

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

Dr. Marino F. A. Santos

1. Associate Laboratory i4HB - Institute for Health and Bioeconomy, NOVA School of Science and Technology, Universidade NOVA de Lisboa, 2819-516 Caparica, Portugal
2. UCIBIO – Applied Molecular Biosciences Unit, Chemistry Department, NOVA School of Science and Technology, Universidade NOVA de Lisboa, 2829-516 Caparica, Portugal

Dr. Luís Passarinha

1. Associate Laboratory i4HB - Institute for Health and Bioeconomy, NOVA School of Science and Technology, Universidade NOVA de Lisboa, 2819-516 Caparica, Portugal
2. UCIBIO–Applied Molecular Biosciences Unit, Chemistry Department, NOVA School of Science and Technology, Universidade NOVA de Lisboa, 2829-516 Caparica, Portugal
3. CICS-UBI–Health Sciences Research Centre, University of Beira Interior, 6201-506 Covilhã, Portugal

Deadline for manuscript submissions

closed (25 February 2025)



Pharmaceuticals

an Open Access Journal
by MDPI

Impact Factor 4.8
CiteScore 7.7
Indexed in PubMed



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Editorial Office
MDPI, Grosspeteranlage 5
4052 Basel, Switzerland
Tel: +41 61 683 77 34
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Editor-in-Chief

Prof. Dr. Amélia Pilar Rauter

Departamento de Química e Bioquímica (DQB) e Centro de Química Estrutural (CQE), Institute of Molecular Sciences, Faculdade de Ciências, Universidade de Lisboa, Lisboa, Portugal

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