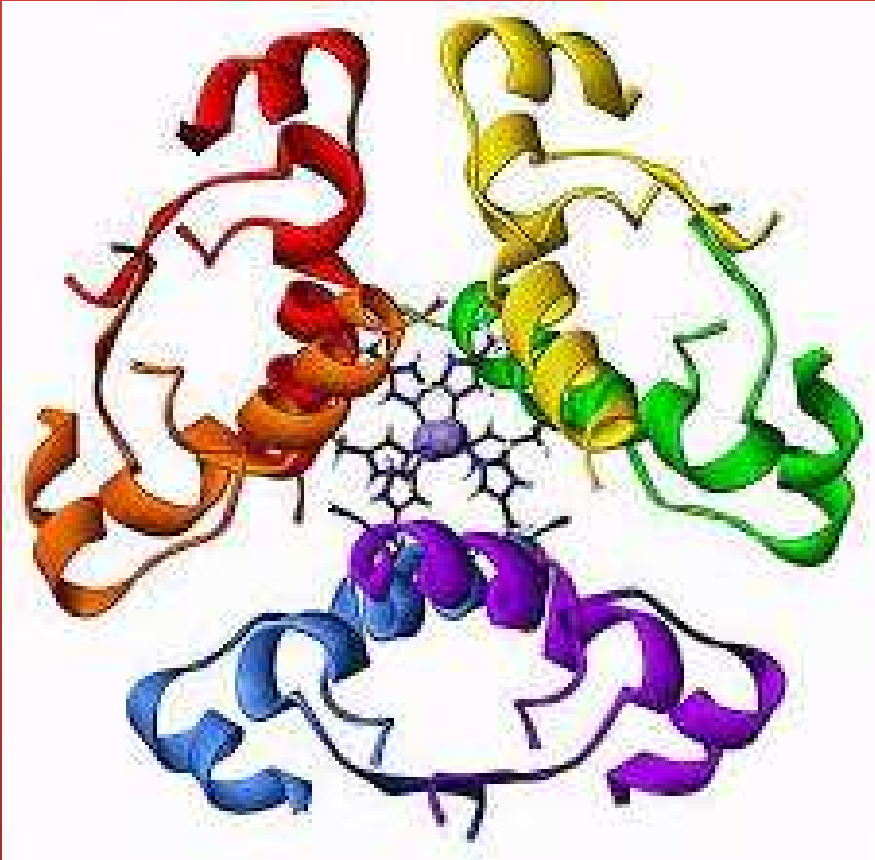


# Proteins



By

**S.K.Sinha , Kota**

# Levels of Protein Structure

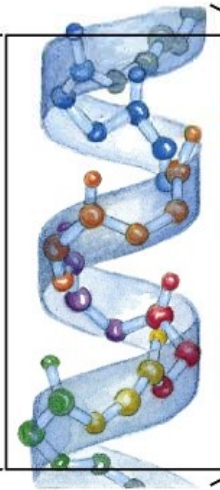
Proteins show 4 levels of structural organisation:

