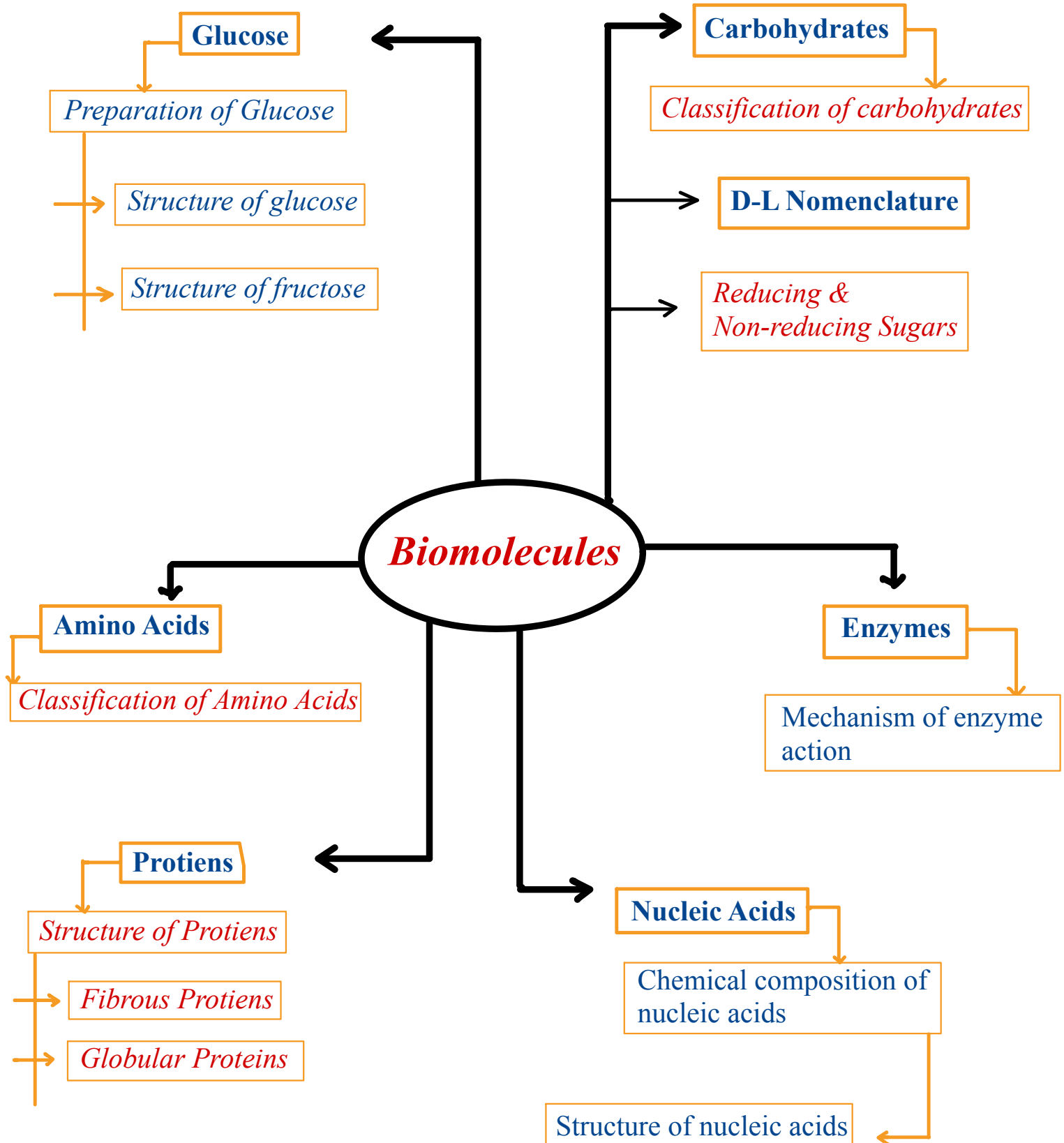


# Biomolecules

## Flow Chart Of Biomolecules



# Biomolecules

→ The branch of chemistry that deals with the molecules involved in living system, is called Biochemistry.

→ Carbohydrates, proteins, vitamins and nucleic acids are some of the major components of our body. These are collectively called Biomolecules.

Carbohydrates :- Carbohydrates are optically active polyhydroxy aldehydes or ketones or substances that will yield these types of compounds on hydrolysis.

↓  
Hydrates of Carbon

$C_x(H_2O)_y$  : General Formula.

Example - :  $C_6(H_2O)_6 \Rightarrow C_6O_6H_{12}$  ( $C_6H_{12}O_6$  : Glucose / Fructose)

## Classification of Carbohydrates :-

→ This classification is based on hydrolysis.

a.] Monosaccharides :- A carbohydrate that can not be hydrolysed further to give simpler unit of polyhydroxy aldehyde or ketones is called a monosaccharide.

For example - : Glucose / Fructose / Ribose

[CBSE 2010] → 1M

b.] Oligosaccharides :- Carbohydrates that produce 2 to 10 monosaccharide units on hydrolysis, are called oligosaccharide.

→ Disaccharide : It produce 2 unit of monosaccharide.

[CBSE 2019] 1M example →  $\frac{\text{Sucrose}}{\text{Lactose}}$  [ Sucrose  $\xrightarrow{\text{Hydrolysis}}$  Glucose + Fructose ]

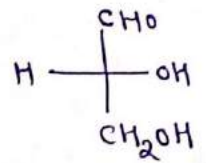
c.] Polysaccharides :- Carbohydrates that produce a large no. of monosaccharide units on hydrolysis are called polysaccharide.

example - : Starch / Cellulose / Glycogen.

