Proteins & Peptides

Peptide Backbone

R H H OH N H

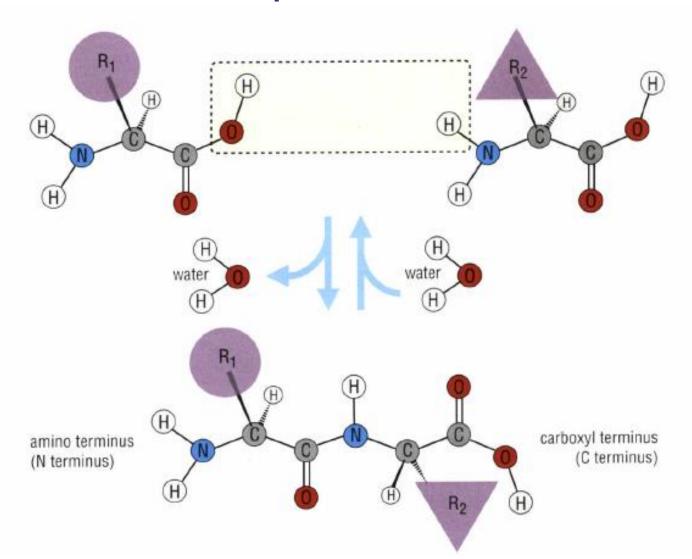
Amino acid sequence

- Hydrophobic side chain: Van der Waals interactions
- Hydrophilic side chain: hydrogen bonding
- Amphoteric side chain
- Helix favoring residues: Ala & Leu

Amide Bond in Peptide Synthesis

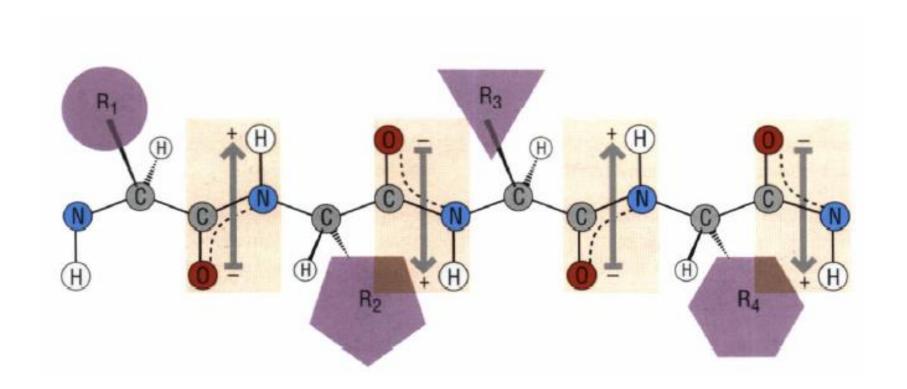
$$H_3C$$
 H_3C
 H_3C

Peptide Bond



Schematic of an Extended Peptide Chain

Consider amide bond tautomery in peptide sequence.



Conformations for Peptide: cis & trans

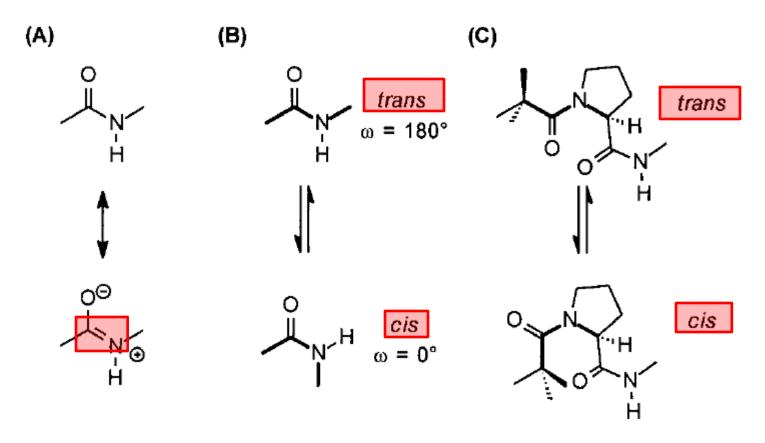


Fig. 2.2 (A) Resonance stabilization and (B) cis/trans isomerization of the peptide bond (C) cis/trans Isomers of a Xaa-Pro bond.

