



Application of Computational Methods and Biomolecular Structural Modeling to the Investigation of Metalloproteins

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Message from the Guest Editors

The structural bioinformatics of metalloproteins is important for providing links between 3D structure and biological function, with particular focus on the role of metal ions and how this role has evolved and become diversified among organisms. In this Special Issue, we wish to cover the most recent advances in all these aspects of computational methods for the study of metalloproteins by hosting a mix of original research articles and short critical reviews.

Deadline for manuscript
submissions:

closed (31 August 2021)





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Message from the Editor-in-Chief

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