

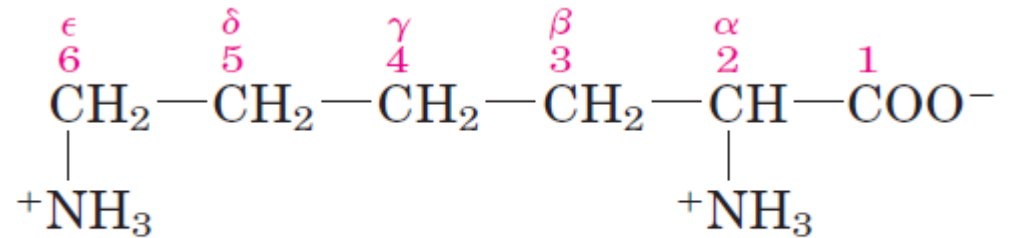
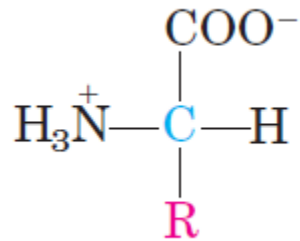
Disciplina: SLC0673

**Aminoácidos, proteínas e
enzimas**

Prof. Dr. Andrei Leitão

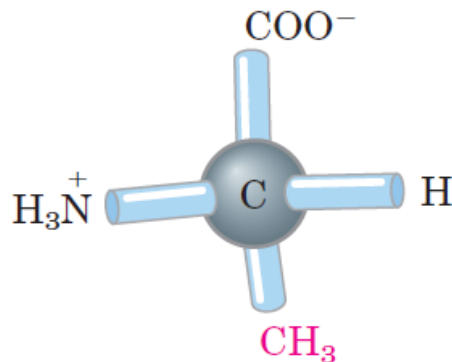
Amino acids

General structure of an amino acid.

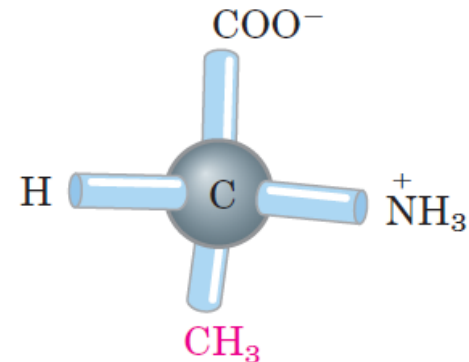


Lysine

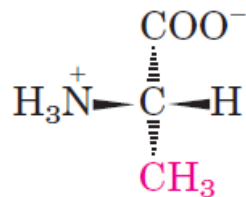
Stereoisomerism



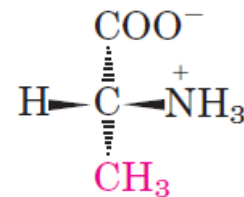
(a) L-Alanine



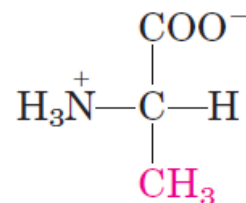
D-Alanine



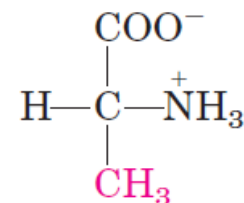
(b) L-Alanine



D-Alanine



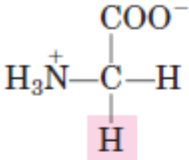
(c) L-Alanine



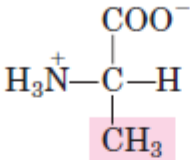
D-Alanine

Nonpolar & aromatic amino acids

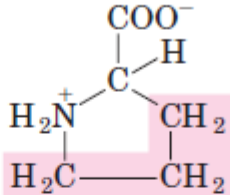
Nonpolar, aliphatic R groups



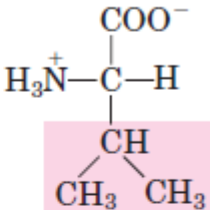
Glycine



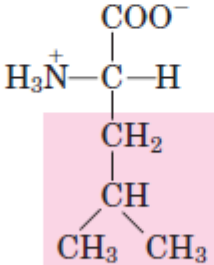
Alanine



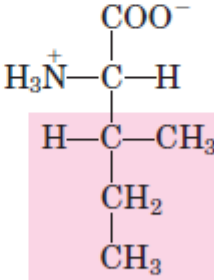
Proline



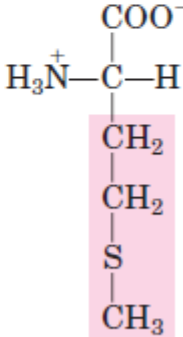
Valine



Leucine

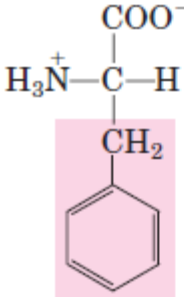


Isoleucine

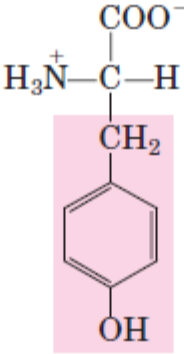


Methionine

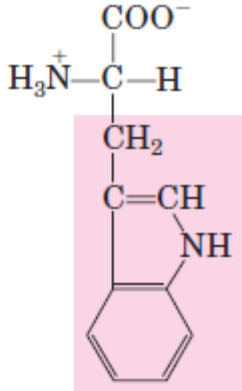
Aromatic R groups



Phenylalanine



Tyrosine



Tryptophan

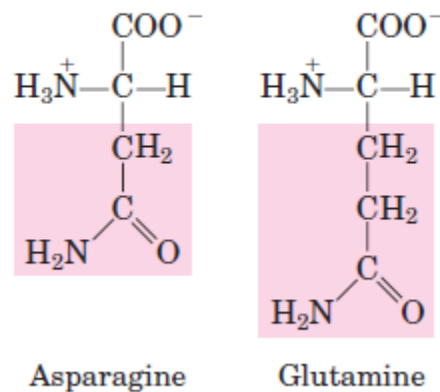
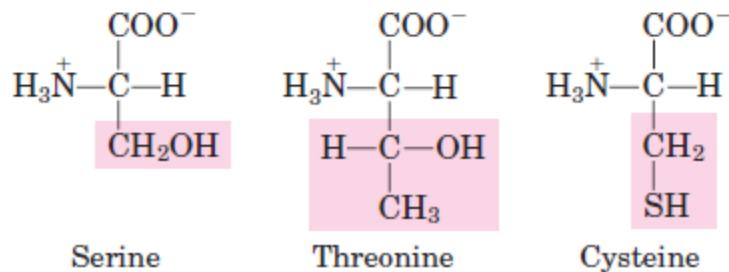
Properties

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

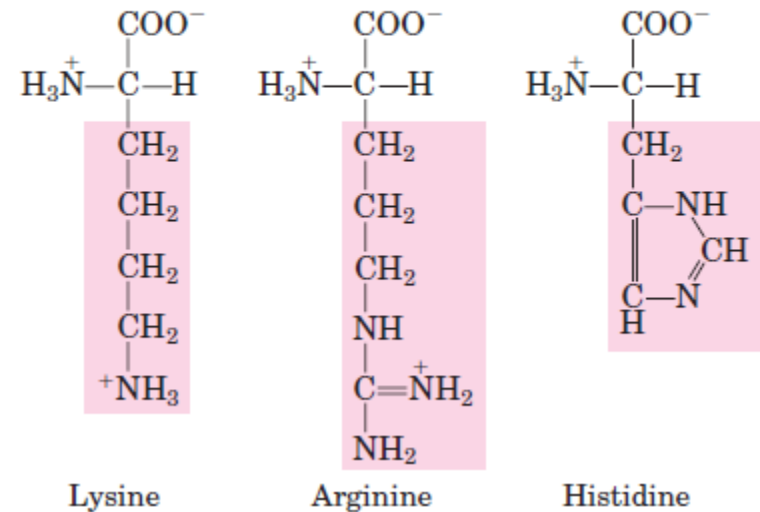
Amino acid	Abbreviation/ symbol	M_r	pK_a values			pI	Hydropathy index*	Occurrence in proteins (%) [†]
			pK_1 (—COOH)	pK_2 (—NH ₃ ⁺)	pK_R (R group)			
Nonpolar, aliphatic R groups								
Glycine	Gly G	75	2.34	9.60		5.97	-0.4	7.2
Alanine	Ala A	89	2.34	9.69		6.01	1.8	7.8
Proline	Pro P	115	1.99	10.96		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 & charged amino acids

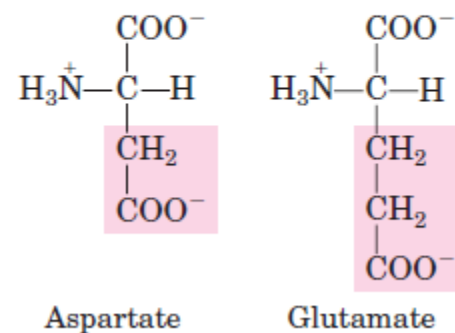
Polar, uncharged R groups



Positively charged R groups



Negatively charged R groups



Properties

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

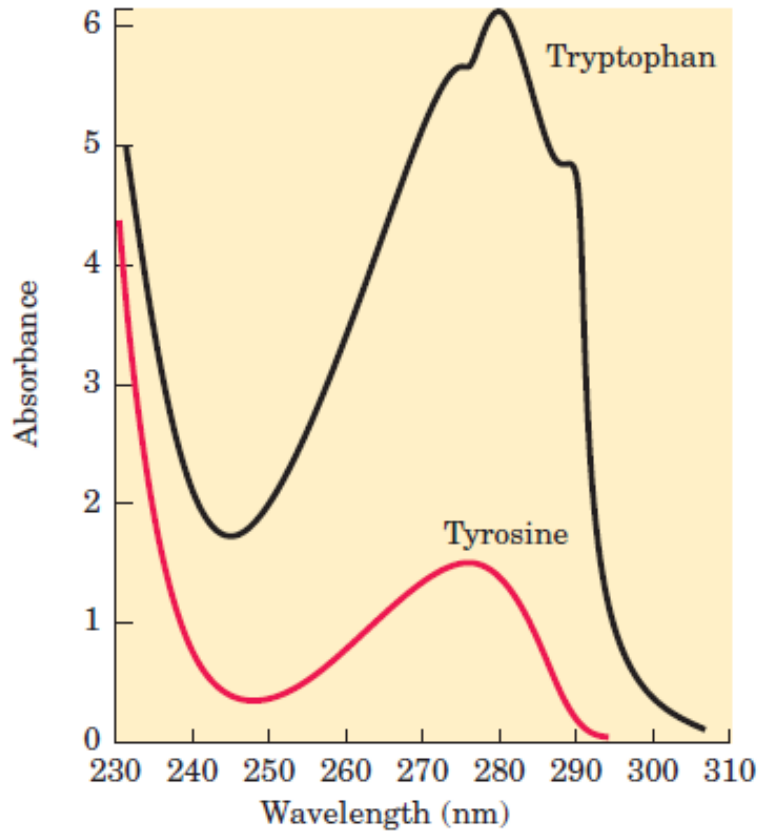
Amino acid	Abbreviation/ symbol	M_r	pK_a values			pI	Hydropathy index*	Occurrence in proteins (%) [†]
			pK_1 (—COOH)	pK_2 (—NH ₃ ⁺)	pK_R (R group)			
Polar, uncharged								
R groups								
Serine	Ser S	105	2.21	9.15		5.68	-0.8	6.8
Threonine	Thr T	119	2.11	9.62		5.87	-0.7	5.9
Cysteine	Cys C	121	1.96	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

*A scale combining hydrophobicity and hydrophilicity of R groups; it can be used to measure the tendency of an amino acid to seek an aqueous environment (- values) or a hydrophobic environment (+ values). See Chapter 11. From Kyte, J. & Doolittle, R.F. (1982) A simple method for displaying the hydropathic character of a protein. *J. Mol. Biol.* **157**, 105-132.

