

Special Issue

Computer Aided Drug Discovery

Message from the Guest Editors

We are pleased to invite you to contribute to the special issue on Computer-Aided Drug Discovery in *Biomolecules*. This collection will highlight advances in MM, MD, AI, vHTS and others to aid CADD. The issue will present the use of CADD in identifying hits and leads, as well as scaffold hopping, to gain an upper hand in terms of intellectual property. The use of CADD for lead optimization, which usually associated with drug development, will also be of interest. Likewise, CADD's application to identification putative adverse consequences over and above those weeded out through the use of algorithms that inform on potential instability, aggregation, and reactivity of compounds will be of interest. Overall, this special issue on CADD will be interested in disseminating any and all aspects of computational approaches and strategies that help discover new drug or drug-like molecules. As an expert working in this area, we welcome your contribution to this Special Issue. You may contribute an original research article or a review. We look forward to receiving your contributions.

Guest Editors

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About the Journal

Message from the Editorial Board

Biomolecules is a multidisciplinary open-access journal that reports on all aspects of research related to biogenic substances, from small molecules to complex polymers. We invite manuscripts of high scientific quality that pertain to the diverse aspects relevant to organic molecules, irrespective of the biological question or methodology. We aim for a competent, fair peer review and rapid publication. Please look at some of the exciting work that has been published in *Biomolecules* so far. We would be delighted to welcome you as one of our authors.

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