

# Protein



▼ **Figure 5.13** An overview of protein functions.

### Enzymatic proteins

**Function:** Selective acceleration of chemical reactions

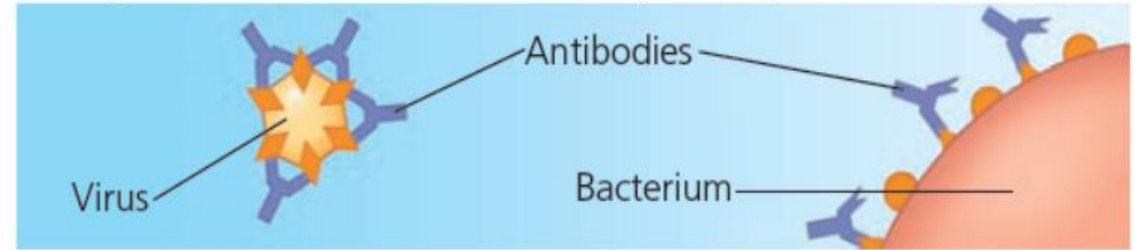
**Example:** Digestive enzymes catalyze the hydrolysis of bonds in food molecules.



### Defensive proteins

**Function:** Protection against disease

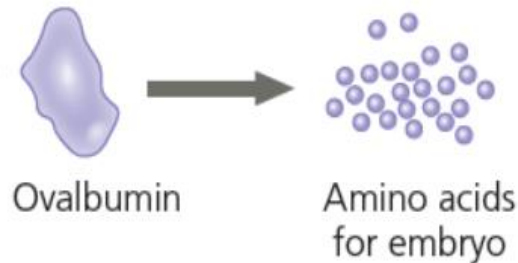
**Example:** Antibodies inactivate and help destroy viruses and bacteria.



### Storage proteins

**Function:** Storage of amino acids

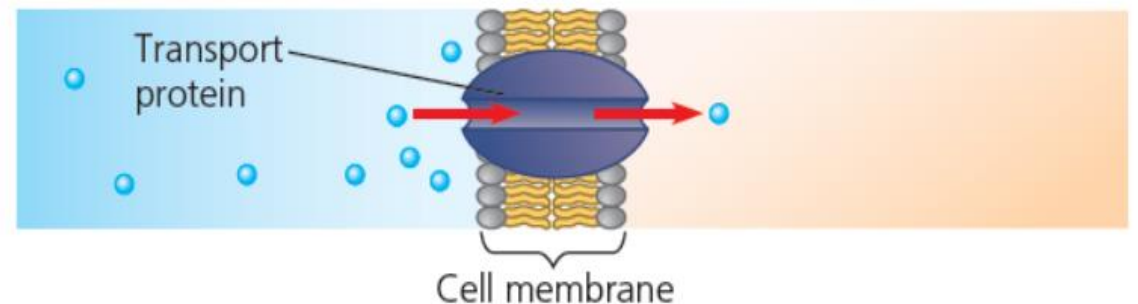
**Examples:** Casein, the protein of milk, is the major source of amino acids for baby mammals. Plants have storage proteins in their seeds. Ovalbumin is the protein of egg white, used as an amino acid source for the developing embryo.



### Transport proteins

**Function:** Transport of substances

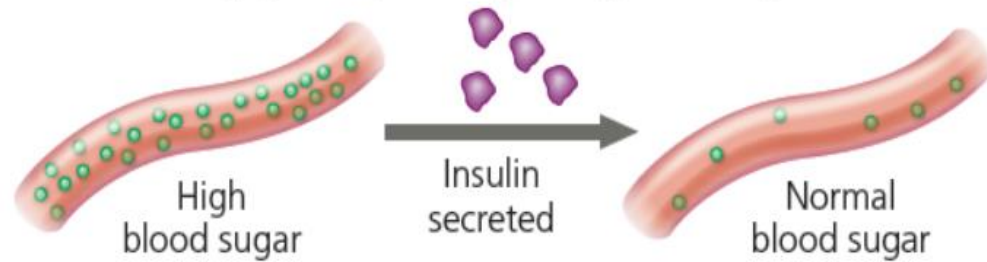
**Examples:** Hemoglobin, the iron-containing protein of vertebrate blood, transports oxygen from the lungs to other parts of the body. Other proteins transport molecules across membranes, as shown here.



## Hormonal proteins

**Function:** Coordination of an organism's activities

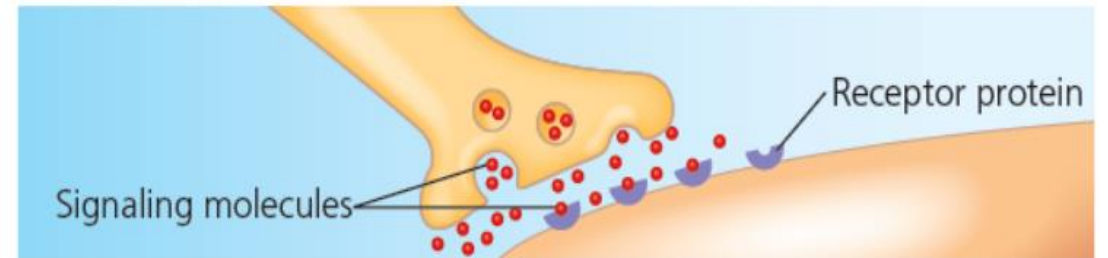
**Example:** Insulin, a hormone secreted by the pancreas, causes other tissues to take up glucose, thus regulating blood sugar concentration.



## Receptor proteins

**Function:** Response of cell to chemical stimuli

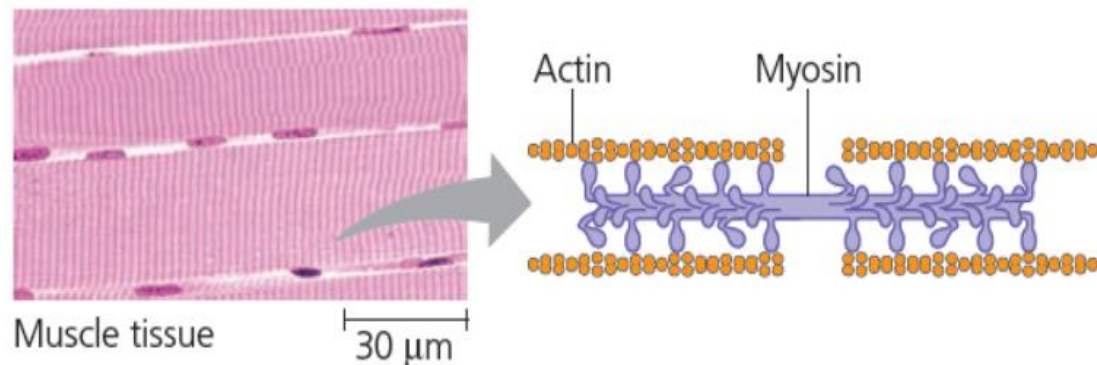
**Example:** Receptors built into the membrane of a nerve cell detect signaling molecules released by other nerve cells.



## 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.

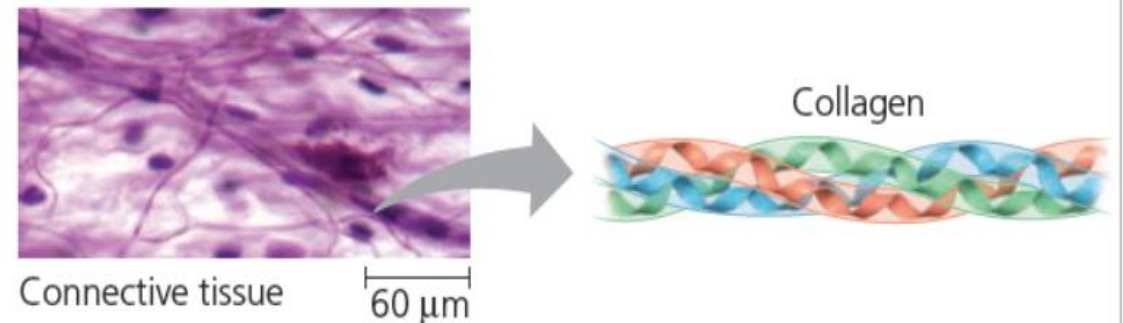


Table 3.2 The Many Functions of Proteins

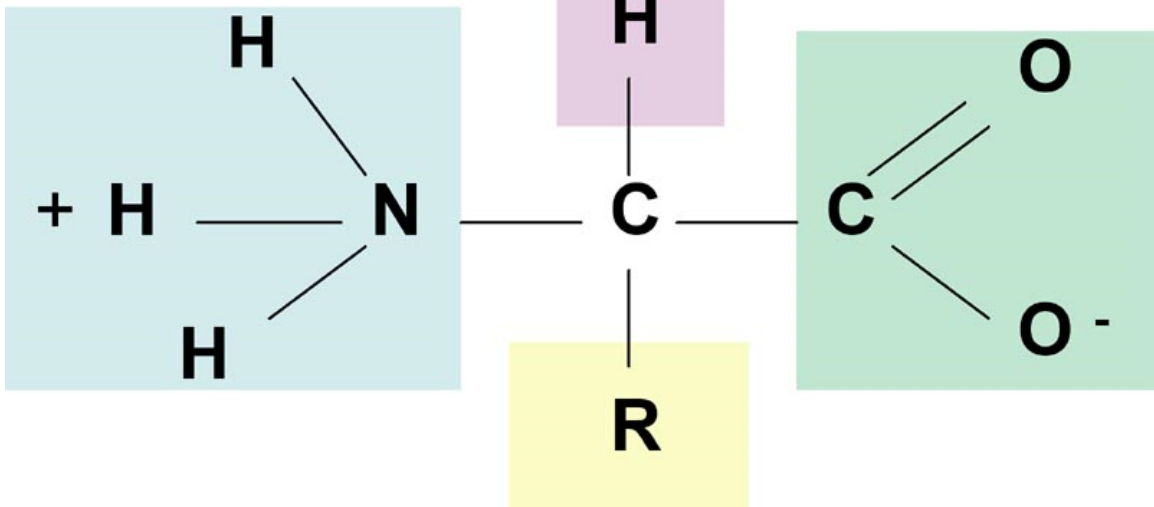
Function	Class of Protein	Examples	Use
Metabolism (Catalysis)	Enzymes	Hydrolytic enzymes Proteases Polymerases Kinases	Cleave polysaccharides Break down proteins Produce nucleic acids Phosphorylate sugars and proteins
Defense	Immunoglobulins	Antibodies	Mark foreign proteins for elimination
Cell recognition	Toxins	Snake venom	Block nerve function
Transport throughout body	Cell surface antigens	MHC proteins	"Self" recognition
	Globins	Hemoglobin Myoglobin	Carries O <sub>2</sub> and CO <sub>2</sub> in blood Carries O <sub>2</sub> and CO <sub>2</sub> in muscle
		Cytochromes	Electron transport
Membrane transport	Transporters	Sodium-potassium pump Proton pump Anion channels	Excitable membranes Chemiosmosis Transport Cl <sup>-</sup> ions
Structure/Support	Fibers	Collagen Keratin Fibrin	Cartilage Hair, nails Blood clot
Motion	Muscle	Actin Myosin	Contraction of muscle fibers Contraction of muscle fibers
Osmotic regulation	Albumin	Serum albumin	Maintains osmotic concentration of blood
Regulation of gene action	Repressors	lac repressor	Regulates transcription
Regulation of body functions	Hormones	Insulin Vasopressin Oxytocin	Controls blood glucose levels Increases water retention by kidneys Regulates uterine contractions and milk production
Storage	Ion binding	Ferritin Casein Calmodulin	Stores iron, especially in spleen Stores ions in milk Binds calcium ions

# Amino Acid Structure

Hydrogen

Amino

Carboxyl

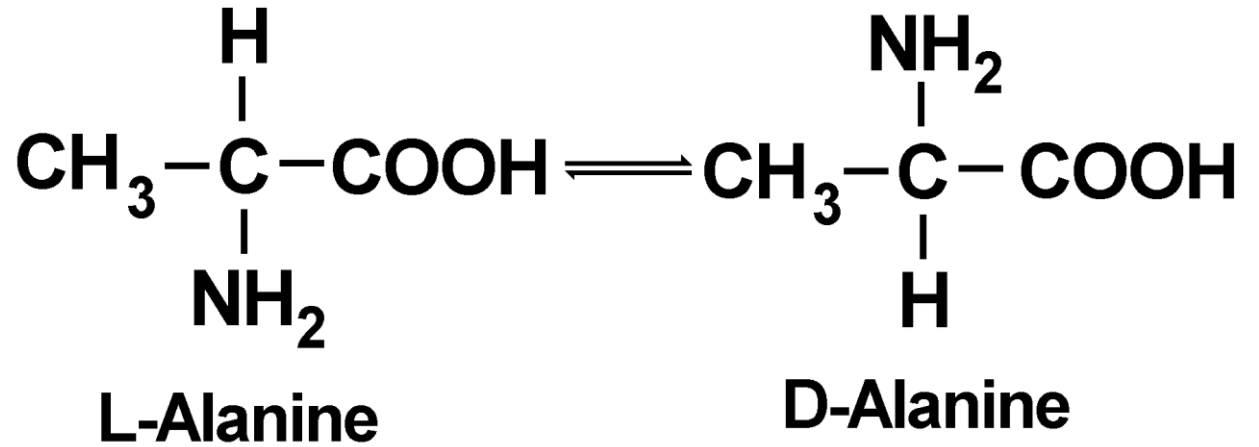
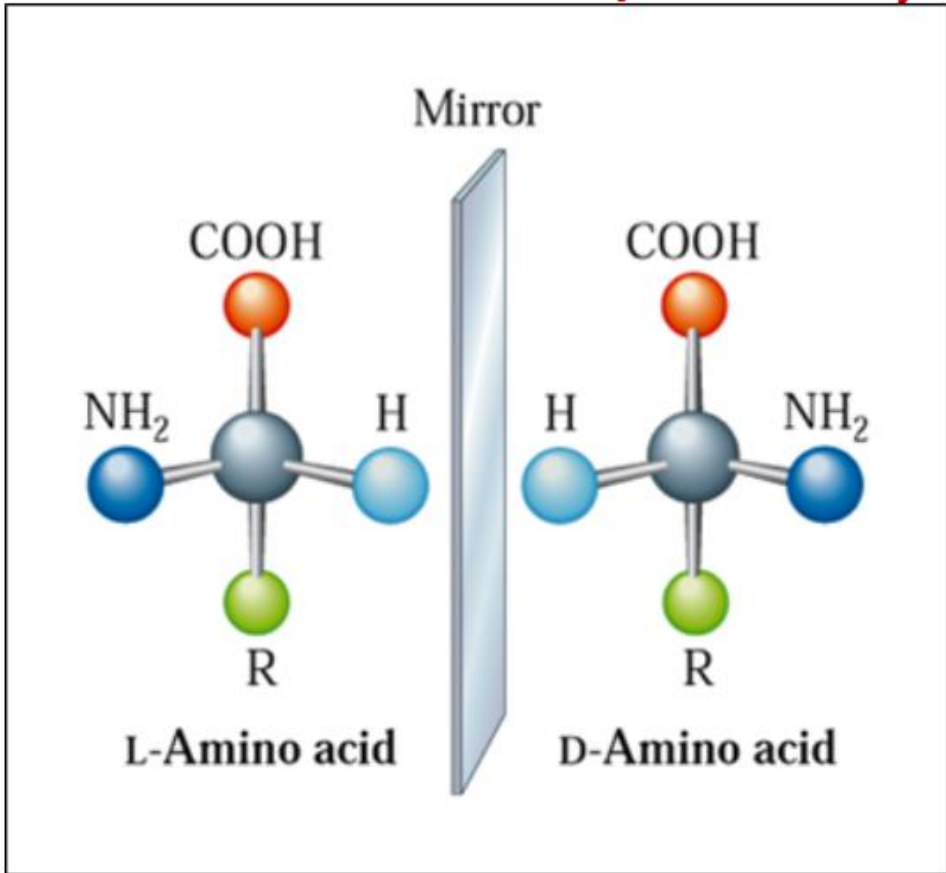


R-group  
(variant)

At the center of the amino acid is an asymmetric carbon atom called the alpha ( $\alpha$ ) carbon.

- Its four different partners are an **amino group**, a **carboxyl group**, a hydrogen atom, and a **variable group**

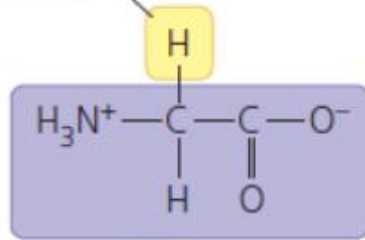
# Handedness/Chirality of Amino Acids



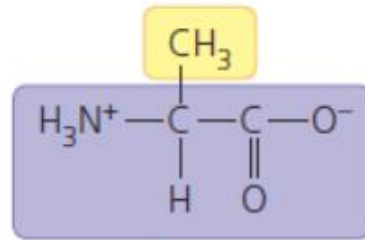
▼ **Figure 5.14** The 20 amino acids of proteins. The amino acids are grouped here according to the properties of their side chains (R groups) and shown in their prevailing ionic forms at pH 7.2, the pH within a cell. The three-letter and one-letter abbreviations for the amino acids are in parentheses. All of the amino acids used in proteins are L enantiomers (see Figure 4.7c).

