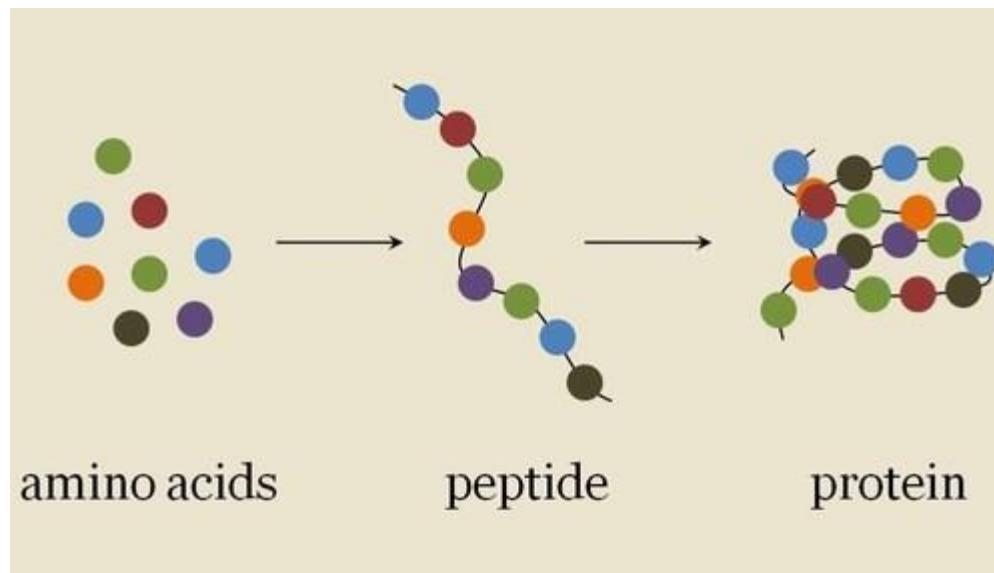
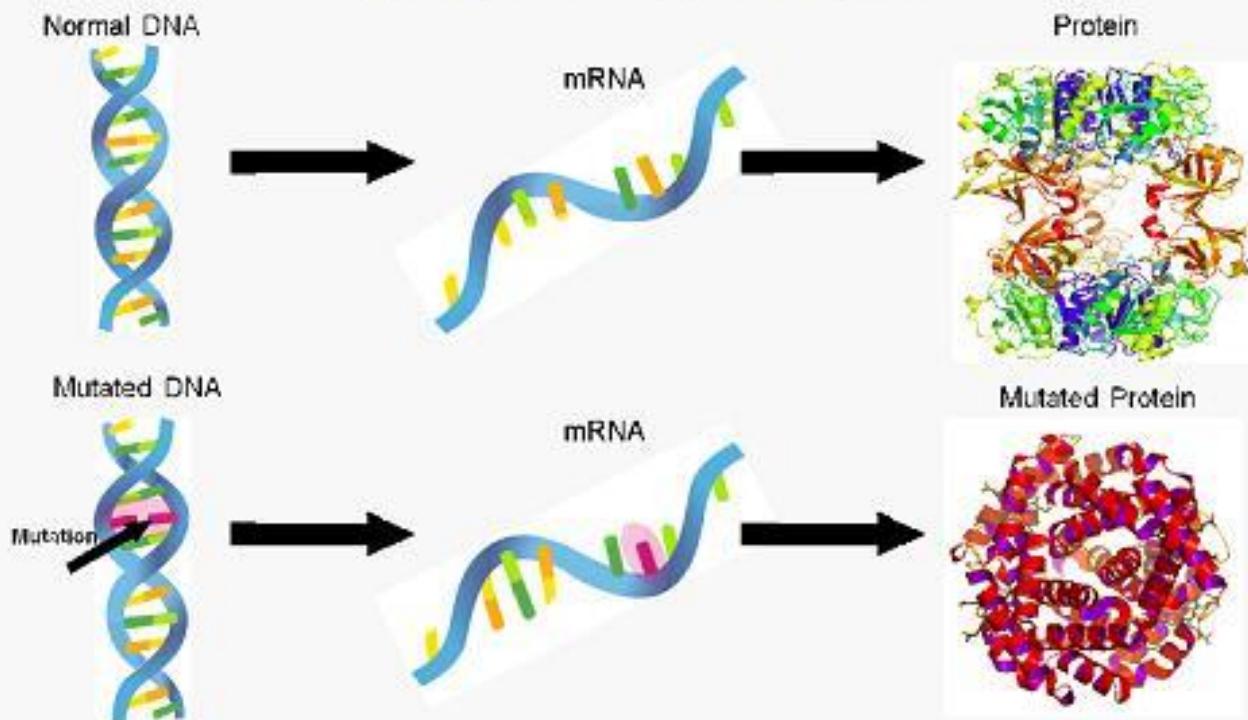


Amino Acids, Peptides, and Proteins



Why Are Mutations Important?

Many Serious Diseases Are Driven by a Change in Protein Structure

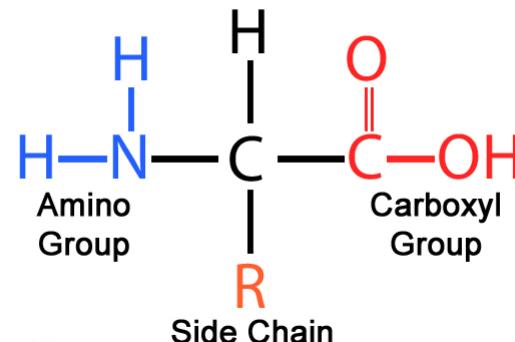


TRANSGENOMIC®
the power of discovery

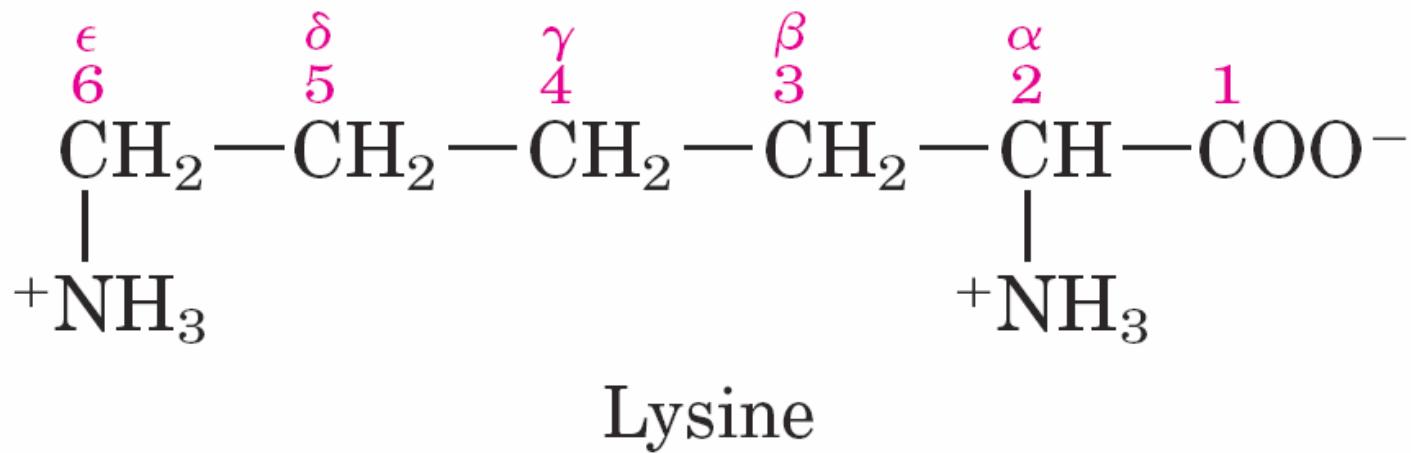
Amino Acid Structure

- Proteins: polymers of amino acids, with each **amino acid residue** joined
- **Amino Acids share common structure**
 - All 20 of the common amino acids are α -amino acids
 - A carboxyl group and an amino group bonded to the same carbon atom (the α carbon).
- Different α –amino acids are distinguished by their different **side chains** (R group).The α -carbon atom is thus a **chiral center**.

Amino Acid Structure

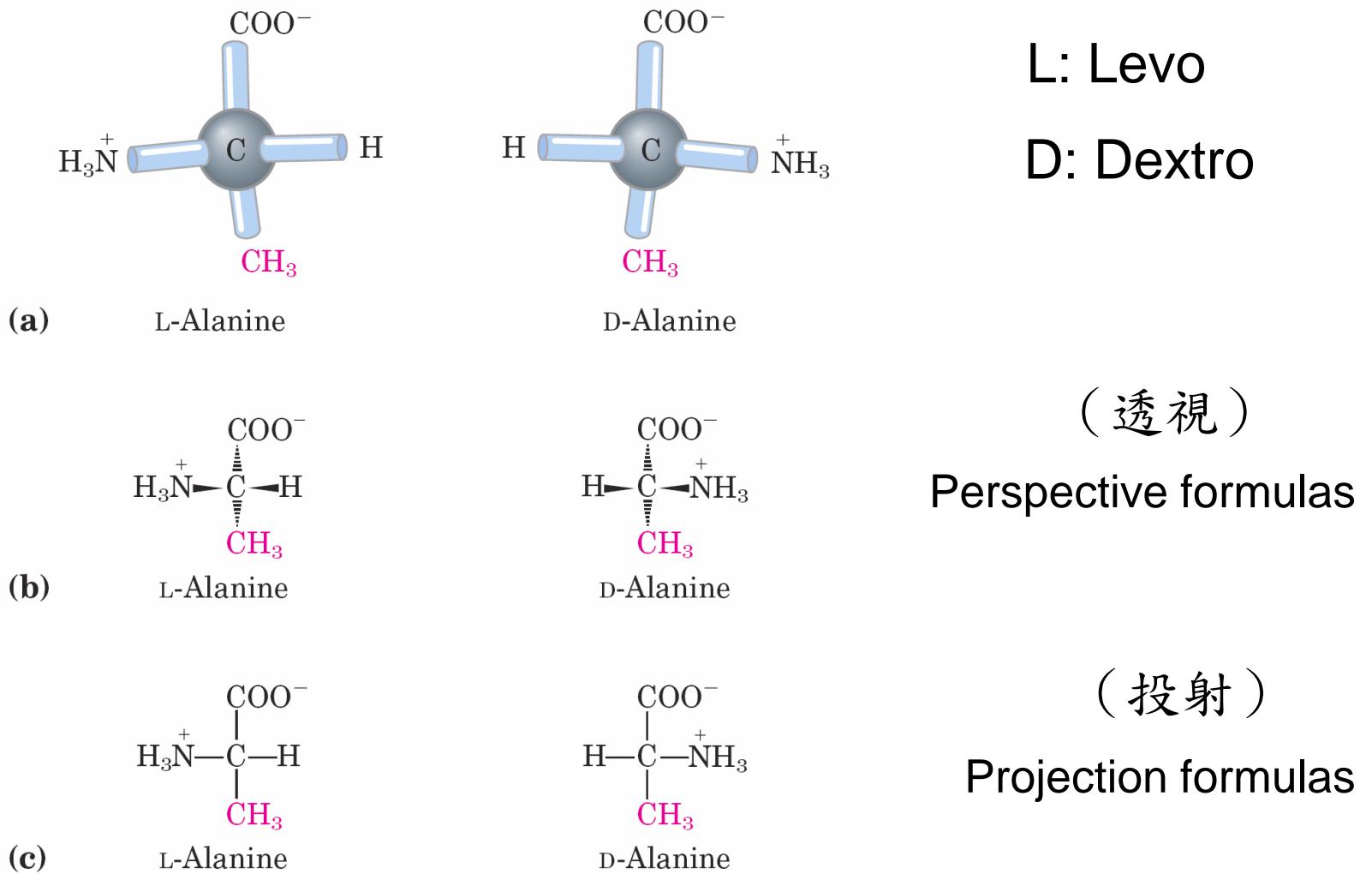


Nomenclature: Carbons



The additional carbons in an R group
are commonly designated β , γ , δ , ϵ .

Amino acids have two stereoisomers



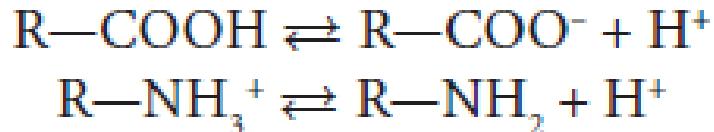
Amino acid can be classified by R group

- The 20 common amino acids can be grouped into five main classes based on the properties of their R groups
- Polarity: or tendency to interact with water at biological pH. The polarity of the R groups varies widely, from nonpolar and hydrophobic (water-insoluble) to highly polar and hydrophilic (water-soluble).
- The 20 common amino acids can also be classified by their side chains:
 - Aliphatic
 - Aromatic
 - Hydroxyl or sulfur-containing
 - Basic
 - Acidic and their amides

TABLE 3-1 Properties and Conventions Associated with the Common Amino Acids Found in Proteins

Amino acid	Abbreviation/ symbol	M_r	pK_a values				Hydropathy index*	Occurrence in proteins (%)†				
			pK_1 (—COOH)	pK_2 (—NH ₃ ⁺)	pK_R (R group)	pI						
Nonpolar, aliphatic												
R groups												
Glycine	Gly G	75	2.34	9.60		5.97	-0.4	7.2				
Alanine	Ala A	89	2.34	9.69		6.01	1.8	7.8				
Proline	Pro P	115	1.99	10.96		6.48	1.6	5.2				
Valine	Val V	117	2.32	9.62		5.97	4.2	6.6				
Leucine	Leu L	131	2.36	9.60		5.98	3.8	9.1				
Isoleucine	Ile I	131	2.36	9.68		6.02	4.5	5.3				
Methionine	Met M	149	2.28	9.21		5.74	1.9	2.3				
Aromatic R groups												
Phenylalanine	Phe F	165	1.83	9.13		5.48	2.8	3.9				
Tyrosine	Tyr Y	181	2.20	9.11	10.07	5.66	-1.3	3.2				
Tryptophan	Trp W	204	2.38	9.39		5.89	-0.9	1.4				
Polar, uncharged												
R groups												
Serine	Ser S	105	2.21	9.15		5.68	-0.8	6.8				
Threonine	Thr T	119	2.11	9.62		5.87	-0.7	5.9				
Cysteine	Cys C	121	1.96	10.28	8.18	5.07	2.5	1.9				
Asparagine	Asn N	132	2.02	8.80		5.41	-3.5	4.3				
Glutamine	Gln Q	146	2.17	9.13		5.65	-3.5	4.2				
Positively charged												
R groups												
Lysine	Lys K	146	2.18	8.95	10.53	9.74	-3.9	5.9				
Histidine	His H	155	1.82	9.17	6.00	7.59	-3.2	2.3				
Arginine	Arg R	174	2.17	9.04	12.48	10.76	-4.5	5.1				
Negatively charged												
R groups												
Aspartate	Asp D	133	1.88	9.60	3.65	2.77	-3.5	5.3				
Glutamate	Glu E	147	2.19	9.67	4.25	3.22	-3.5	6.3				

pKa – The acid strength of weak acids (代表一種酸解離氫離子的能力)



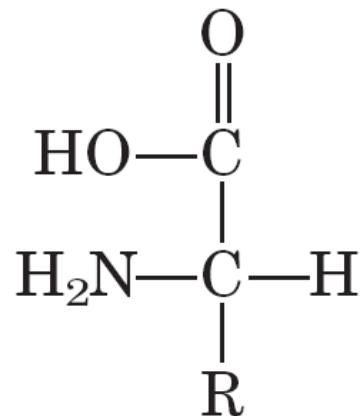
pI(isoelectric point)- the pH value at which number of positive charges equals the number of negative charges

isoelectric pH (pI) of alanine

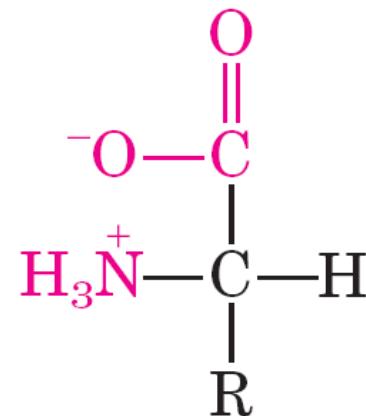
$$pI = \frac{pK_1 + pK_2}{2} = \frac{2.35 + 9.69}{2} = 6.02$$

hydropathy index - a measure of polarity of an amino acid residue.

