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AMINO ACIDS AND PROTEINS

Concept Covered

- Amino Acids
- ✤ D-/ L- Isomerism
- Classification of Amino Acids
- Properties of Amino Acids
- Peptide/Protein
- Structure of Peptide/ Protein

Amino Acids

The building blocks of peptides and proteins

- Amino acids are monomers of proteins Each amino acid contains
- $\boldsymbol{\diamondsuit}$ a central carbon atom referred to as the Ca(C-alpha) atom
- * an amino group (-NH2)
 * a carboxyl group (-COOH)
 * a side chain (-R)
 * a single hydrogen atom

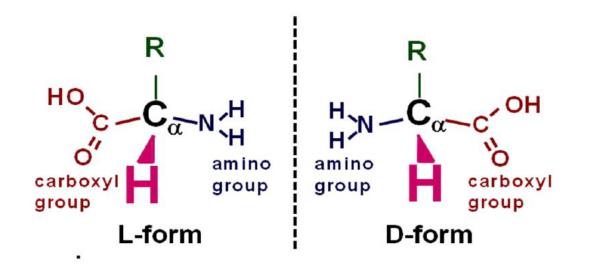
 $H_3N - C - COO^{-1}$

In the physiological pH range, both the carboxylic acid and the amino groups of -amino acids are completely ionized

Zwitterionic form of the α -amino acids

- All Amino Acids are Chiral Compound
- A chiral molecule is a type of molecule that has a non-superposable mirror image. The presence of an asymmetric carbon atom is normally the reason a molecule is chiral and this property is also referred to as asymmetry.
- only one amino acid Gly is achiral meaning that the four atoms attached to the central Cα are not all different
- The R-group represents what is called a side chain which varies from one amino acid to another.
- This R-group is what makes each amino acid unique in nature.
- This group of atoms may have just carbon and hydrogen or in addition to these atoms may contain heteroatoms such as nitrogen, oxygen and/or sulfur.
- This specific combination of atoms will give rise to a particular side chain with specific properties that will define the characteristics of the amino acid.

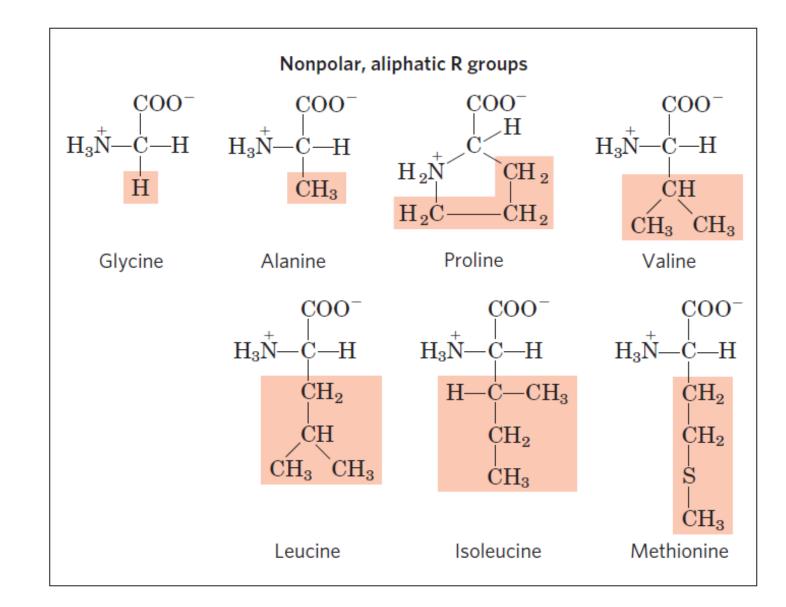
- ✤ An important feature of amino acid structure is the stereochemistry.
- Each amino acid have two enantiomers of the , L- amino acid and D- amino acid
 e.g., L-alanine and D-alanine. They are also called optical isomers.
- Each of these molecules are mirror images of one another.
- The L-form is the isomer mostly seen in organisms.
- D-amino acids are not found in proteins naturally
- ✤ D- amino acids are found in bacterial cell walls



- The normal practice by chemists is to use what is called an R, S system to distinguish between enatiomers of molecules. However, many biochemical systems – mostly amino acids and sugars – use the D and L system based on the earliest method for identifying enantiomers from glyceraldehyde.
- The glyceraldehyde molecule contains three carbon atoms with one form referred to as D and the other as L based on their ability to rotate polarized light either to the right (dexter) or left (laevus).
- The enantiomers of the amino acids are designated on the basis of their similarity to the D and L forms of glyceraldehyde.

