

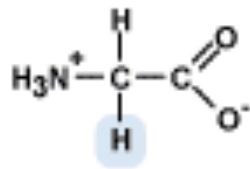
Ch18: Amino Acids and Proteins

Protein Functions

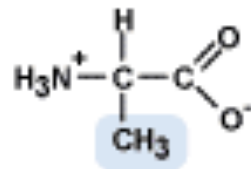
TABLE 18.2 Classification of Proteins by Function

Type	Function	Example
Enzymes	Catalyze biochemical reactions	<i>Amylase</i> —begins digestion of carbohydrates by hydrolysis
Hormones	Regulate body functions by carrying messages to receptors	<i>Insulin</i> —facilitates use of glucose for energy generation
Storage proteins	Make essential substances available when needed	<i>Myoglobin</i> —stores oxygen in muscles
Transport proteins	Carry substances through body fluids	<i>Serum albumin</i> —carries fatty acids in blood
Structural proteins	Provide mechanical shape and support	<i>Collagen</i> —provides structure to tendons and cartilage
Protective proteins	Defend the body against foreign matter	<i>Immunoglobulin</i> —aids in destruction of invading bacteria
Contractile proteins	Do mechanical work	<i>Myosin and actin</i> —govern muscle movement

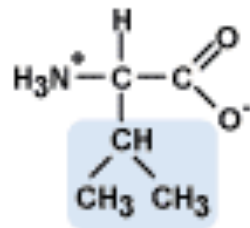
Amino Acids: Neutral Nonpolar Side Chains



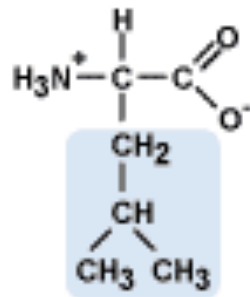
Glycine (Gly)



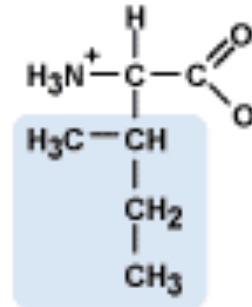
Alanine (Ala)



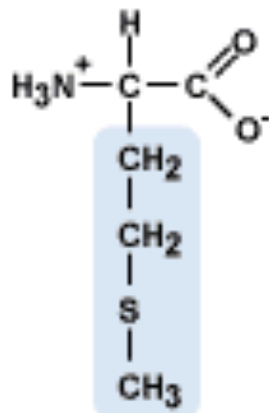
Valine (Val)



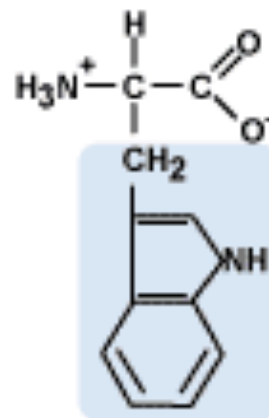
Leucine (Leu)



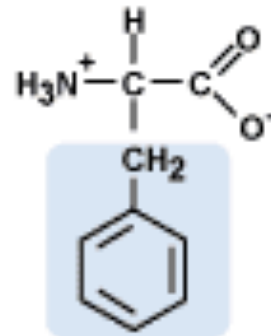
Isoleucine (Ile)



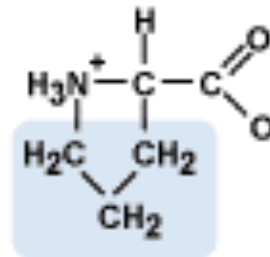
Methionine (Met)



Tryptophan (Trp)

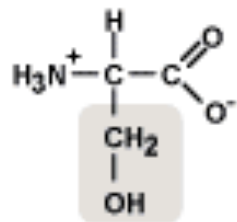


Phenylalanine (Phe)

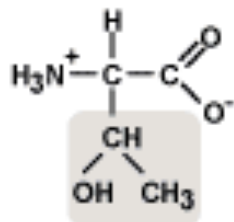


Proline (Pro)

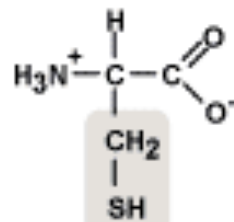
Amino Acids: Neutral Polar Side Chains



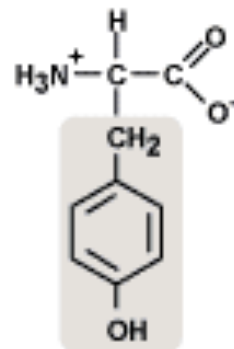
Serine (Ser)



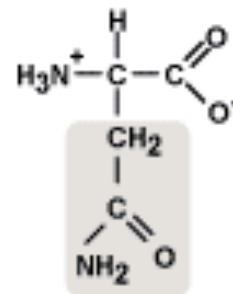
Threonine (Thr)



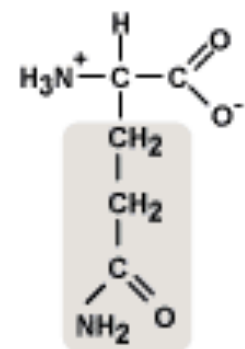
Cysteine (Cys)



Tyrosine (Tyr)

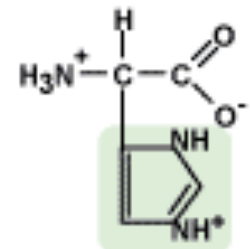
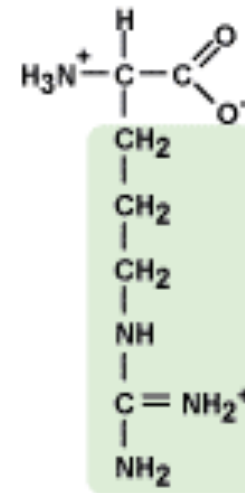
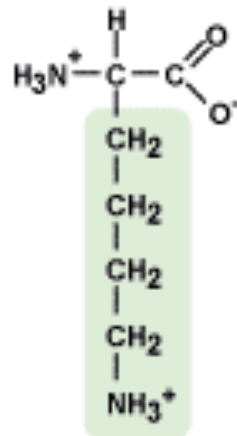
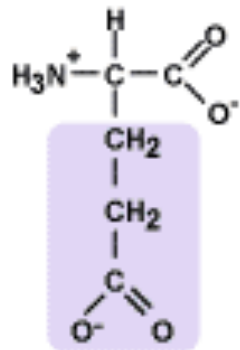
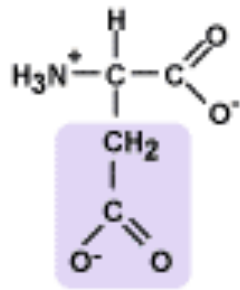


Asparagine (Asn)



Glutamine (Gln)

Amino Acids: Ionizable Side Chains

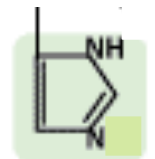
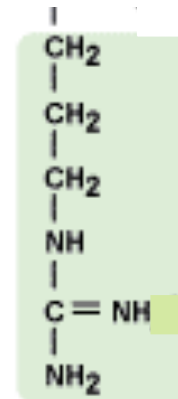
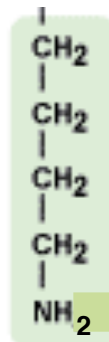
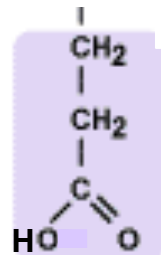
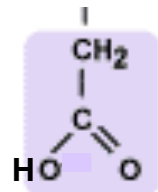


