BINF 731

Protein Structure Analysis

http://binf.gmu.edu/vaisman/binf731/

Iosif Vaisman

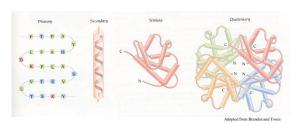
2023

Secondary Structure: Computational Problems

Secondary structure characterization

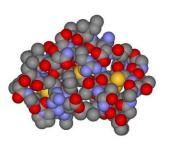
Secondary structure assignment Secondary structure prediction Protein structure classification

Protein Structure Hierarchy

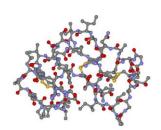


- •Primary the sequence of amino acid residues
- •Secondary ordered regions of primary sequence (helices, beta-sheets, turns)
- •Tertiary the three-dimensional fold of a protein subunit
- •Quaternary the arrangement of subunits in oligomers.

Protein representations



Protein representations

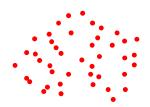


Protein representations



Protein representations

Protein representations





Protein representations

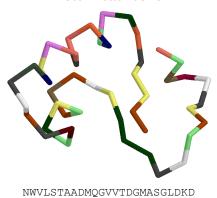
Protein representations

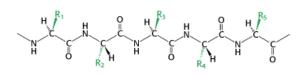


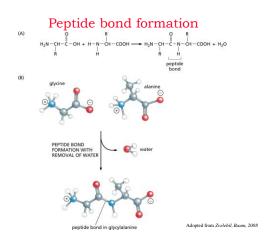


Protein backbone

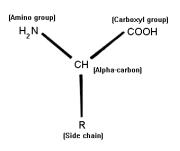
Protein backbone



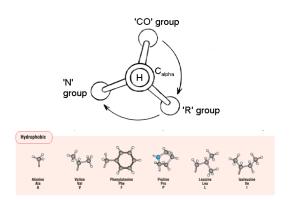




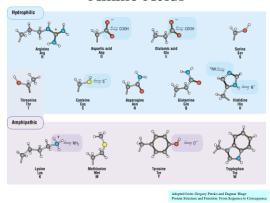
Amino Acid Residue



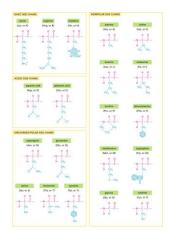
Amino Acid Residue



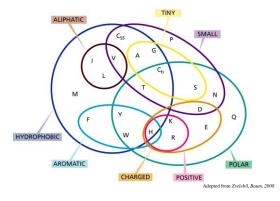
Amino Acids



Amino Acids

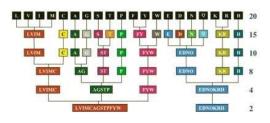


Amino Acid Residue Clustering

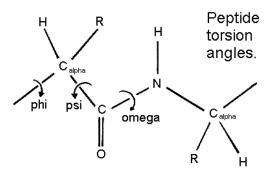


Adopted from Zvelebil, Baum, 2008

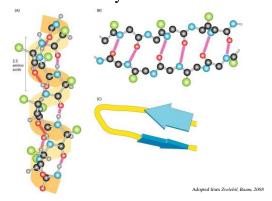
Amino Acid Residue Clustering



Adopted from: L.R.Murphy et al., 2000



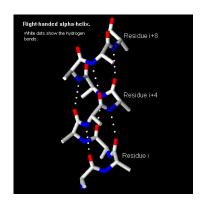
Secondary Structures



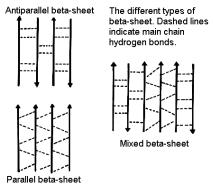
Secondary Structure (Helices)

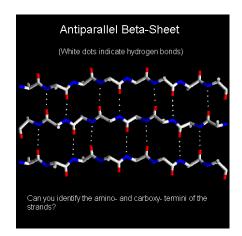


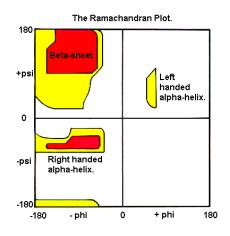
Helix

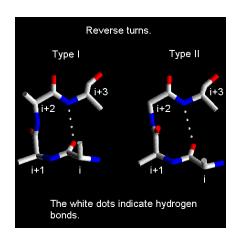


Secondary Structure (Beta-sheets)

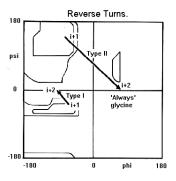


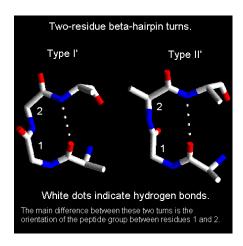




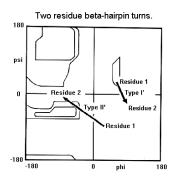


Reverse Turns on a Ramachandran Plot





Beta-hairpin Turns on a Ramachandran Plot



Side-Chain Atom Nomenclature

C delta C gamma C alpha L alpha L alpha C alpha C two amino acid side chains to indicate the atom naming convention. C delta C psilon1 C delta C delta C pamma C delta C psilon1 C psilon1 C psilon2 C psilon1 C psilon

Side-Chain Torsional Angles