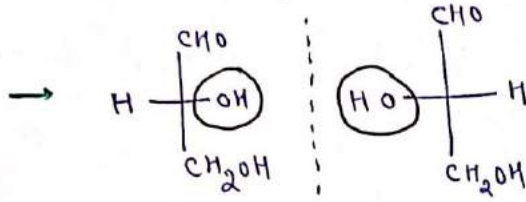
→ Polysaccharides are not sweet in taste. Hence they are also called non-sugars.

## D-L Nomenclature :-

Standard for this Nomenclature :



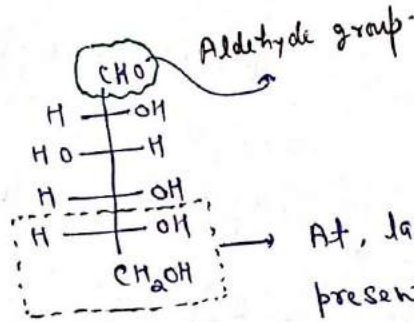
Glyceraldehyde



D-Glyceraldehyde Mirror L-Glyceraldehyde

→ (+) and (-) represents dextrorotatory nature and levorotatory nature of a compound, means that optical active nature can be defined by ⊕ or ⊖. But remember that D and L have no relation with optical activity of a compound, they represents only configuration of a compound.

→ Structure of Glucose :-



At, last chiral carbon -OH group is present at right side ⇒ D-Glucose.

## Reducing and Non-reducing Sugars :- [Delhi 2010c] (1M)

→ Reducing Sugars :- All those carbohydrates which reduce Tollen's reagent and Fehling reagent are called reducing sugars. [CBSE 2010] (1M)

→ All monosaccharides are reducing sugars. (example → Glucose and Fructose)

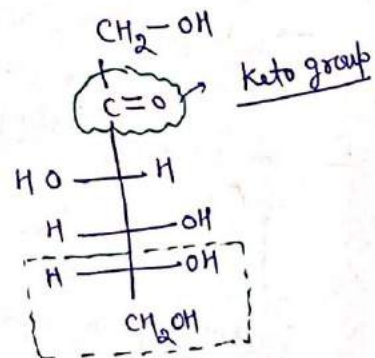
→ Non-reducing Sugars :- Carbohydrates which can not reduce Tollen's reagent and Fehling solution are called non-reducing sugars.

For example → Sucrose

## Classification of monosaccharides :-

Aldose :- Monosaccharide containing aldehyde group.

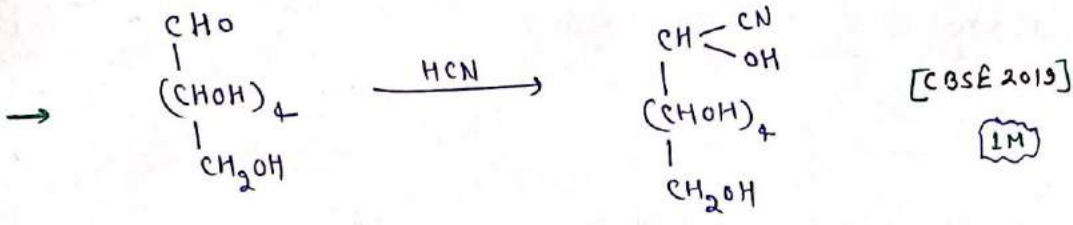
Ketose :- Monosaccharide containing keto group.



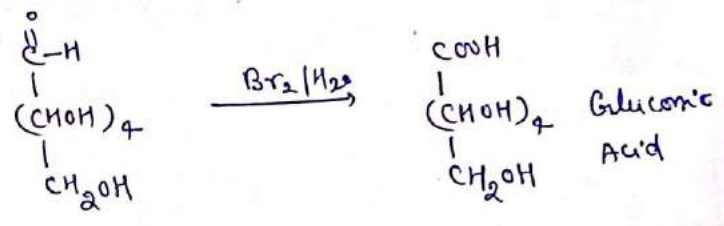
D-Fructose

→ Glucose is an example of Aldose, while Fructose is an example of Ketose.

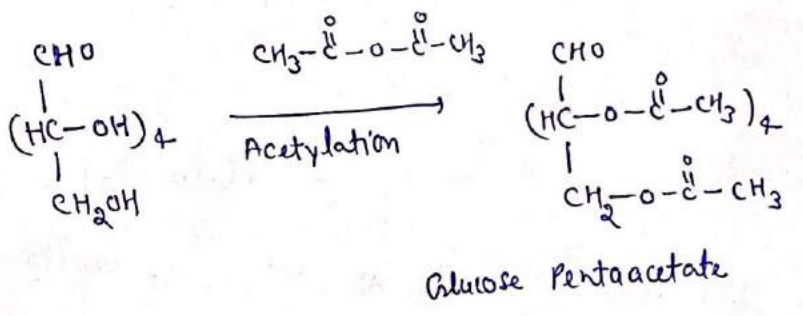




c.] Presence of aldehyde group :-  
 [CBSE 2019/2018] (1M)

