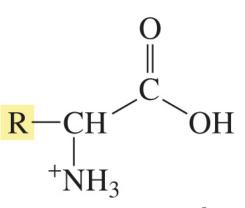
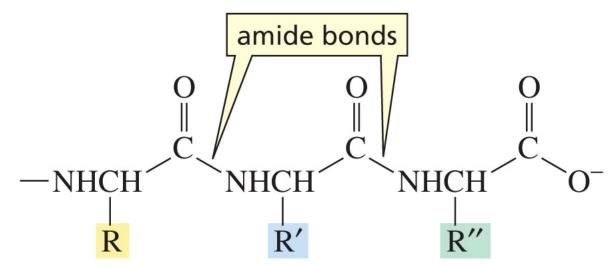
### 01. Amino Acids

### **Biomolecules**

- Protein
- Carbohydrate
- Nucleic acid
- Lipid



a protonated  $\alpha$ -aminocarboxylic acid an amino acid



amino acids are linked together by amide bonds

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peptide ---- protein

di-, tri-, oligo-

#### **Table 22.1 Examples of the Many Functions of Proteins in Biological Systems** Structural proteins These proteins impart strength to biological structures or protect organisms from their environment. For example, collagen is the major component of bones, muscles, and tendons; keratin is the major component of hair, hooves, feathers, fur, and the outer layer of skin. Protective proteins Snake venoms and plant toxins protect their owners from predators. Blood-clotting proteins protect the vascular system when it is injured. Antibodies and peptide antibiotics protect us from disease. Enzymes Enzymes are proteins that catalyze the reactions that occur in living systems. Some of the hormones, such as insulin, that Hormones regulate the reactions that occur in living

systems are proteins.

Proteins with physiological functions

These proteins are responsible for physiological functions such as the transport and storage of oxygen in the body, the storage of oxygen in the muscles, and the contraction of muscles.

### fibrous proteins

proteins



globular proteins

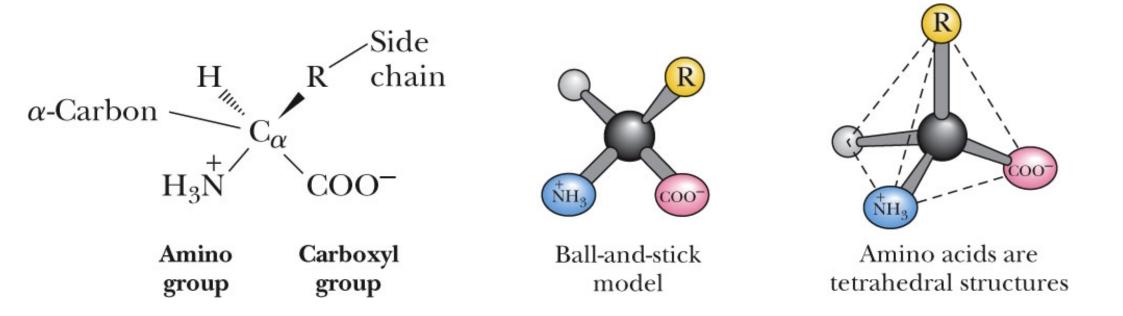


Figure 4.1 Anatomy of an amino acid. Except for proline and its derivatives, all of the amino acids commonly found in proteins possess this type of structure.

Glycine (Gly, G)

Alanine (Ala, A)

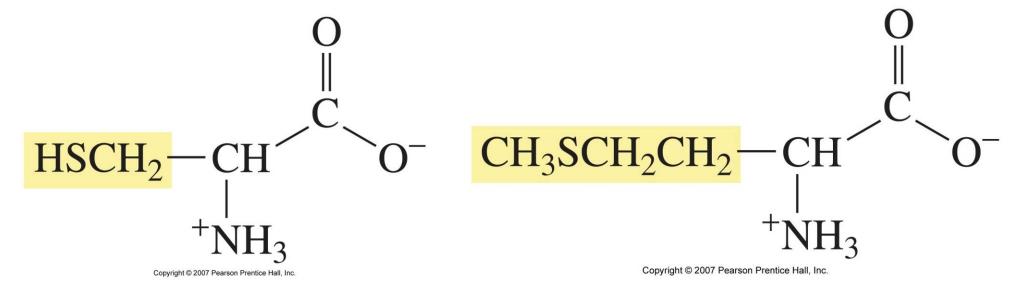
Valine (Val, V)\*

Leucine (Leu, L)\*

Isoleucine (Ile. I)\* 7

### Serine (Ser, S)

### Threonine (Thr, T)\*



Cysteine (Cys, C)

cystine

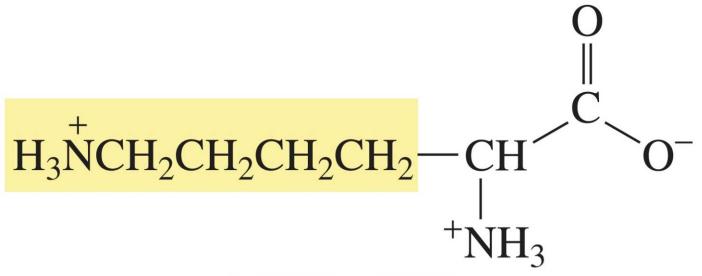
Methionine (Met, M)\*

#### Aspartate (Asp, D)

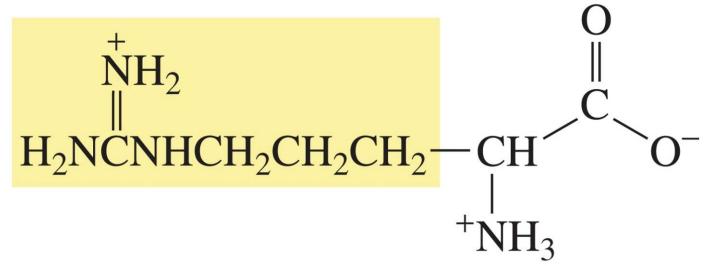
### Glutamate (Glu, E)

Asparagine (Asn, N)

Glutamine (Gln, Q) 9



### Lysine (Lys, K)\*

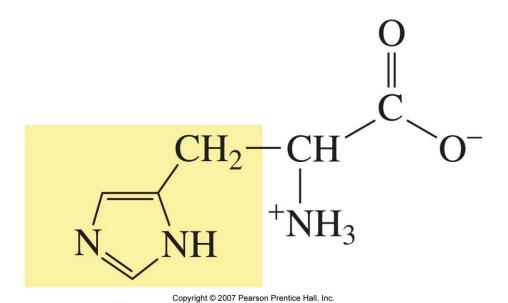


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**Arginine (Arg, R)\*** 

$$\begin{array}{c} O \\ \\ C \\ C \\ C \\ C \\ C \\ C \\ C \\ C \\ C \\ C \\ C$$

### Phenylalanine (Phe, F)\*

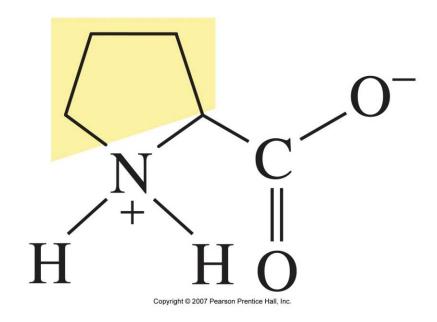


**Histidine (His, H)\*** 

### Tyrosine (Tyr, Y)

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Tryptophan (Trp, W)\*



Proline (Pro, P)

Hydrophobic (A, G, I, L, F, V, P)

Hydrophilic (D, E, R, S, T, C, N, Q, H)

Amphipathic (K, M, W, Y)

Essential amino acids: V, L, I, T, M, K, R, F, H, W



### We'll see some of these in later chapters

- Selenocysteine in many organisms
- Pyrrolysine in several archaeal species
- Hydroxylysine, hydroxyproline collagen
- Carboxyglutamate blood-clotting proteins
- Pyroglutamate in bacteriorhodopsin
- GABA, epinephrine, histamine, serotonin act as neurotransmitters and hormones
- Phosphorylated amino acids a signaling device





