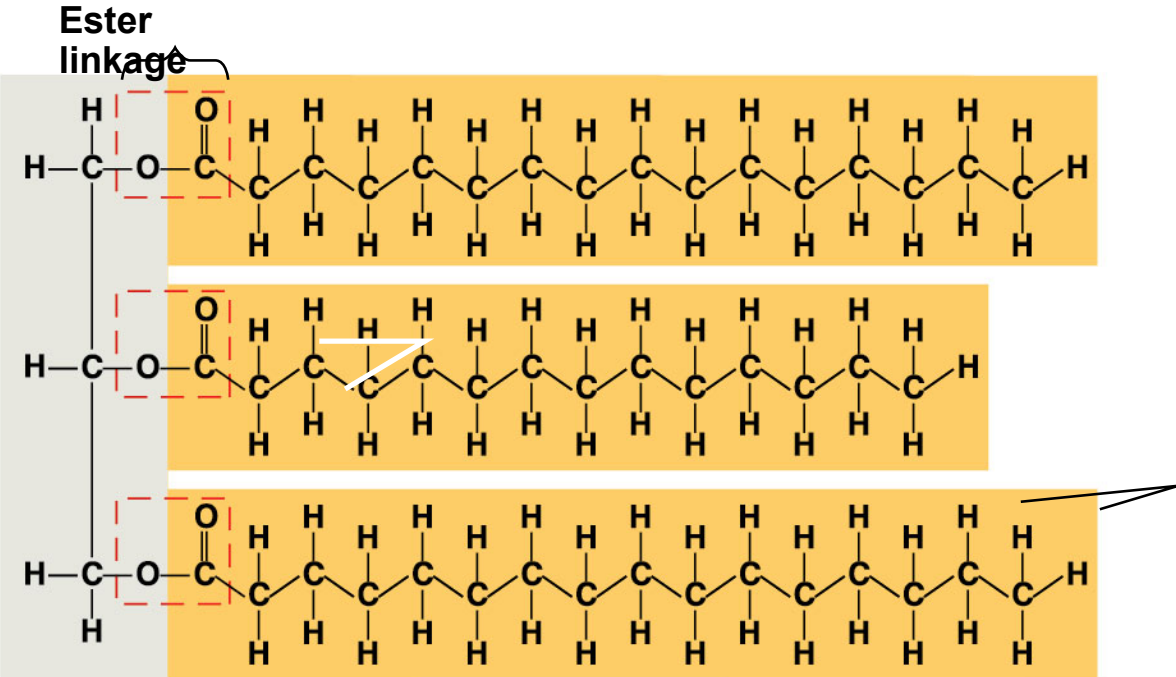


Figure 5.9



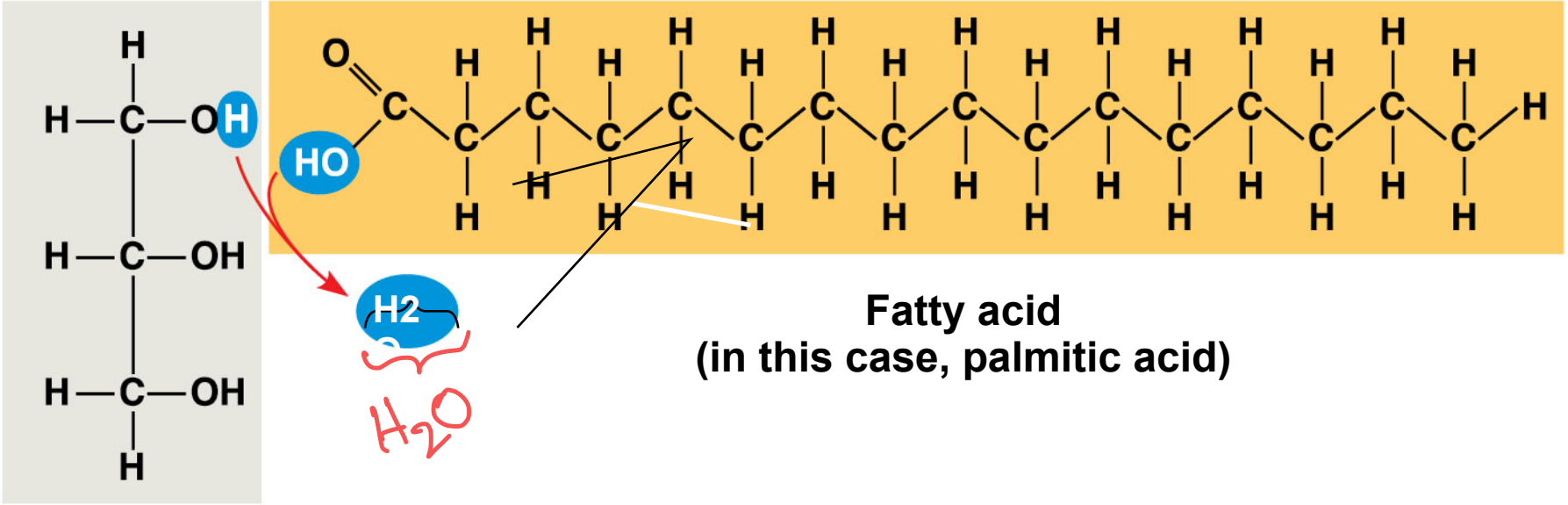
Glycero

(a) One of three dehydration reactions in the synthesis of a fat



(b) Fat molecule (triacylglycerol)

Figure 5.9a



Fatty acid
(in this case, palmitic acid)

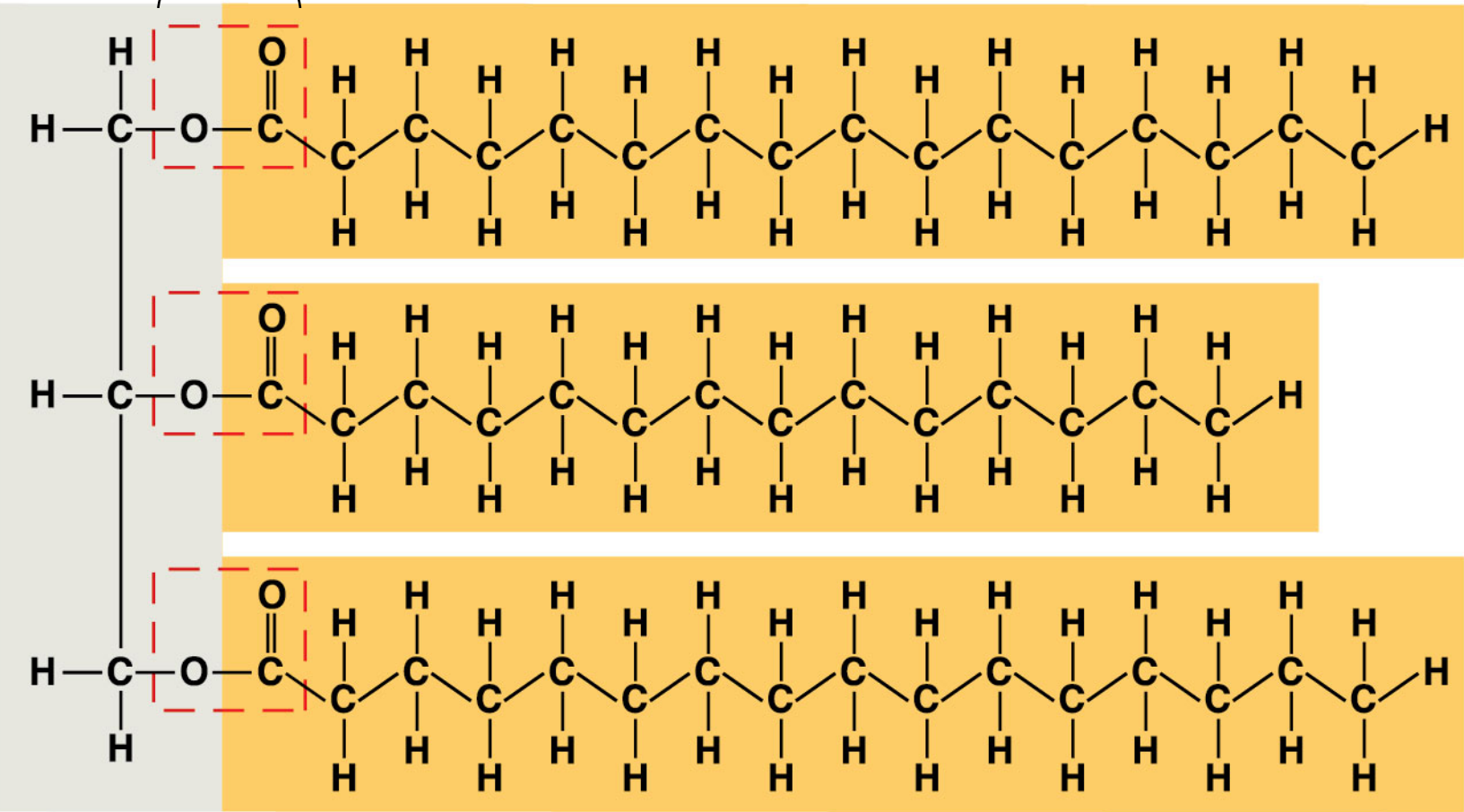
Glycero

(a) One of three dehydration reactions in the synthesis of a fat

Figure 5.9b

Hydroxyl — carboxyl linkage

Ester linkage



(b) Fat molecule (triacylglycerol)



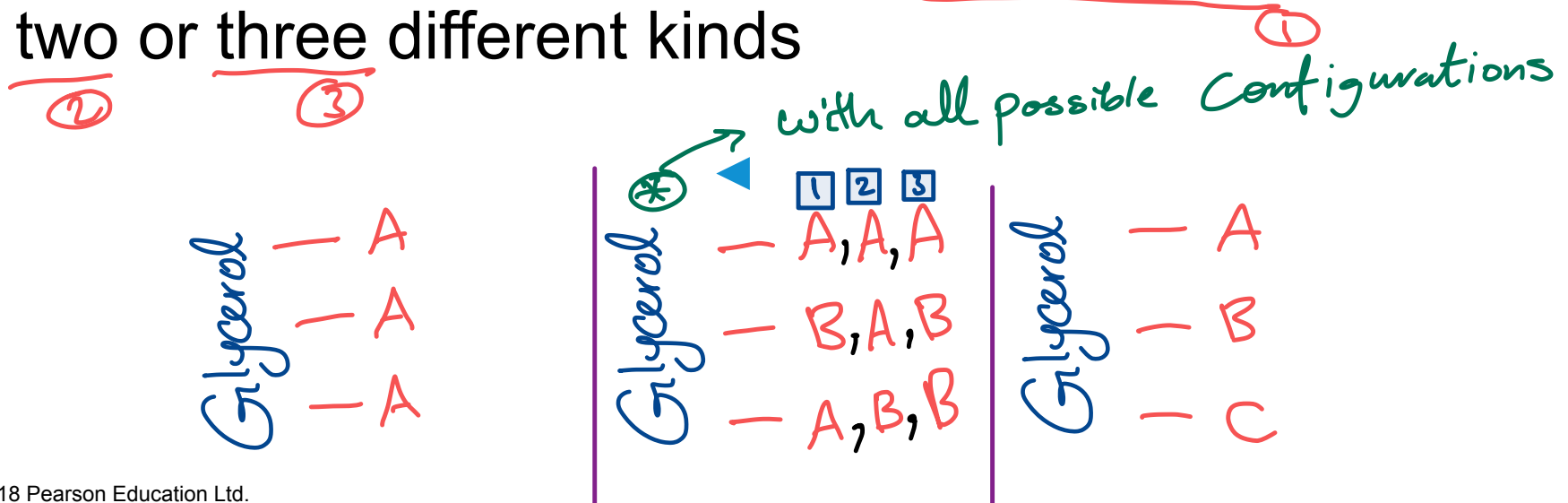
C-H : non-polar covalent bond.



Fats separate from water because water molecules hydrogen-bond to each other and exclude the fats

In a fat, three fatty acids are joined to glycerol by an ester linkage, creating a triacylglycerol, or triglyceride

The fatty acids in a fat can be all the same or of two or three different kinds



- Fatty acids vary in length (number of carbons) and in the number and locations of double bonds
- Saturated fatty acids have the maximum number of hydrogen atoms possible and no double bonds
- Unsaturated fatty acids have one or more double bonds

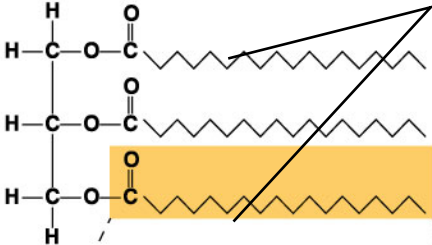


Figure 5.10

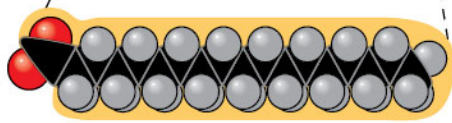
(a) Saturated fat



Structural formula of a saturated fat molecule



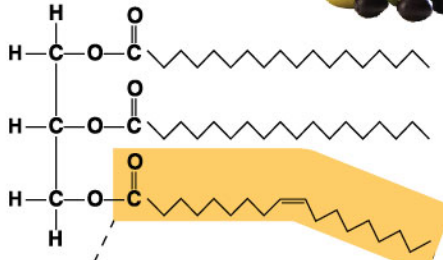
Space-filling model of stearic acid, a saturated fatty acid



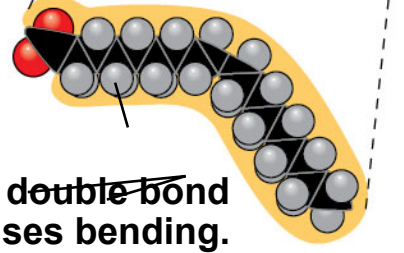
(b) Unsaturated fat



Structural formula of an unsaturated fat molecule



Space-filling model of oleic acid, an unsaturated fatty acid

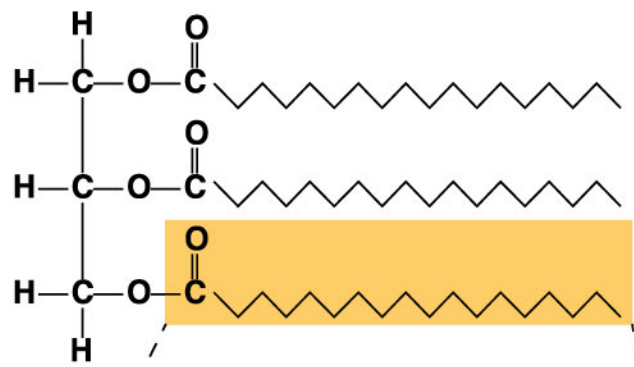


Cis double bond causes bending.

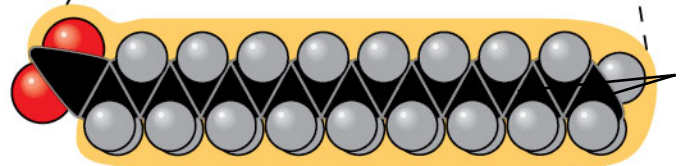
(a) Saturated fat



**Structural formula
of a saturated fat
molecule**



**Space-filling model of
stearic acid, a
saturated fatty acid**

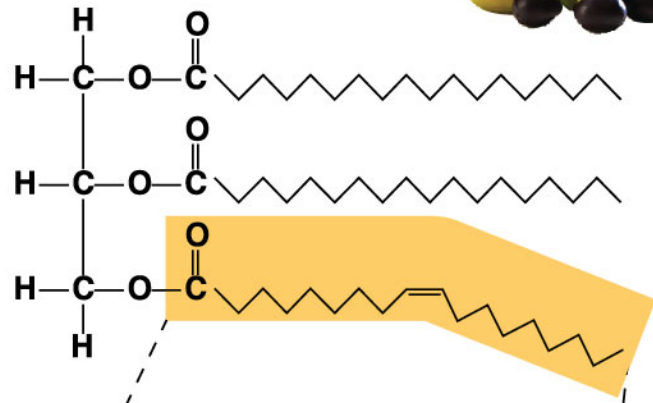


⊗ High Hydrophobic interactions
between saturated fats.

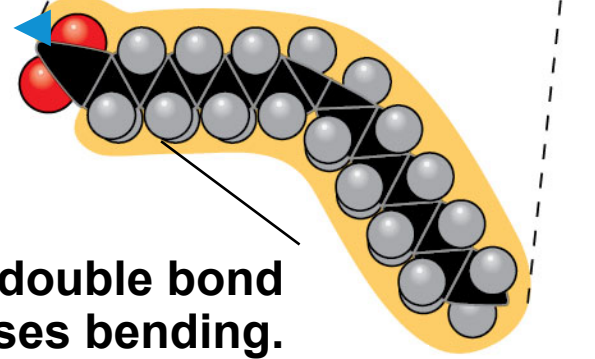
(b) Unsaturated fat



Structural formula of an unsaturated fat molecule



Space-filling model of oleic acid, an unsaturated fatty acid



Cis double bond causes bending.

() Low Hydrophobic interactions Between unsaturated fats.*

- **Fats** made from saturated fatty acids are called saturated fats and are solid at room temperature
- Most animal fats are saturated
- **Fats** made from unsaturated fatty acids are called unsaturated fats or oils and are liquid at room temperature
- **Plant fats** and **fish fats** are usually unsaturated

⊕ levels of Triacylglycerol are reported when blood is tested for lipids.