### Nonpolar side chains; hydrophobic

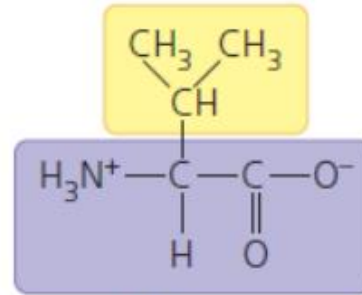
Side chain  
(R group)



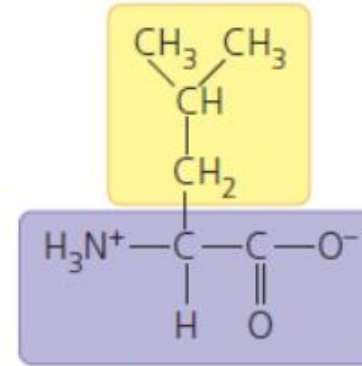
Glycine  
(Gly or G)



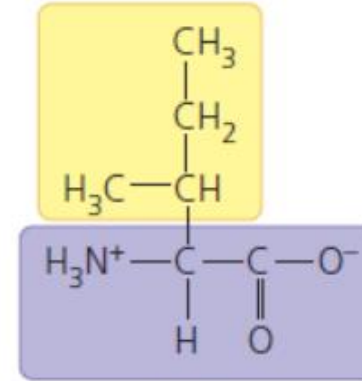
Alanine  
(Ala or A)



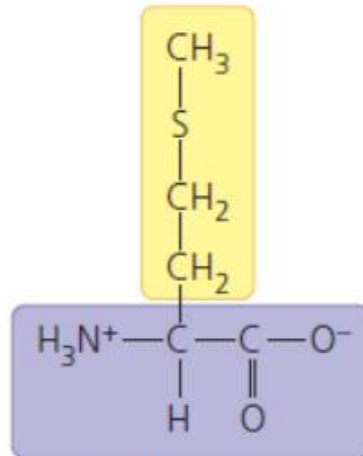
Valine  
(Val or V)



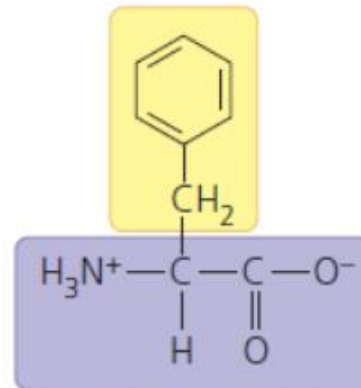
Leucine  
(Leu or L)



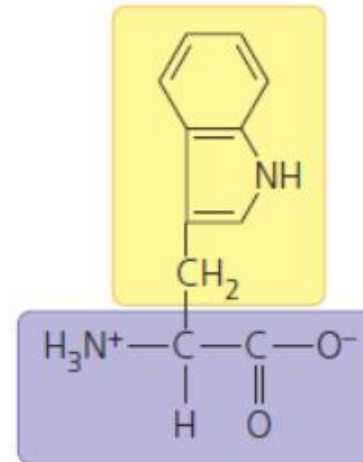
Isoleucine  
(Ile or I)



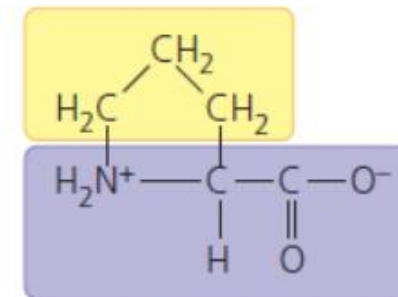
Methionine  
(Met or M)



Phenylalanine  
(Phe or F)



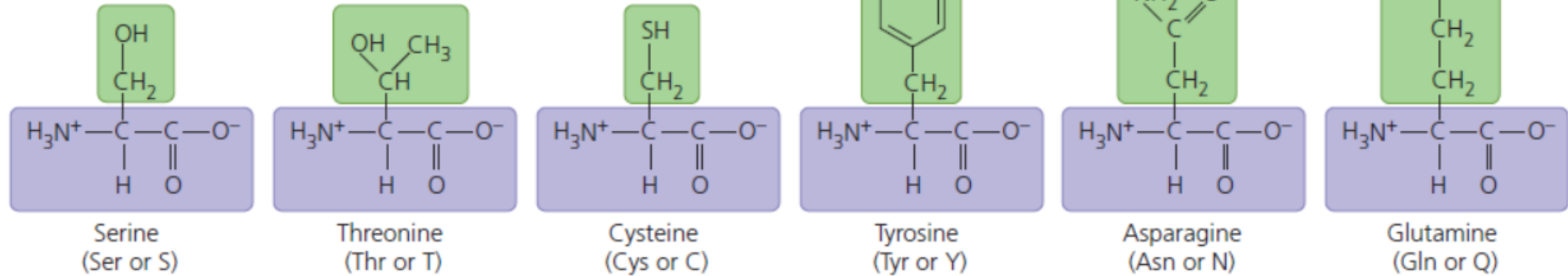
Tryptophan  
(Trp or W)



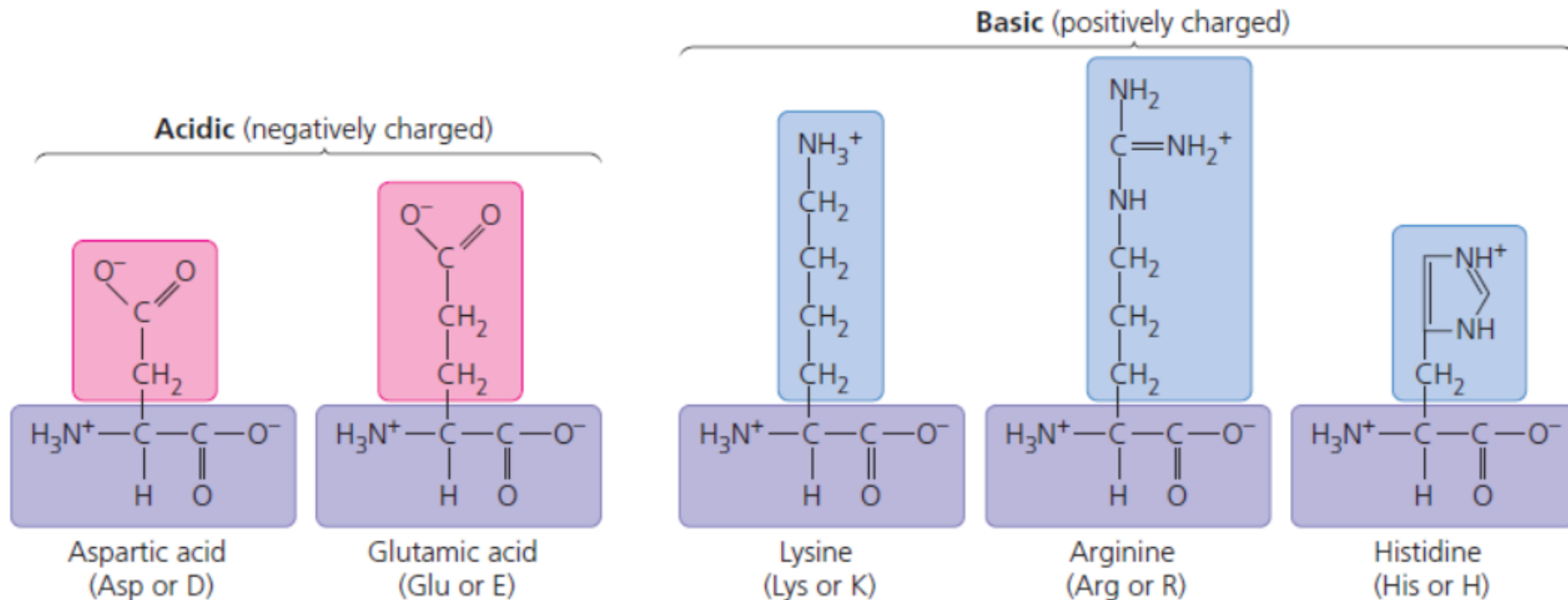
Proline  
(Pro or P)

## Polar side chains; hydrophilic

Since cysteine is only weakly polar, it is sometimes classified as a nonpolar amino acid.



## Electrically charged side chains; hydrophilic



**TABLE 18–1****Nonessential and Essential Amino Acids  
for Humans and the Albino Rat**

<b>Nonessential</b>	<b>Conditionally essential*</b>	<b>Essential</b>
<b>Alanine</b>	<b>Arginine</b>	<b>Histidine</b>
<b>Asparagine</b>	<b>Cysteine</b>	<b>Isoleucine</b>
<b>Aspartate</b>	<b>Glutamine</b>	<b>Leucine</b>
<b>Glutamate</b>	<b>Glycine</b>	<b>Lysine</b>
<b>Serine</b>	<b>Proline</b>	<b>Methionine</b>
	<b>Tyrosine</b>	<b>Phenylalanine</b>
		<b>Threonine</b>
		<b>Tryptophan</b>
		<b>Valine</b>

**\*Required to some degree in young, growing animals, and/or sometimes during illness.**

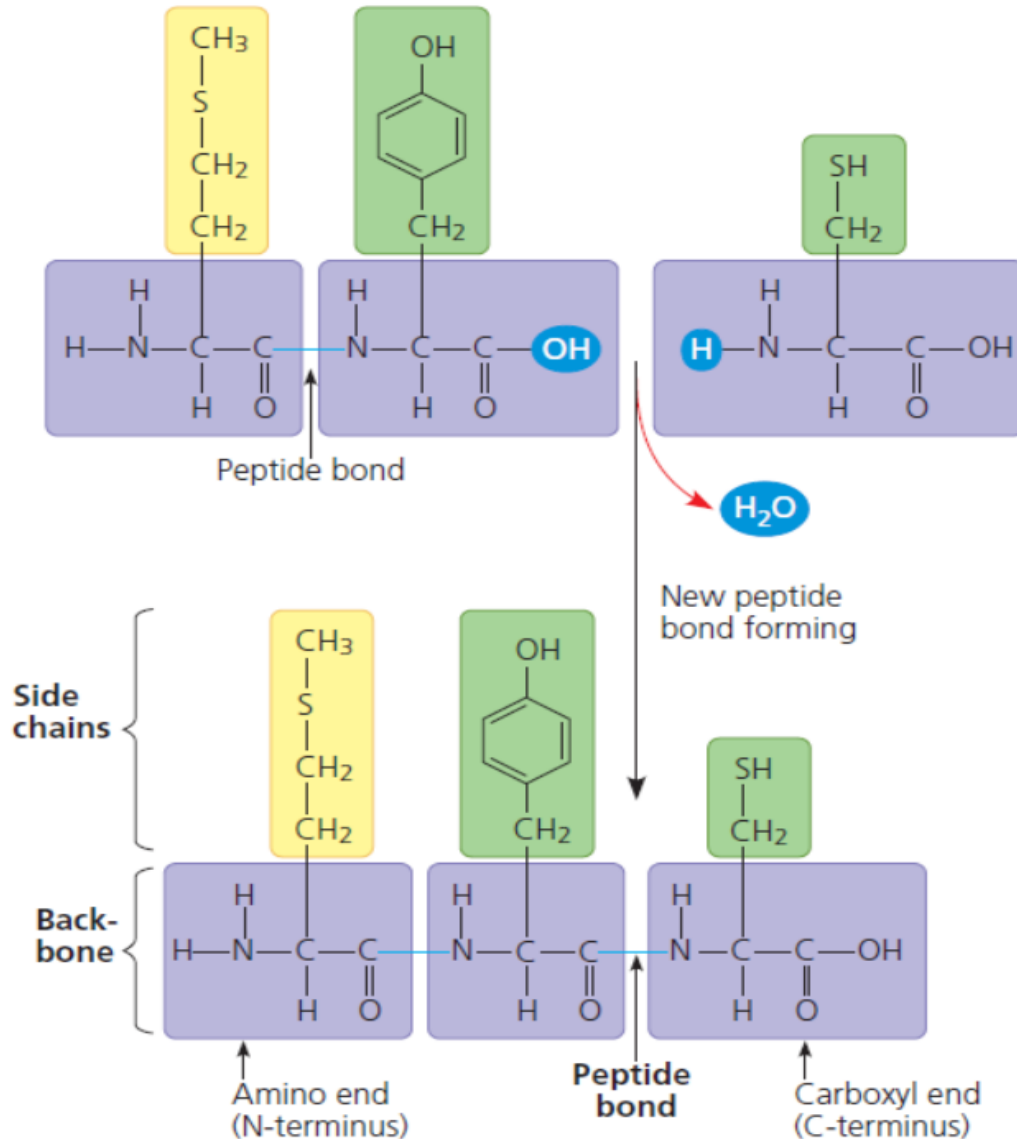
**Table 18-1**

*Lehninger Principles of Biochemistry, Fifth Edition*

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# Polypeptides (Amino Acid Polymers)

## Making a Polypeptide Chain



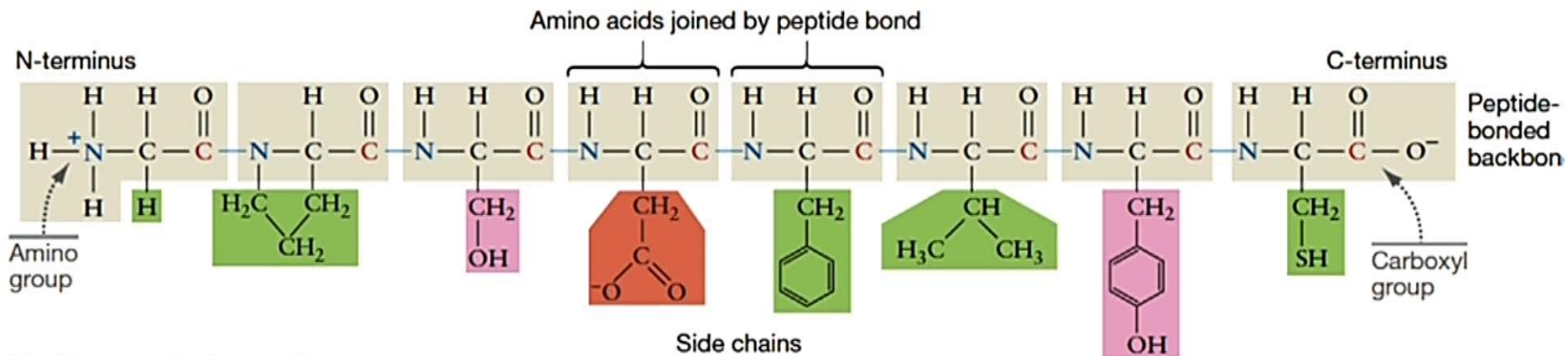
Note: dehydration synthesis.

Note: carboxyl group of one end attaches to amino group of another.

Note: peptide bond is formed.

Note: repeating this process builds a polypeptide.

(a) Chain of amino acid residues



(b) Residue numbering system

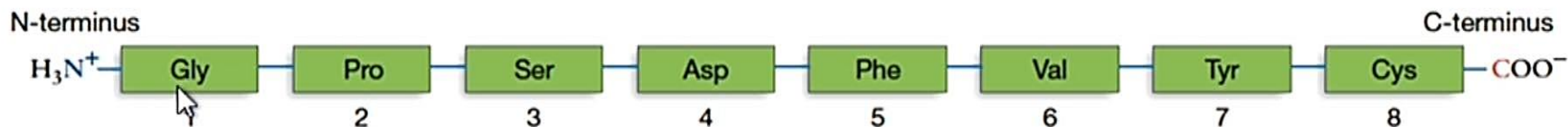
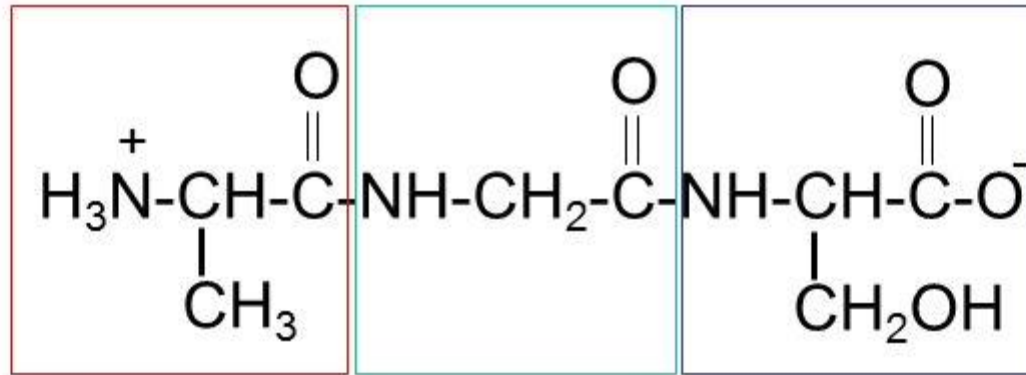


Figure 3.6 Amino Acids Polymerize to Form Chains.

# Naming of peptides

- Begin from the N terminal.
- Drop “-ine” and it is replaced by “-yl”.
- Give the full name of amino acid at the C terminal.

