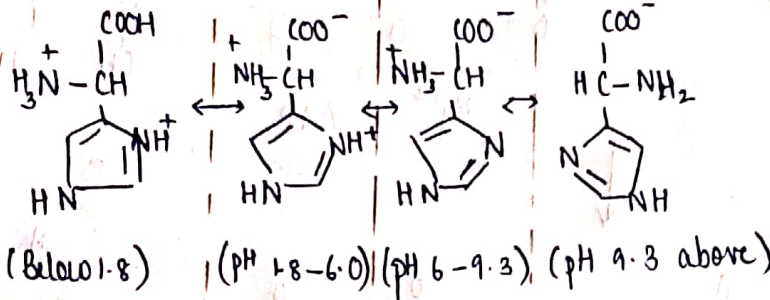
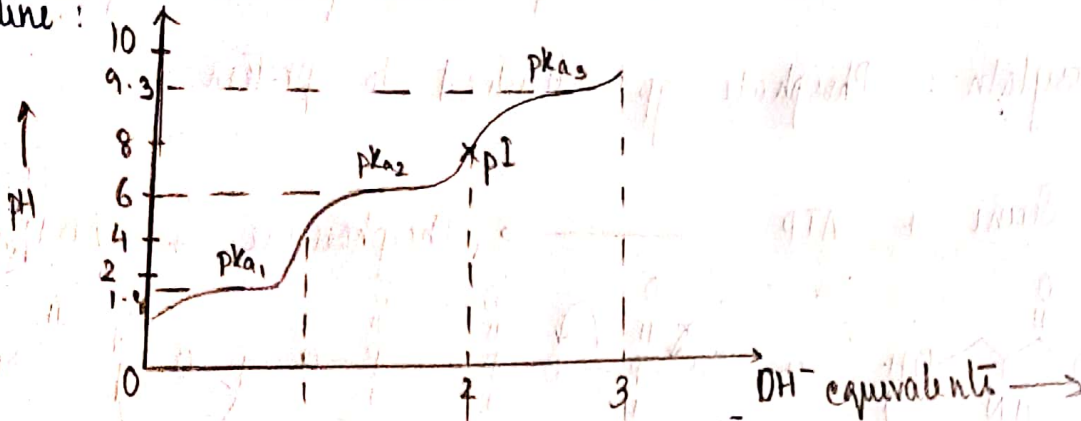


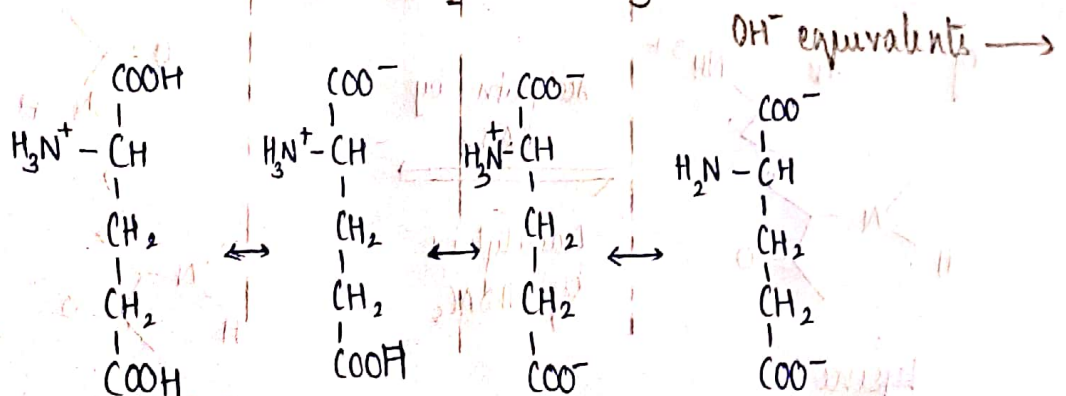
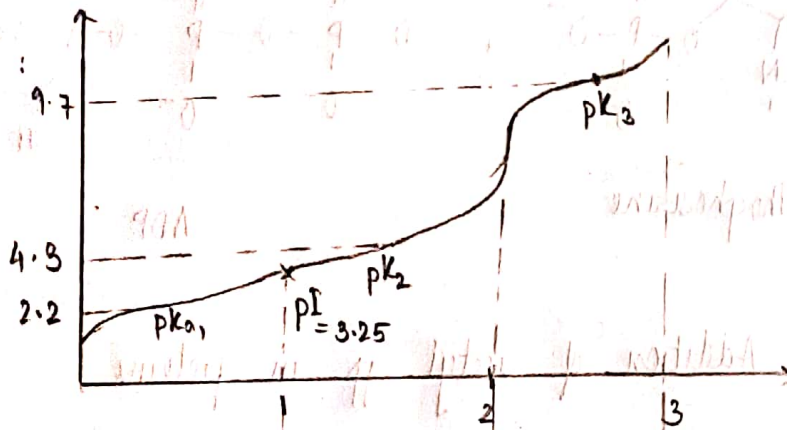
Biomolecular Structures Assignment 1