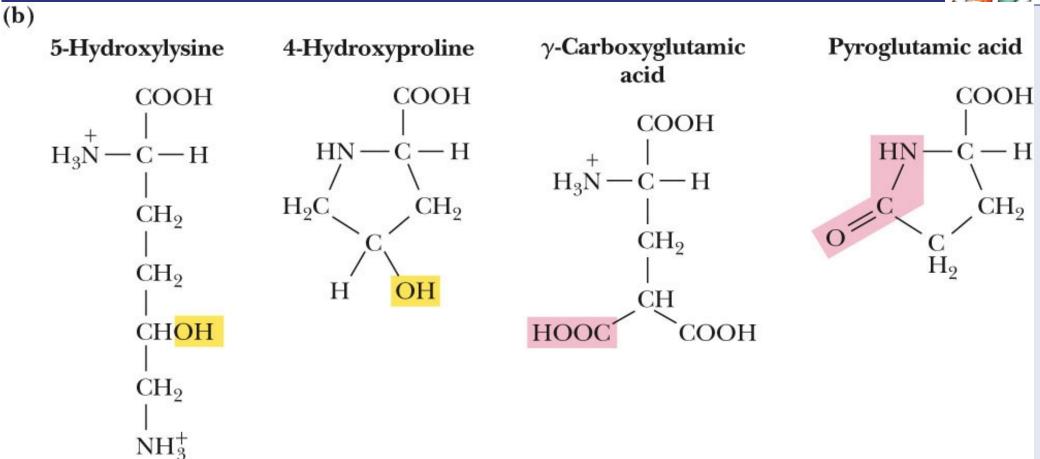
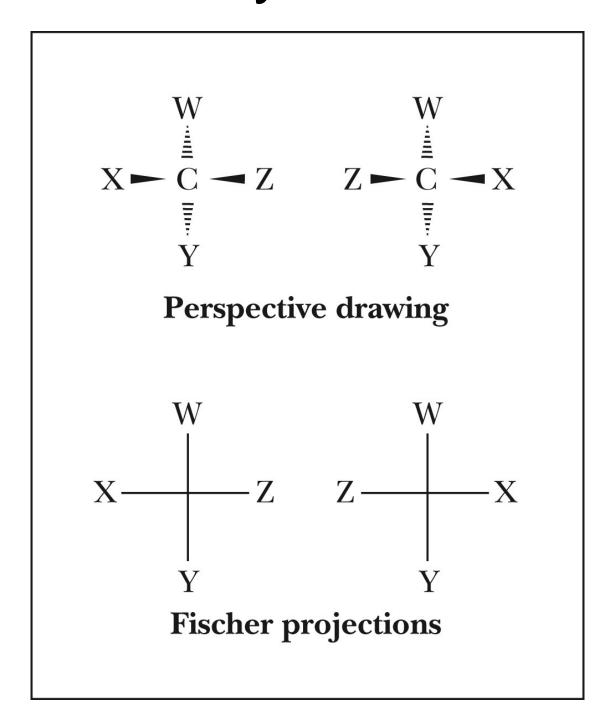


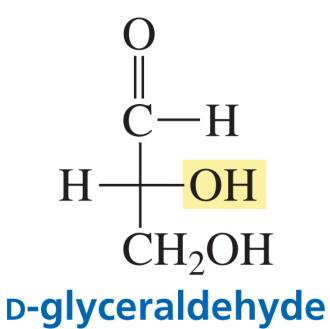
Figure 4.4 (b) Some amino acids are less common, but nevertheless found in certain proteins. Hydroxylysine and hydroxyproline are found in connective-tissue proteins; carboxyglutamate is found in blood-clotting proteins; pyroglutamate is found in bacteriorhodopsin (see Chapter 9).

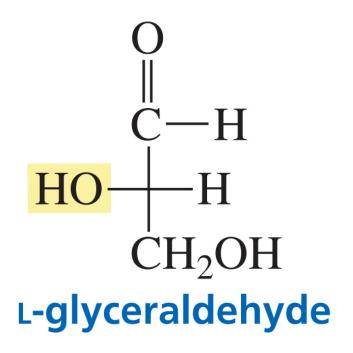


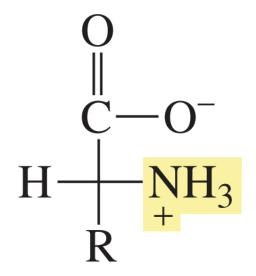
Figure 4.4 (c) Several amino acids that act as neurotransmitters and hormones.

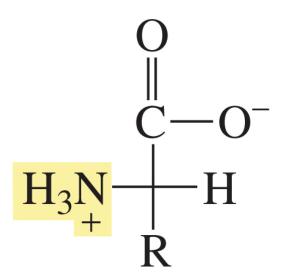
## Stereochemistry of Amino Acids











**D-amino** acid

L-amino acid

## The Murchison Meteorite – Discovery of Extraterrestrial Handedness

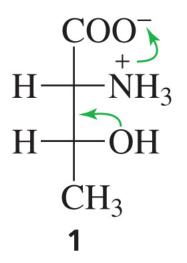


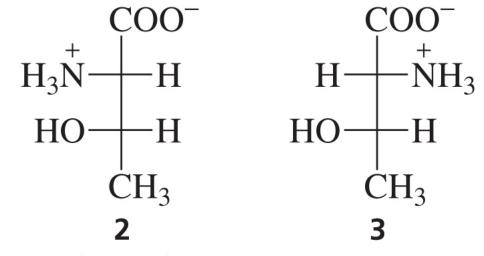
Amino acids found in the Murchison meteorite.

Why do L-amino acids predominate in biological systems? What process might have selected L-amino acids over their D- counterparts?

The meteorite found near Murchison, Australia may provide answers. Certain amino acids found in the meteorite have been found to have L-enantiomeric excesses of 2% to 9%.

### Which is (2*S*,3*R*)-threonine?





 $H_3$ N-H H OH  $CH_3$   $\mathbf{4}$ 

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- Amino Acids are Weak Polyprotic Acids
- The degree of dissociation depends on the pH of the medium

• 
$$H_2A^+ + H_2O \rightarrow HA^0 + H_3O^+$$

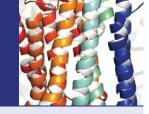
$$K_{a1} = \frac{[HA^0][H_30^+]}{[H_2A^+]}$$



The second dissociation (the amino group in the case of glycine):

