



## Clinical Proteomics: Third Edition

Guest Editors:

**Dr. Edwin Lasonder**

Department of Applied Sciences,  
Faculty of Life and Health  
Sciences, Northumbria  
University, Newcastle-Upon-Tyne  
NE1 8ST, UK

**Dr. Vikram Sharma**

School of Biomedical Sciences,  
University of Plymouth, Plymouth  
PL4 8AA, UK

Deadline for manuscript  
submissions:

**closed (30 June 2023)**

### Message from the Guest Editors

Dear Colleagues,

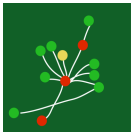
In the post-genome era, ‘omics’ technologies are key to our understanding of human disease because these technologies provide a global approach rather than studying single genes/proteins by traditional biochemical approaches. This Special Issue is dedicated to covering the area of clinical proteomics and translational mass-spectrometry-based proteomics research. This typically involves protein profiling of human cells and tissues in health and disease state for the detection of diagnostic and prognostic biomarkers, or for developing mechanistic insights into disease pathogenicity. An important advantage of the global proteome approach is that it provides an overall catalogue of differentially expressed proteins in body fluids, cells, and tissues that define disease phenotypes in patients, which can contribute to individualized treatments.

In this Special Issue of *Proteomes*, we are looking forward to original research studies on proteome analysis of disease-related samples and welcome the submission of review articles covering recent developments in clinical proteomics.

Dr. Edwin Lasonder

*Guest Editor*





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School of Life Sciences and  
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Australia

### Prof. Dr. Jens R. Coorsen

Institute for Globally Distributed  
Open Research and Education  
(IGDORE), St. Catharines, ON L2M  
4X2, Canada

## Message from the Editorial Board

*Proteomes* is an international, peer-reviewed, open access journal that was first published in 2013 by MDPI. *Proteomes* addresses all aspects of proteome analysis with a special focus on the quantification and characterisation of the proteome at the level of proteoforms. We encourage submission of articles that accurately quantify and characterise the proteome, as well as new and updated methods and technologies that enhance the accurate quantification and characterisation of the proteome and thereby provide evidence directly facilitating the understanding of biological mechanisms. Articles emphasising a multi/ transdisciplinary approach combining different omics techniques are welcomed.

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*Proteomes* Editorial Office  
MDPI, Grosspeteranlage 5  
4052 Basel, Switzerland

Tel: +41 61 683 77 34  
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