SHREEVA
2018113011

1) Histidine:

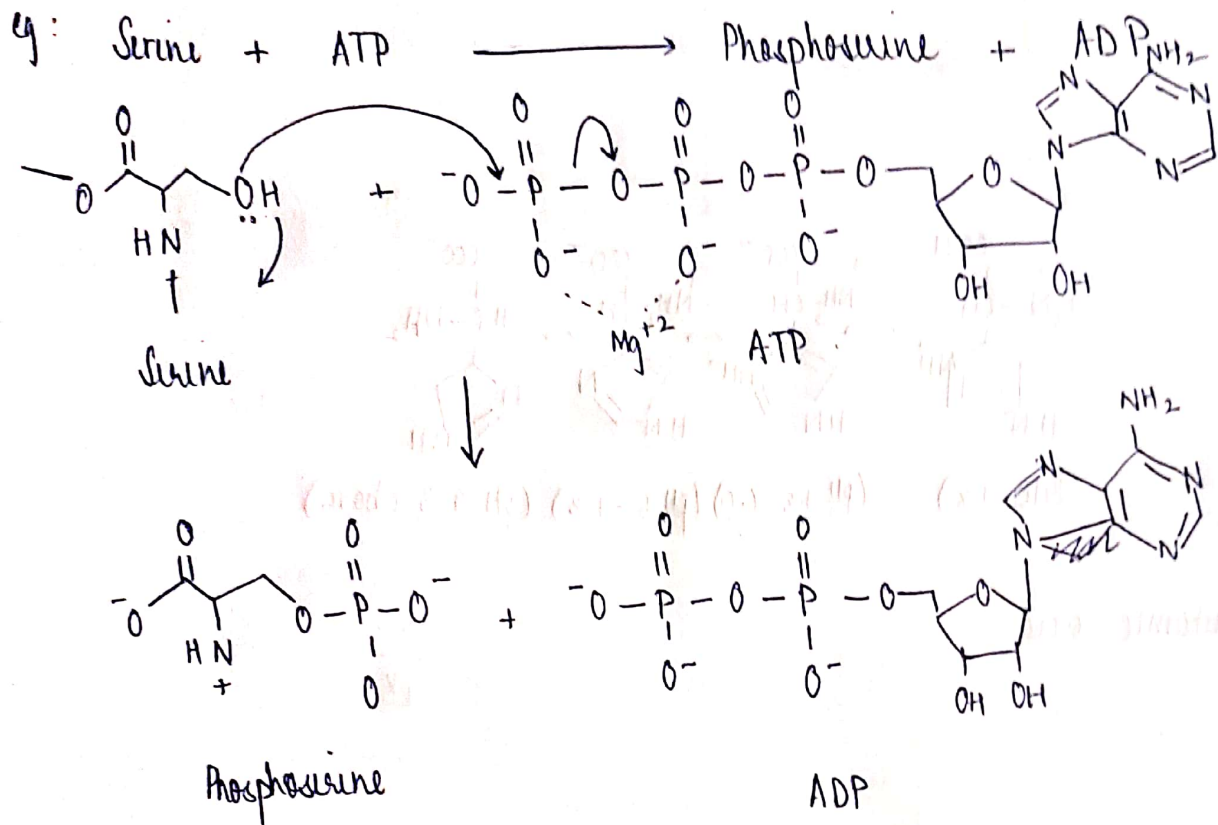


Glutamic acid:

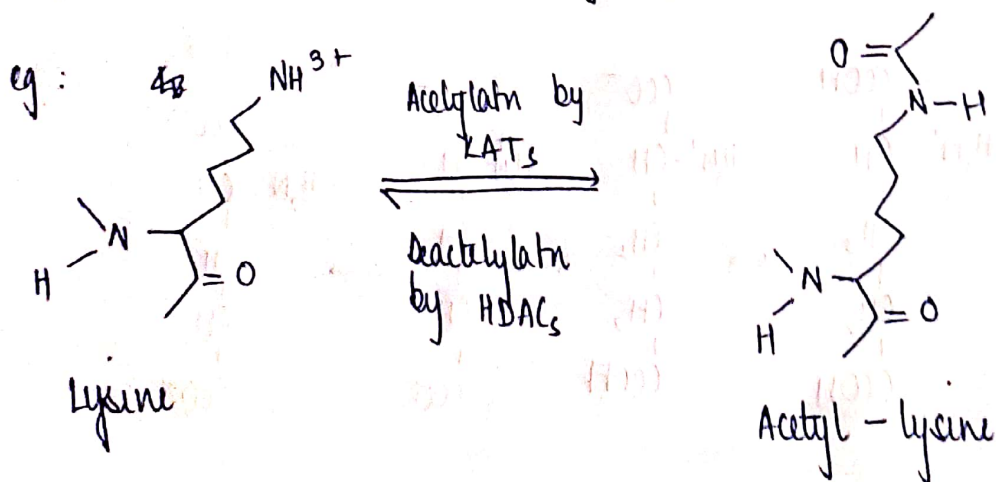


2) The chemical changes in a protein that occur after a protein has been produced