Primary structure



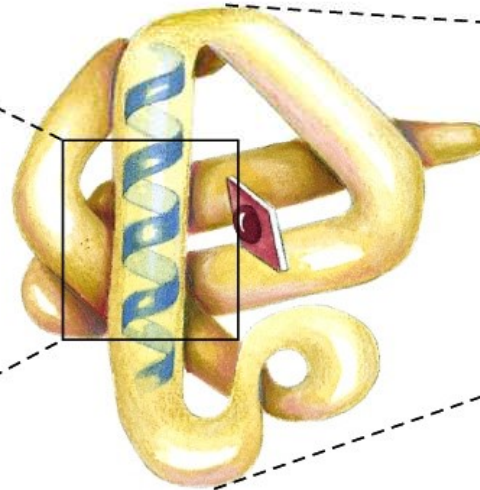
Amino acid residues

Secondary structure



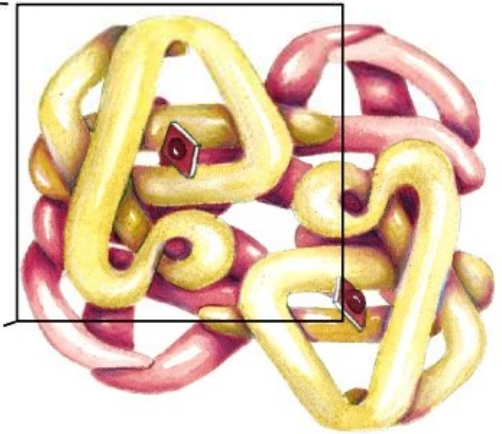
$\alpha$  Helix

Tertiary structure



Polypeptide chain

Quaternary structure



Assembled subunits

Sinha Sir Kota

## Levels of Protein Structure

**Proteins show 4 levels of structural organisation:**

**1. Primary structure = amino acid sequence**

- **Determined by the genetic code of the mRNA.**

**2. Secondary structure = folding and twisting of a single polypeptide chain.**

- **Result of weak H-bond and electrostatic interactions.**
- **e.g.,  $\alpha$ -helix (coiled) and  $\beta$ -pleated sheet (zig-zag).**

**Sinha Sir Kota**

## Levels of Protein Structure

**3. Tertiary structure = three dimensional shape (or conformation) of a polypeptide chain.**

- **Function of R groups contained in the polypeptide.**

**4. Quaternary structure = association between polypeptides in multi-subunit proteins (e.g. hemoglobin).**

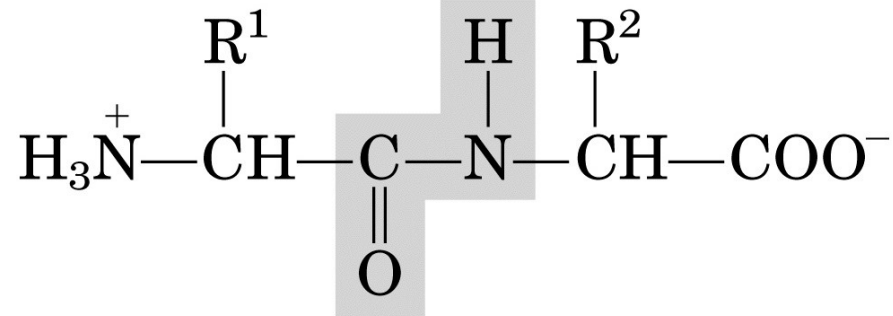
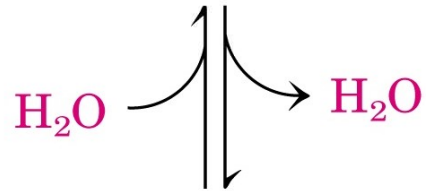
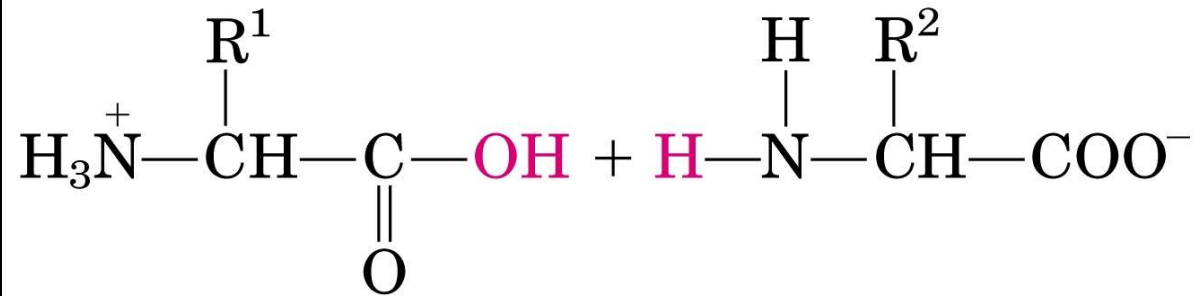
- **Occurs only with two or more polypeptides.**