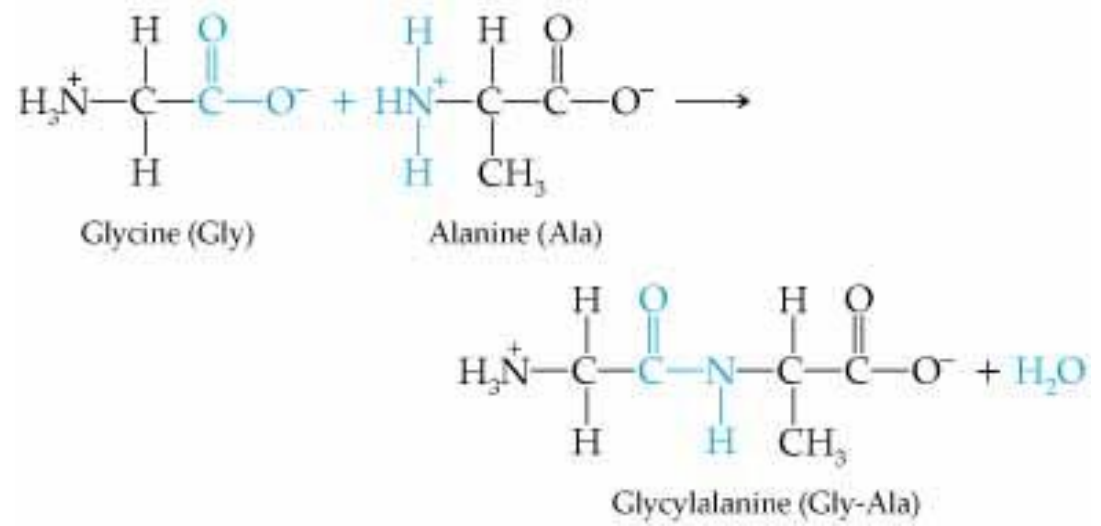


From alanine  
alanyl

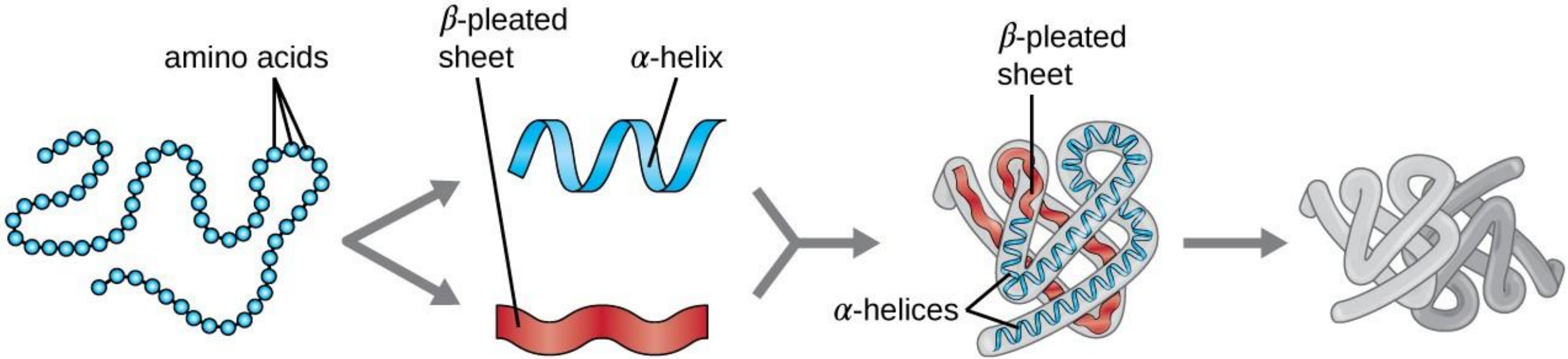
From glycine  
glycyl

From serine  
serine

Alanylglycylserine  
(Ala-Gly-Ser)



# Levels of Protein Structure



## Primary Protein Structure

Sequence of a chain of amino acids

## Secondary Protein Structure

Local folding of the polypeptide chain into helices or sheets

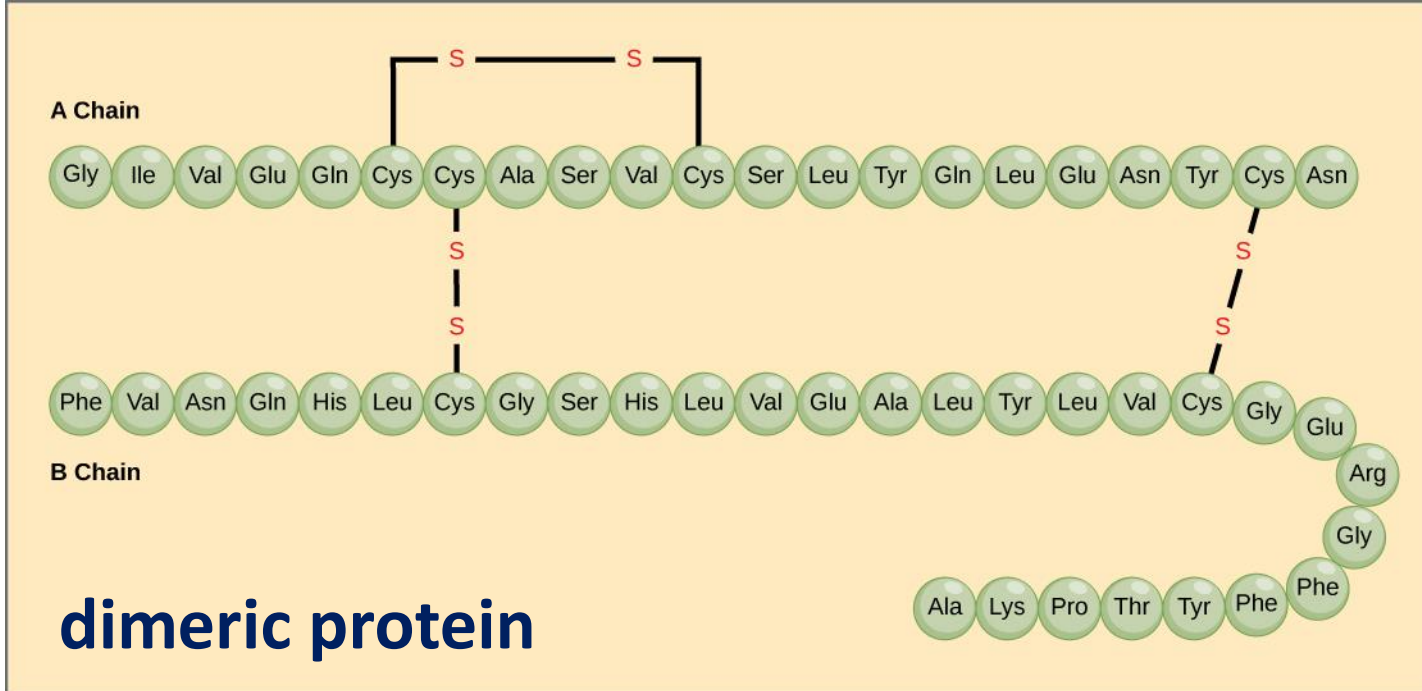
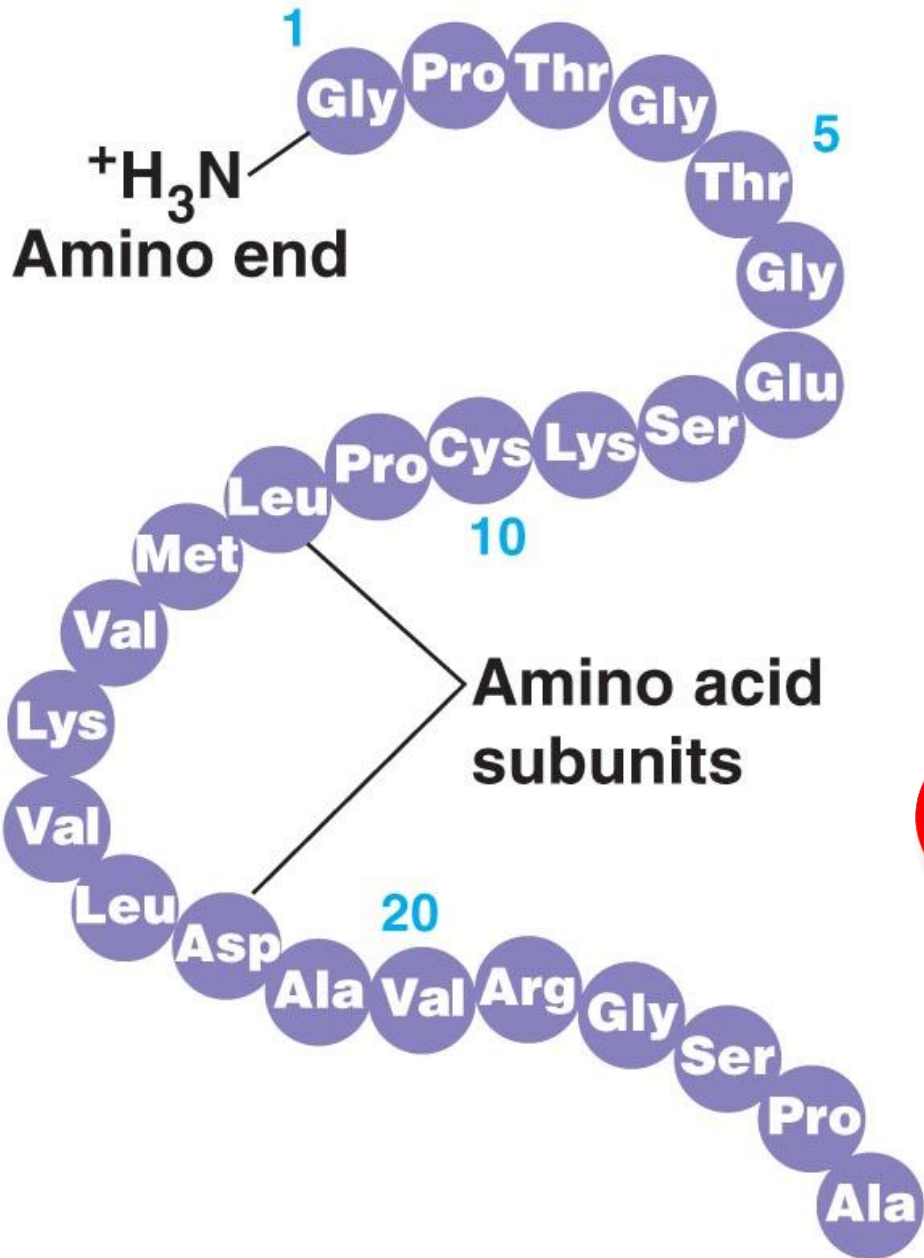
## Tertiary Protein Structure

three-dimensional folding pattern of a protein due to side chain interactions

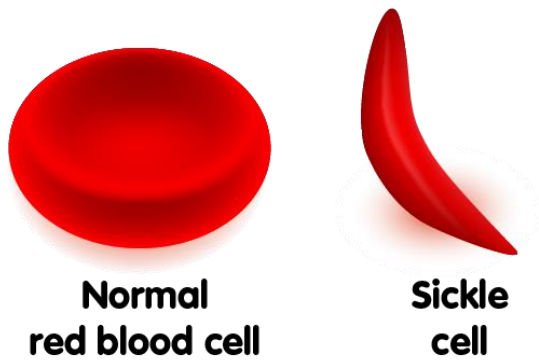
## Quaternary Protein Structure

protein consisting of more than one amino acid chain

# Primary Structure

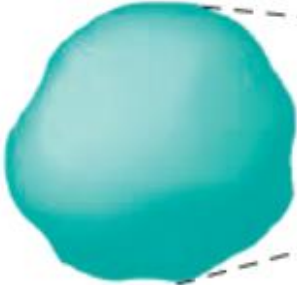
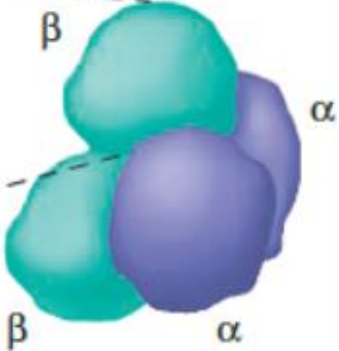
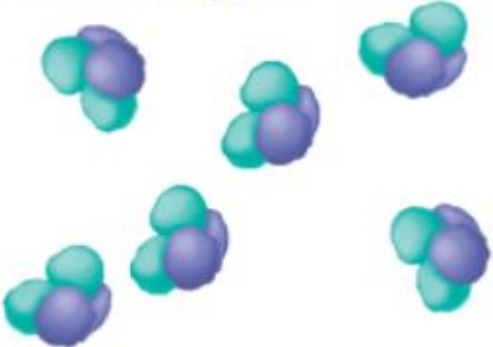
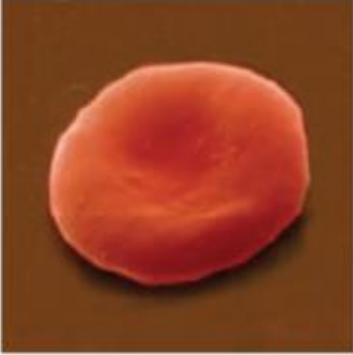
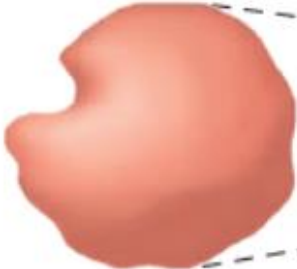
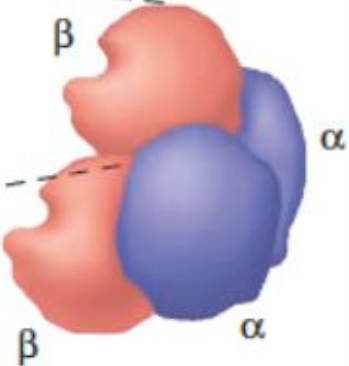
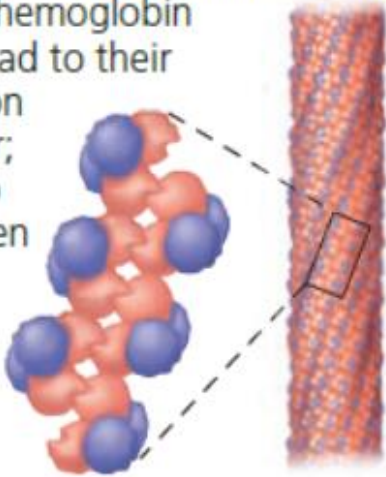



## Effect of Amino Acid Change—Sickle Cell Anemia



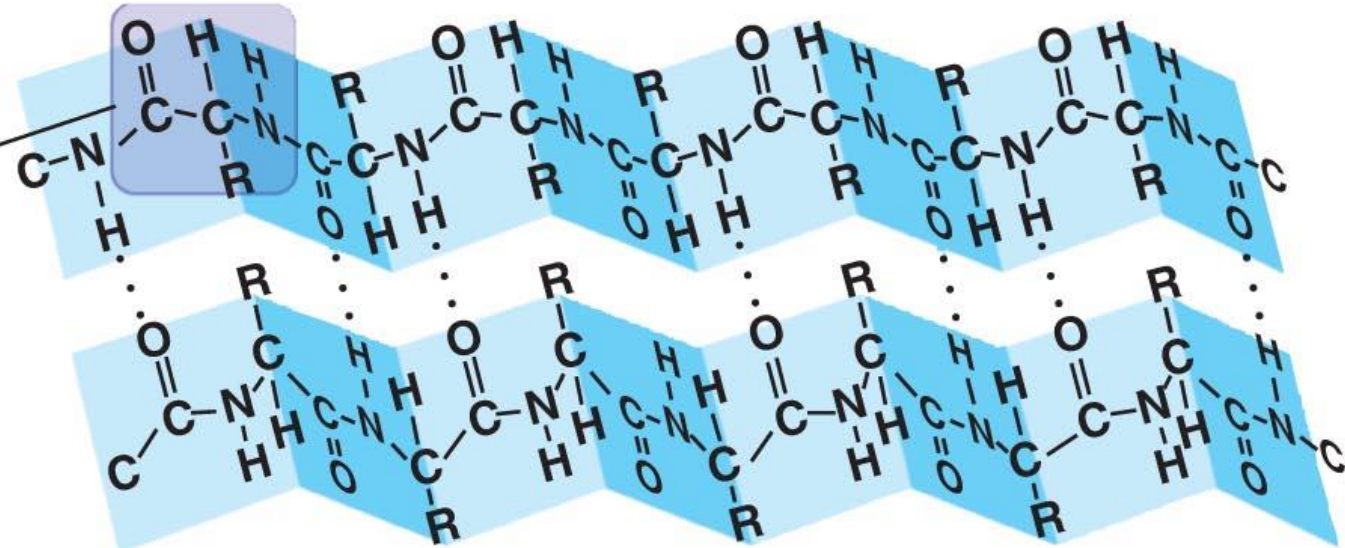
Normal hemoglobin $\beta$ chain						
Valine	Histidine	Leucine	Threonine	Proline	Glutamic acid	Glutamic acid
Sickle cell anemia hemoglobin $\beta$ chain						
Valine	Histidine	Leucine	Threonine	Proline	Valine	Glutamic acid



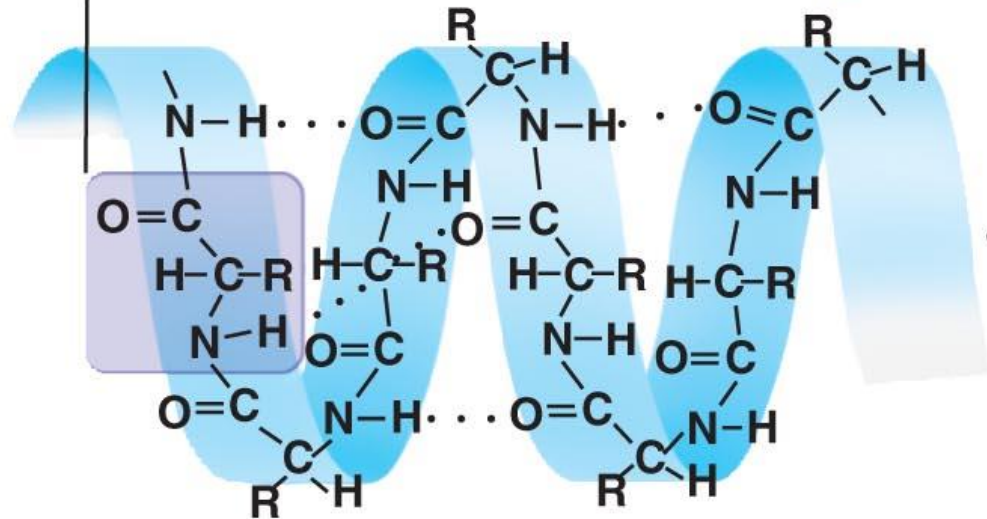
	Primary Structure	Secondary and Tertiary Structures	Quaternary Structure	Function	Red Blood Cell Shape
Normal hemoglobin	1 Val 2 His 3 Leu 4 Thr 5 Pro 6 Glu 7 Glu	Normal $\beta$ subunit 	Normal hemoglobin 	Normal hemoglobin proteins do not associate with one another; each carries oxygen. 	Normal red blood cells are full of individual hemoglobin proteins.  5 $\mu\text{m}$
Sickle-cell hemoglobin	1 Val 2 His 3 Leu 4 Thr 5 Pro 6 Val 7 Glu	Sickle-cell $\beta$ subunit 	Sickle-cell hemoglobin 	Hydrophobic interactions between sickle-cell hemoglobin proteins lead to their aggregation into a fiber; capacity to carry oxygen is greatly reduced. 	Fibers of abnormal hemoglobin deform red blood cell into sickle shape.  5 $\mu\text{m}$