Amino acids can act as acids and bases



Nonionic
form

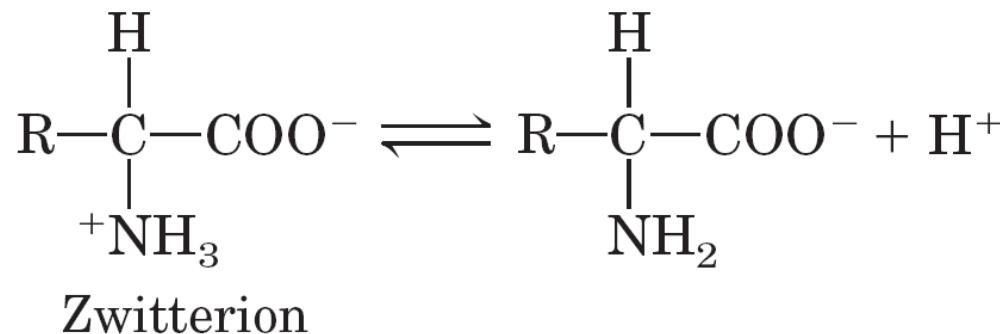


Zwitterionic
form

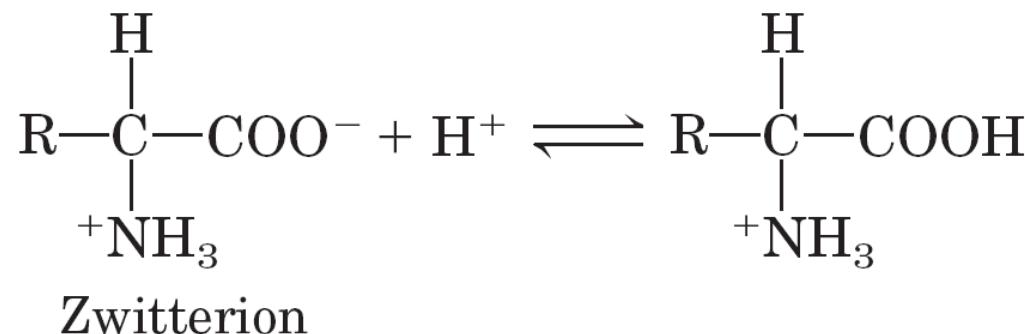
The nonionic form does not occur in significant amounts in aqueous solutions.
The zwitterion predominates at neutral pH.

Amino acids can act as acids and bases

as an acid (proton donor)

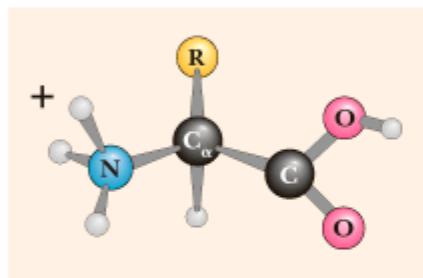


or a base (proton acceptor)

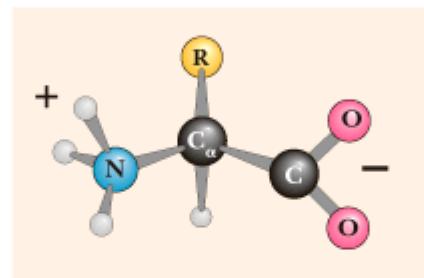


Isoelectric point (pI)

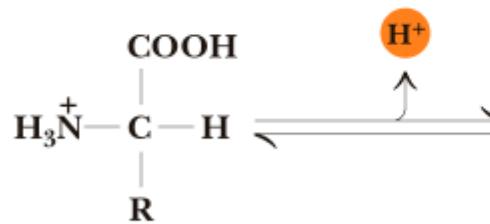
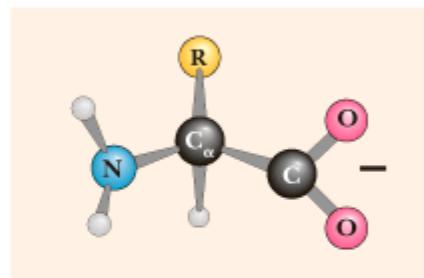
pH 1 Net charge +1



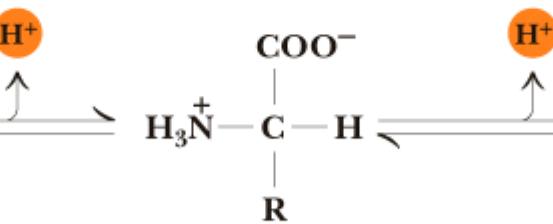
pH 7 Net charge 0



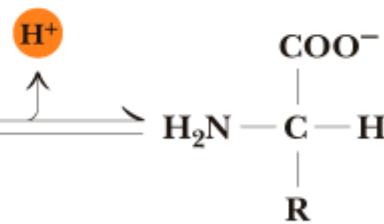
pH 13 Net charge -1



Cationic form

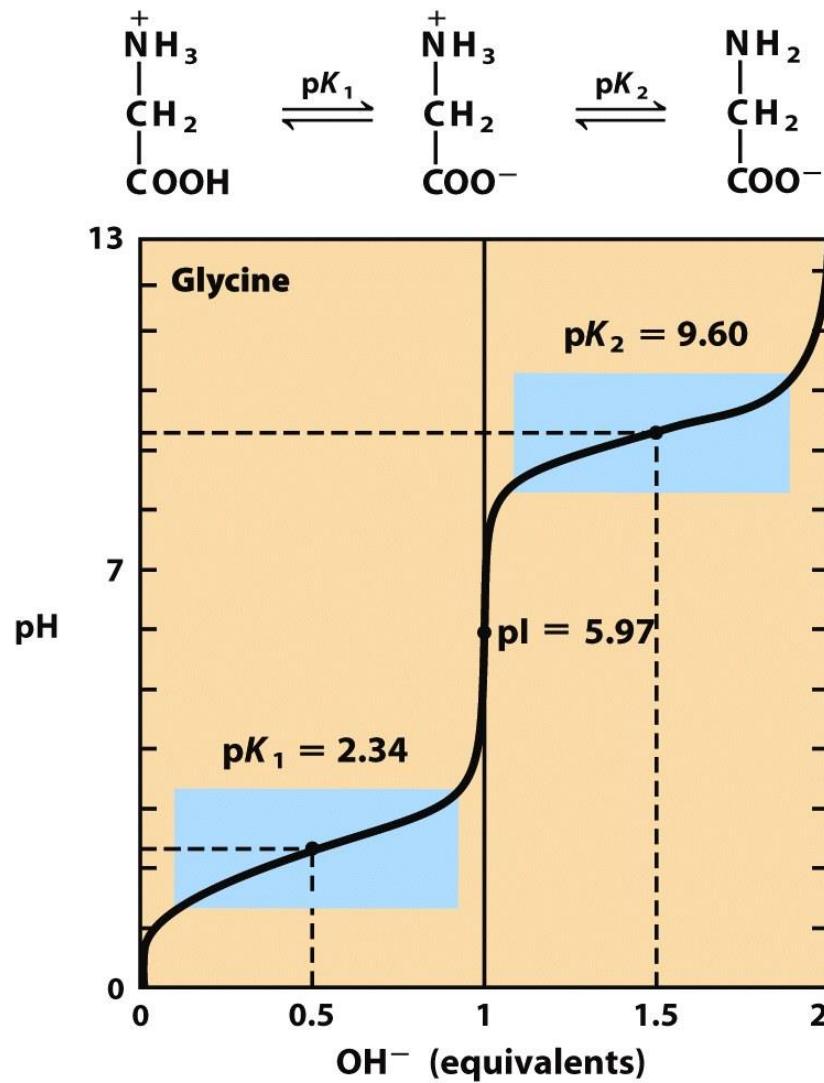


Zwitterion (neutral)



Anionic form

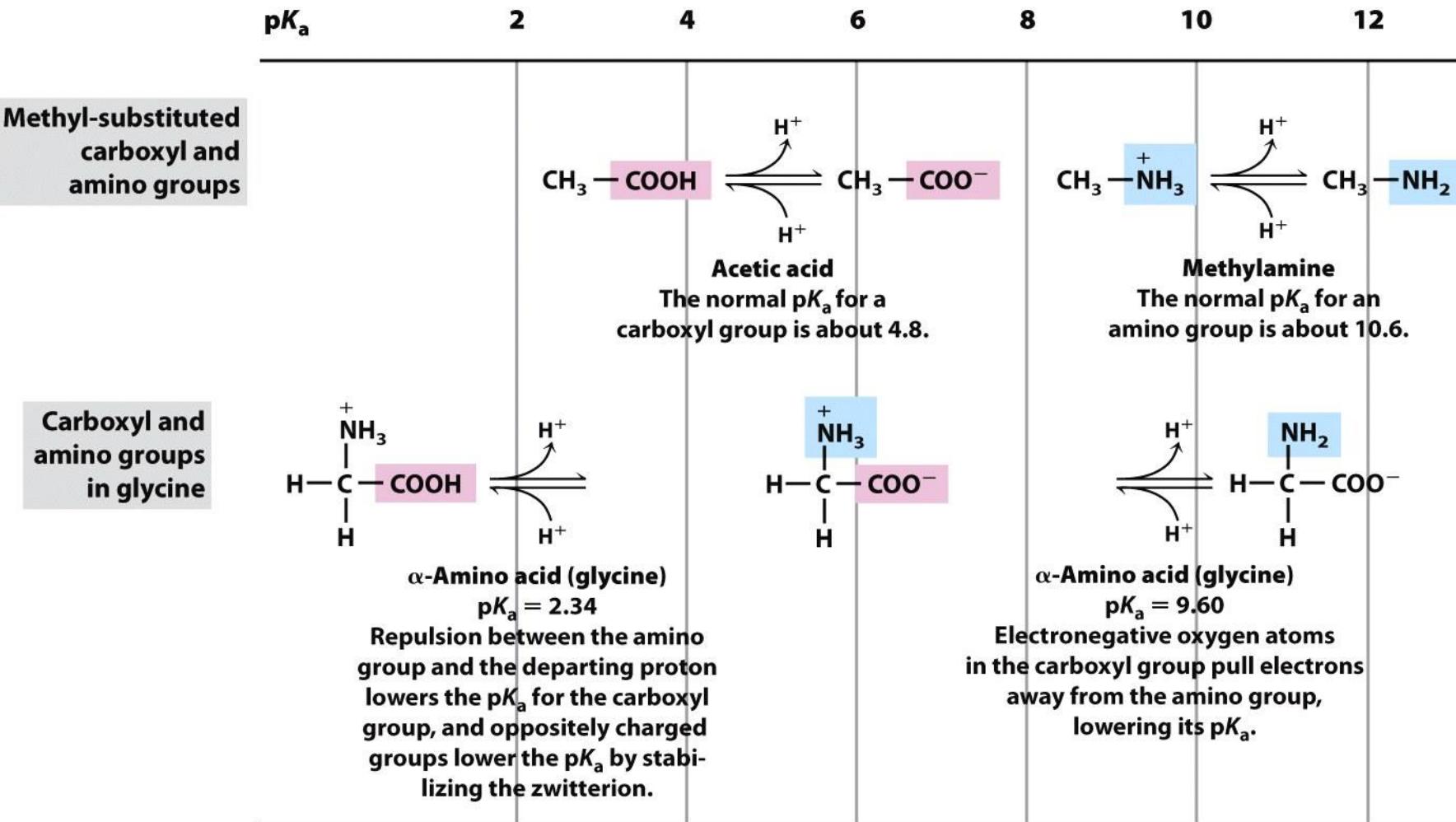
Amino acids have characteristic titration curves



$$\begin{aligned} pI &= \frac{1}{2} (pK_1 + pK_2) \\ &= \frac{1}{2} (2.34 + 9.6) \\ &= 5.97 \end{aligned}$$

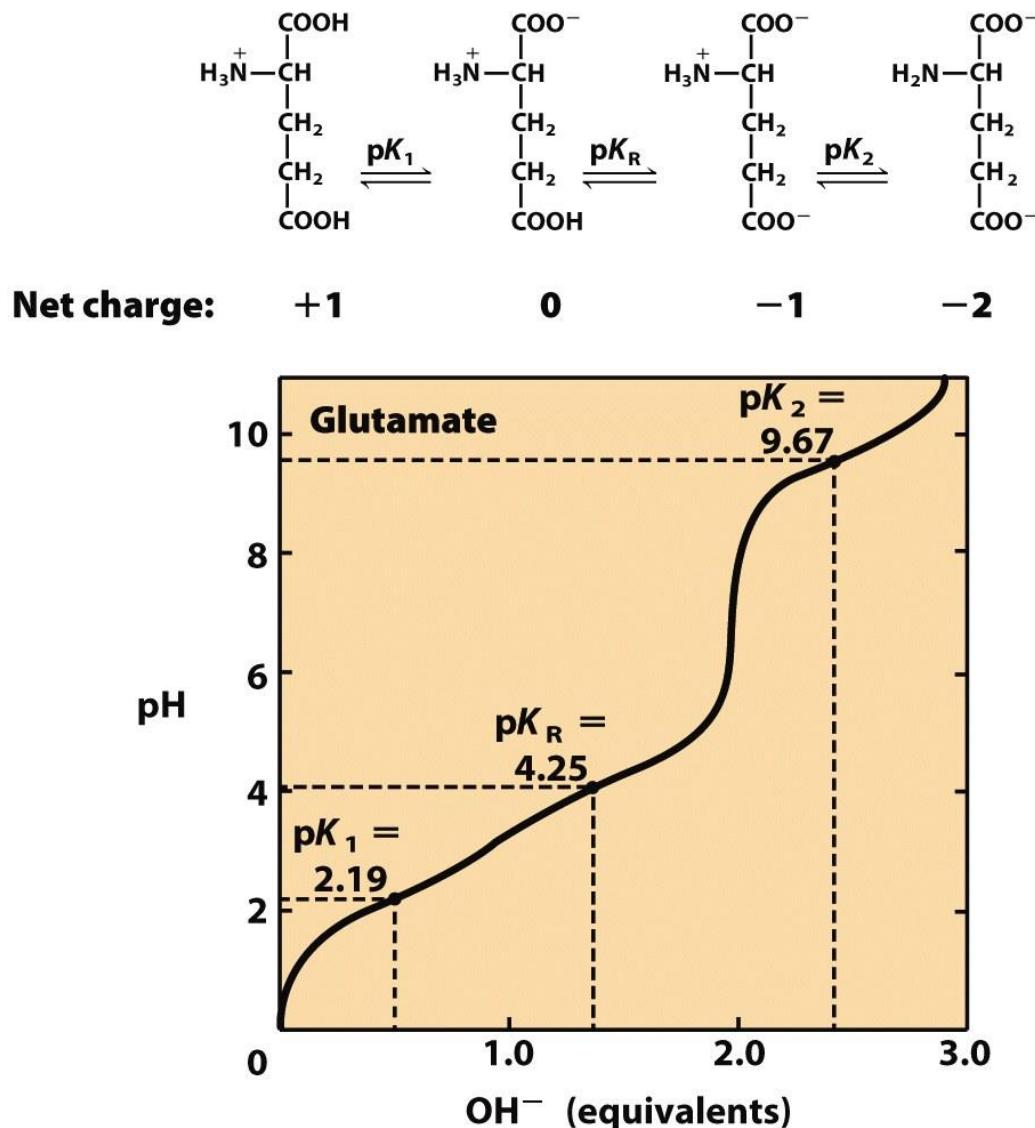
Figure 3-10
Lehninger Principles of Biochemistry, Fifth Edition
© 2008 W.H. Freeman and Company

Effect of the chemical environment on pK_a



The pK_a values for the ionizable groups in glycine are lower than those for simple, methyl-substituted amino and carboxyl groups. These downward perturbations of pK_a are due to intramolecular interactions. They can be caused by chemical groups that happen to be positioned nearby.