Acidic

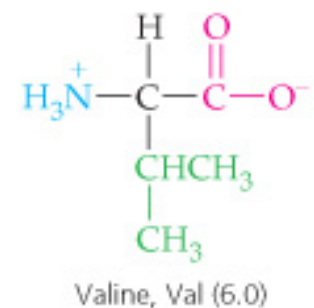
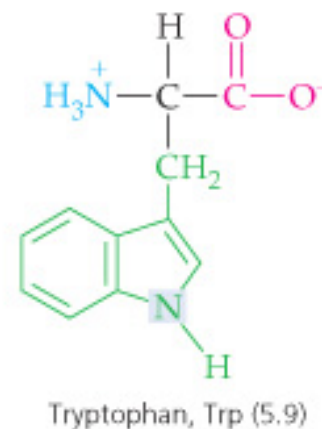
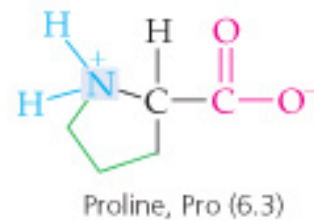
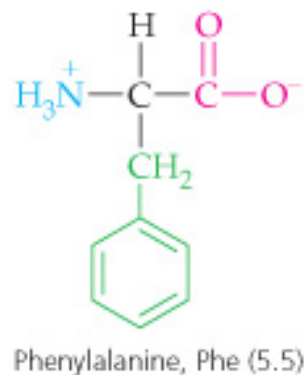
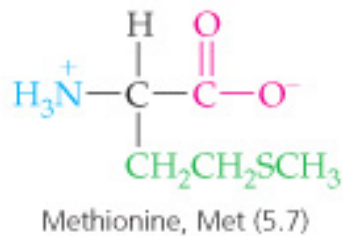
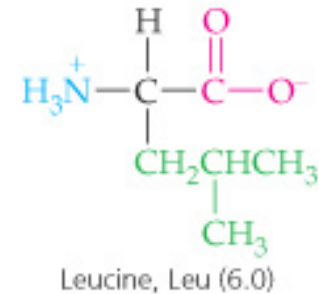
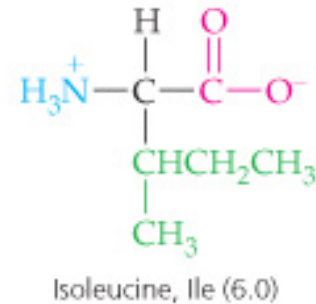
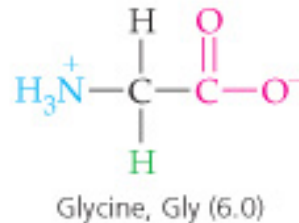
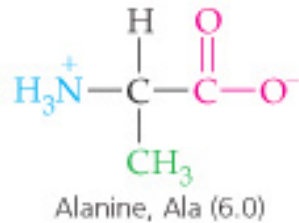
Aspartic Acid (Asp) Glutamic Acid (Glu)

Basic

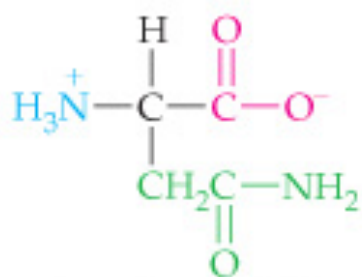
Lysine (Lys) Arginine (Arg) Histidine (His)



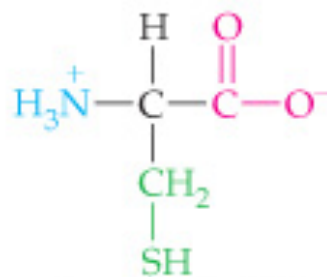
Amino Acids: Neutral Nonpolar Side Chains



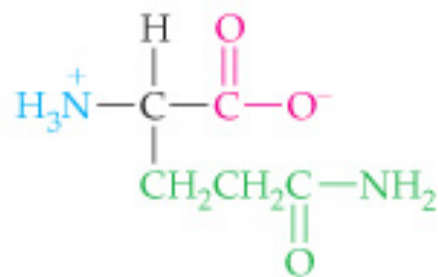
Amino Acids: Neutral Polar Side Chains



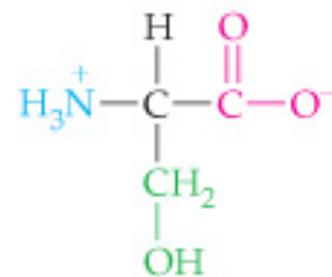
Asparagine, Asn (5.4)



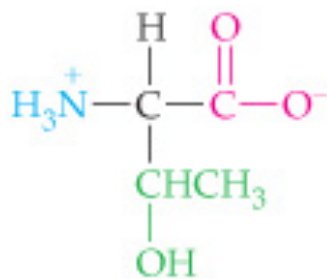
Cysteine, Cys (5.0)



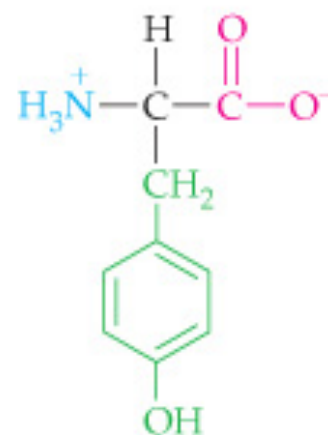
Glutamine, Gln (5.7)



Serine, Ser (5.7)

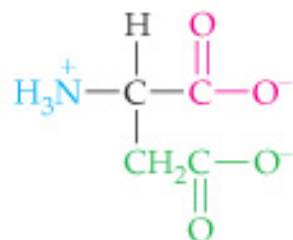


Threonine, Thr (5.6)

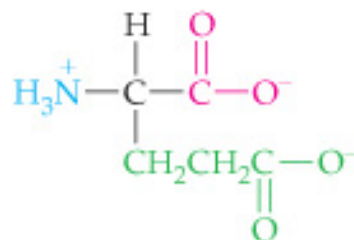


Tyrosine, Tyr (5.7)

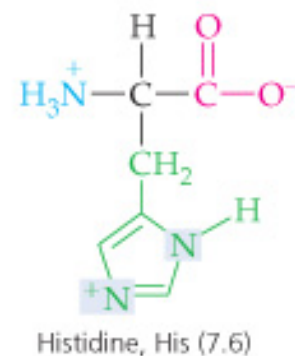
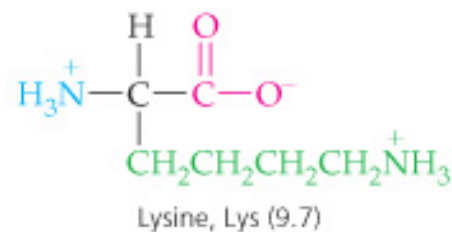
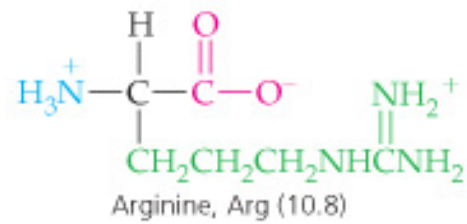
Amino Acids: Ionizable Side Chains



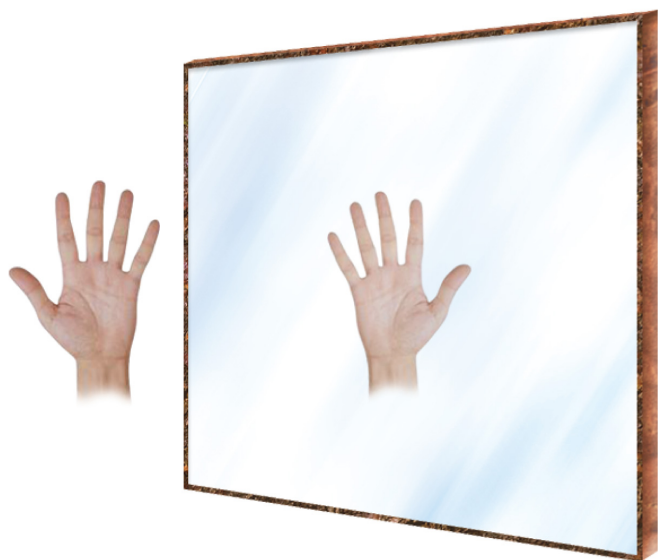
Aspartic acid, Asp (3.0)
(Aspartate)



Glutamic acid, Glu (3.2)
(Glutamate)



Handedness (Chirality)



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Hand is not superimposable on its mirror image → CHIRAL