Three Types of Bond Angles in Peptide Sequence

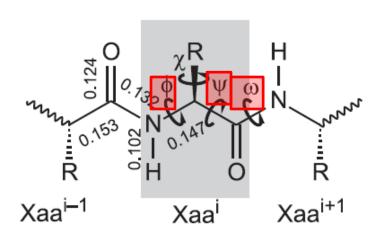


Fig. 2.1 Torsion angles φ , ψ , ω , and χ^1 and bond lengths of the amino acid Xaa in a peptide.

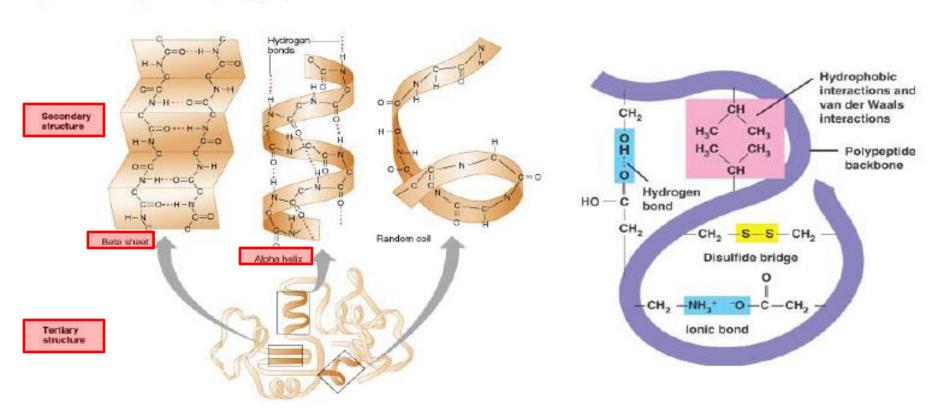
Structures Introduced for Proteins

Primary: sequence of amino acids

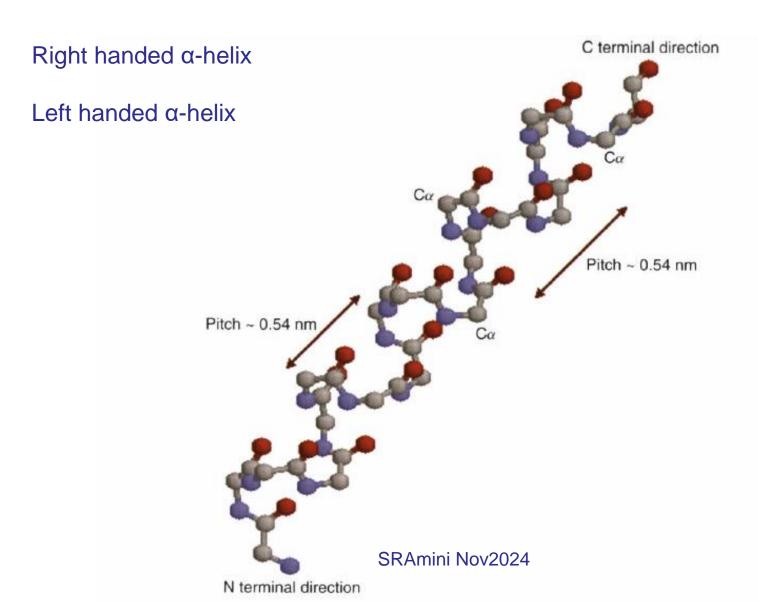
Secondary: backbone; Hydrogen bonds

Tertiary: side chain interactions

Quaternary: 2+ polypeptides



α-Helix as Secondary Structure of Protein



β-sheet as Secondary Structure of Protein

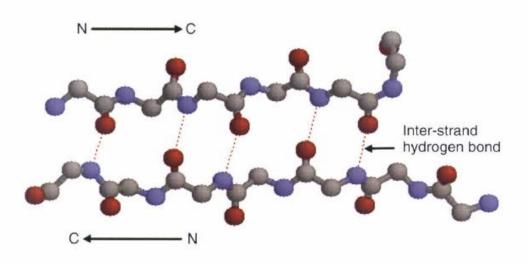


Figure 3.11 Two adjacent β strands are hydrogen bonded to form a small element of β sheet. The hydrogen bonds are inter-strand between neighbouring CO and NH groups. Only the heavy atoms are shown in this diagram for clarity

List of Essential Amino Acids