# Secondary Structure

$\beta$  pleated sheet



Examples of amino acid subunits



$\alpha$  helix



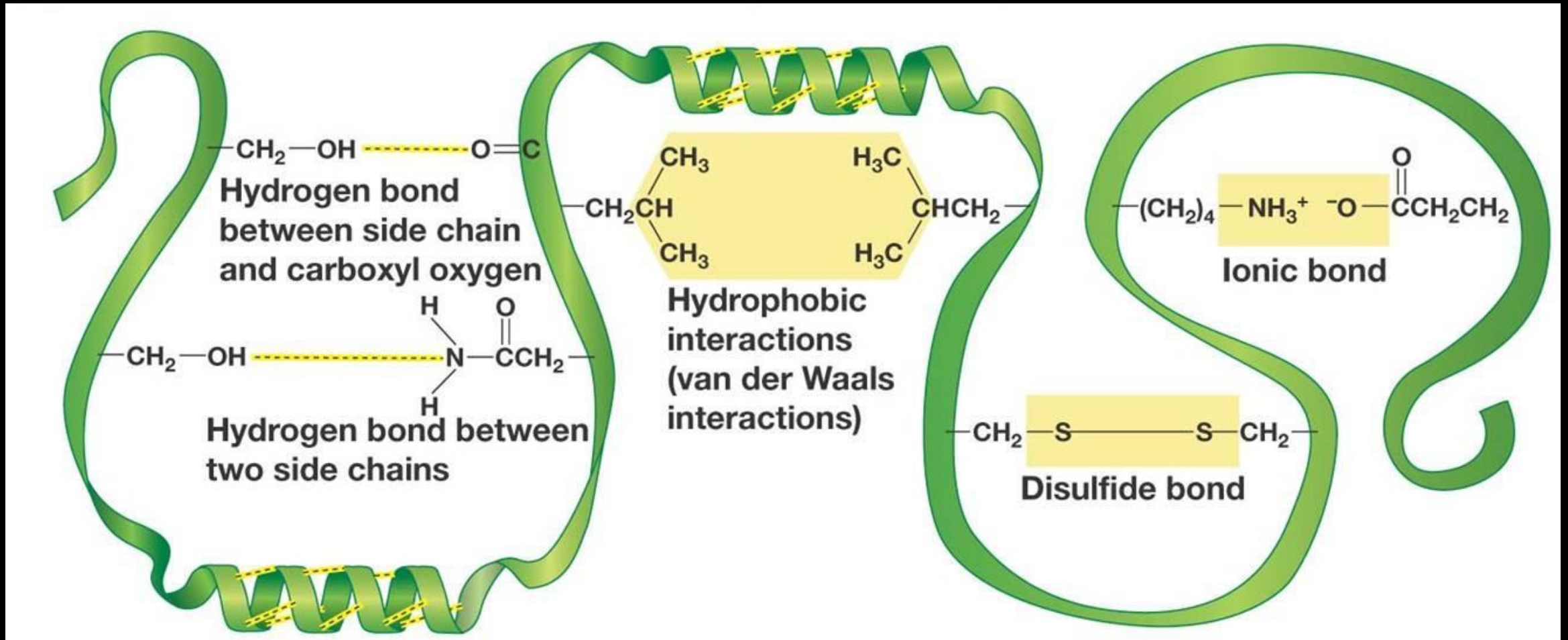


a.  
**FIGURE 3.18** Fibrous proteins.

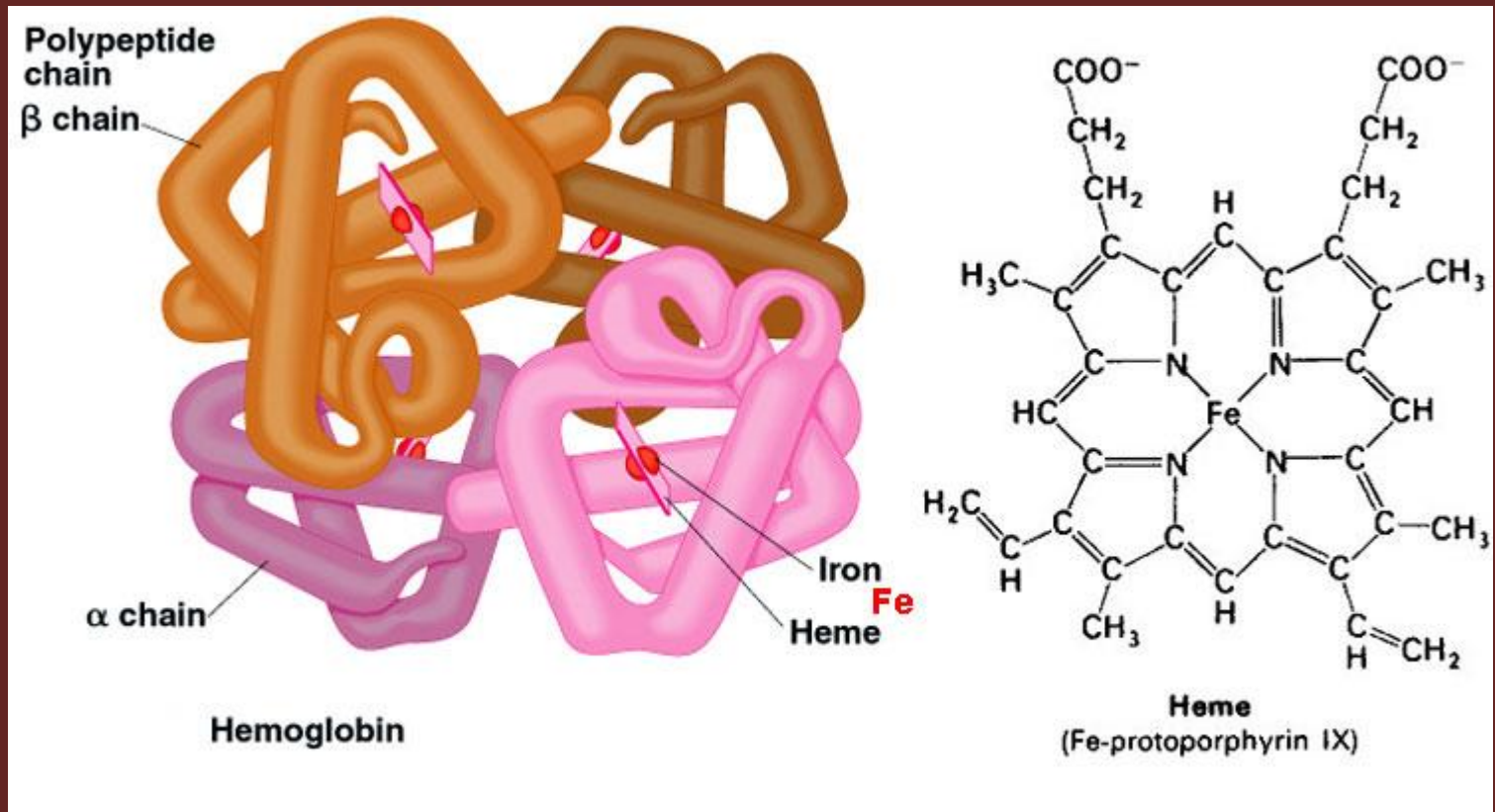
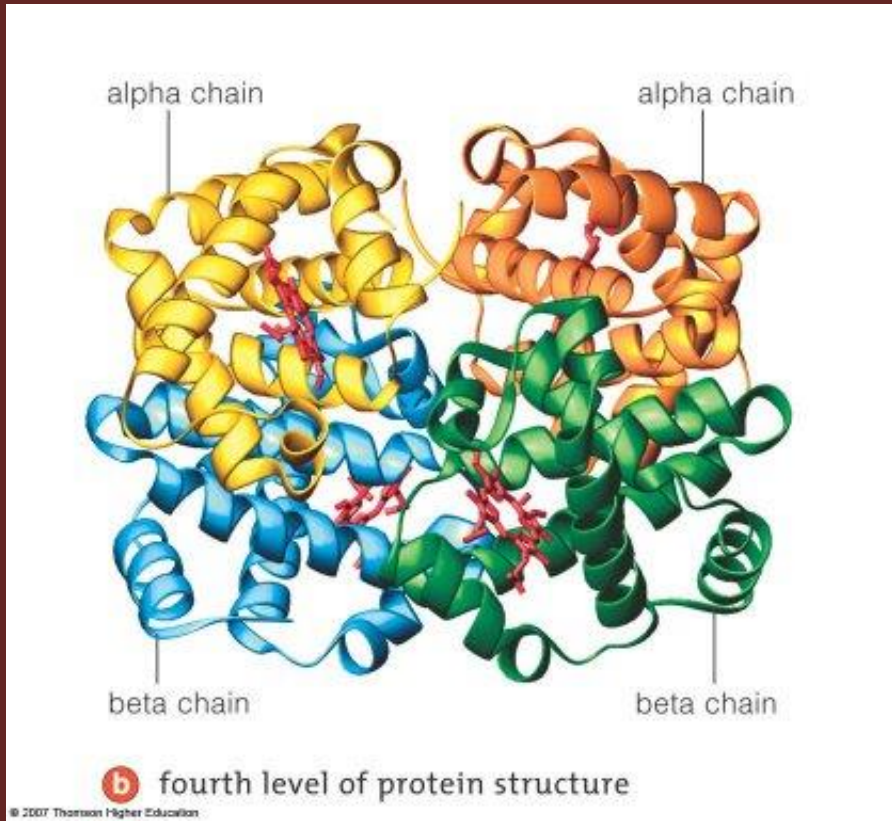
b.