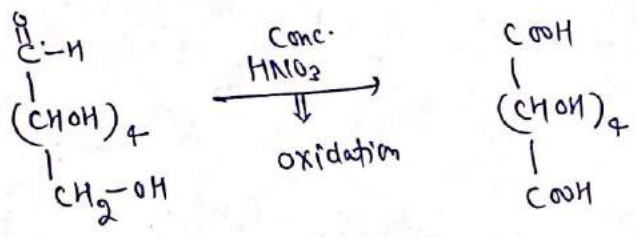


d.] Presence of 5 OH groups :-  
 [CBSE 2019] (1M)



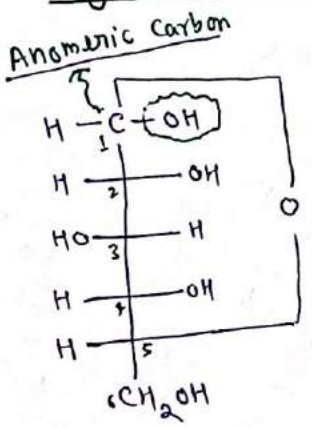
e.] Presence of 1° alcoholic group (-OH) :-

[CBSE 2018] (1M)  
 Conc. HNO<sub>3</sub> oxidises only aldehyde and 1° alcohol.

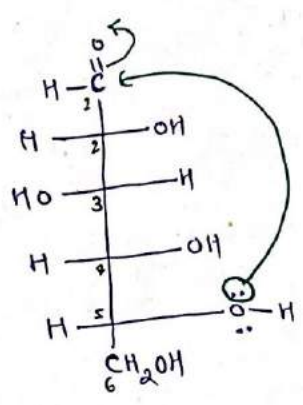


Cyclic Structure of glucose :-

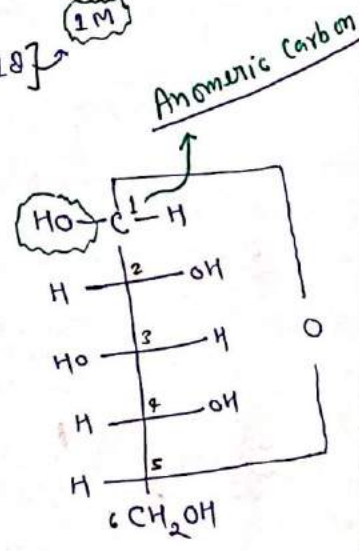
[Delhi 2012/2011 / CBSE 2018] (1M)



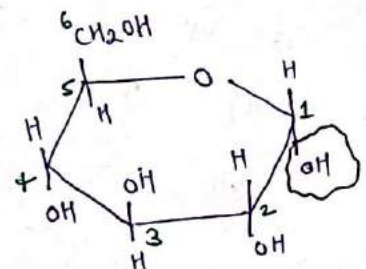
α-D-Glucose



D-Glucose



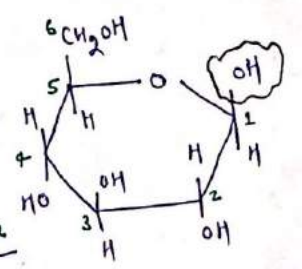
β-D-Glucose




α-D-Glucopyranose

Haworth Projection

β-D-Glucopyranose



→ In Haworth Projection, the six membered cyclic structure of glucose is called Pyranose structure. (In analogy with Pyran ).

→ Anomers—: Anomers are isomers that differ in the configuration at the acetal or hemiacetal carbon atom of a sugar in its cyclic form.

For example —:  $\alpha$ -D-Glucose and  $\beta$ -D-Glucose are Anomers.