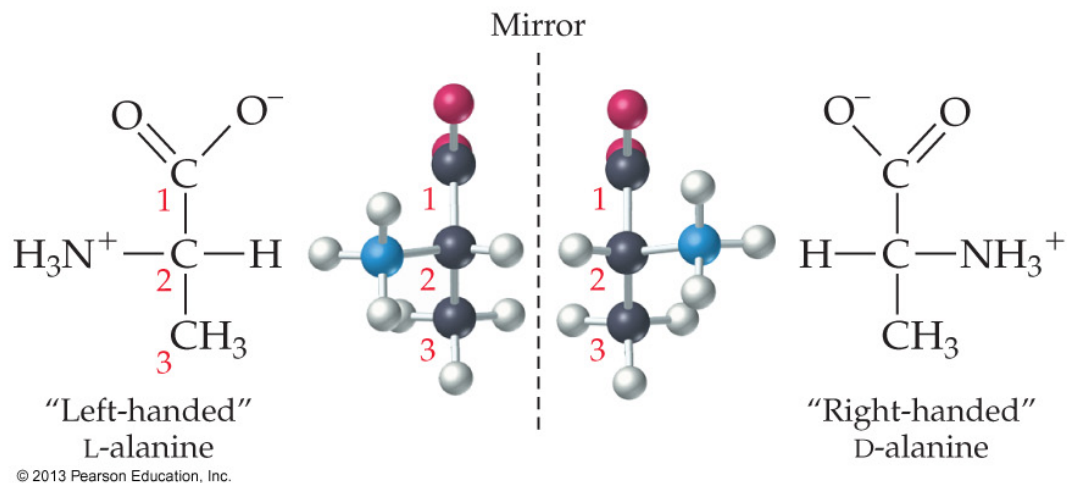


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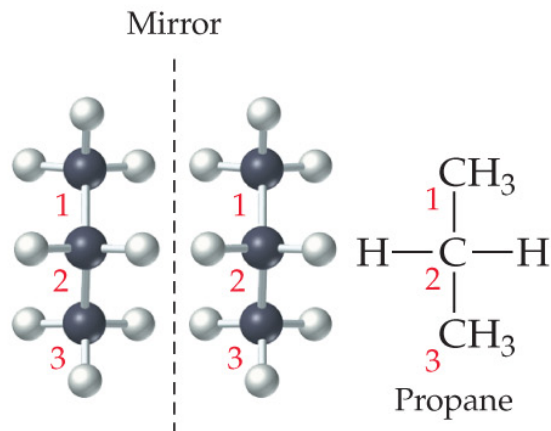
Chair is superimposable on its mirror image → ACHIRAL

Chirality

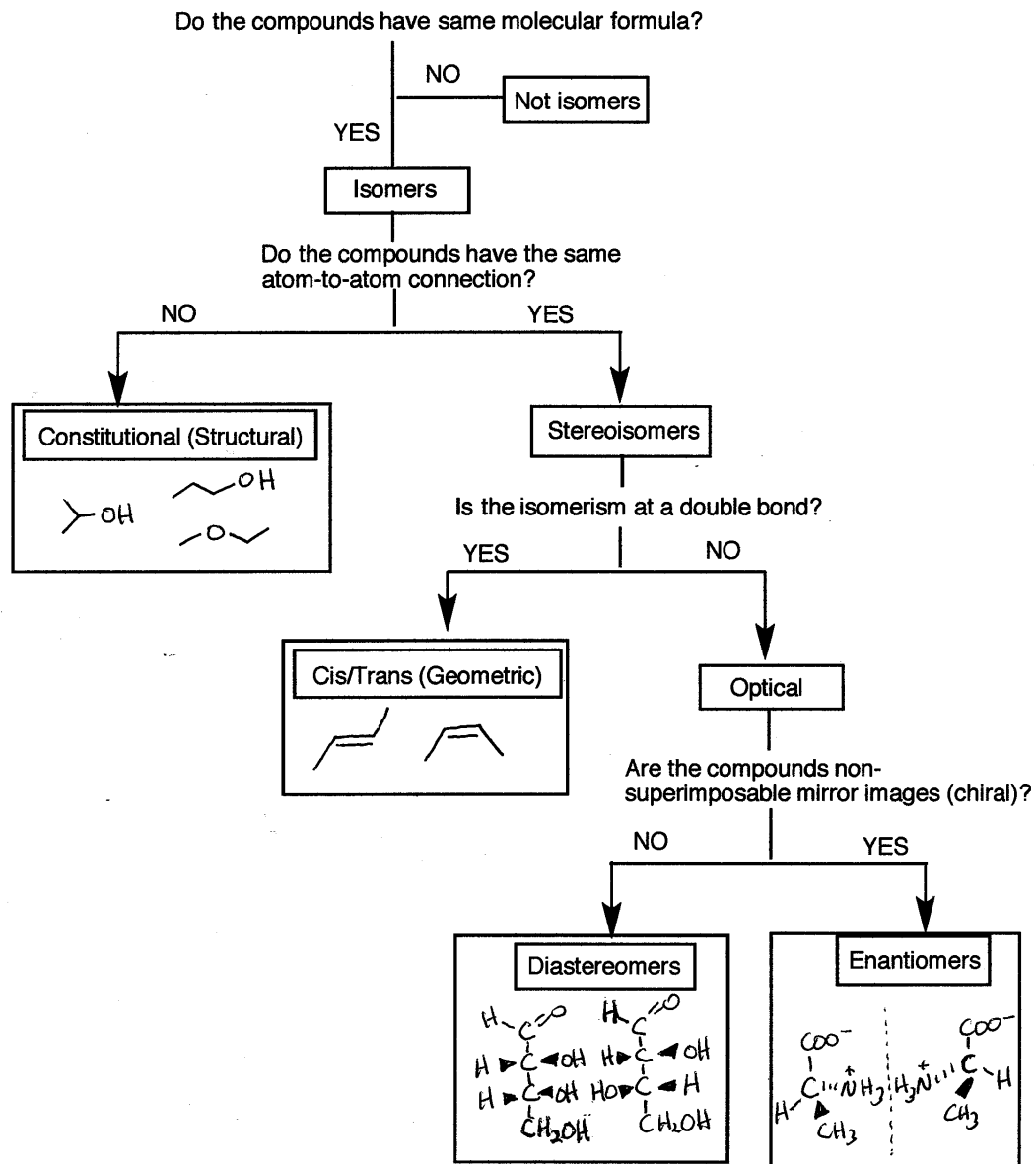
Alanine: Chiral



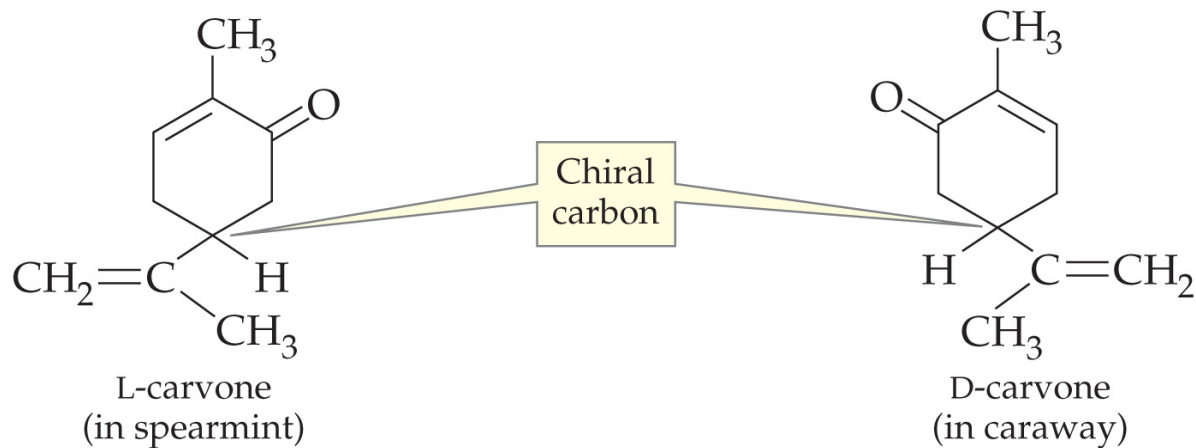
Propane: Achiral



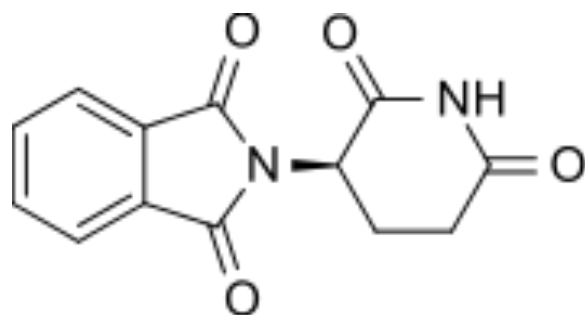
Isomer Types



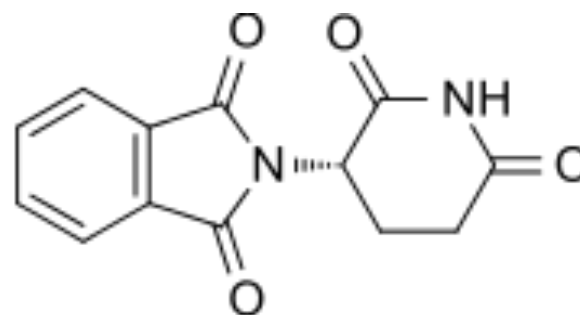
Enantiomers of Carvone



Case of Thalidomide



(R)-thalidomide



(S)-thalidomide

1957- Sold as sedative, nausea-reliever for pregnant women.
Racemic mix resulted in infants born with limb deformation