Titration curves for glutamate



$$pI = \frac{2.19 + 4.25}{2}$$

Figure 3-12a

Lehninger Principles of Biochemistry, Fifth Edition

© 2008 W. H. Freeman and Company

Titration curves for histidine

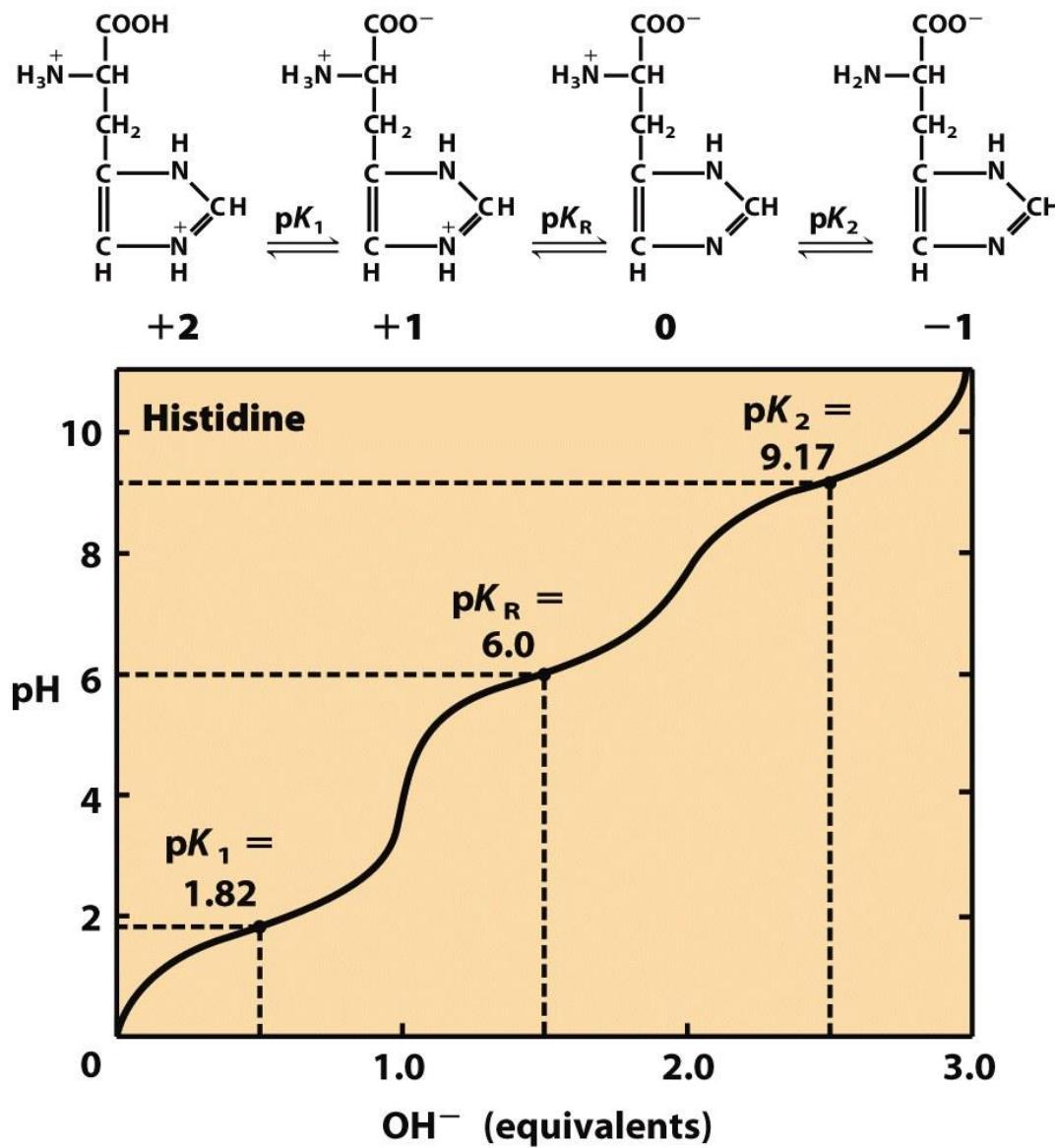


Figure 3-12b

Lehninger Principles of Biochemistry, Fifth Edition

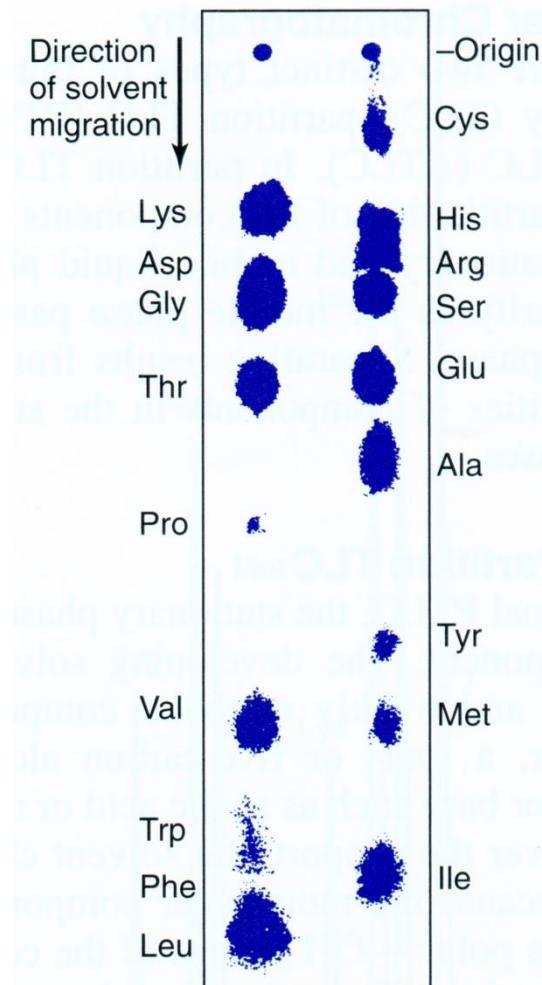
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Typical range of pK values for ionizable groups in proteins

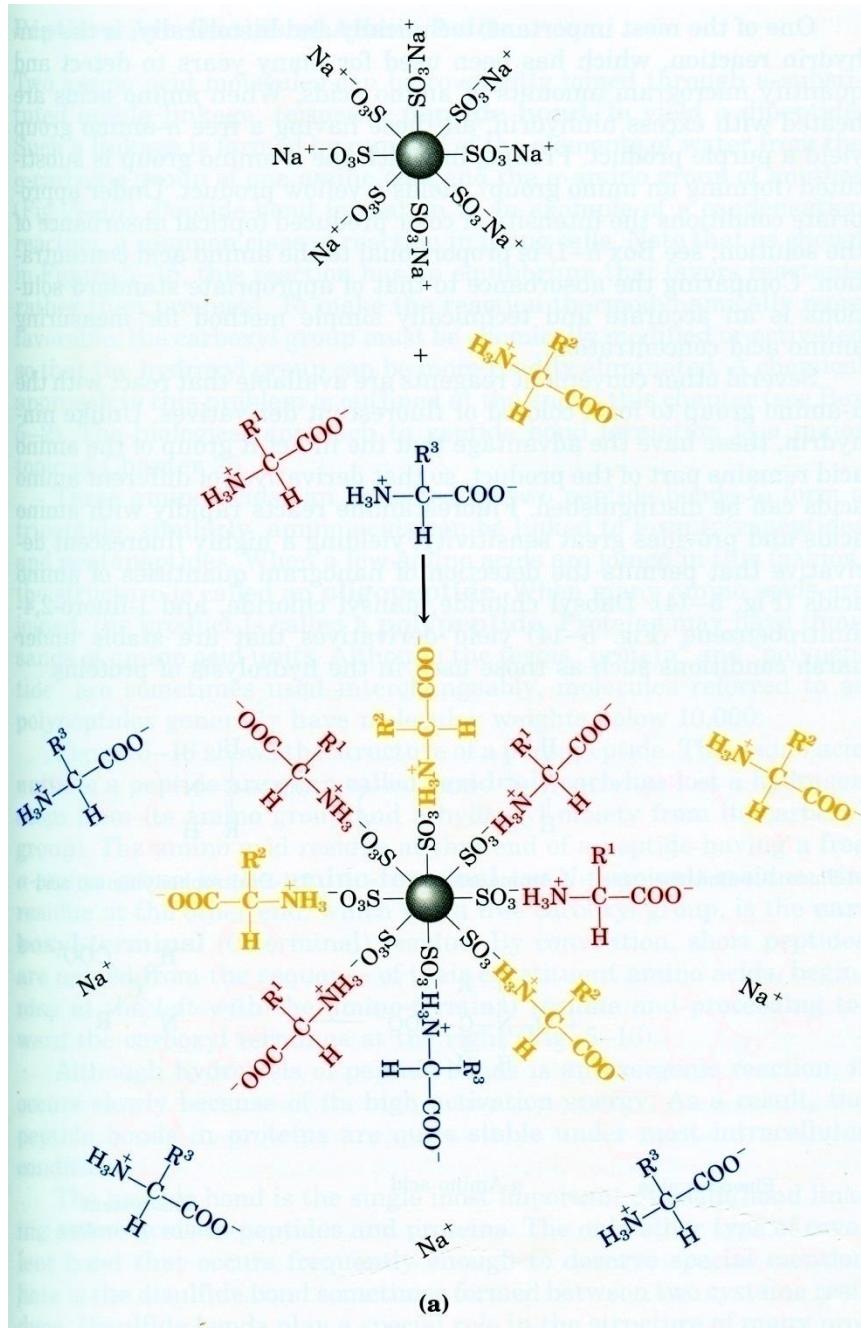
Dissociating Group	pK Range
α -Carboxyl	3.5–4.0
Non- α COOH of Asp or Glu	4.0–4.8
Imidazole of His	6.5–7.4
SH of Cys	8.5–9.0
OH of Tyr	9.5–10.5
α -Amino	8.0–9.0
ε -Amino of Lys	9.8–10.4
Guanidinium of Arg	~12.0

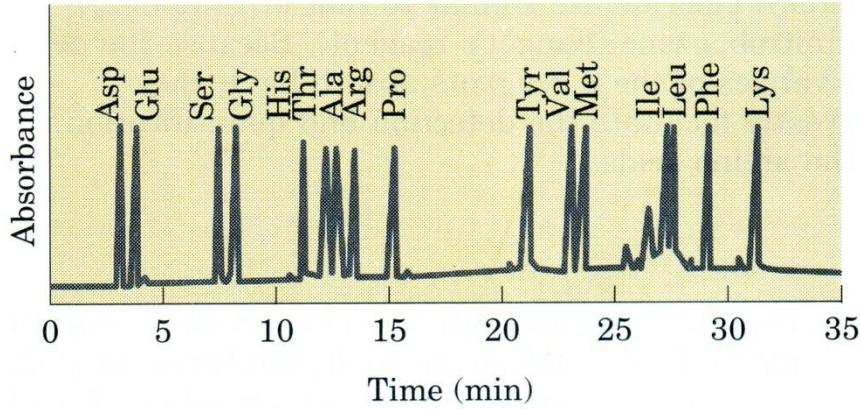
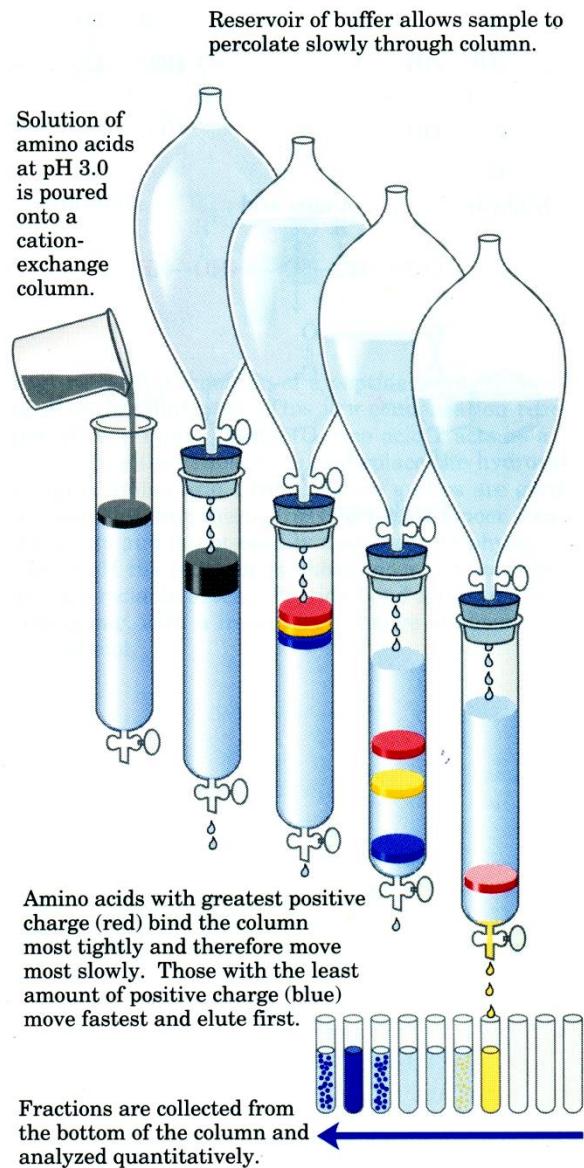
Various techniques separate amino acids