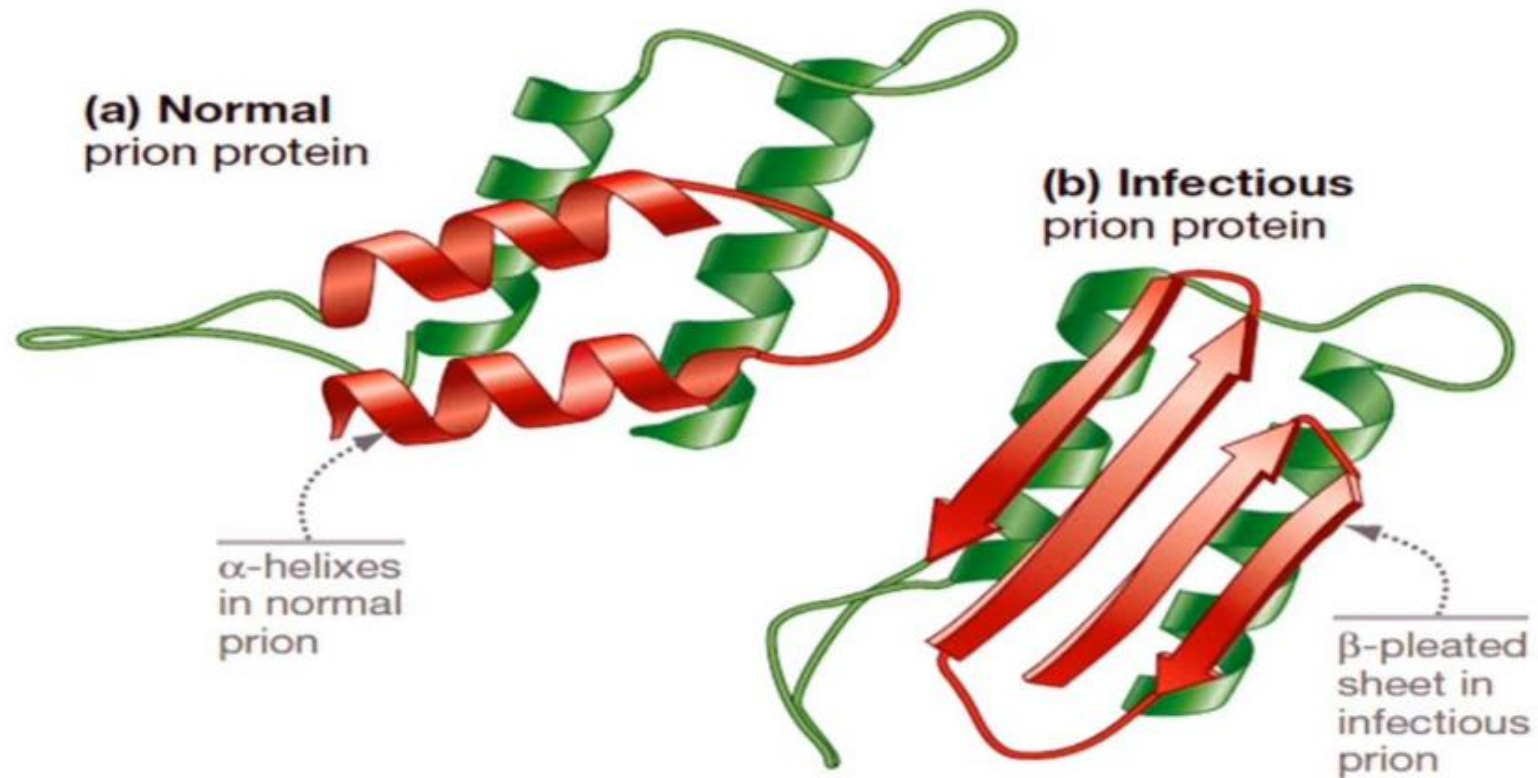


# Tertiary structure



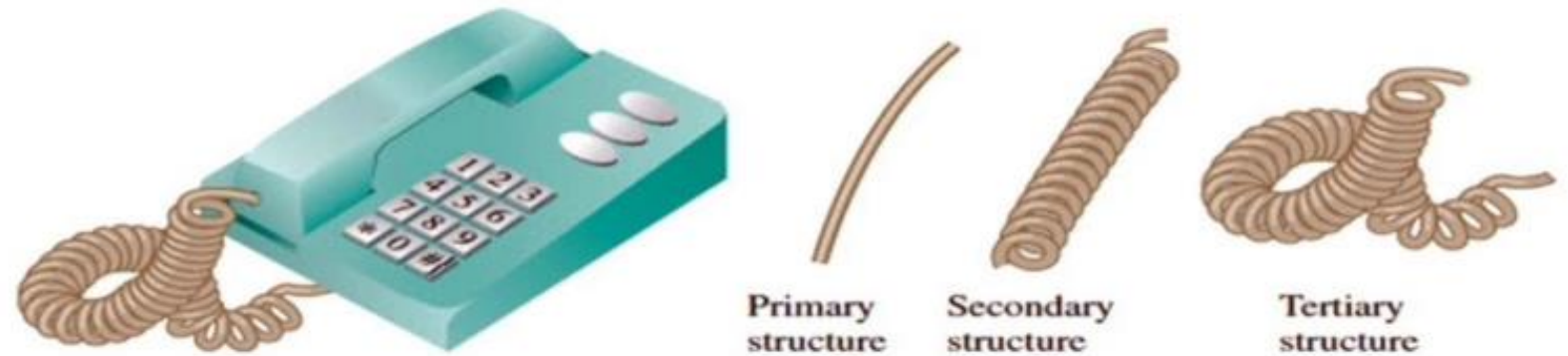
# quaternary structure





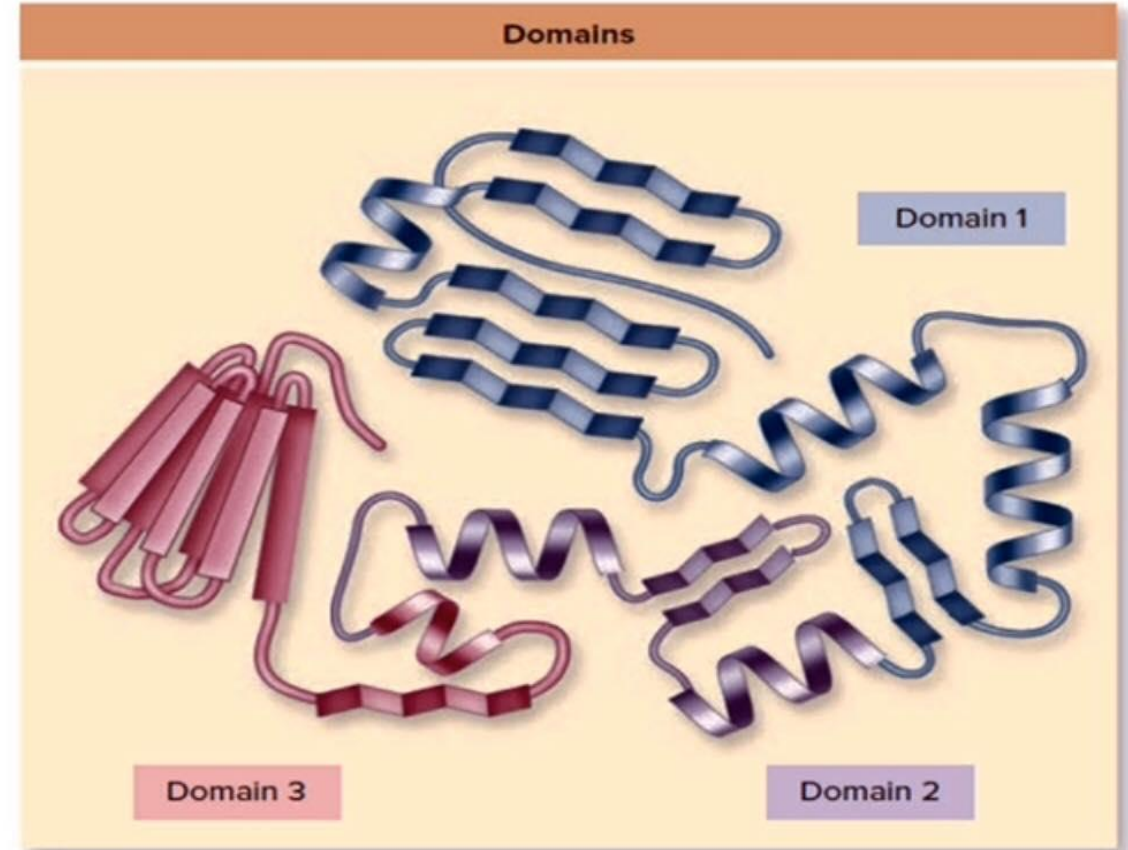
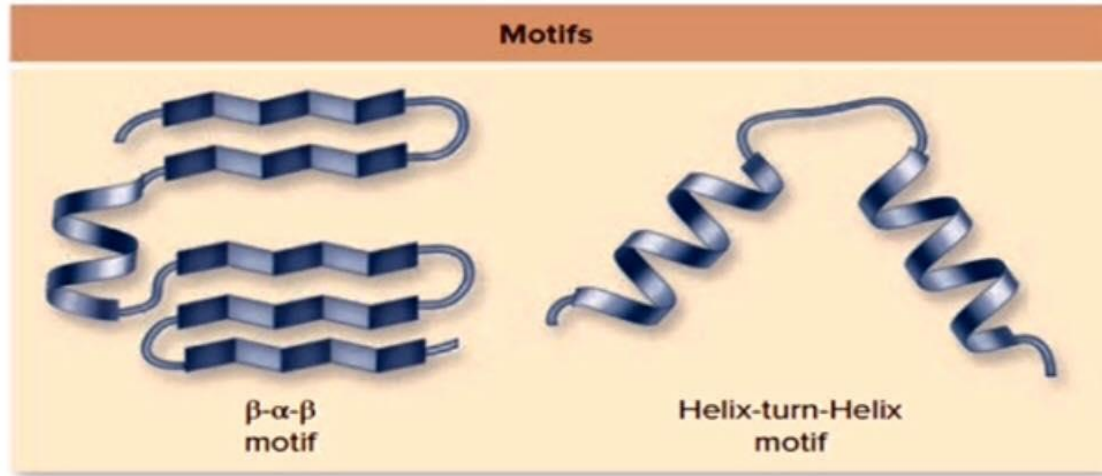
**Figure 3.14 Prion Infectivity Is Linked to Structure.** Ribbon diagram of (a) a normal, noninfectious prion protein with  $\alpha$ -helices; and (b) the infectious form with  $\beta$ -pleated sheets that causes mad cow disease in cattle.

**Figure 20.9** A telephone cord has three levels of structure. These structural levels are a good analogy for the first three levels of protein structure.



**TABLE 16.6** Summary of Structural Levels in Proteins

Structural Level	Characteristics
Primary	Peptide bonds join amino acids in a specific sequence in a polypeptide.
Secondary	The coiled $\alpha$ helix, $\beta$ -pleated sheet, or a triple helix forms by hydrogen bonding between peptide bonds along the chain.
Tertiary	A polypeptide folds into a compact, three-dimensional shape stabilized by interactions between R groups of amino acids to form a biologically active protein.
Quaternary	Two or more protein subunits combine to form a biologically active protein.



**Figure 3.16 Motifs and domains.** The elements of secondary structure can combine, fold, or crease to form motifs. These motifs are found in different proteins and can be used to predict function. Proteins also are made of larger domains, which are functionally distinct parts of a protein. The arrangement of these domains in space is the tertiary structure of a protein.

agents: pH, temp, ionic strength, solubility

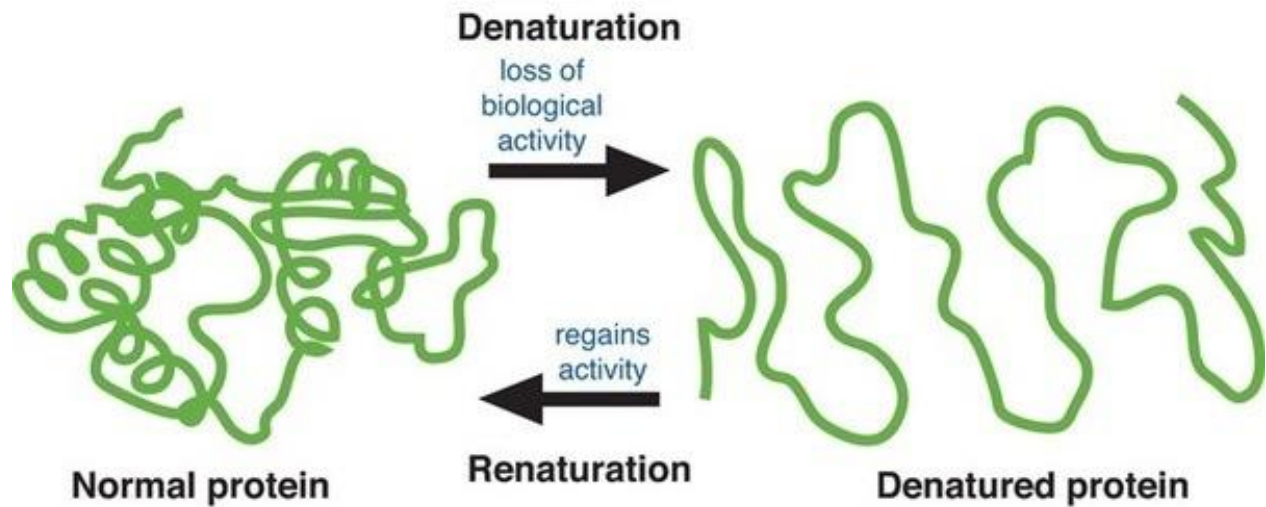
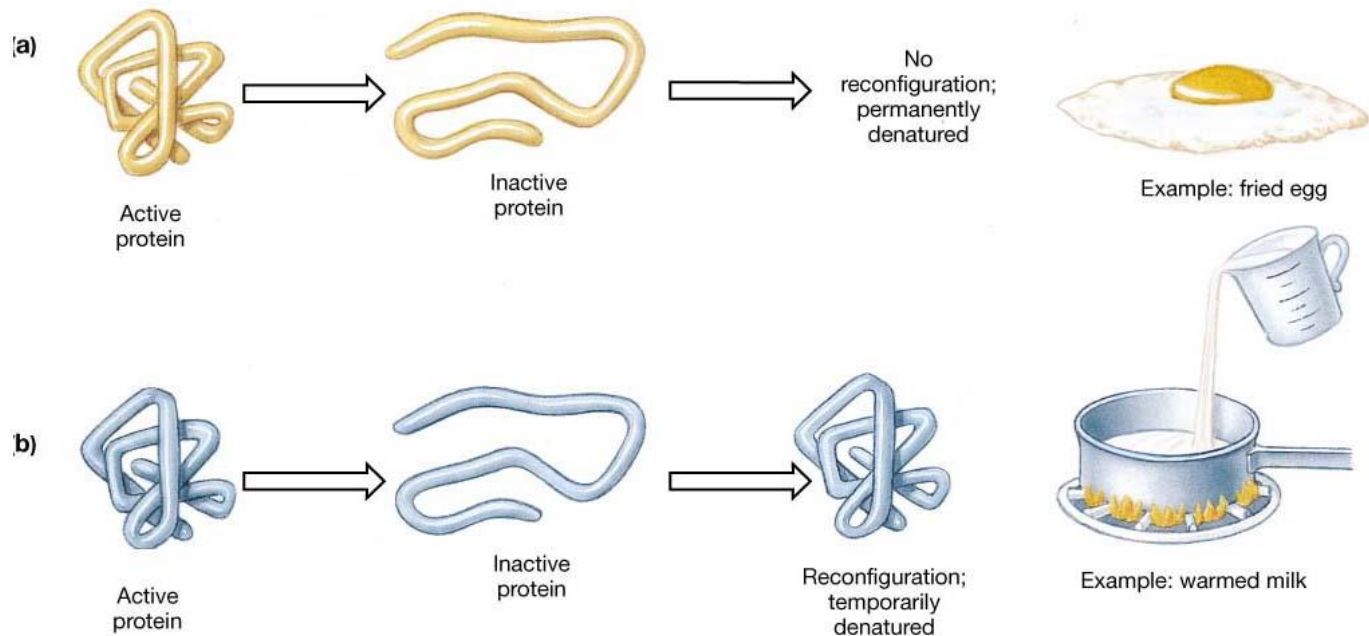
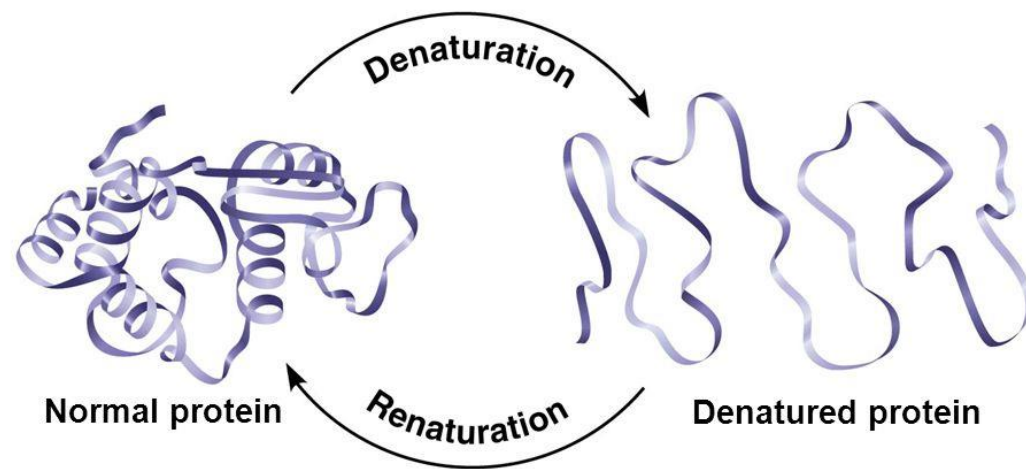


Figure 3.23-3



## ตารางที่ 3.11 กลไกการเสียสภาพของโปรตีนจำแนกตามสิ่งที่ทำให้เสียสภาพ

สิ่งทำให้เสียสภาพ (denaturing agent)	กลไกการเสียสภาพ (mode of action)
ความร้อน	ทำให้พันธะไฮโดรเจนระหว่างโมเลกุลสั้นสะเทือนอย่างรุนแรง จนทำให้เกิดการตกตะกอน เช่น การทอดไข่ดาว
รังสีไมโครเวฟ	ทำให้เกิดการสะเทือนของโมเลกุลอย่างรุนแรงจนทำให้พันธะไฮโดรเจนแตกออก
รังสีอัลตราไวโอเล็ต	ทำให้พันธะไฮโดรเจนระหว่างโมเลกุลสั้นสะเทือนอย่างรุนแรง เช่น ผิวไหม้จากแสงแดด
การเขย่าอย่างรุนแรง (violent shaking)	ทำให้โมเลกุลโปรตีนทรงกลมยืดยาวออกจนพันกันยุ่ง เช่น การตีไข่ขาวให้เป็นครีมในการทำขนมเบี้องหรือเมอร์แรงก์ (meringue)
สารซักฟอก (detergent)	มีผลต่อพันธะของโซ่ข้าง (หมู่ R)
ตัวทำลายอินทรีย์ (เช่น เอทานอล แอซีโตน)	มีผลต่อพันธะของโซ่ข้าง (หมู่ R) เนื่องจากตัวทำลายอินทรีย์ทำลายพันธะไฮโดรเจนและทำให้โปรตีนเสียสภาพอย่างรวดเร็ว เช่น การทำลายแบคทีเรียด้วยเอทานอลร้อยละ 70
กรดแก่ และเบสแก่	ทำลายพันธะไฮโดรเจน และสะพานเกลือ (salt bridge) จนทำให้เกิดปฏิกิริยาสลายพันธะเพปไทด์ของโปรตีน
เกลือของโลหะหนัก (เช่น เกลือของ $Hg^{2+}$ , $Ag^+$ , $Pb^{2+}$ )	โลหะหนักจับกับหมู่ซัลไฟไฮดริลและเกิดเป็นเกลือที่เป็นพิษ (poisonous salt)
สารรีดิวซ์	รีดิวซ์พันธะไดซัลไฟด์ของโปรตีนเกิดเป็นกรดอะมิโนที่มีหมู่ซัลไฟไฮดริล

ตารางที่ 3.12 จุดประจุไฟฟ้าสมดุล (pI) ของโปรตีนบางชนิด

โปรตีน	pI	โปรตีน	pI	โปรตีน	pI
เพปซิน	< 1.0	บีตา-แลกโทโกลบูลิน	5.2	โคโมทริปซินเจน	9.5
อัลบูมินในไข่ขาว	4.6	อินซูลิน	5.4	ไซโทโครม ซี	10.7
อัลบูมินในซีรัม	4.9	ฮีโมโกลบิน	6.8	ฮีสโตน	10.8
ยูรีเอส	5.0	ไมโอโกลบิน	7.0	ไลโซไซม์	11.0

pI > pH ของสารละลาย

net charge (-)

pI < pH ของสารละลาย

net charge (+)

pI = pH ของสารละลาย

net charge (0)

โปรตีนผลักกัน  
ไม่ตกตะกอน

โปรตีนรวมกัน  
ตกตะกอน

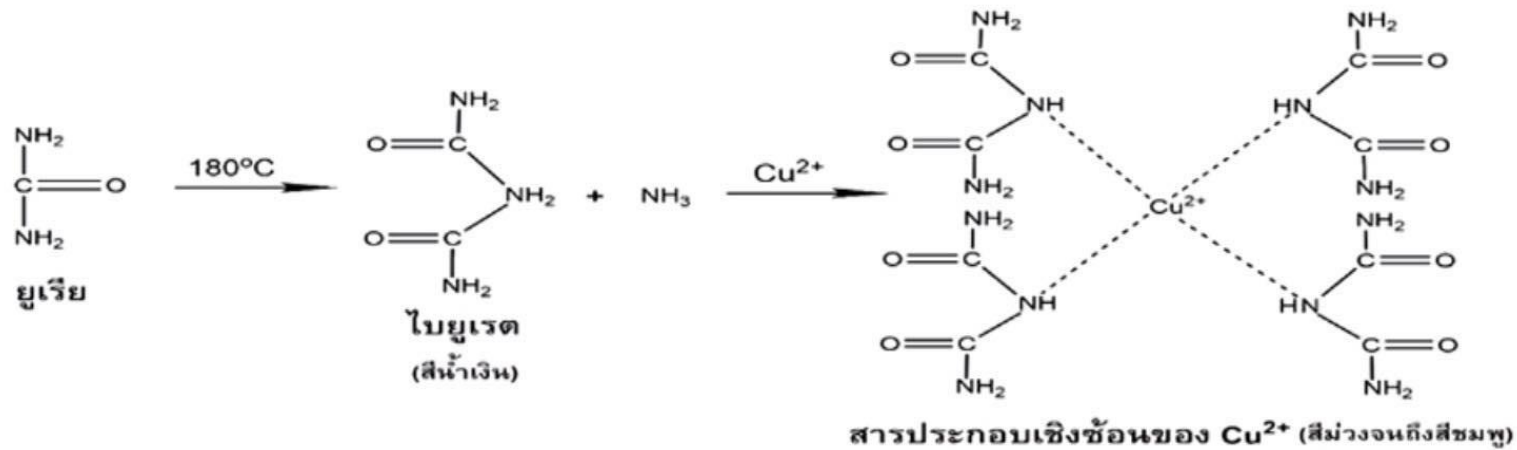
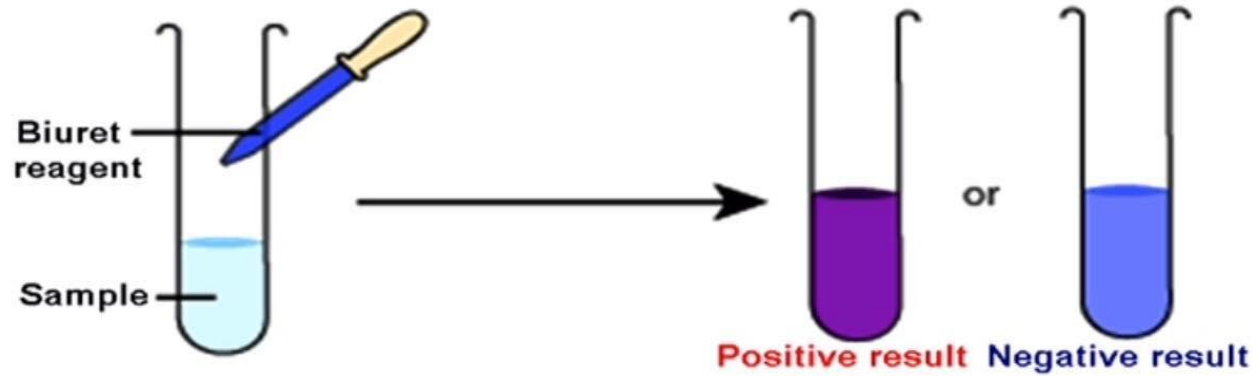
กรดอะมิโนแต่ละชนิดมีค่าความเป็นกรด-เบสที่ทำให้ประจุมีค่าสมมูลหรือมีจำนวนประจุบวกและประจุลบเท่ากัน เรียกค่าความเป็นกรด-เบสนี้ว่า จุดไอโซอิเล็กทริก หรือจุดประจุไฟฟ้าสมมูล (isoelectric point, pI)