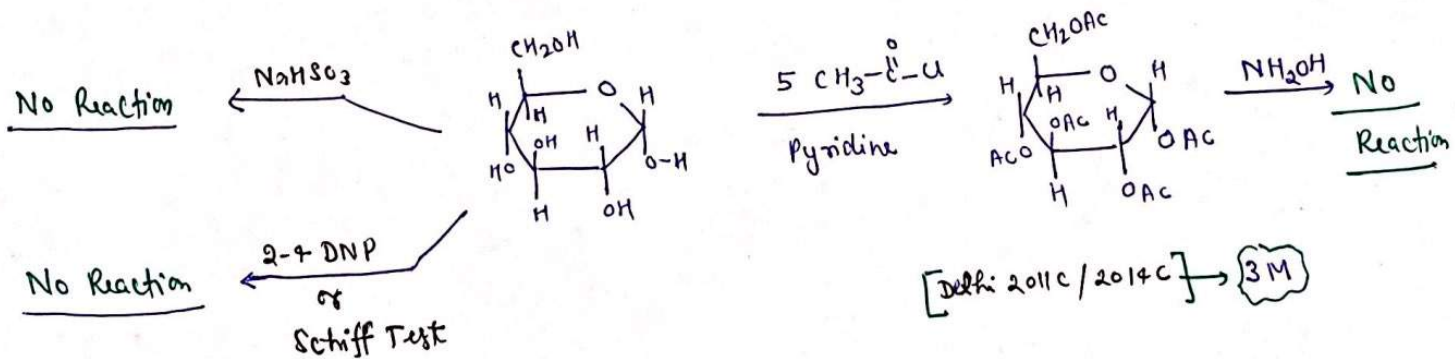
CBSE 2014

↓  
1M

Cyclic structure of glucose : Supporting Evidence :- [CBSE 2010C/2011/2011C] 2M

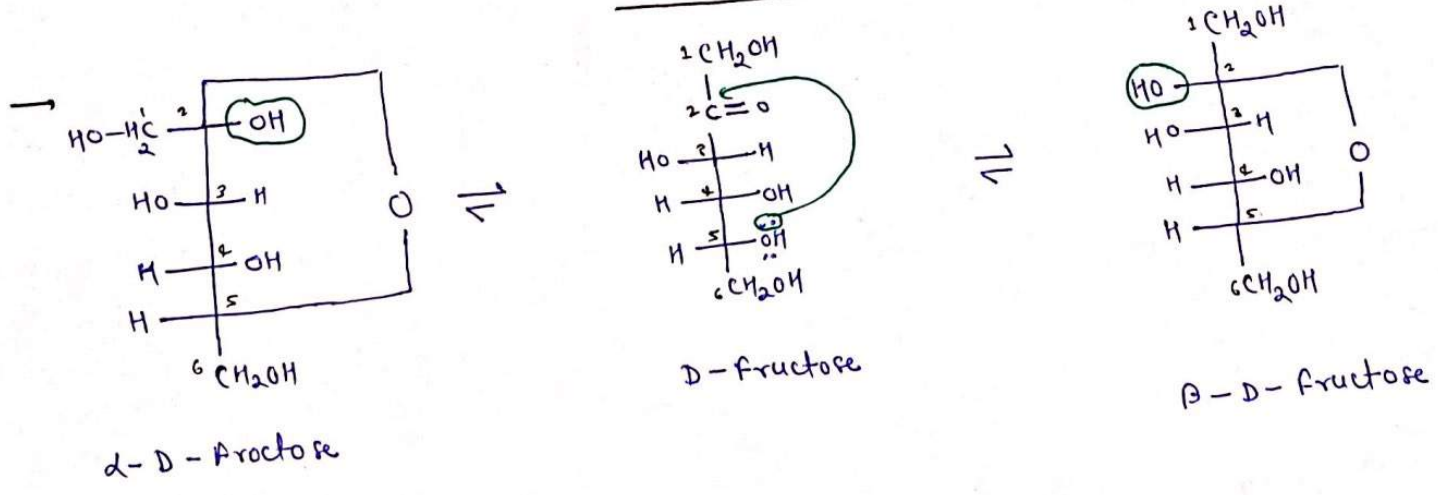
(i) Despite having aldehyde group, glucose does not give 2,4-DNP test, Schiff test and it does not form adduct with  $\text{NaHSO}_3$ .

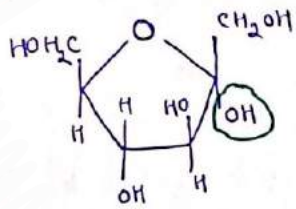
(ii) Pentaacetate of glucose does not react with  $\text{NH}_2\text{-OH}$  indicating the absence of free  $\text{-CHO}$  group.



(iii) Glucose is found to exist in two different crystalline forms which are named as  $\alpha$  and  $\beta$ . They both have different melting point and different temperature for crystallisation.

Structure of Fructose —:

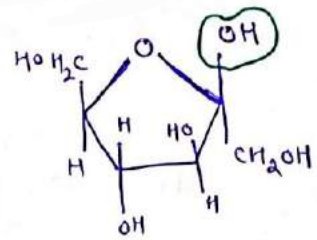




$\alpha$ -D-Fructofuranose



Furan

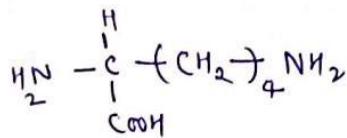


$\beta$ -D-Fructofuranose



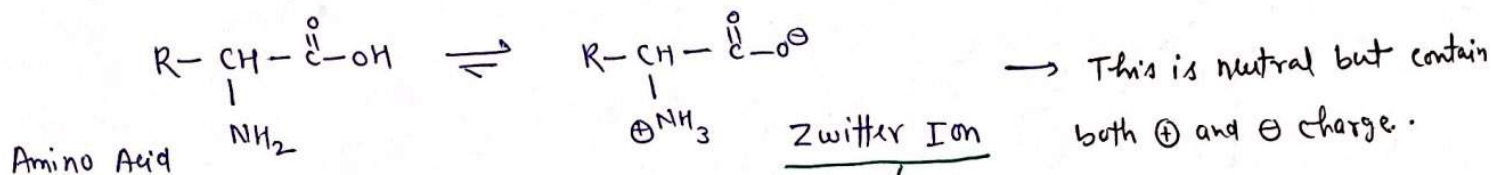
c. → Basic Amino Acid -: More no. of amino group than carboxyl group.

For example → Lysine

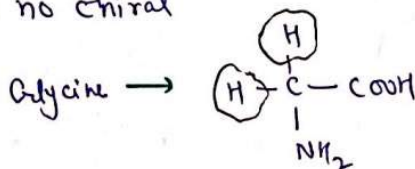


NOTE -: Amino acids are crystalline solids. These are water soluble and behave like salts rather than simple amines or carboxylic acids.

Zwitter Ion -: Due to presence of both acidic (carboxyl group -COOH) and basic (-NH<sub>2</sub> group) in the same molecule, in aqueous solution -COOH group can lose a proton and -NH<sub>2</sub> group can accept a proton giving rise to a dipolar ion. This dipolar ion is known as zwitter ion. [CBSE 2011C] (1M)



→ All α-amino acids are optically active except Glycine. Because there is no chiral carbon in glycine.



