

Principles Of Protein X Ray Crystallography

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Abstract In Chapter 1 you learned how crystals of a protein can be grown and you observed a diffraction pattern. The crystalline form of a protein is required to determine the protein's structure...

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The crystals of a pure protein are exposed to X-ray beam and X-ray is diffracted by atoms present in a protein crystal. Depending on the organization of atoms within a crystal and the number of electrons in the atoms, an X-ray beam is diffracted into many specific directions.

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x-ray Crystallography of Proteins The crystals that form are frozen in liquid nitrogen and taken to the synchrotron which is a highly powered tunable x-ray source. They are mounted on a goniometer and hit with a beam of x-rays. Data is collected as the crystal is rotated through a series of angles.

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