

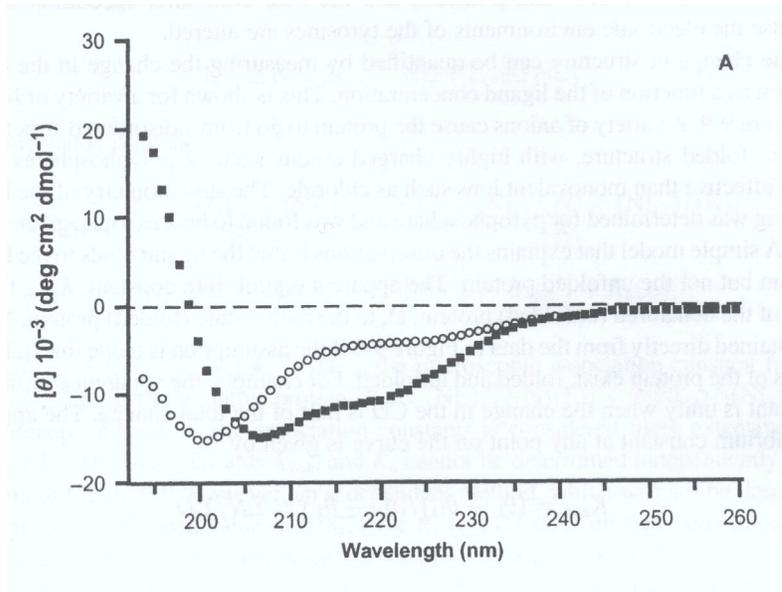
1) A crystal mounted on a diffractometer 150mm away from the detector is exposed to X-rays of wavelength 1.54Å. A reflection is located at 15.4mm from the location of the direct beam. Assume  $n=1$ . **A)** What is the lattice spacing along this axis? **B)** Is this spacing compatible with a large protein that has a radius of 30Å? Explain your reasoning.

$\tan(2\theta)=15.4/150$  to get  $\theta$

Bragg's law to  $2d\sin\theta=n\lambda$  to determine  $d$

$d=15\text{Å}$  so a protein with radius of 30Å will not fit.

2) Circular dichroism of a protein measured in the presence (black squares) and absence (open circles) of a small compound is shown below. Explain what is happening upon binding of the small compound. NOTE: the small compound is not optically active



The protein undergoes a transition from mostly random coil to alpha helical upon binding of a small molecule.

3A) Describe the lattice type and symmetry axes for each of the following

a)  $P2_1$  Primitive lattice (molecules on every corner) with two fold screw axis along Y and translations along X and Z

b)  $I422$  Body centered lattice (molecules on every corner and in the middle of the Unit cell) with a four-fold axis on X and two-fold axes along Y and Z

c)  $C222$  C center lattice (molecules on every corner and in the middle of two opposite sides) with two fold axes along X, Y, and Z

3B) Give the short abbreviations for the following types of lattices

d) A lattice with molecules on every corner and every face with two fold axes of rotation on every axis.  $F222$

e) A lattice that has molecules ONLY at every corner, 4 fold screw axis with a  $\frac{3}{4}$  unit cell translation along X, a two fold screw axis along Y, and a two fold axis along Z.  $P4_32_12$