• 
$$HA^0 + H_2O \rightarrow A^- + H_3O^+$$

$$K_{a2} = \frac{[A^{-}][H_{3}O^{+}]}{[HA^{0}]}$$



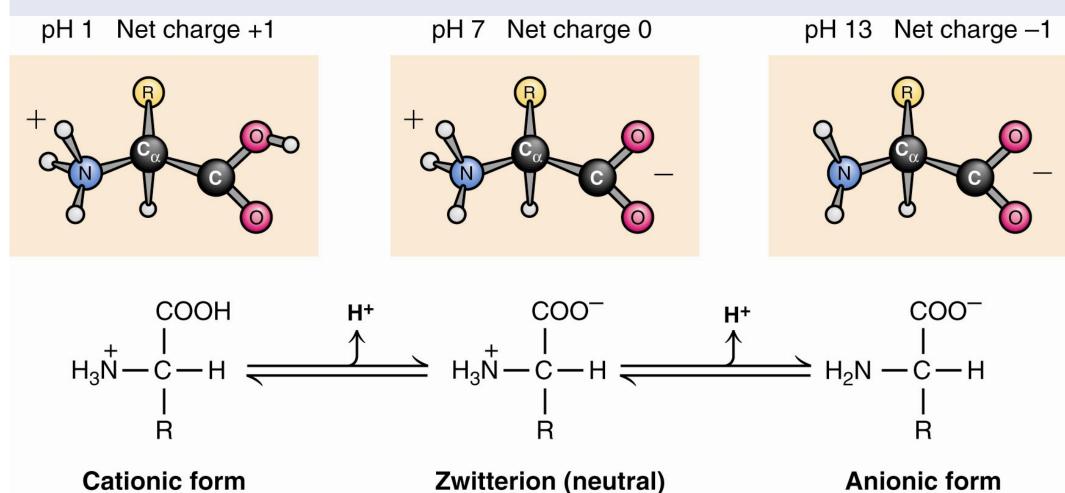
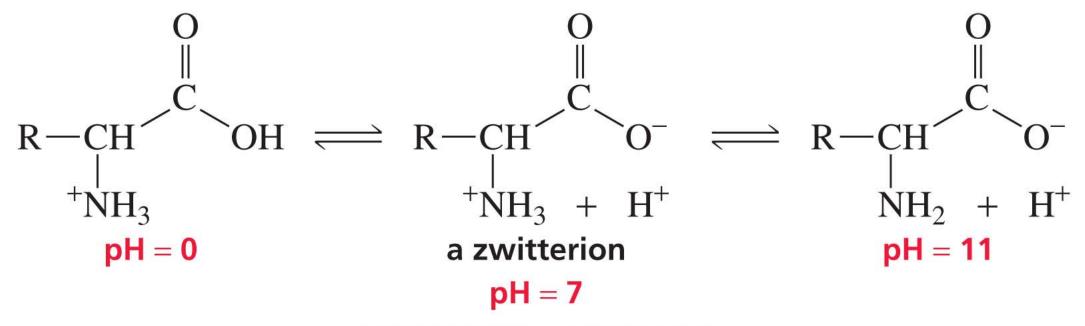
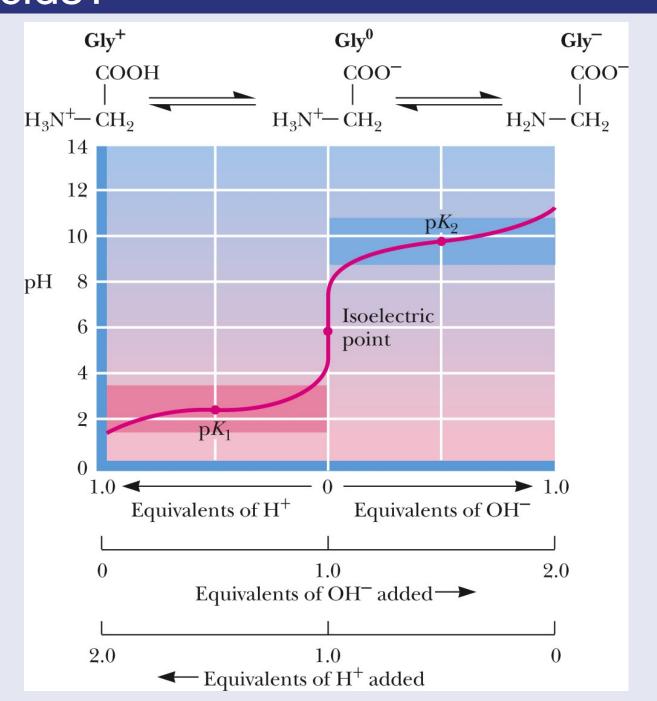


Figure 4.5 The ionic forms of the amino acids, shown without consideration of any ionizations on the side chain.



The adjacent  $\alpha$ -amino group makes the  $\alpha$ -COOH group more acidic.





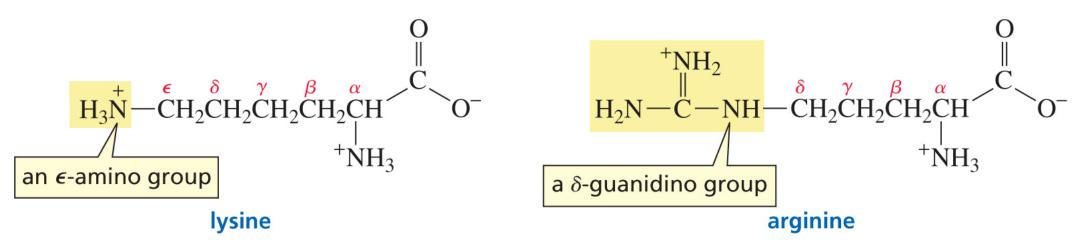
### pK<sub>a</sub> Values of the Amino Acids



## You should know these numbers and know what they mean

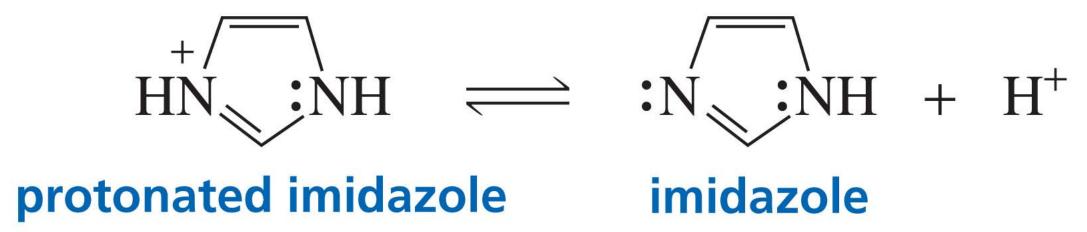
- Alpha carboxyl group  $pK_a = 2$
- Alpha amino group  $pK_a = 9$
- These numbers are approximate, but entirely suitable for our purposes.

Table 22.3 The pK <sub>a</sub> Values of Amino Acids			
Amino acid	$pK_a lpha$ -COOH	${}_{m{lpha}}^{m{K_a}}$ ${}_{m{lpha}}^{m{NH_3}^+}$	p $K_a$ side chain
Alanine	2.34	9.69	_
Arginine	2.17	9.04	12.48
Asparagine	2.02	8.84	_
Aspartic acid	2.09	9.82	3.86
Cysteine	1.92	10.46	8.35
Glutamic acid	2.19	9.67	4.25
Glutamine	2.17	9.13	_
Glycine	2.34	9.60	_
Histidine	1.82	9.17	6.04
Isoleucine	2.36	9.68	_
Leucine	2.36	9.60	_
Lysine	2.18	8.95	10.79
Methionine	2.28	9.21	_
Phenylalanine	2.16	9.18	_
Proline	1.99	10.60	_
Serine	2.21	9.15	_
Threonine	2.63	9.10	_
Tryptophan	2.38	9.39	_
Tyrosine	2.20	9.11	10.07
Valine	2.32	9.62	_



#### Protonated at physiological pH

$$pKa = 6.0$$



### pK<sub>a</sub> Values of the Amino Acids



## You should know these numbers and know what they mean

- Arginine, Arg, R: pK<sub>a</sub>(guanidino group) = 12.5
- Aspartic Acid, Asp, D: pK<sub>a</sub> = 3.9
- Cysteine, Cys, C: pK<sub>a</sub> = 8.3
- Glutamic Acid, Glu, E: pK<sub>a</sub> = 4.3
- Histidine, His, H:  $pK_a = 6.0$

### pK<sub>a</sub> Values of the Amino Acids



## You should know these numbers and know what they mean

- Lysine, Lys, K: pK<sub>a</sub> = 10.5
- Serine, Ser, S: pK<sub>a</sub> = 13
- Threonine, Thr, T: pK<sub>a</sub> = 13
- Tyrosine, Tyr, Y: pK<sub>a</sub> = 10.1

### Another Sample Calculation



What is the pH of a lysine solution if the side chain amino group is 3/4 dissociated?

$$pH = 10.5 + \log_{10} \frac{[3]}{[1]}$$

- pH = 10.5 + (0.477)
- pH = 10.977 = 11.0
- Note that, when the group is ¾ dissociated, ¾ is dissociated and ¼ is not; thus the ratio in the log term is ¾ over ¼ or 3/1.

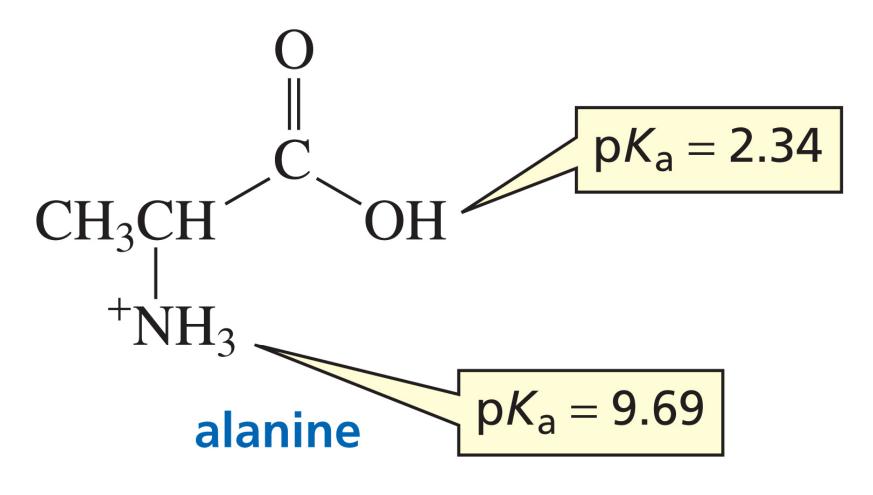
$$K_{a} = \frac{[H_{3}O^{+}][A^{-}]}{[HA]} = K_{eq}[H_{2}O]$$