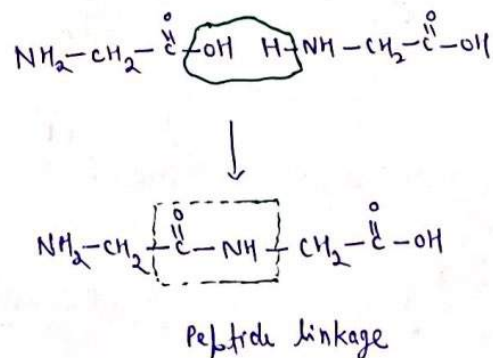
This can react with both acid and base. So, has amphoteric character. [CBSE 2015] (1M) 2019

## Proteins

→ Proteins are most abundant biomolecules of the living system.

→ Proteins are the polymers of α-amino groups and they are connected to each other by peptide bond or peptide linkage. (1M) [CBSE 2014C / CBSE 2013 / CBSE 2015 / Delhi 2014] 2016

chemically, peptide linkage is an amide linkage formed between -COOH group of one α-amino acid and -NH<sub>2</sub> group of other α-amino acid formed by the loss of water molecule.

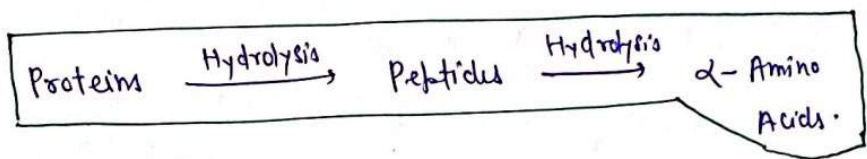


→ Dipeptide :- Combination of 2 amino acid by peptide bond is known as dipeptide<sup>-de</sup>.

→ Similarly, a tripeptide contains 3 amino acids linked by 2 peptide linkages.

→ Polypeptide :- Combination of 10 or more than 10 amino acids by peptide bonds, is known as polypeptide. (1M) [Delhi 2010]

→ Protein is a polypeptide.



[Case 2010/2016/Delhi 2019] → (1M)

Classification of Proteins :- Two types on the basis of their molecular shape.

a.) Fibrous Proteins :- When polypeptide chains run parallel and are held together by hydrogen and disulphide bonds, then fibre like structure is formed.

→ Such proteins are insoluble in water.

→ Example :- Keratin [hair/wool/silk] and myosin [present in muscles].

b.) Globular Proteins :- The chains of polypeptides coil around to give a spherical shape. These are usually soluble in water.

→ Example :- Insulin and albumins.

→ Structure and shape of proteins can be studied at four different levels

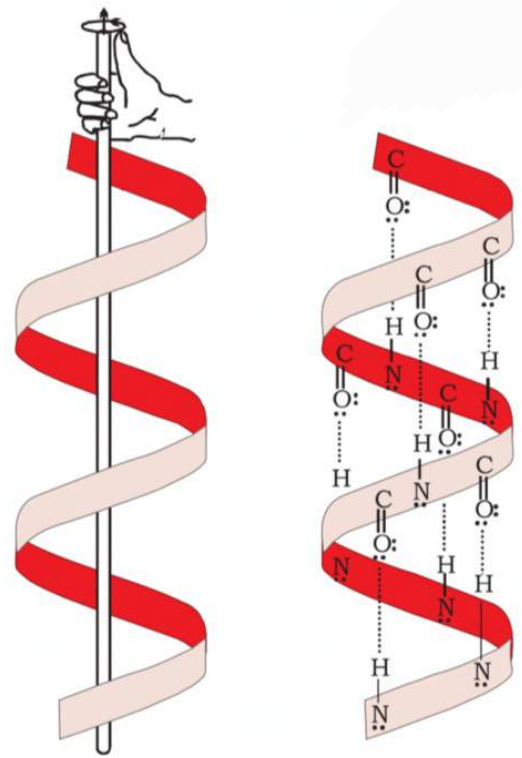
→ Primary, secondary, tertiary and quaternary, each level being more complex than previous one.

a.) Primary Structure of proteins :- In a protein molecule, one or more polypeptide chains may be present. Each polypeptide chain in a protein is linked together in a specific sequence of amino acids. This sequence of amino acids is termed as primary structure of proteins. [Case 2015] (1M)

b.) 2<sup>o</sup> Structure of proteins :- It refers to the shape in which a long polypeptide chain can exist. They are found to exist in two different types of structures → α-Helix & β-Sheet.

→ These structures arise due to regular folding of backbone of polypeptide chain due to hydrogen bonding between  $-C=O-$  and  $-NH-$  groups of peptide bond.

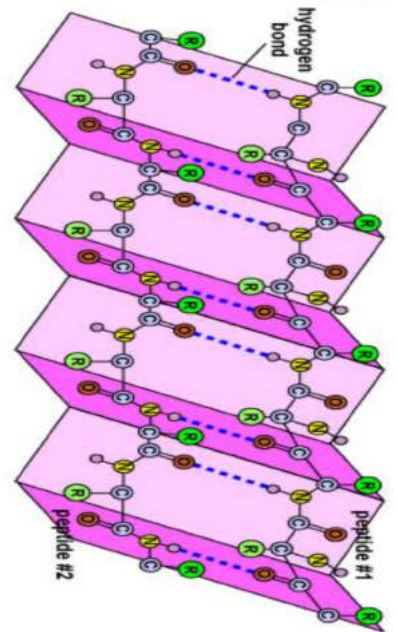
$\alpha$ -Helix -: It is one of the most common ways in which a polypeptide chain forms all possible hydrogen bonds by twisting into a right handed screw (helix). This hydrogen bond is in between  $-NH-$  group of each amino acid to the  $-C=O-$  group of an adjacent turn of helix. [Dutt 2013 / CBSE 2010] 1M



$\beta$ -pleated sheet -: In  $\beta$ -structure, all peptide chains are stretched out to maximum extent and then laid side by side (which are held together by intermolecular hydrogen bonding).