Paper chromatography



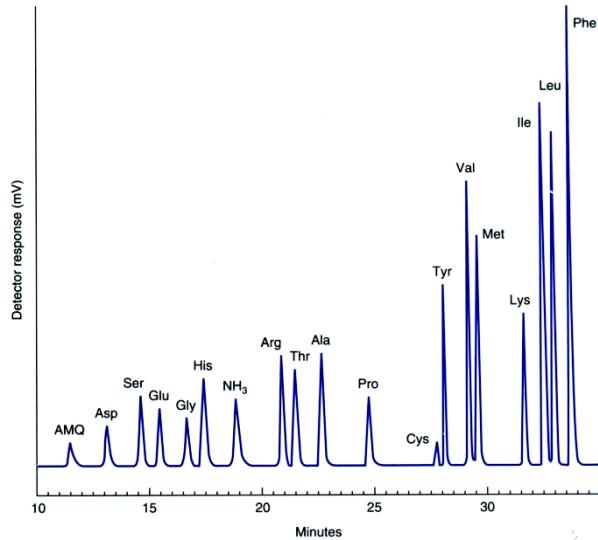
Ion exchange chromatography separates amino acids by electric charge





Analysis of amino acids

High Performance liquid chromatography (HPLC)



Amino Acid Analyzer



The 20 standard amino acids of proteins

Amino acids with aliphatic side chains

G: R=-H

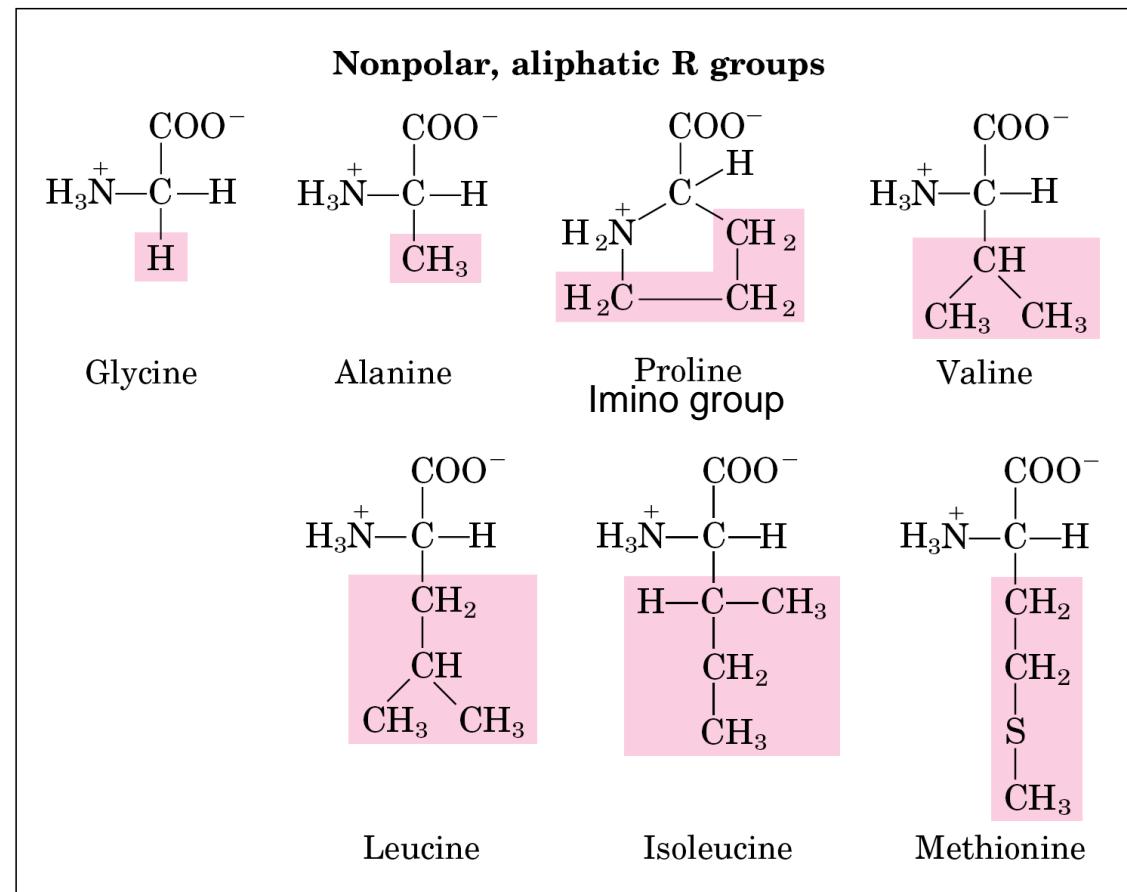
A: R=-CH₃

V: R=-C^H₃
CH₃

L: R=-CH₂-CH^H₃
CH₃

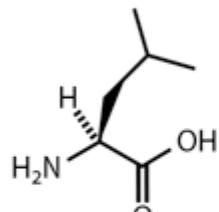
M: R=-CH₂-CH₂-S-CH₃

I: R=-CH^H₃
CH₂-CH₃

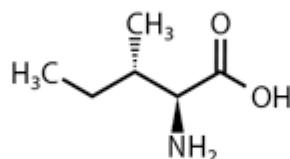


Small amino acids- glycine and alanine
Branched chain amino acids-valine,₂₁
leucine and isoleucine

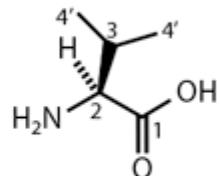
BRANCHED CHAIN AMINO ACIDS



LEUCINE



ISOLEUCINE



VALINE

(Leucine) 2-Amino-4-methylpentanoic acid
(Isoleucine) 2-Amino-3-methylpentanoic acid
(Valine) 2-amino-3-methylbutanoic acid



Improve exercise performance
and reduce muscle breakdown

Of the essential amino acids,
three account for as much as
33% of muscle tissue – leucine,
isoleucine, and valine.

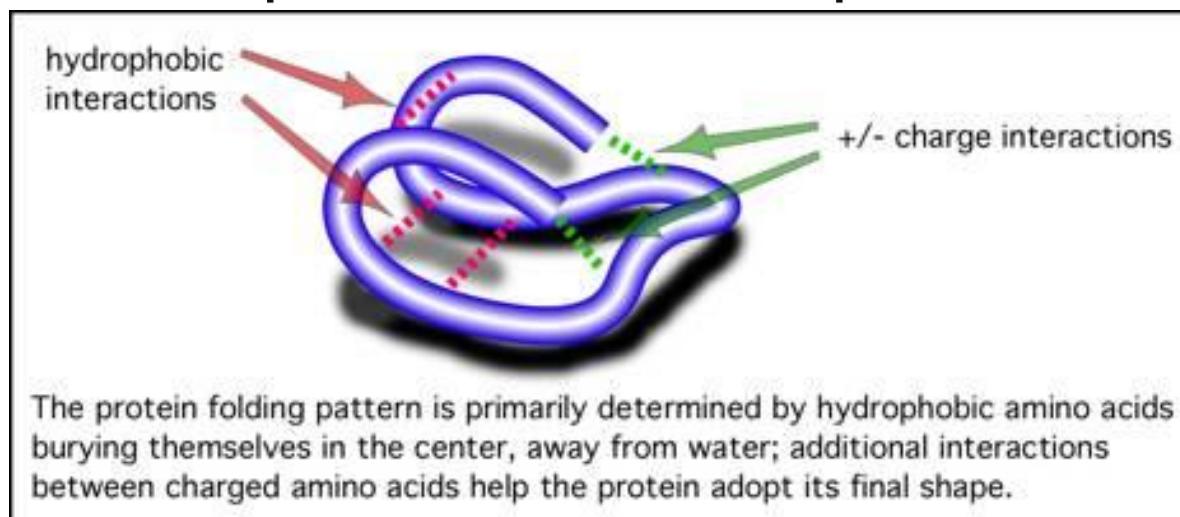
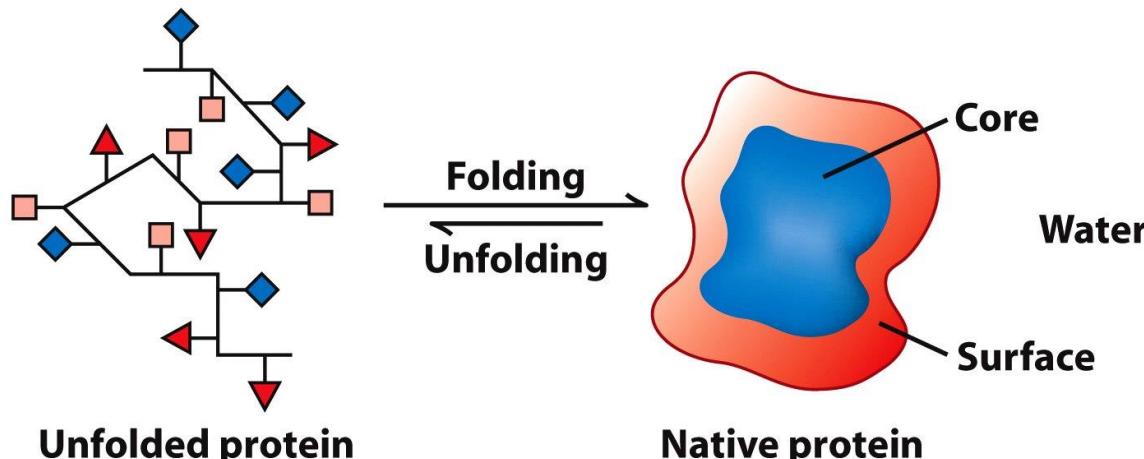


Classification of the L- α -amino acids of proteins based on their relative hydrophilicity or hydrophobicity

Hydrophobic	Hydrophilic	
Alanine	Arginine	Histidine
Isoleucine	Asparagine	Lysine
Leucine	Aspartic acid	Serine
Methionine	Cysteine	Threonine
Phenylalanine	Glutamic acid	
Proline	Glutamine	
Tryptophan	Glycine	
Tyrosine		
Valine		

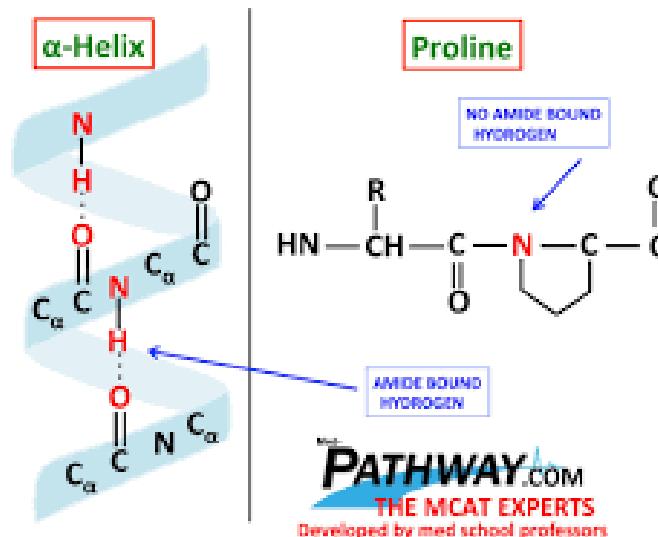
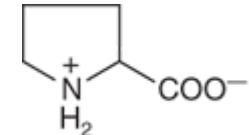
Properties of the Amino Acid Side Chains

- The more hydrophobic amino acids such as **isoleucine** are usually found *within the core of a protein molecule, where they are shielded from water.*

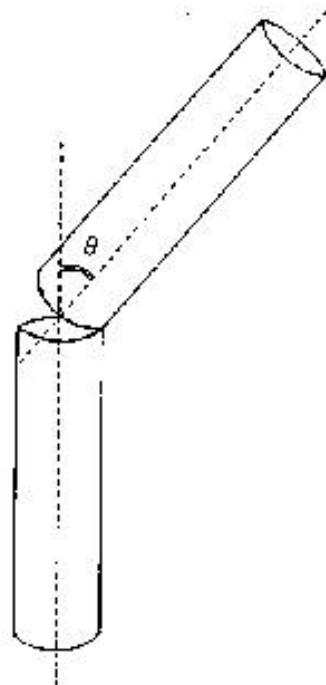


Properties of the Amino Acid Side Chains

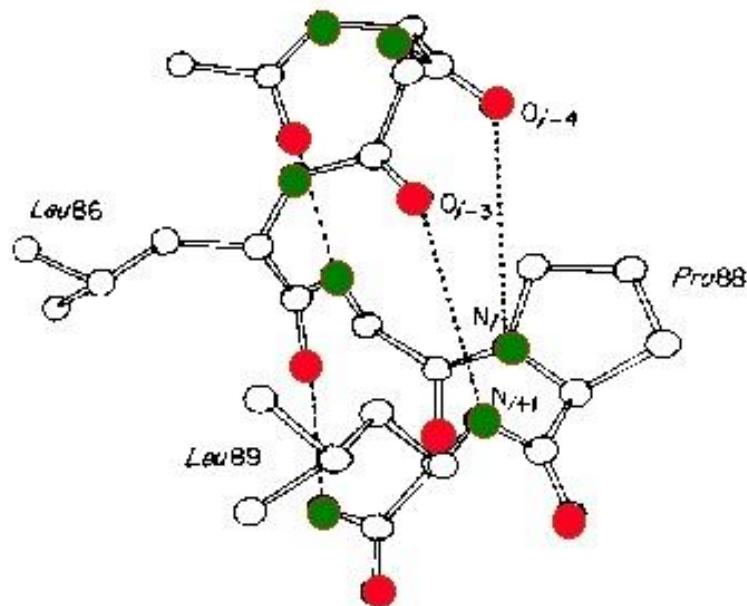
- **Proline:** only amino acid which the side chain forms a covalent bond with the α -amino group.
- The rigid ring of proline: protein structure where the protein must fold back on itself (so-called “turns”).
- Proline acts as a structural disruptor in the middle of regular secondary structure elements such as alpha helices and beta sheets.