$$pK_a = -log K_a$$
  
the Henderson–Hasselbalch equation

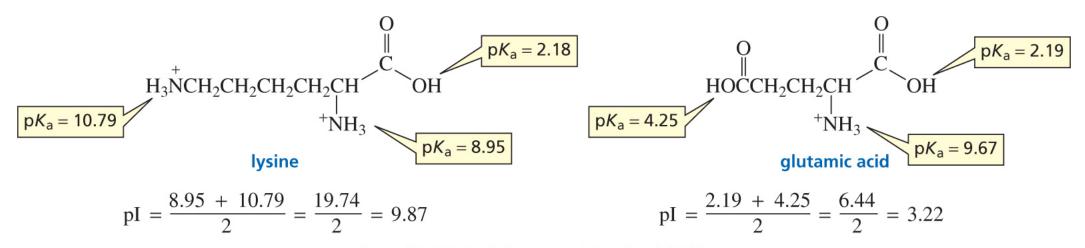
$$pK_a = pH + log \frac{[HA]}{[A^-]}$$

If pH < p $K_a$ , acidic form If pH > p $K_a$ , basic form

#### Isoelectric point (IP): no net charge



$$pI = \frac{2.34 + 9.69}{2} = \frac{12.03}{2} = 6.02$$



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### Reactions of Amino Acids

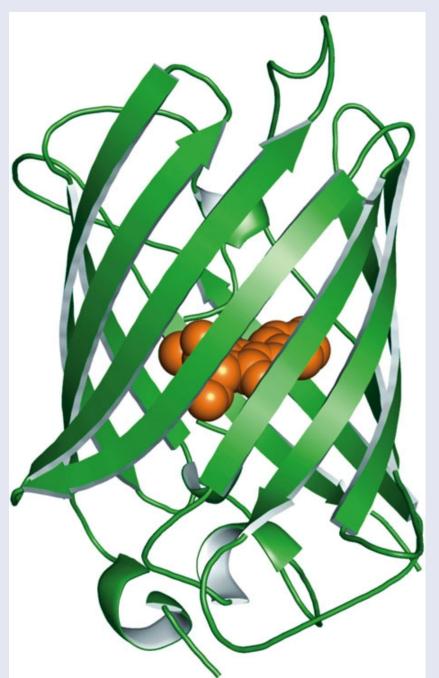


- Carboxyl groups form amides & esters
- Amino groups form Schiff bases and amides
- Edman reagent (phenylisothiocyanate) reacts with the  $\alpha$ -amino group of an amino acid or peptide to produce a phenylthiohydantoin (PTH) derivative.
- Side chains show unique reactivities
  - Cys residues can form disulfides and can be easily alkylated
  - Few reactions are specific to a single kind of side chain

### Green Fluorescent Protein



A jellyfish (Aequorea victoria) native to the northwest Pacific Ocean contains a green fluorescent protein. GFP is a naturally fluorescent protein. Genetic engineering techniques can be used to "tag" virtually any protein, structure, or organelle in a cell. The GFP chromophore lies in the center of a  $\beta$  -barrel protein structure.



## Green Fluorescent Protein

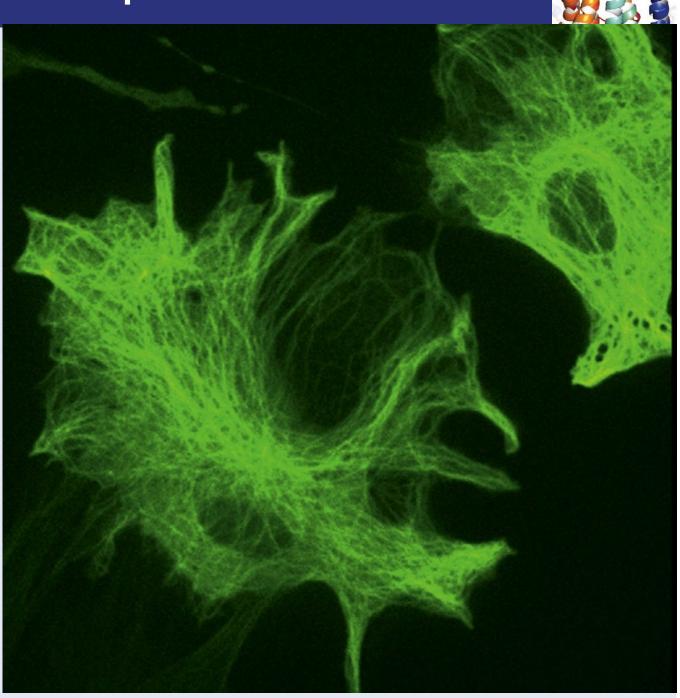


Phe-Ser-Tyr-Gly-Val-Gln 
$$O_2$$
  $O_2$   $O_2$   $O_3$   $O_4$   $O_4$   $O_4$   $O_5$   $O_4$   $O_5$   $O_6$   $O_7$   $O_8$   $O_8$ 

The prosthetic group of GFP is an oxidative product of the sequence –FSYGVQ-.

## Yellow fluorescent protein

Amino acid substitutions in GFP can tune the color of emitted light. Shown here is an image of African green monkey kidney cells expressing yellow fluorescent protein (YFP) fused to  $\alpha$  tubulin, a cytoskeletal protein.



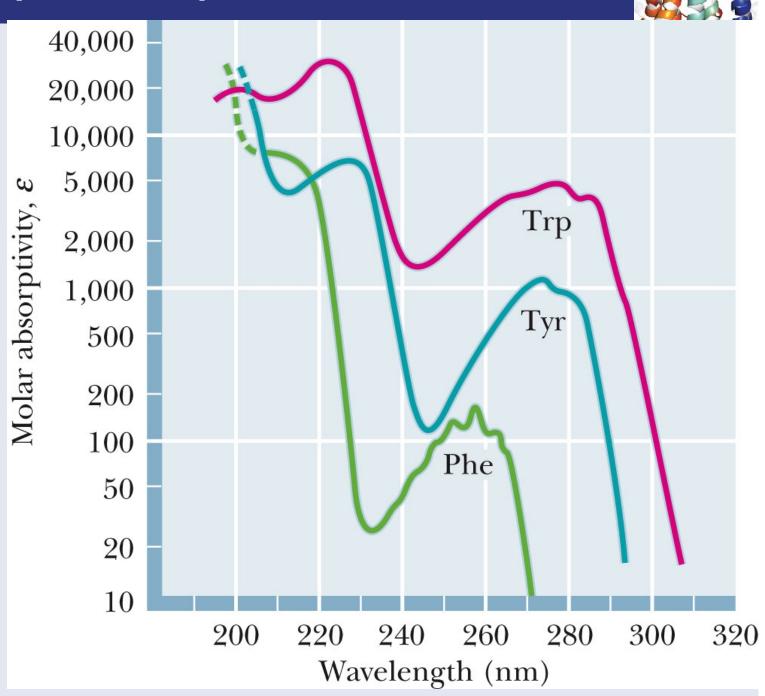
## Spectroscopic Properties



- All amino acids absorb at infrared wavelengths
- Only Phe, Tyr, and Trp absorb UV
- Absorbance at 280 nm is a good diagnostic device for amino acids
- NMR spectra are characteristic of each residue in a protein, and high resolution NMR measurements can be used to elucidate three-dimensional structures of proteins

## Spectroscopic Properties

Figure 4.10 The UV spectra of the aromatic amino acids at pH 6.