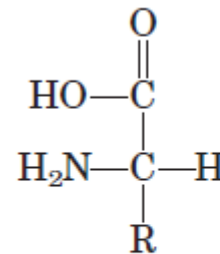
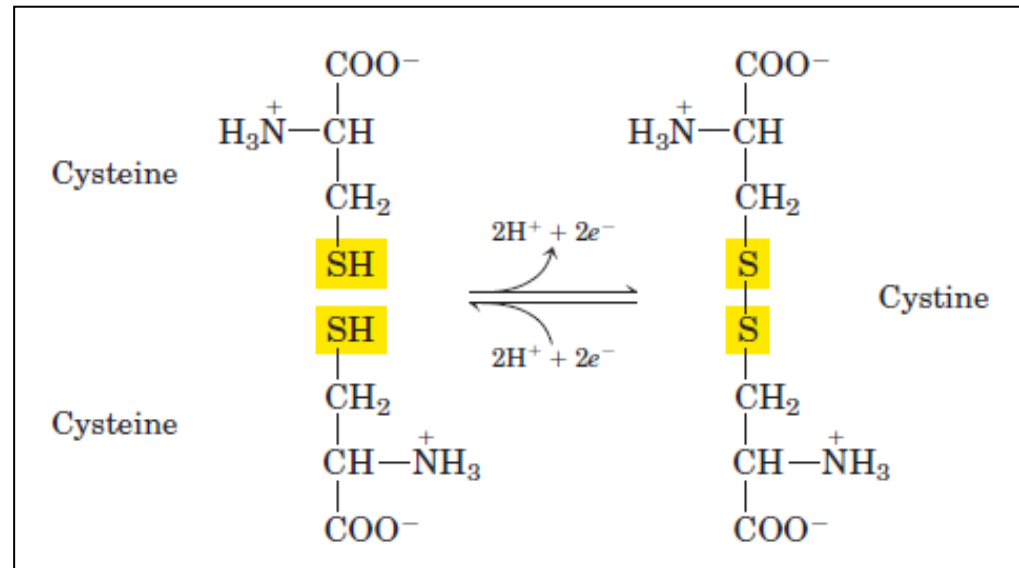
[†]Average occurrence in more than 1,150 proteins. From Doolittle, R.F. (1989) Redundancies in protein sequences. In *Prediction of Protein Structure and the Principles of Protein Conformation* (Fasman, G.D., ed.), pp. 599-623, Plenum Press, New York.

Properties

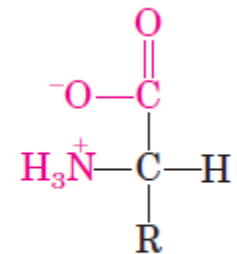
Absorption of ultraviolet light by aromatic amino acids.



Disulfide bonds



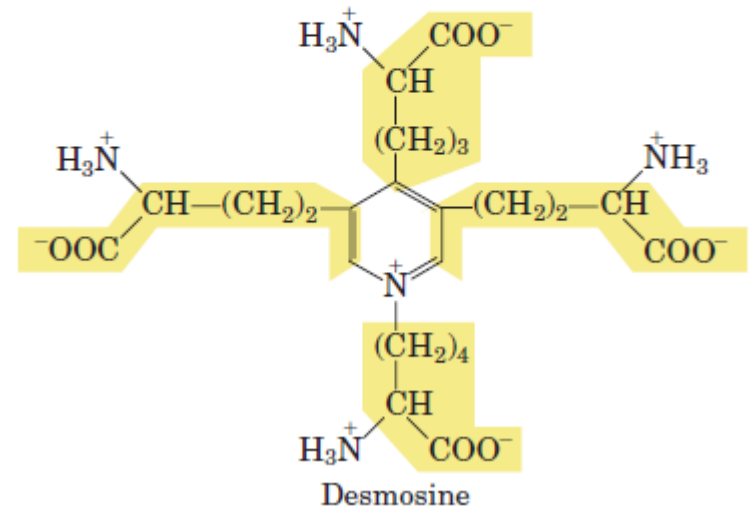
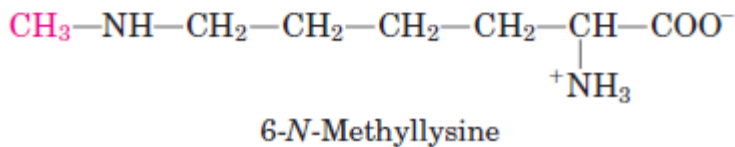
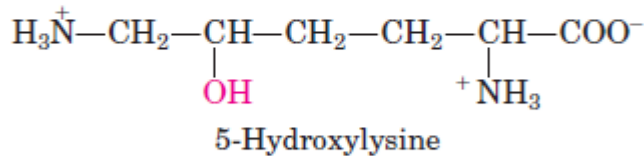
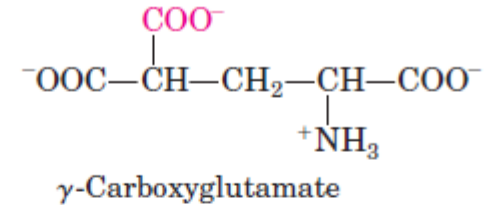
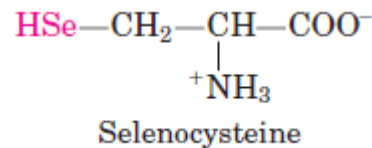
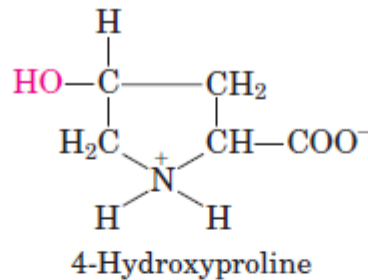
Nonionic form



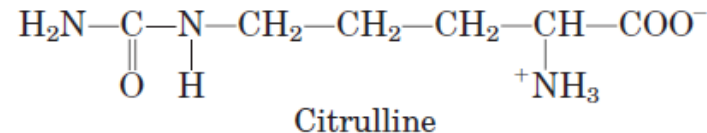
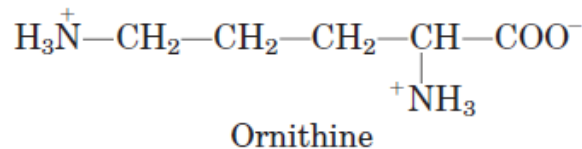
Zwitterionic form

Other types of amino acids

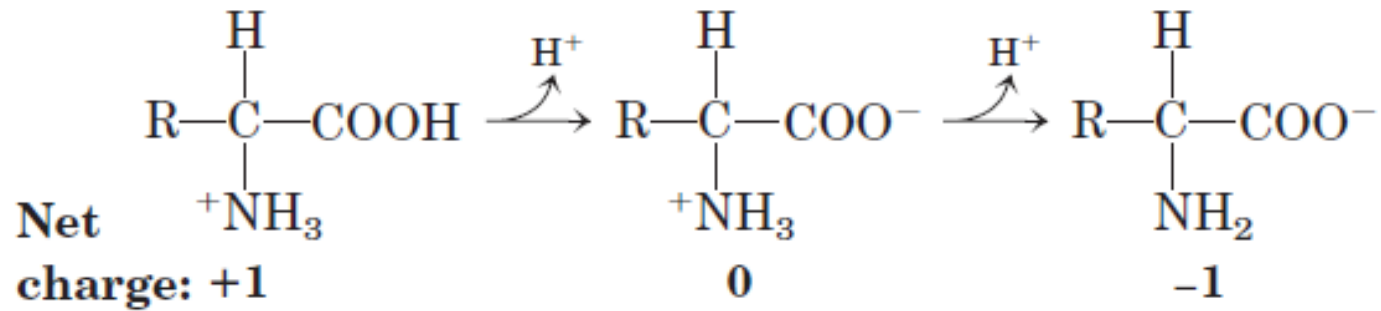
Uncommon amino acids.



Precursors



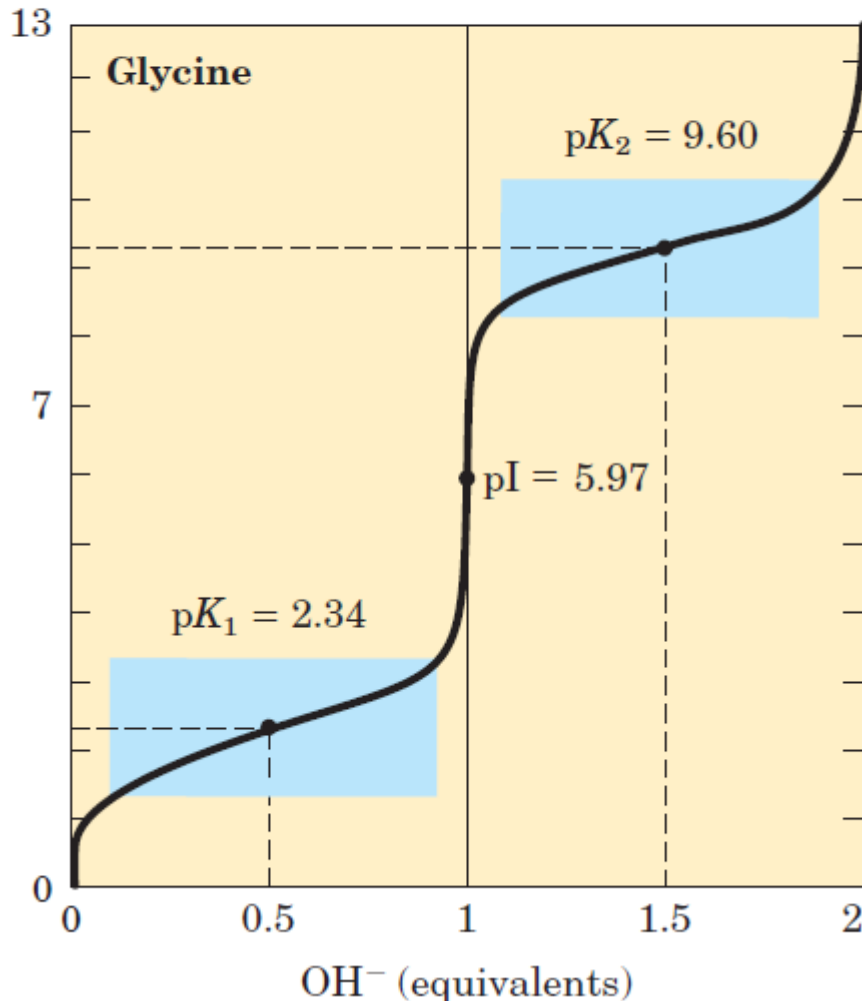
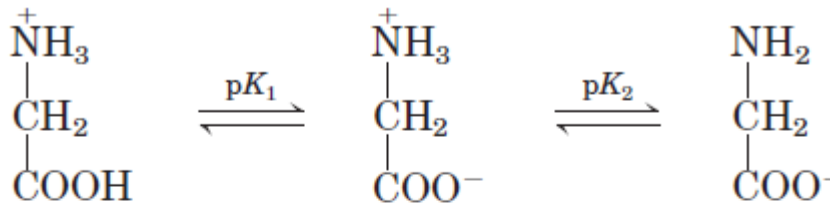
Zwitterions