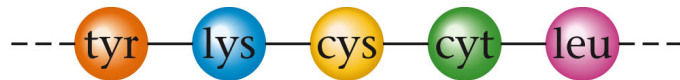
Levels of Protein Structure

1. Primary
2. Secondary
3. Tertiary
4. Quaternary

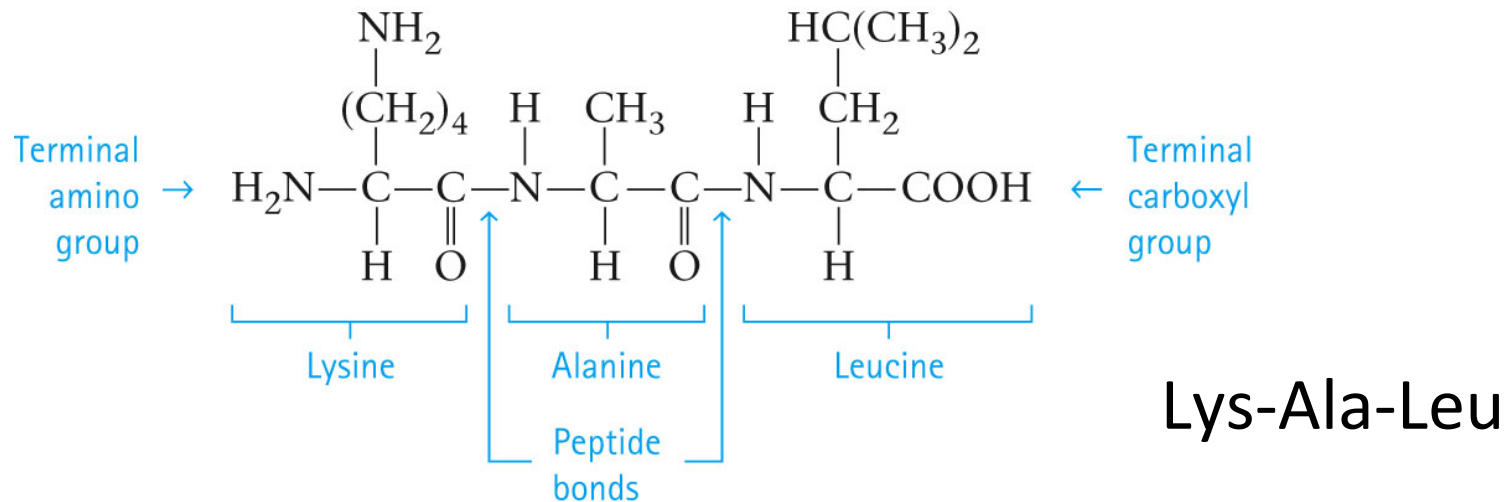
Protein structure and function are intimately linked!

Primary Structure of Proteins

Primary structure: the sequence of the amino acids (residues) in the chain.



- Read sequence from amino end → carboxyl end.



Primary Structures of Two Peptides

Oxytocin: Hormone that triggers contraction of uterus and milk secretion

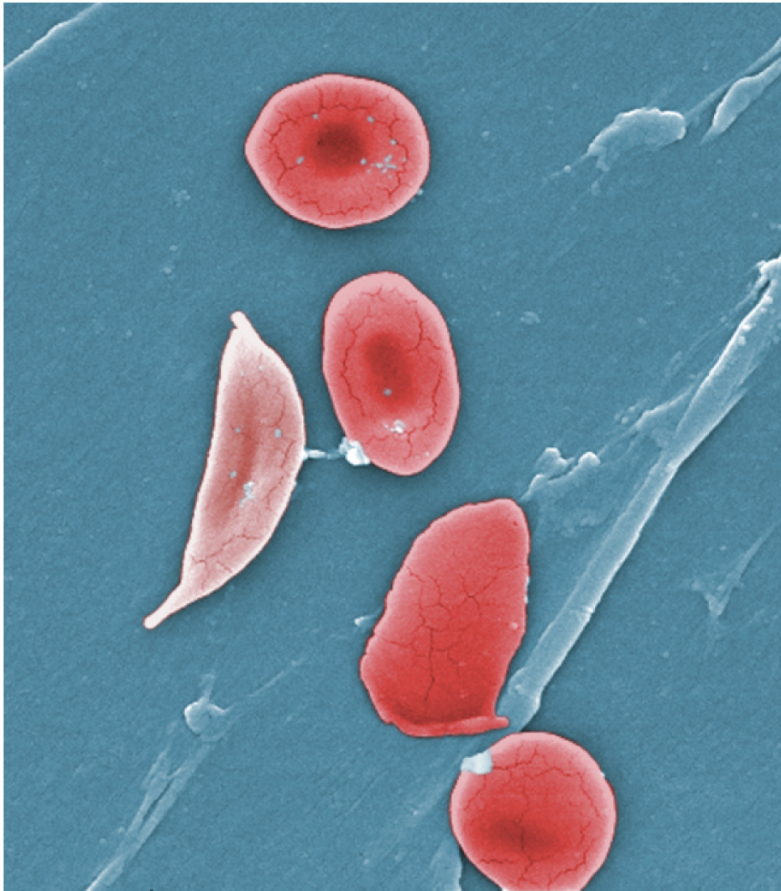
cys-tyr-**ile**-gin-asn-cys-pro-**leu**-gly

Vasopressin: Hormone that raises blood pressure and regulates kidney function

cys-tyr-**phe**-gin-asn-cys-pro-**arg**-gly

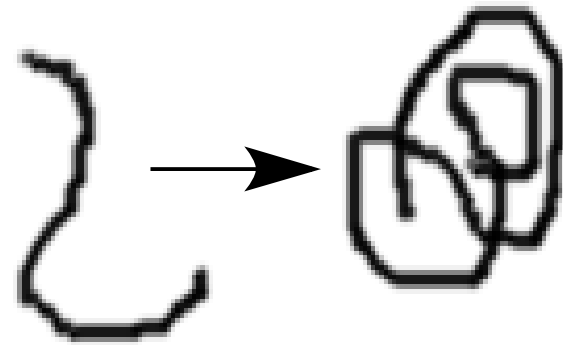
Sickle Cell Anemia

Result of a single Glu → Val mutation in hemoglobin



Higher-Level Structures in Proteins

- **Secondary, Tertiary, and Quaternary structures** have to do with how peptides “fold” into 3-dimensional structure.
- Protein chains fold into very specific 3D structures that are finely tuned to their functions.