- A diet rich in saturated fats may contribute to cardiovascular disease through plaque deposits
- Hydrogenation is the process of converting unsaturated fats to saturated fats by adding hydrogen
- Hydrogenating vegetable oils also creates unsaturated fats with *trans* double bonds
- These **trans fats** may contribute more than saturated fats to cardiovascular disease

trans fats are worse than Saturated fats

- The major function of fats is energy storage
- Humans and other mammals store their long-term food reserves in adipose cells
- Adipose tissue also cushions vital organs and insulates the body

Adipose \equiv Fatty

A gram of fat stores more than double the amount of energy of a gram of carbohydrates.

phospholipids

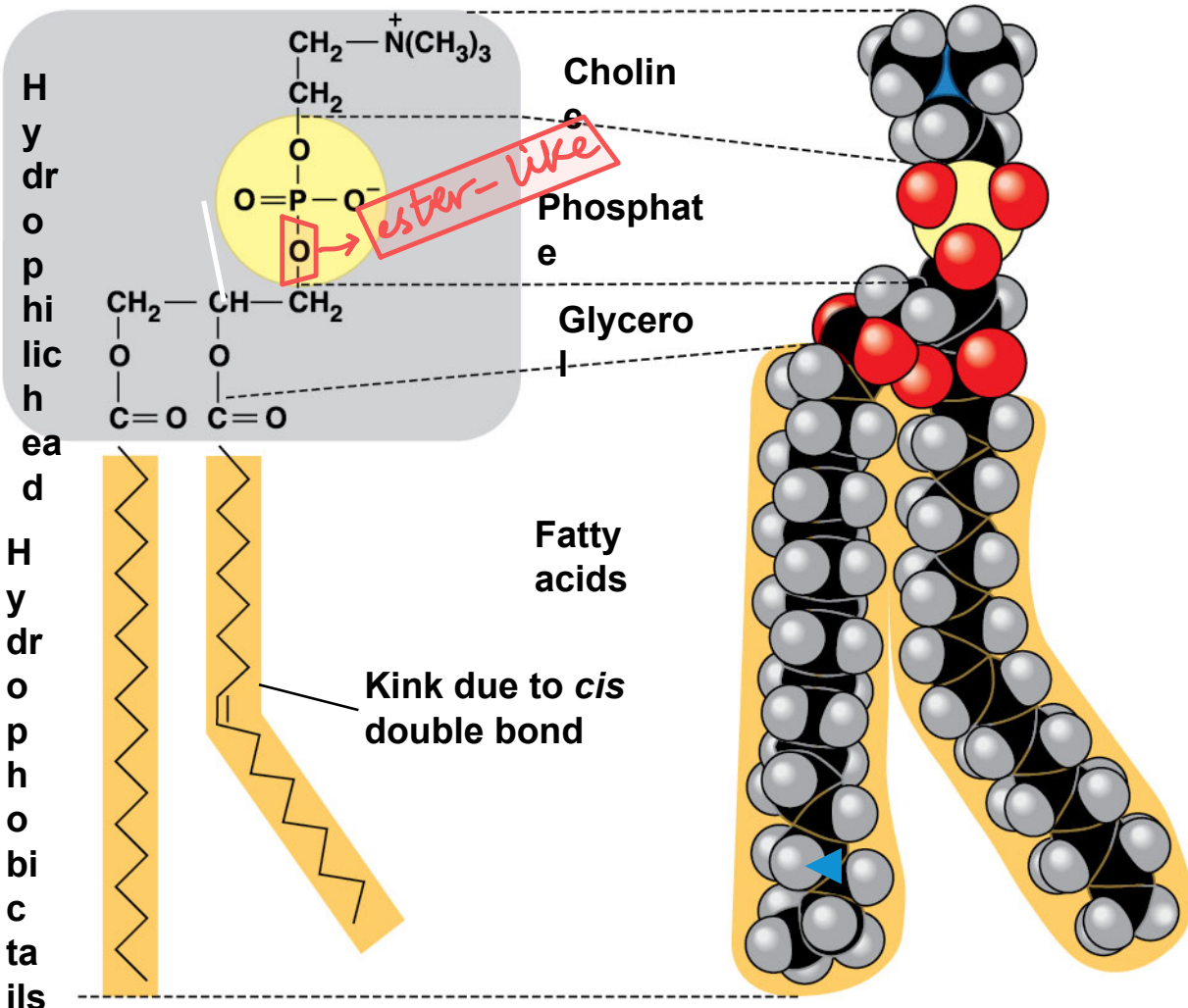
- In a **phospholipid**, two fatty acids and a phosphate group are attached to glycerol
- The two fatty acid tails are **hydrophobic**, but the phosphate group and its attachments form a **hydrophilic head**

Choline or other groups

⊗ interaction between glycerol and a phosphate group is called

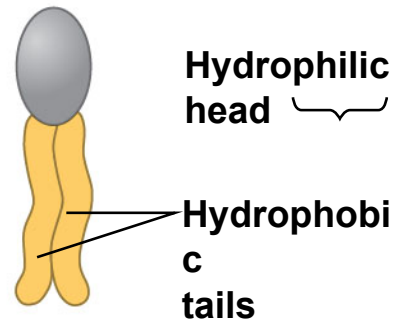
α phosphoester linkage

Figure 5.11

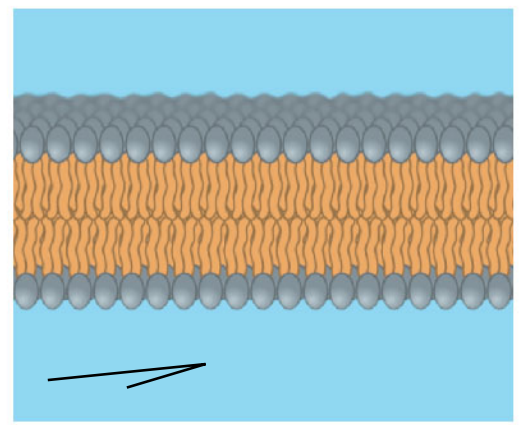


(a) Structural formula

(b) Space-filling model



(c) Phospholipid symbol



(d) Phospholipid bilayer

⊗ Next 3 slides are fake: images
 send to back to see highlights

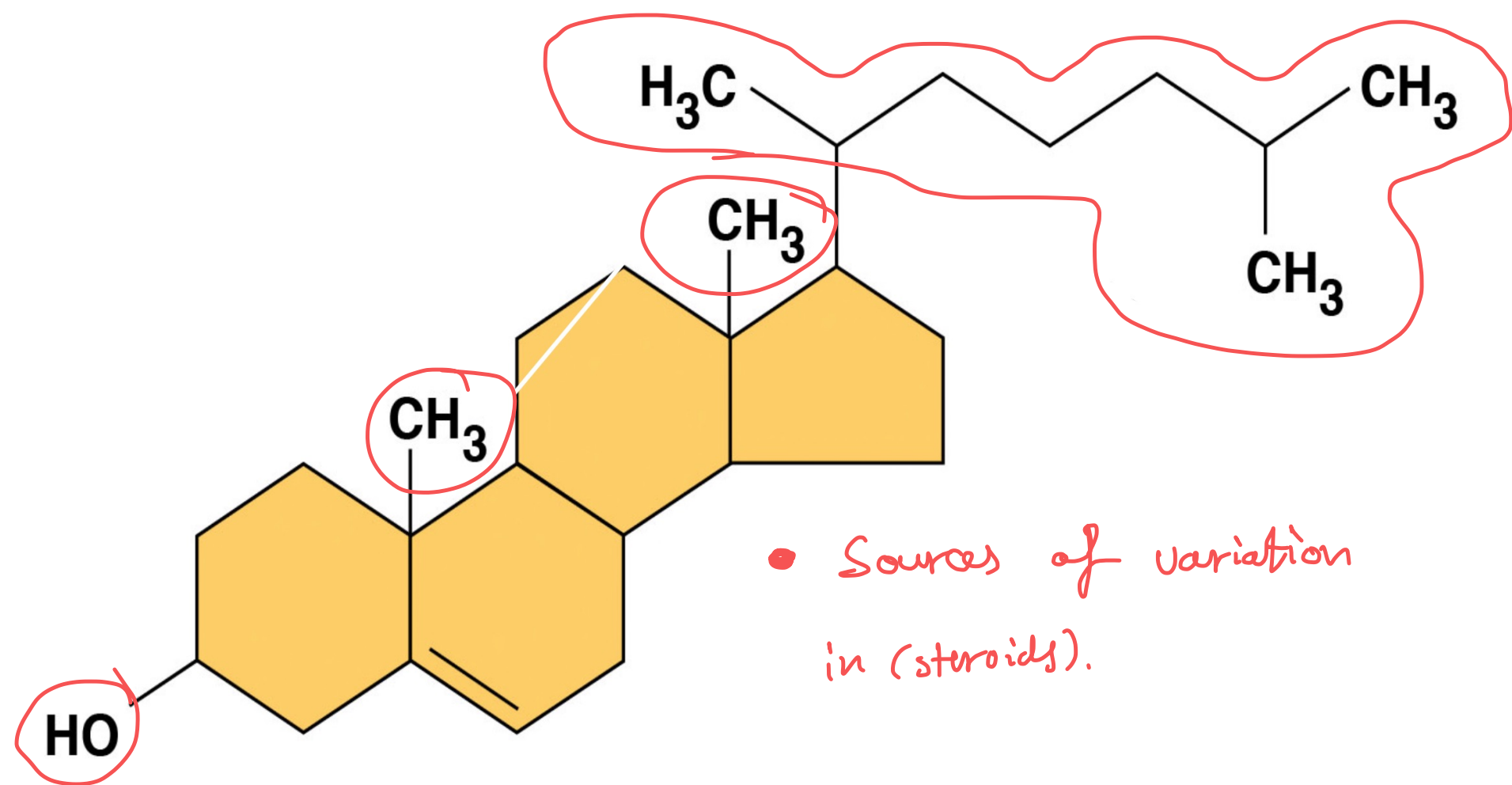
- When phospholipids are added to water, they self-assemble into double-layered sheets called bilayers
- At the surface of a cell, phospholipids are also arranged in a bilayer, with the hydrophobic tails pointing toward the interior
- The phospholipid bilayer forms a boundary between the cell and its external environment

steroids

- **Steroids** are lipids characterized by a carbon skeleton consisting of four fused rings
- **Cholesterol**, a type of steroid, is a component in animal cell membranes and a precursor from which other steroids are synthesized. *“raw material”*
- A high level of cholesterol in the blood may contribute to cardiovascular disease ⊗

⊗ *Atherosclerosis* : a vascular disease caused by
High levels of : Saturated fat or Cholesterol.

Figure 5.12



• Sources of variation in (steroids).

Concept 5.4: Proteins include a diversity of structures, resulting in a wide range of functions

- Proteins account for more than 50% of the dry mass of most cells
- Some proteins speed up chemical reactions } enzymes
- Other protein functions include defense^①, storage^②, transport^③, cellular communication^④, movement^⑤, and structural support^⑥

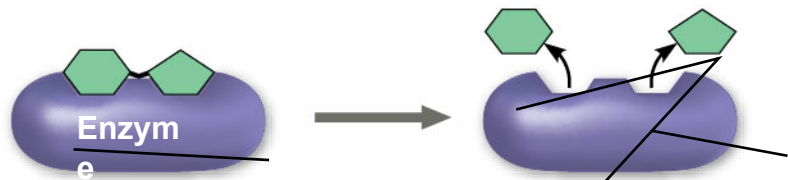


Figure 5.13a

Enzymatic proteins

Function: Selective acceleration of chemical reactions

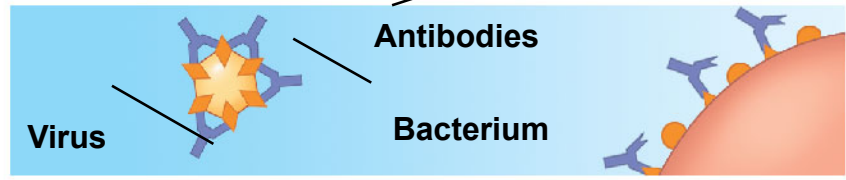
Example: Digestive enzymes catalyze the hydrolysis of bonds in food molecules.



Defensive proteins

Function: Protection against disease

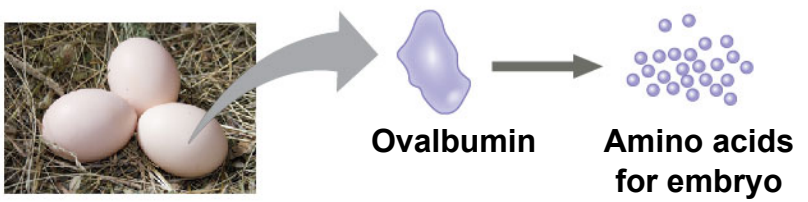
Example: Antibodies inactivate and help destroy viruses and bacteria.



Storage proteins

Function: Storage of amino acids

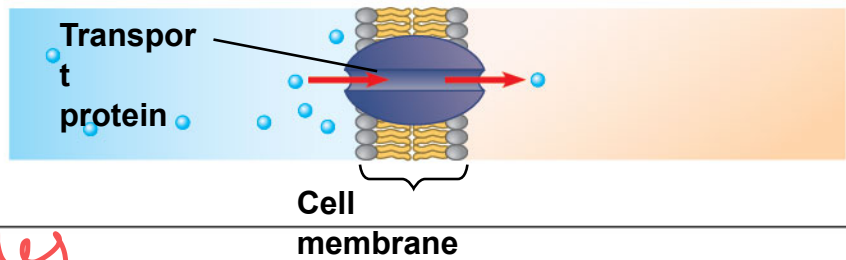
Examples: Casein, the protein of milk, is the major source of amino acids for baby mammals. Plants have storage proteins in their seeds. Ovalbumin is the protein of egg white, used as an amino acid source for the developing embryo.



Transport proteins

Function: Transport of substances

Examples: Hemoglobin, the iron-containing protein of vertebrate blood, transports oxygen from the lungs to other parts of the body. Other proteins transport molecules across membranes, as shown here.

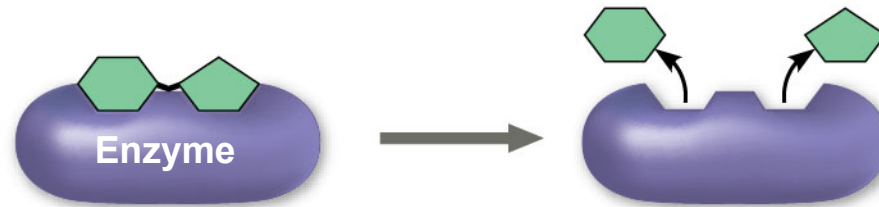


** see specific slides (SSS)*

Enzymatic proteins

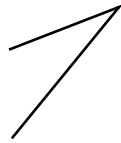
Function: Selective acceleration of chemical reactions

Example: Digestive enzymes catalyze the hydrolysis of bonds in food molecules.



Such as maltase

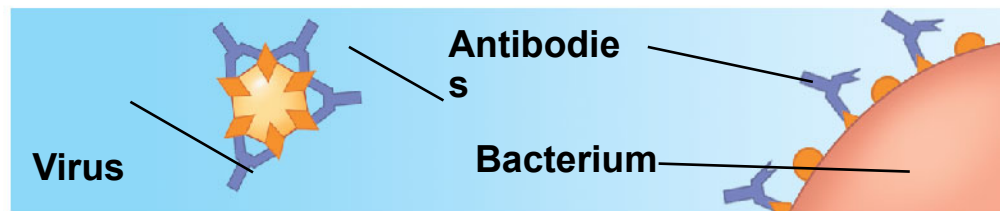
catalyzes (maltose + H₂O → 2 glucose)



Defensive proteins

Function: Protection against disease

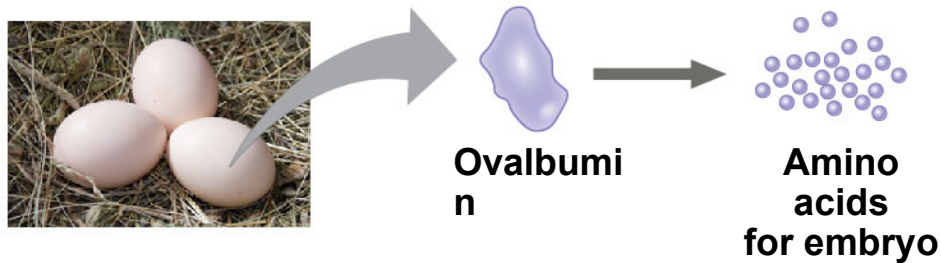
Example: Antibodies inactivate and help destroy viruses and bacteria.



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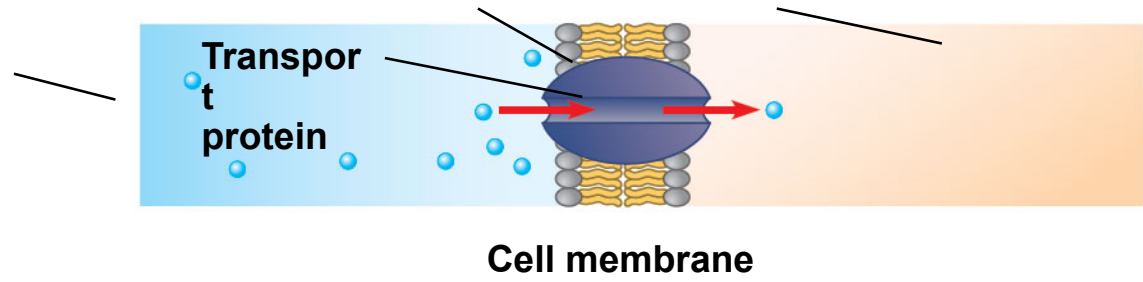
Examples: Casein, the protein of milk, is the major source of amino acids for baby mammals. Plants have storage proteins in their seeds. Ovalbumin is the protein of egg white, used as an amino acid source for the developing embryo.



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Function: Transport of substances

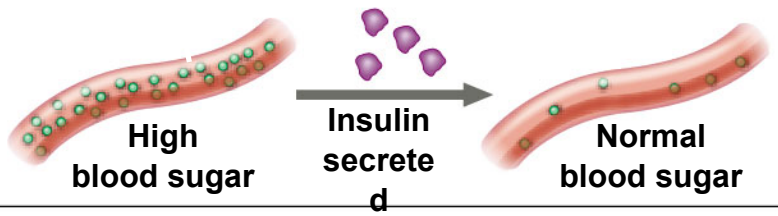
Examples: Hemoglobin, the iron-containing protein of vertebrate blood, transports oxygen from the lungs to other parts of the body. Other proteins transport molecules across membranes, as shown here.



Hormonal proteins

Function: Coordination of an organism's activities

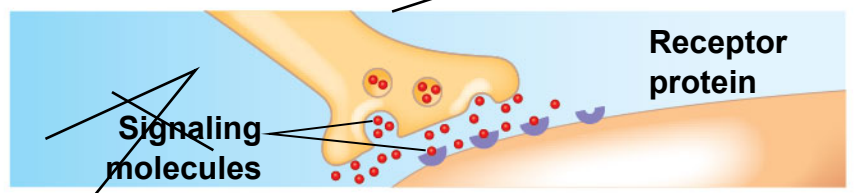
Example: Insulin, a hormone secreted by the pancreas, causes other tissues to take up glucose, thus regulating blood sugar concentration.



Receptor proteins

Function: Response of cell to chemical stimuli

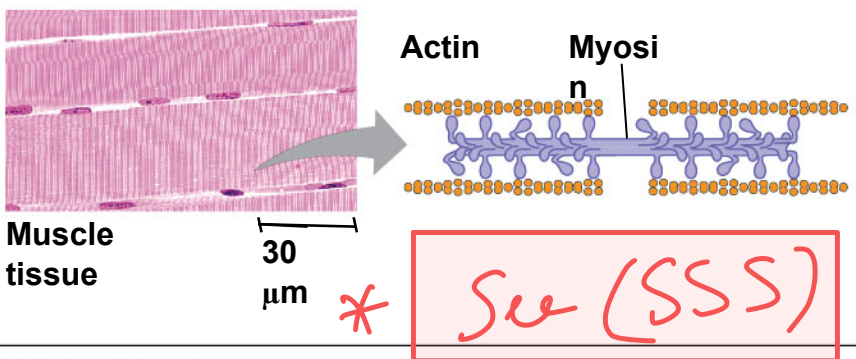
Example: Receptors built into the membrane of a nerve cell detect signaling molecules released by other nerve cells.



Contractile and motor proteins

Function: Movement

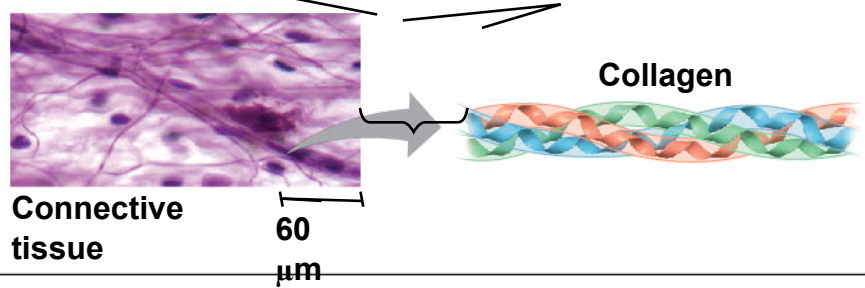
Examples: Motor proteins are responsible for the undulations of cilia and flagella. Actin and myosin proteins are responsible for the contraction of muscles.



Structural proteins

Function: Support

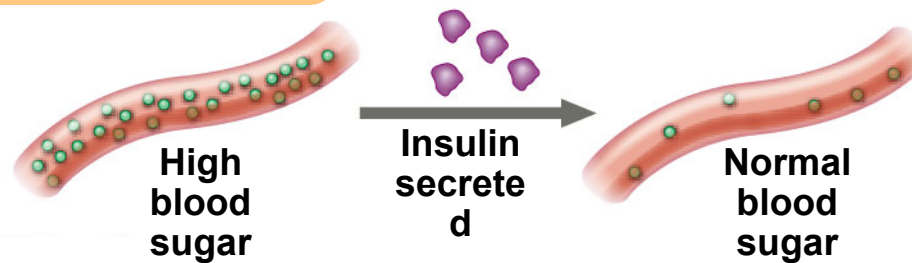
Examples: Keratin is the protein of hair, horns, feathers, and other skin appendages. Insects and spiders use silk fibers to make their cocoons and webs, respectively. Collagen and elastin proteins provide a fibrous framework in animal connective tissues.