are called post-translational modifications (PTMs). It is the covalent addition of certain functional groups to proteins.

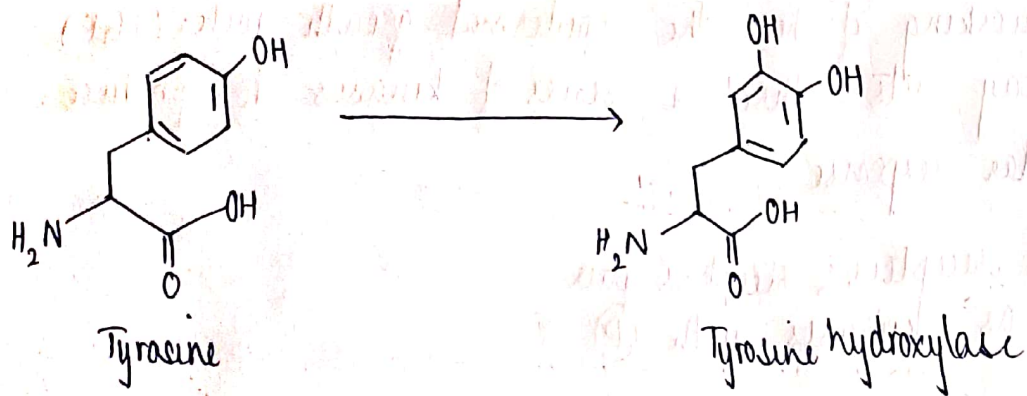
① Phosphorylation: Phosphate group attached to protein.