Amphoters

0.1 M glycine at 25 °C.

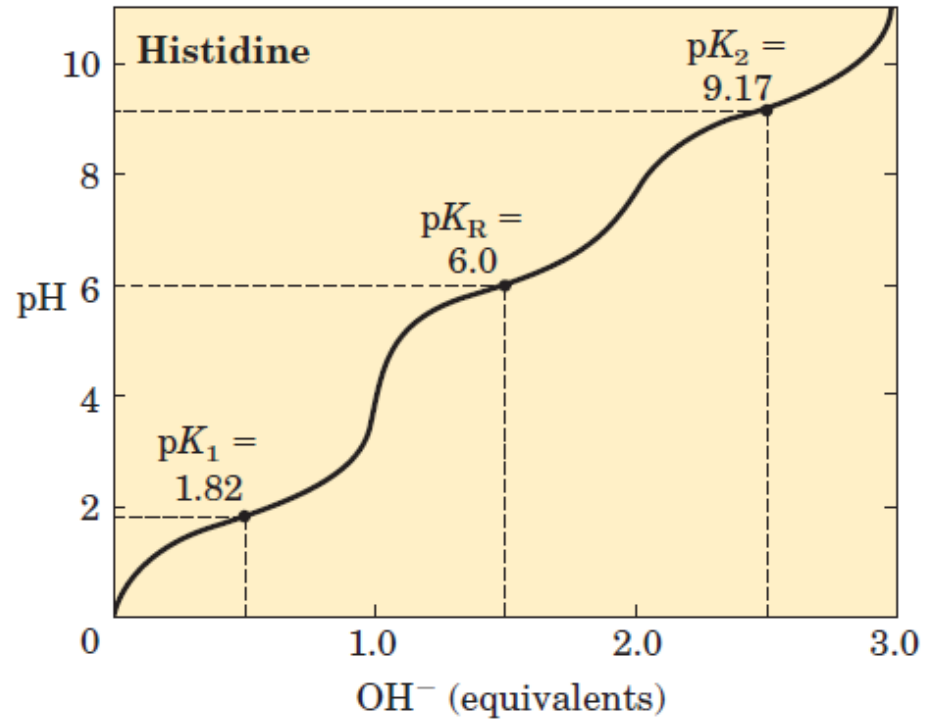
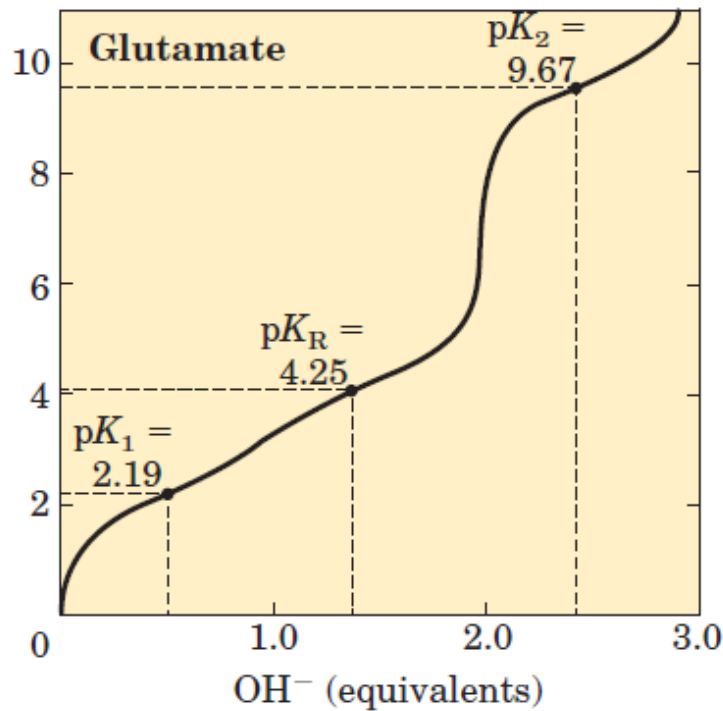
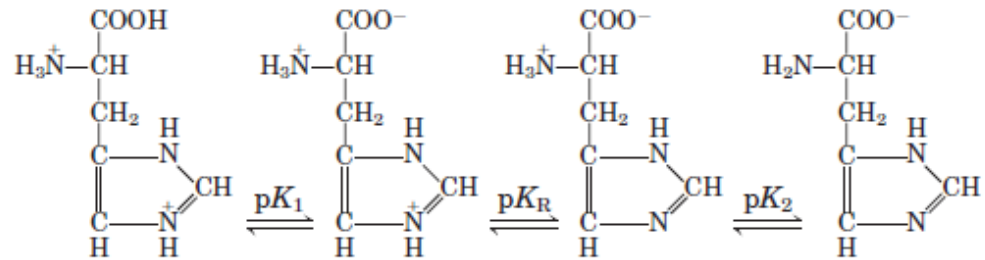
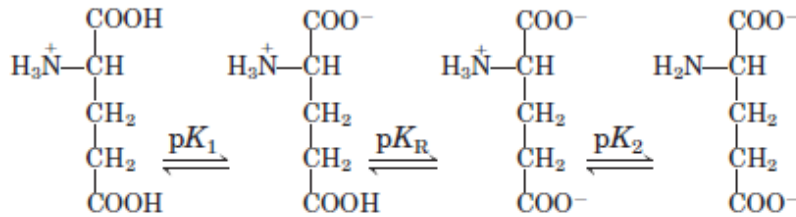
Titration curve



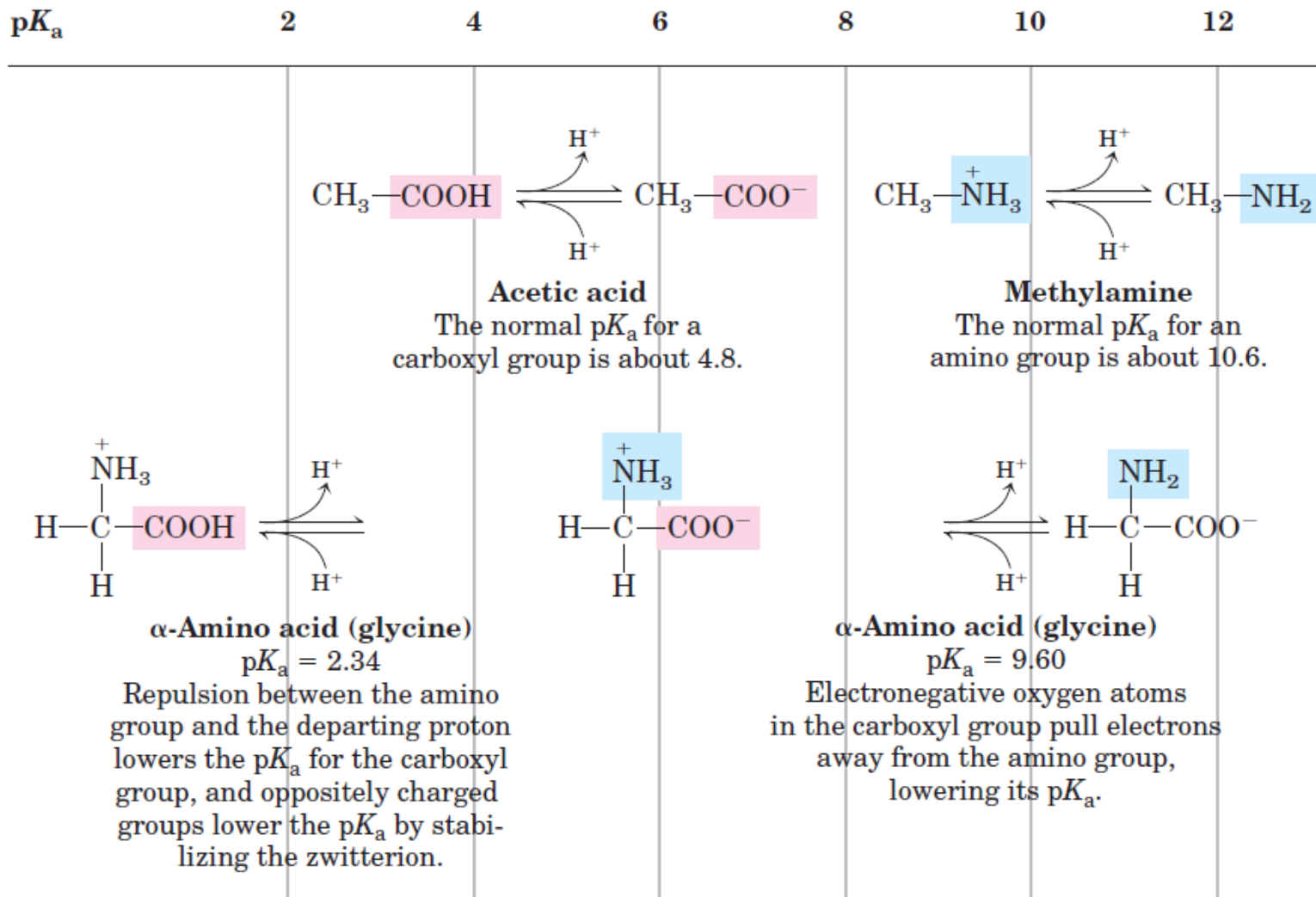
isoelectric point or isoelectric pH, designated **pI**.

$$\text{pI} = \frac{1}{2} (\text{p}K_1 + \text{p}K_2) = \frac{1}{2} (2.34 + 9.60) = 5.97$$

