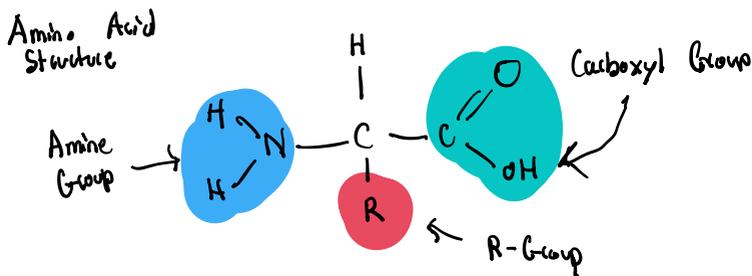


# Biochemistry

Proteins are organized in 4 levels of structure, which differ in their size/scale

## #1: Primary Structure (Smallest)

- Composed of amino acids, which are the building blocks of proteins, held together by peptide bonds
- Dictates the other 3 levels of structure



R-Groups, or Side Chains, determine an amino acid's properties; they are what make amino acids distinctive from one another and vary

### Beta-Branched Amino Acids

Includes!

- Leucine
- Isoleucine
- Valine (all nonpolar)

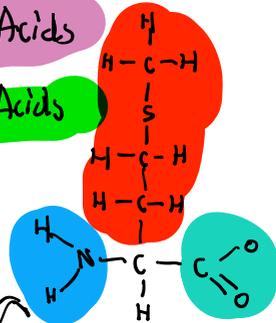
Contain long hydrocarbon branches, leading to the formation of Beta-Sheets

### Basic Amino Acids

- Lysine (Lys)
- Arginine (Arg)
- Histidine (His)

### Nonpolar / Hydrophobic Amino Acids

- Glycine (Gly)
- Alanine (Ala)
- Valine (Val)
- Leucine (Leu)
- Isoleucine (Ile)
- Methionine (Met)
- Phenylalanine (Phe)
- Tryptophan (Trp)
- Proline (Pro)



Above! Structure of Methionine, which is the start codon in protein synthesis (AUG)

### Aromatic Amino Acids

Includes!

- Phenylalanine
- Tyrosine
- Tryptophan

Contains stable six-carbon benzene rings that stabilize the molecule

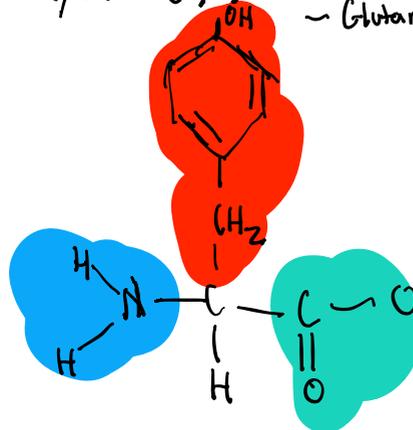


### Acidic Amino Acids

- Aspartic Acid
- Glutamic Acid

### Polar Side Chains / Hydrophilic

- Serine (Ser)
- Threonine (Thr)
- Cysteine (Cys)
- Tyrosine (Tyr)
- Asparagine (Asn)
- Glutamine (Gln)



Every amino acid chain ends at a carboxyl and amino group. They are referred to as a C-Terminus and N-Terminus respectively

# #2: Secondary Structure (2<sup>nd</sup> Smallest)

- Arises from interactions between amine and Carboxyl groups

## $\alpha$ - Helices

→ A secondary structure that arises from Hydrogen Bonding

R-Groups don't play a major role in the formation of alpha-helices, but some can break down helix structures, including

- Proline
- Glycine

## $\beta$ - Sheets

→ Another major type of secondary structure

- $\beta$ -pleated sheets form from Hydrogen bonds between parallel parts of a polypeptide chain
- Leads to stability in the molecule

Example:



# #3: Tertiary Structure

↳ stems from interactions between R-groups of amino acids

- Can coincide w/ 2<sup>nd</sup> Structure, they depend on different interactions

Hydrophobic Interactions → Caused by nonpolar/polar R-groups

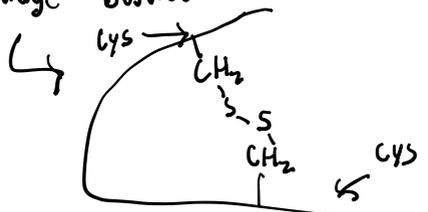
- Polar amino acids → Hydrophobic
- Nonpolar amino acids → Hydrophobic

IMFs between R-Groups determine protein structure

R-Group interactions also include:

- **Ionic Bonding**: Depends on the environmental pH, which determines whether amino acids undergo an acid-base reaction

- **Covalent Bonding**: The Cysteine (polar) amino acid interacts with each other, forming a disulfide bridge between its R-Groups



Amino Acids can undergo chemical modification through phosphorylation, including these 3...

- Serine
- Threonine
- Tyrosine (tyrosine kinase)

#4:

## Quaternary Structure

↳ Arises from interactions between multiple polypeptides  
→ Some proteins DON'T have a quaternary structure

One protein that makes up stronger tissue is collagen → together in a triple helix

Another one is hemoglobin → transports Oxygen in red blood cells  
↳ two  $\alpha$  and two  $\beta$  subunits