Hormonal proteins

Function: Coordination of an organism's activities

Example: Insulin, a hormone secreted by the pancreas, causes other tissues to take up glucose, thus regulating blood sugar concentration.

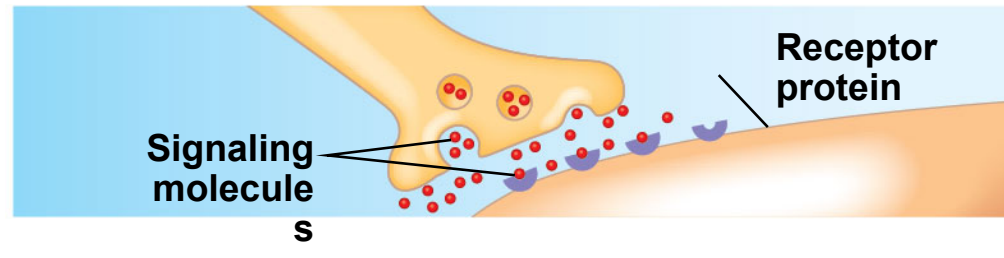




Receptor proteins

Function: Response of cell to chemical stimuli

Example: Receptors built into the membrane of a nerve cell detect signaling molecules released by other nerve cells.



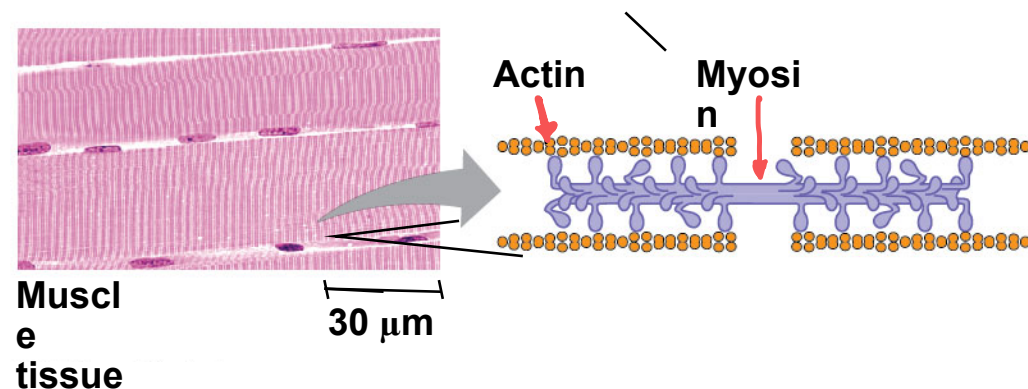


Contractile and motor proteins

Function: Movement

Examples: Motor proteins are responsible for the undulations of cilia and flagella.

Actin and myosin proteins are responsible for the contraction of muscles.



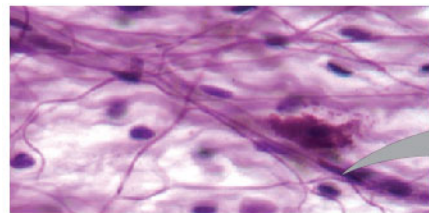
Structural proteins

Function: Support

Examples: Keratin is the protein of hair, horns, feathers, and other skin appendages.

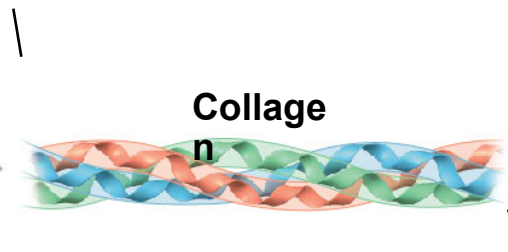
Insects and spiders use silk fibers to make their cocoons and webs, respectively.

Collagen and elastin proteins provide a fibrous framework in animal connective tissues.



Connective tissue

60 μm



- Enzymes are proteins that act as **catalysts** to speed up chemical reactions
- Enzymes can perform their functions repeatedly, functioning as **workhorses** that carry out the processes of life

Because they are not consumed during the Chemical Reaction.

- Proteins are all constructed from the same set of 20 amino acids
- Polypeptides are unbranched polymers built from these amino acids
- A protein is a biologically functional molecule that consists of one or more polypeptides

Primary structure

Amino Acid Monomers

- **Amino acids** are **organic molecules** with **amino** and **carboxyl groups**
- **Amino acids** differ in their properties **due to differing side chains**, called **R groups**

⊕ all amino acids are in Ionized Form
since aqueous solutions and pH and other conditions
are mostly suitable for.

for ex:

Acidic A. Acids \Rightarrow (-)

Basic \Leftarrow \Rightarrow (+)

carboxyl group \Rightarrow COO^-

Amino group \Rightarrow NH_3^+

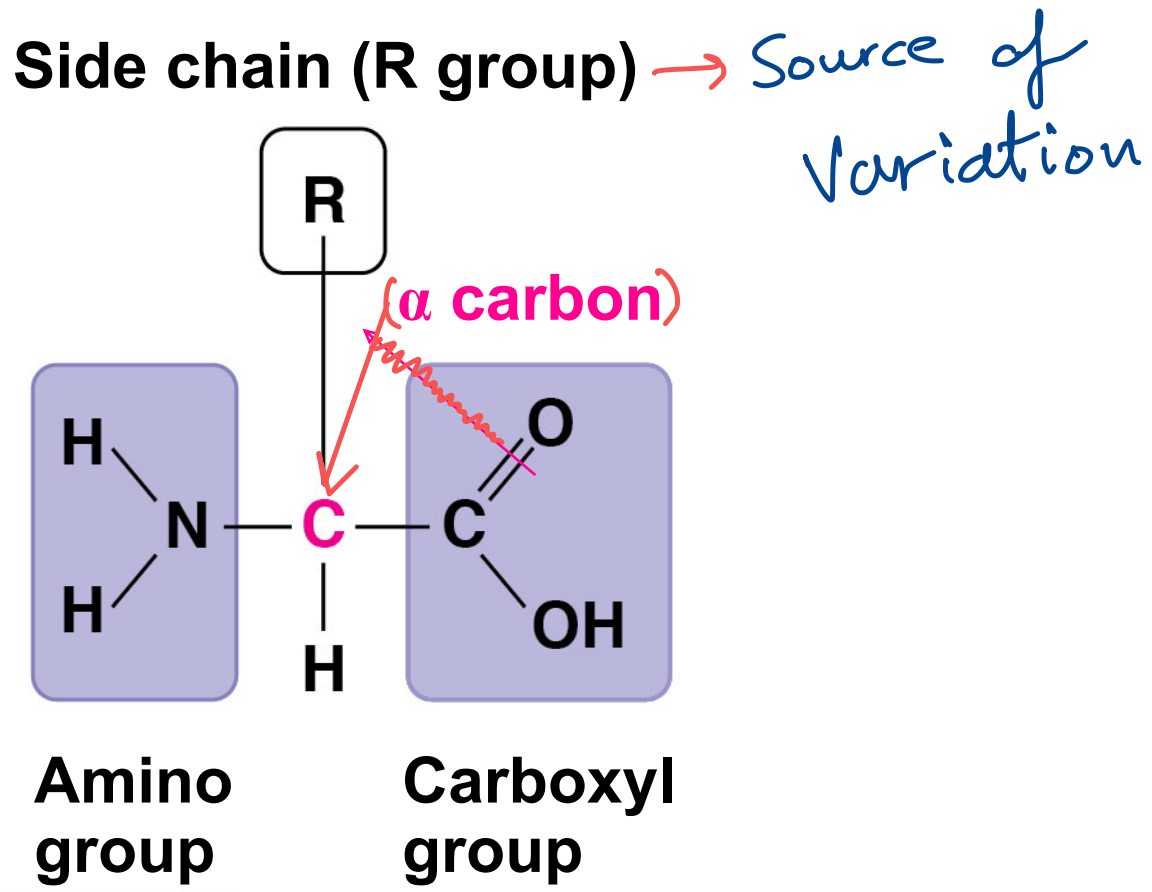
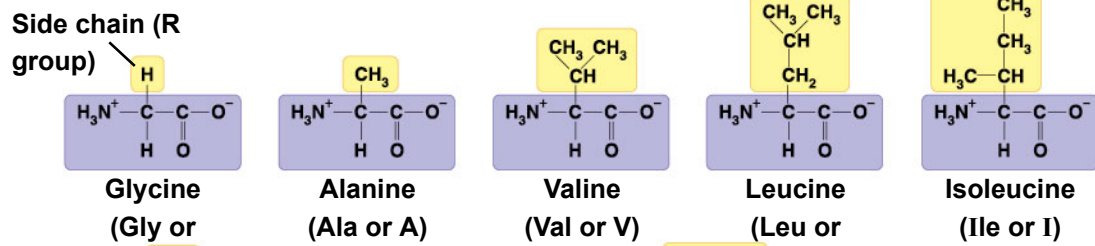
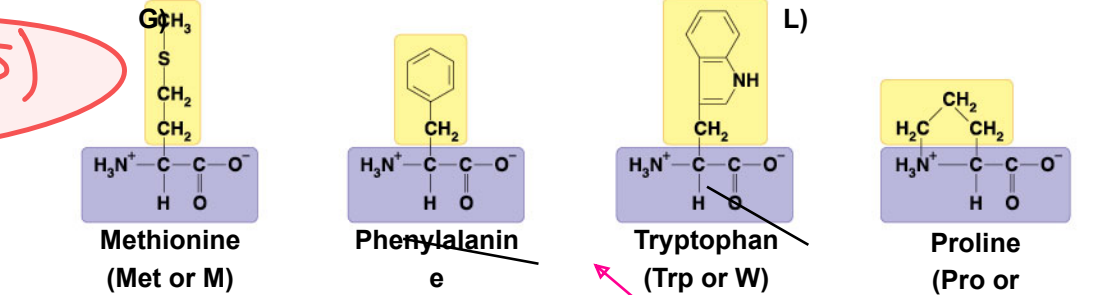


Figure 5.14

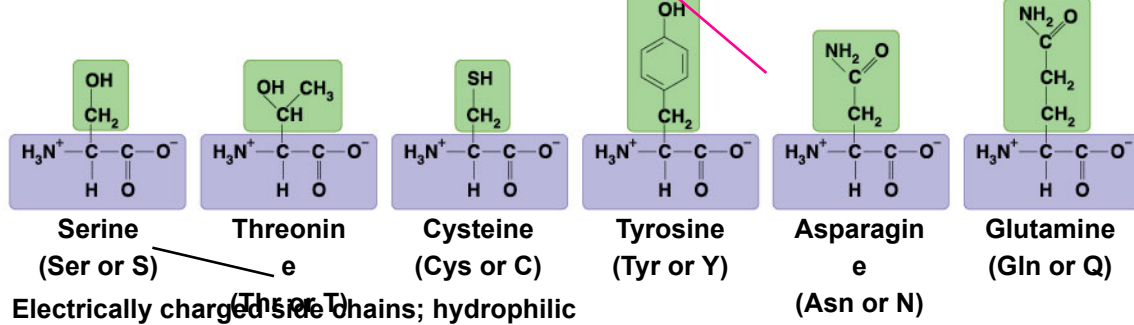
Nonpolar side chains; hydrophobic



See (SSS)

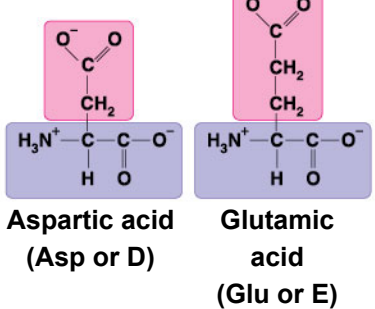


Polar side chains; hydrophilic



Electrically charged side chains; hydrophilic

Acidic (negatively charged)



Basic (positively charged)

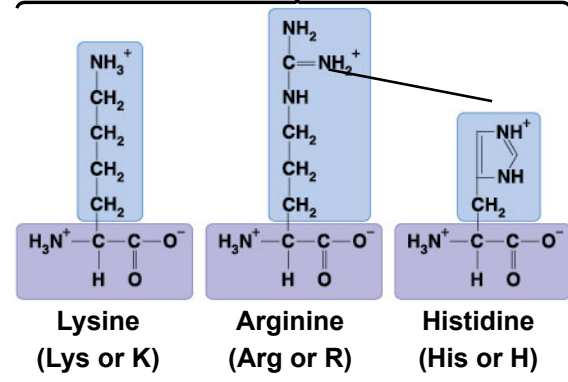
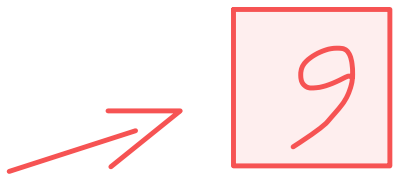
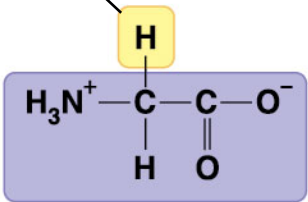


Figure 5.14a

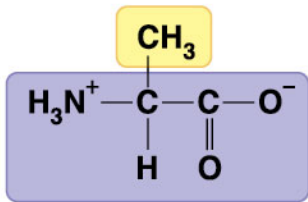


Nonpolar side chains; hydrophobic

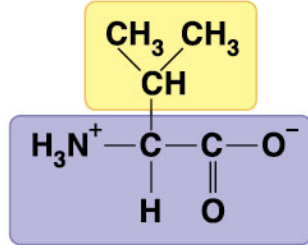
Side chain (R group)



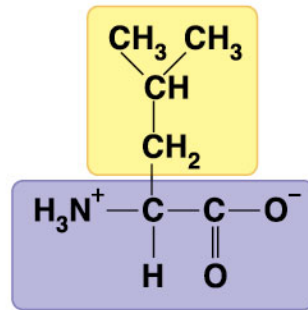
Glycine
(Gly or G)



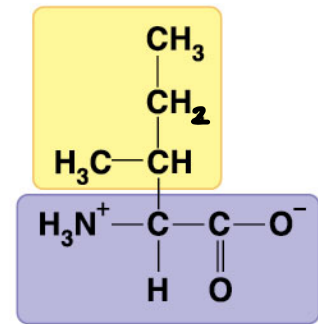
Alanine
(Ala or A)



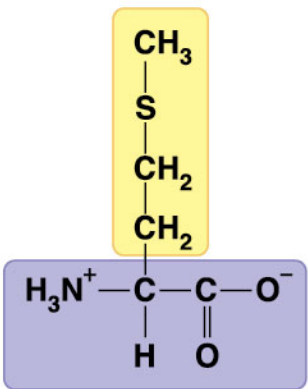
Valine
(Val or V)



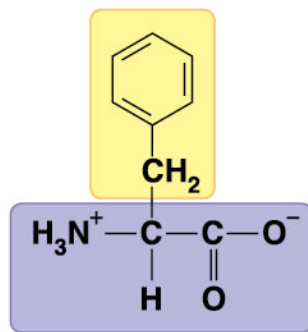
Leucine
(Leu or L)



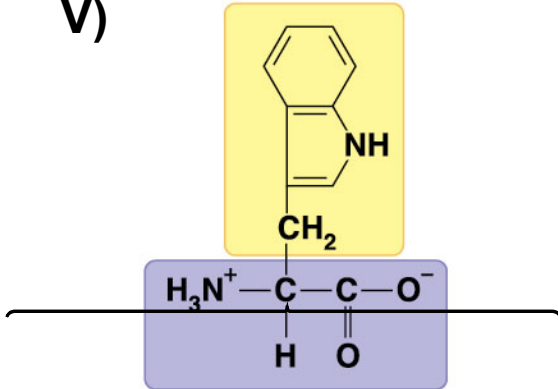
Isoleucine
(Ile or I)



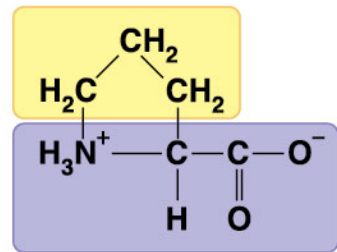
Methionine
(Met or M)



Phenylalanine
(Phe or F)

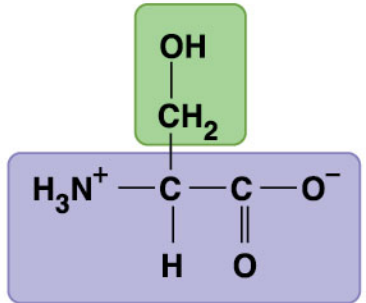


Tryptophan
(Trp or W)

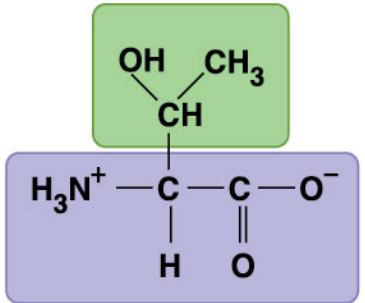


Proline
(Pro or P)

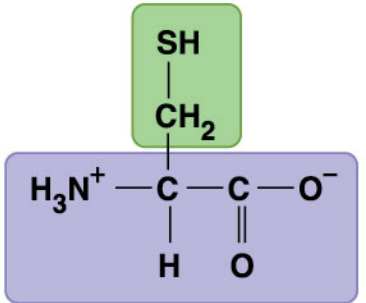
Polar side chains; hydrophilic



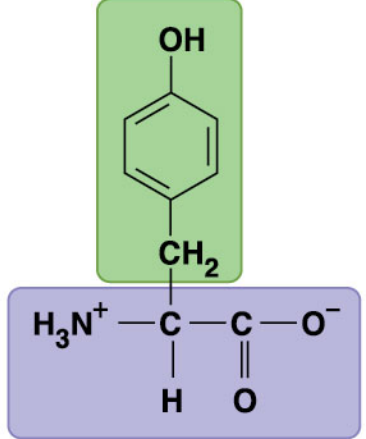
Serine
(Ser or S)



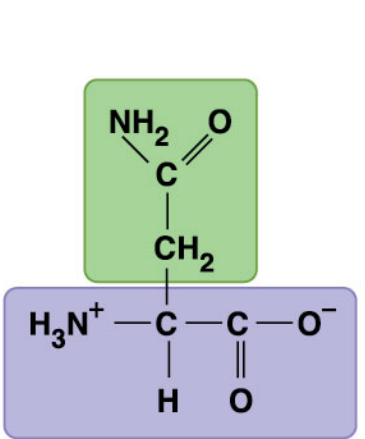
Threonine
(Thr or T)



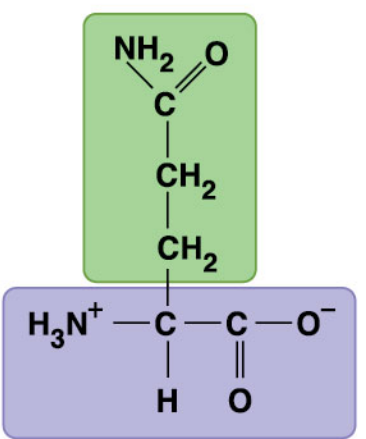
Cysteine
(Cys or C)



Tyrosine
(Tyr or Y)



Asparagine
(Asn or N)



Glutamine
(Gln or Q)

