

Structural Effects in Proteins

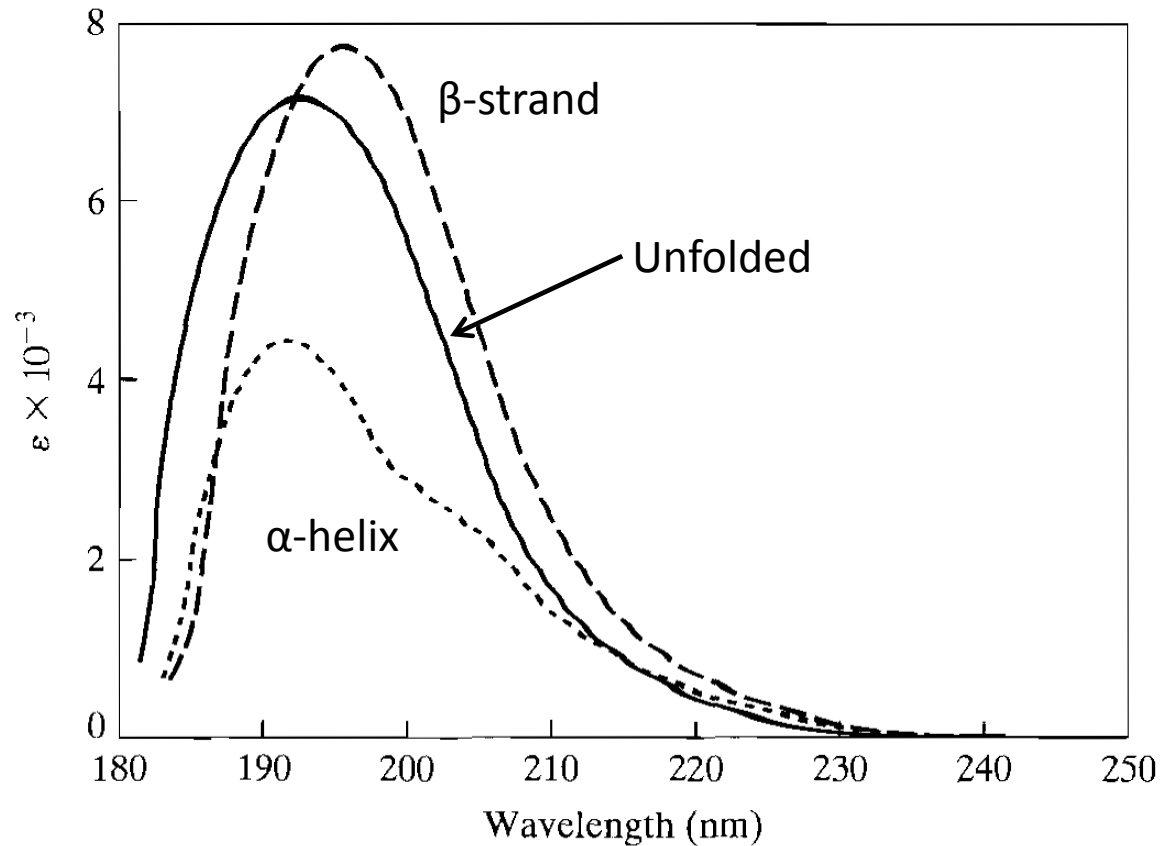


Figure 9.16 The electronic absorption spectra for poly-L-lysine hydrochloride in aqueous solution as a random coil at pH 6.0, 25°C (—); α helix at pH 10.8, 25°C (---); β strand at pH 10.8, 52°C (-.-). [Adapted from K. Rosenheck and P. Doty (1961), *Proc. Natl. Acad. Sci. USA* **47**, 1775–1785.]