Proline is a helix-breaker but participates in formation of other motifs

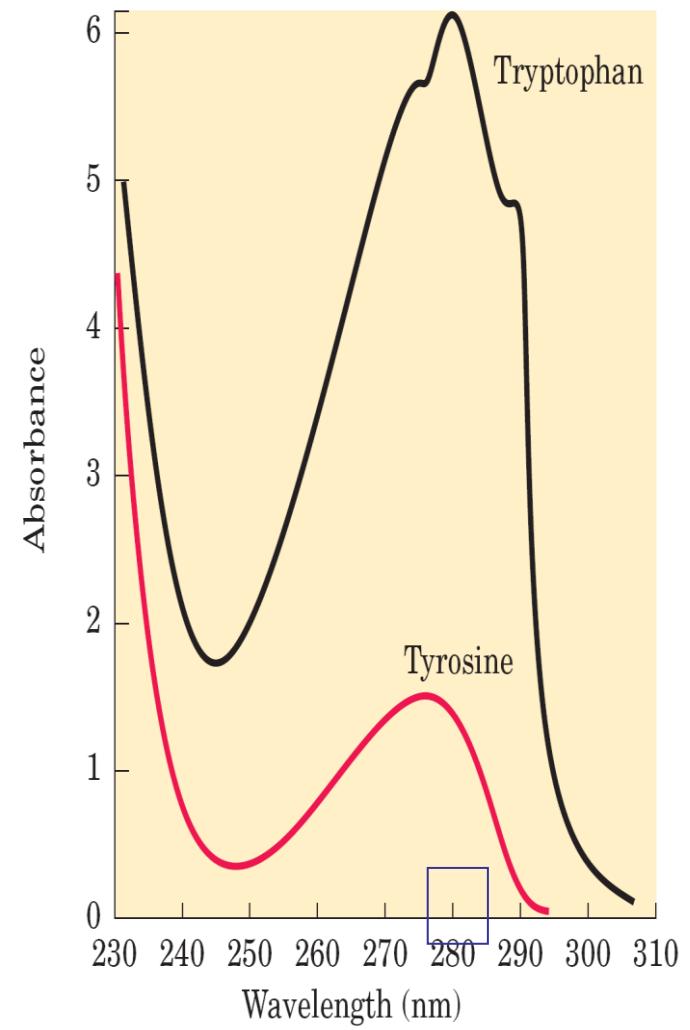
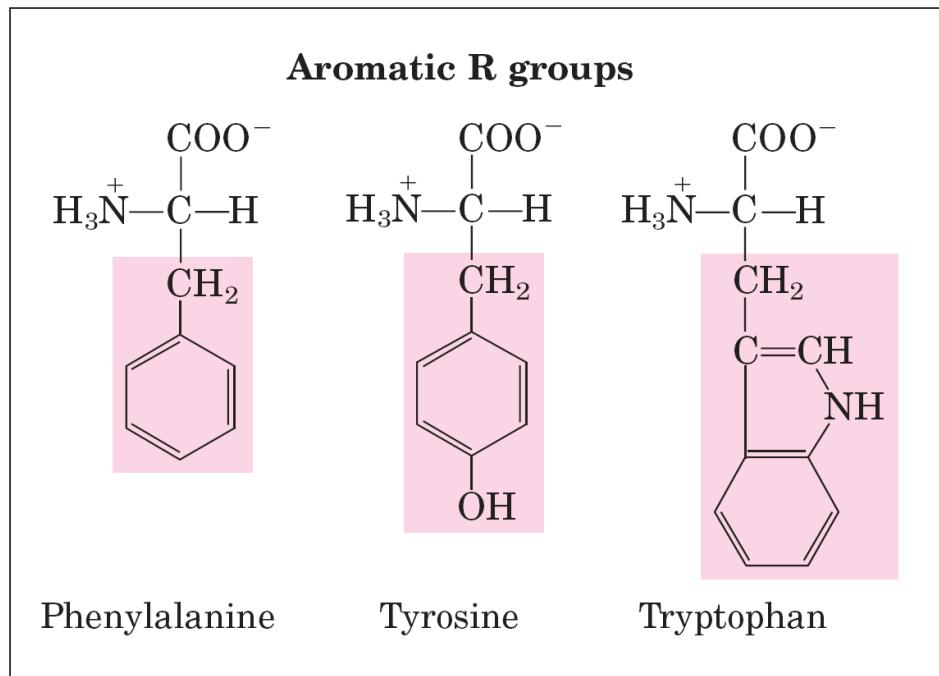


In helices



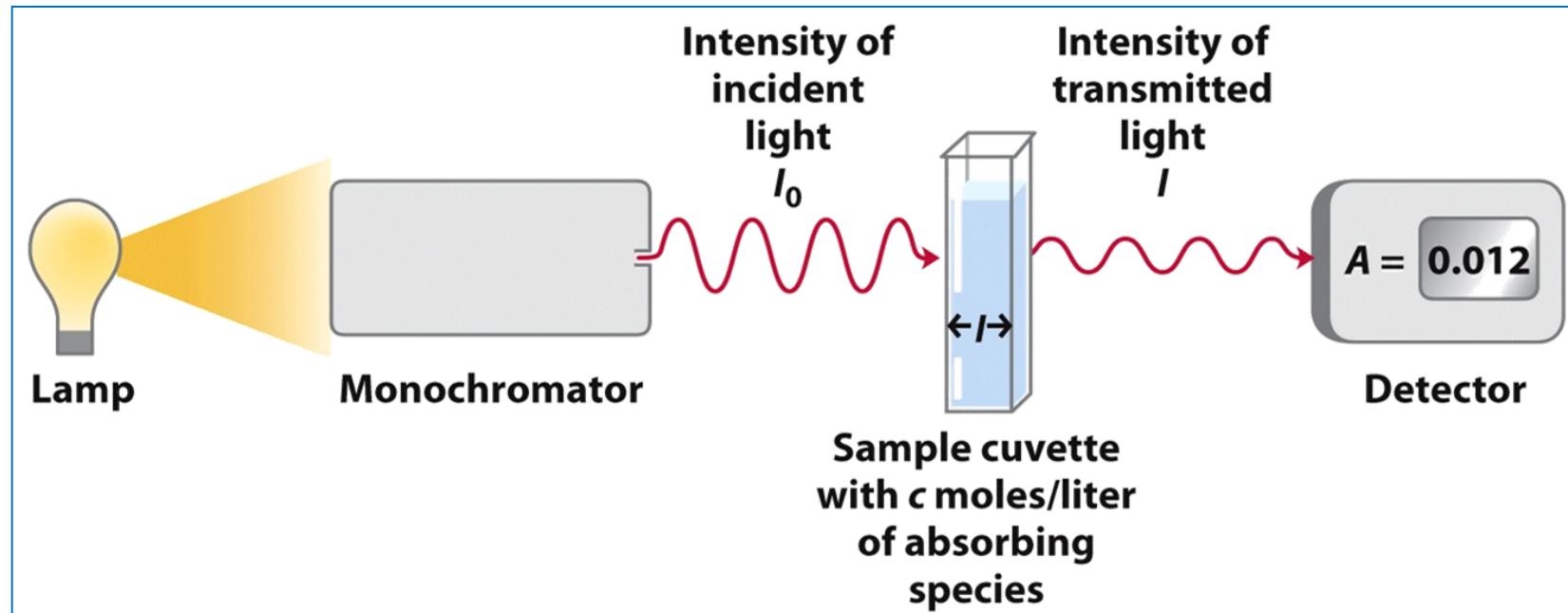
In turns

Aromatic amino acids



Phenylalanine: one of the most hydrophobic amino acids.

The principle components of a spectrophotometer



Lambert-Beer Law

$$\log \frac{I_0}{I} = \varepsilon c l$$

I_0 = Absorbance (A)= Optical density (OD)

I_0 = Intensity of the incident light

I = Intensity of the transmitted light

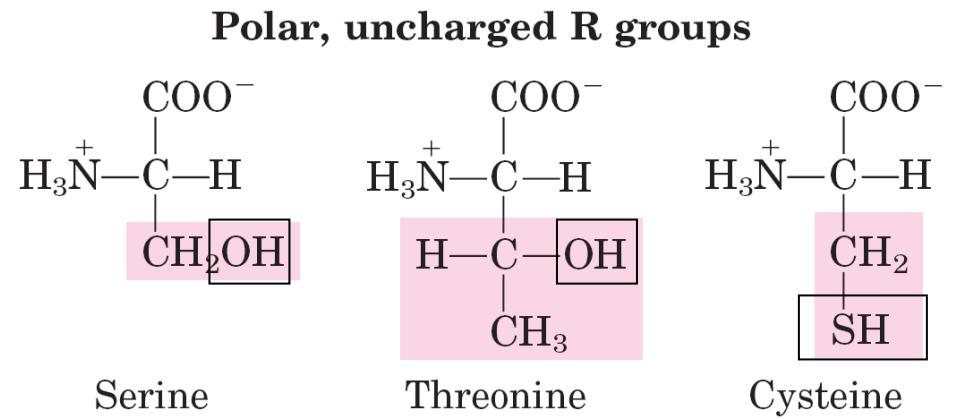
ε = molecular absorption coefficient

c = concentration

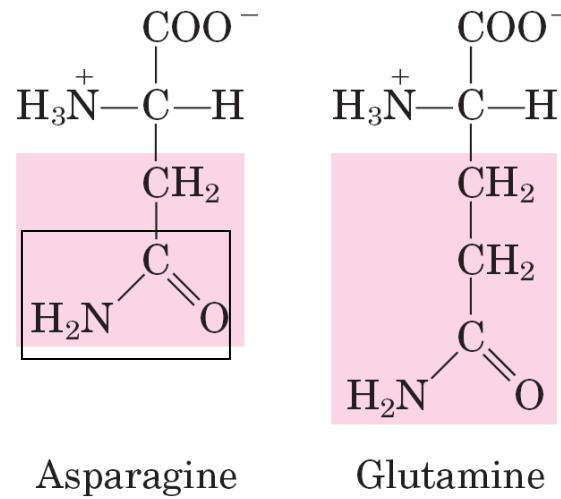
l = path length $OD_{280} = 1.2 \Rightarrow 1 \text{ mg/ml}$

Amino acids with hydroxy or sulfur containing side chains

Hydroxy amino acids



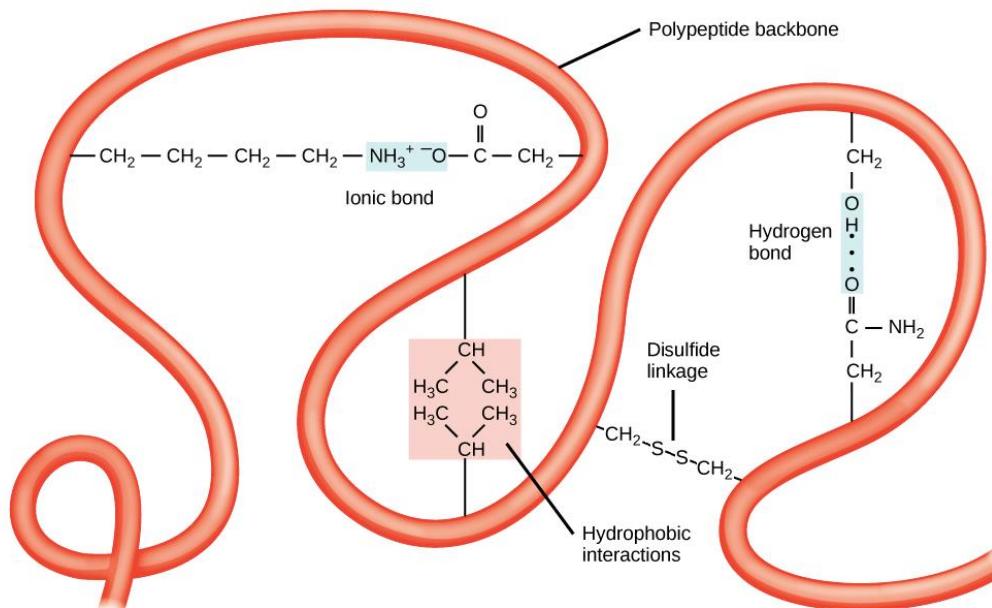
hydroxy group sulfhydryl group



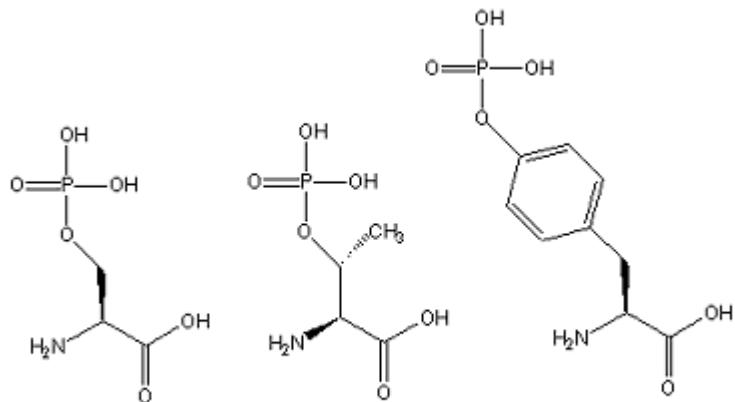
Amide group

Properties of the Amino Acid Side Chains

- The –OH group of serine and the –SH group of cysteine are good nucleophiles and often play key roles in enzyme activity.
- Asn and Gln are hydrophilic and tend to be on the surface of a protein molecule, in contact with the surrounding water.

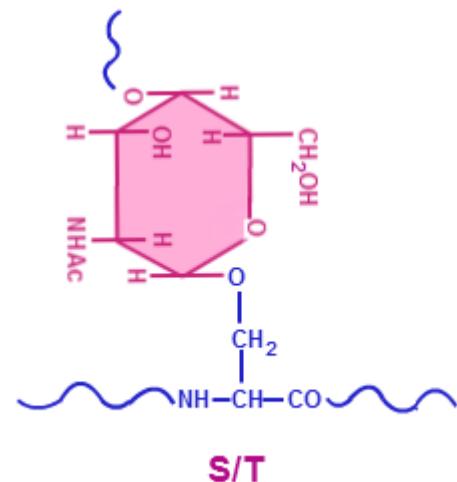


Post translational modification



Phospho Serine, Threonine and Tyrosine amino acids

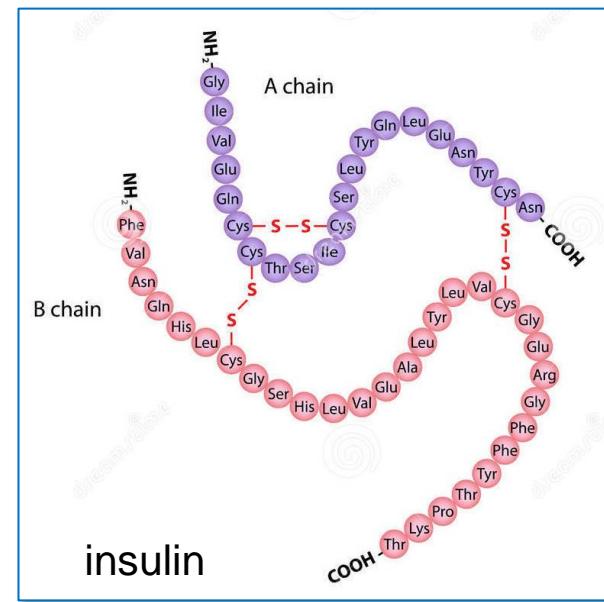
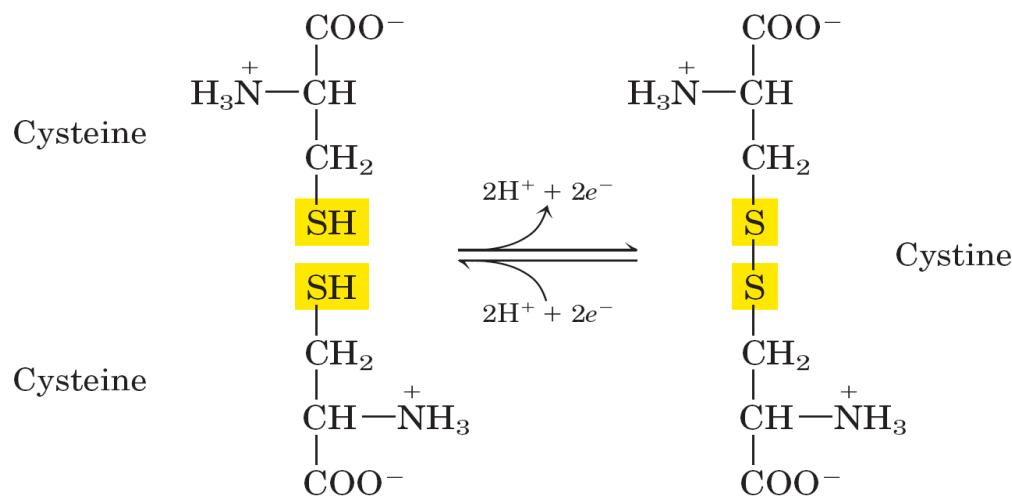
phosphorylation



