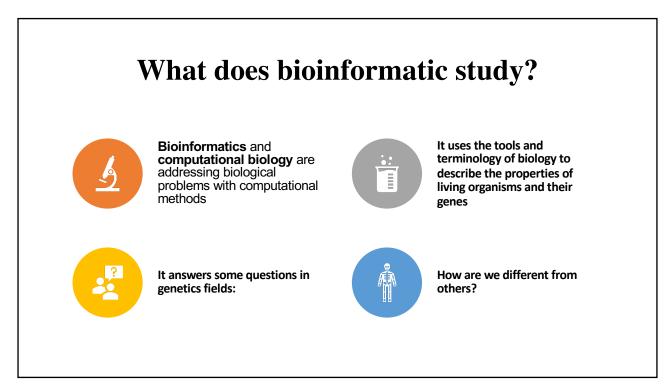
Introduction to Bioinformatics I

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Bioinformatics:

Genetics:

- DNA and RNA analysis and Structure.
- Includes techniques.
- PCR, Cloning and mutagenesis.
- Statistics of different variants of species

Proteins:

- **Expression and purification of protein.**
- □ Structural and functional of proteins.
- □ Chemistry biology of the protein.
- □ Biophysics.
- NMR, X-ray Crystallography, Electron microscopy and enzymes and others' kinetics
- □ Interactions and affinities.
- □ Drug design and discovery

Organism Organ Cell What we do Live Organelle consist of ? Hepatocyte What are the small Mitochondrion lolecules Human molecules that Citrate synthase code us? DN Ubiquinon

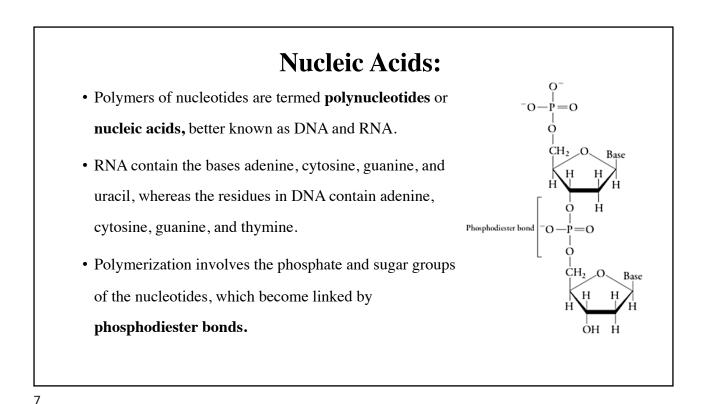
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The Storage of Genetic Information: What is the meaning of the terms?

- Bioinformatics are driven by by simple information that are built up from the sequences of DNA, RNA and proteins (which are the **Storage of Genetic Information**)
- DNA consists of four nucleotides that store genetic information.
- The base sequence encodes the necessary information to generate proteins.
- The entirety of genomic DNA in any organism is known as a genome.
- The total pool of mRNA in any organism is referred to as a transcriptome.
- The entire pool of proteins in any organism is referred to as the proteome.
- A genome comprises genes that contain the information to build proteins.
- Genome is the entire DNA sequence of an organism.

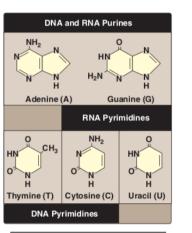
What are Macro (proteins) and micro (DNA)biomolecules ??

- DNA and RNA ?? Which are polymers of nucleotides, each of which consists of a purine or pyrimidine base, deoxyribose or ribose, and phosphate.
- Amino acids ??? Are coded by nucleotides>>>
- Proteins ?? Poly amino acids
- Note: Micro- is a prefix which may be applied to word when describing something that is small scale. Macro- is a prefix which means large.



Purine or pyrimidine base

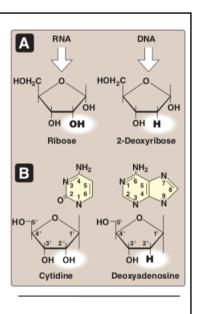
- Nucleotides are composed of a nitrogenous base, a pentose monosaccharide, and one, two, or three phosphate groups.
- The nitrogen- containing bases belong to two families of compounds: the purines and the pyrimidines.
- Both DNA and RNA contain the same purine bases: adenine (A) and guanine (G). Both DNA and RNA contain the pyrimidine cytosine (C), but they differ in their second pyrimidine base: DNA contains thymine (T), whereas RNA contains uracil (U). T and U differ in that only T has a methyl group



Purines and pyrimidines commonly found in DNA and RNA.

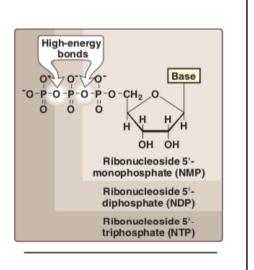
Nucleosides

- The addition of a pentose sugar to a base produces a nucleoside.
- If the sugar is ribose, a ribonucleotide is produced; if the sugar is 2-deoxyribose, a deoxyribonucleoside is produced
- Nucleotides
- The addition of one or more phosphate groups to a nucleoside produces a nucleotide.