→ The structure resembles the pleated folds of drapery and therefore is known as  $\beta$ -pleated sheet.

c.) Tertiary structure of proteins -: It represents further folding of secondary structure. It gives rise to two major molecular shapes → fibre and globular.

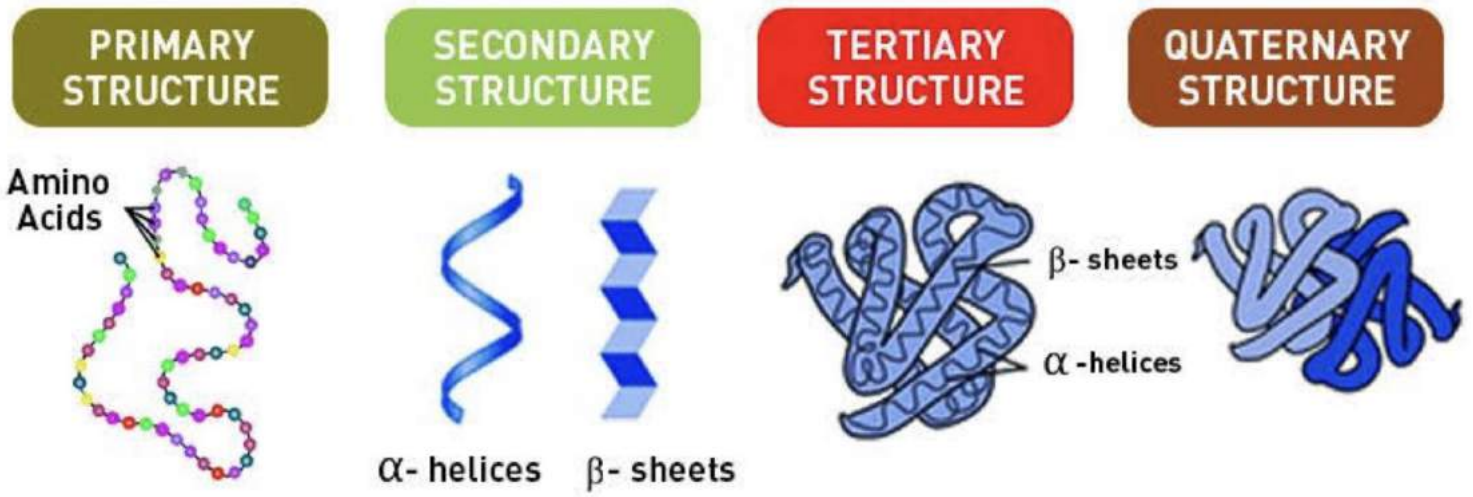


→ Stability of this structure depends on H-bonding, disulphide linkages, Vander Waals force of attraction and electrostatic forces of attraction.

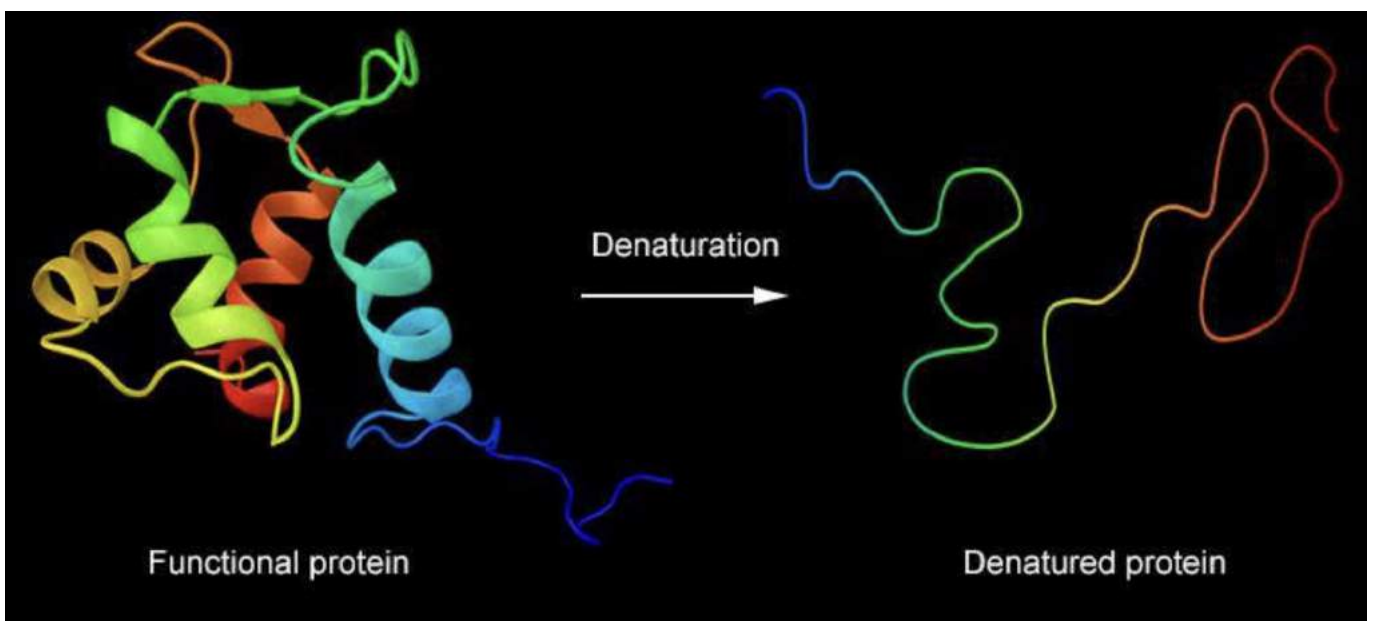
d.) Quaternary structure of proteins :- The spatial arrangement of sub units of proteins [which are composed of two or more polypeptide chains] with respect to each other is known as quaternary structure.