## Peptide Bonds

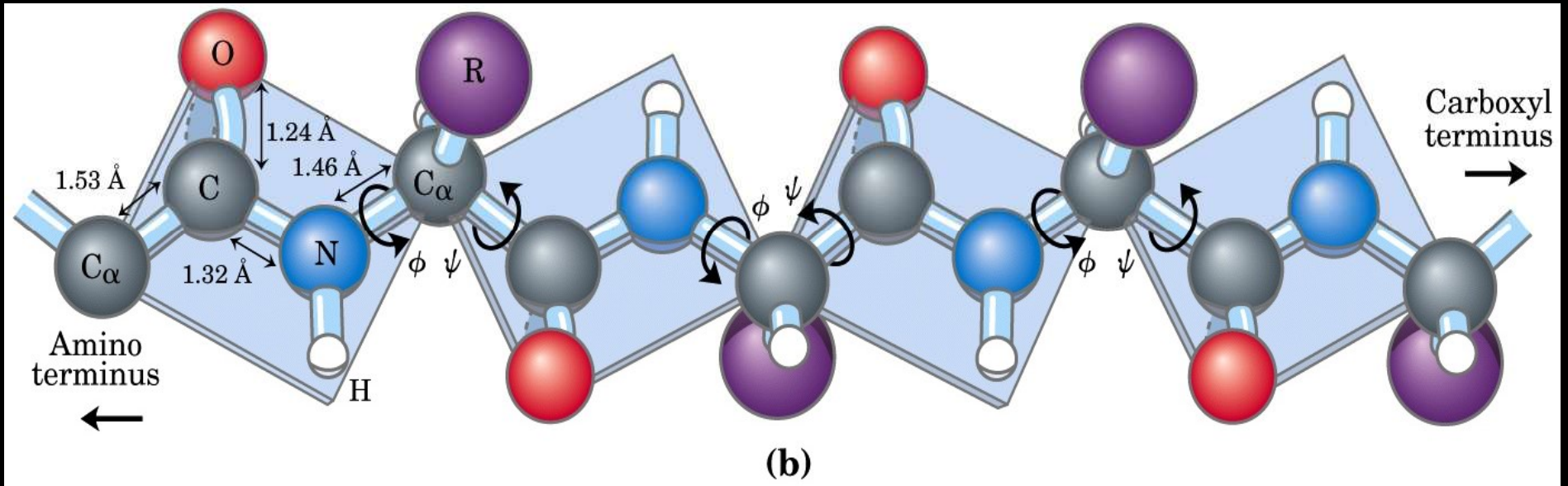
**– $\alpha$ -carboxyl of one amino acid is joined to  $\alpha$ -amino of a second amino acid (with removal of water)**