S/T

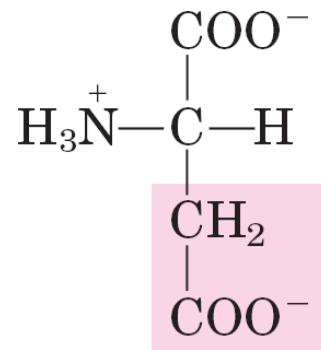
glycosylation

Formation of a disulfide bond by the oxidation of two molecules of cysteine

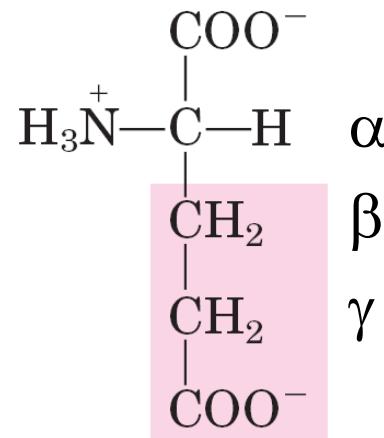


Acidic amino acids

Negatively charged R groups



Aspartate

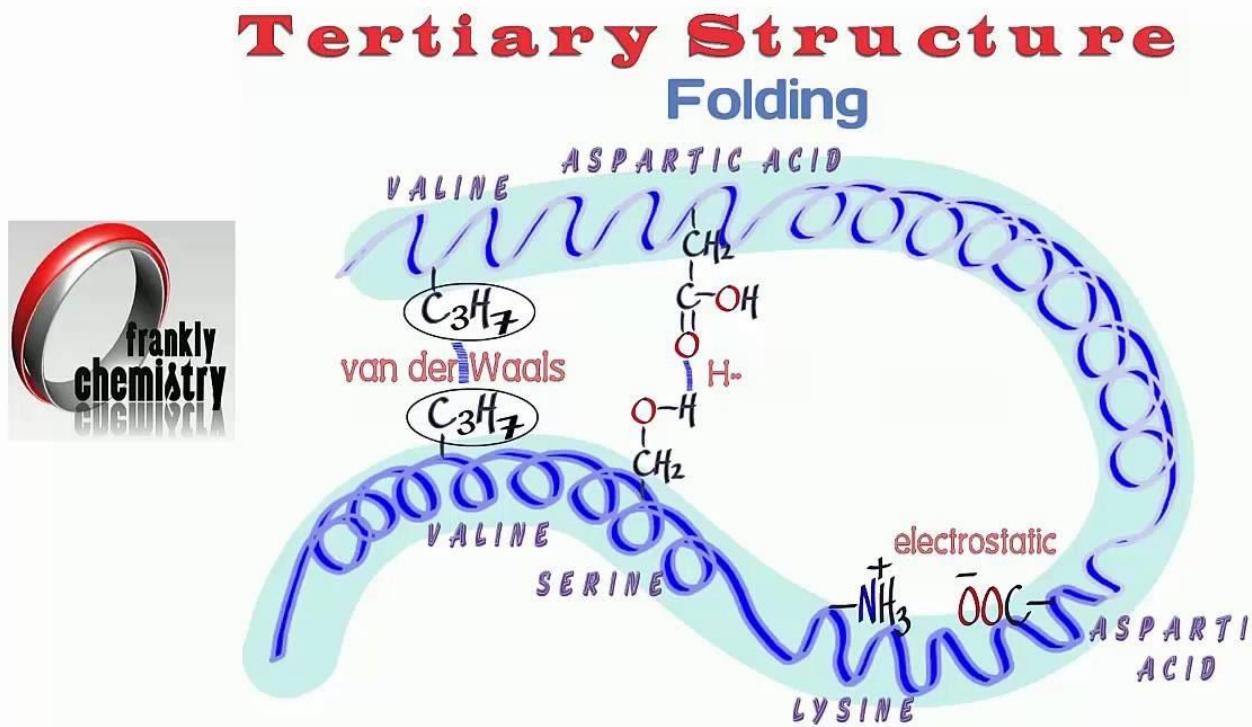


Glutamate

COOH= carboxy group

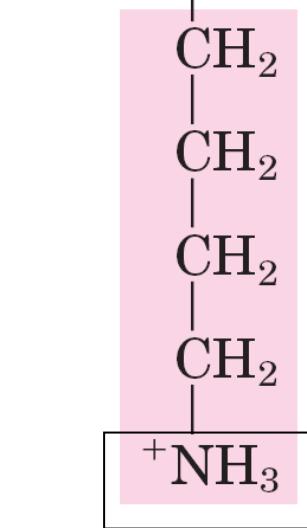
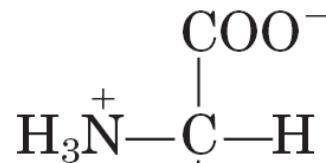
Properties of the Amino Acid Side Chains

- Aspartic acid ($pK_a = 3.9$) and glutamic acid ($pK_a = 4.2$) typically carry negative charges at pH 7.

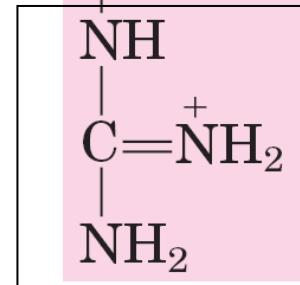
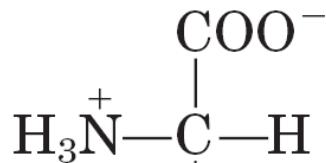


Basic amino acids

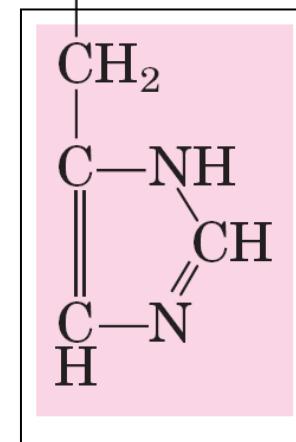
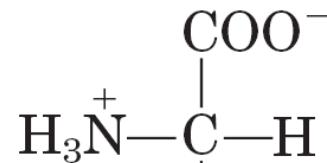
Positively charged R groups



Lysine



Arginine



Histidine

amino group

guanidino group

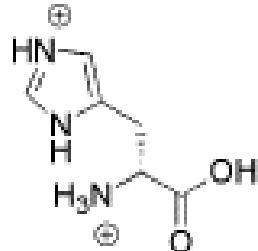
Imidazole group 36

Properties of the Amino Acid Side Chains

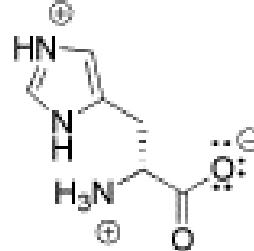
- Lysine ($pK_a = 10.0$) and arginine ($pK_a = 12.5$) are the more basic amino acids.
- Their side chains are almost always positively charged under physiological conditions.

Histidine

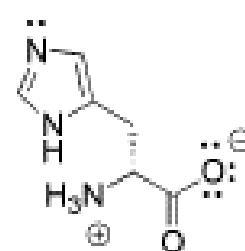
- Histidine contains a side chain imidazole. At the physiological pH (pH ~7.4), the side chains of histidine is only partially charged



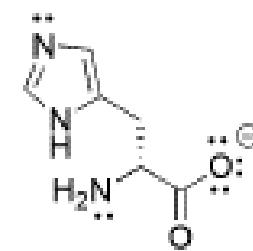
at pH = 1



at pH = 4

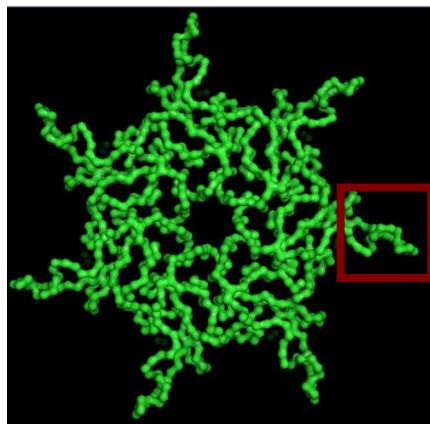


at pH = 8

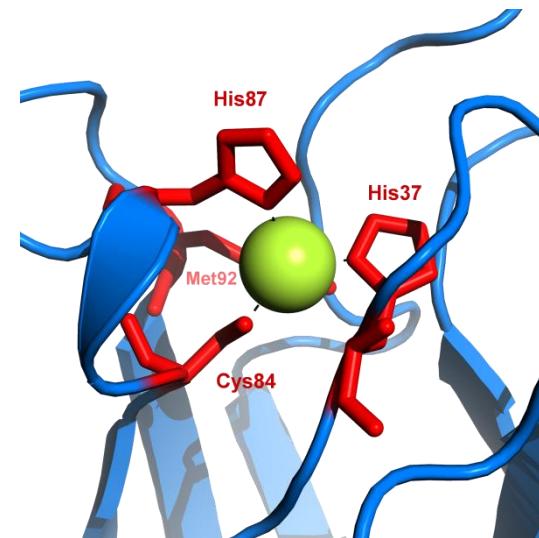
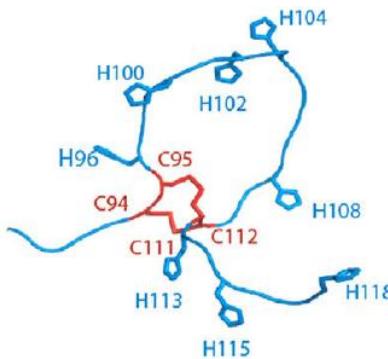


at pH = 11

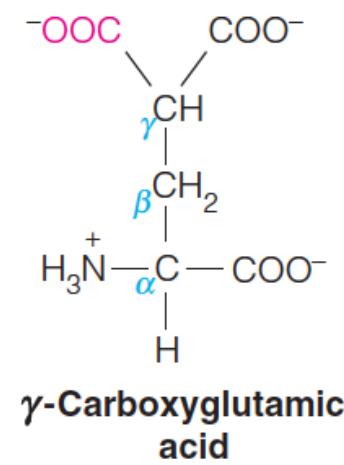
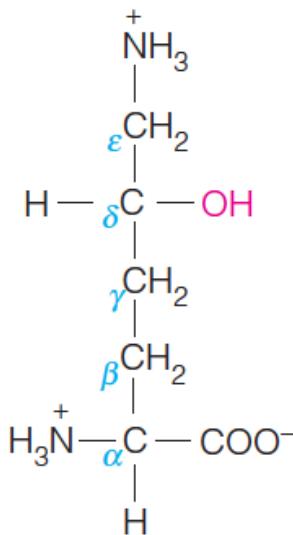
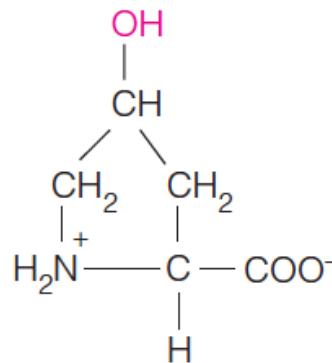
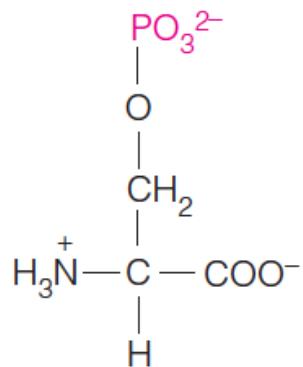
- The imidazole side chain of histidine is a common coordinating ligand in metalloproteins and is a part of catalytic sites in certain enzymes.



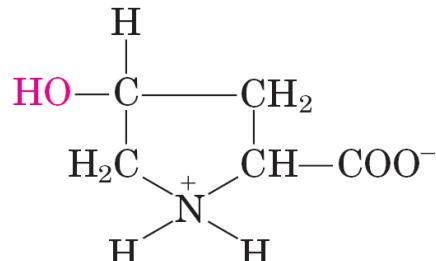
GroES of *H. pylori*



Amino acids can undergo ***post-translational modification*** resulting in modified amino acids with unique properties

