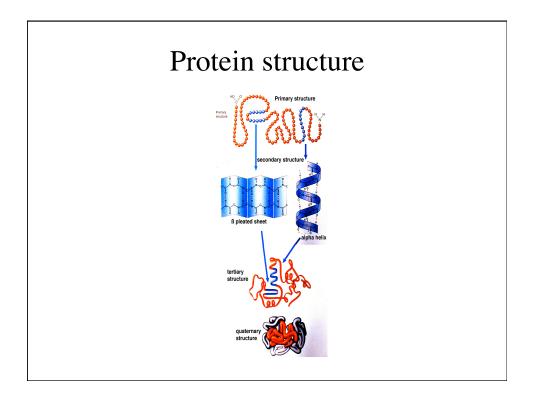
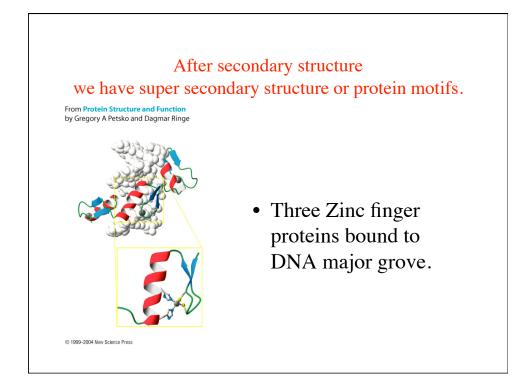
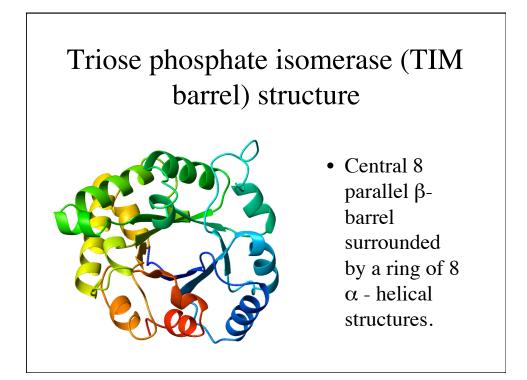
Tertiary and Quaternary Structure of Proteins

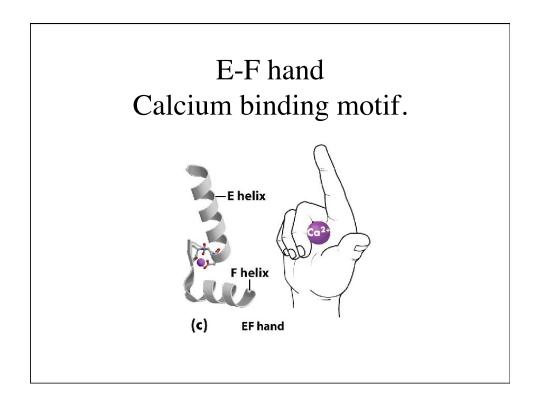
Manickam Sugumaran

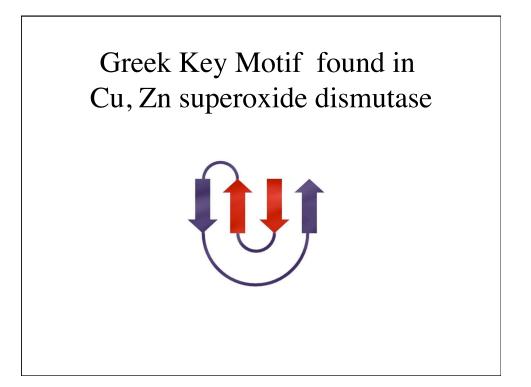
Department of Biology University of Massachusetts Boston, MA 02125





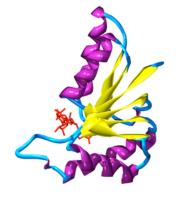






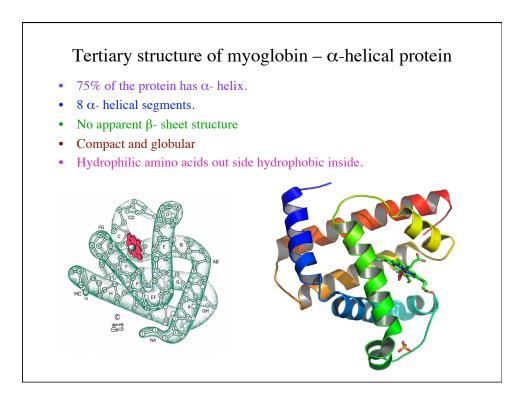
NAD binding fold in lactate dehydrogenase

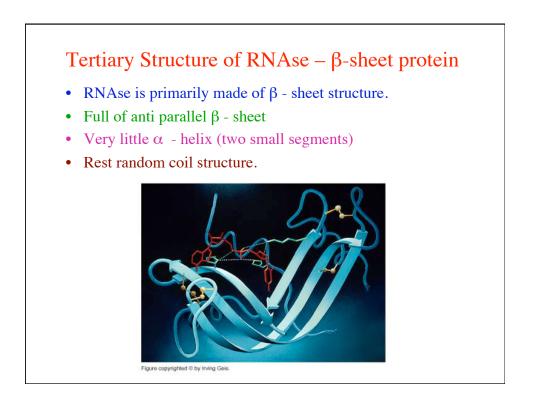
• Red molecule is NAD that is bound to lactate dehydrogenase



Tertiary structure Tertiary structure deals with the spatial relationship of amino acids that are present far away from each other

- Gives a protein its biological activity.
- Proteins having entirely different amino acid sequence can still have the same tertiary structure. Example myoglobin & hemoglobin.
- Proteins have very similar amino acid structure may have the same tertiary structure also.