**- only  $\alpha$ -carboxyl and  $\alpha$ -amino groups are used, not R-group carboxyl or amino groups.**

## Peptide bond formation



# The peptide bond is planar

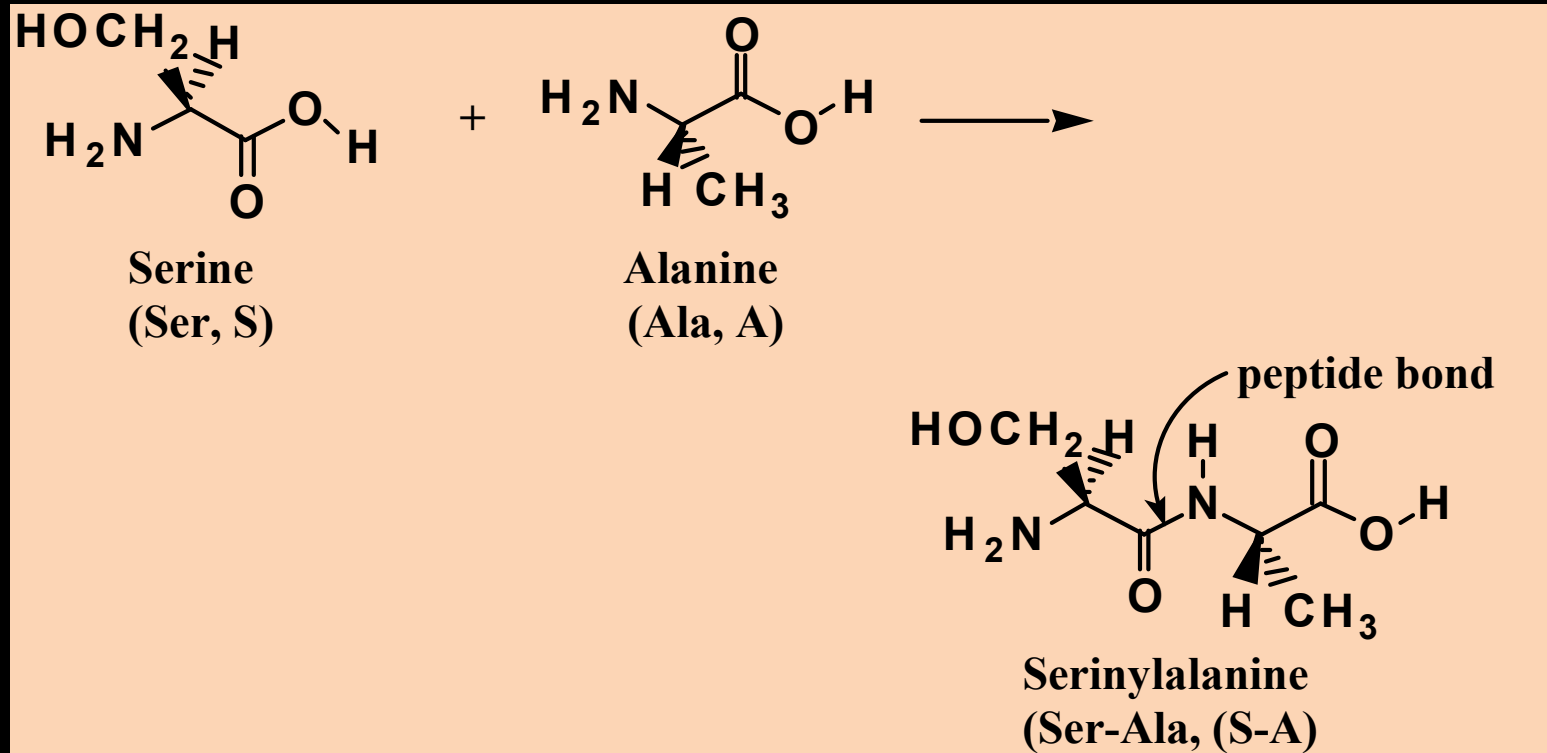


## Polypeptides & Proteins

**Peptide bond:** The special name given to the amide bond between the  $\alpha$ -carboxyl group of one amino acid and the  $\alpha$ -amino group of another.



## Serinylalanine (Ser-Ala)



## Peptides

**Dipeptide:** A molecule containing two amino acids joined by a peptide bond.

## Peptides

**Tripeptide:** A molecule containing three amino acids joined by peptide bonds.

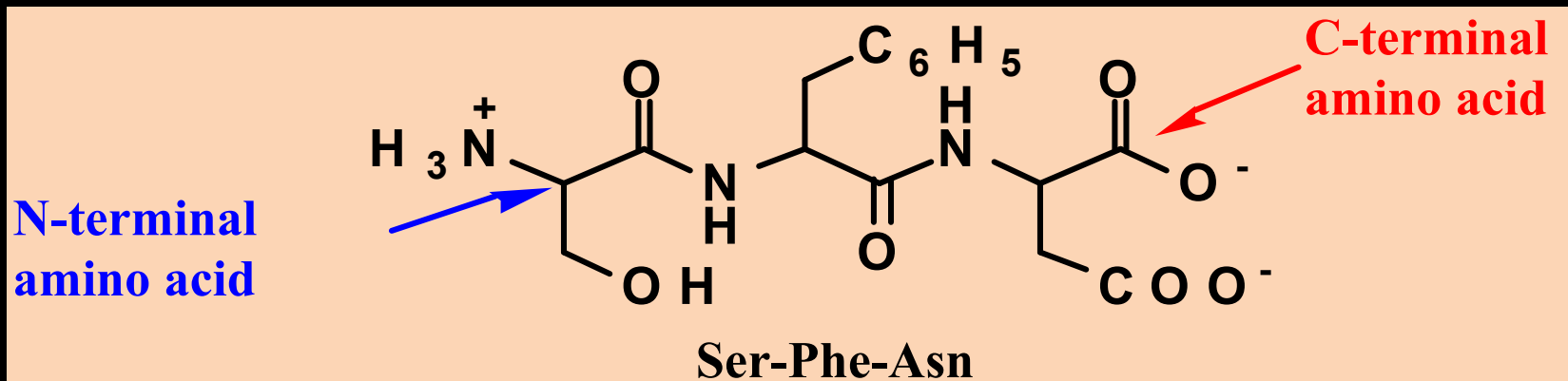
**Polypeptide:** A macromolecule containing many amino acids joined by peptide bonds.