Amino Acids Can Be Classified by R Group

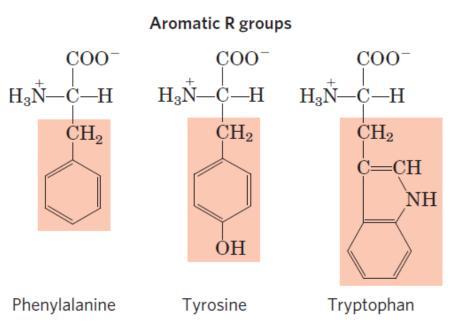
- Nonpolar, Aliphatic R Groups
 e.g. alanine, valine, leucine, isoleucine, Glycine, Methionine, Proline
- Methionine, one of the two sulfurcontaining amino acids, has a slightly nonpolar thioether group in its side chain.
- Proline has an aliphatic side chain with a distinctive cyclic structure. The secondary amino (imino) group of proline residues is held in a rigid conformation that reduces the structural flexibility of polypeptide regions containing proline

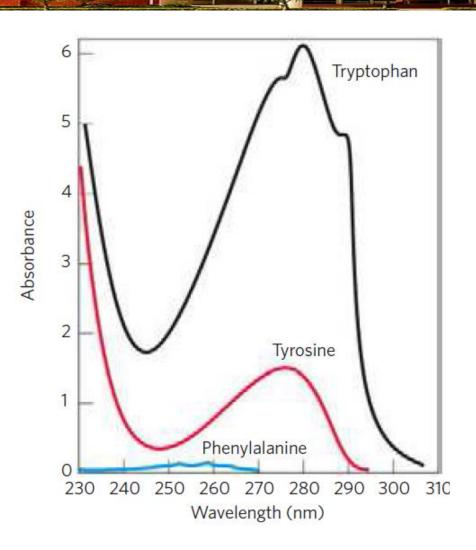


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Aromatic R Groups

- Phenylalanine, tyrosine, and tryptophan, are relatively nonpolar (hydrophobic).
- All can participate in hydrophobic interactions.
- The hydroxyl group of tyrosine can form hydrogen bonds, and it is an important functional group in some enzymes.
- Tyrosine and tryptophan are significantly more polar than phenylalanine, because of the tyrosine hydroxyl group and the nitrogen of the tryptophan indole ring.
- Tryptophan and tyrosine, and to a much lesser extent phenylalanine, absorb ultraviolet light