Titration curves

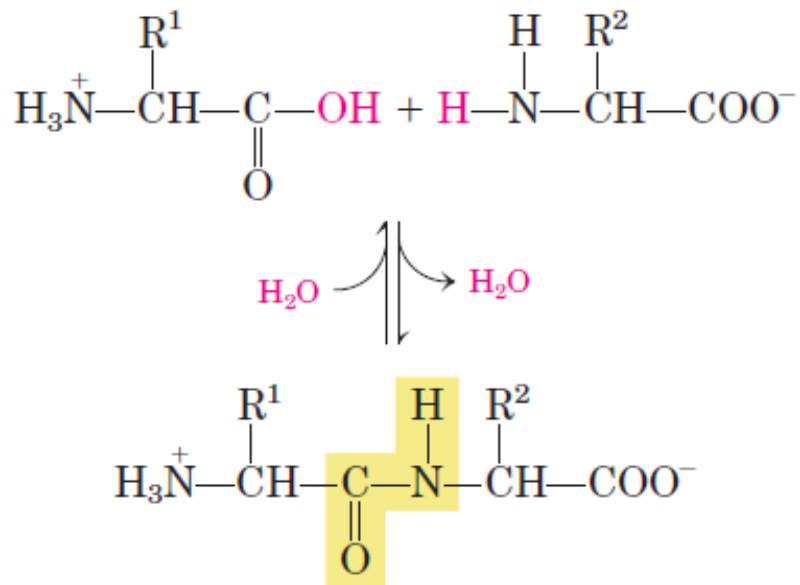


Effect of the chemical environment

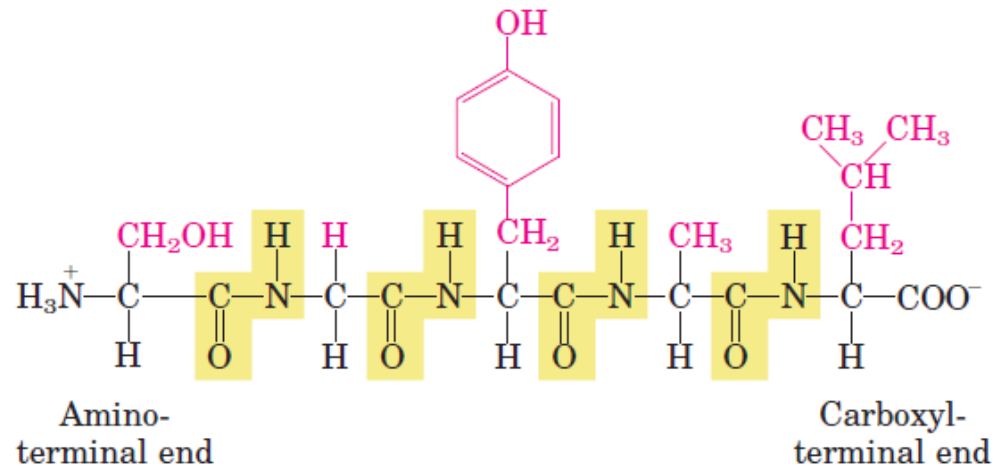


Peptides

Condensation reaction

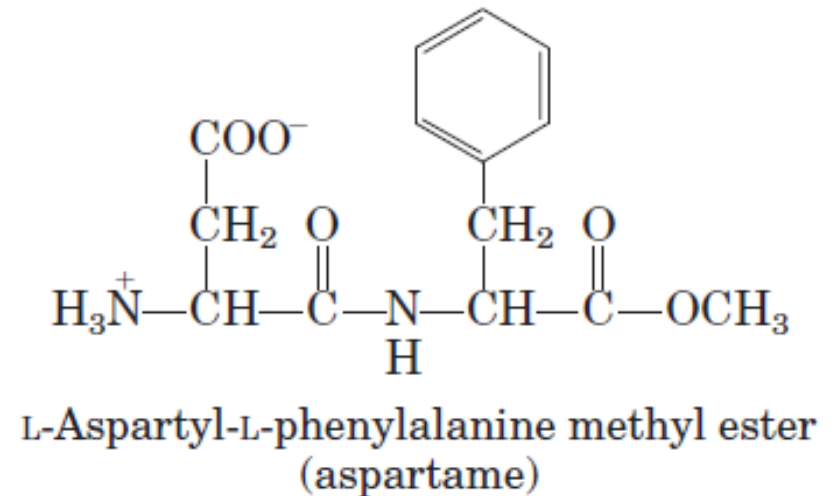
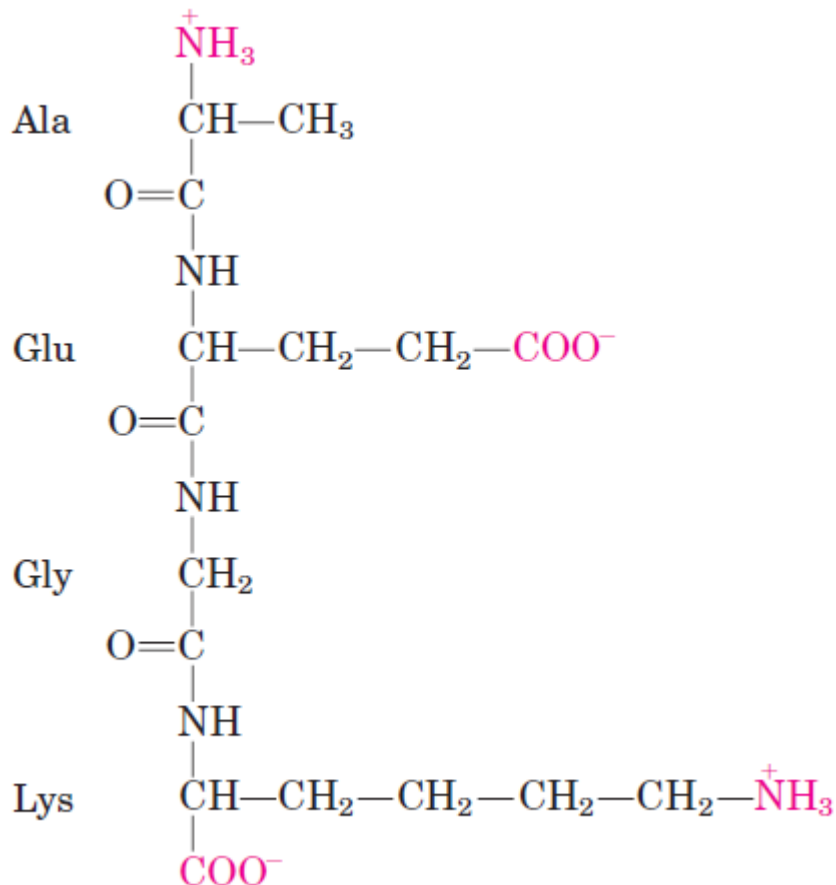


Pentapeptide



Peptides

Alanylglutamylglycyllysine.



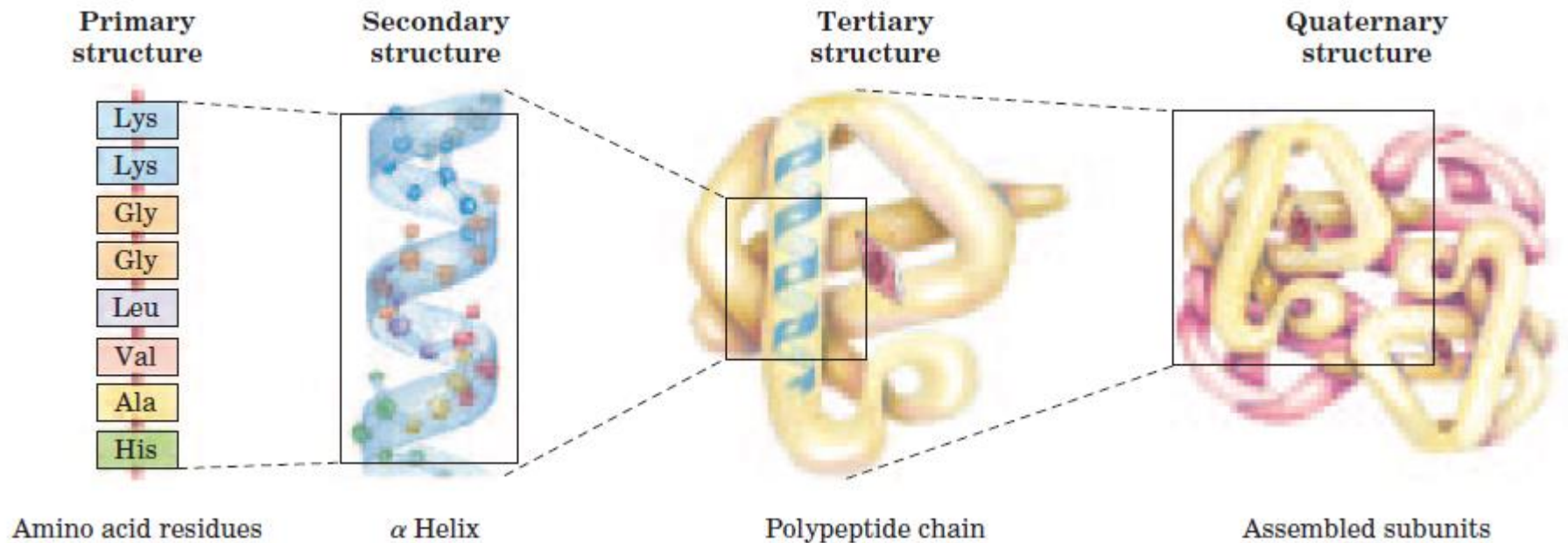
The groups ionized at pH 7.0 are in red.

Proteins

TABLE 3-2 Molecular Data on Some Proteins

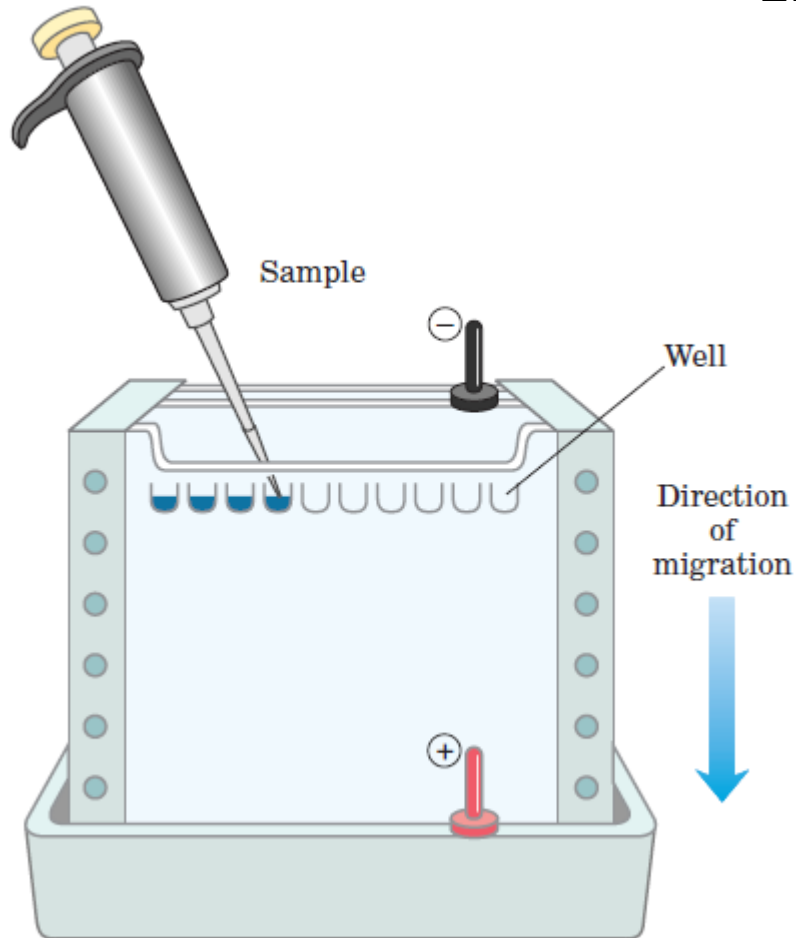
	<i>Molecular weight</i>	<i>Number of residues</i>	<i>Number of polypeptide chains</i>
Cytochrome c (human)	13,000	104	1
Ribonuclease A (bovine pancreas)	13,700	124	1
Lysozyme (chicken egg white)	13,930	129	1
Myoglobin (equine heart)	16,890	153	1
Chymotrypsin (bovine pancreas)	21,600	241	3
Chymotrypsinogen (bovine)	22,000	245	1
Hemoglobin (human)	64,500	574	4
Serum albumin (human)	68,500	609	1
Hexokinase (yeast)	102,000	972	2
RNA polymerase (<i>E. coli</i>)	450,000	4,158	5
Apolipoprotein B (human)	513,000	4,536	1
Glutamine synthetase (<i>E. coli</i>)	619,000	5,628	12
Titin (human)	2,993,000	26,926	1