Interactions that Stabilize 3D Protein Structure

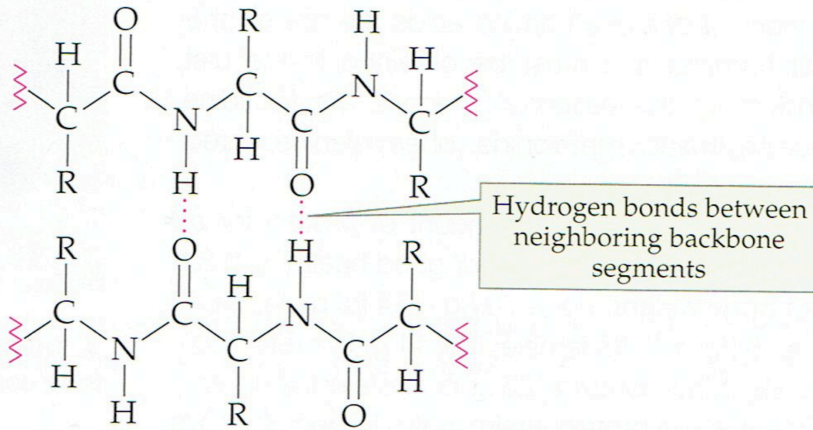
Noncovalent

- Hydrogen bonding
- Ionic interactions
- Hydrophobic interactions

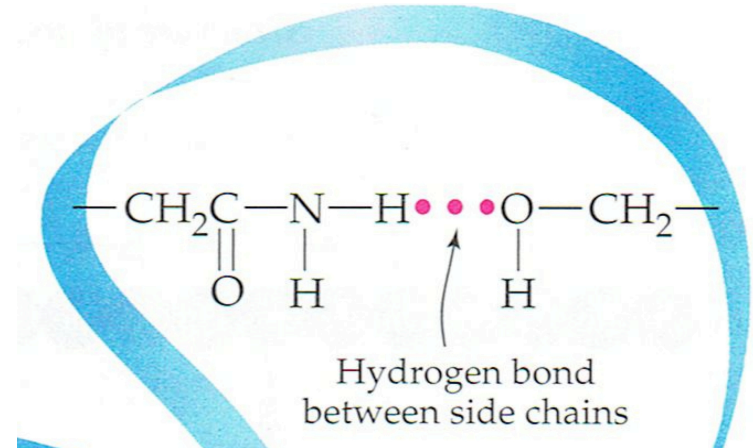
Covalent

- Disulfide bonds

Hydrogen Bonding

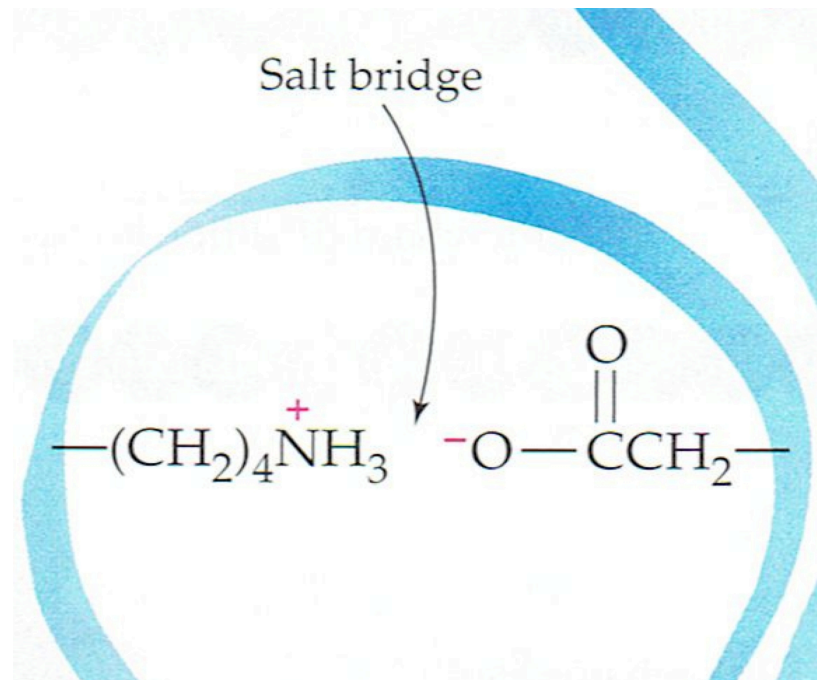


Along backbone



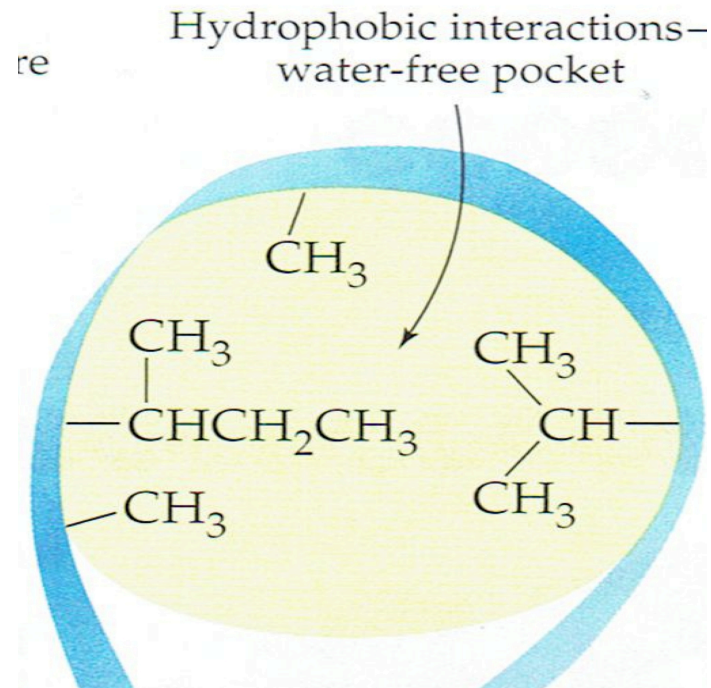
Between side chains

Ionic Interactions (salt bridge)



Between ionized acidic and basic side chains

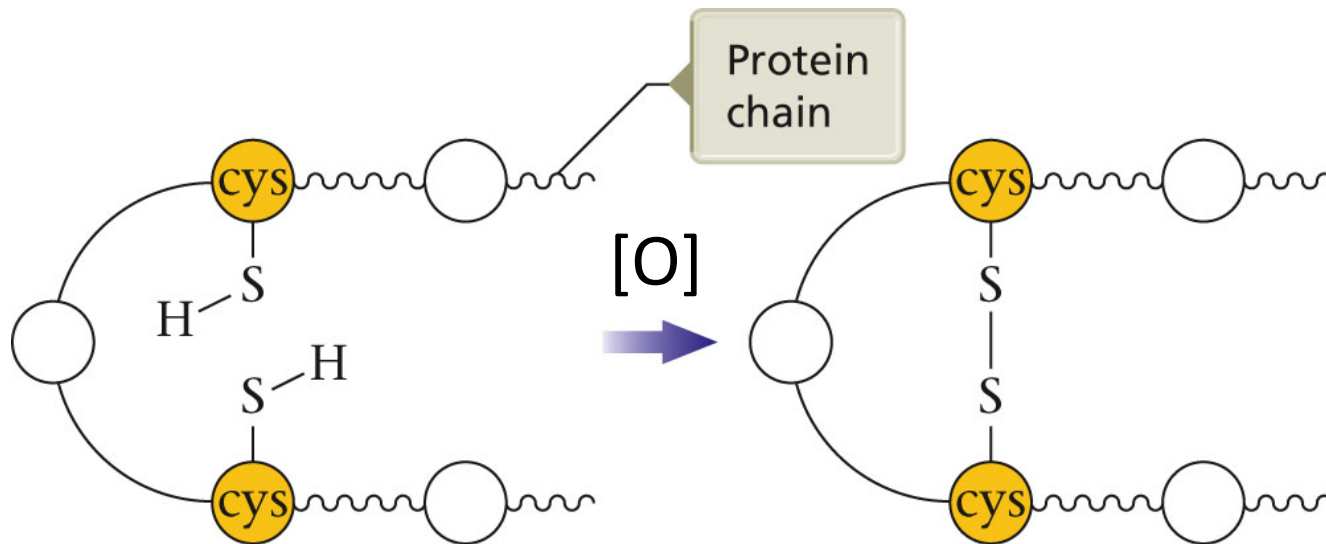
Hydrophobic Interactions



Between nonpolar side chains

Disulfide Bond (S-S bond)