กรดอะมิโน	pI	กรดอะมิโน	pI	กรดอะมิโน	pI
ไกลซีน	5.97	เฟนิลอะลานีน	5.48	ไทโรซีน	5.66
อะลานีน	6.01	ทรีปโตเฟน	5.88	กรดแอสพาร์ติก	2.77
เวอรีน	5.97	เซรีน	5.68	กรดกลูตามิก	3.22
ลิวซีน	5.98	ทรีโอนีน	5.87	ไลซีน	9.74
ไอโซลิวซีน	6.02	ซิสเทอีน	5.07	อาร์จินีน	10.76
โพรลีน	6.48	แอสพาราจีน	5.41	ฮิสทีดีน	7.59
เมไทโอนีน	5.74	กลูตามีน	5.65		

# Biuret test

## การทดสอบโปรตีน

ทดสอบสารที่มีพันธะเพปไทด์ (-CO-NH-) ตั้งแต่ 2 ตำแหน่งขึ้นไป ได้แก่ ไตรเพปไทด์ โอลิโกเพปไทด์ และโปรตีนที่มีโครงสร้างซับซ้อน



## Sulfur test

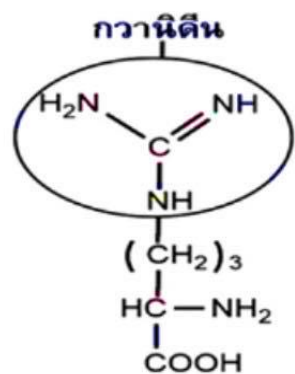
ทดสอบกรดอะมิโนที่มีกำมะถันเป็นองค์ประกอบ ได้แก่ ซีสเทอีน และซิสทีน ยกเว้นเมไทโอนีน



ตะกอนดำ

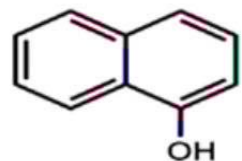
## Sakaguchi reaction

ใช้ทดสอบกรดอะมิโนอาร์จินีนที่มีโซ่ข้าง เป็นกวานิดีน (guanidine)

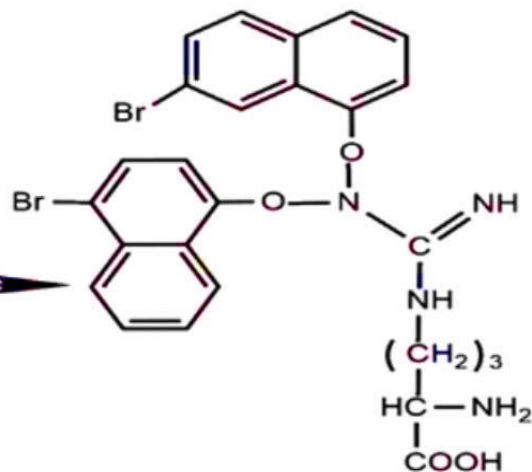
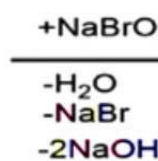


+

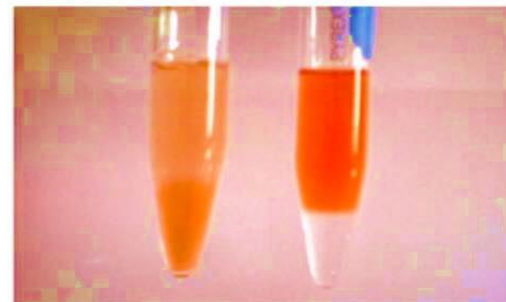
2



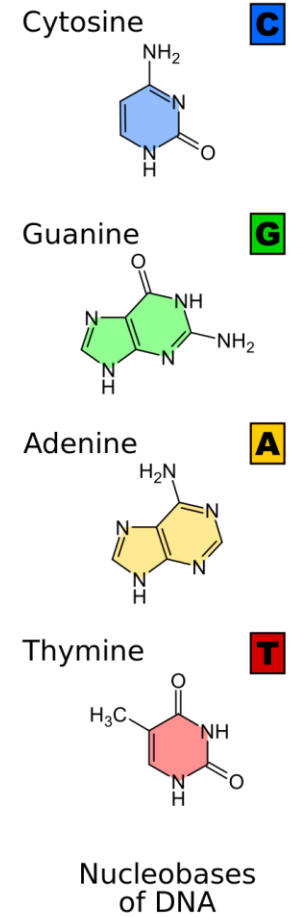
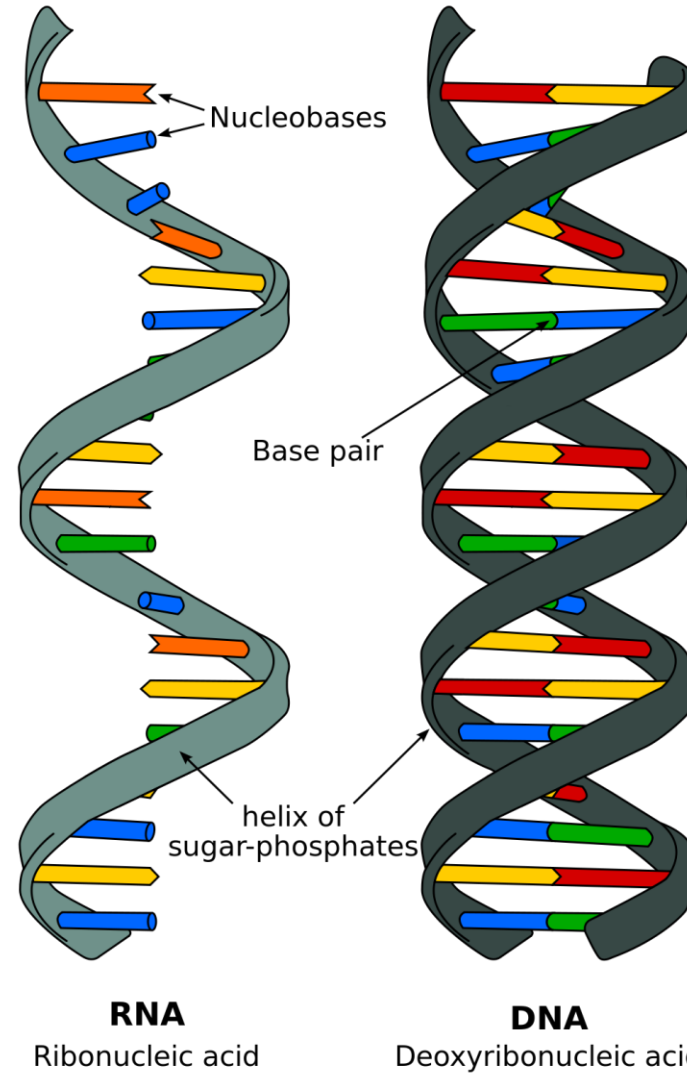
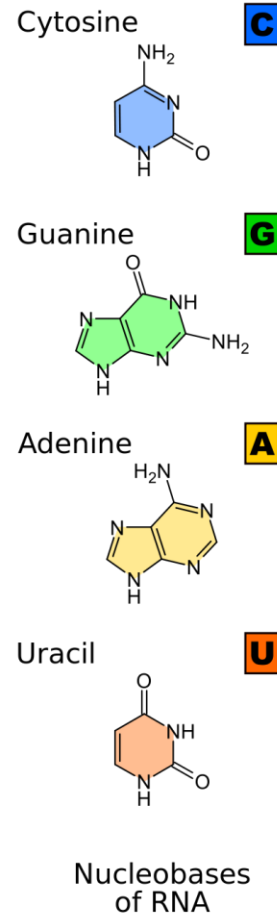
แอลฟา-นัพทอล



สารประกอบเชิงซ้อนสีแดง

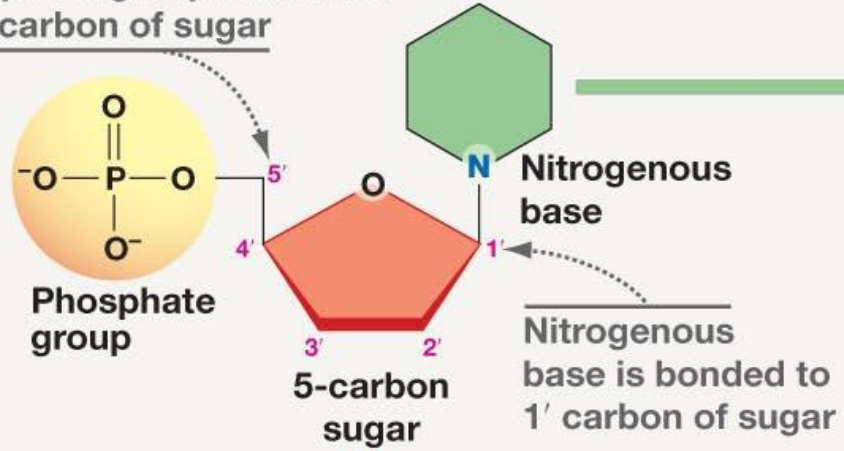


# Nucleic acids

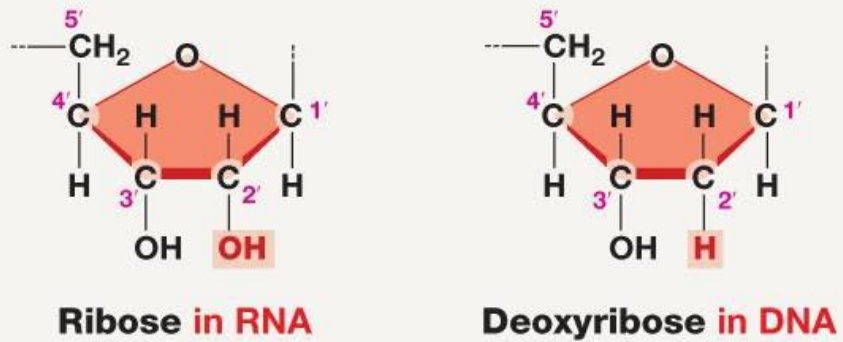


### (a) Nucleotide

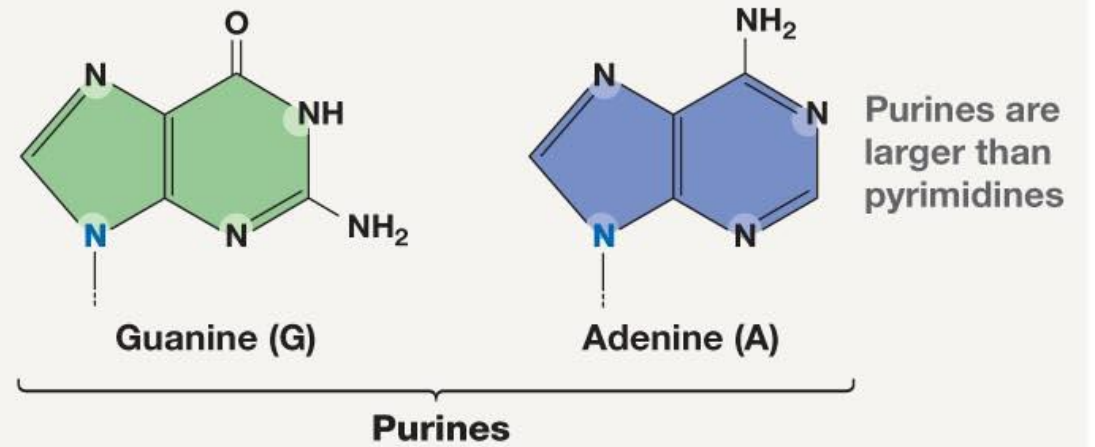
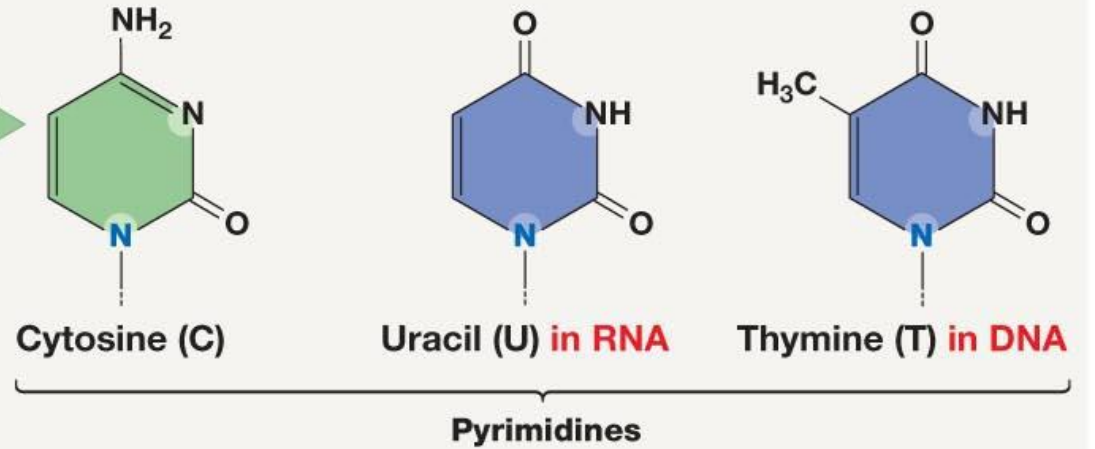
Phosphate group is bonded to 5' carbon of sugar



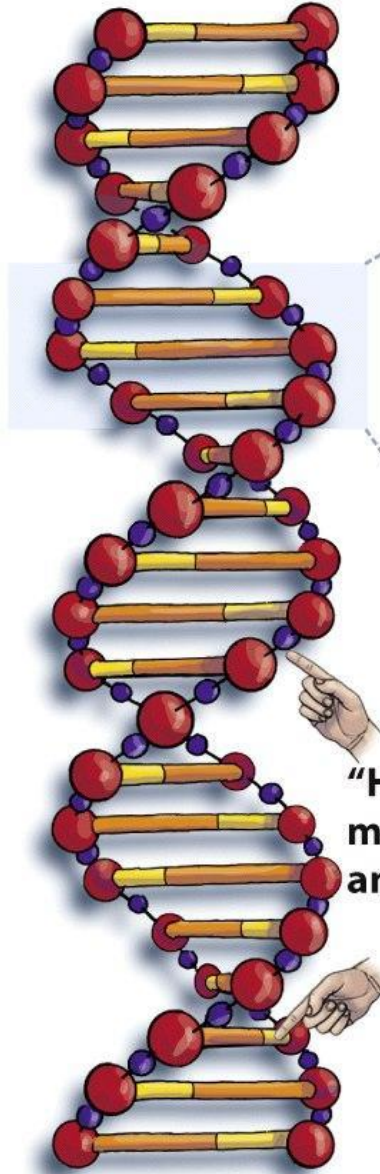
### (b) Sugars



### (c) Nitrogenous bases



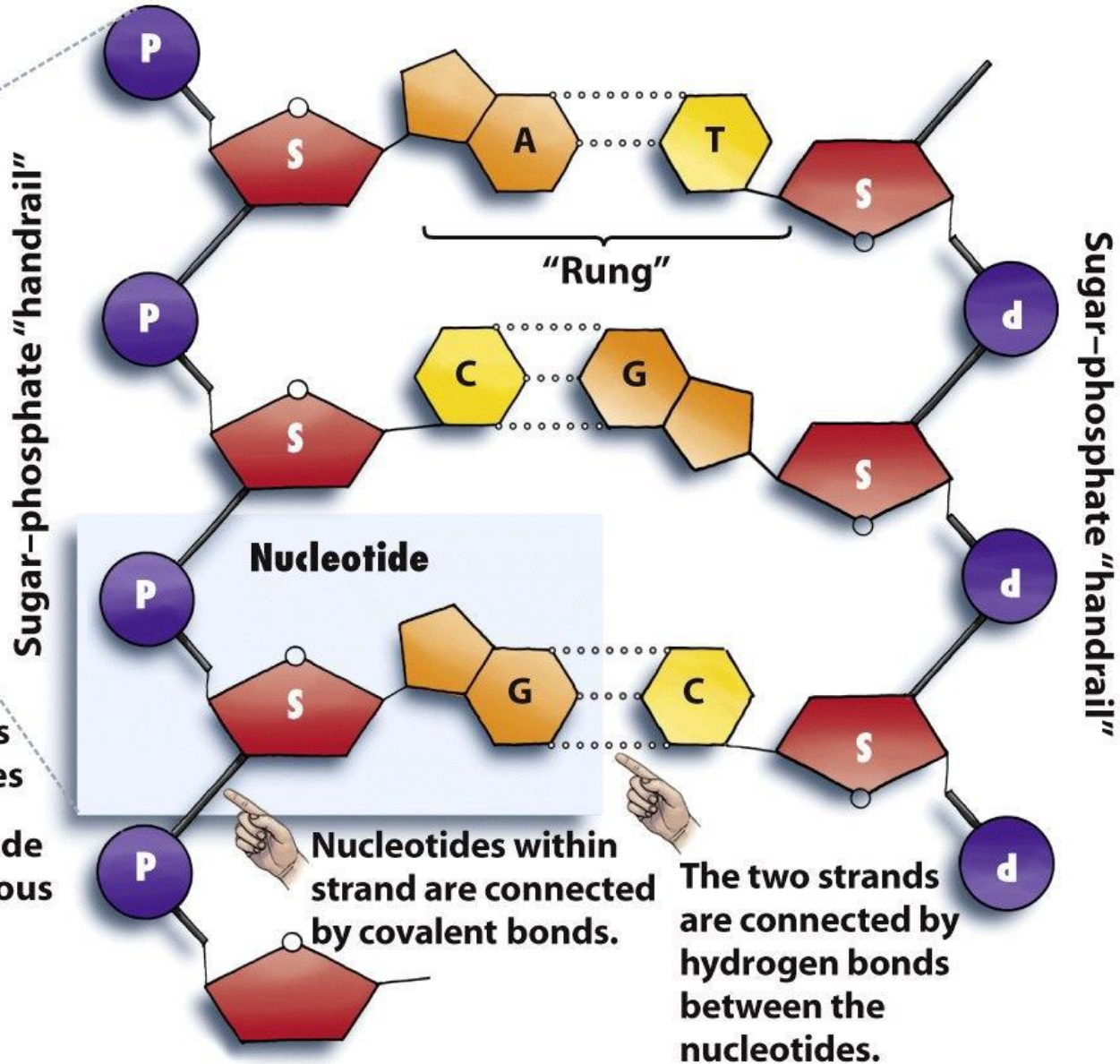
DNA double helix is made of two strands.