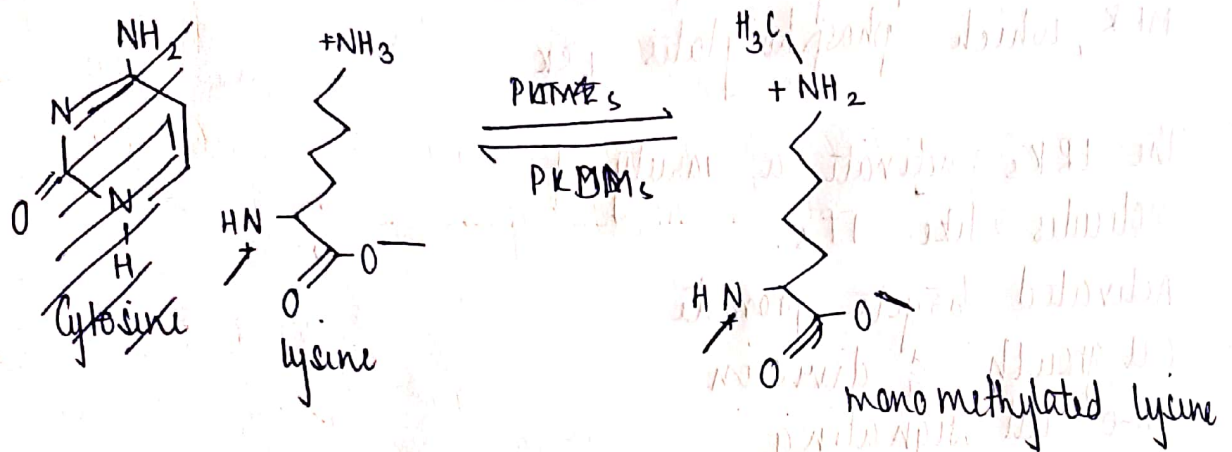
② Acetylation: Addition of acetyl group in proteins.



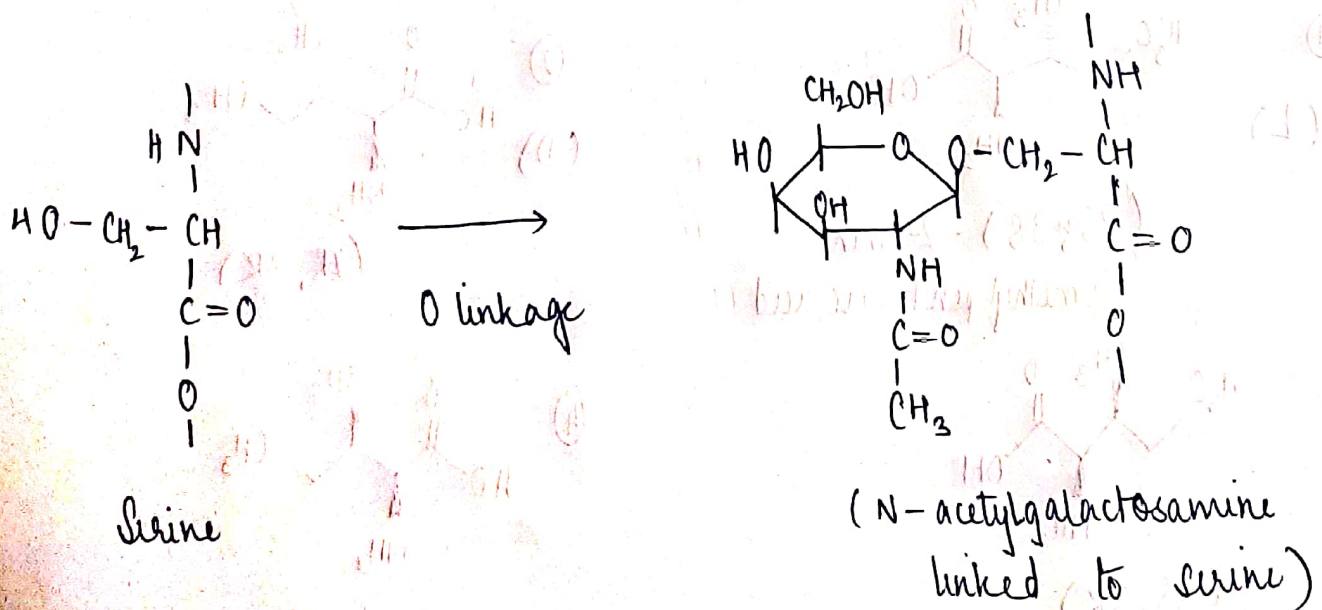
③ Hydroxylation: Addition of hydroxyl group:



④ Methylation: Addition of methyl gp



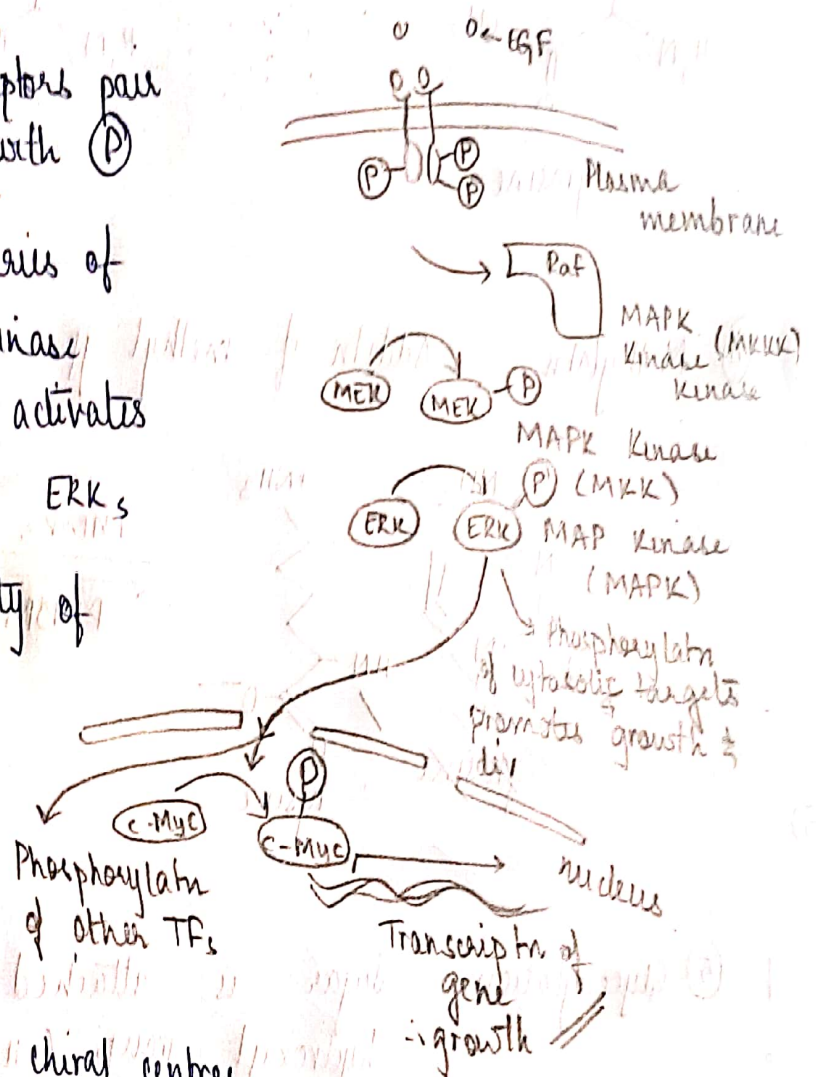
⑤ Glycosylation: Sugar is attached thro' a oxygen on a hydroxyl group (OH) to serine (O-linkage)