Proteins



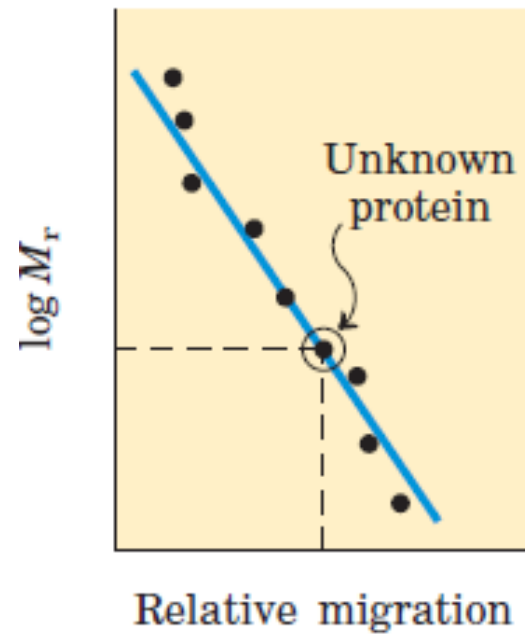
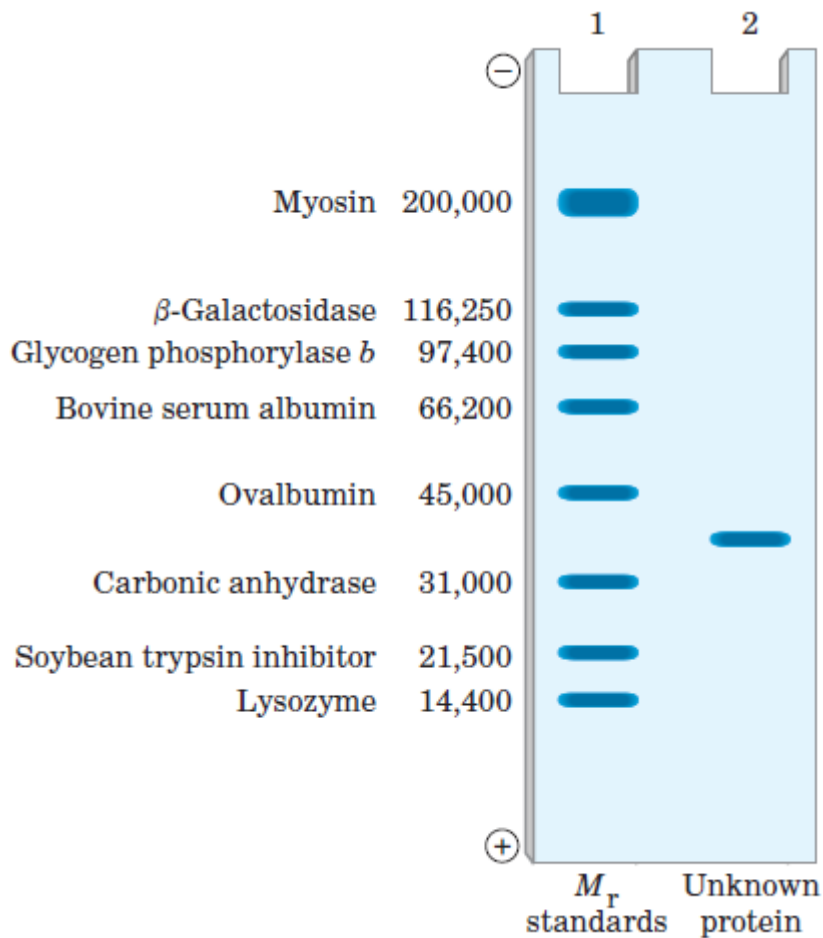
Separation of proteins

Electrophoresis



Separation of proteins

Electrophoresis



Alignment of protein sequences

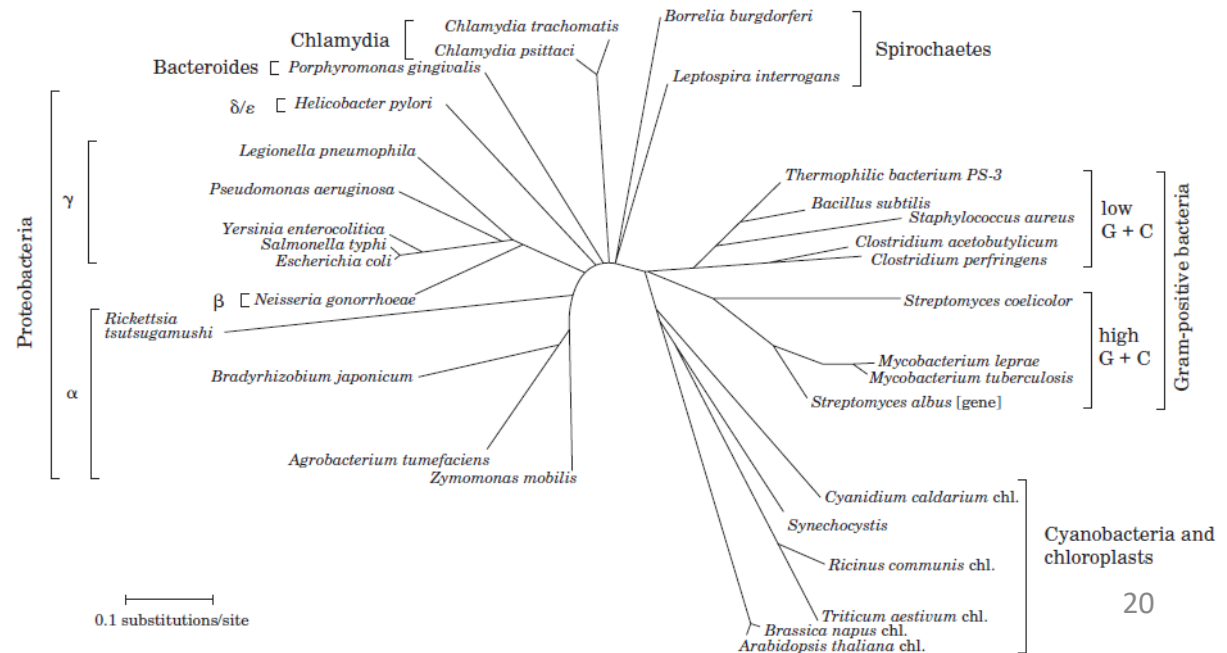
E. coli TGNRTIAVYDLGGGTFDISIIEIDEVDGEKTFEVLATNGDTHLGGEDFDSRLIHYL
B. subtilis DEDQTILLYDLGGGTFDVSILELGDG TFEVRSTAGDNRLGGDDFDQVIIDHL

Gap

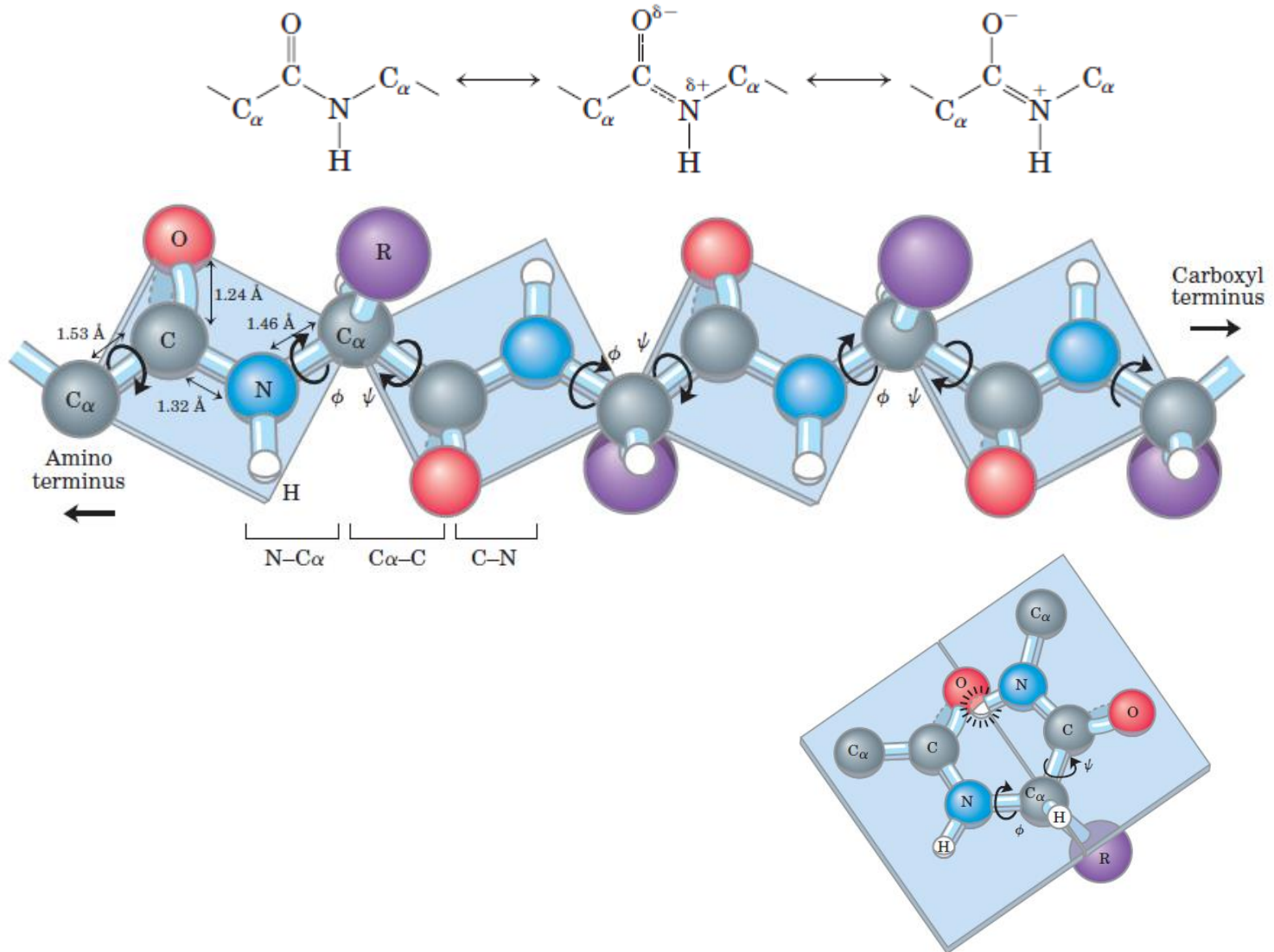
Archaeobacteria	}	<i>Halobacterium halobium</i>	IGHVDH GKST MVGR	LLYETG SVPEHV	IEQH
		<i>Sulfolobus solfataricus</i>	IGHVDH GKST LVGR	LLMDRG FIDEKT	VKEA
Eukaryotes	}	<i>Saccharomyces cerevisiae</i>	IGHVDS GKST TTGHL	LIYKCG GIDKRT	IEKF
		<i>Homo sapiens</i>	IGHVDS GKST TTGHL	LIYKCG GIDKRT	IEKF
Gram-positive bacterium		<i>Bacillus subtilis</i>	IGHVDH GKST MVGR		ITTV
Gram-negative bacterium		<i>Escherichia coli</i>	IGHVDH GKST LTAA		ITTV

Signature sequence

Evolutionary tree derived from amino acid sequence comparisons.