"Handrails" made of sugars and phosphates

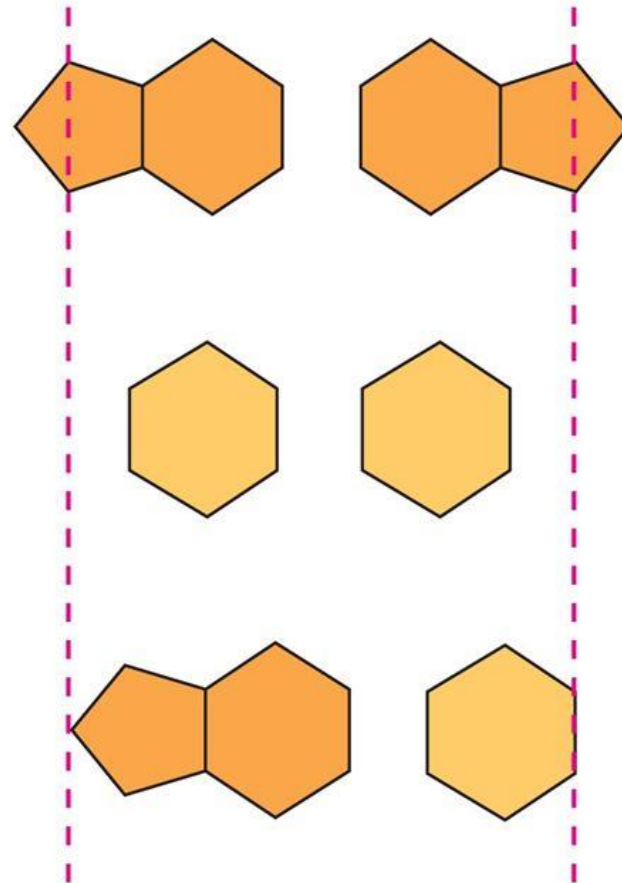
"Rungs" made of nitrogenous bases

Each strand is a chain of of antiparallel nucleotides.



A purine-purine pair is too wide and a pyrimidine-pyrimidine pair too narrow to account for the 2-nm diameter of the double helix.

LE 16-UN298



Purine + purine: too wide

Pyrimidine + pyrimidine: too narrow

Purine + pyrimidine: width consistent with X-ray data

# Chargaff's Rule: Rule of Base Pairing

**Table 8.1**

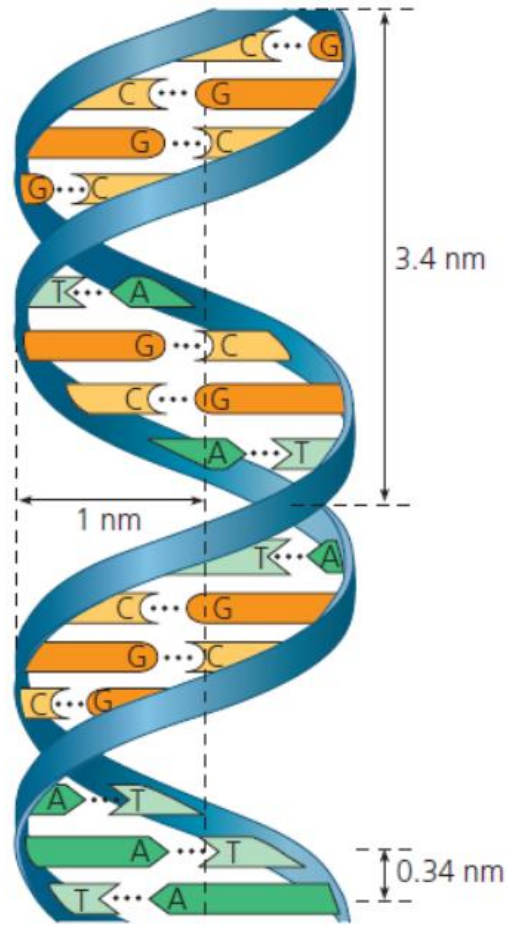
**Base composition of DNA from different sources and ratios of bases**

Source of DNA					Ratio		
	A	T	G	C	A/T	G/C	(A + G)/ (T + C)
<i>E. coli</i>	26.0	23.9	24.9	25.2	1.09	0.99	1.04
Yeast	31.3	32.9	18.7	17.1	0.95	1.09	1.00
Sea urchin	32.8	32.1	17.7	18.4	1.02	0.96	1.00
Rat	28.6	28.4	21.4	21.5	1.01	1.00	1.00
Human	30.3	30.3	19.5	19.9	1.00	0.98	0.99

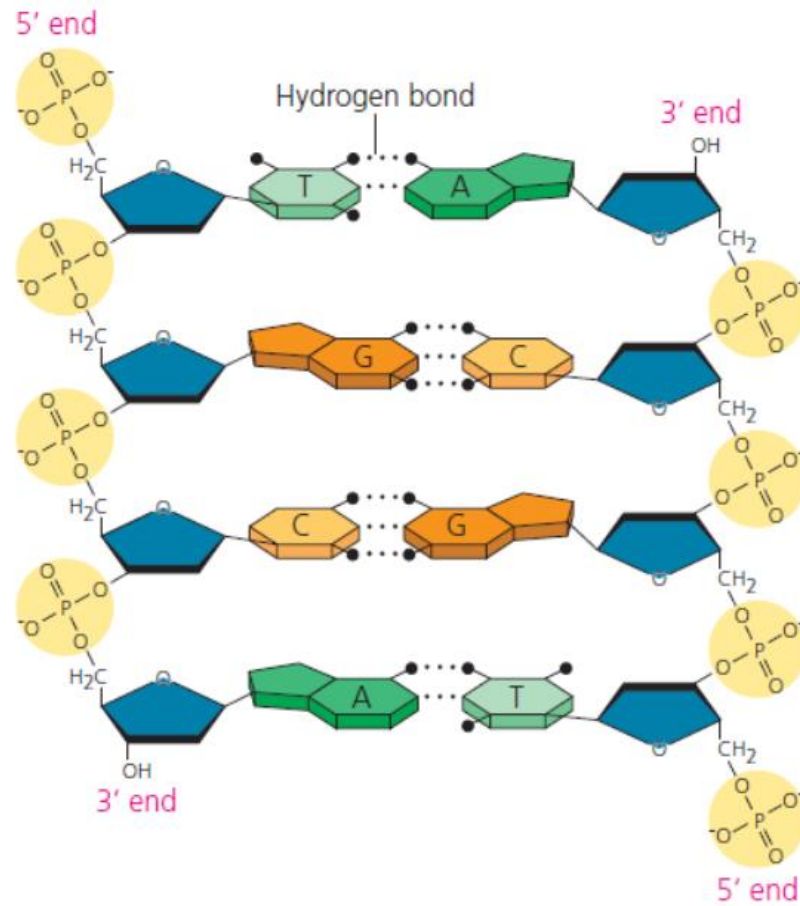
**All DNA,  $A=T$  &  $G=C$ ,  
that is,  $A+G/T+C = 1$**

## Names of Nucleosides and Nucleotides

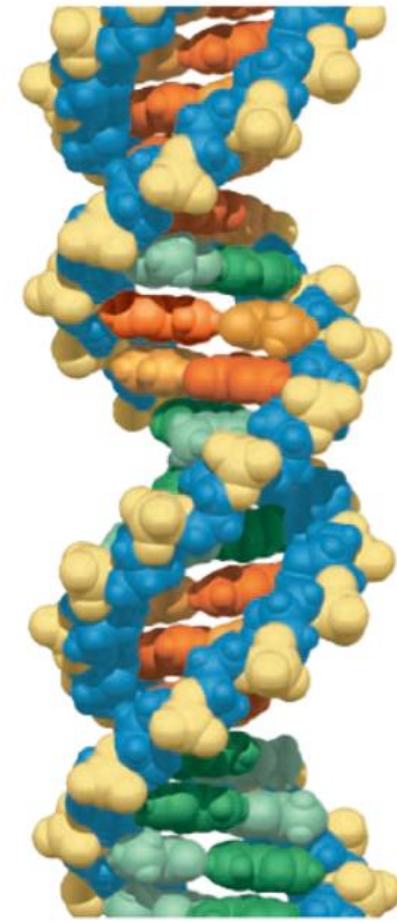
Base	Nucleosides	Nucleotides
<b>RNA</b>		
Adenine (A)	Adenosine (A)	Adenosine 5'-monophosphate (AMP)
Guanine (G)	Guanosine (G)	Guanosine 5'-monophosphate (GMP)
Cytosine (C)	Cytidine (C)	Cytidine 5'-monophosphate (CMP)
Uracil (U)	Uridine (U)	Uridine 5'-monophosphate (UMP)
<b>DNA</b>		
Adenine (A)	Deoxyadenosine (A)	Deoxyadenosine 5'-monophosphate (dAMP)
Guanine (G)	Deoxyguanosine (G)	Deoxyguanosine 5'-monophosphate (dGMP)
Cytosine (C)	Deoxycytidine (C)	Deoxycytidine 5'-monophosphate (dCMP)
Thymine (T)	Deoxythymidine (T)	Deoxythymidine 5'-monophosphate (dTMP)



**(a) Key features of DNA structure.** The “ribbons” in this diagram represent the sugar-phosphate backbones of the two DNA strands. The helix is “right-handed,” curving up to the right. The two strands are held together by hydrogen bonds (dotted lines) between the nitrogenous bases, which are paired in the interior of the double helix.

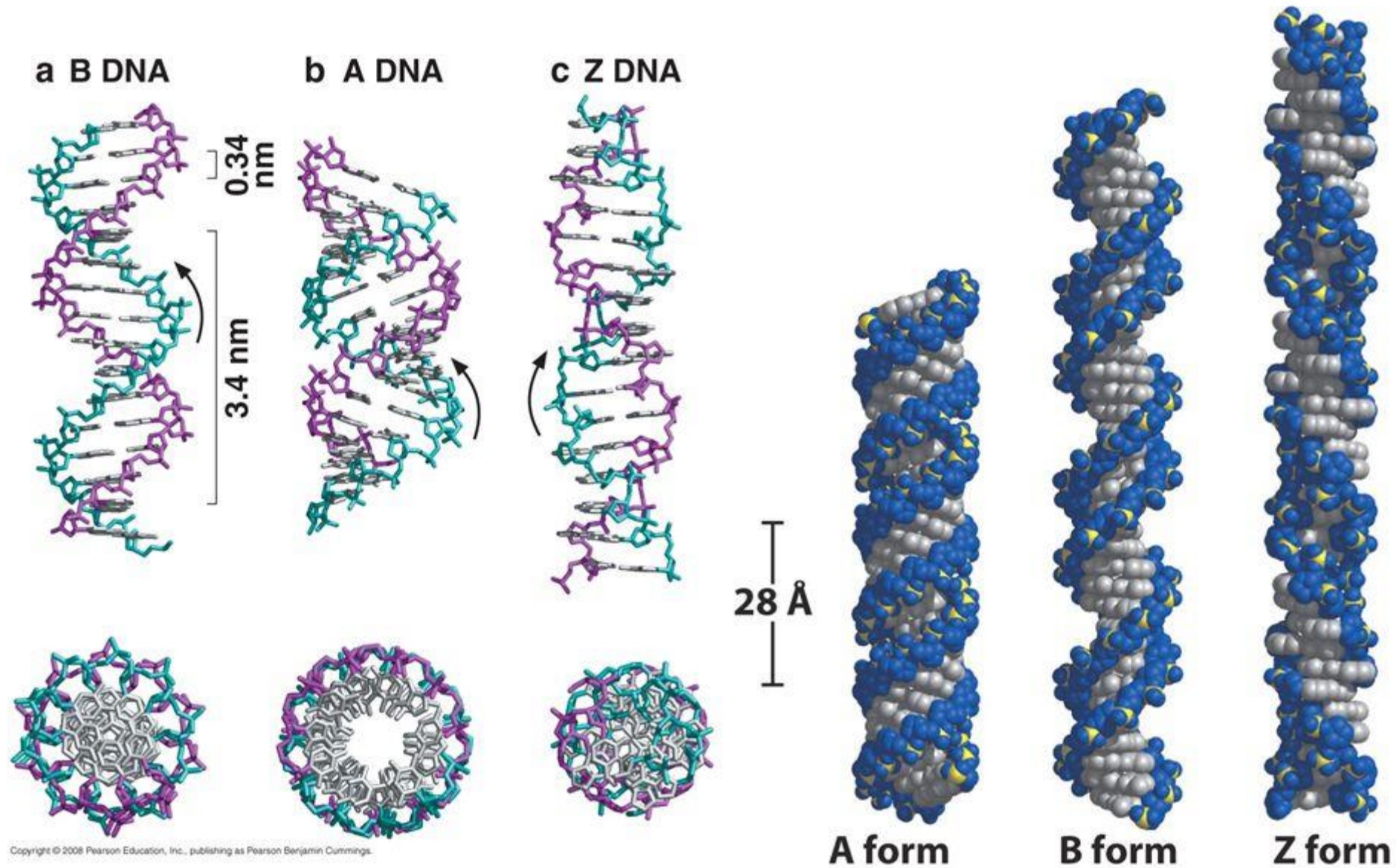


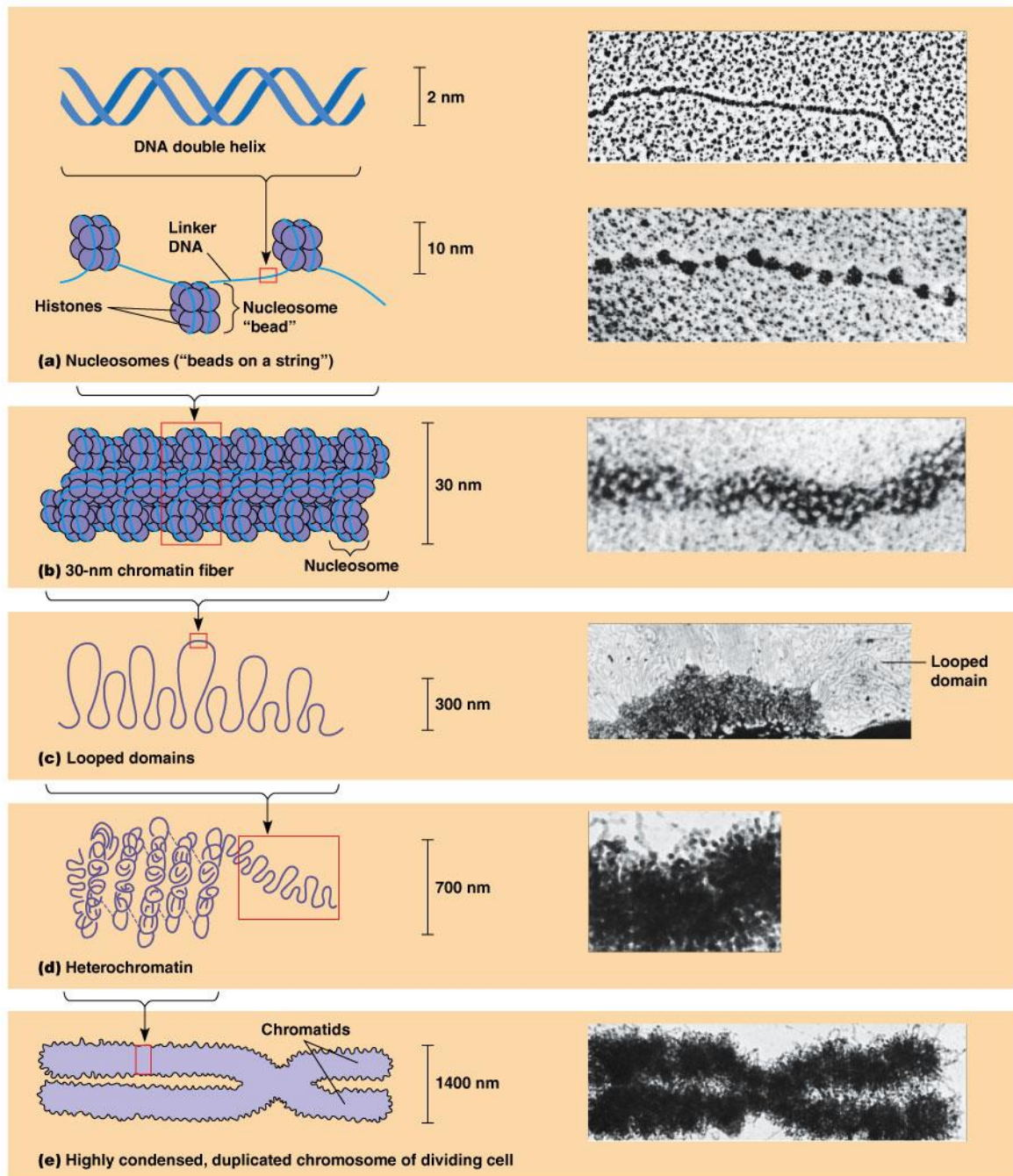
**(b) Partial chemical structure.** For clarity, the two DNA strands are shown untwisted in this partial chemical structure. Strong covalent bonds link the units of each strand, while weaker hydrogen bonds between the bases hold one strand to the other. Notice that the strands are antiparallel, meaning that they are oriented in opposite directions, like the lanes of a divided highway.



