

## Special Issue

# Structural and Other Proteomics Approaches in Drug Discovery

### Message from the Guest Editors

Proteins are the primary biological targets for numerous drugs. As such, the relevance of high-throughput structural proteomics and other proteomics-related approaches in the drug discovery is easily comprehensible at different levels including target identification, target validation, potential drug toxicity and protein-protein or protein-ligand interactions. In addition to the classic experimental proteomics techniques (*e.g.*, mass spectrometry), the emergence of new *in silico* methods (*e.g.*, bioinformatics and computational modelling) as well as artificial intelligence (AI) tools can provide valuable insights to improve the current knowledge and contribute to the design of new useful drugs. In this sense, the present Special Issue entitled “Structural and other Proteomics Approaches in Drug Discovery” aims to collect reviews that consider the current state of the art and future prospects in this field, as well as original research articles reflecting. We are looking forward to receiving your valuable contributions.

### Guest Editors

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### Deadline for manuscript submissions

closed (25 February 2025)



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Impact Factor 4.8  
CiteScore 7.7  
Indexed in PubMed



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## About the Journal

### Message from the Editor-in-Chief

Because of your expertise in the field of drug sciences, I kindly invite you to consider publishing your current work, in the form of a research article or a review, in the open access electronic journal *Pharmaceuticals*.

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We hope to handle your contribution to *Pharmaceuticals* soon.

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### Editor-in-Chief

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