## Peptides

**Protein:** A biological macromolecule of molecular weight 5000 g/mol or greater, consisting of one or more polypeptide chains.

## Writing Peptides

By convention, peptides are written from the left, beginning with the free - $\text{NH}_3^+$  group and ending with the free - $\text{COO}^-$  group on the right.

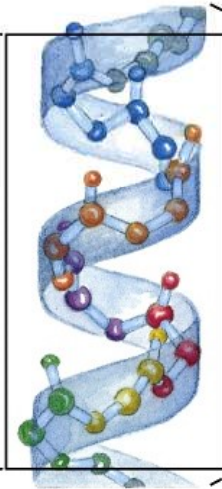


# Secondary structure

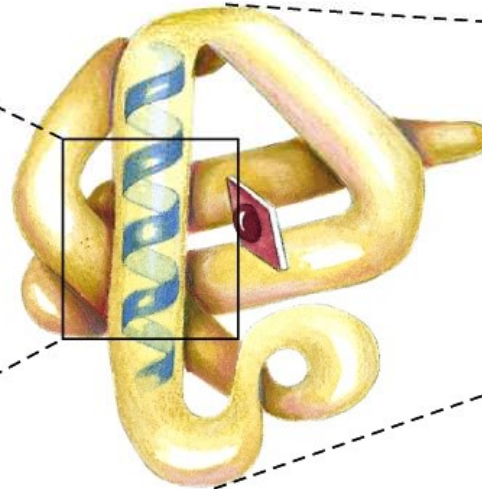
Primary structure



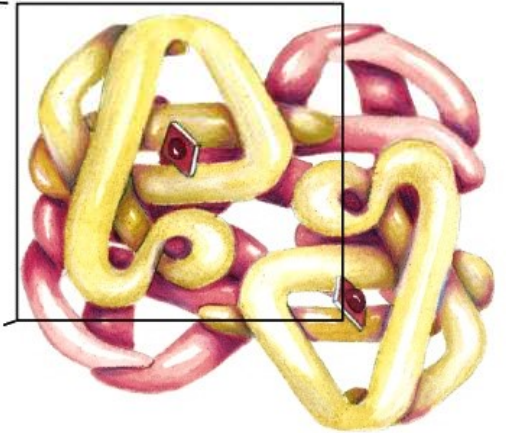
Secondary structure



Tertiary structure