Structural Effects in DNA

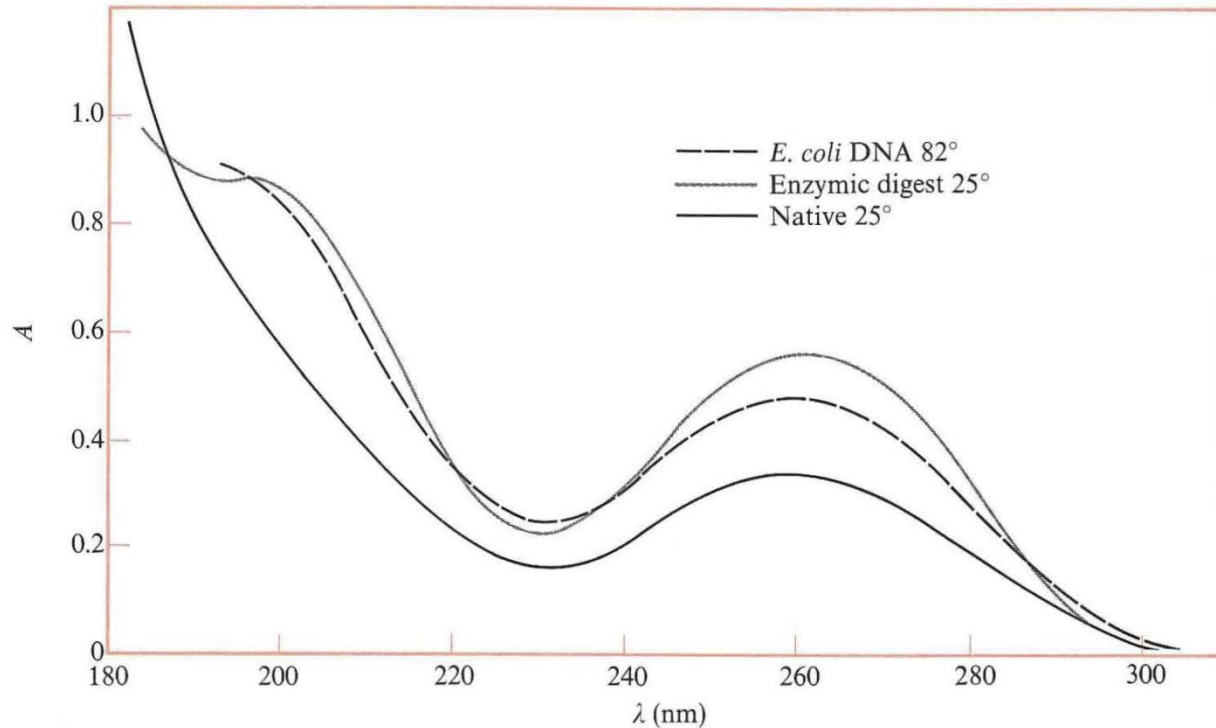