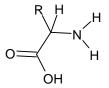
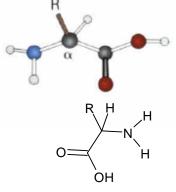


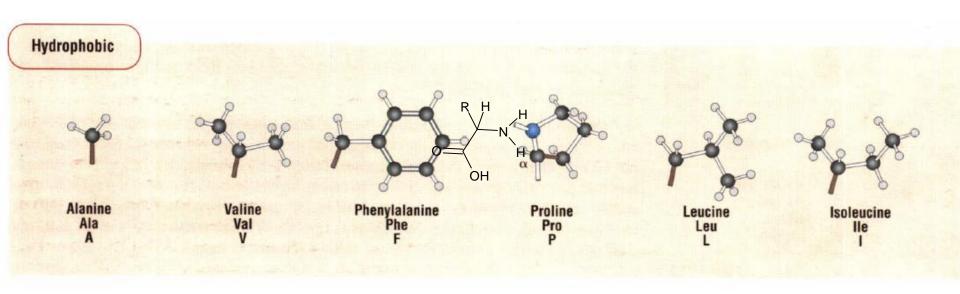
Table 2.1 The pK values for the α -carboxyl, α -amino groups and side chains found in the individual amino acids

Amino acid	pK_1	pK_2	pK_R	Amino acid	pK_1	pK_2	pK_R
Alanine	2.4	9.9	-	Leucine	2.3	9.7	
Arginine	1.8	9.0	12.5	Lysine	2.2	9.1	10.5
Asparagine	2.1	8.7	-	Methionine	2.1	9.3	-
Aspartic Acid	2.0	9.9	3.9	Phenylalanine	2.2	9.3	-/-
Cysteine	1.9	10.7	8.4	Proline	2.0	10.6	_
Glutamic Acid	2.1	9.5	4.1	Serine	2.2	9.2	_
Glutamine	2.2	9.1	-	Threonine	2.1	9.1	-
Glycine	2.4	9.8		Tyrosine	2.2	9.2	10.5
Histidine	1.8	9.3	6.0	Tryptophan	2.5	9.4	-
Isoleucine	2.3	9.8	-	Valine	2.3	9.7	

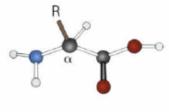
Adapted from Dawson, R.M.C, Elliot, W.H., & Jones, K.M. 1986 Data for Biochemical Research, 3rd edn. Clarendon Press Oxford.

