

IIT Guwahati

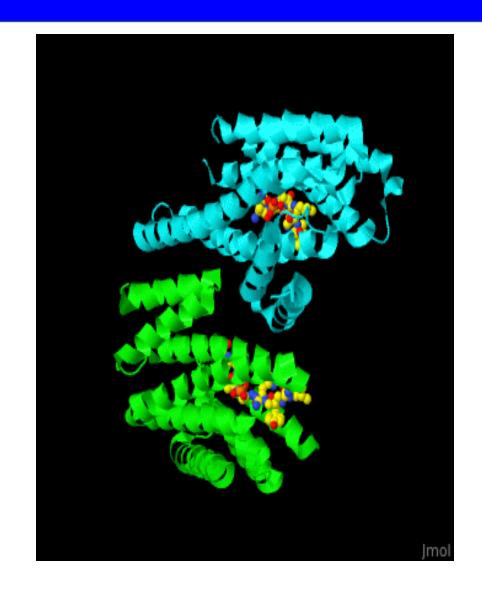
Lecture 5

Course BT 631

Protein Structure function and Crystallography

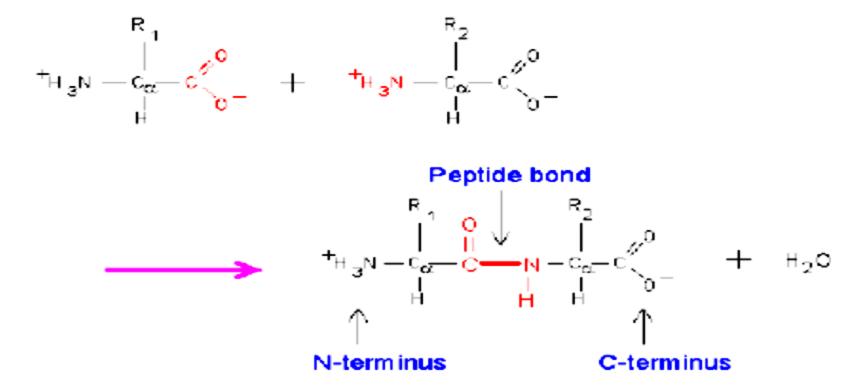
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Peptide Bond

- Amino acids are linked by forming amide bond between α -carboxylic group of one amino acid and α Amino group of other amino acid, it is called Peptide bond.
- When two amino acids are joined together by a peptide bond it is called as Dipeptide.



Formation of Peptide Bond

Types of Peptide Bond

- 1. Linear Peptide: It is formed between α -Carboxylic group of one amino acid and α -Amino group of other amino acid e.g. Most of the peptide bonds
- 2. Semi-cyclic Peptide Bond: It is formed between α-Carboxylic group or α-Amino group of an amino acid and its side chain containing Amino group or carboxylic group, respectively. This type of peptide bond is known as semi-cyclic peptide bond. e.g. Bacitracin A
- 3. Cyclic Peptide Bond: It is formed between N-terminal amino group and C-terminal carboxyl group of amino acid of the same polypeptide chain. This type of peptide bond is known as cyclic peptide bond. e.g. Tyrocidine is naturally occurring cyclo-decapeptide. Gramicidin S is a decapeptide. Vallinomycin (antibiotic) is also cyclo-peptide.

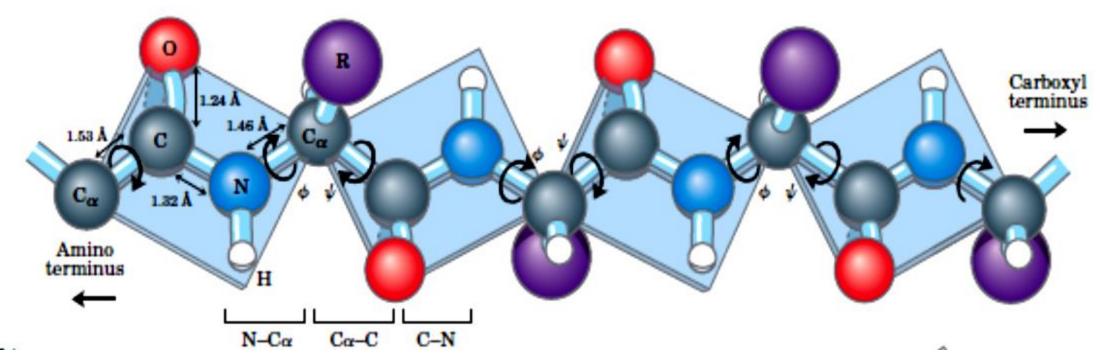
- 1. Peptide bond is rigid and planner.
- 2. Peptide bond has Partial Double Bond characteristics (Double bond is due to Resonance). On average, a peptide bond length is 1.32 Å as compared to 1.45 Å for an ordinary C-N bond. Whereas, the average bond length of a C=N double bond is 1.25 Å.