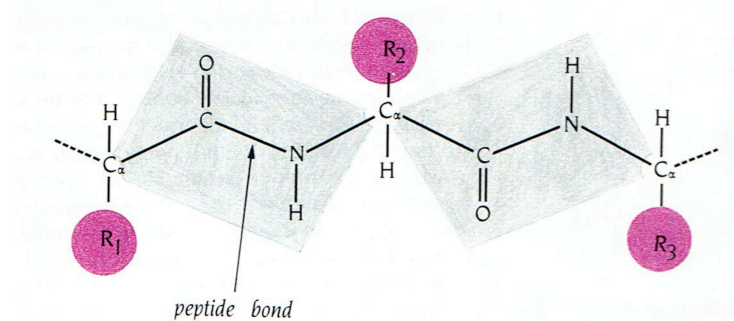
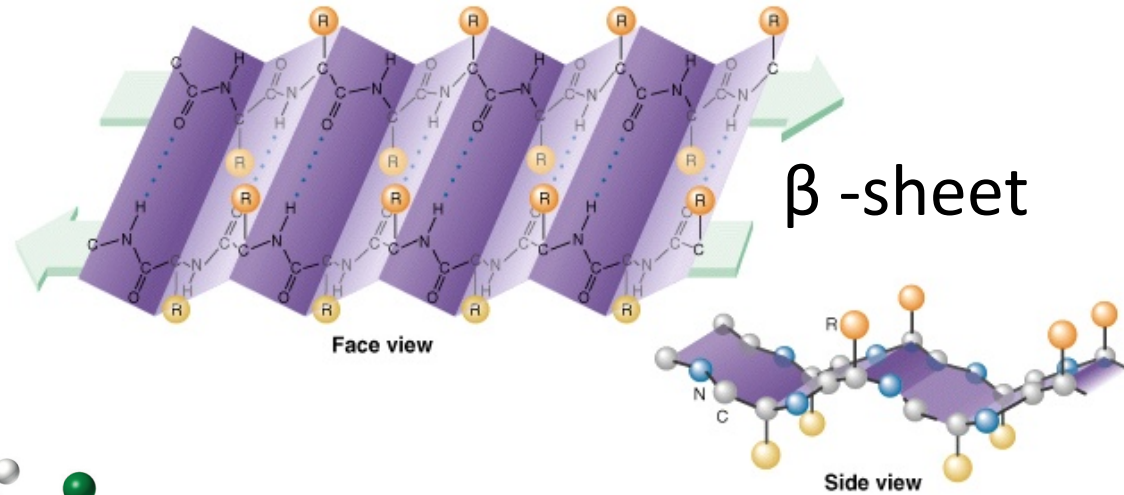
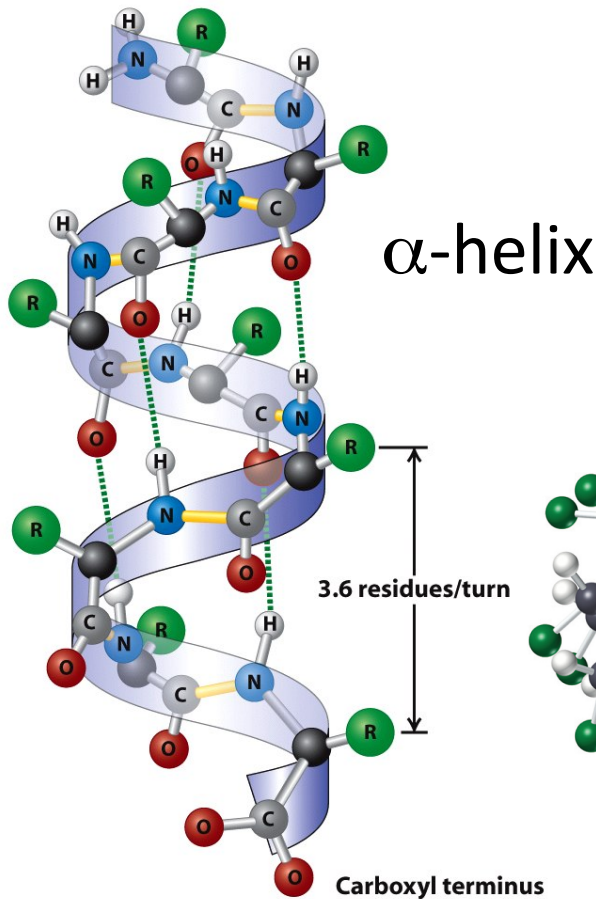
- **Disulfide Bond:** Covalent linkage between two cysteine side chains



Secondary Structure

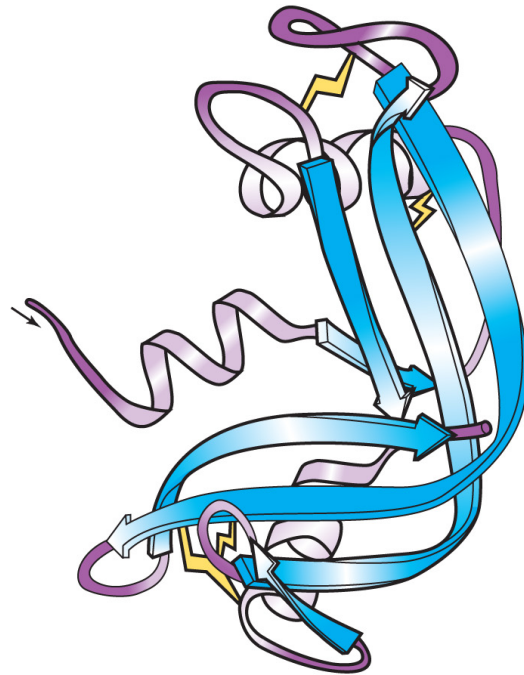
Secondary structure: regular 3D arrangement of polypeptide backbone held together by hydrogen bonding between backbone atoms, plus loops or coils

Amino terminus



Tertiary Structure of Proteins

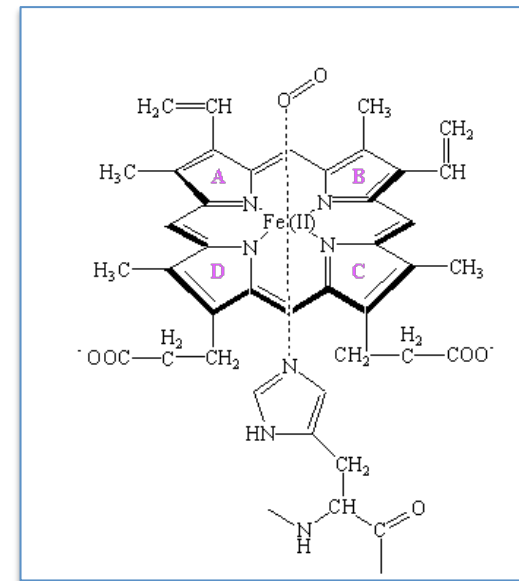
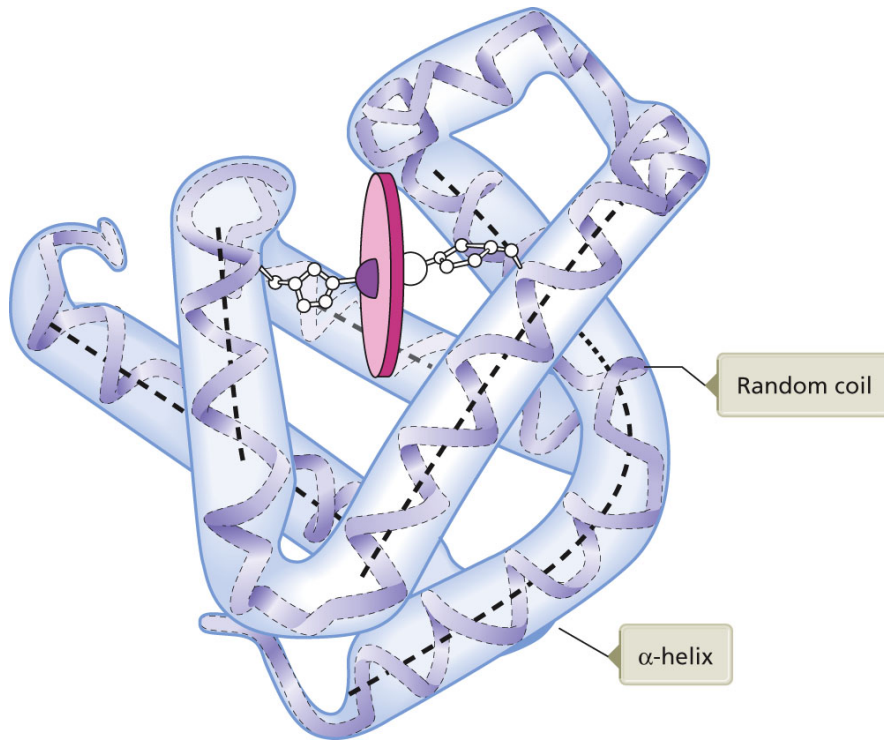
- **Tertiary structure**: the overall 3-dimensional shape that results from folding of the protein; mainly dependent on side chain interactions, often far apart on polypeptide chain



Ribonuclease

Ribonuclease

Tertiary Structure



heme

Myoglobin (a conjugated protein)

Simple vs. Conjugated Proteins

- Simple proteins: Contain only amino acids
 - Eg. Ribonuclease
- Conjugated proteins: Have non-amino acid component(s)
 - Eg. myoglobin