Denaturation of Proteins -:

Native Protein → Protein found in a biological system with a unique three dimensional structure (3D) and biological activity is called native protein.



## Desaturation of Proteins



→ When a native form of protein is subjected to a physical change (like change in temperature) or chemical change (like change in pH) hydrogen bonds are disturbed. Due to this unfolding of proteins or uncoiling of helix happens and protein loses its biological activity. This is called Denaturation of protein.

→ During denaturation 2<sup>o</sup>/3<sup>o</sup> structures are destroyed but 1<sup>o</sup> structure remains intact. [Delhi 2013/2014] [CBSE 2010/2015] (1M)

Example → The coagulation of egg white on boiling → upon boiling the egg, denaturation followed by coagulation occurs. The water present in egg gets adsorbed / absorbed in coagulated proteins through Hydrogen bonding. [Delhi 2010] (1M)

→ Curdling of milk

## Nucleic Acids

→ Nucleus of a living cell is responsible for the transmission of inherent characters.

→ The particles in nucleus of cell (responsible for heredity), are called chromosomes.

→ Chromosomes are made up of proteins and nucleic acids.

→ [Deoxyribonucleic acid] DNA ← Nucleic Acids → RNA [Ribonucleic Acid] [Delhi 2012/2011 | CBSE 2011] (1M)

### Chemical composition of nucleic acids —:

→ Nucleic Acid  $\xrightarrow{\text{Hydrolysis}}$  Pentose sugar + Phosphoric acid + base

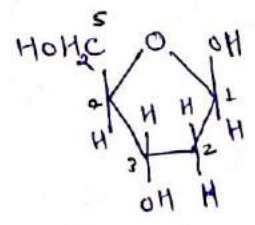
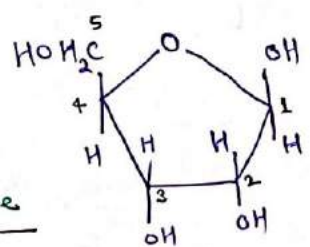
→ DNA ⇒  $\beta$ -D-2-deoxyribose + Phosphoric acid + [A G C T]

→ RNA ⇒  $\beta$ -D-ribose + Phosphoric acid + [A G C U]

↑  
Nitrogen containing heterocyclic compounds.

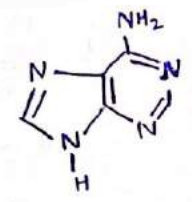
Pentose sugar →

$\beta$ -D-ribose

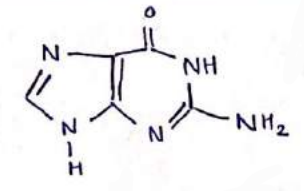


$\beta$ -D-2-deoxyribose

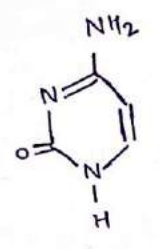
→ Adenine (A)



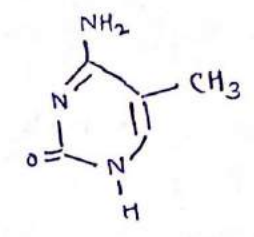
→ Guanine (G)



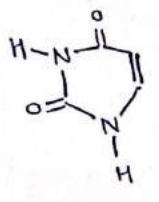
→ Cytosine (C)



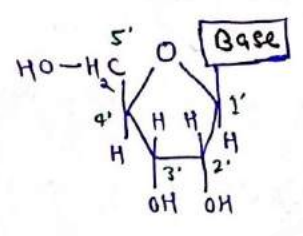
→ Thymine (T)



→ Uracil (U)