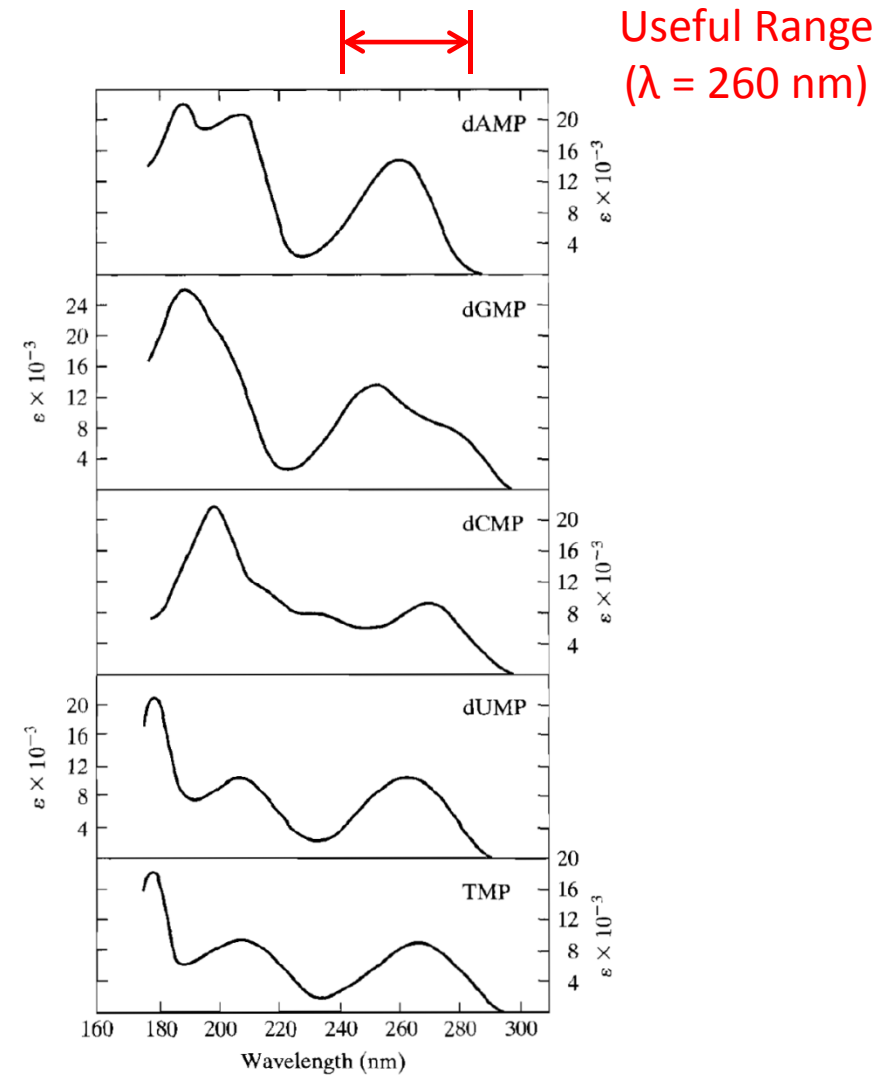
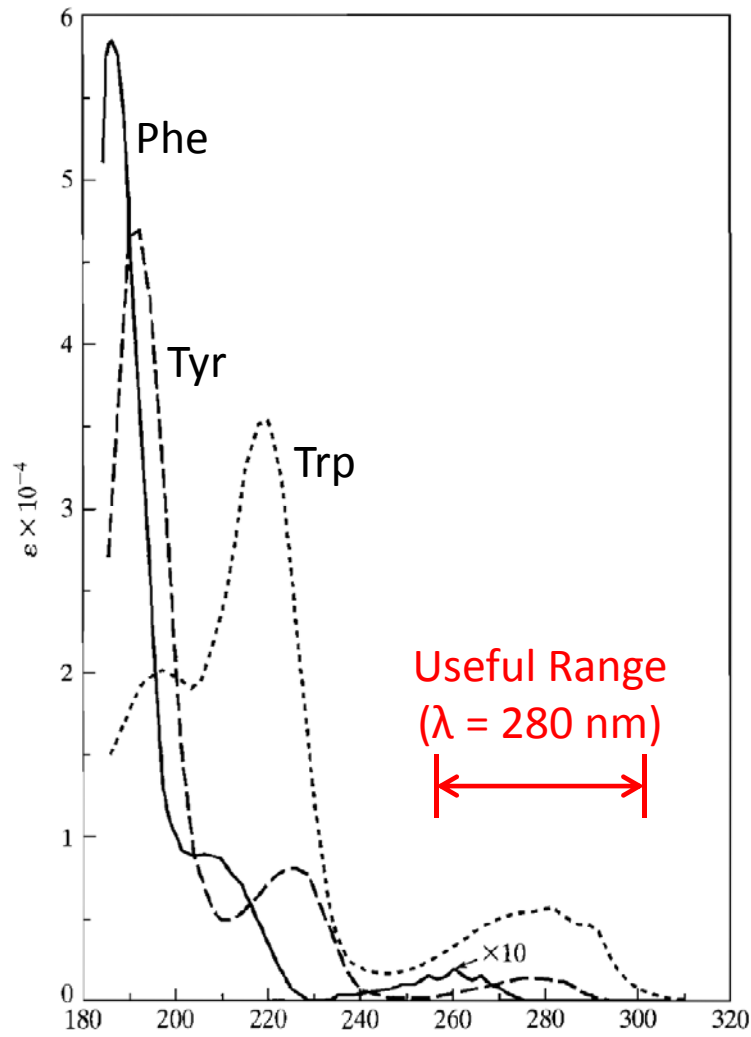