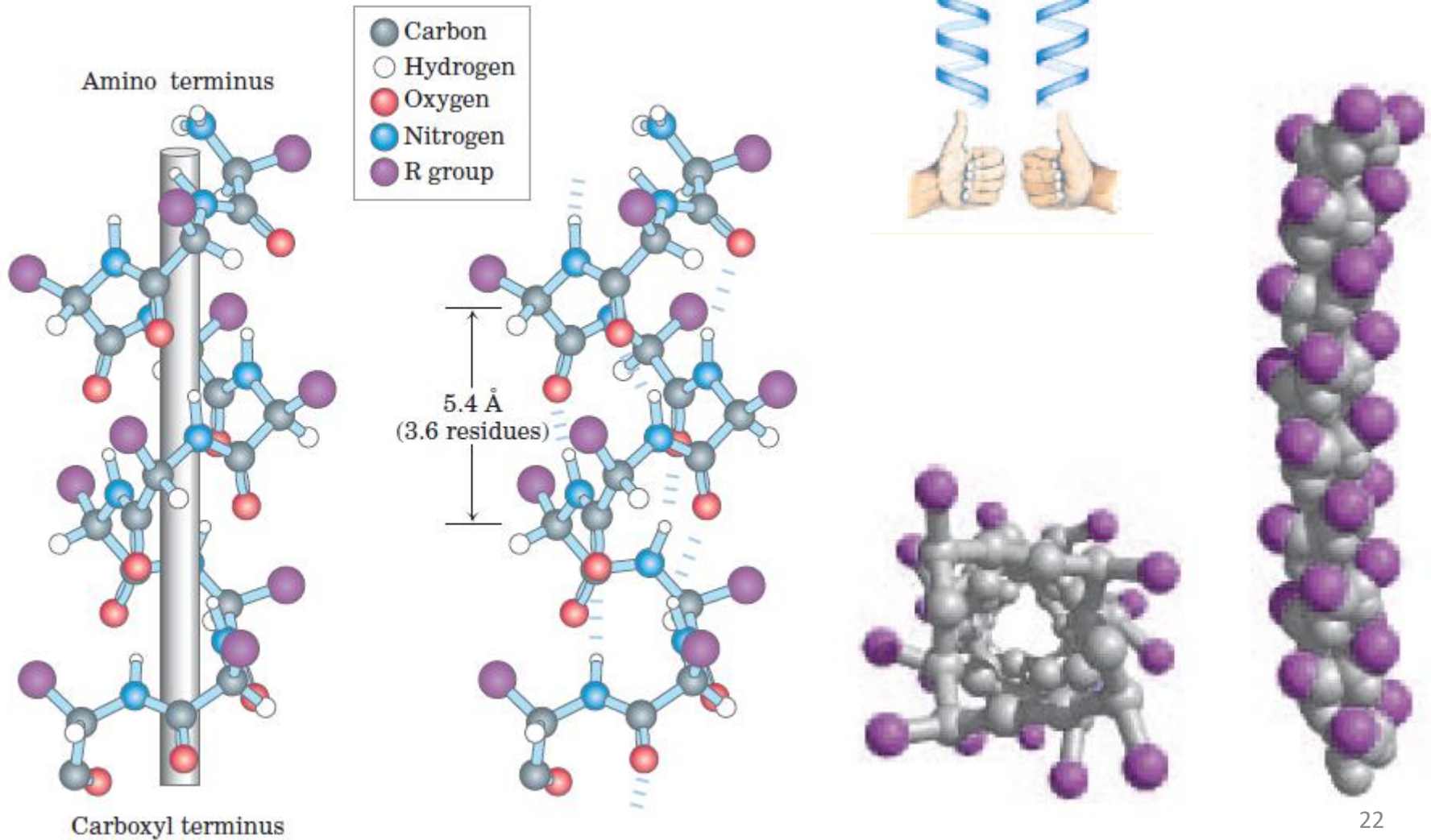


3D structure of proteins



3D structure of proteins

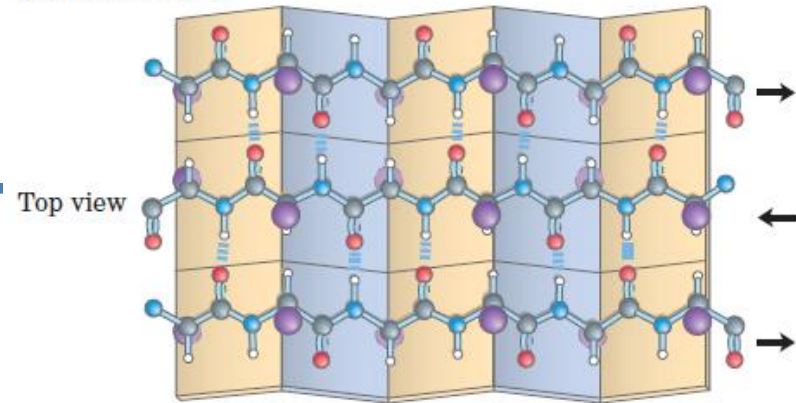
Four models of the α helix



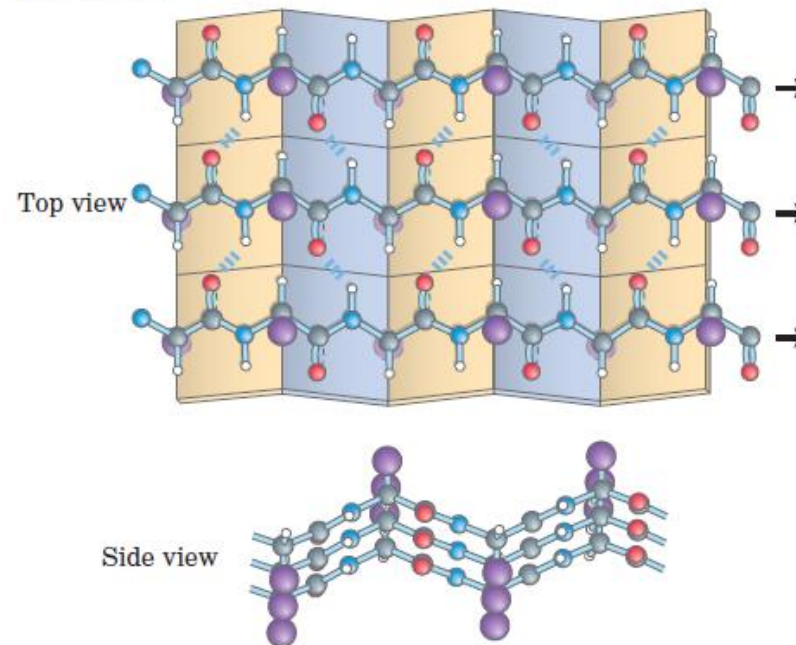
3D structure of proteins

β sheet

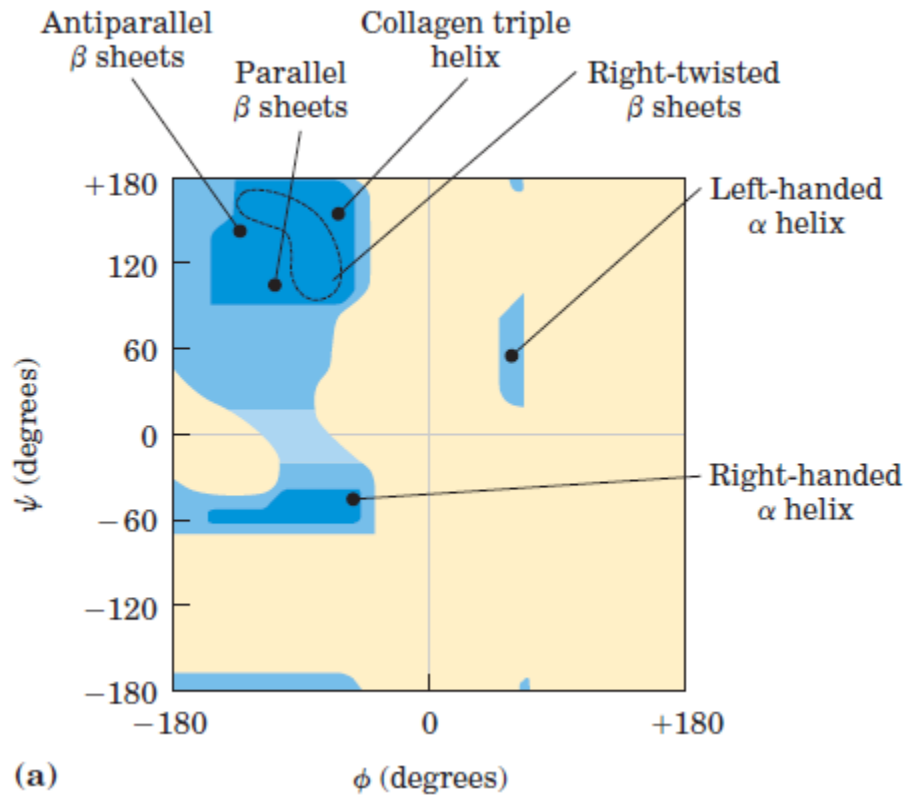
(a) Antiparallel



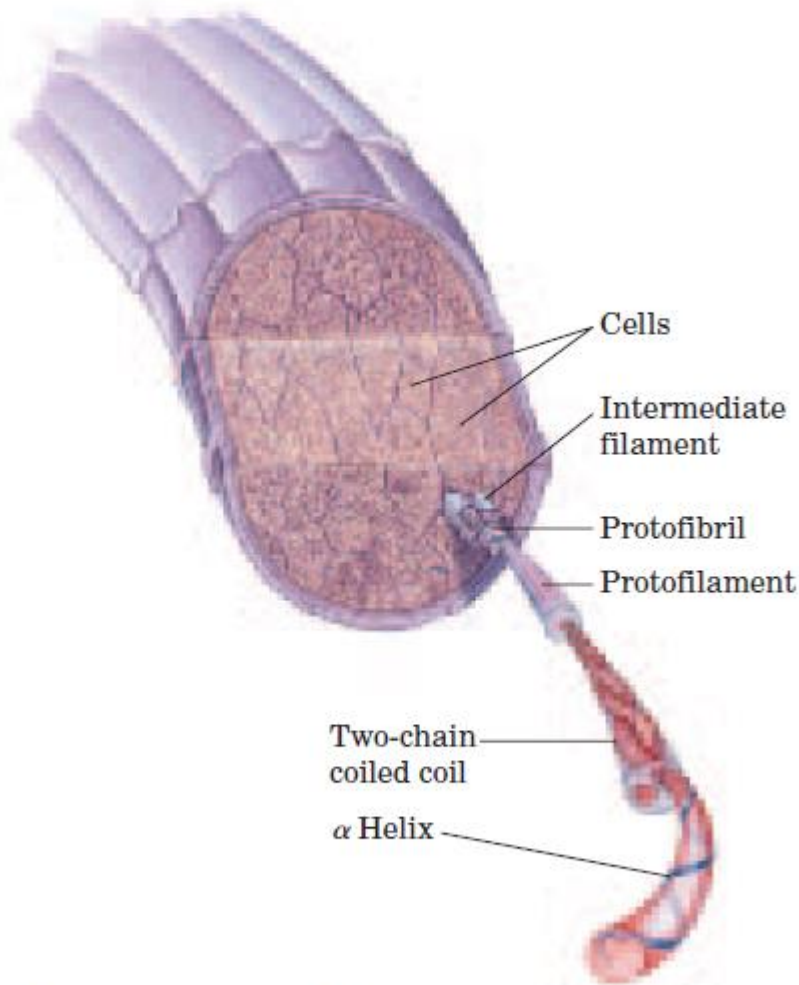
(b) Parallel



Ramachandran plot

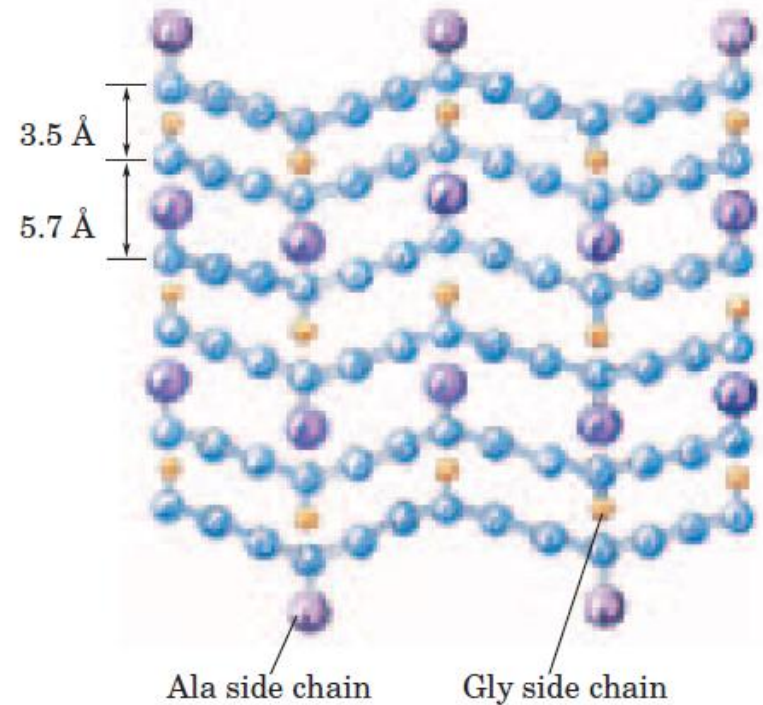


Examples



(b) Cross section of a hair

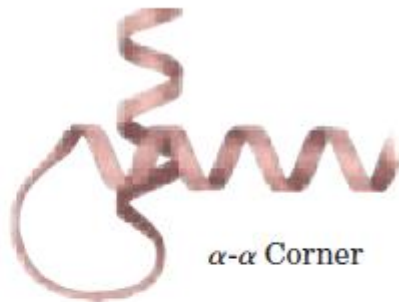
Structure of silk



Common arrangements



(a) β - α - β Loop



α - α Corner



(c) Right-handed connection
between β strands



Left-handed connection
between β strands
(very rare)



(b) Typical connections
in an all- β motif



Crossover connection
(not observed)



(d) β Barrel



Twisted β sheet

Enzymes

TABLE 6-3 International Classification of Enzymes

<i>No.</i>	<i>Class</i>	<i>Type of reaction catalyzed</i>
1	Oxidoreductases	Transfer of electrons (hydride ions or H atoms)
2	Transferases	Group transfer reactions
3	Hydrolases	Hydrolysis reactions (transfer of functional groups to water)
4	Lyases	Addition of groups to double bonds, or formation of double bonds by removal of groups
5	Isomerases	Transfer of groups within molecules to yield isomeric forms
6	Ligases	Formation of C—C, C—S, C—O, and C—N bonds by condensation reactions coupled to ATP cleavage

Note: Most enzymes catalyze the transfer of electrons, atoms, or functional groups. They are therefore classified, given code numbers, and assigned names according to the type of transfer reaction, the group donor, and the group acceptor.