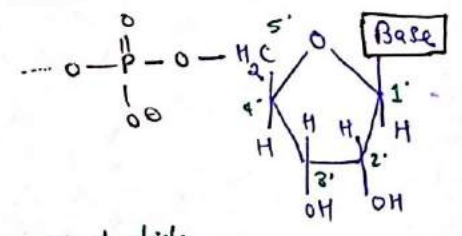
→ Nucleoside = Sugar + base



Delhi 2010 & 2014  
← Nucleoside

→ Nucleotide = phosphate + Nucleoside

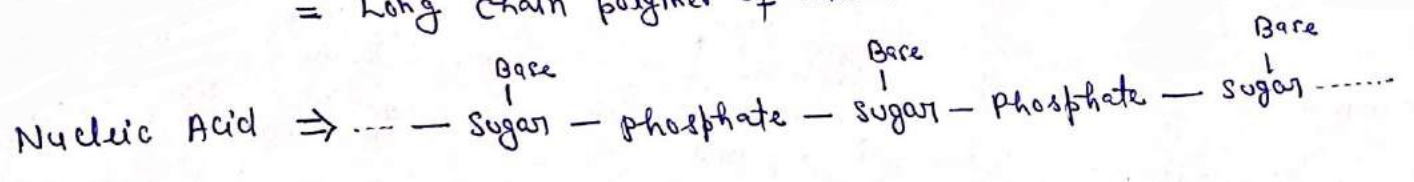
= phosphate + sugar + base



Nucleotide

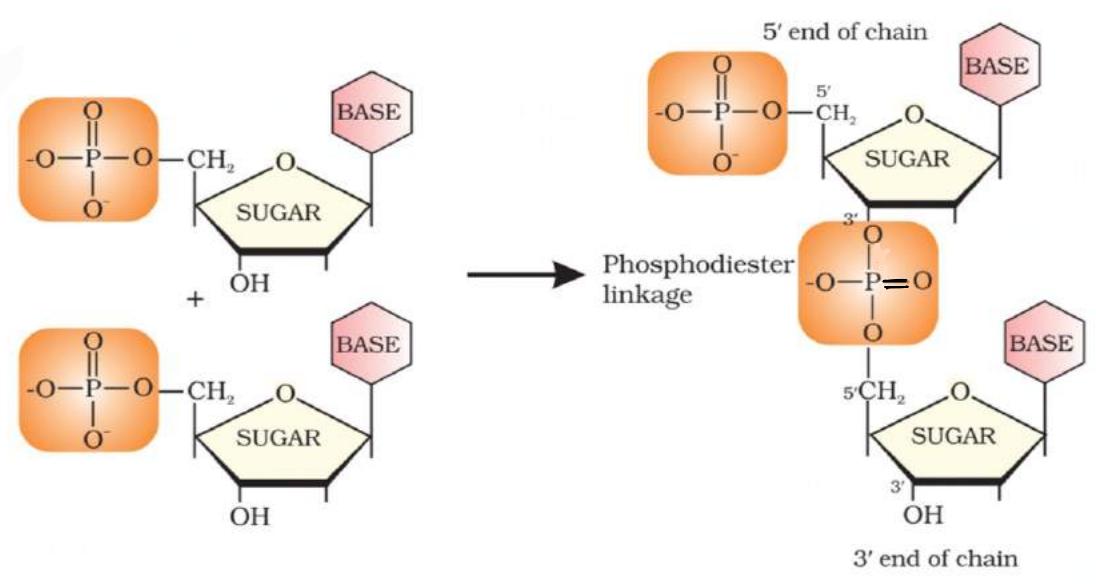
→ Nucleic Acid = Many nucleotides = Polynucleotides

= Long chain polymer of nucleotides.

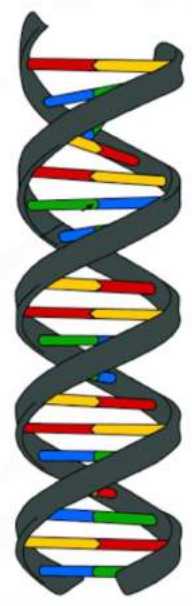
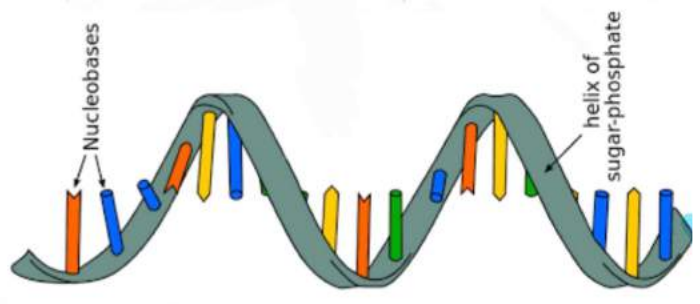
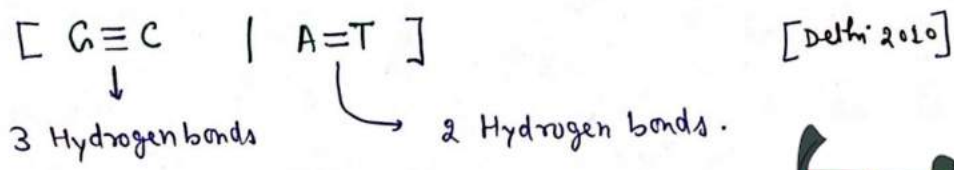


→ In a nucleotide, base is connected to 1' carbon of sugar and phosphate is connected to 5' carbon of sugar.

→ Nucleotides are joined together by phosphodiester linkage between 5' and 3' carbon atoms of the pentose sugar.



→ Double Strand helix structure for DNA :- The two strands are complementary to each other because the hydrogen bonds are formed between specific pairs of bases. Adenine forms hydrogen bonds with Thymine, whereas Cytosine forms hydrogen bonds with Guanine.



RNA :- Structure :- Single stranded helix.

- RNA molecules are of 3 types.
- (i) messenger RNA [m-RNA]
  - (ii) Ribosomal RNA [r-RNA]
  - (iii) Transfer RNA [t-RNA]
- [Delhi 2013] (1M)

Biological functions of Nucleic Acids :- DNA is the chemical basis of heredity and may be regarded as the reserve of genetic information. Another important function of nucleic acids is the protein synthesis in the cell.