## Spectroscopic Properties



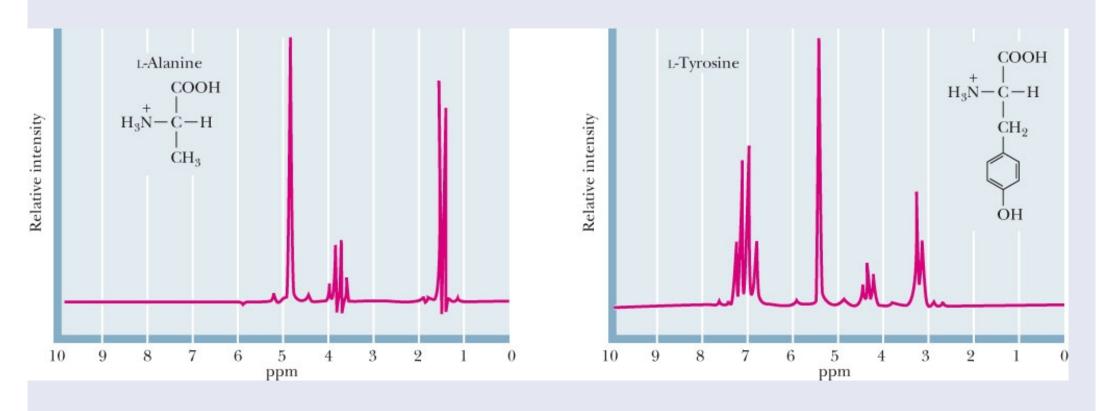
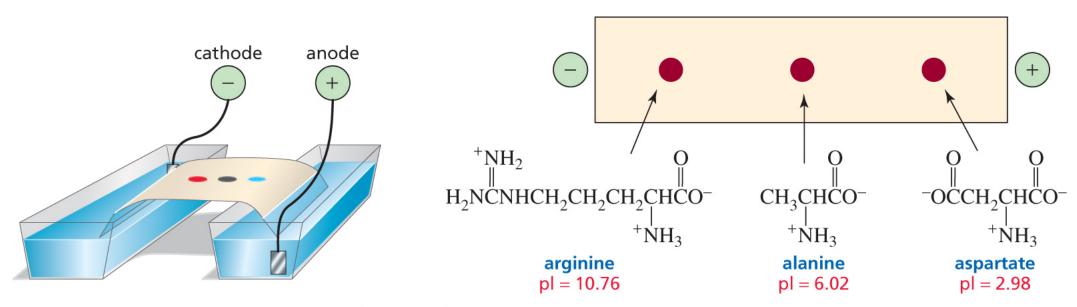


Figure 4.11 Proton NMR spectra of several amino acids.

### Separation of AA

# **Analytical separation Preparative separation**

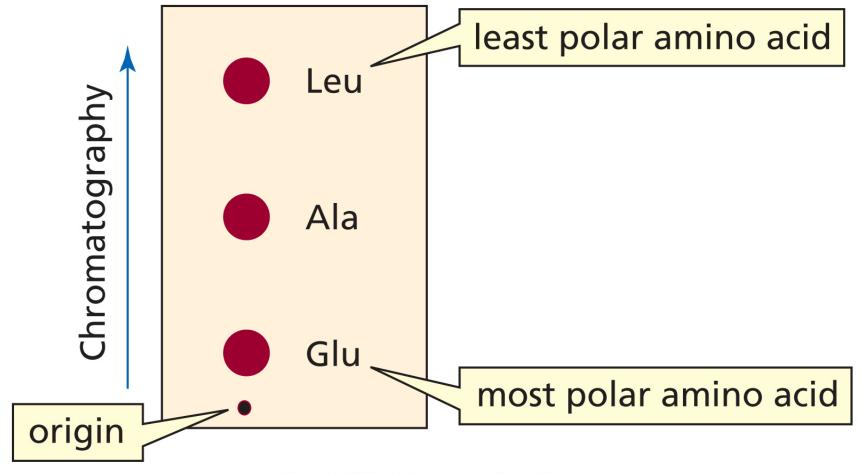
#### Electrophoresis (based on charge)



#### Ninhydrin test (AA is purple)

#### Paper Chromatography and Thin-Layer Chromatography

#### (based on polarity)



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#### Solvent: H<sub>2</sub>O/AcOH/BuOH

MIT 5.301 Chemistry Laboratory Techniques http://www.youtube.com/watch?v=EUn2skAAjHk

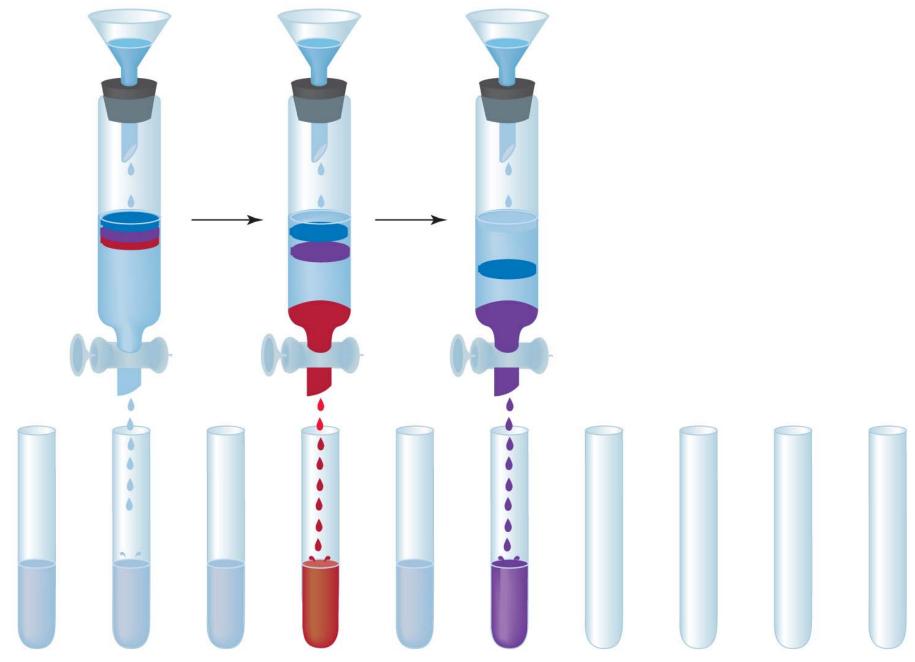
#### **Ion-Exchange Chromatography**

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Cation-exchange resin

Nonpolar nature of the column caused it to retain nonpolar amino acids longer than polar amino acids.

49

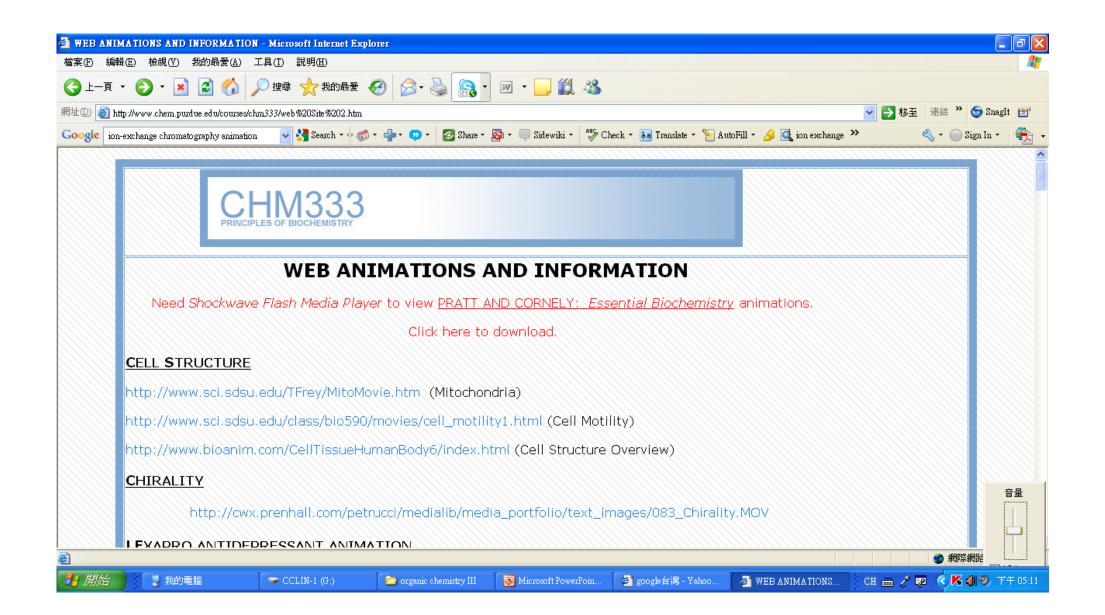


Fractions sequentially collected

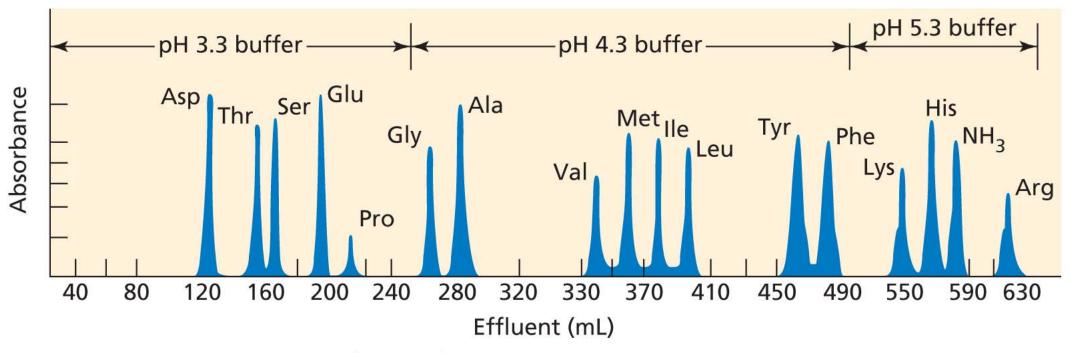
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#### http://www.chem.purdue.edu/courses/chm333/web%20Site%202.htm