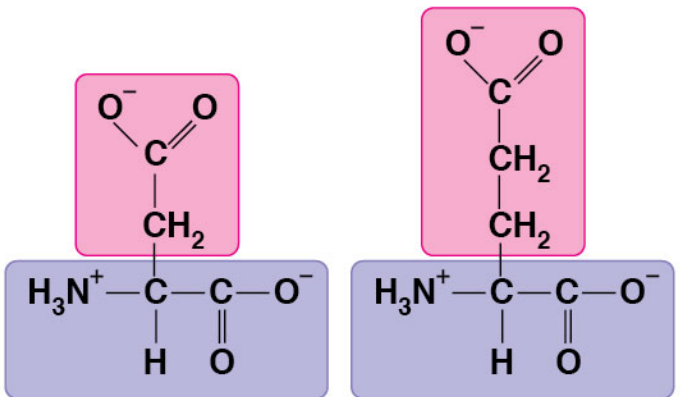
Figure 5.14c

Electrically charged side chains; hydrophilic

After reacting

Acidic (negatively charged)

After reacting

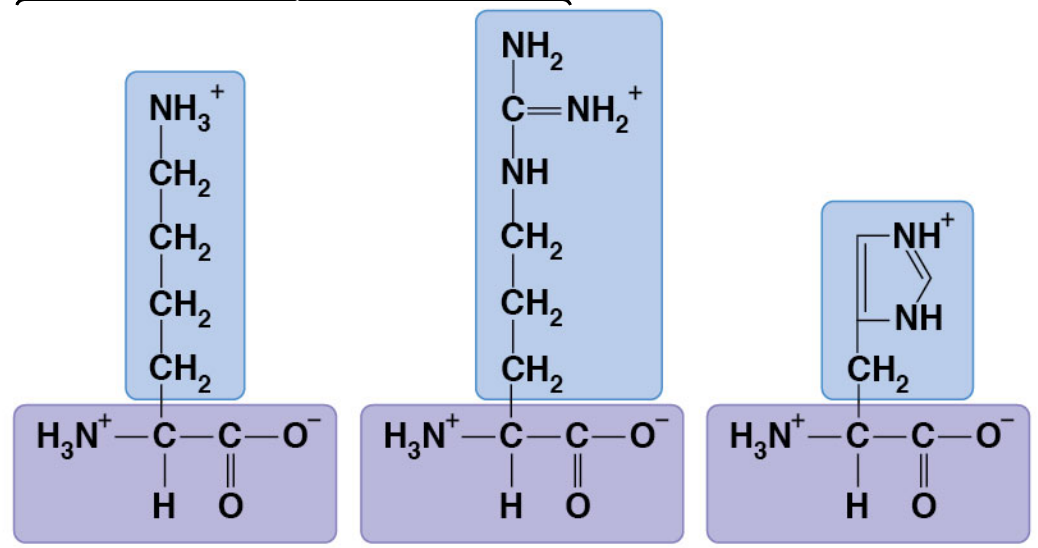


Aspartic acid (Asp or D)

Glutamic acid (Glu or E)

2

Basic (positively charged)



Lysine (Lys or K)

Arginine (Arg or R)

Histidine (His or H)

3

Polypeptides (Amino Acid Polymers)

- Amino acids are linked by covalent bonds called **peptide bonds** ⇒ *Carboxyl — Amino linkage*
- A polypeptide is a polymer of amino acids
- Polypeptides range in length from a few to more than 1,000 monomers
- Each polypeptide has a unique linear sequence of amino acids, with a carboxyl end (C-terminus) and an amino end (N-terminus)
- Amino acids vary in water solubility according to the R group.

Figure 5.15

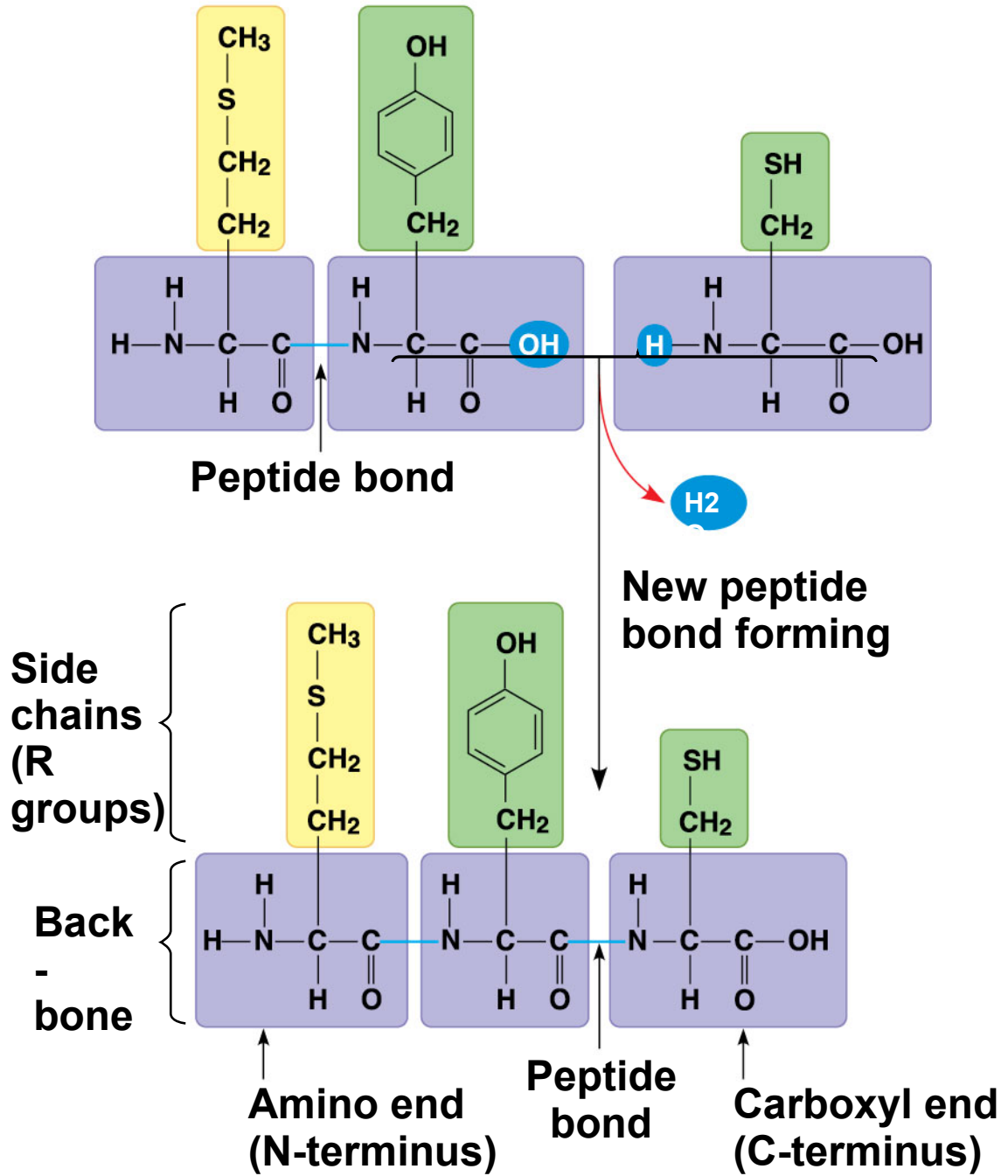


Figure 5.15a

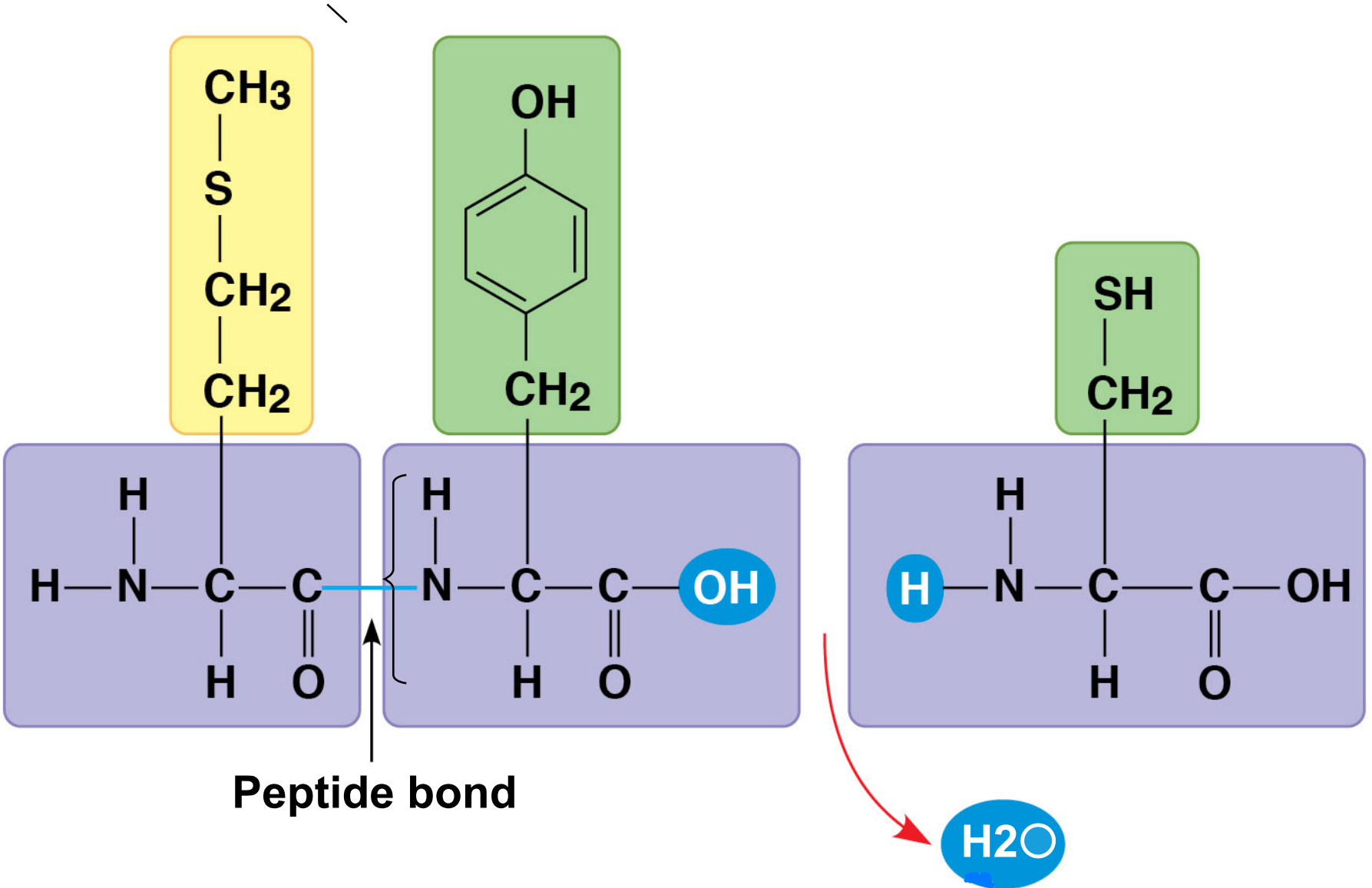
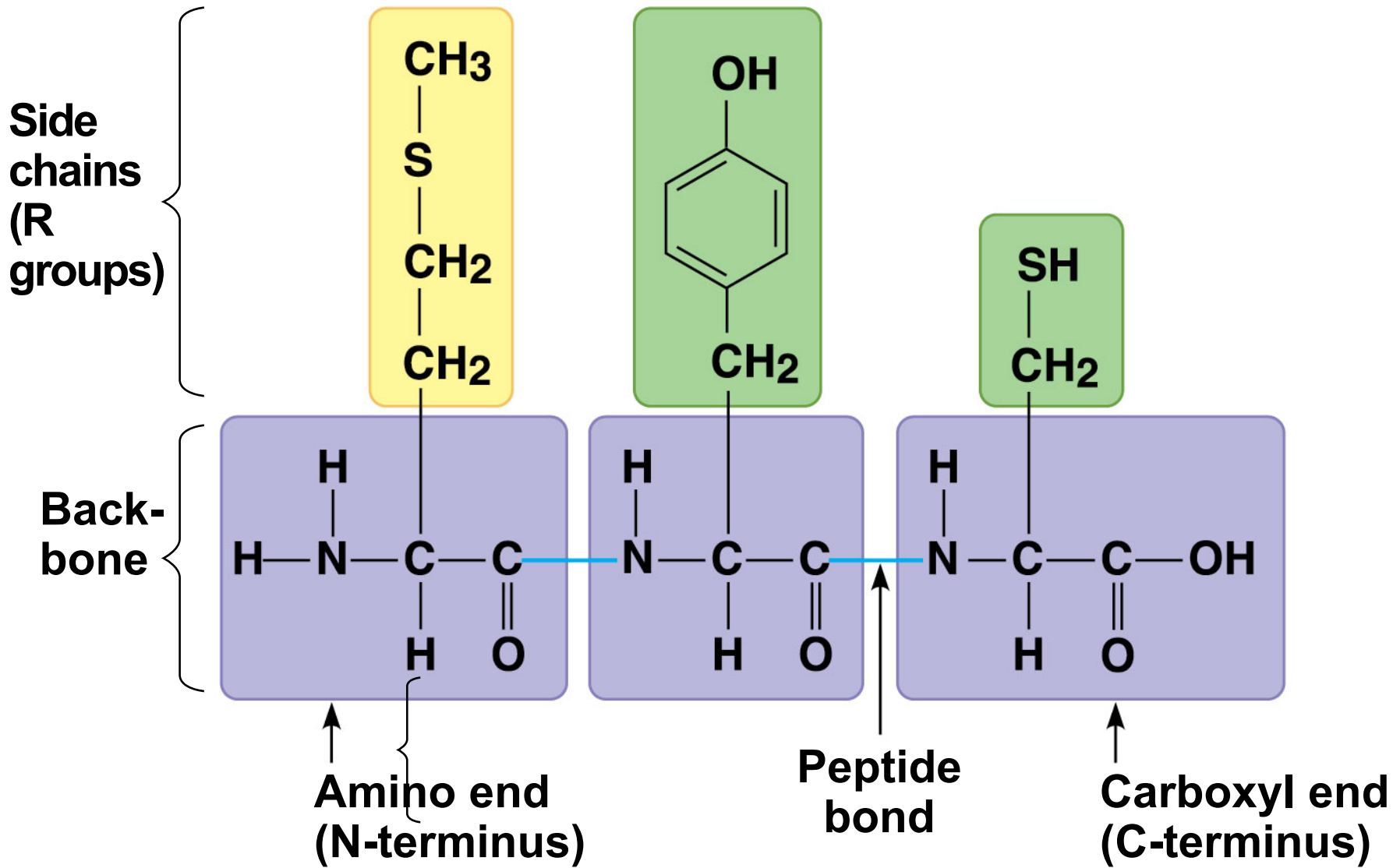


Figure 5.15b

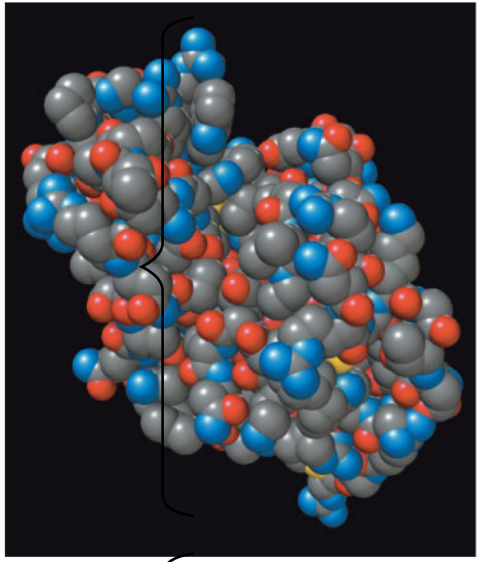


Protein Structure and Function

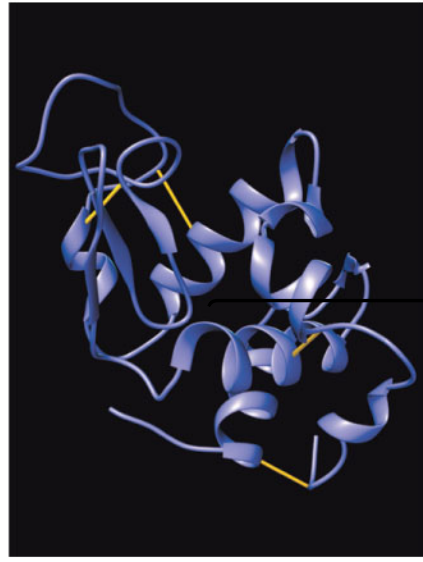
- The specific activities of proteins result from their intricate three-dimensional architecture
- A functional protein consists of one or more polypeptides precisely twisted, folded, and coiled into a unique shape

Figure 5.16

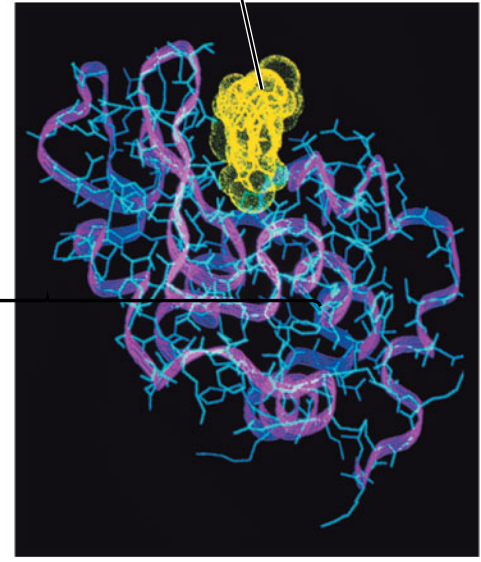
Structural Models



Space-filling model

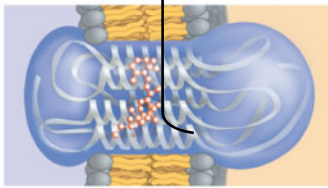


Ribbon model

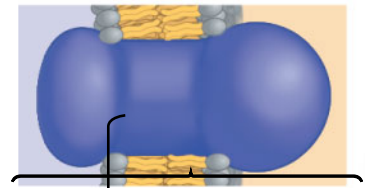


Wire-frame model (blue)

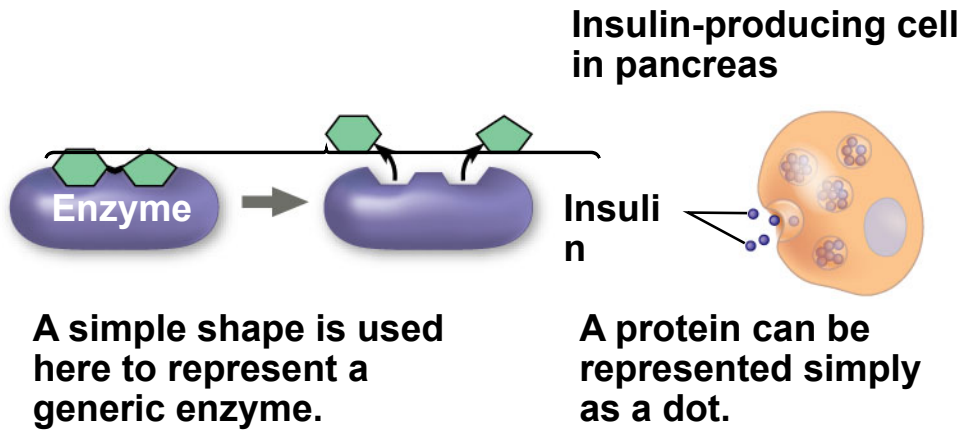
Simplified Diagrams



A transparent shape shows the overall shape of the molecule and some internal details.