Figure 7-14

*Spectrum of DNA as a function of temperature. The enzymic digest should result in a solution of only mononucleotides and possibly a few short oligomers. Note that the high-temperature spectrum is fairly close to the digested spectrum. [After D. Voet, W. B. Gratzer, R. A. Cox, and P. Doty, *Biopolymers* 1:193 (1963).]*

Concentration of DNA and Protein



Websites for Estimating ϵ

- Proteins
 - ProtParam
<http://web.expasy.org/protparam/>
- DNA (Double and Single Stranded)
 - IDT Biophysics
<http://biophysics.idtdna.com/UVSpectrum.html>
 - Crude ϵ for ds plasmid DNA: $0.02 \left(\frac{\mu\text{g}}{\text{mL}}\right)^{-1} \text{cm}^{-1}$
- RNA
 - Harder to estimate, because of structure, but see
<http://www.scripps.edu/california/research/dna-protein-research/forms/biopolymercalc2.html>

Solvent Effects

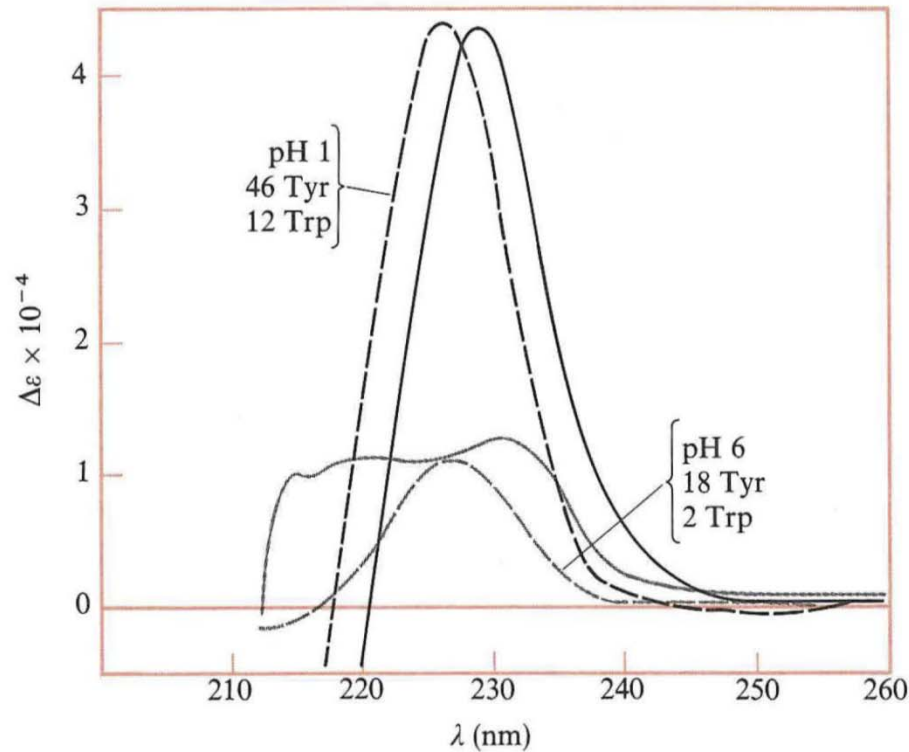


Figure 7-16

Solvent-perturbation difference spectrum, produced by treating rabbit-muscle aldolase with 20% ethylene glycol. The observed spectra (*solid curves*) are compared with calculated difference spectra (*dashed curves*) generated using data obtained on individual amino acids. Clearly, the protein has many fewer exposed aromatic groups at pH 6 than it does at pH 1. [After J. W. Donovan, *J. Biol. Chem.* 244:1961 (1969).]

IR Spectroscopy

Figure 7-1

Energy levels of a small molecule. Selected rotational sublevels of the vibrational levels of each of two electronic states are shown. Transitions corresponding to electronic (e), vibrational (v), and rotational (r) spectra are indicated.