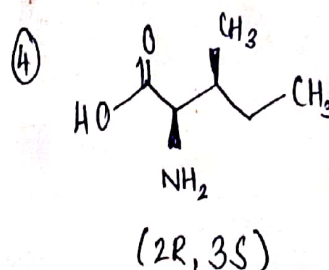
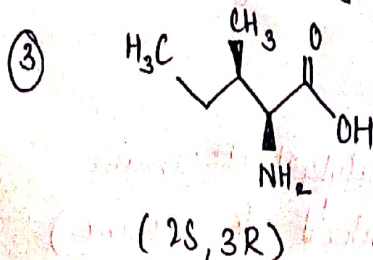
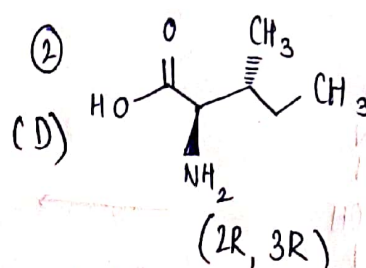
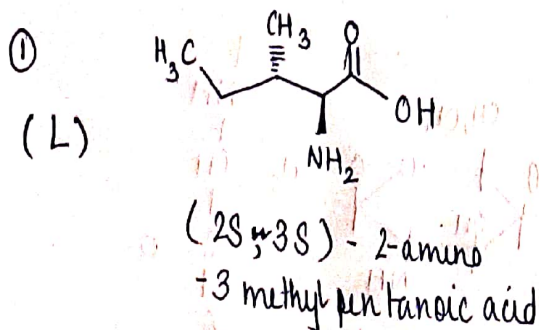
3) MAPK signaling cascade (Phosphorylation):

The working of how the epidermal growth factor (EGF) pathway acts thro' a series of kinases to produce cellular response.

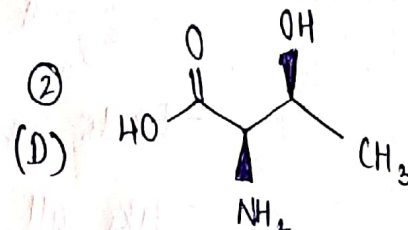
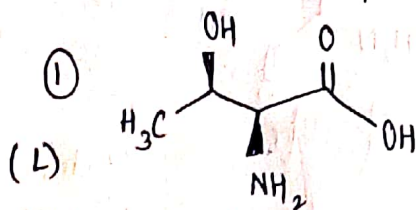
- ① GF bind to receptors, receptors pair up & act as kinases with (P)
 - ② Activated receptors trigger series of events & activate Raf kinase
 - ③ Active Raf phosphorylates & activates MEK, which phosphorylates ERKs
 - ④ The ERKs activate a variety of molecules like TFs.
- ∴ Activated targets promote cell growth & division thro' cell signaling using phosphorylation.



4) Isomers: C_2 , C_3 are chiral centres.

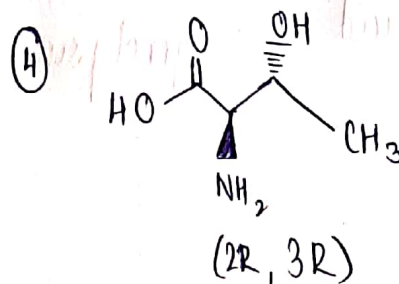
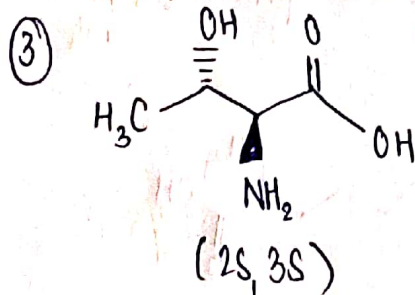


② Threonine : C_2, C_3 are chiral centres.



