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Protein Crystallography

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closed (1 July 2020)

Message from the Guest Editors

Dear Colleagues,

Protein X-ray crystallography has played dominant roles and will continue to contribute greatly to structural biology despite the recent technical revolutions in cryo-EM and XFEL (X-ray free electron laser). Structural biology dissects bio-macromolecules and their complexes at the atomic resolution, thus giving the best mechanistic connections and understanding between physiochemical structures and biological phenomena. Structural biology does not only deal with the well-ordered bio-macromolecules, but also studies flexible and disordered proteins, and phase separation mechanisms caused by some of the disordered proteins. This Special Issue of "Protein Crystallography" will cover all aspects of structural biology relevant to X-ray and electron crystallography.

Keywords

- Protein preparation and crystallization
- X-ray and electron diffraction
- Structural determination and analyses
- Structural biology
- Rational drug design
- Bio-macromolecule dynamics and interactions
- Bio-macromolecule design
- Disordered proteins and phase separation

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Editor-in-Chief

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Message from the Editor-in-Chief

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