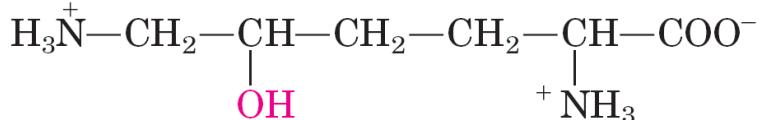


Nonstandard amino acids



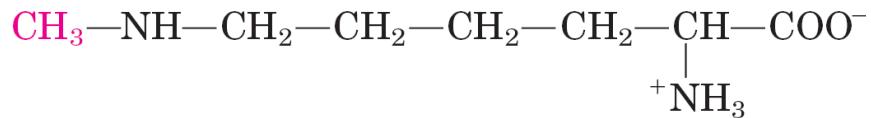
4-Hydroxyproline

plant cell wall connective tissue



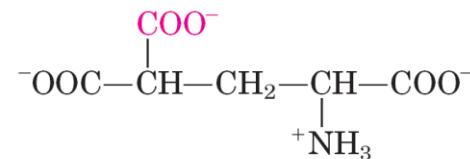
5-Hydroxylysine

collagen



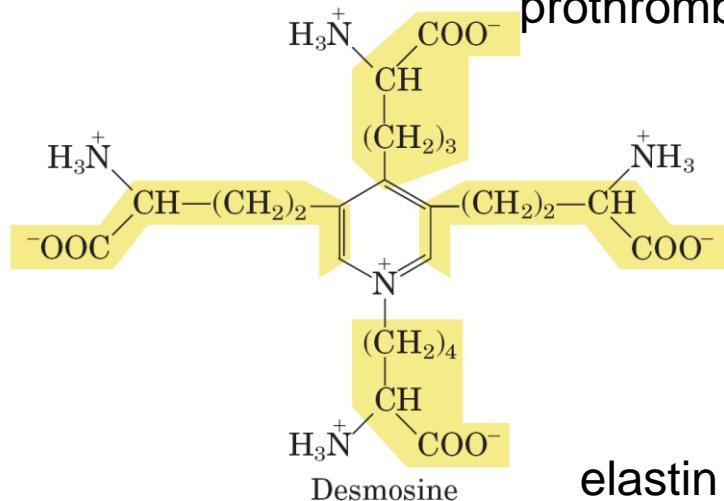
6-N-Methyllysine

myosin



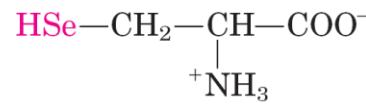
γ-Carboxyglutamate

blood-clotting protein
prothrombin



elastin

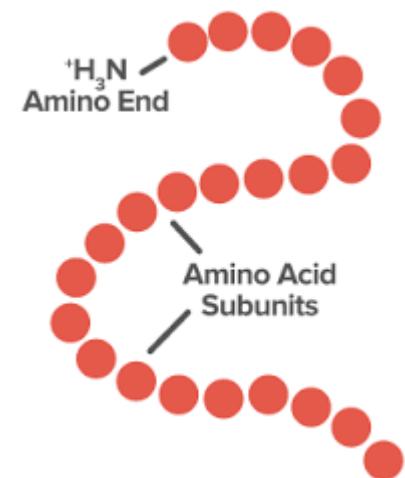
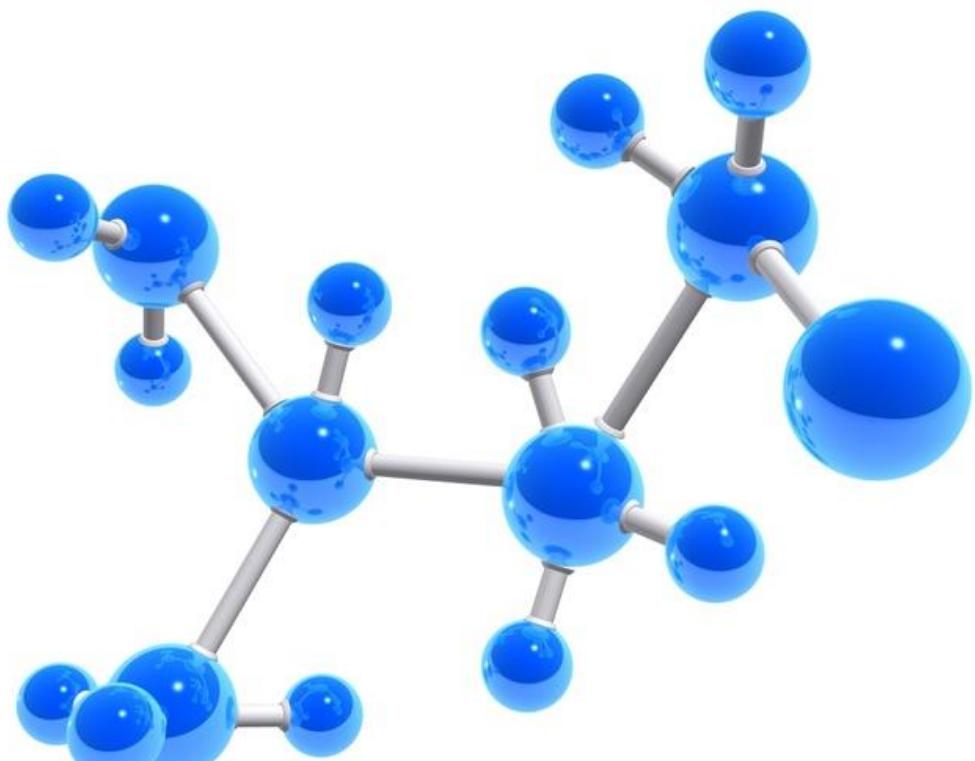
Desmosine



Selenocysteine

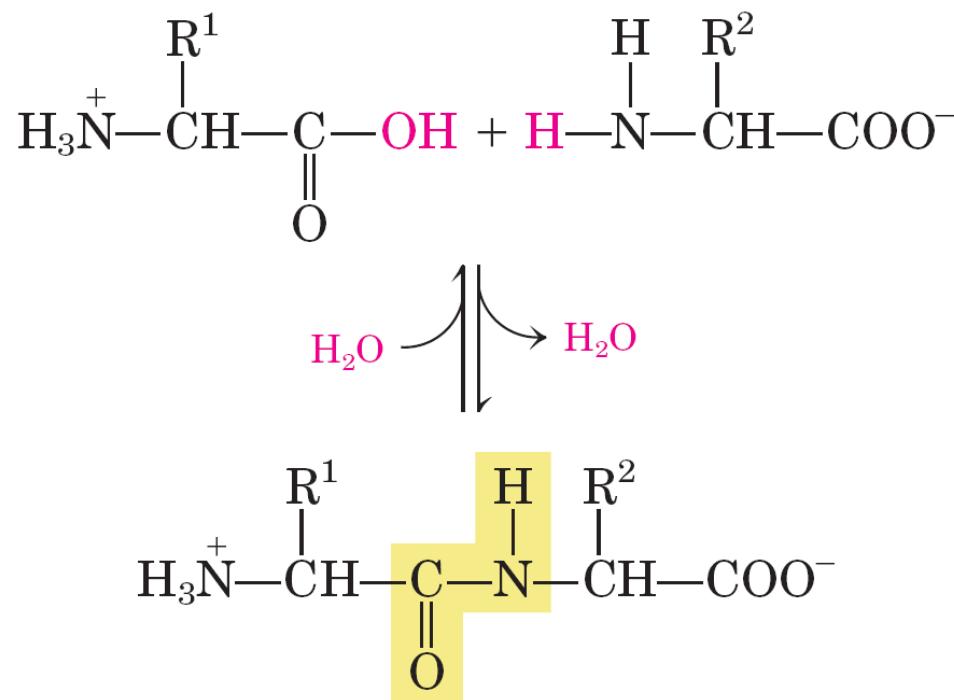
glutathione peroxidase

Peptides



Peptides: polymers of amino acids

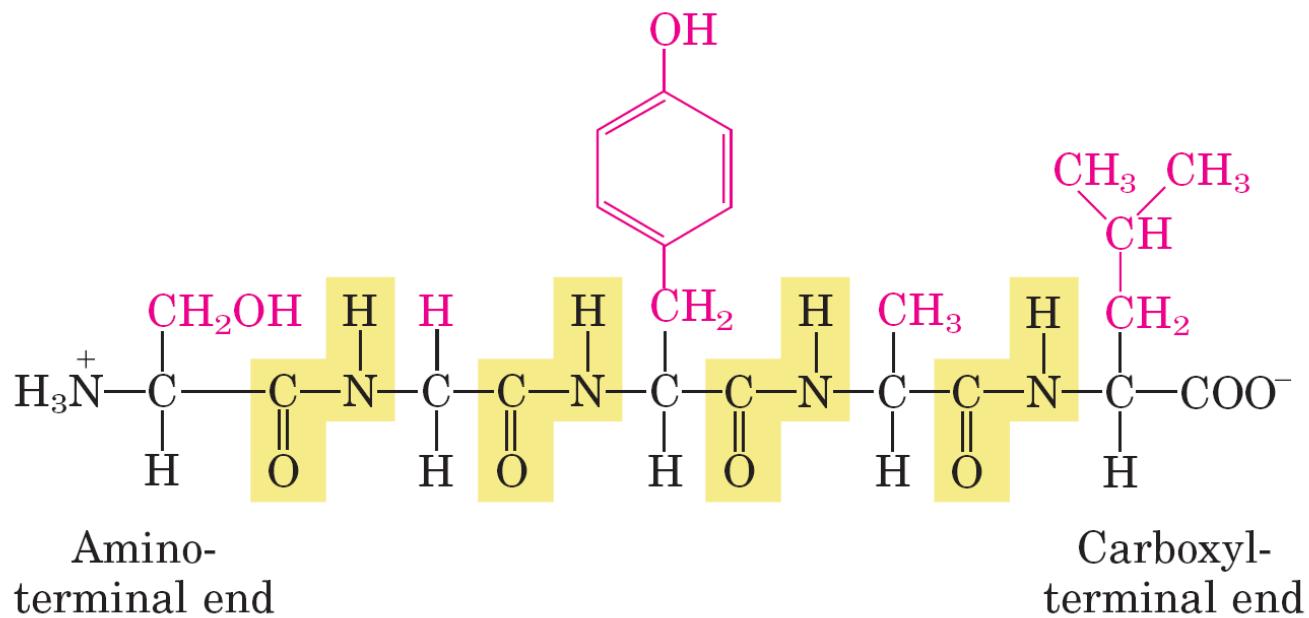
Peptide bond: two amino acid molecules can be covalently joined through a substituted amide linkage



Formation of a peptide bond by condensation

Oligopeptide: a few amino acids are joined, the structure is called an oligopeptide

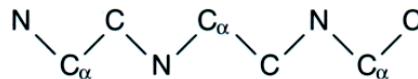
Polypeptide: many amino acids are joined, the product is called polypeptide



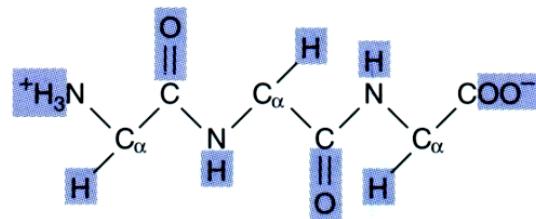
The pentapeptide serylglycyltyrosylalanylleucine, or Ser-Gly-Tyr-Ala-Leu

How to draw the peptide structure

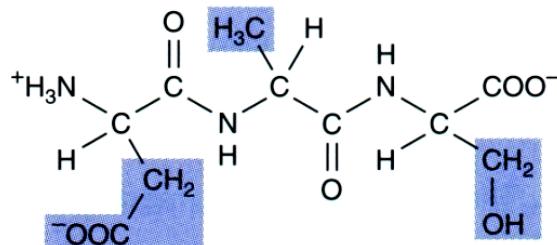
Write a zigzag formed from the repeating sequence of main chain, or “backbone,” atoms: α -nitrogen, α -carbon, carbonyl carbon.



Complete the amino and carboxyl terminals, add a hydrogen to each α -carbon and to each peptide nitrogen, and add oxygen to the carbonyl carbon.



Add appropriate R groups (shaded) to each α -carbon atom.



Abbreviations

Glu-Ala-Lys-Gly-Tyr-Ala

E A K G Y A

	pK_a values		
Amino acid	pK_1 ($-COOH$)	pK_2 ($-NH_3^+$)	pK_R (R group)
Nonpolar, aliphatic R groups			
Glycine			
Glycine	2.34	9.60	
Alanine	2.34	9.69	
Proline	1.99	10.96	
Valine	2.32	9.62	
Leucine	2.36	9.60	
Isoleucine	2.36	9.68	
Methionine	2.28	9.21	
Aromatic R groups			
Phenylalanine	1.83	9.13	
Tyrosine	2.20	9.11	10.07
Tryptophan	2.38	9.39	
Polar, uncharged R groups			
Serine	2.21	9.15	
Threonine	2.11	9.62	
Cysteine	1.96	10.28	8.18
Asparagine	2.02	8.80	
Glutamine	2.17	9.13	
Positively charged R groups			
Lysine	2.18	8.95	10.53
Histidine	1.82	9.17	6.00
Arginine	2.17	9.04	12.48
Negatively charged R groups			
Aspartate	1.88	9.60	3.65
Glutamate	2.19	9.67	4.25

Net Electric Charge of Peptides



9.67 4.25 6.0 12.48 2.34

+3 +2 +1 0 -1 -2

$$\begin{array}{ccccc} 2.34 & 4.25 & 6.0 & 9.67 & 12.48 \\ \text{pK}_1 & \text{pK}_2 & \text{pK}_3 & \text{pK}_4 & \text{pK}_5 \end{array}$$

pH=3	+2
pH=8	0
pH=11	-1

$$pI = \frac{6.0 + 9.67}{2} = 7.835$$

Many peptides have physiologic activity

A **neuropeptide** is a peptide that is active in association with neural tissue.

A **lipopeptide** is a peptide that has a lipid connected to it, and **pepducins** are new modulators of G protein–coupled receptors (GPCRs)

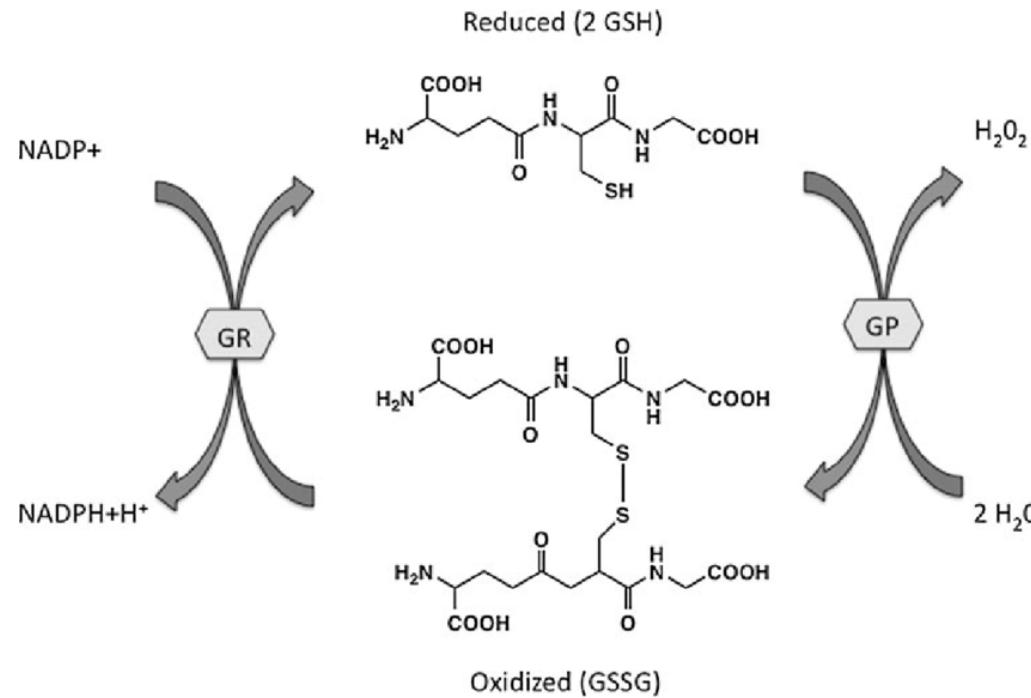
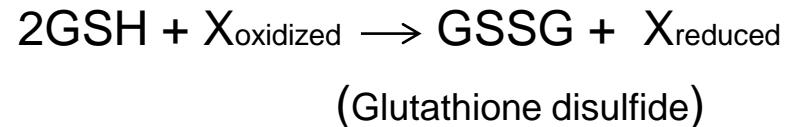
A **peptide hormone** is a peptide that acts as a hormone.

A **proteose** is a mixture of peptides produced by the hydrolysis of proteins.