Quaternary structure



Amino acid residues

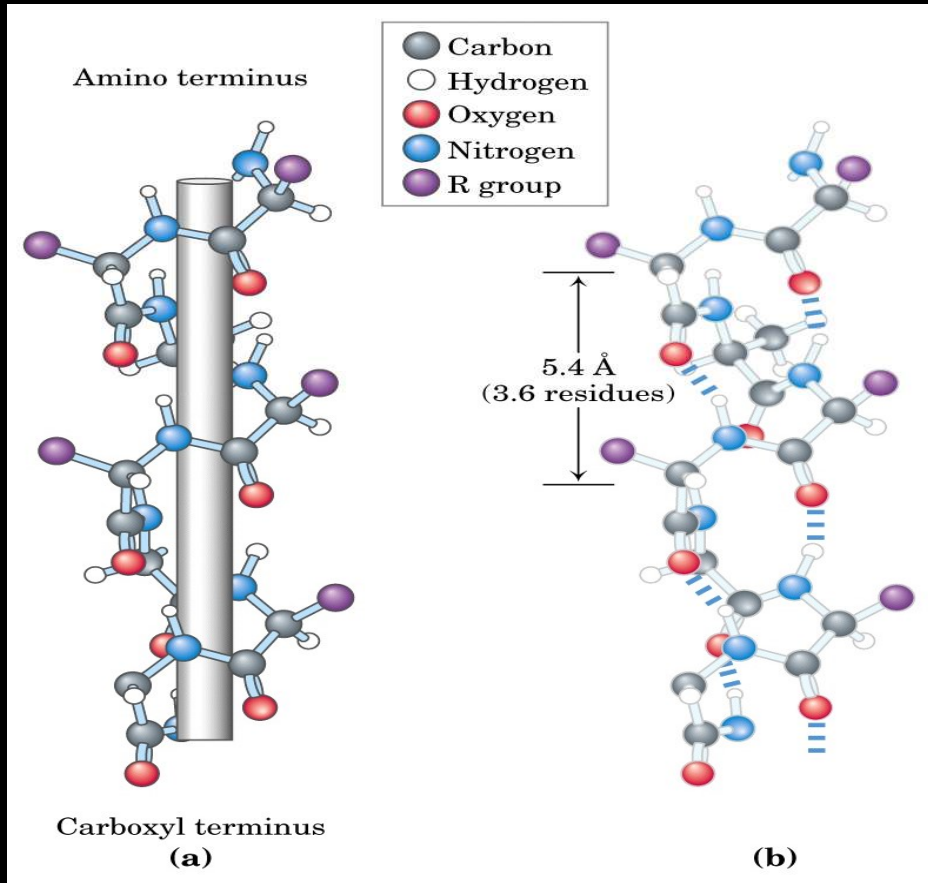
$\alpha$  Helix

Polypeptide chain

Assembled subunits

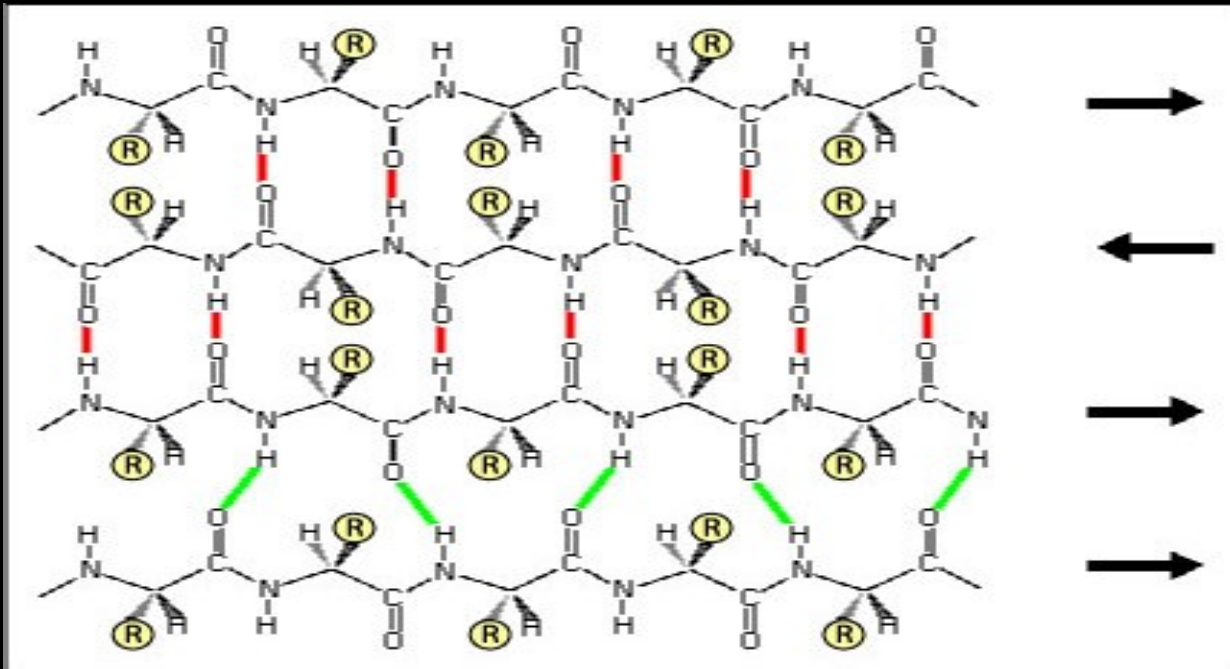
Sinha Sir Kota

# The $\alpha$ -helix



- In the  $\alpha$ -helix, the carbonyl oxygen of residue “i” forms a hydrogen bond with the amide of residue “i+4”.
- Although each hydrogen bond is relatively weak in isolation, the sum of the hydrogen bonds in a helix makes it quite stable.
- The propensity of a peptide for forming an  $\alpha$ -helix also depends on its sequence.