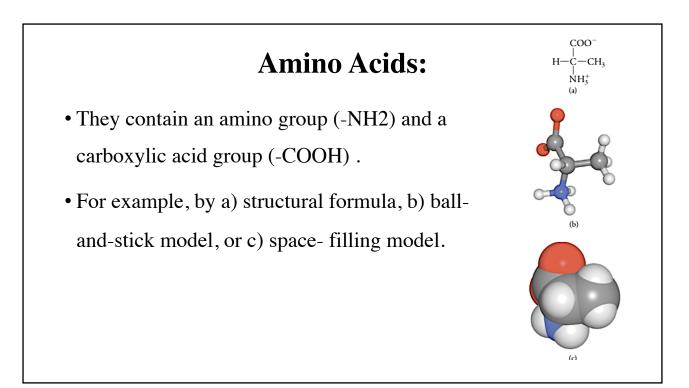


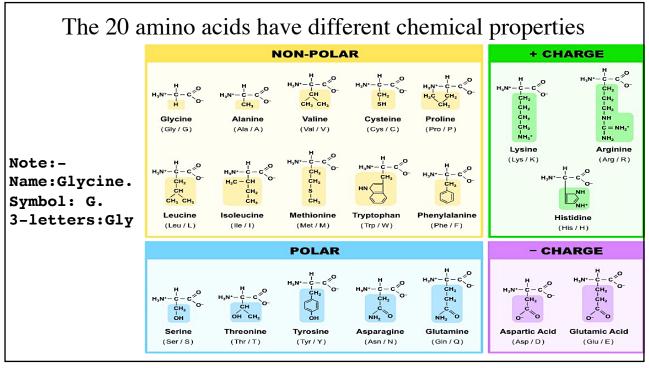
A. Pentoses found in nucleic acids. B. Examples of the numbering systems for purine- and pyrimidinecontaining nucleosides.

- The first phosphate group is attached by an ester linkage to the 5'-OH of the pentose.
 Such a compound is called a nucleoside 5'-phosphate or a 5'-nucleotide.
- The second and third phosphates are each connected to the nucleotide by a "high-energy" bond.



Ribonucleoside monophosphate, diphosphate, and triphosphate.



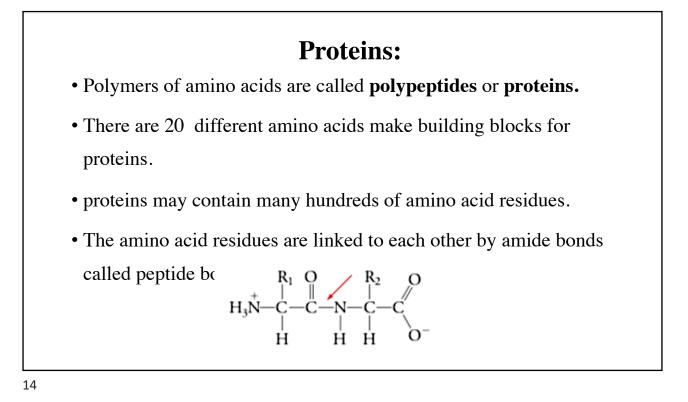


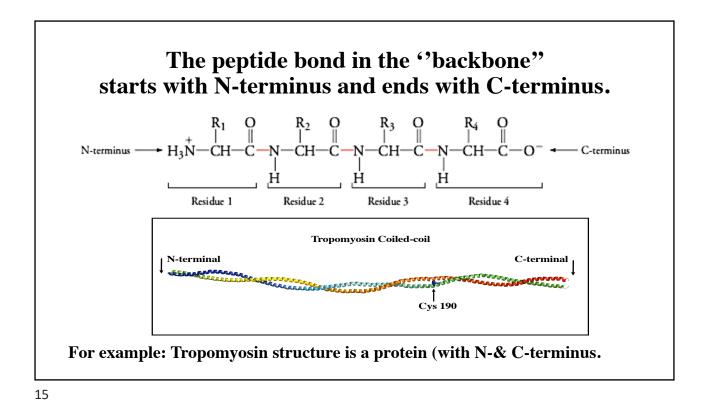
- Amino Acids with Hydrophobic Side Chains: Sidechains interact very weakly or not with water. e.g. alanine (Ala).
- Amino Acids with Polar Side Chains: The sides interact with water because they contain hydrogen-bonding groups. E.g. Serine (Ser). Cysteine (Cys) has a thiol group: can form a disulphide bond:

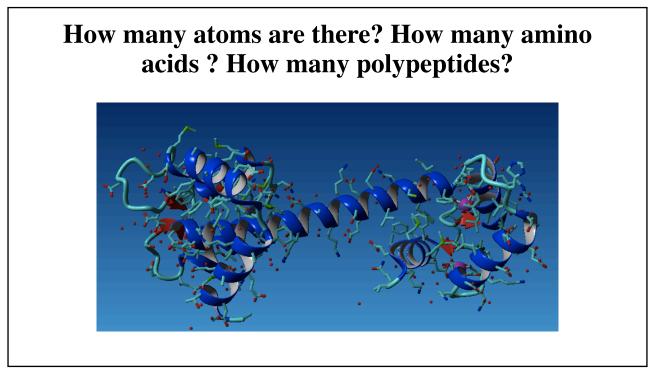
 $\begin{array}{cccc} COO^- & H^+ & COO^- & COO^- & COO^- & COO^- & H^+ & COO^- & H^+ & H^- & H^$

