

Special Issue

Functionally Relevant Macromolecular Interactions of Disordered Proteins

Message from the Guest Editor

It is common that most proteins function in folded form. Another significant portion of proteins or protein segments spend a part- or sometimes most- of their time in an unstructured/disordered form. They generally fold only temporarily-typically on the surface of another protein or other macromolecule during their biochemical activity. This phenomenon has been widely studied in the past decade. However, we are expecting a great deal of new information about the functional relevance of this coupled folding and binding for this issue of IJMS.

Guest Editor

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The International Journal of Molecular Sciences (*IJMS*, ISSN 1422-0067) is an open access journal, which was established in 2000. The journal aims to provide a forum for scholarly research on a range of topics, including biochemistry, molecular and cell biology, molecular biophysics, molecular medicine, and all aspects of molecular research in chemistry. *IJMS* publishes both original research and review articles, and regularly publishes special issues to highlight advances at the cutting edge of research. We invite you to read recent articles published in *IJMS* and consider publishing your next paper with us.

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