Thus, partial double bond restricts the rotation around this bond. This leads to the six atoms being coplanar.

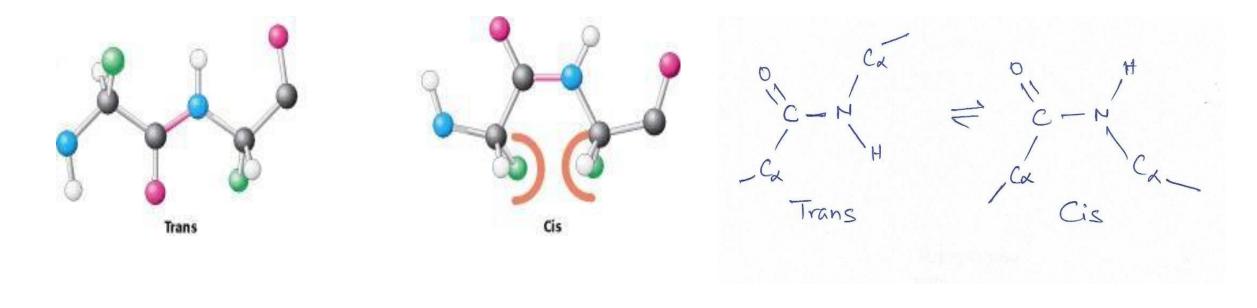
The carbonyl oxygen has a partial negative charge and the amide nitrogen a partial positive charge, setting up a small electric dipole.

Virtually all peptide bonds in proteins occur in this trans configuration; an exception is noted in Figure 4–8b.

(a)



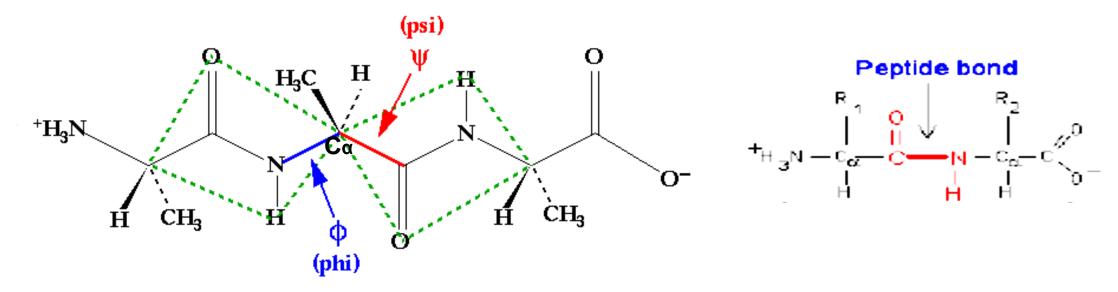
3. All peptide bonds are Trans in nature. Because Trans Bonds are 2 kcal more stable, with an exception of X-Proline peptide bond (Where X is any amino acid behind proline).



4. Rotation around peptide bond (C-N) is not possible (because of partial double bond character). Only the $C\alpha$ -N and $C\alpha$ -C bonds rotation are possible.

The rotational (dihedral) angle between $C\alpha$ -N is called as Φ , while Dihedral angle between $C\alpha$ -C is called as Ψ .

The dihedral Angle between $C\alpha$ -R is called χ and N-C is called ω .



 C_a -C=1.53 Å; N- C_a =1.46 Å; C=O=1.24 Å; C-N (Rigid) = 1.32 Å;

Alanine tripeptide: distance from residue i to $(i+1) = C_{\alpha} - C_{\alpha} = 3.8 \text{ Å}$ A fully extended polypeptide chain has $\Phi = \Psi = 180^{\circ}$