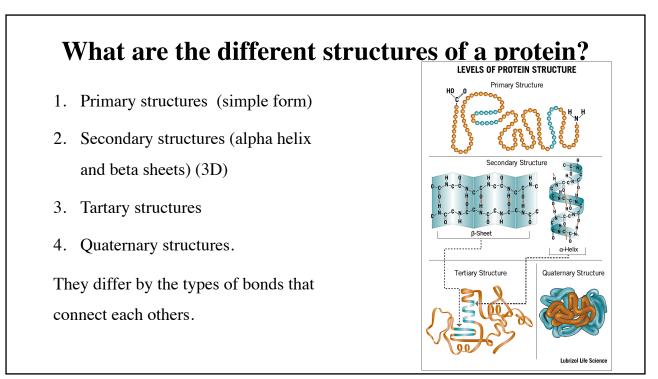
- Amino Acids with Charged Side Chains (4 amino acids only): Side chains are always charged under physiological conditions.
- (Asp) and (Glu), are acidic (COOH)
- (Lys) and (Arg) are positively charged (NH₃)

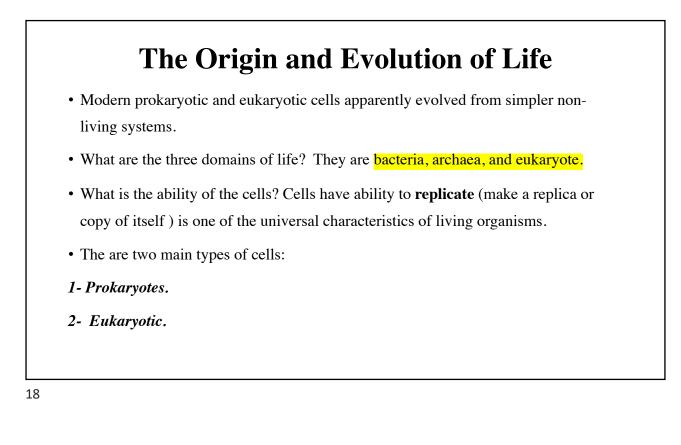




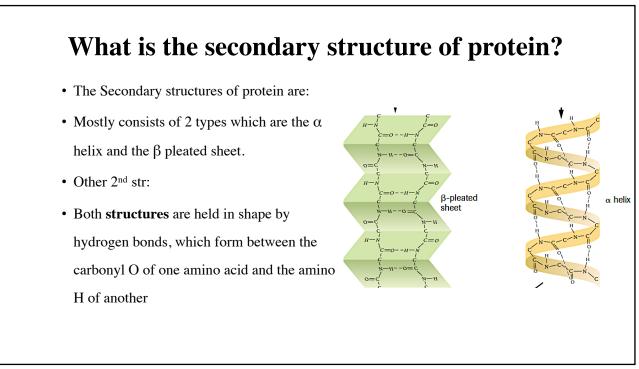






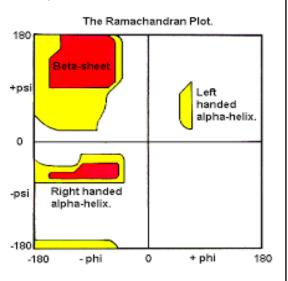


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The chain conformation of a polypeptide can be determined by?

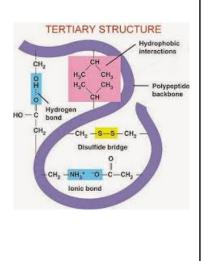
- It can be determined by the torsion angles around the Cα–N binding (φ) and the Cα–C binding (ψ) of the constituent amino acid residues.
- A Ramachandran plot is a conformation chart of those values that are sterically possible for φ and ψ.
- It determines the alpha helices and the beta sheets contents of a protein.



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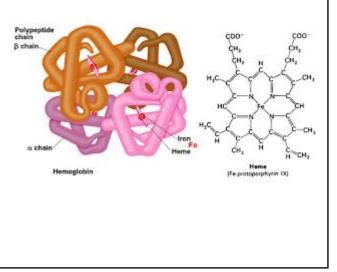
What are the tertiary & quaternary structures of protein?

- Tertiary: It is 3D structures shape of protein.
- It has a single polypeptide chain "backbone" with one or more protein secondary structures that form the protein domain.
- Bonds?



What is the Quaternary structure of proteins?

- Quaternary:
- It is an arrangement of multiple folded protein subunits in a multisubunit complex.
- It involves at least 2 polypeptides (domains).
- It can be a dimer, tetramer, homo or hetero protein.



Next lecture,

- We will continue explaining biomolecules, the bases bioinformatics study....
- We will talk about DNA and RNA molecules...