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X-ray crystallography is used to determine the structure of large biomolecules such as proteins. such as proteins, do not occur naturally as crystals.

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Protein X-ray crystallography is a technique The three components needed to complete an X-ray crystallography analysis are a protein crystal, a source of x-rays

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Abstract

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X-ray Crystallography. Most of the structures included in the PDB archive were determined using X-ray crystallography. For this method, the protein is purified and

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