(1S, 3R) - 2-amino-3-hydroxybutanoic acid

(2R, 3S)



⑤ Essential amino acids : Cannot be made by the human body & hence must be retrieved from outside (diet)

Non essential " " : Can be made within the body.

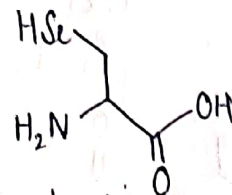
Non essential (11)

Essential (9)

- ① ALA
- ② ARG
- ③ ASN
- ④ ASP
- ⑤ CYS
- ⑥ GLY
- ⑦ PRO
- ⑧ SER
- ⑨ TYR
- ⑩ GLN
- ⑪ GLU

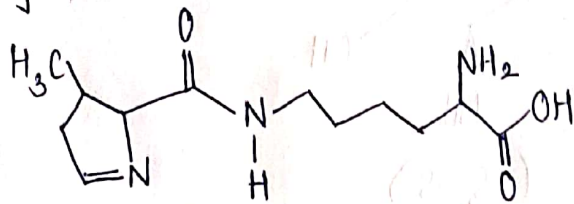
- ① HIS
- ② ILE
- ③ LEU
- ④ LYS
- ⑤ MET
- ⑥ PHE
- ⑦ THR
- ⑧ TRP
- ⑨ VAL

6) 21st Amino acid : - Selenocysteine (Sec)



- Exists in all 3 domains of life (codon: UGA)
- Like all amino acids has its own tRNA.
- Selenium analogue of Cysteine (Se replaces S).
- Imp for fundamental cellular processes. (metabolic rate reg.)

22nd amino acid : - Pyrrolysine (Pyl)



- Like all amino acids has its own tRNA & aminoacyl-tRNA synthase (aARS) (codon: UAG)
- Very rare, found in 7 organisms

1) This is because carbon atoms make up the backbone of many important molecules in ~~our~~ our body like DNA, RNA sugars, proteins & fats. i.e organic molecules. Since carbon is the only element that can form 4 chem. bonds to other atoms \therefore it's the right size. ~~to accommodate~~ (4 valency) [unlike Si atoms \therefore large]. Along with this it has the ability to form long C-C chains, straight & branched (complex). C atom can form $=$ & \equiv bonds also. Chains & rings can also be used to form bonds - millions of org. comp. exist with C as main atom.

1) Isoionic pt : pt is the pH at which zwitterion molecule has equal no. of +ve & -ve charges & no adjacent ionic species

→ Isoionic pt of 0.1 M ALA :

$$[H] = \sqrt{\frac{k_1 k_2 C + k_1 k_w}{k_1 + C}}$$

$$k_1 = 10^{-2.34}$$

$$k_2 = 10^{-9.69}$$

$$k_w = 10^{-14}$$

$$C = 0.1$$

$$= \sqrt{\frac{10^{-2.34} + 10^{-9.69} - 1}{10^{-2.34} + 10^{-14}}}$$

$$= 9.45 \times 10^{-7}$$

$$\therefore pH = -\log [H^+]$$

$$= 6.02$$

$$\therefore \text{Isoionic pt} = \underline{\underline{6.02}}$$

9) a) 2.27 ribbon : - Similar to α also called α_{II} ribbon (α_{II} struct.)

- N-H...O \angle differ from st. \angle by $60^\circ - 66^\circ$

- \angle b/w N-H & N...O are $40^\circ - 48^\circ$

- unacceptable structure all over.