ATP:glucose phosphotransferase

(website)

[E.C. number] 2.7.1.1

TABLE 6–1 Some Inorganic Elements That Serve as Cofactors for Enzymes

Cu^{2+}	Cytochrome oxidase
Fe^{2+} or Fe^{3+}	Cytochrome oxidase, catalase, peroxidase
K^{+}	Pyruvate kinase
Mg^{2+}	Hexokinase, glucose 6-phosphatase, pyruvate kinase
Mn^{2+}	Arginase, ribonucleotide reductase
Mo	Dinitrogenase
Ni^{2+}	Urease
Se	Glutathione peroxidase
Zn^{2+}	Carbonic anhydrase, alcohol dehydrogenase, carboxypeptidases A and B

Cofactor & coenzymes

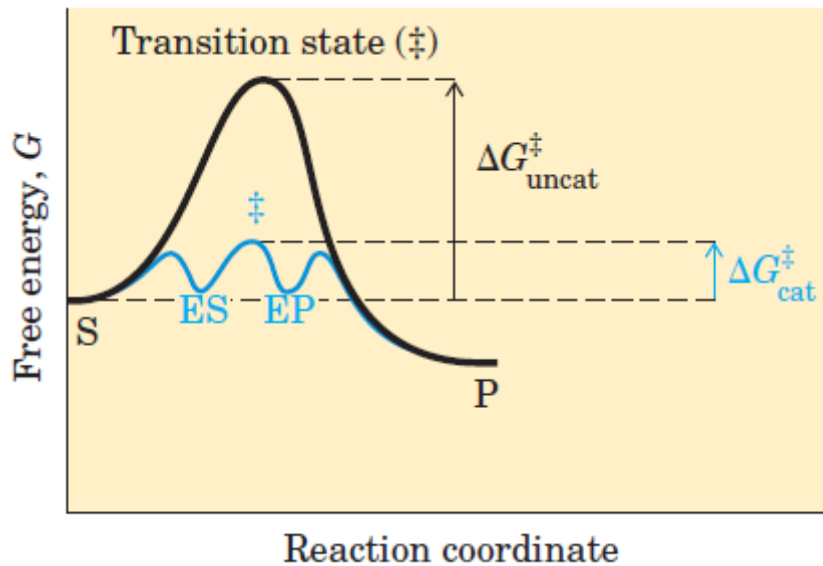
TABLE 6–2 Some Coenzymes That Serve as Transient Carriers of Specific Atoms or Functional Groups

<i>Coenzyme</i>	<i>Examples of chemical groups transferred</i>	<i>Dietary precursor in mammals</i>
Biotin	CO_2	Biotin
Coenzyme A	Acyl groups	Pantothenic acid and other compounds
5'-Deoxyadenosylcobalamin (coenzyme B_{12})	H atoms and alkyl groups	Vitamin B_{12}
Flavin adenine dinucleotide	Electrons	Riboflavin (vitamin B_2)
Lipoate	Electrons and acyl groups	Not required in diet
Nicotinamide adenine dinucleotide	Hydride ion ($:\text{H}^-$)	Nicotinic acid (niacin)
Pyridoxal phosphate	Amino groups	Pyridoxine (vitamin B_6)
Tetrahydrofolate	One-carbon groups	Folate
Thiamine pyrophosphate	Aldehydes	Thiamine (vitamin B_1)

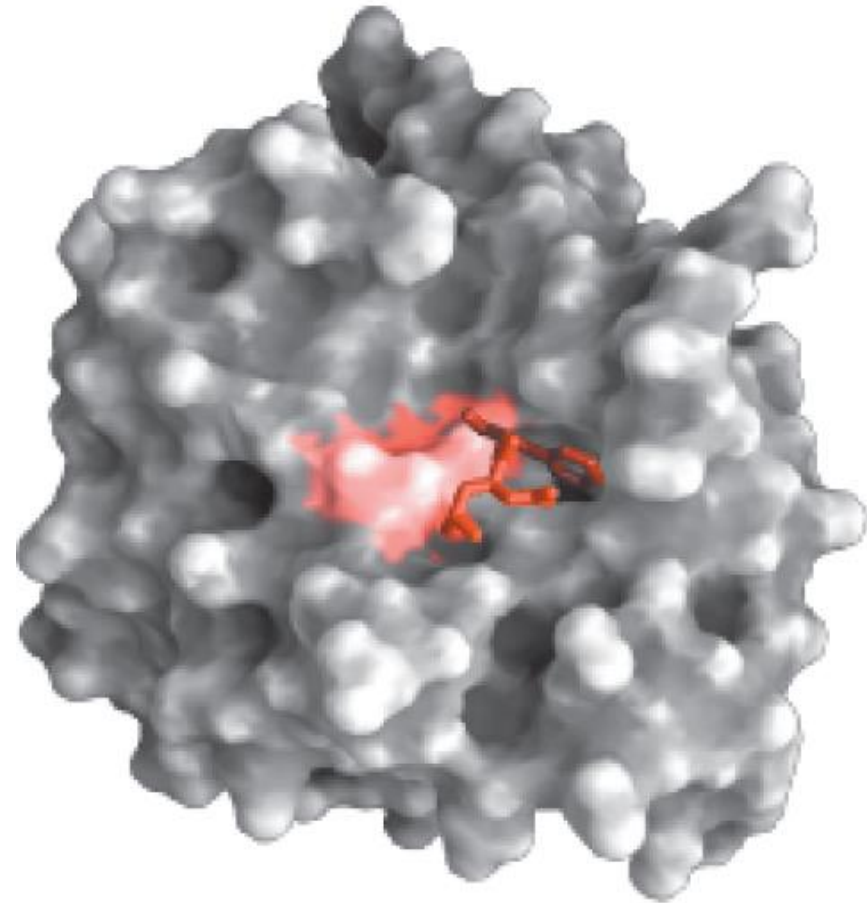
Catalysis and active site



E, S, and P represent the enzyme, substrate, and product; ES and EP are transient complexes of the enzyme with the substrate and with the product.



Reaction coordinate diagram for a chemical reaction.