Hydrophobic α-Amino Acids

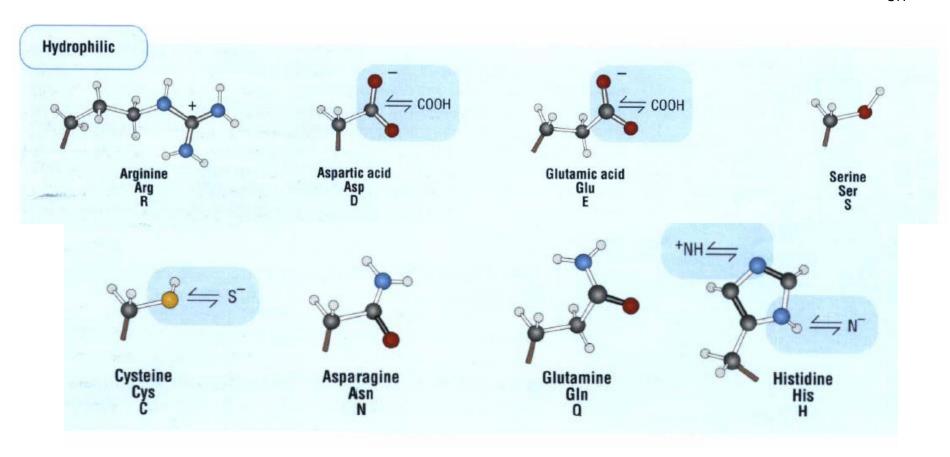




Hydrophilic α-Amino Acids

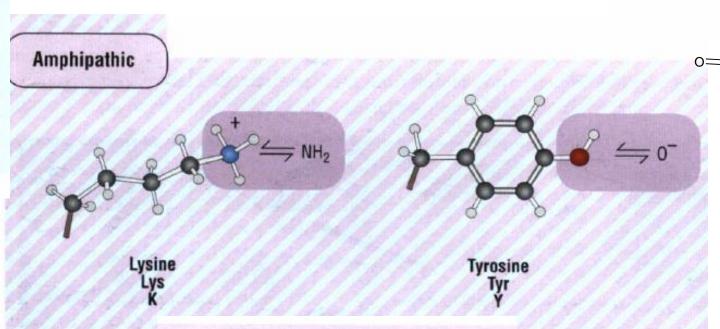


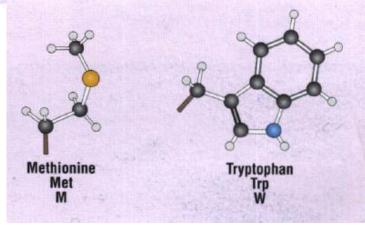
$$O \longrightarrow H$$



Some Other α-Amino Acids







ÔН

Amino acid	Property of individual amino acid residues	Ball and stick representation of each amino acid
Alanine	Non-polar side chain.	Assemble websites and the
A	Small side chain volume.	
Ala	Van der Waals volume = 67 Å ^{3*}	to the same of the
M _r 71.09	Frequency in proteins = 7.7 %	6000
	Surface area = 115 Å^2	
	Unreactive side chain	6-0
Arginine	Positively charged side chain at pH 7.0. pK	and the state of t
A BA MARIANA		
SERVICE OF THE PROPERTY.	for guanidino group in proteins ~12.0	
R	for guanidino group in proteins ~ 12.0 Van der Waals volume = 167 Å ³	
R Arg		La Contraction of the Contractio
R Arg M _r 156.19	Van der Waals volume = 167 Å ³	Service of the servic

Glycine

G

Gly

M_r 57.05

Uncharged, small side chain.