# Protein structure: sheets



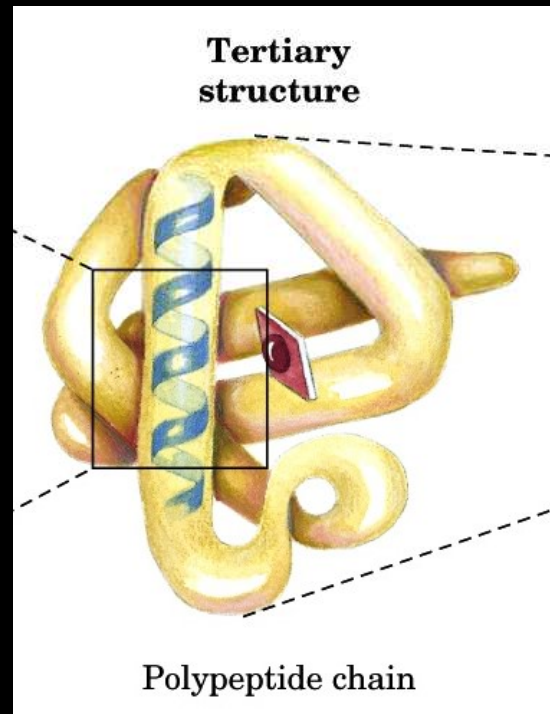
→  
←  
→  
→

- notice the difference in H-bonding pattern between parallel and anti-parallel beta-sheets

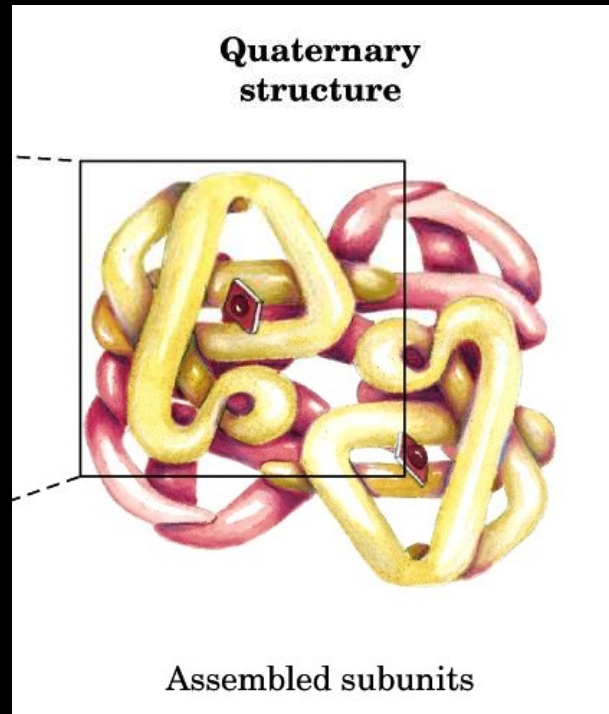
- also notice orientation of side chains relative to the sheets



# Tertiary structure



# Quaternary structure



# Types of interactions

<b>interaction</b>	<b>nature</b>	<b>bond length</b>	<b>“bond” strength</b>	<b>example</b>
<b>ionic (salt bridge)</b>	electrostatic	1.8-4.0 Å (3.0-10 Å for like charges)	1-6 kcal/mol	positive: K, R, H, N-terminus negative: D, E, C-terminus
<b>hydrophobic</b>	entropy	-	2-3	hydrophobic side chains (M,I,L,V,F,W,Y,A,C,P)
<b>H-bond</b>	H-bonding	2.6-3.5	2-10	H donor, O acceptor
<b>van der Waals</b>	attraction/ repulsion	2.8-4.0	<1	closely-spaced atoms; if too close, repulsion
<b>aromatic-aromatic</b>	$\pi-\pi$	4.5-7.0	1-2	F,W,Y (stacked)
<b>aromatic-amino group</b>	H-bonding	2.9-3.6	2.7-4.9	N-H donor to F,W,Y

# **Protein-solvent interactions**

## **hydrophilic amino acids**

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- **these amino acids tend to interact extensively with solvent in context of the folded protein; the interaction is mostly ionic and H-bonding**