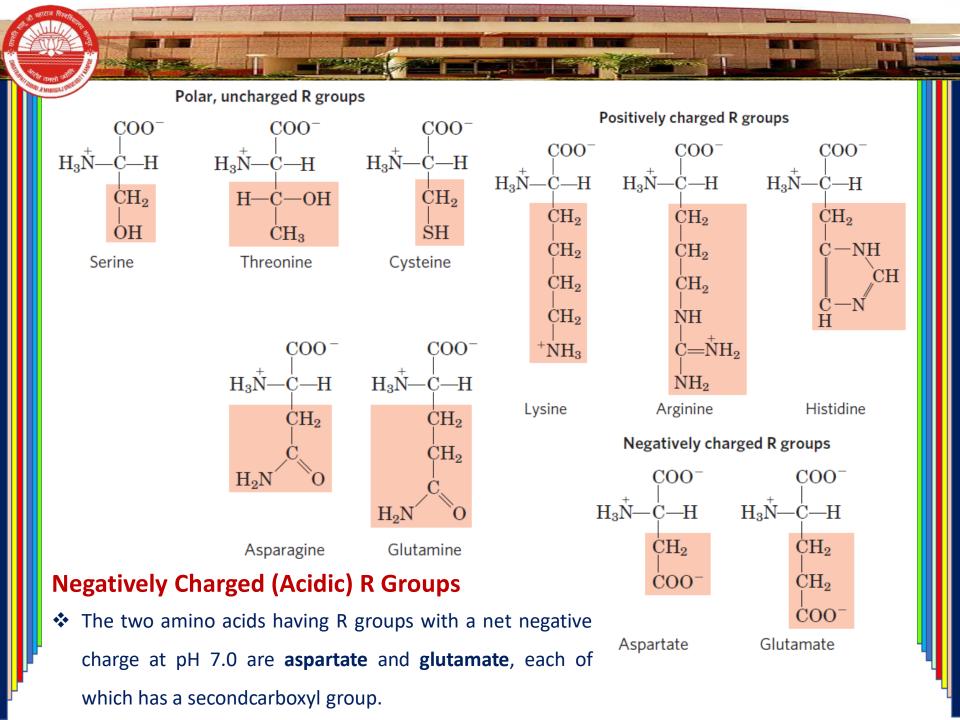
LINE

N D

2

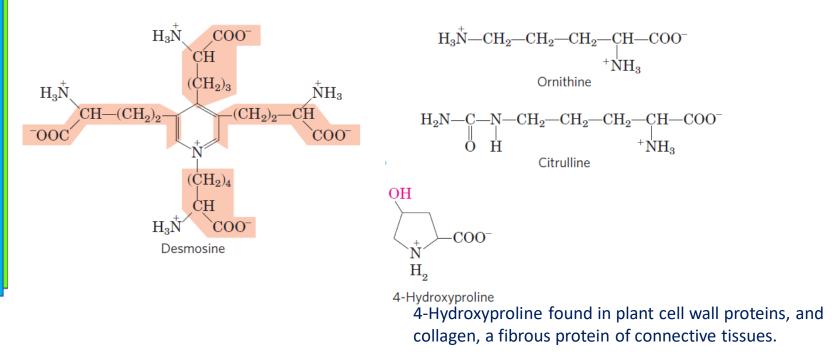
Polar, Uncharged R Groups

- The R groups of these amino acids are more soluble in water, or more hydrophilic, than those of the nonpolar amino acids, because they contain functional groups that form hydrogen bonds with water. This class of amino acids includes serine, threonine, cysteine, asparagine, and glutamine.
- The polarity of serine and threonine is contributed by their hydroxyl groups, and that of asparagine and glutamine by their amide groups.
- Cysteine is readily oxidized to form a covalently linked dimeric amino acid called cystine, in which two cysteine molecules or residues are joined by a disulfide
 Positively Charged (Basic) R Groups
- * The amino acids in which the R groups have significant positive charge at pH 7.0 are lysine, which has a second primary amino group at the ε position on its aliphatic chain; arginine, which has a positively charged guanidinium group; and histidine, which has an aromatic imidazole group.



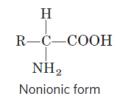
Uncommon amino acids.

- Some uncommon amino acids found in proteins. All are derived from common amino acids.
- Desmosine is formed from four Lys residues
- Ornithine and citrulline, which are not found in proteins, are intermediates in the biosynthesis of arginine and in the urea cycle.



Amino Acids Have Characteristic Titration Curves

- Acid-base titration involves the gradual addition or removal of protons
- The two ionizable groups of glycine, the carboxyl group and the amino group, are titrated with a strong base such as NaOH.
- The plot has two distinct stages, corresponding to deprotonation of two different groups on glycine.
- Each of the two stages resembles in shape the titration



Η $R - C - COO^{-}$ $+NH_{2}$

Zwitterionic form

$$\begin{array}{ccc} H & H \\ | \\ R - C - COO^{-} \rightleftharpoons R - C - COO^{-} + H^{+} \\ + NH_{3} & NH_{2} \\ \\ Zwitterion \\ as acid \end{array}$$

$$\begin{array}{ccc} H & H \\ H & H \\ R - C - COO^{-} + H^{+} \rightleftharpoons R - C - COOH \\ + NH_{3} & + NH_{3} \\ Zwitterion \\ as base \end{array}$$

- Titration of an amino acid. Shown here is the titration curve of 0.1 M glycine at 25°C. The ionic species predominating at key points in the titration are shown above the graph.
- The shaded boxes, centered at about pK1 5 2.34 and pK2 5 9.60, indicate the regions of greatest buffering power. Note that 1 equivalent of OH 5 0.1 M NaOH added.