Often found in turn regions of proteins or regions of conformational flexibility

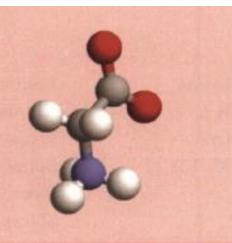
No chiral centre; due to two hydrogens

attached to Ca centre

Van der Waals volume = 48 Å^3

Frequency in proteins = 7.4%

Surface area = 75 Å^2



Asparagine Polar, but uncharged, side chain

N Van der Waals volume = 148 Å^3

As $\frac{1}{2}$ Frequency in proteins = 4.3 %

 M_r 114.11 Surface area = 160 Å²

Polar side chain will hydrogen bond

Relatively small side chain volume leads to

this residue being found relatively

frequently in turns

Aspartate Negatively charged side chain

D pK for side chain of ~ 4.0

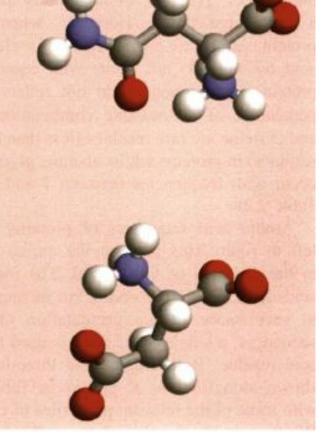
Asp Van der Waals volume = 67 Å^3

 M_r 115.09 Frequency in proteins = 5.2 %

Surface area = 150 Å^2

Charged side chain exhibits electrostatic

interactions with positively charged groups.



Amino acid	Property of individual amino acid residues	Ball and stick representation of each amino acid
Cysteine	Side chain contains thiol (SH) group.	
C	Van der Waals volume = 86 Å ³	
Cys	Frequency in proteins = 2.0 %	
M _r 103.15	Surface area = 135 Å^2	
	Thiol side chain has pK in isolated amino acid of \sim 8.5 but in proteins varies 5–10	
	Thiol group is very reactive	
Glutamine	Polar but uncharged side chain	
Q	Van der Waals volume = 114 Å ³	2000
Gln	Frequency in proteins = 4.1 %	
M _r 128.12	Surface area = 180 Å^2	-
	Polar side chain can hydrogen bond	0 0 9
Glutamate	Negatively charged side chain.	
E	Van der Waals volume = 109 Å ³	
Glu	Frequency in proteins = 6.2 %	
Mr 129.12	Surface area = 190 Å^2	× 0-
	Side chain has pK of \sim 4.5.	

Amino acid	Property of individual amino acid residues	Ball and stick representation of each amino acid
Histidine	Imidazole side chain	
H	Van der Waals volume = 118 Å^3	
His	Frequency in proteins = 2.3 %	
M _r 137.14	Surface area = 195 Å^2	
	The side chain exhibits a p $K \sim 6.0$ in model peptides but in proteins can vary from $4-10$	- Contraction
Isoleucine	Hydrophobic side chain exhibiting non-polar	
I	based interactions but generally unreactive	
Ile	Van der Waals volume = 124 Å ³	
M _r 113.16	Frequency in proteins = 5.3% Surface area = 175 Å^2	C.E.E.
Leucine	Hydrophobic side chain	
L	Van der Waals volume = 124 Å^3	
Leu	Frequency in proteins = 8.5 %	000
M _r 113.16	Surface area = 170 Å ²	

Lysine

K

Lys

Mr 128.17

Positively charged side chain

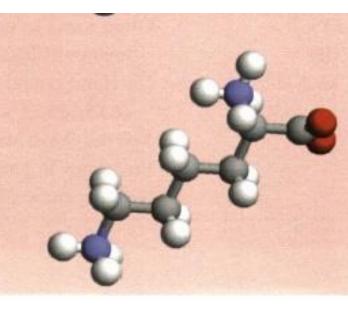
Van der Waals volume = 135 Å^3

Frequency in proteins = 5.9 %

Surface area = 200 Å^2

Side chain is basic with pK of ~ 10.5 .

Shows ionic interactions



Amino acid	Property of individual amino acid residues	Ball and stick representation of each amino acid
Methionine M Met M _r 131.19	Sulfur containing hydrophobic side chain The sulfur is unreactive especially when compared with thiol group of cysteine Van der Waals volume = 124 Å ³ Frequency in proteins = 2.4 % Surface area = 185 Å ²	
Phenylalanine F Phe M _r 147.18	Hydrophobic, aromatic side chain Phenyl ring is chemically unreactive in proteins. Exhibits weak optical absorbance around 280 nm Van der Waals volume = 135 Å ³ Frequency in proteins = 4.0 % Surface area = 210 Å ²	A de la