- N...O distance is $> 0.1 \text{ \AA}$ more shorter than reliable value $\rightarrow (-2.52 \text{ \AA})$

b) ① Left handed polyproline II: (PP II)

- $\phi = -78^\circ$, $\psi = +146^\circ$
- Most are shorter than 5 residues
- Have perfect 3-fold & rotational sym.
- correltn bto hydrophobicity of residues i & $i+3$

② Left handed polyglycine II: (PG II)

- ~~Basis of Gly: Gly differ~~
- Do not form a self contained H bonding network (same for PP II)
- $\phi = -77^\circ$, $\psi = +145^\circ$
- An extended 3, helix is hydrogen bonded to 6th neighbouring chains packed in hexagonal array.
- CO & NH H-bond stabilise (inter molecular)

c) Collagen triple helix:

- Confers strict amino acid ~~structure~~ seq. constraints requiring a $(Gly-X-Y)_n$ repeating pattern
- Requires high content of imino acids.
- Has supercoiled^{triple} helix structure.
- X & Y were found to be usually hydroxyproline (OH gp of this stabilises the collagen helix stabilising proline via stereoelectronic effects & does not participate in H bonding.)

10) α helix \rightarrow $5.4 \text{ \AA} \rightarrow 3.6 \text{ residues (one turn)}$

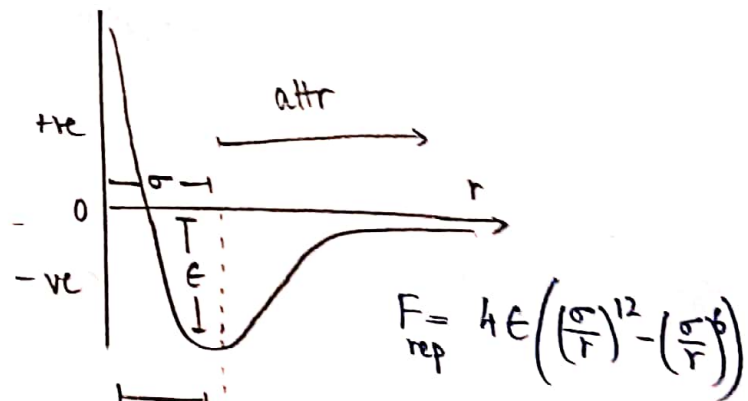
$\alpha \rightarrow 30 \text{ residues}$

$$\alpha = \frac{30 \times 5.4}{3.6}$$

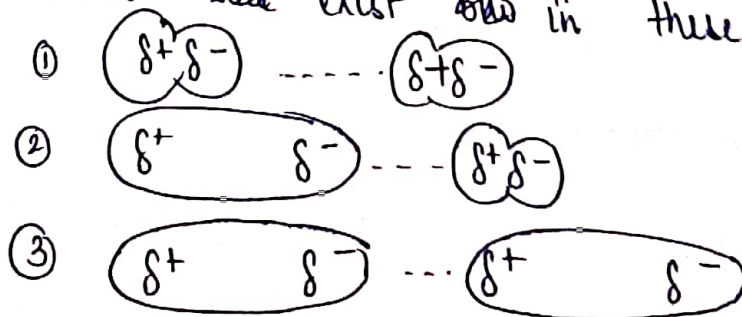
$$= 45 \text{ \AA}$$

- 11) - Vanderwaals forces are the cum total of non covalent forces that depend on distance b/w entities
- They can be repulsive or attr based on dist & the interacting nonbonded atoms/molecules.
- Vanderwaal F are weak electrostatic & are mainly attractive.

- attr when interactn b/w entities are induced dipoles
- repulsion when clouds overlap or too close.



& Since VDW forces exist b/w in these following cases :



& are weak electrostatic F \therefore mostly attr.

Coulombic forces on the other hand are much stronger & develop b/w charged atoms (already charged) or ions.

$F = \frac{Q_1 Q_2}{4\pi\epsilon_0 r^2}$: - Nature & mgn depend on signs of charges, medium, distance b/w the entities involved.

$\therefore Q_1 > 0 \quad Q_2 > 0 \quad \text{or} \quad Q_1 < 0 \quad Q_2 < 0$

$\Rightarrow F = +ve \quad \therefore \text{repulsion}$

$Q_1 < 0 \quad \& \quad Q_2 < 0 \quad \text{or} \quad \text{vice versa}$

$\Rightarrow F = -ve \quad \therefore \text{attraction}$

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