# **Protein-solvent interactions**

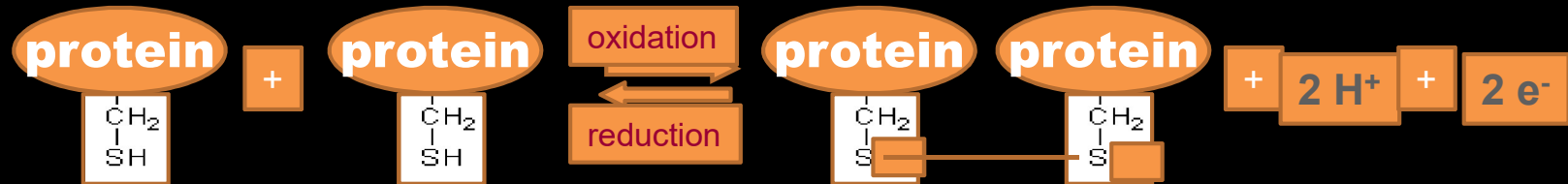
## **hydrophobic amino acids**

**-these tend to form the 'core' of the protein, *i.e.*, are buried within the folded protein; some hydrophobic residues can be entirely (or partially) exposed**

## **small neutral amino acids**

**- less preference for being solvent-exposed or not**

# The disulfide bond



- **Disulfide bond formation is a covalent modification; the oxidation reaction can either be intramolecular (within the same protein) or inter-molecular (within different proteins)**

## Protein denaturants

- **high temperatures**
  - ***cause protein unfolding, aggregation***
- **low temperatures**
  - ***some proteins are sensitive to cold denaturation***
- **heavy metals (e.g., lead, cadmium, etc.)**
  - ***highly toxic; efficiently induce the 'stress response'***

## Protein denaturants

- **proteotoxic agents (e.g., alcohols, cross-linking agents, etc.)**
- **oxygen radicals, ionizing radiation**
  - ***cause permanent protein damage***
- **chaotropes (urea, guanidine hydrochloride, etc.)**
  - ***highly potent at denaturing proteins; often used in protein folding studies***



# Classes of proteins

## Functional definition:

**Enzymes:** Accelerate biochemical reactions

**Structural:** Form biological structures

**Transport:** Carry biochemically important substances

**Defense:** Protect the body from foreign invaders

# Classes of proteins

## Structural definition:

**Globular:** Complex folds, irregularly shaped tertiary structures

**Fibrous:** Extended, simple folds - generally structural proteins

## Cellular localization definition:

**Membrane:** In direct physical contact with a membrane; generally water insoluble.

**Soluble:** Water soluble; can be anywhere in the cell.

# Classes of proteins

## Conjugated Proteins

	<b>Prosthetic group</b>	<b>Example</b>
<b>Lipoproteins</b>	<b>Lipids</b>	<b><math>\beta_1</math>-Lipoprotein of blood</b>
<b>Glycoproteins</b>	<b>Carbohydrates</b>	<b>Immunoglobulin G</b>
<b>Phosphoproteins</b>	<b>Phosphate groups</b>	<b>Casein of milk</b>
<b>Hemoproteins</b>	<b>Heme (iron porphyrin)</b>	<b>Hemoglobin</b>
<b>Flavoproteins</b>	<b>Flavin nucleotides)</b>	<b>Succinate dehydrogenase</b>
<b>Metalloproteins</b>	<b>Iron</b>	<b>Ferritin</b>
	<b>Calcium</b>	<b>Calmodulin</b>
	<b>Copper</b>	<b>Plastocyanin</b>

## Use of Amino Acids

**Aspartame** (aspartyl-phenylalanine-1-methyl ester) is an artificial sweetener.

**5-HTP** (5-hydroxytryptophan) has been used to treat neurological problems associated with PKU (phenylketonuria), as well as depression.

**L-DOPA** (L-dihydroxyphenylalanine) is a drug used to treat Parkinsonism.

**Monosodium glutamate** is a food additive to enhance flavor.

## **Classification of Proteins**

**Proteins can be classified into two types on the basis of their molecular shape.**

**(a) Fibrous proteins :**

**(b) Globular proteins :**

## Use of Amino Acids

**(a) Fibrous proteins :** The polypeptide chains run parallel and are held together by hydrogen and disulphide bonds, then fibre– like structure is formed.

Such proteins are generally insoluble in water.

**Examples:**

- 1. keratin (present in hair, wool, silk)**
- 2. Myosin (present in muscles),**

## **Use of Amino Acids**

### **(b) Globular proteins:**

**This structure results when the chains of polypeptides coil around to give a spherical shape. These are usually soluble in water.**

### **Examples:**

- 1. Insulin**
- 2. albumins**