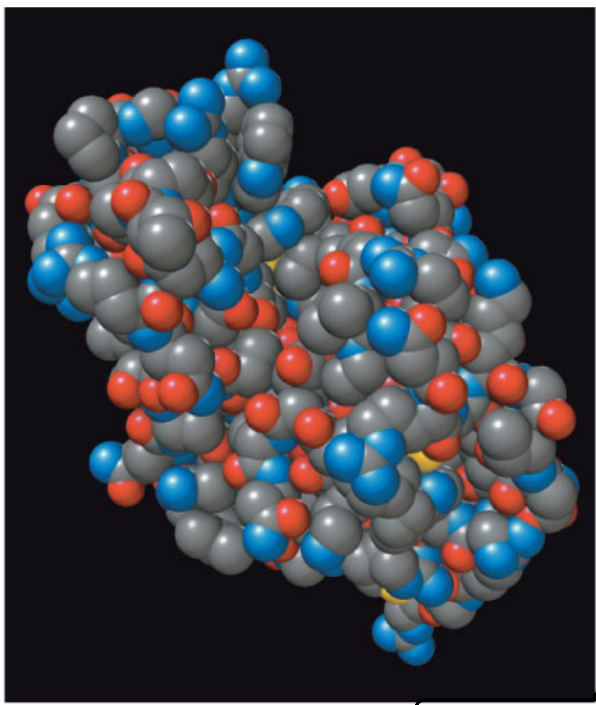


A solid shape is used when structural details are not needed.

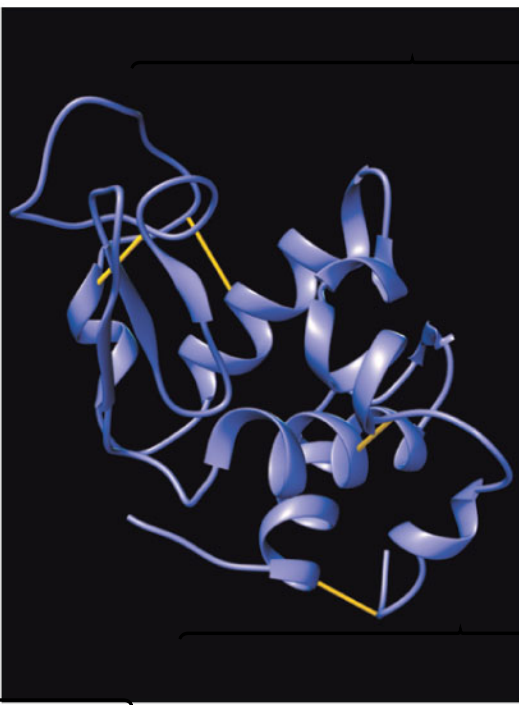


Structural Models

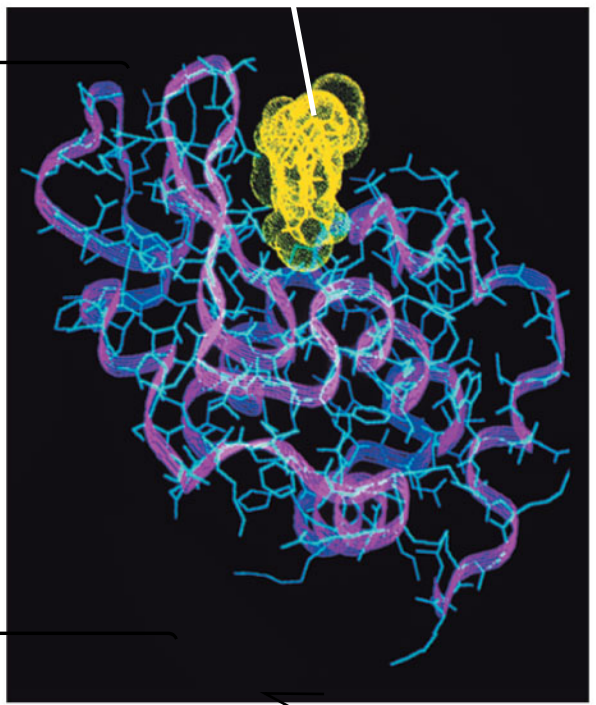
Target molecule (on bacterial cell surface) bound to lysozyme



Space-filling model

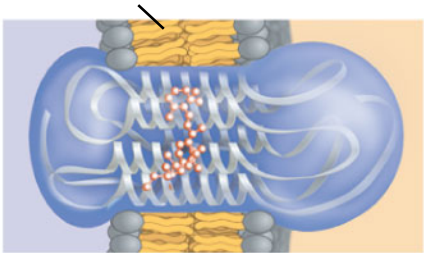


Ribbon model

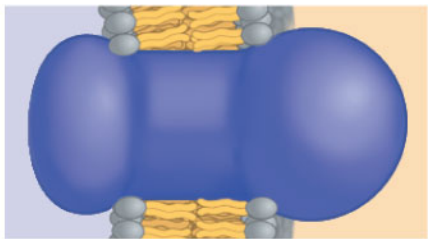


Wire-frame model (blue)

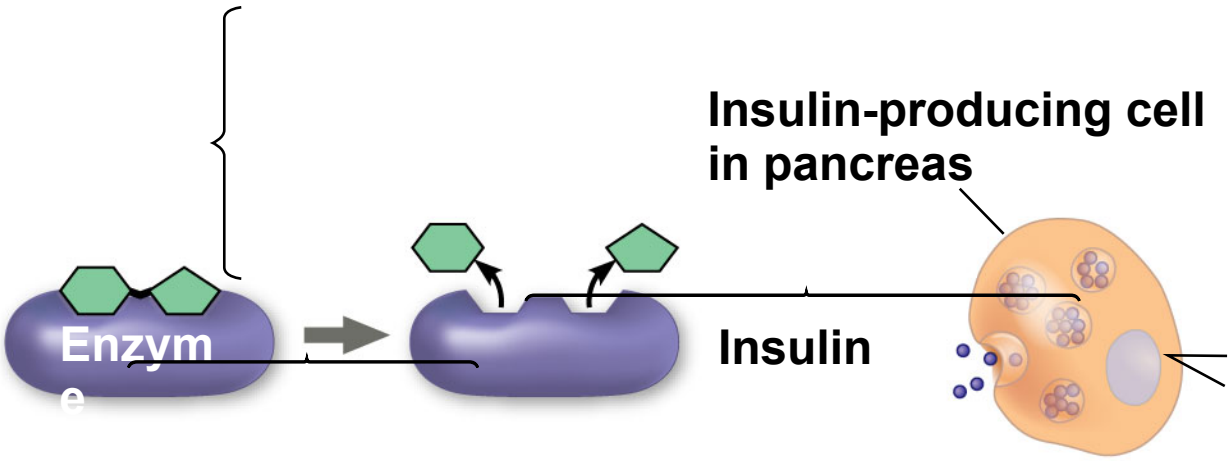
Simplified Diagrams



A transparent shape shows the overall shape of the molecule and some internal details.



A solid shape is used when structural details are not needed.



A simple shape is used here to represent a generic enzyme.

A protein can be represented simply as a dot.


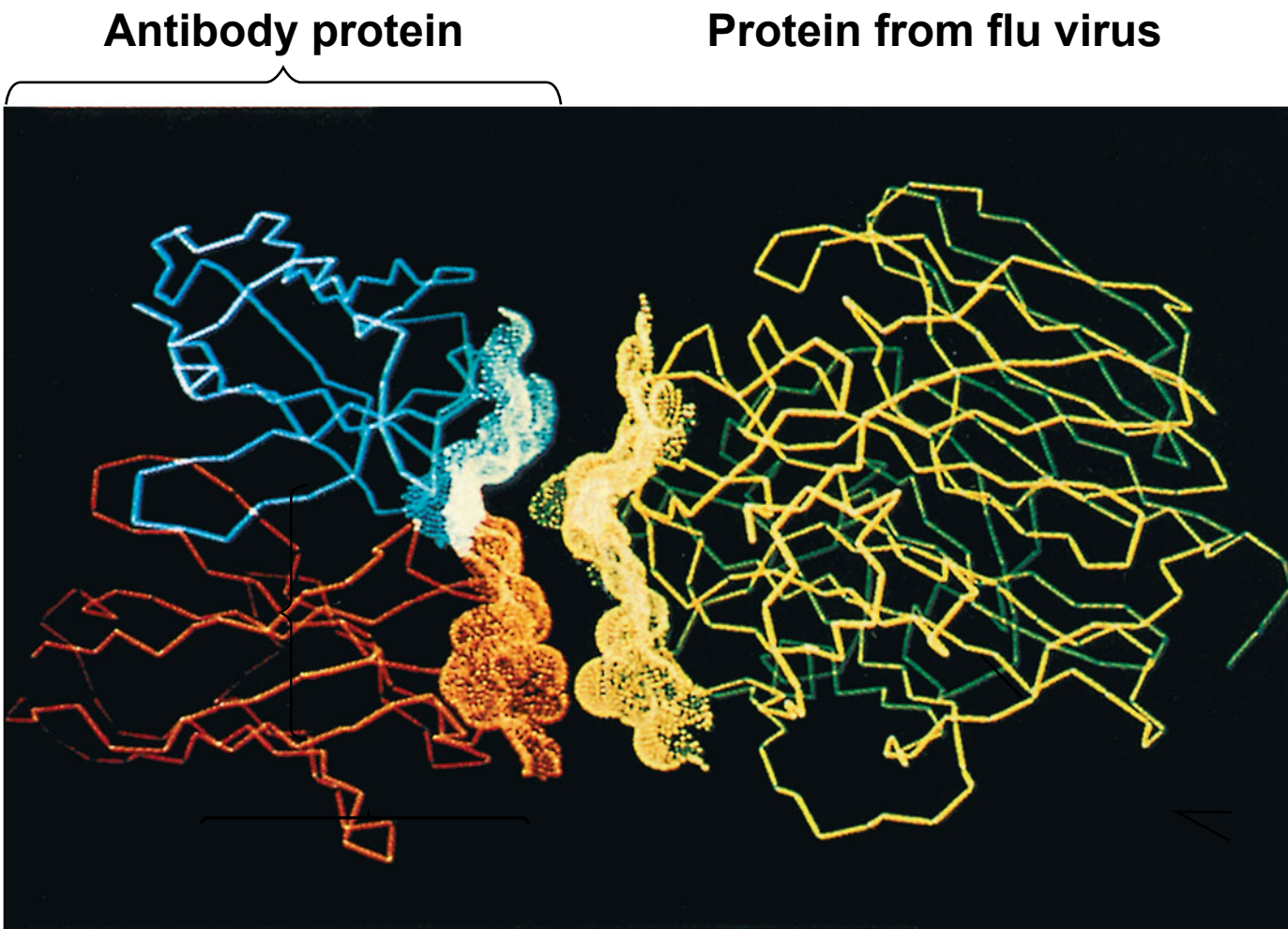
- The sequence of amino acids determines a protein's three-dimensional structure
 - A protein's structure determines how it works
 - The function of a protein usually depends on its ability to recognize and bind to some other molecule
- 

Figure 5.17



our Levels of Protein Structure

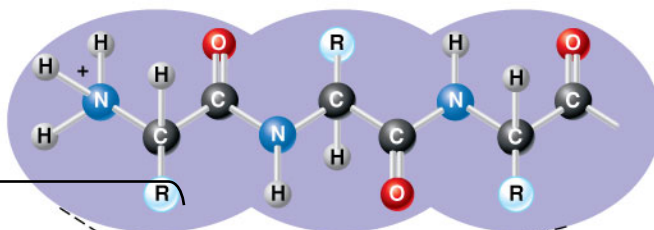
- The primary structure of a protein is its unique sequence of amino acids
- Secondary structure, found in most proteins, consists of coils and folds in the polypeptide chain
- Tertiary structure is determined by interactions among various side chains (R groups)
- Quaternary structure results when a protein consists of multiple polypeptide chains

Figure 5.18a

Primary Structure

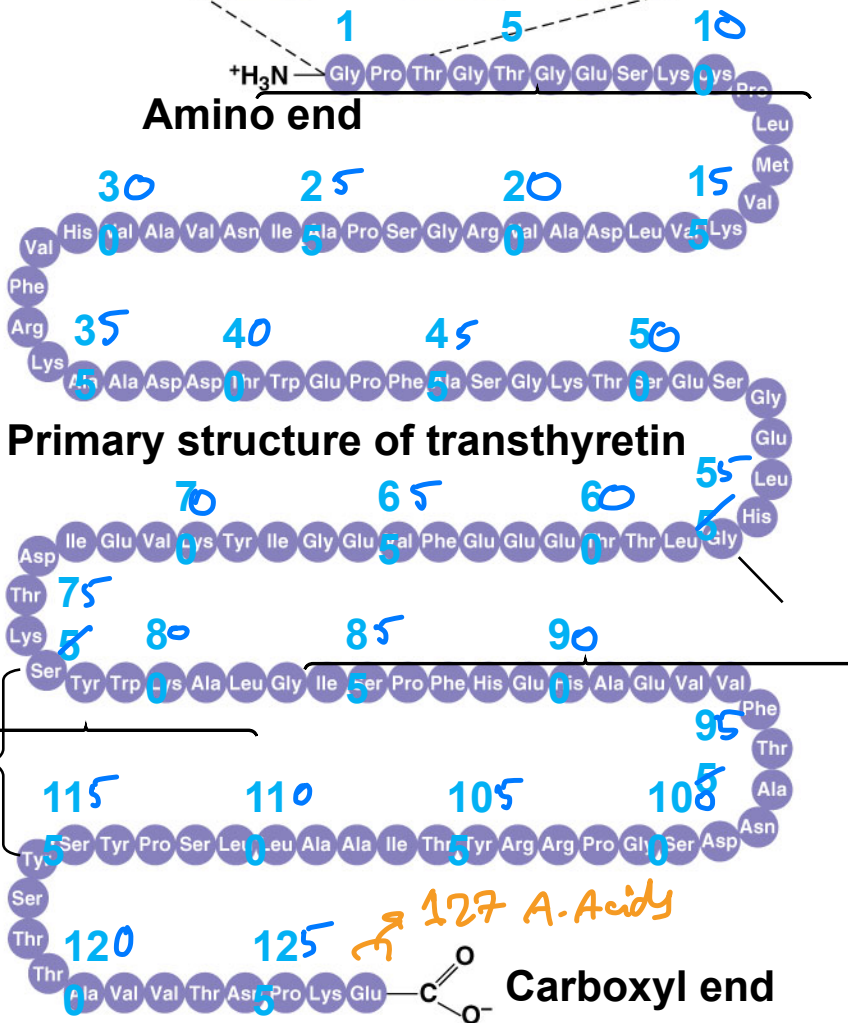
\otimes Transthyretin:
4 polypeptide chains

Amino acids



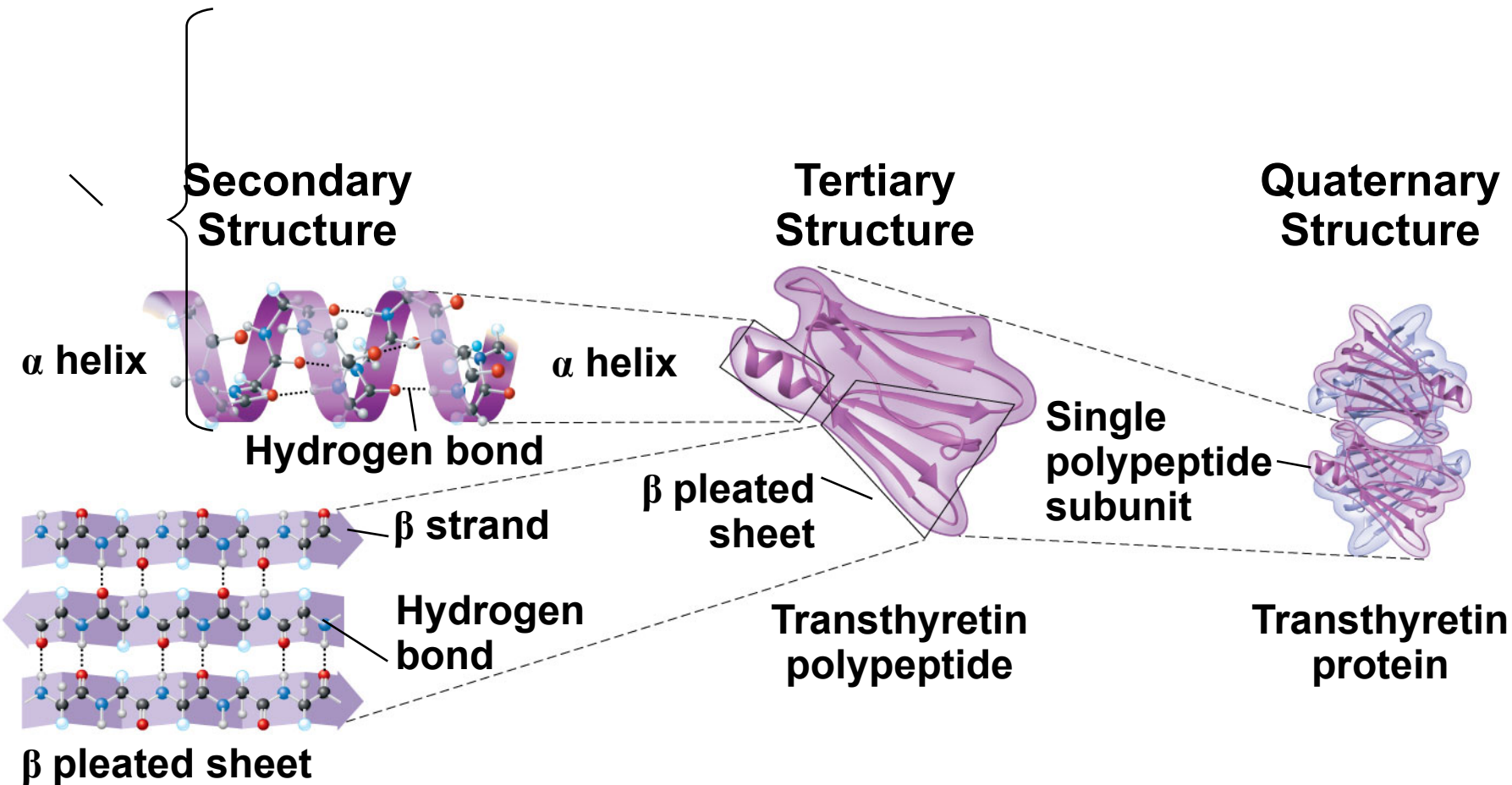
127 A.A. acids per chain

transfers Vitamin A & one of thyroid hormones



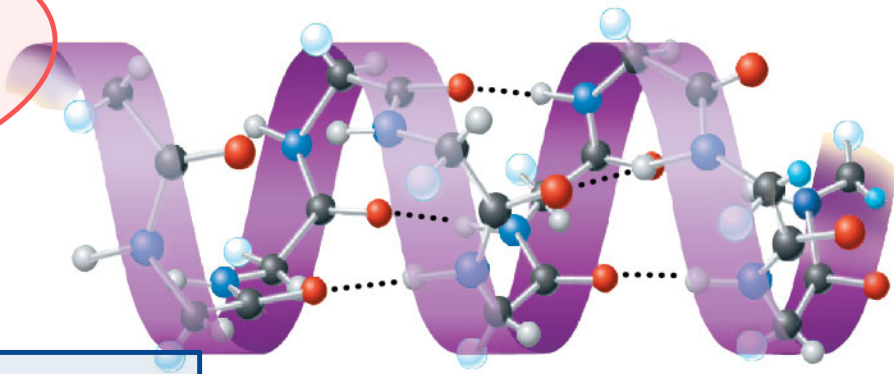
127 A.A. acids

Figure 5.18b



Secondary Structure

H-Bonds with every 4th A-Acid.