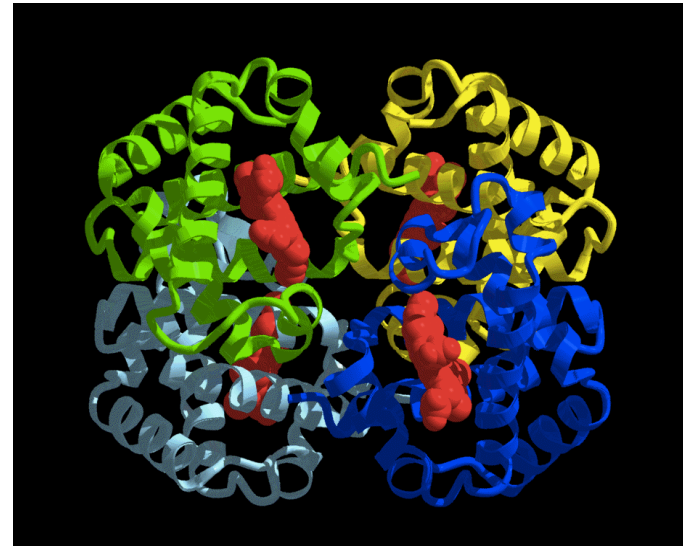
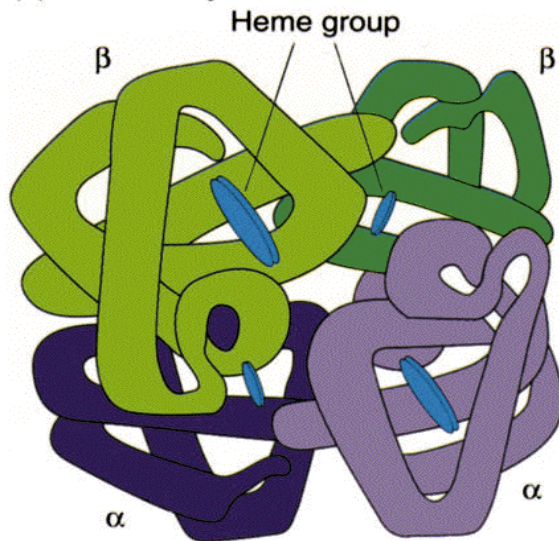
Examples of Conjugated Proteins

TABLE 18.5 Some Examples of Conjugated Proteins

Class of Protein	Nonprotein Part	Examples
Glycoproteins	Carbohydrates	Glycoproteins in cell membranes (Section 21.9)
Lipoproteins	Lipids	High- and low-density lipoproteins that transport cholesterol and other lipids through the body (Section 24.2)
Metalloproteins	Metal ions	The enzyme cytochrome oxidase, necessary for biological energy production, and many other enzymes
Phosphoproteins	Phosphate groups	Milk casein, which provides essential nutrients to infants
Hemoproteins	Heme	Hemoglobin (transports oxygen) and myoglobin (stores oxygen)
Nucleoproteins	RNA (ribonucleic acid)	Found in cell ribosomes, where they take part in protein synthesis

Quaternary Structure

- Quaternary structure: two or more folded protein chains (subunits) held together by noncovalent interactions

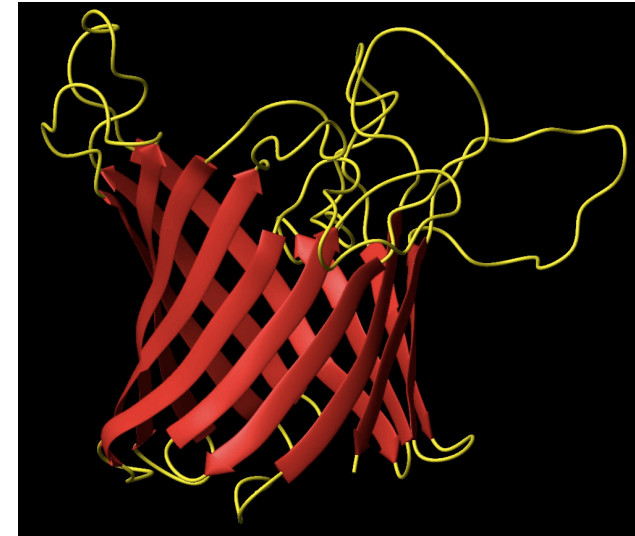
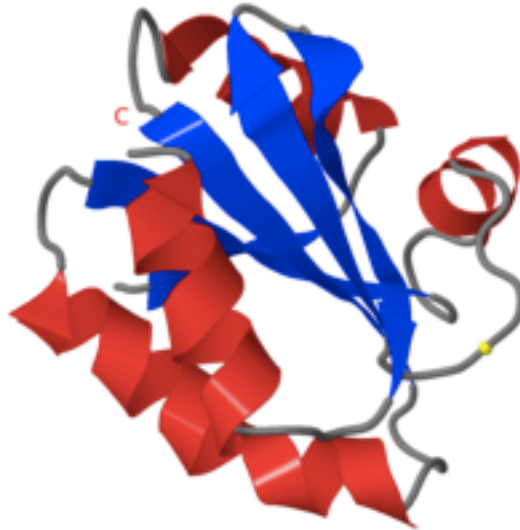
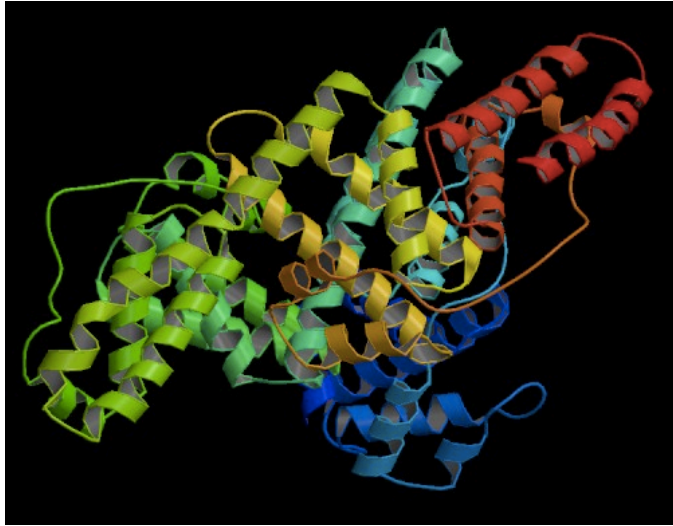


Hemoglobin: O₂ carrier in red blood cells

Broad Categories of Protein Shapes

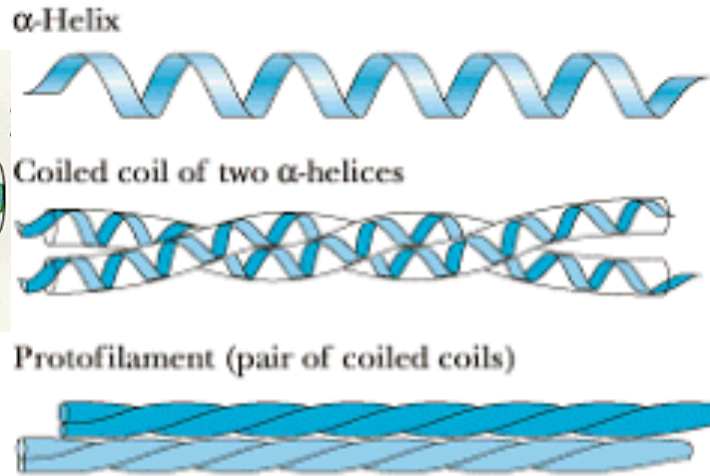
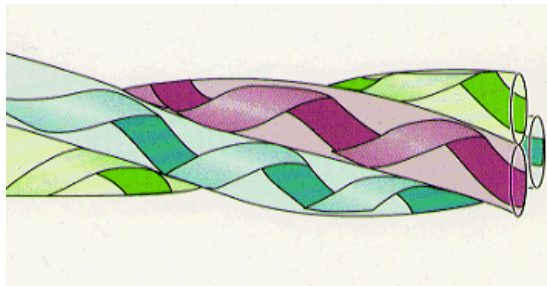
- **Globular** – roughly spherical; “worker” molecules with many different functions; typically water-soluble
- **Fibrous** – tough, water-insoluble, composed of fibers and sheets; give structural integrity and strength; main components of muscle, hair, cartilage, etc.