Proline Cyclic ring forming hydrophobic side chain The cyclic ring limits conformational Pro flexibility around N-Ca bond Mr 97.12 In a polypeptide chain lacks amide hydrogen and cannot form backbone hydrogen bonds Van der Waals volume = 90 Å3 Frequency in proteins = 5.1%Surface area = 145 Å^2 Serine Polar but uncharged side chain. Contains hydroxyl group (-OH) that hydrogen bonds Ser Oxygen atom can act as potent nucleophile in Mr 87.08 some enzymes Van der Waals volume = 73 Å^3 Frequency in proteins = 6.9%Surface area = 115 Å^2

Amino acid	Property of individual amino acid residues	Ball and stick representation of each amino acid
Threonine T Thr M _r 101.11	Polar but uncharged side chain. Contains hydroxyl group (-OH) Hydrogen bonding side chain Van der Waals volume = 93 Å ³ Frequency in proteins = 5.9 % Surface area = 140 Å ²	
Tryptophan W Trp M _r 186.21	Large, hydrophobic and aromatic side chain Almost all reactivity is based around the indole ring nitrogen Responsible for majority of near uv absorbance in proteins at 280 nm Van der Waals volume = 163 Å ³ Frequency in proteins = 1.4 % Surface area = 255 Å ²	

