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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.

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crystals that form are frozen in liquid

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Crystallography  
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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