Structures of Globular Proteins



- Protein structure is optimized for protein function:
- Solubility (polar and nonpolar side chains)
 - Binding sites

Structures of Fibrous Proteins

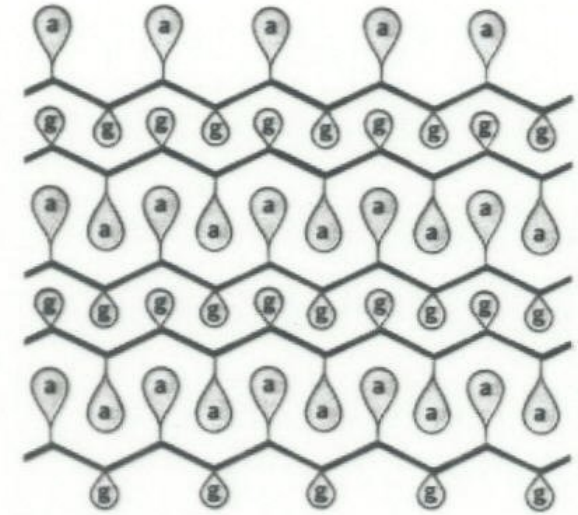


Collagen

- Connective tissues like cartilage, ligament, tendons, skin
- Triple helix cable

Keratin

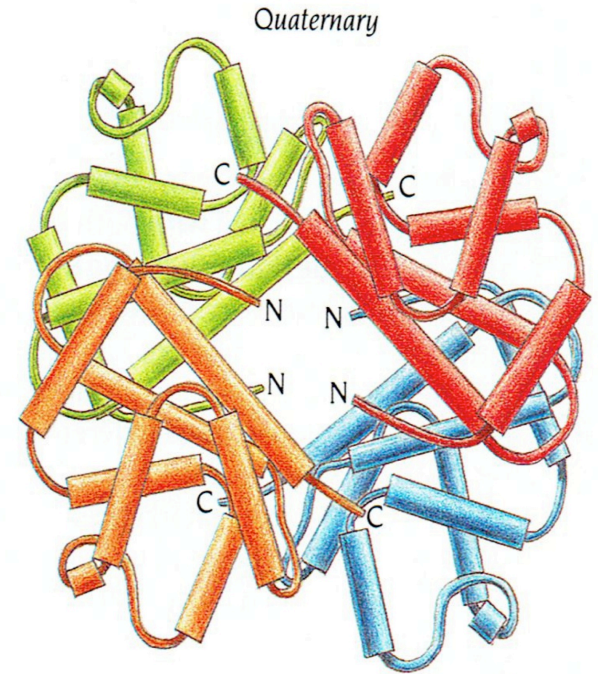
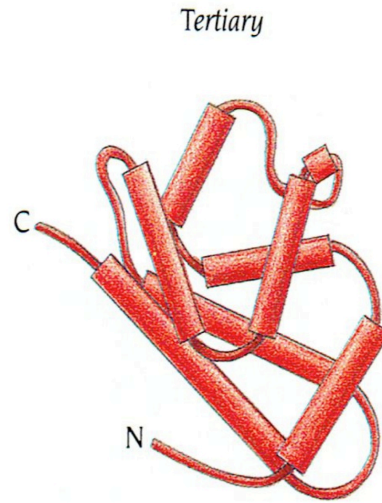
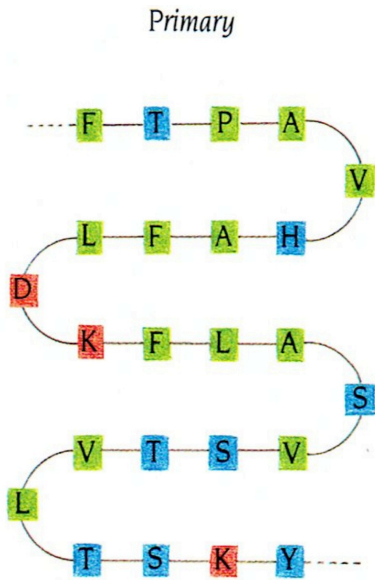
- Hair, horn, nail
- Helix of helices



Silk

- Stacked layers of pleated sheets

Levels of Protein Structure



Primary

Secondary

Tertiary

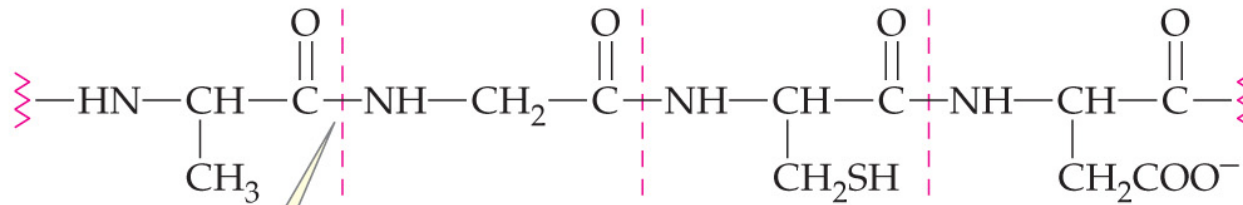
Quaternary

Example Problem: Identify level of structure

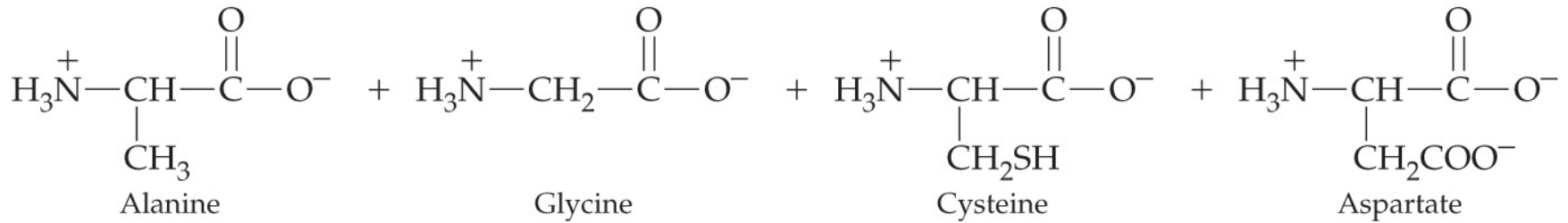
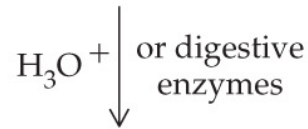
Q: Identify as secondary, tertiary, or quaternary structure.

1. The protein α -hemolysin is made of 7 subunits associated together.
2. Polypeptide backbone hydrogen-bond to make a β -sheet.
3. In the protein albumin, α -helices pack together with all the nonpolar residues in the interior.

Protein Hydrolysis

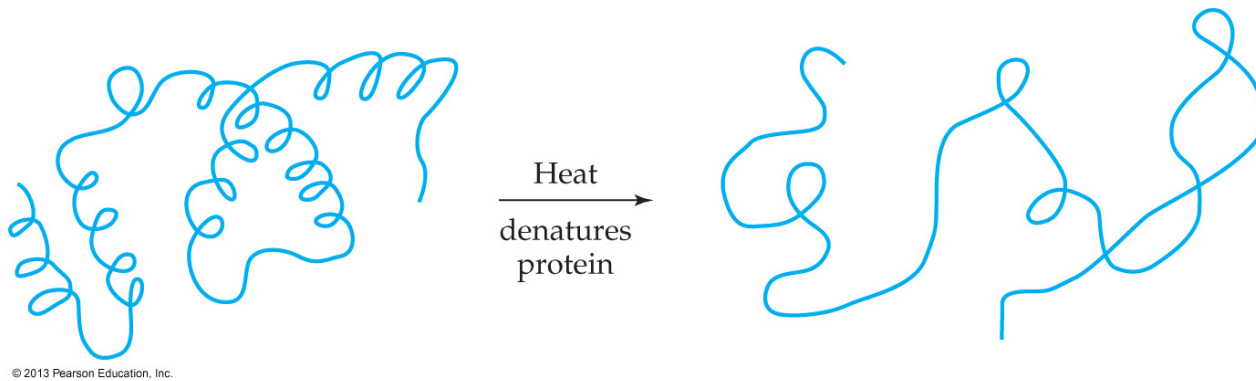


Peptide bond
broken by
hydrolysis



Protein Denaturation

- **Denaturation:** loss of secondary, tertiary, and quaternary protein structure (protein “unfolding”)
 - Primary structure is intact.
 - Protein function lost.



Protein Denaturing Agents

- **Heat**—Disrupts side-chain interactions (eg. frying egg, cooking collagen into gelatin)
- **Mechanical agitation**—(eg. beating egg whites)
- **Detergents**—Disrupt hydrophobic interactions.
- **Organic compounds**—Polar solvents interfere with hydrogen bonding; nonpolar solvents disrupt hydrophobic interactions.
- **pH change**—Disrupt salt bridges
- **Inorganic salts**—High concentrations can disturb salt bridges.

Most denaturation is irreversible.