#### After addition of ninhydrin

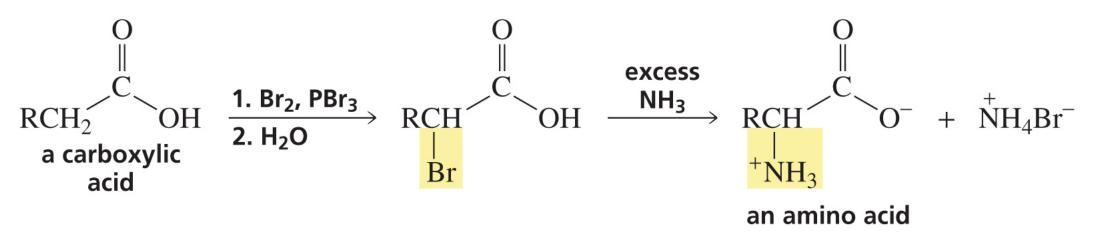


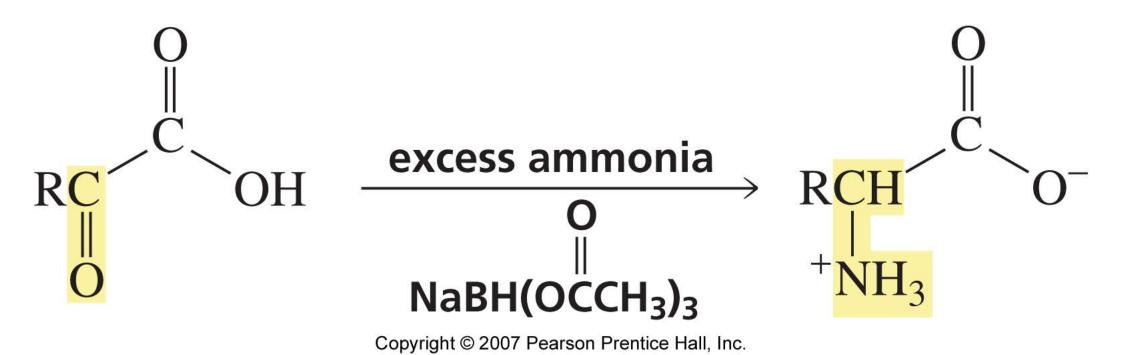
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At 570 nm

#### **Synthesis of Amino acids**

#### Hell-Volhard-Zelinski reaction





$$C_{2}H_{5}OC-CH-COC_{2}H_{5} + N: K^{+} \longrightarrow N-C-COC_{2}H_{5}$$

$$\alpha\text{-bromomalonic ester} \quad \text{potassium phthalimide} \quad N\text{-phthalimidomalonic ester}$$

$$COCC_{2}H_{5} \longrightarrow CH_{3}OH_{2}OC$$

$$CH_{3}OH_{2}OCC_{2}H_{5} \longrightarrow CCC_{2}CCC_{2}H_{5} \longrightarrow CCC_{2}CCC_{2}H_{5} \longrightarrow CCC_{2}CCC_{2}CCC_{2}H_{5} \longrightarrow CCC_{2}C$$

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an amino acid

phthalic acid

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acetic acid

an amino acid

#### Strecker synthesis

#### Kinetic resolution

In **kinetic resolution**, two <u>enantiomers</u> show different <u>reaction</u> rates in a <u>chemical reaction</u>, thereby creating an excess of the less reactive enantiomer. This excess goes through a maximum and disappears on full completion of the reaction. Kinetic resolution is a very old concept in <u>organic chemistry</u> and can be used in the <u>organic synthesis</u> of chiral molecules. It has been surpassed by other methods.

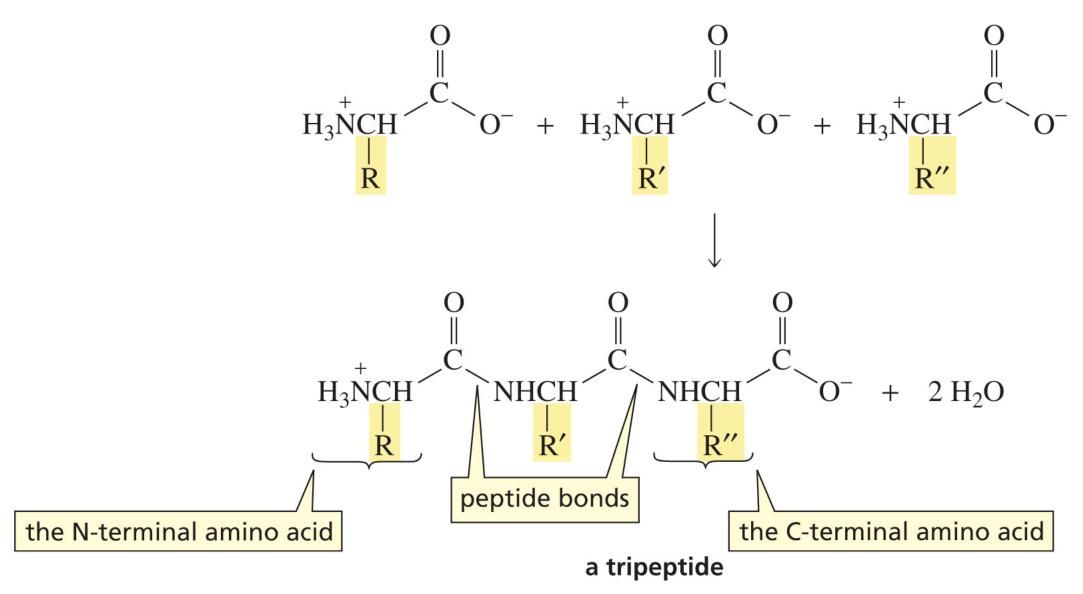
$$\begin{array}{c} O \\ H_2NCH \\ R \\ D\text{-amino acid} \\ L\text{-amino acid} \\ \\ N\text{-acetyl-L-amino acid} \\ \\ N\text{-acetyl-L-amino acid} \\ \\ N\text{-acetyl-L-amino acid} \\ \\ N\text{-acetyl-D-amino acid} \\ N\text{-acetyl-D$$

# 4.7 What is the Fundamental Structural Pattern in Proteins?



- Proteins are unbranched polymers of amino acids
- Amino acids join head-to-tail through formation of covalent peptide bonds
- Peptide bond formation results in release of water
- The peptide backbone of a protein consists of the repeated sequence –N-C<sub>α</sub>-C<sub>o</sub>-
- "N" is the amide nitrogen of the amino acid
- "C<sub>α</sub>" is the alpha-C of the amino acid
- "C<sub>o</sub>" is the carbonyl carbon of the amino acid

#### peptide bond (amide bond)



Glu, Cys, His, Val, Ala

the pentapeptide contains the indicated amino acids, but their sequence is not known