α helix

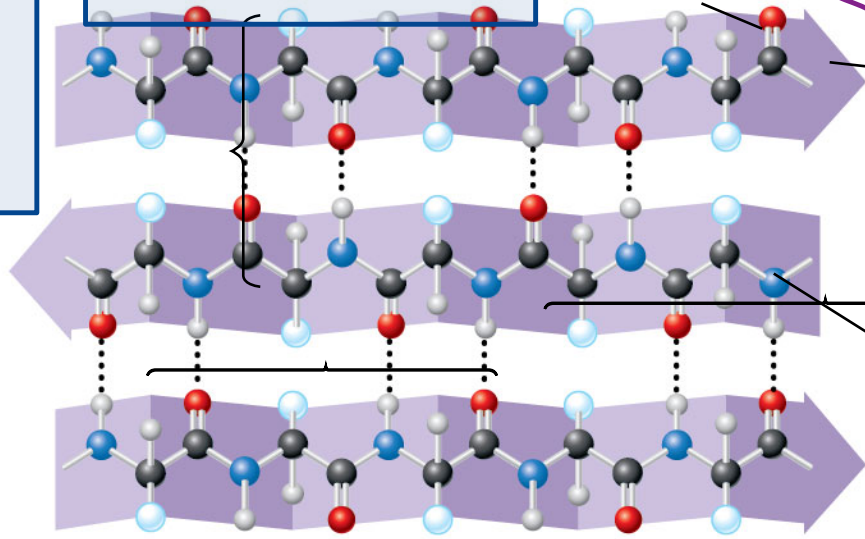
Between Back bones

Hydrogen bonds

Btwn $O^{\delta-}$ et $H^{\delta+}$ Carbonyl Amin

Bonds Between R-chains are in further structures

tertiary & quaternary

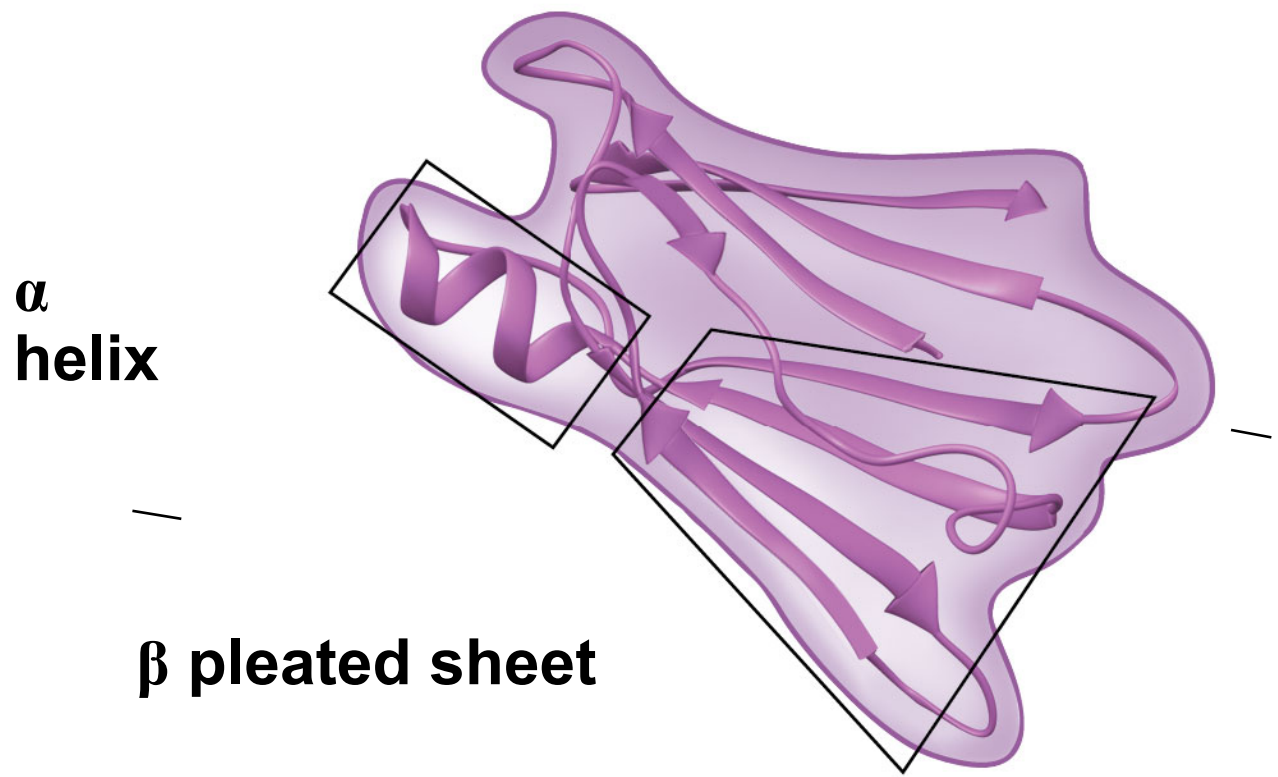


β strand

Hydrogen bond

β pleated sheet

Tertiary Structure



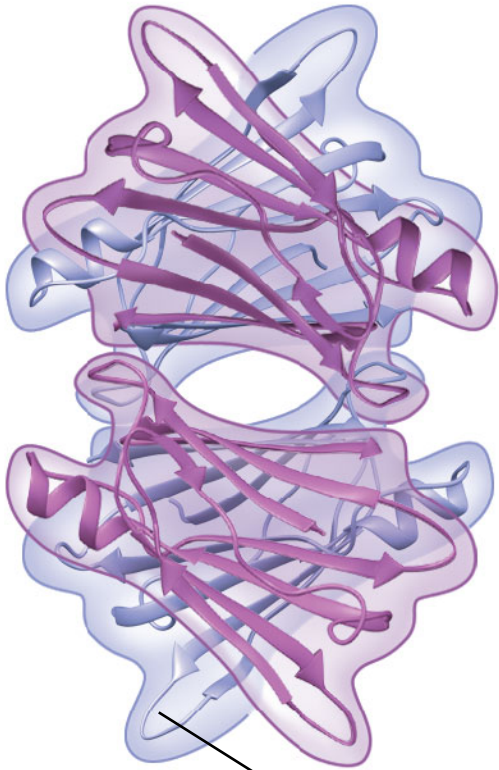
α
helix

β pleated sheet

Transthyretin
polypeptide

Quaternary Structure

Single polypeptide subunit



Transthyretin protein

Figure 5.18c

+ (mainly)

⊕ Roughly :

Globular Proteins : many β p. sheets

Fibrous Proteins : many α Helices



⊕ α -keratin (protein of hair) has many α -Helices
So do silk-protein & other structural proteins.

Figure 5.18d

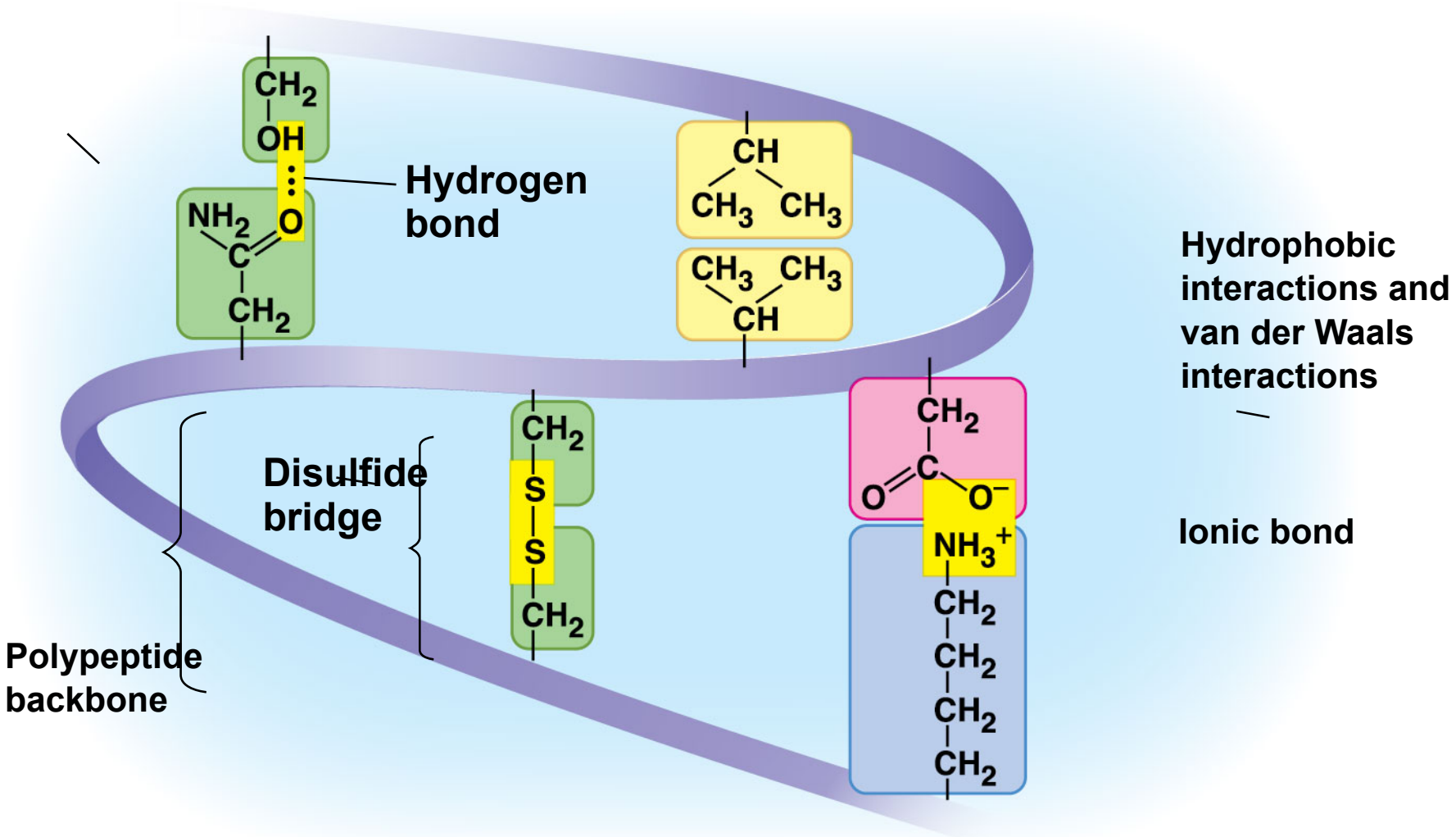


Figure 5.18e

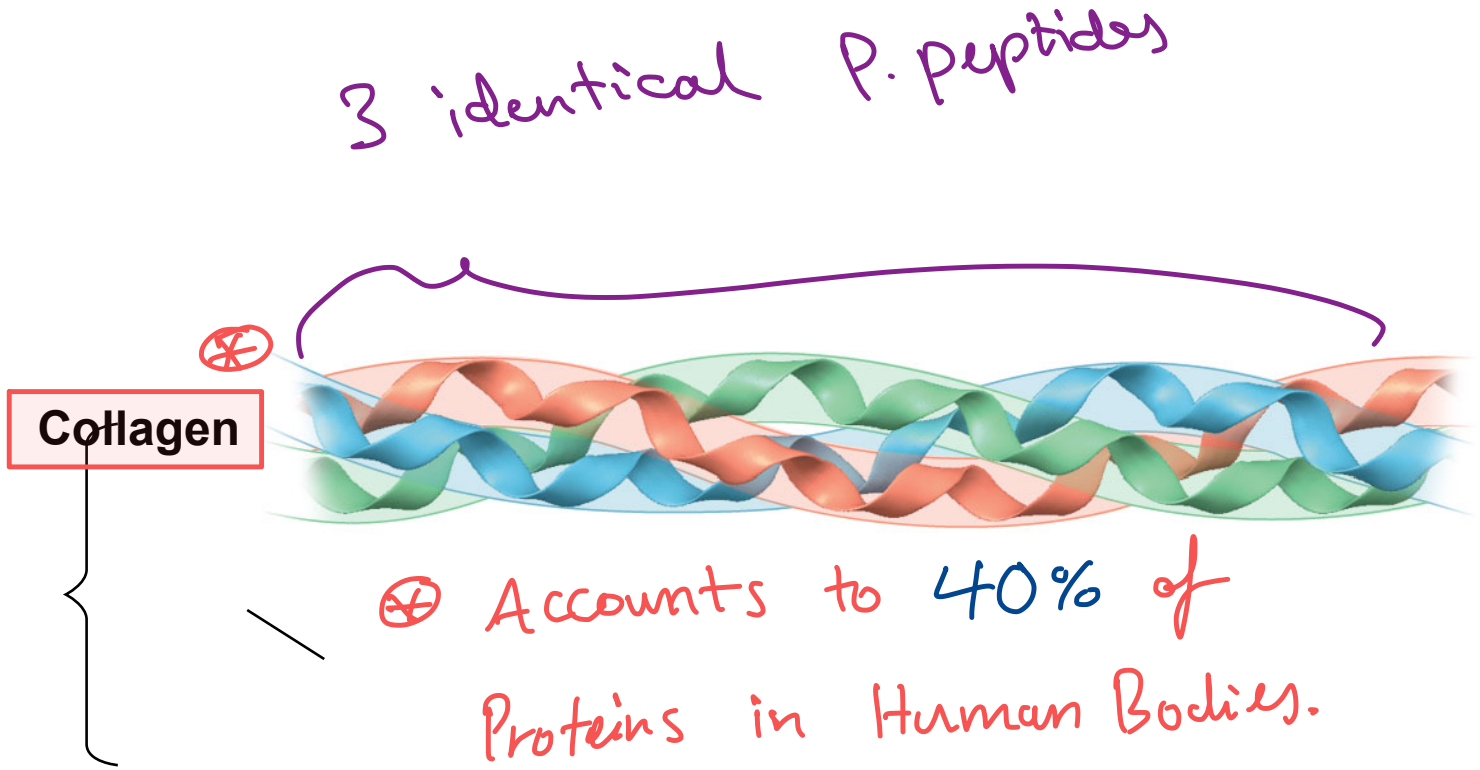
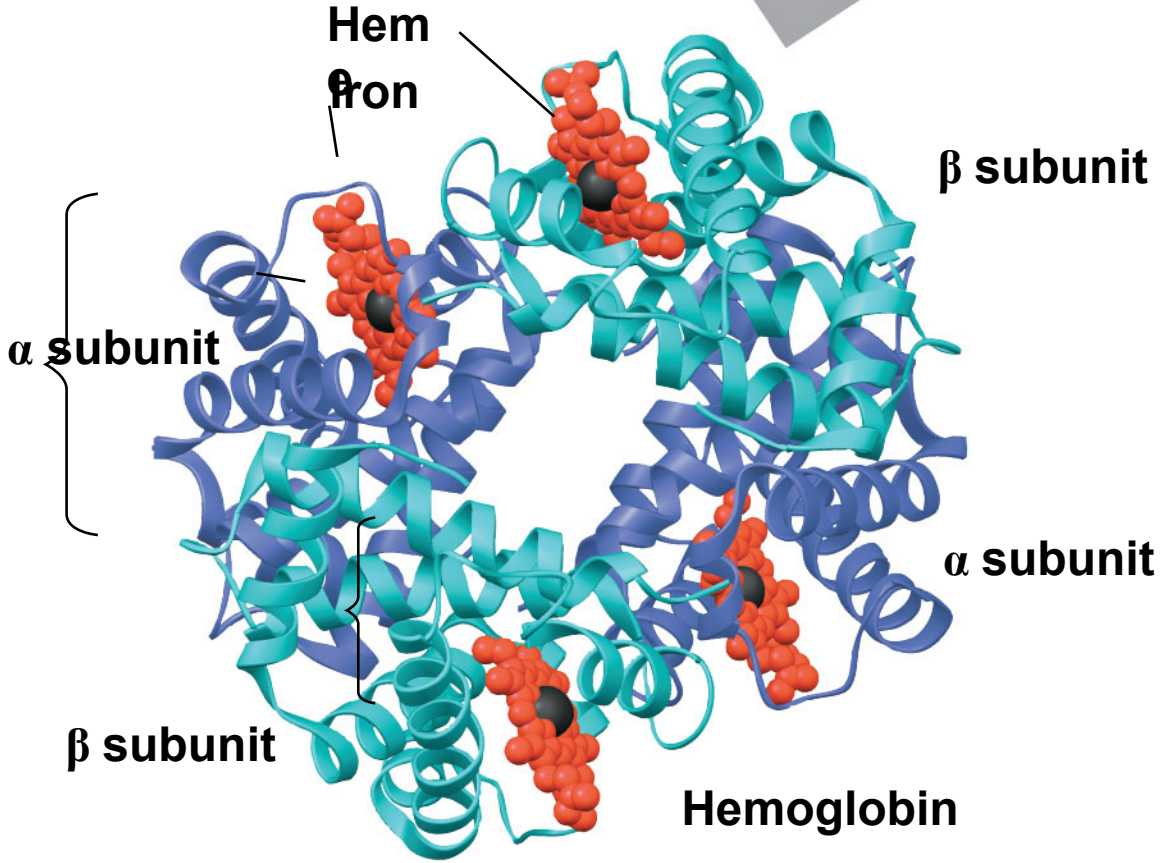


Figure 5.18f

Hemoglobin is not functional unless all four chains are assembled

Quaternary structure must be formed for hemoglobin to be functional



- The **primary structure** of a protein is its sequence of amino acids
- Primary structure is like the order of letters in a long word
- Primary structure is determined by inherited genetic information

}

- The coils and folds of secondary structure result from hydrogen bonds between repeating constituents of the polypeptide backbone
- Typical secondary structures are a coil called an α helix and a folded structure called a β pleated sheet

- **Tertiary structure**, the overall shape of a polypeptide, results from interactions between R groups, rather than interactions between backbone constituents
- These interactions include hydrogen bonds, ionic bonds, hydrophobic interactions, and van der Waals interactions
- Strong covalent bonds called disulfide bridges may reinforce the protein's structure

* Hydrophobic interaction as a name is misleading; when water exclude non-polar R chains they cluster when they come close together, V der W's work.