Tyrosine Aromatic side chain

Y Van der Waals volume = 141 Å^3

Tyr Frequency in proteins = 3.2%

 M_r 163.18 Surface area = 230 Å²

Phenolic hydroxyl group ionizes at pH values

around pH 10

Aromatic ring more easily substituted than

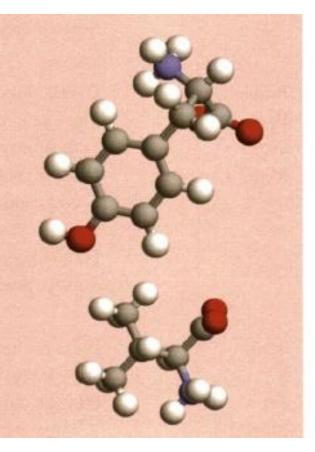
that of phenylalanine

Valine Hydrophobic side chain

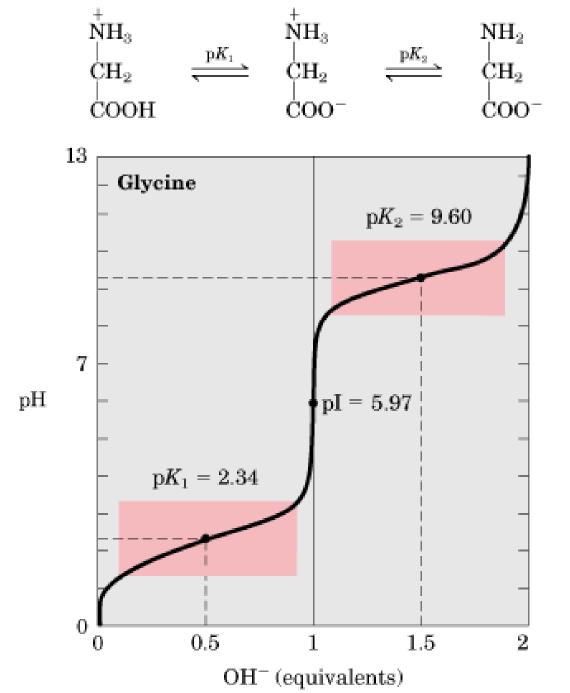
V Van der Waals volume = 105 Å^3

Val Frequency in proteins = 6.6 %

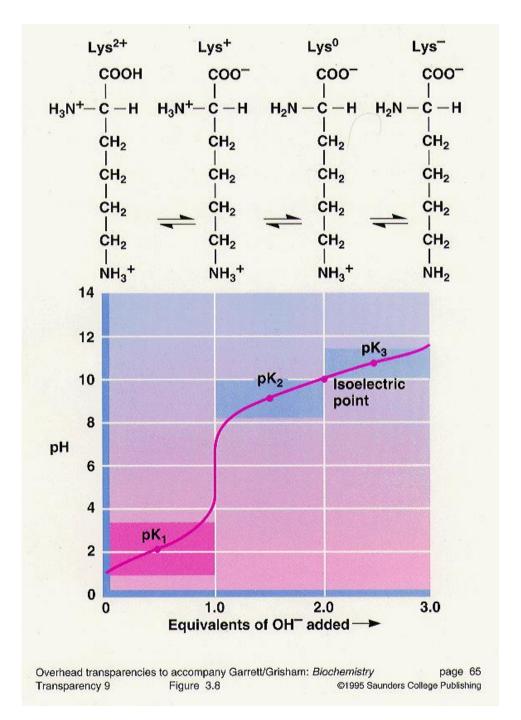
 M_r 99.14 Surface area = 155 Å²

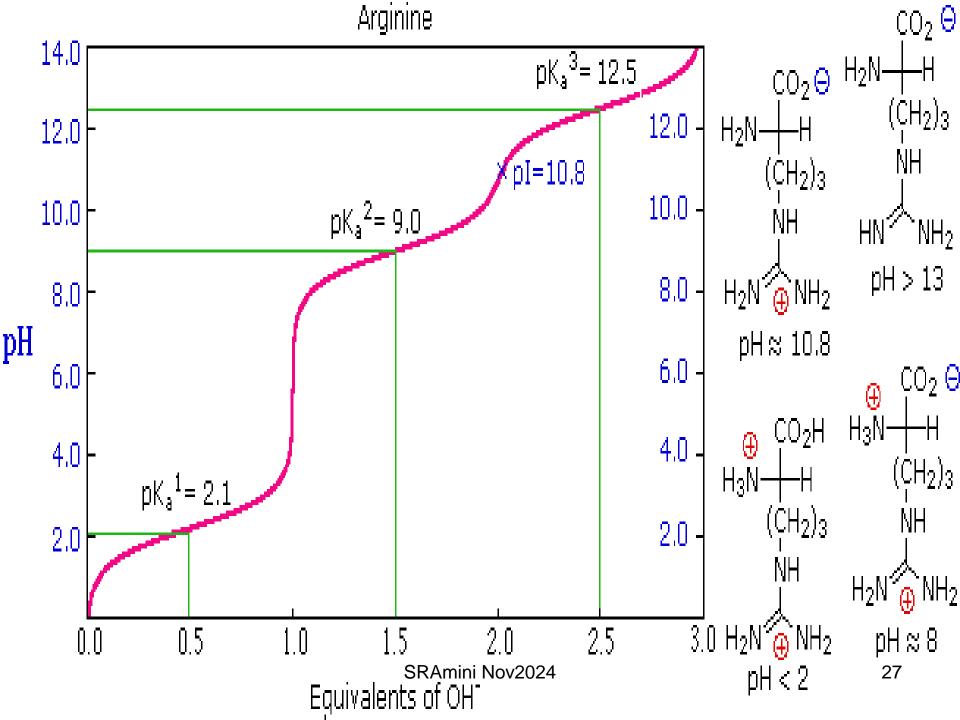


Quantitative
Study via
Titration of
Gly as an
α-Amino Acids
with NaOH
as Titrant



Quantitative Study via Titration of Lys as an α-Amino Acids with NaOH as Titrant





Protein Analysis

- Purification:
- ✓ Electrophoresis: regarding pK_a & pI (isoelectric pH)
- ✓ chromatography

- Structure Elucidation:
- √ X- ray
- ✓ Circular Dichroism (CD)
- **✓** NMR

Protein Types

Simple proteins

- Conjugated proteins:
- ✓ nucleoproteins
- √ glycoproteins
- ✓ phosphoproteins
- ✓ lipoproteins
- ✓ metalloproteins
- √ chromoproteins