(PDB)

Enzymatic catalysis



$$V = k[S]$$

$$K'_{\text{eq}} = \frac{[P]}{[S]}$$

$$\Delta G'^{\circ} = -RT \ln K'_{\text{eq}}$$

$$R = 8.315 \text{ J/mol} \cdot \text{K}$$

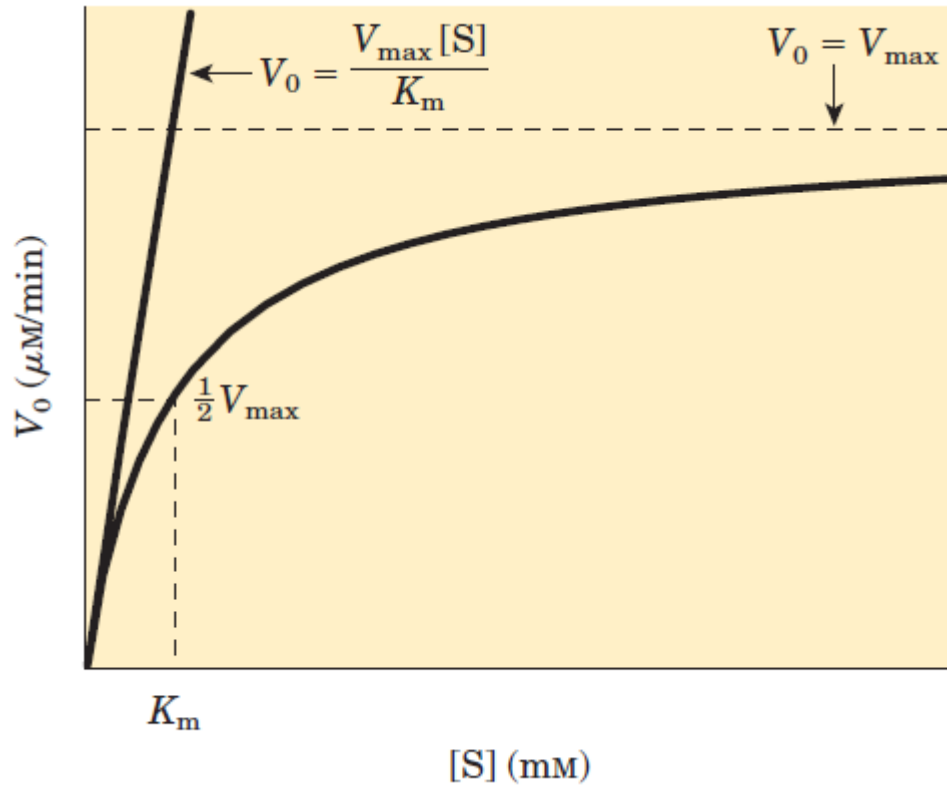
$$T = 298 \text{ K (25 }^{\circ}\text{C)}$$

TABLE 6-4 Relationship between K'_{eq} and $\Delta G'^{\circ}$

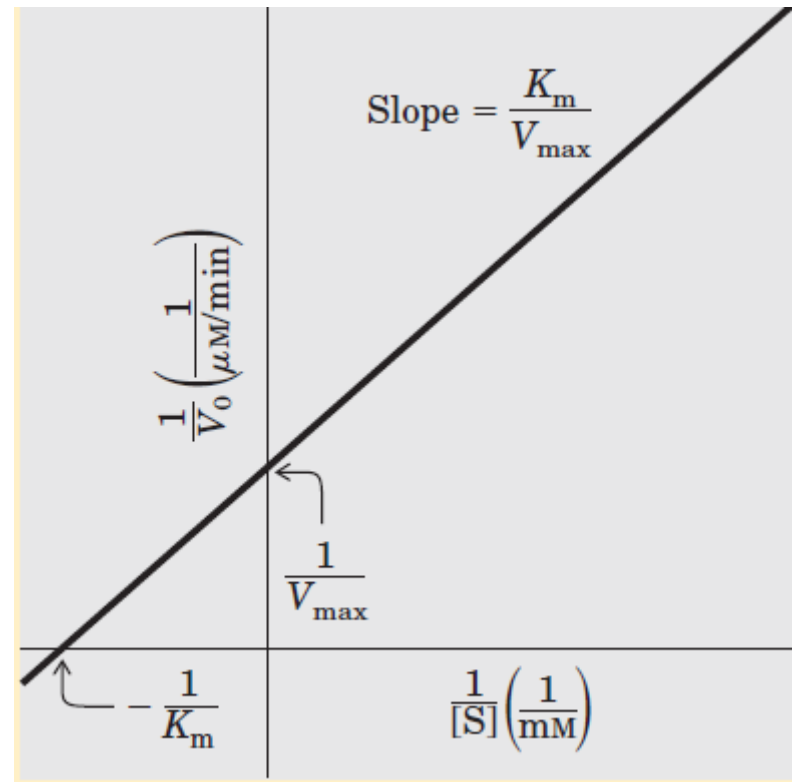
K'_{eq}	$\Delta G'^{\circ}$ (kJ/mol)
10^{-6}	34.2
10^{-5}	28.5
10^{-4}	22.8
10^{-3}	17.1
10^{-2}	11.4
10^{-1}	5.7
1	0.0
10^1	-5.7
10^2	-11.4
10^3	-17.1

Note: The relationship is calculated from $\Delta G'^{\circ} = -RT \ln K'_{\text{eq}}$ (Eqn 6-3).

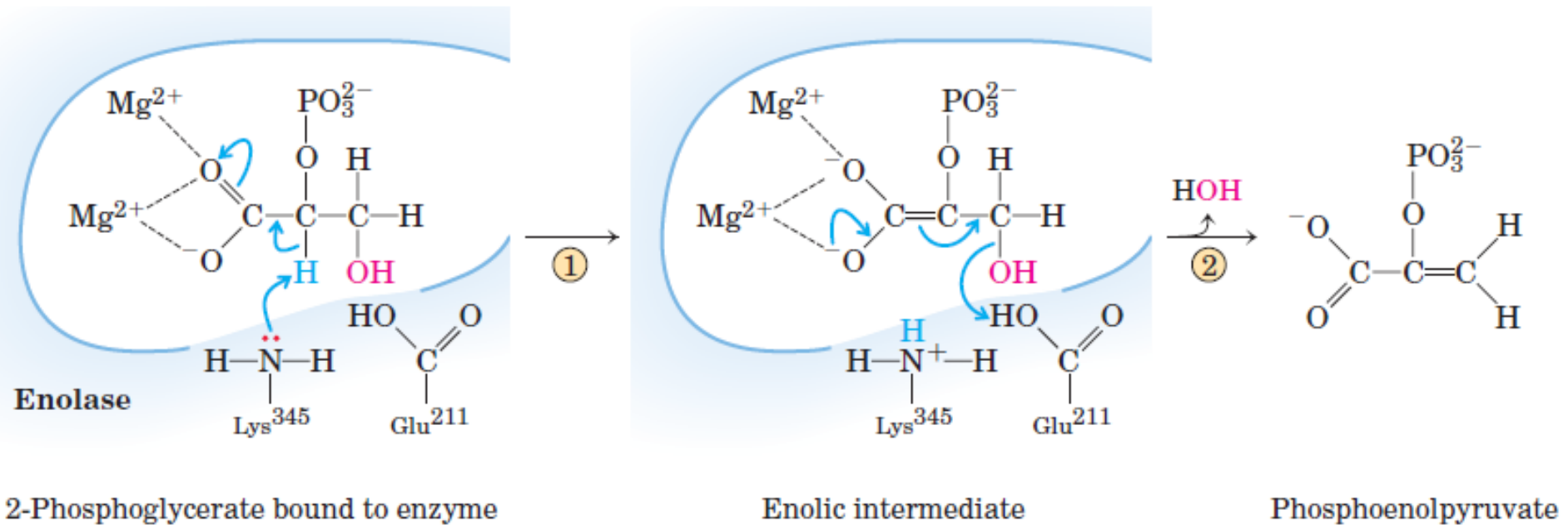
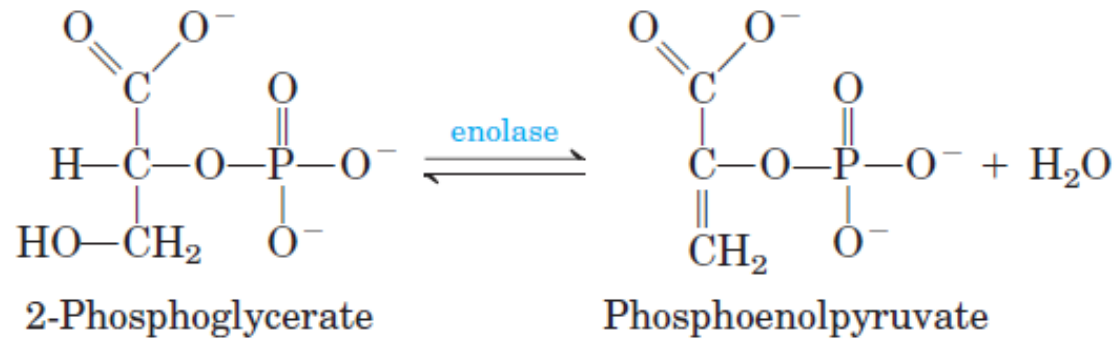
Enzymatic catalysis



A double-reciprocal or Lineweaver-Burk plot.



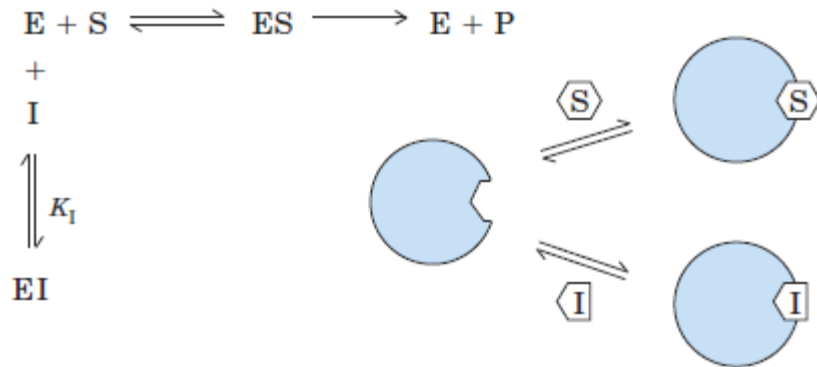
Example of the reaction mechanism



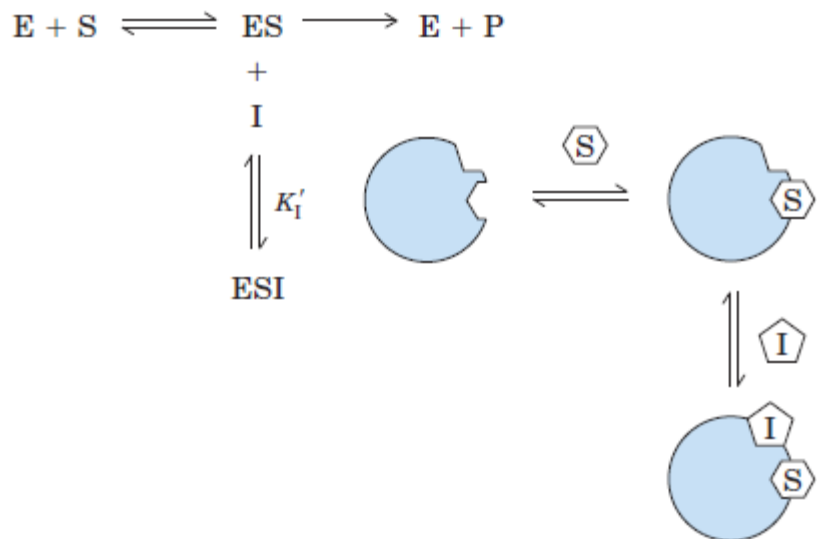
Enzymatic inhibitors

Three types of reversible inhibition.

(a) Competitive inhibition



(b) Uncompetitive inhibition



(c) Mixed inhibition