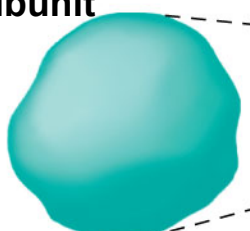
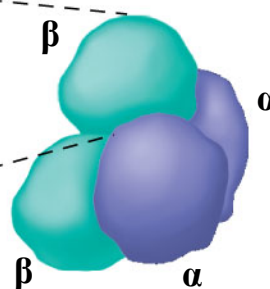
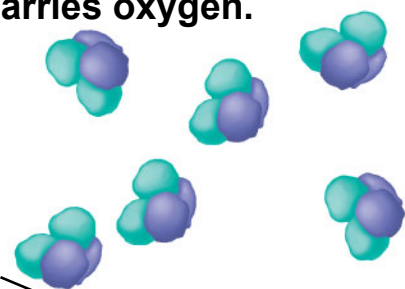
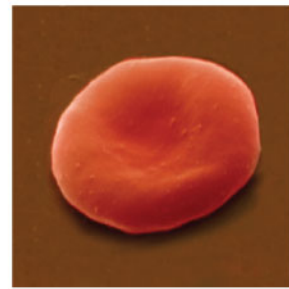
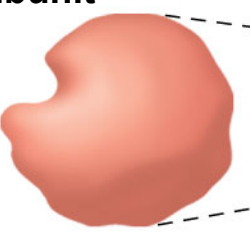
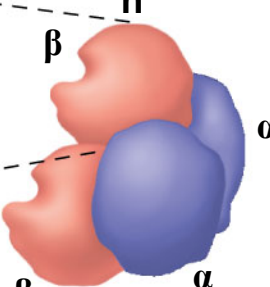
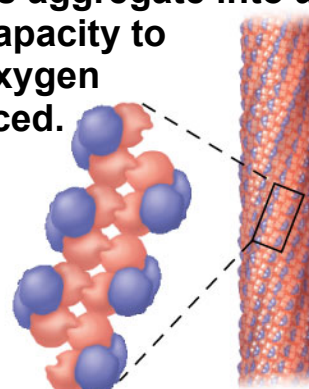

- **Quaternary structure** results when two or more polypeptide chains form one macromolecule
- Collagen is a fibrous protein consisting of three polypeptides coiled like a rope
- Hemoglobin is a globular protein consisting of four polypeptides: two α and two β subunits

Sickle-Cell Disease: A Change in Primary Structure

- A slight change in primary structure can affect a protein's structure and ability to function
- **Sickle-cell disease**, an inherited blood disorder, results from a single amino acid substitution in the protein hemoglobin
- The abnormal hemoglobin molecules cause the red blood cells to aggregate into chains and to deform into a sickle shape

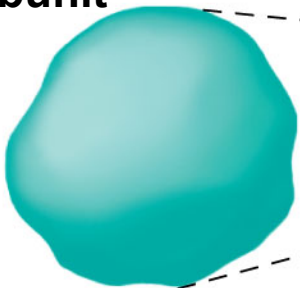
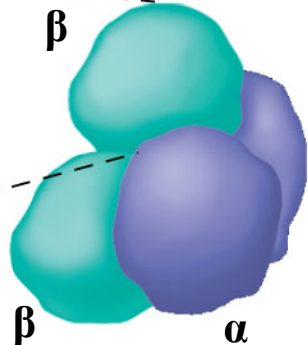
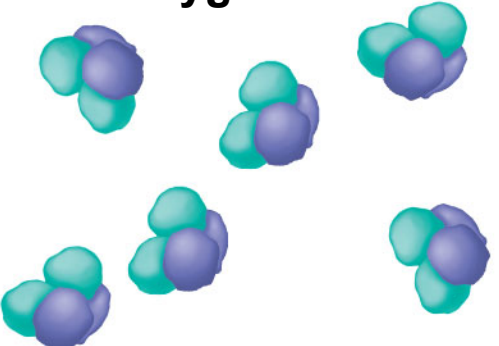


Figure 5.19

	Primary Structure	Secondary and Tertiary Structures	Quaternary Structure	Function	Red Blood Cell Shape
Normal	1 Val 2 His 3 Leu 4 Thr 5 Pro 6 Glu 7 Glu	Normal β subunit 	Normal hemoglobin 	Proteins do not associate with one another; each carries oxygen. 	 5 μm
Sickle-cell	1 Val 2 His 3 Leu 4 Thr 5 Pro 6 Val 7 Glu	Sickle-cell β subunit 	Sickle-cell hemoglobin 	Proteins aggregate into a fiber; capacity to carry oxygen is reduced. 	 5 μm

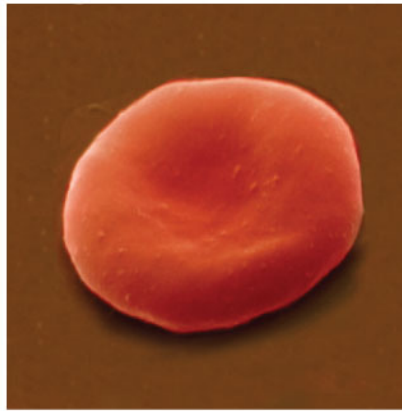
see (SSS)

Figure 5.19a

	Primary Structure	Secondary and Tertiary Structures	Quaternary Structure	Function
Normal	1 Val 2 His 3 Leu 4 Thr 5 Pro 6 Glu 7 Glu	Normal β subunit 	Normal hemoglobin 	Proteins do not associate with one another; each carries oxygen. 
		<i>Acidic</i>		

Quaternary structures remain separated from one another to maximize gas transportation and work properly.

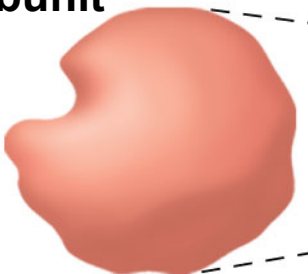
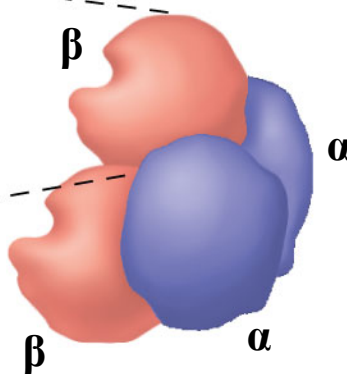
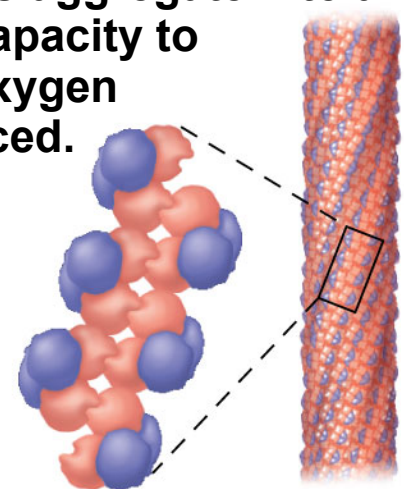
Normal Red - Blood Cell.



5
μm



Figure 5.19b

	Primary Structure	Secondary and Tertiary Structures	Quaternary Structure	Function
Sickle-cell	1 Val 2 His 3 Leu 4 Thr 5 Pro 6 Val 7 Glu	Sickle-cell β subunit  <i>Hydrophobic Non-polar.</i>	Sickle-cell hemoglobin 	Proteins aggregate into a fiber; capacity to carry oxygen is reduced. 

Abnormal sickle-cell.



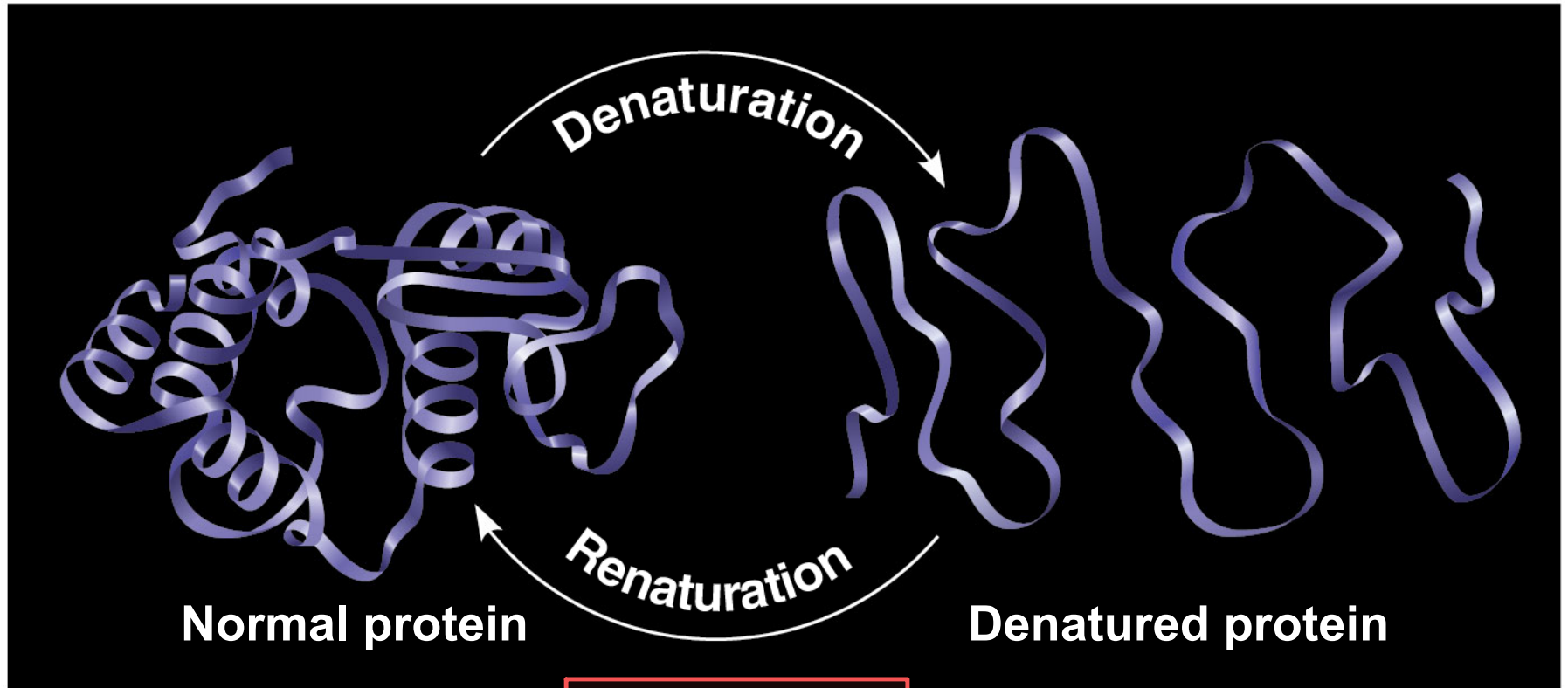
5
μm

What Determines Protein Structure?

- In addition to primary structure, **physical** and **chemical** conditions can affect structure
- Alterations in pH, salt concentration, temperature, or other environmental factors can cause a protein to unravel
- This loss of a protein's native structure is called **denaturation** ⊗
- A denatured protein is biologically inactive

⊗ Denaturation may occur if proteins are placed in a non-polar solvent ⇒ they refold so the non-polar [hydrophobic parts] face the outside (solvent).

Figure 5.20



Renaturation is not always possible

it depends on conditions and other factors.

Protein Folding in the Cell

- It is hard to predict a protein's structure from its primary structure
- Most proteins probably go through several stages on their way to a stable structure
- Diseases such as Alzheimer's^①, Parkinson's^②, and mad cow disease are associated with misfolded proteins^③
+ { cystic fibrosis }^④

- Scientists use **X-ray crystallography** to **determine a protein's structure**
- Another method is **nuclear magnetic resonance (NMR) spectroscopy**, which **does not require protein crystallization**
- **Bioinformatics** is **another approach** to **prediction of protein structure from amino acid sequences**

⊕ Intrinsically disordered Proteins ⊗

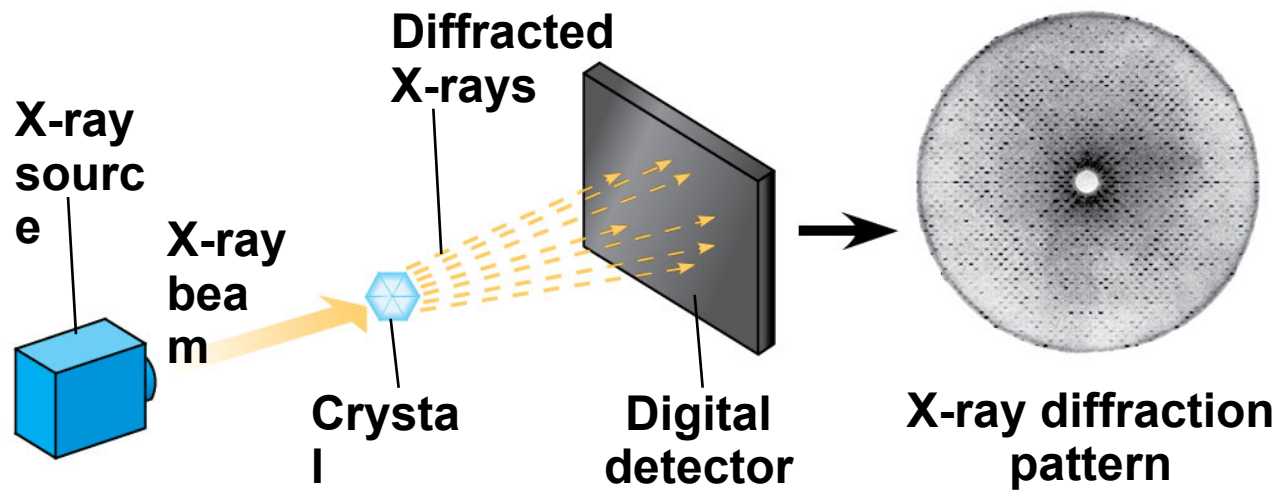
account for (20-30)% of mammalian Proteins

⊗ don't have a specific 3-D structure

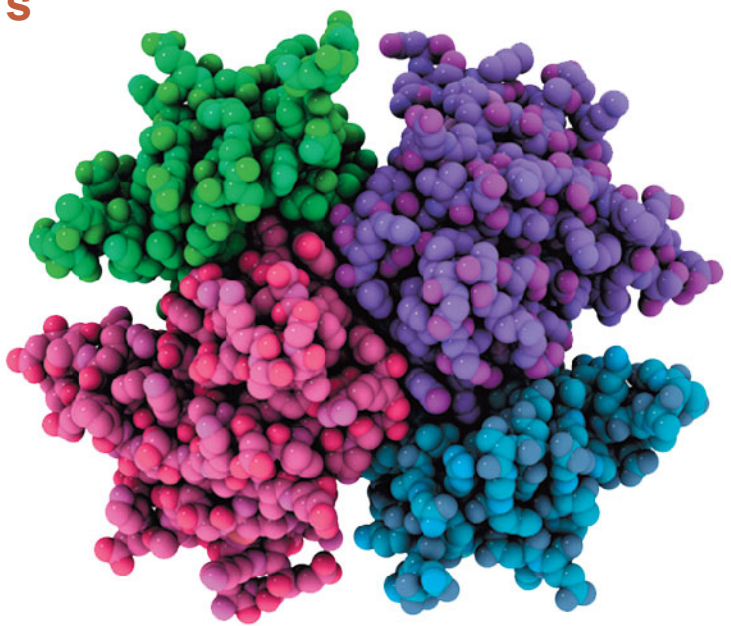
→ Variability helps with variable functions.

Figure 5.21

Technique



Results



Concept 5.5: Nucleic acids ¹store, ²transmit, and ³help express hereditary information

- The amino acid sequence of a polypeptide is programmed by a unit of inheritance called a gene
- Genes consist of DNA, a nucleic acid made of monomers called nucleotides

⊕ each chromosome contains (non-duplicated)

[1] DNA molecule

The Roles of Nucleic Acids

- There are two types of nucleic acids
 - Deoxyribonucleic acid (DNA)
 - Ribonucleic acid (RNA)
- DNA provides directions for its own replication
- DNA directs synthesis of messenger RNA (mRNA) and, through mRNA, controls protein synthesis
- This process is called gene expression

DNA → RNA → Proteins

← which carries most of the Biological functions.