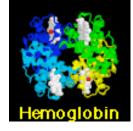
Quaternary structure

- Few protein have more than one protein chains in them that are not usually connected.
- Example Hemoglobin. $\alpha_2\beta_2$ subunits can be easily dissociated.
- Quaternary structure usually ascribes regulatory properties to a protein.

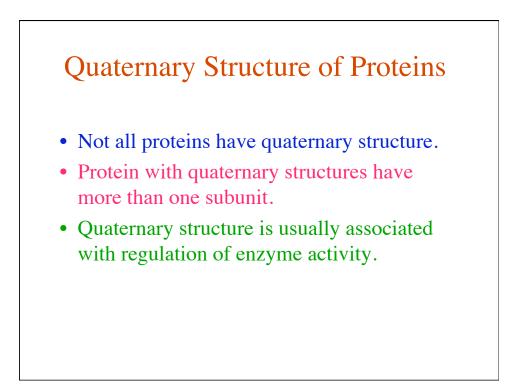
Myoglobin and hemoglobin

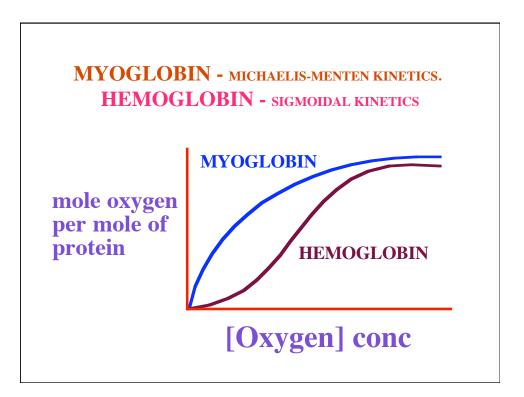


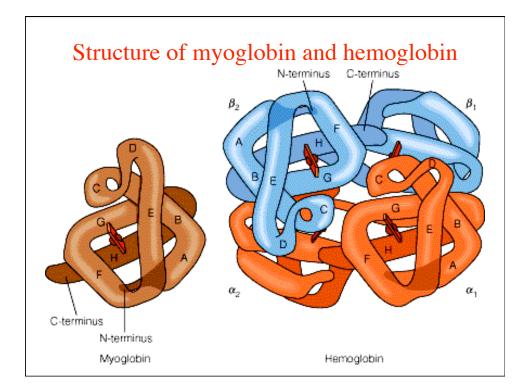
Monomer M.wt 17,000 kDa Single polypeptide No subunit structure



Tetramer M.Wt 64,000 kDa Four polypeptide Two kinds of subunit $(\alpha_2\beta_2)$



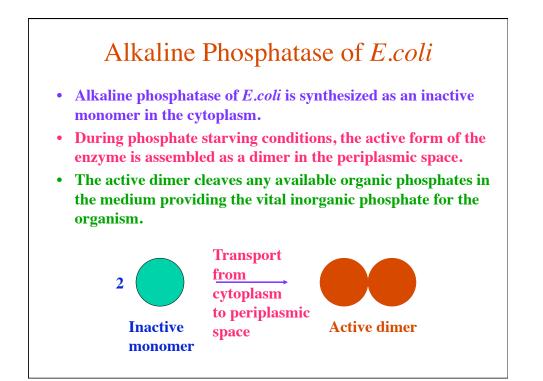


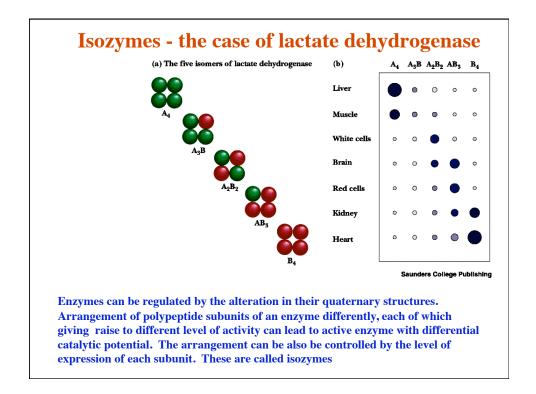


Differences between myoglobin and hemoglobinMyoglobinHemoglobin

- Storage protein
- Monomeric in nature.
- No subunit structure
- M.M. Kinetics.
- pH has no drastic effect.
- CO₂ No effect.
- Diphosphoglycerate No effect
- Only one form.
- NO has no effect

- Transport protein
- Tetrameric in nature
- Two kind of subunits, $\alpha_2\beta_2$.
- Sigmoidal kinetics.
- pH inhibits O₂ binding.
- CO₂ inhibits O₂ binding.
- Diphosphoglycerate inhibits O₂ binding.
- Exists in two forms
- Nitric oxide binding





References

- L. Jia, C. Bonaventura, J. Bonaventura and J. S. Stamler. S-nitroso hemoglobin: a dynamic activity of blood involved in vascular control. Nature. 380, 221-226 (1996).
- Jane. S. Richardson. Introduction: Protein motifs. The FASEB Journal 8, 1237-1239 (1994).