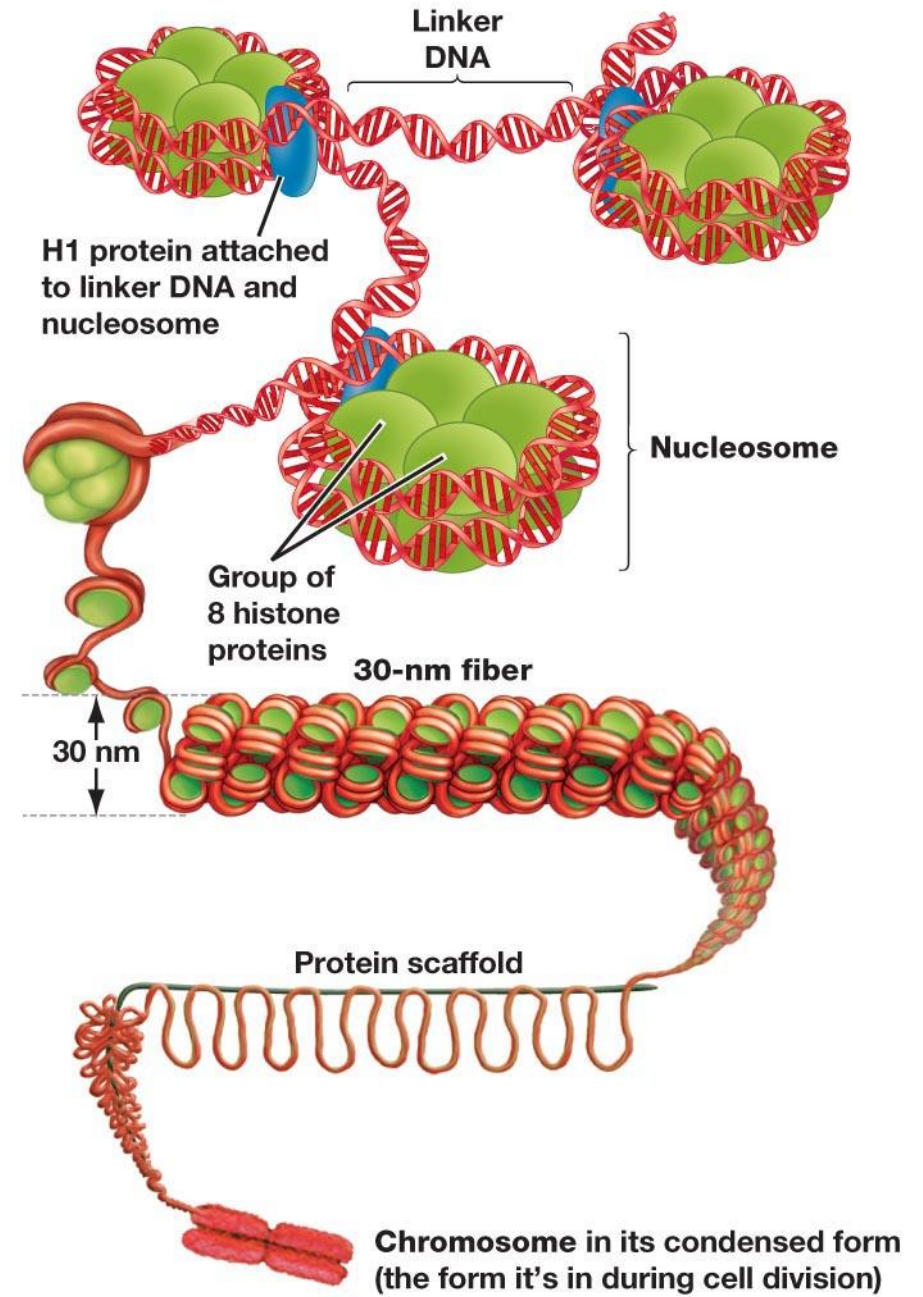
**(c) Space-filling model.** The tight stacking of the base pairs is clear in this computer-generated, space-filling model. Van der Waals interactions between the stacked pairs play a major role in holding the molecule together.

# The double helix exists in multiple conformations





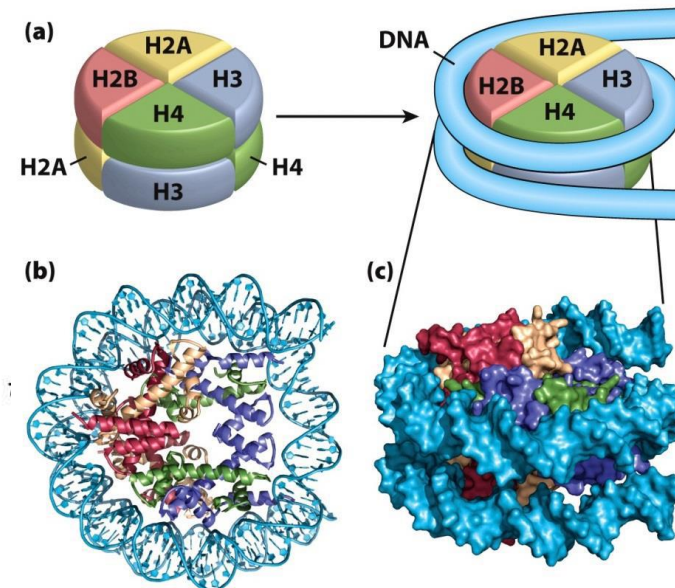
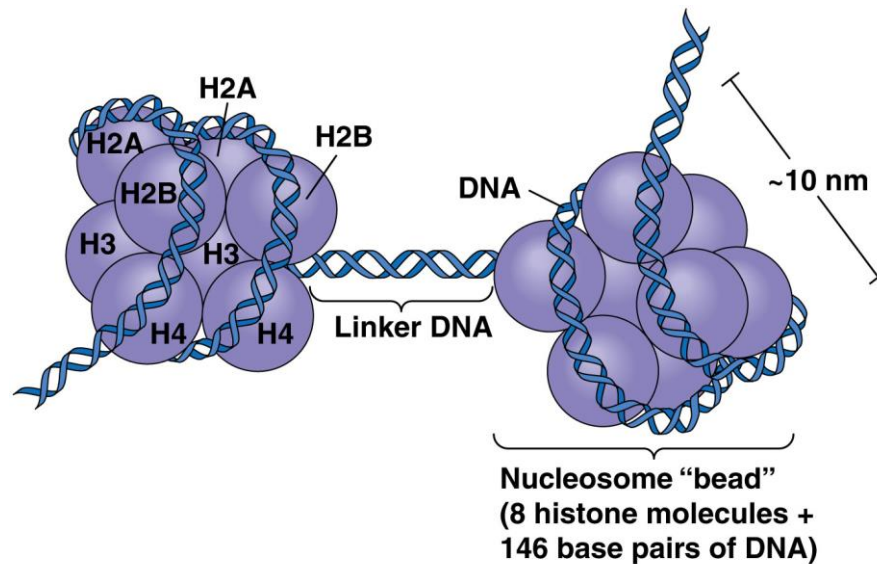
**(b) Nucleosome structure**



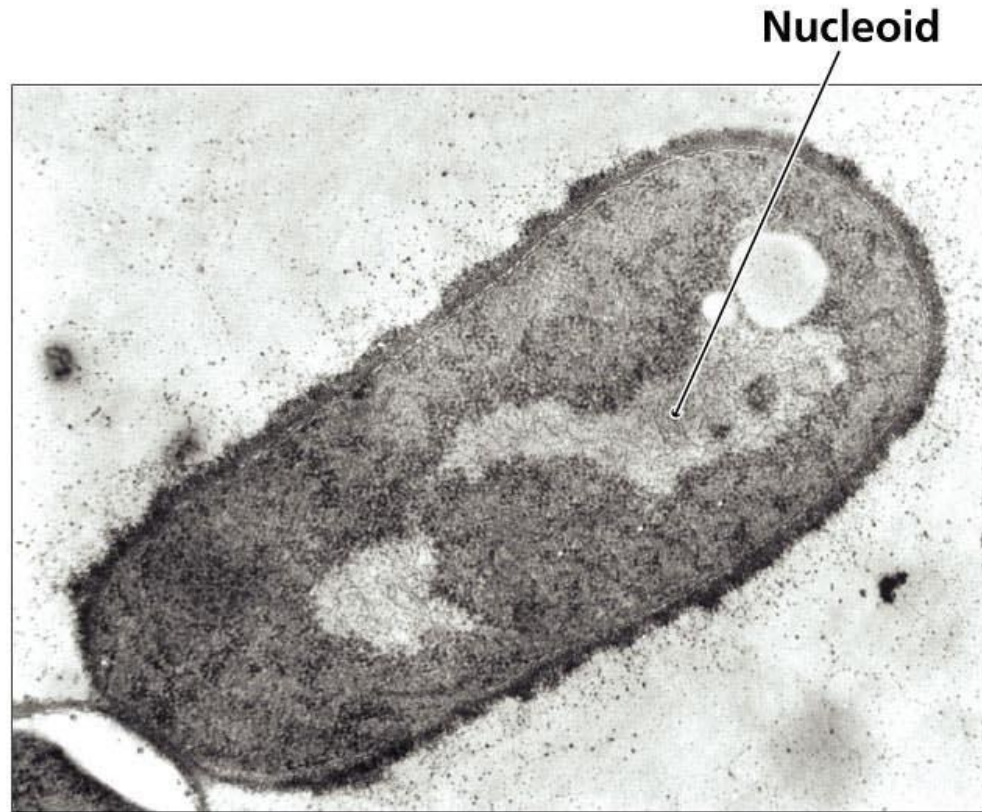
**TABLE 5.4** The Major Histone Proteins

Histone <sup>a</sup>	Molecular weight	Number of amino acids	Percentage lysine + arginine
H1	22,500	244	30.8
H2A	13,960	129	20.2
H2B	13,774	125	22.4
H3	15,273	135	22.9
H4	11,236	102	24.5

<sup>a</sup> Data are for rabbit (H1) and bovine histones.



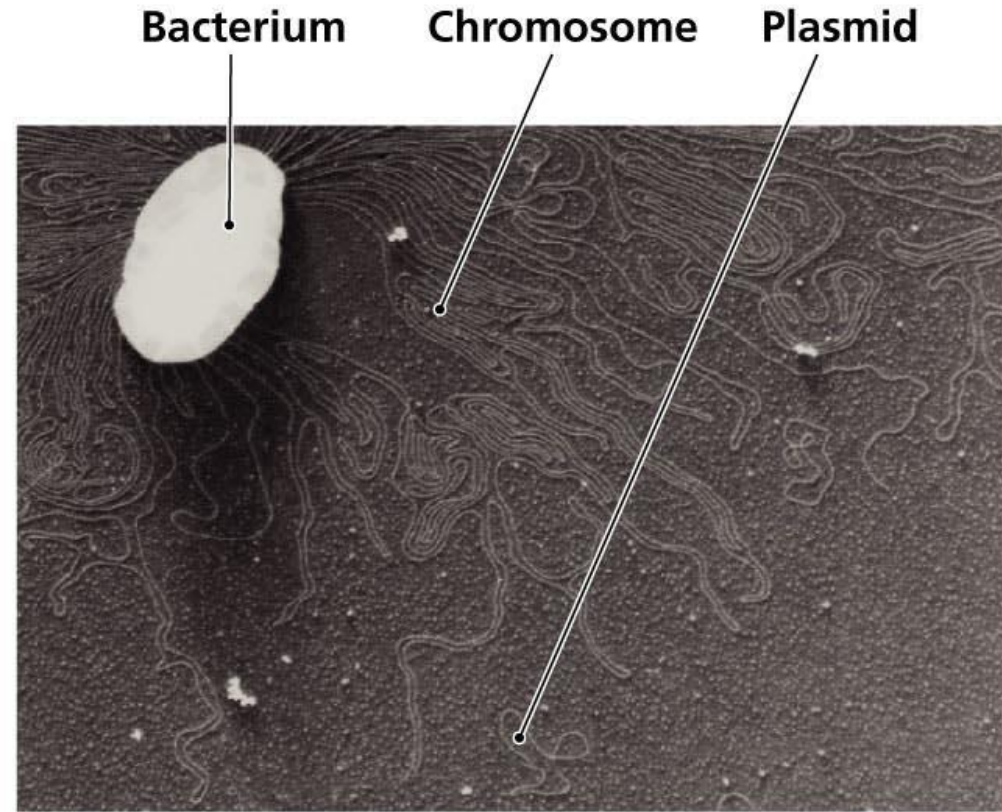
# Bacterial chromosome



(a)

TEM

0.5  $\mu\text{m}$

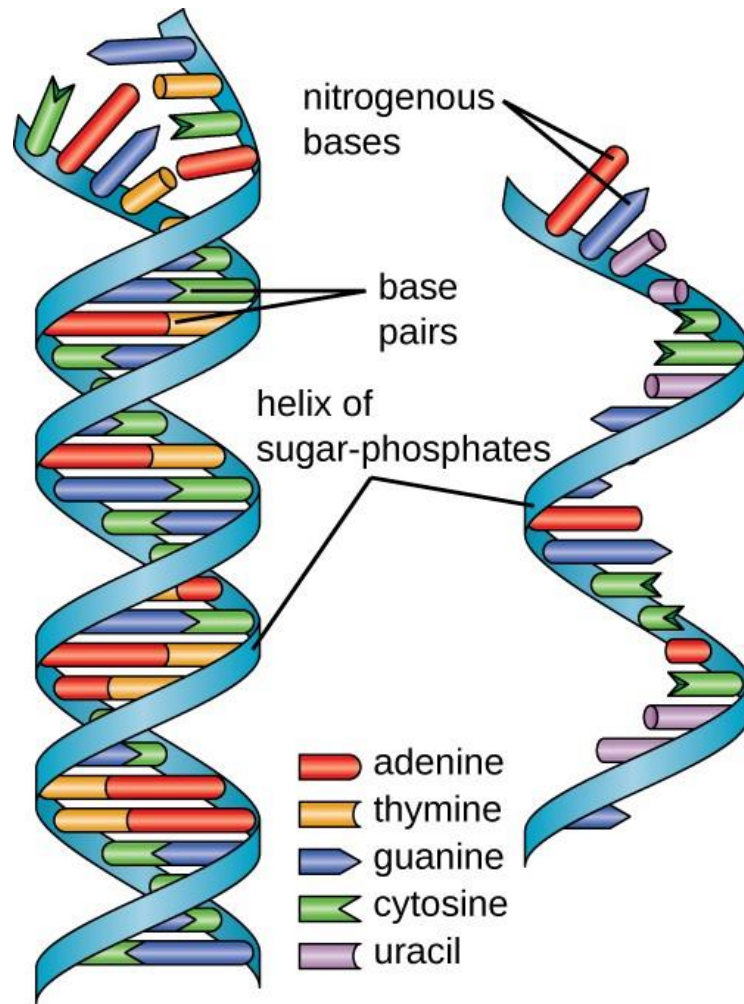


(b)

SEM

1  $\mu\text{m}$

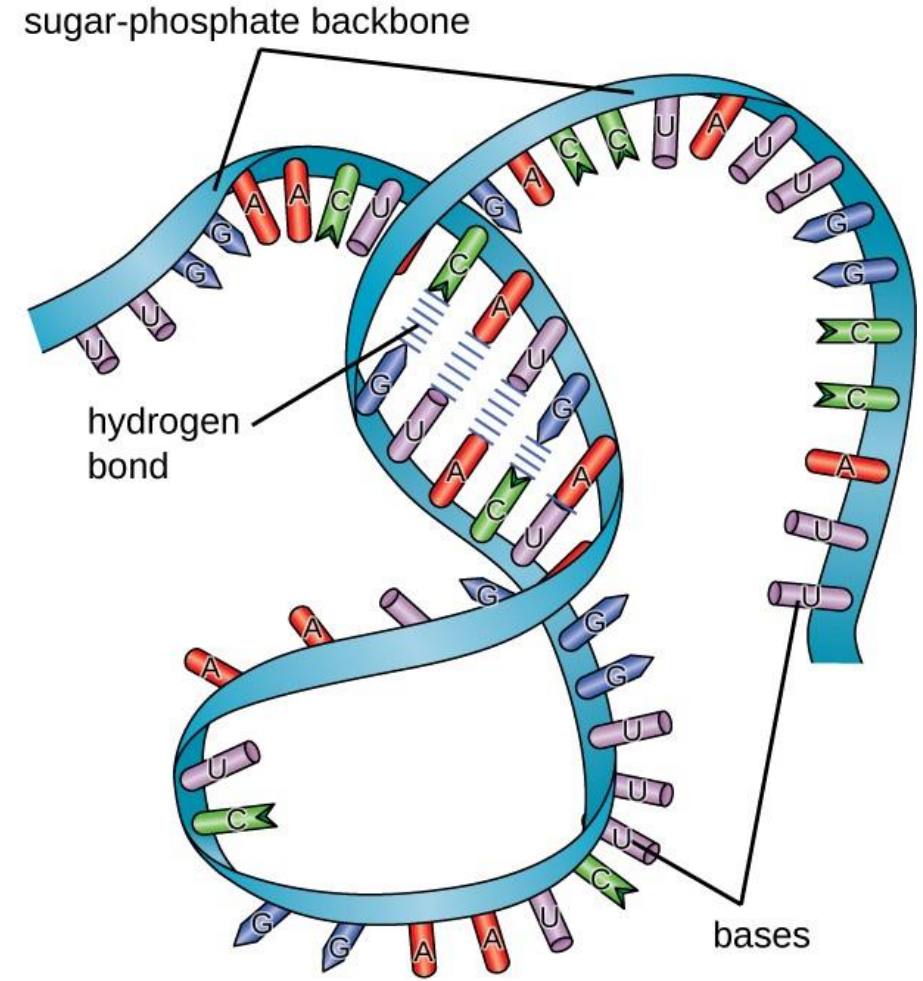
# RNA



DNA

RNA

(a)



(b)

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## Types of RNA

The three main types of RNA are:

Messenger RNA



Ribosomal RNA



Transfer RNA