Figure 5.22_1

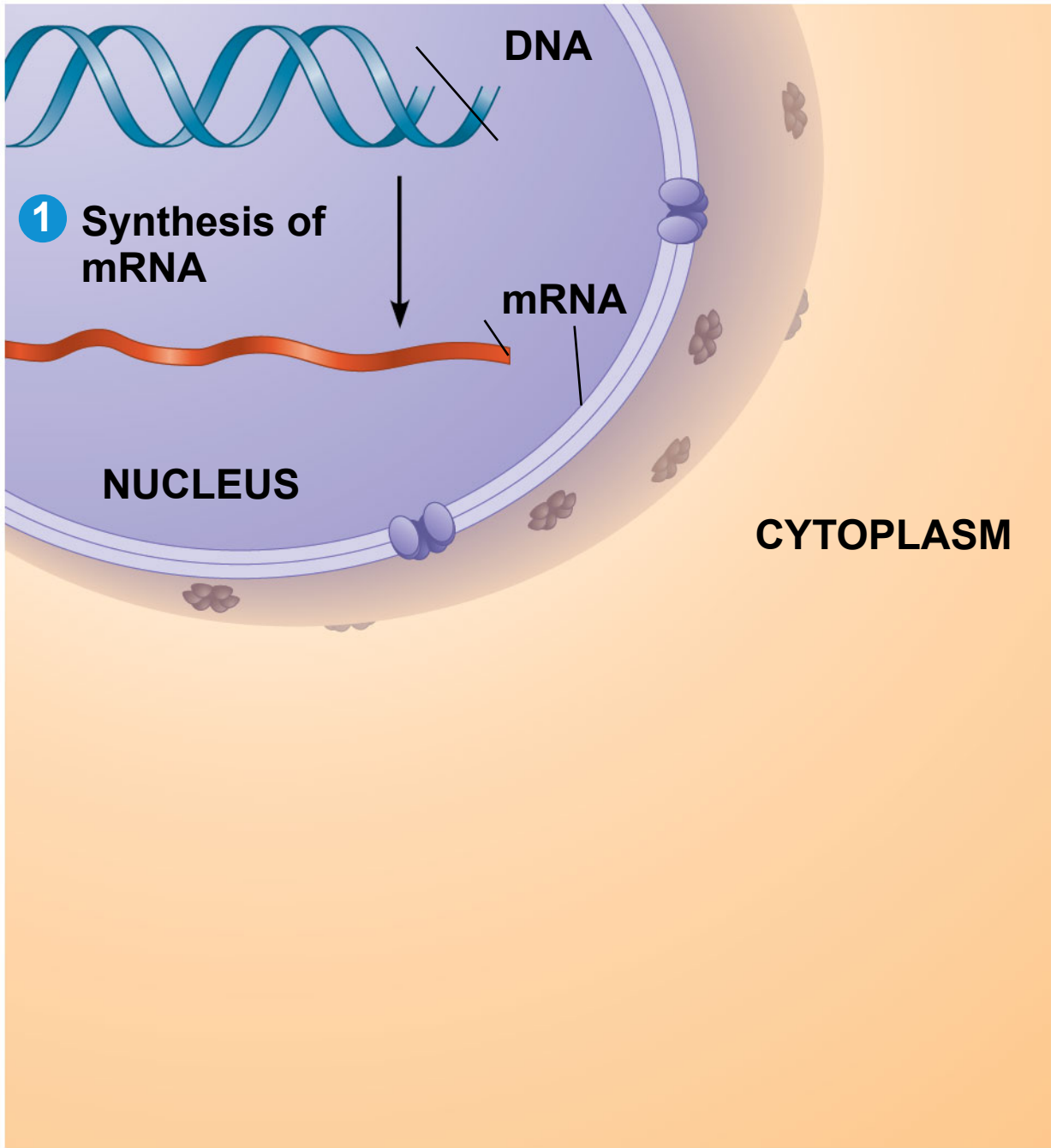


Figure 5.22_2

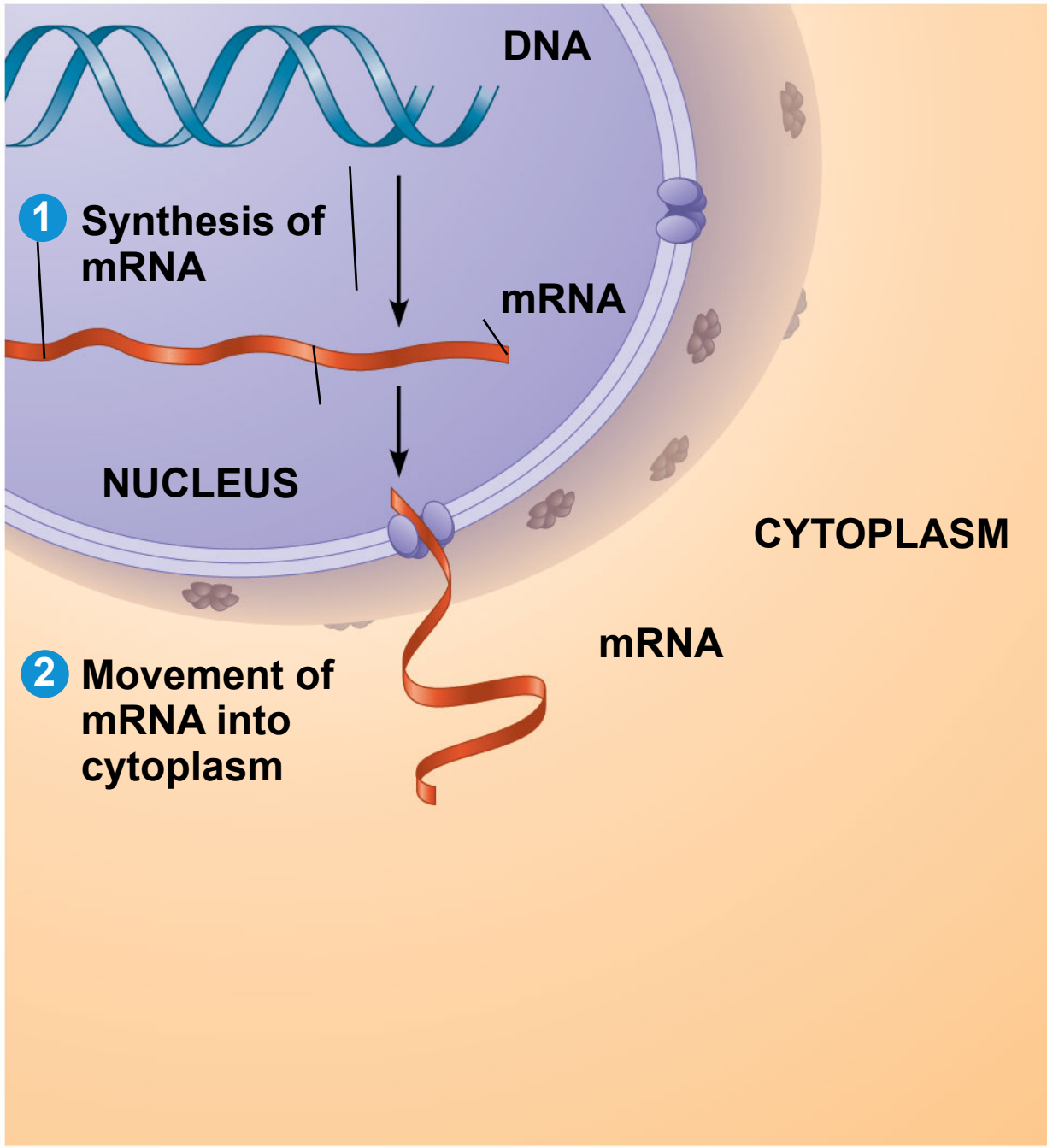
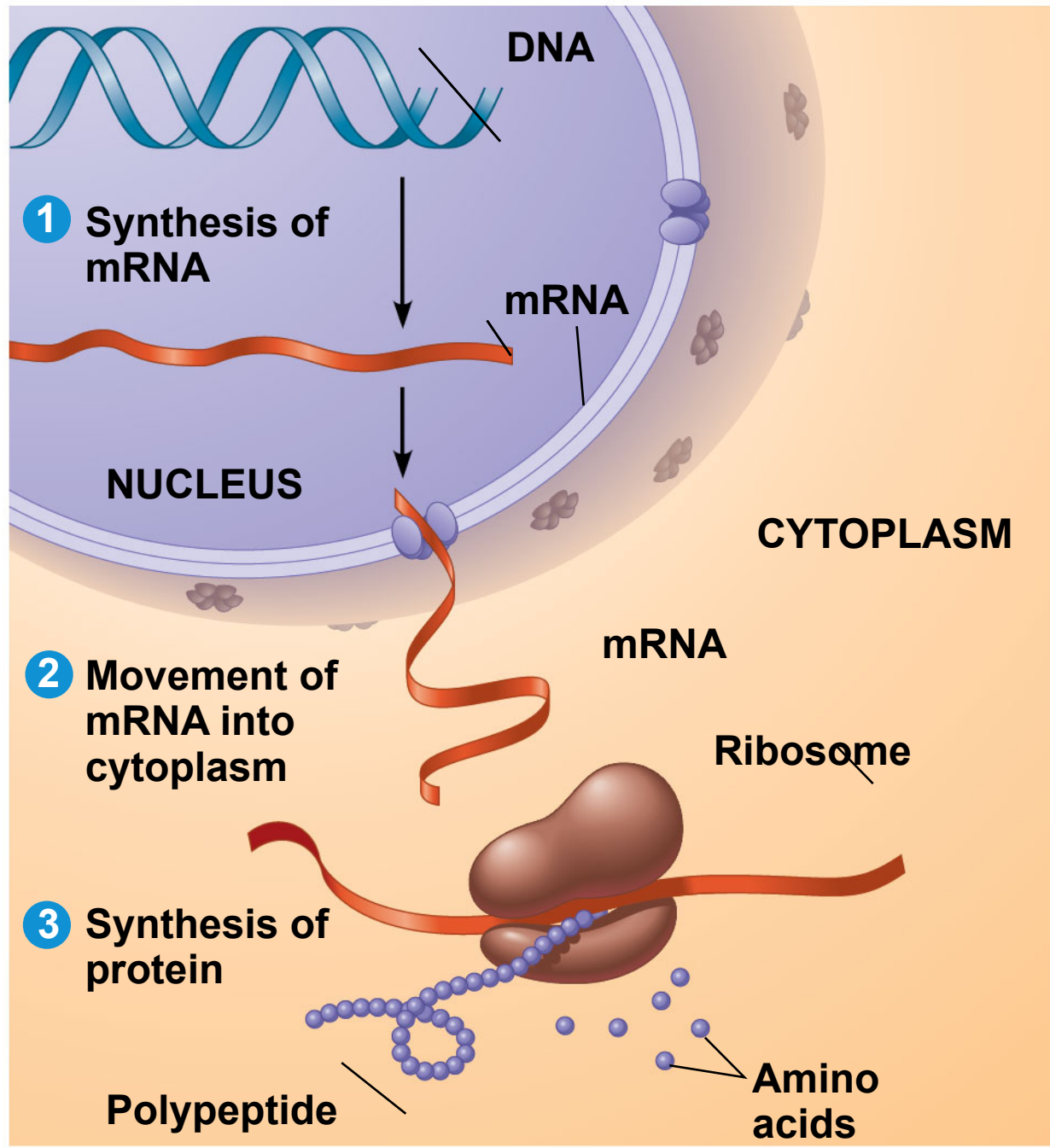


Figure 5.22_3



- Each gene along a DNA molecule directs synthesis of a messenger RNA (mRNA)
- The mRNA molecule interacts with the cell's protein-synthesizing machinery to direct production of a polypeptide
- The flow of genetic information can be summarized as DNA → RNA → protein

The Components of Nucleic Acids

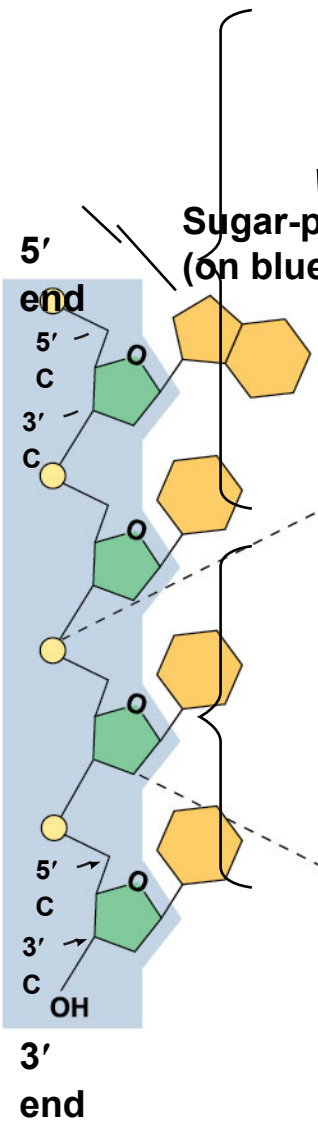
- Nucleic acids are polymers called **polynucleotides**
- Each polynucleotide is made of monomers called **nucleotides**
- Each nucleotide consists of a nitrogenous base, a pentose sugar, and one or more phosphate groups
- The portion of a nucleotide without the phosphate group is called a **nucleoside**

- Nucleoside = nitrogenous base + sugar
- There are two families of nitrogenous bases
 - **Pyrimidines** (cytosine, thymine, and uracil) have a single six-membered ring
 - **Purines** (adenine and guanine) have a six-membered ring fused to a five-membered ring
- In DNA, the sugar is deoxyribose; in RNA, the sugar is ribose

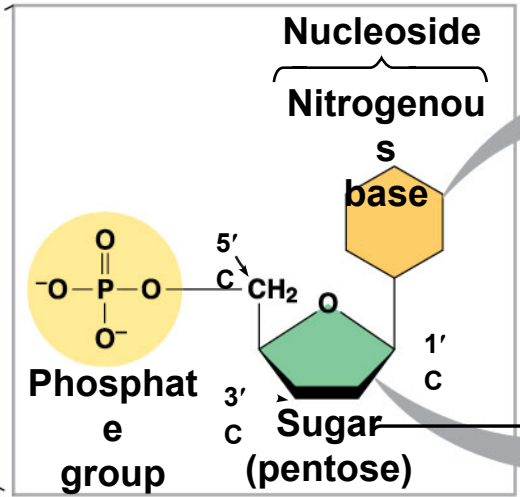
● Nucleotide = nucleoside + phosphate group

← (1, 2, 3) groups.
most common

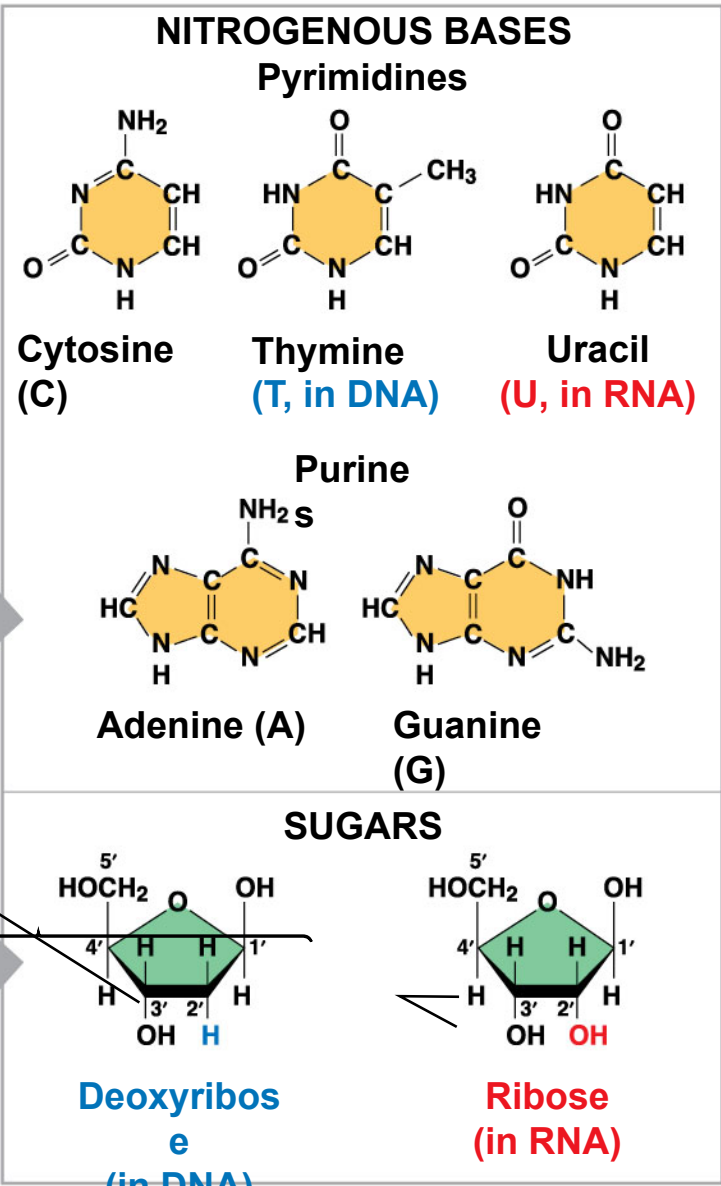
See (SSS)



Sugar-phosphate backbone (on blue background)



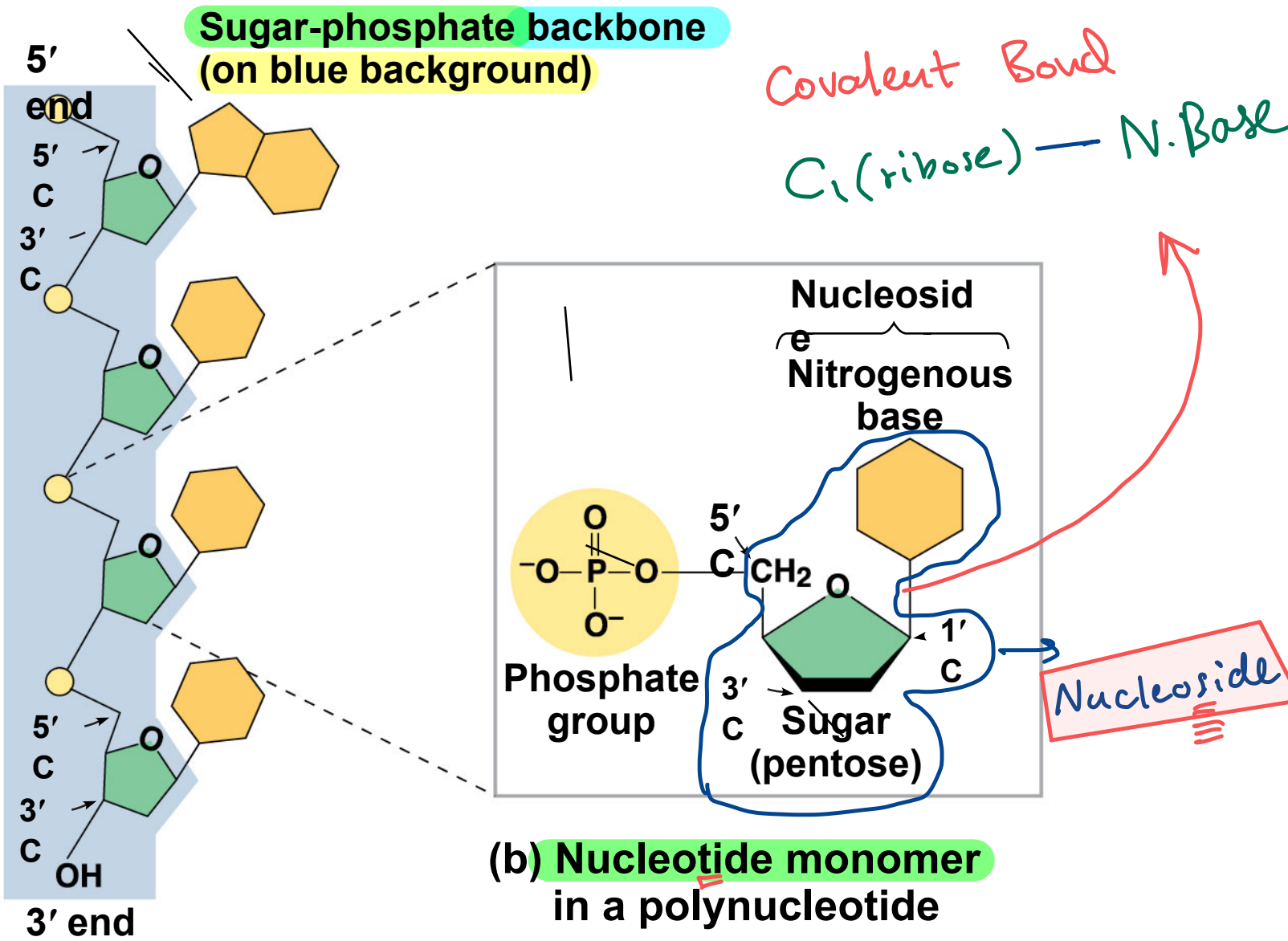
(b) Nucleotide monomer in a polynucleotide



(c) Nucleoside components

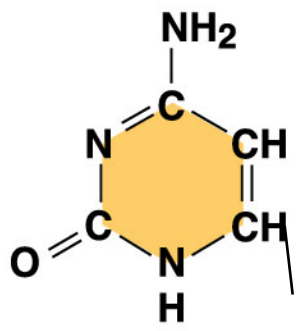
(a) Polynucleotide, or nucleic acid

Figure 5.23a

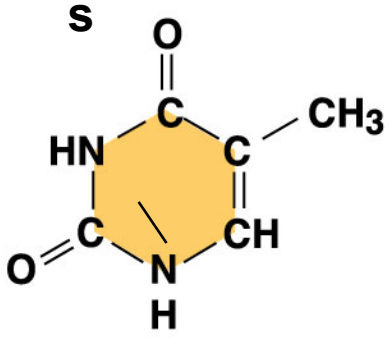


NITROGENOUS BASES

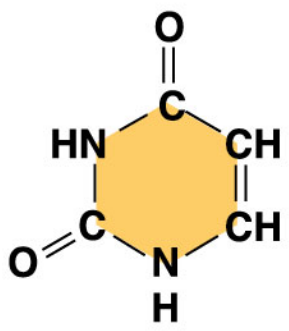
Pyrimidine



Cytosine (C)

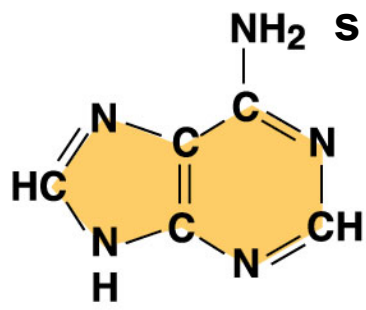


Thymine
(T, in DNA)

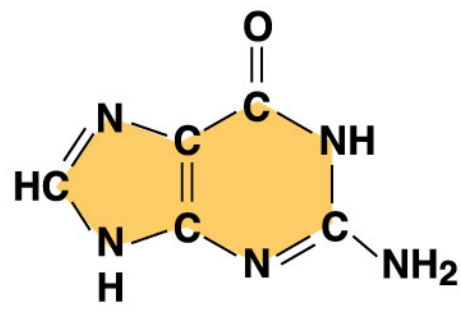


Uracil
(U, in RNA)

Purine



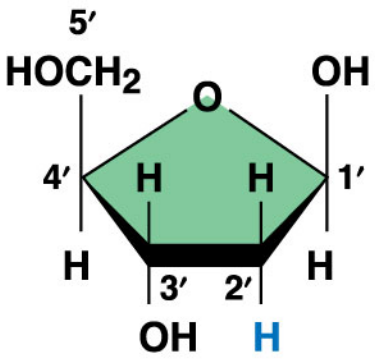
Adenine (A)



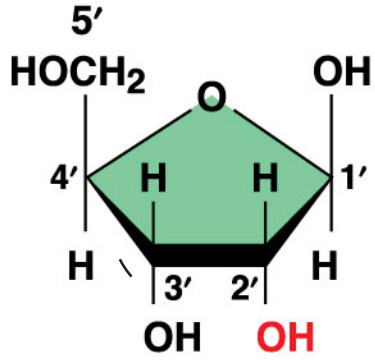
Guanine (G)

(c) Nucleoside components

SUGARS



Deoxyribose
(in DNA)



Ribose
(in RNA)

(c) Nucleoside components

C_{2'} ⇒ determines the type of Ribose.

C_{3'} ⇒ OH links with P-group from another nucleotide

C_{5'} ⇒ OH links with P-group from the same nucleotide.

Nucleotide Polymers

- Nucleotides are linked together by a phosphodiester linkage to build a polynucleotide
- A phosphodiester linkage consists of a phosphate group that links the sugars of two nucleotides
- These links create a backbone of sugar-phosphate units with nitrogenous bases as appendages
- The sequence of bases along a DNA or mRNA polymer is unique for each gene

The Structures of DNA and RNA Molecules

- DNA molecules have two polynucleotides spiraling around an imaginary axis, forming a **double helix**
- The backbones run in opposite $5' \rightarrow 3'$ directions from each other, an arrangement referred to as **antiparallel**
- One DNA molecule includes many genes

-
-

- Only certain bases in DNA pair up and form hydrogen bonds: adenine (A) always with thymine (T), and guanine (G) always with cytosine (C)
- This is called complementary base pairing
- This feature of DNA structure makes it possible to generate two identical copies of each DNA molecule in a cell preparing to divide

- RNA, in contrast to DNA, is single-stranded
- Complementary pairing can also occur between two RNA molecules or between parts of the same molecule ^① _②
- In RNA, thymine is replaced by uracil (U), so A and U pair
- While DNA always exists as a double helix, RNA molecules are more variable in form

Figure 5.24