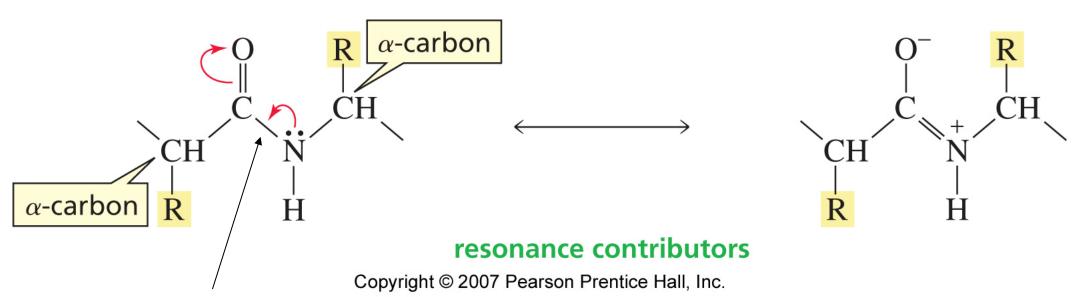
Val-Cys-Ala-Glu-His

the amino acids in the pentapeptide have the indicated sequence

## The Peptide Bond



- Is usually found in the trans conformation
- Has partial (40%) double bond character
- Is about 0.133 nm long shorter than a typical single bond but longer than a double bond
- Due to the double bond character, the six atoms of the peptide bond group are always planar
- N partially positive; O partially negative



40% double-bond character

# 4.7 What is the Fundamental Structural Pattern in Proteins?



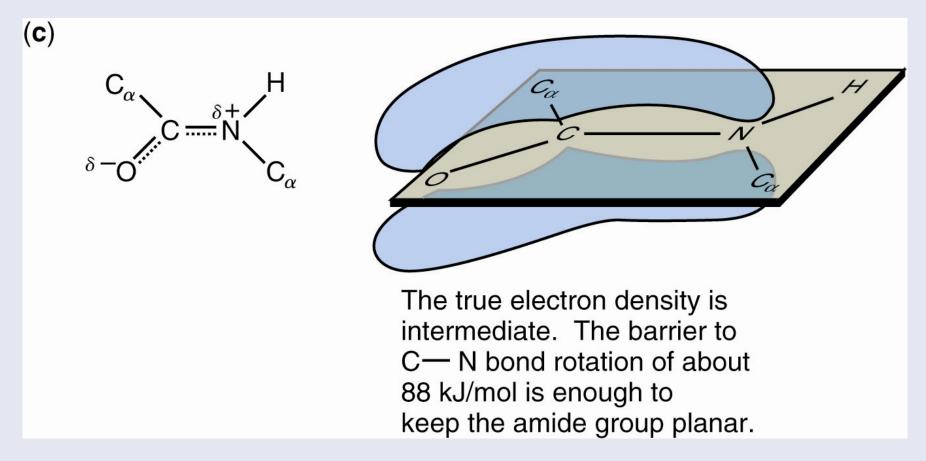


Figure 4.16 (c) The peptide bond is best described as a resonance hybrid of the forms shown on the two previous slides.

## The Peptide Bond



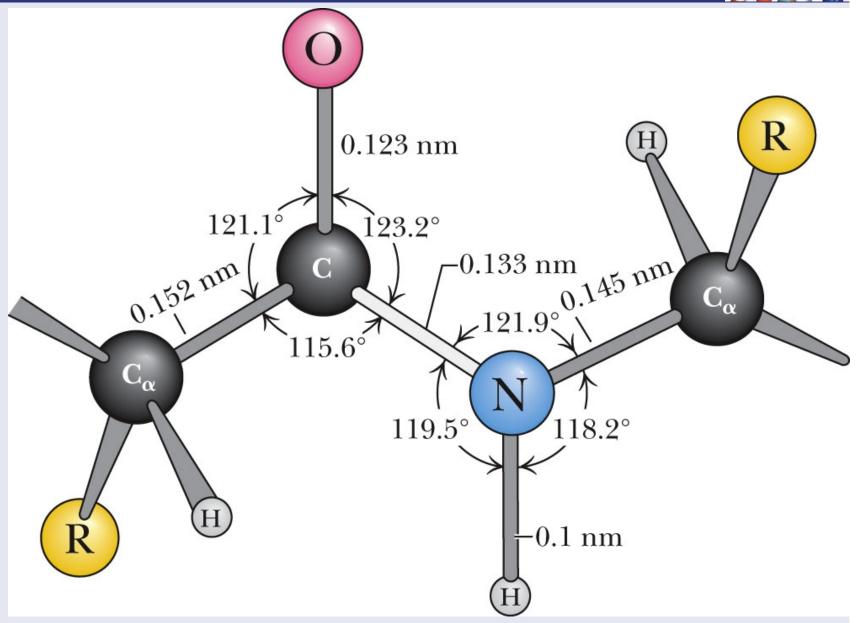
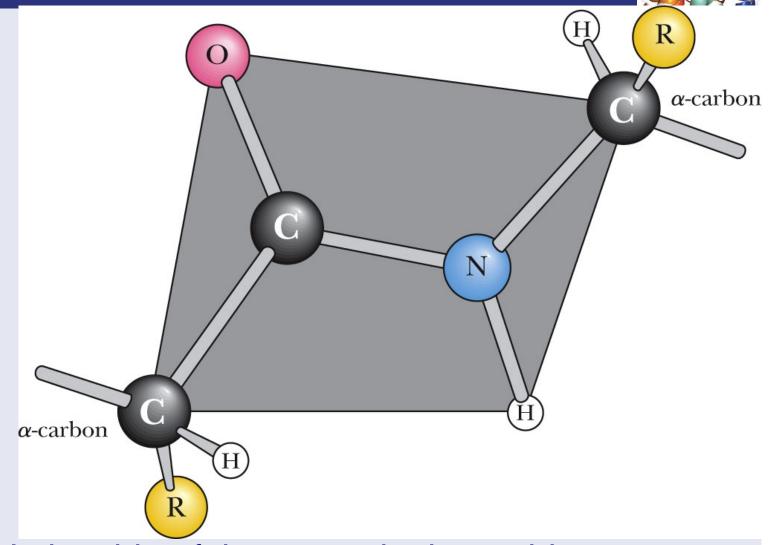
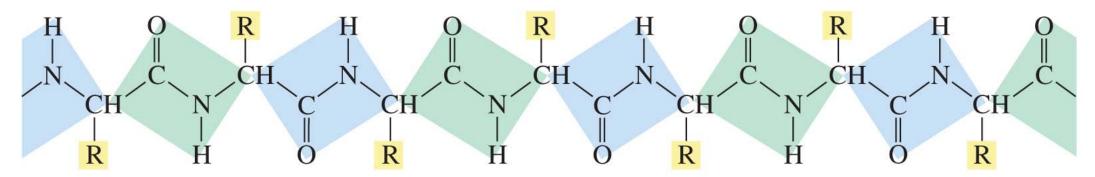


Figure 4.15 The *trans* conformation of the peptide bond.

# 4.7 What is the Fundamental Structural Pattern in Proteins?



The coplanar relationship of the atoms in the amide group is highlighted here by an imaginary shaded plane lying between adjacent  $\alpha$ -carbons.



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2 R—SH a thiol

mild oxidation

 $Br_2$  (or  $I_2$ )

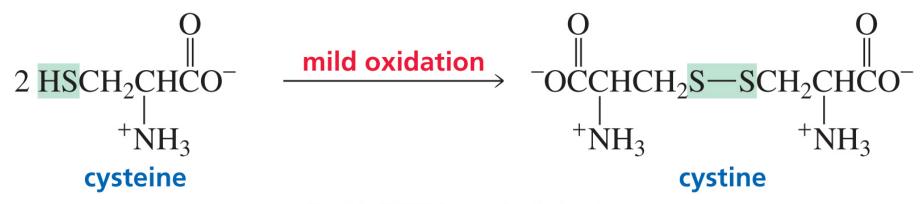
RS—SR a disulfide

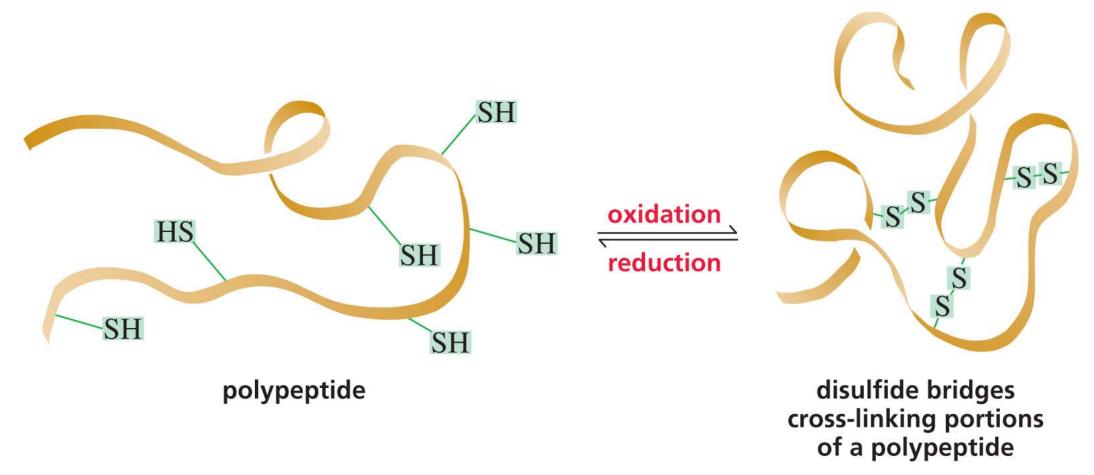
#### mechanism for oxidation of a thiol to a disulfide