Many peptides have physiologic activity

a. Glutathione (GSH)

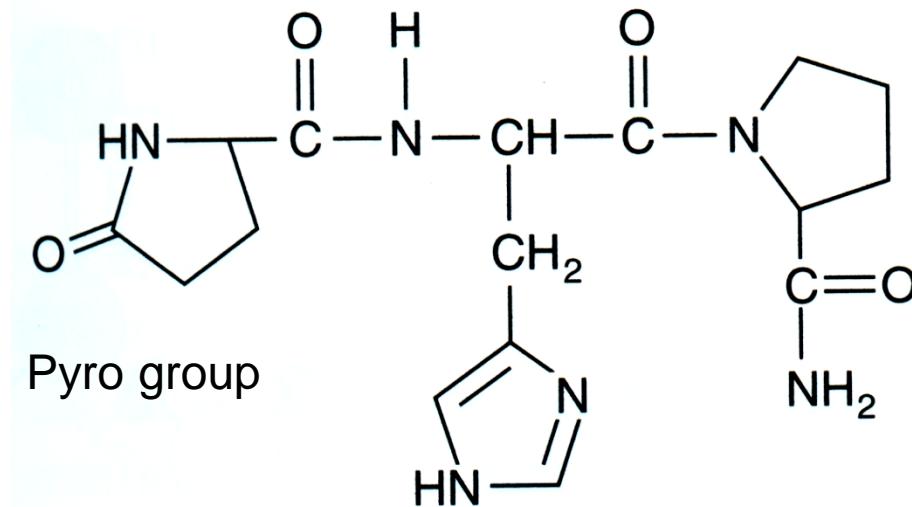
γ -Glu-Cys-Gly (tripeptide)



Glutathione (GSH) is an important antioxidant agent.

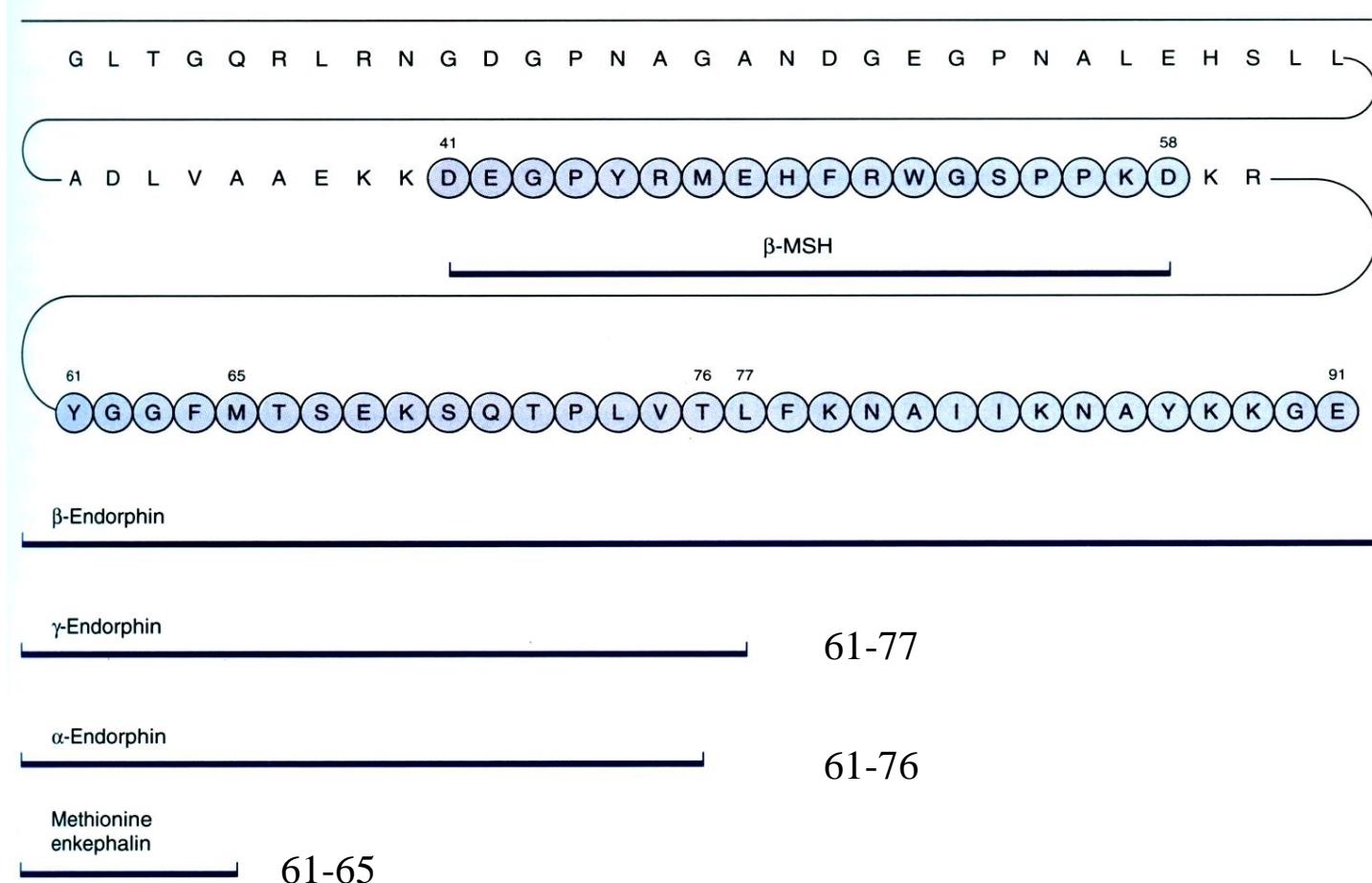
b. TRH (Thyrotropin-releasing hormone)

<Glu-His-Pro-NH₂

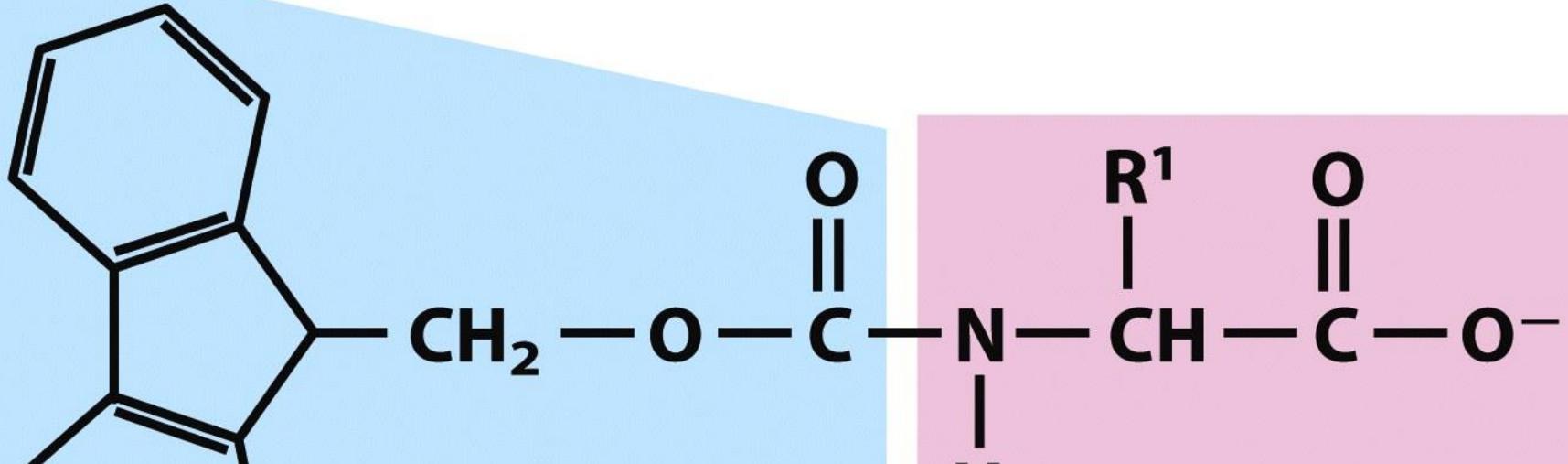


由下視丘（Hypothalamus）所製造的肽類激素（Peptide Hormone），能夠調節腦下垂體前葉分泌促甲狀腺激素（甲促素，thyroid stimulating hormone、TSH）與催乳素

c. β -lipotropin



腦垂體分泌的一種激素，可分解脂肪



Fmoc

**Amino acid
residue**

Figure 3-29a

Lehninger Principles of Biochemistry, Fifth Edition

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Solid phase synthesis of peptide



R. Bruce Merrifield
1921–2006

Unnumbered p 101
Lehninger Principles of Biochemistry, Fifth Edition
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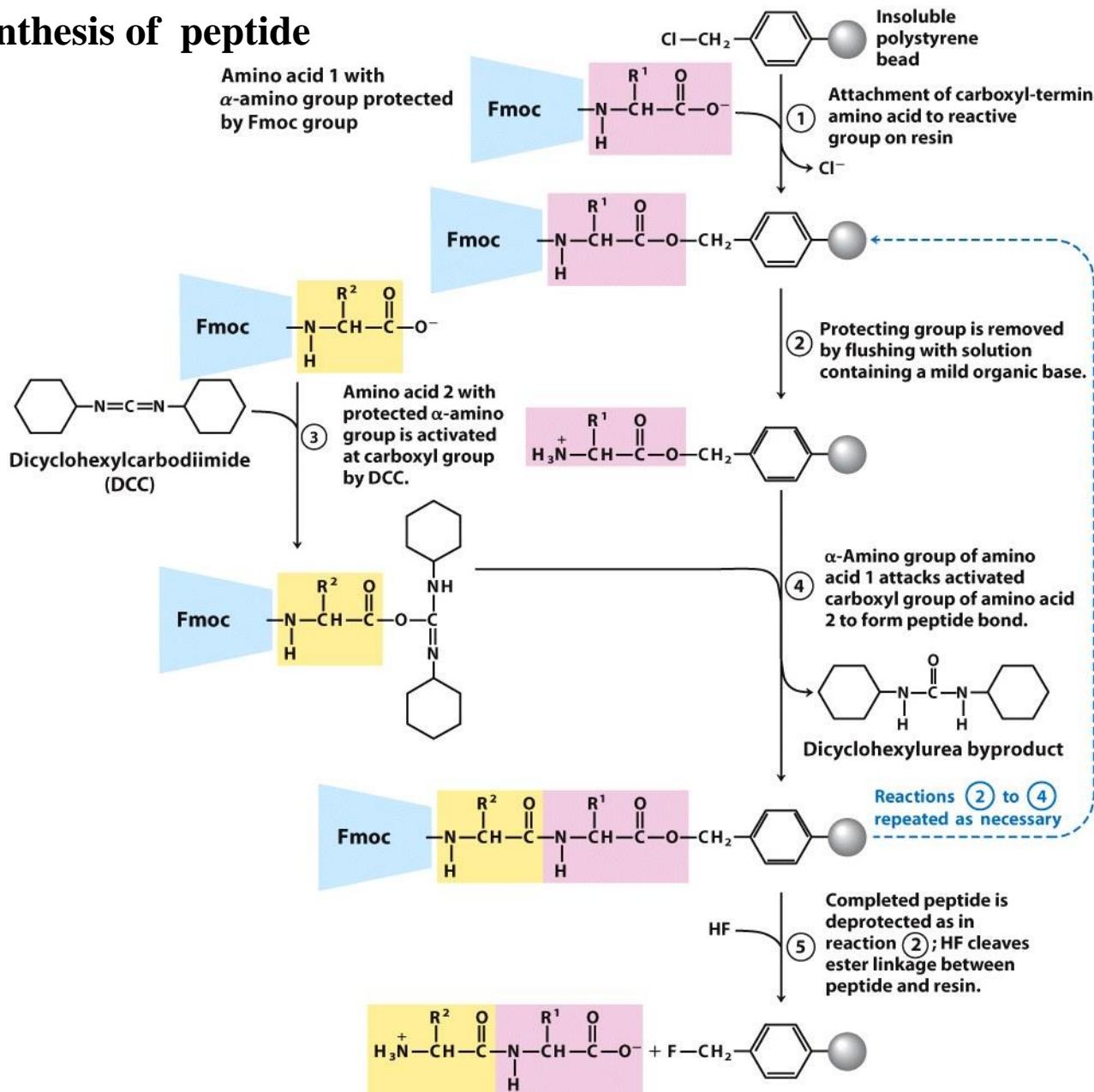
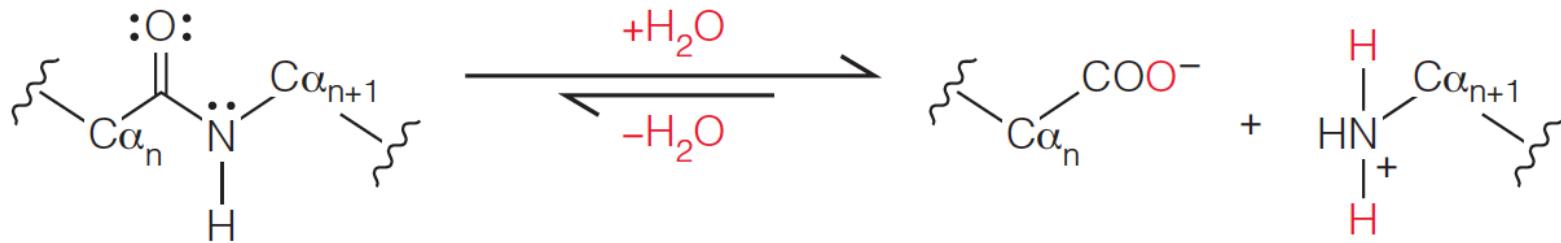


Figure 3-29b

Lehninger Principles of Biochemistry, Fifth Edition

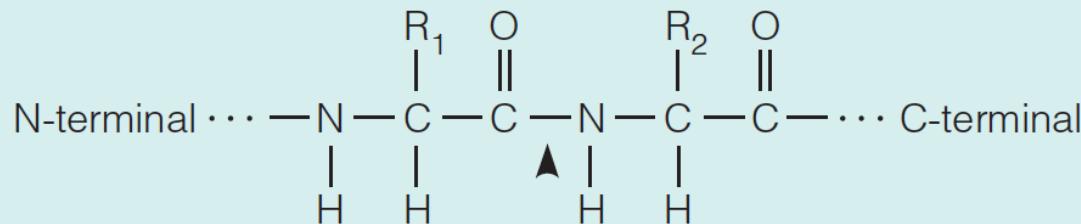
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Peptides and the peptide bond



- While hydrolysis of peptide bonds is thermodynamically favored in aqueous solutions, the reaction is exceedingly slow at physiological pH and temperature
- Peptide bond hydrolysis can be achieved by:
 - Strong mineral acid (e.g., 6 M HCl) cleaves all peptide bonds (including the Asn and Gln amide bonds)
 - Chemicals that cleave at specific sites (e.g., CNBr cleaves at Met)
 - Proteolytic enzymes (proteases) that cleave at specific sites

Sequence specificities of some proteolytic enzymes



Enzyme	Preferred Site ^a	Source
Trypsin	R ₁ = Lys, Arg 53	From digestive systems of animals, many other sources
Chymotrypsin	R ₁ = Tyr, Trp, Phe, Leu	Same as trypsin
Thrombin	R ₁ = Arg	From blood; involved in coagulation
V-8 protease	R ₁ = Asp, Glu	From <i>Staphylococcus aureus</i>
Prolyl endopeptidase	R ₁ = Pro	Lamb kidney, other tissues
Subtilisin	Very little specificity	From various bacilli
Carboxypeptidase A	R ₂ = C-terminal amino acid	From digestive systems of animals
Thermolysin	R ₂ = Leu, Val, Ile, Met	From <i>Bacillus thermoproteolyticus</i>

Separation of peptides