RS—SR a disulfide



2 R—SH a thiol





## "Peptides"



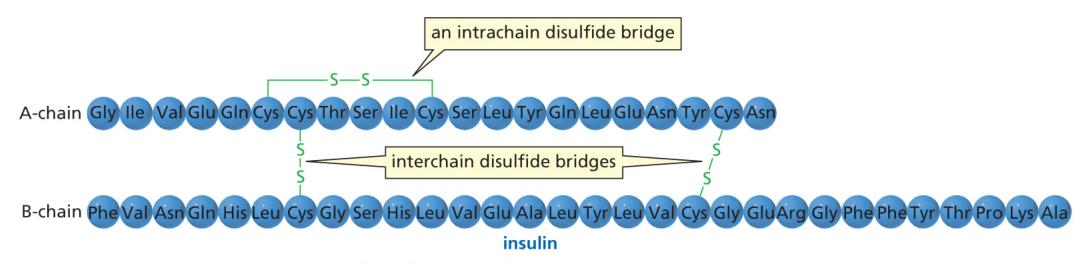
- Short polymers of amino acids
- Each unit is called a residue
- 2 residues dipeptide
- 3 residues tripeptide
- 12-20 residues oligopeptide
- many polypeptide

### "Protein"



#### One or more polypeptide chains

- One polypeptide chain a monomeric protein
- More than one multimeric protein
- Homomultimer one kind of chain
- Heteromultimer two or more different chains
- Hemoglobin, for example, is a heterotetramer
- It has two alpha chains and two beta chains



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#### Enkephalins synthesized by the body to control pain

Tyr-Gly-Gly-Phe-Leu leucine enkephalin

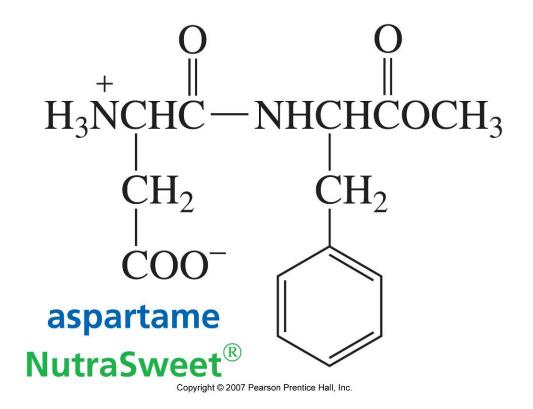
Tyr-Gly-Gly-Phe-Met methionine enkephalin

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morphine

#### **Peptide hormones**

**Bradykinin** inhibits the inflammation of tissues **Vasopressin** controls blood pressure **Oxyocin** induces labor in pregnant women by stimulating the uterine muscle



# 200 times sweeter than sucrose D-amino acid is bitter.

## Destroy harmful oxidizing agents in the body

## The Sequence of Amino Acids in a Protein



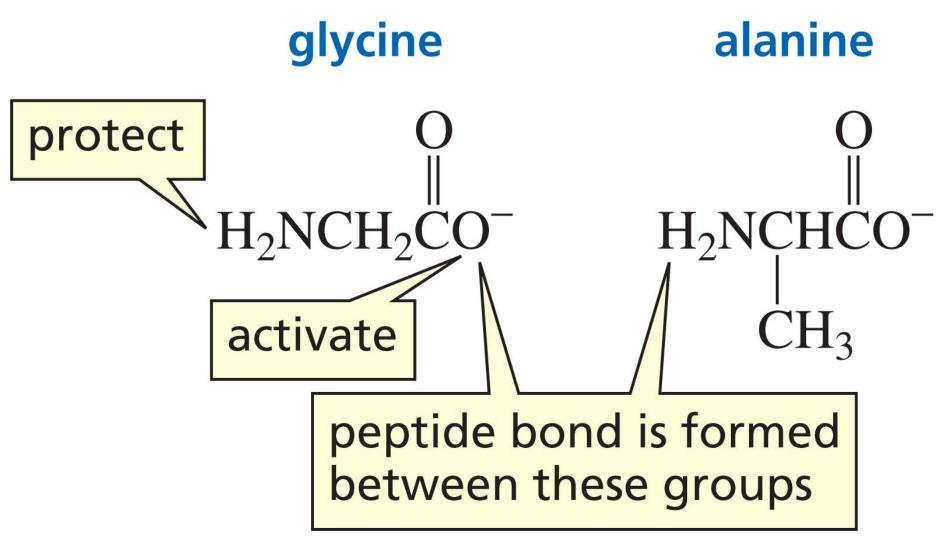
- Is a unique characteristic of every protein
- Is encoded by the nucleotide sequence of DNA
- Is thus a form of genetic information
- Is read from the amino terminus to the carboxyl terminus

## **Chemical Synthesis of peptide**

## Solid Phase Synthesis of Peptides

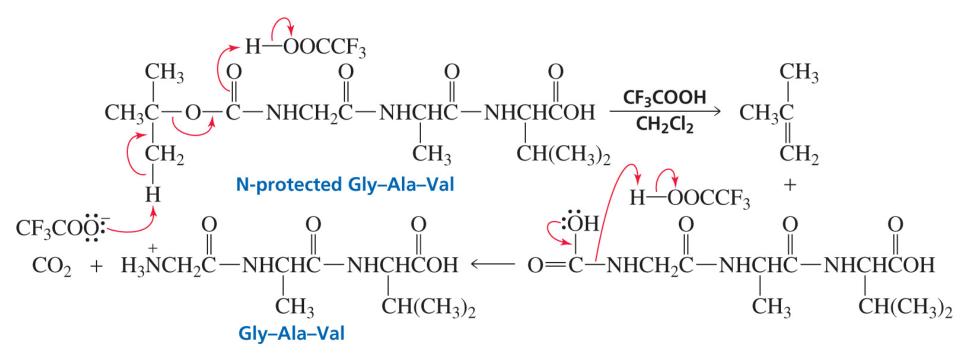


- R. Bruce Merrifield and his collaborators pioneered the solid-phase synthesis of polypeptides in the laboratory
- Carboxy terminus of a nascent peptide is covalently anchored to an insoluble resin
- After each addition of a residue, the resin particles are collected by filtration
- Automation and computer control now permit synthesis of peptides of 30 residues or more



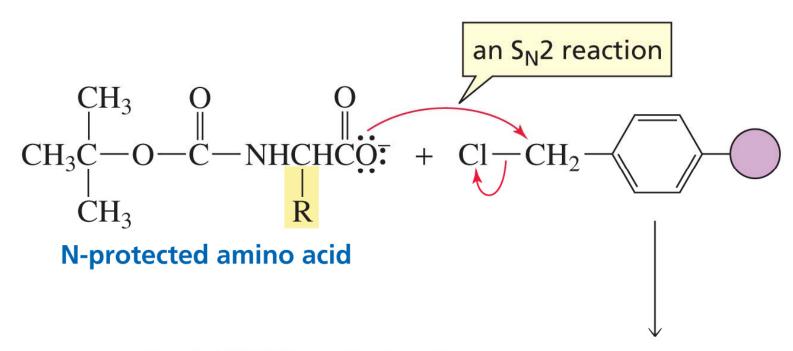
#### Boc

$$\begin{array}{c} \text{CH}_3 \text{ O} \\ \text{CH}_4 \text{ O} \\ \text{CH}_4 \text{ O} \\ \text{CH}_5 \text{ O} \\ \text{CH}_5 \text{ O} \\ \text{CH}_5 \text{ O} \\ \text{CH}_6 \text{ O} \\ \text{CH}_7 \text{ O} \\ \text{CH}_7$$



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#### Merrifield automated solid-phase synthesis of a tripeptide



#### **Automated solid-phase peptide synthesis**

