



Proteomics and Its Applications in Cancers 2.0

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Message from the Guest Editor

Dear Colleagues,

The search, determination, separation, quantitative, and qualitative identification of protein molecules that play a role in providing sensitivity or directly in the formation of a disease are the main tasks of proteomics. The most popular is the comparative analysis of proteomes in the field of cancer research. The detection of changes in proteoform profiles associated with pathology enables the selection of the most characteristic markers of the diseases and their use for inclusion in the developed barcodes, which represent powerful tools for monitoring the course of cancer and the efficacy and safety of therapeutic agents. However, the analysis of large data sets obtained in various conditions and in different tissues and organs is now coming to the fore. In this Special Issue, we would like to gather contributions reporting very recent advances in the proteomics study of cancer. We invite you to participate in this Special Issue.





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Message from the Editor-in-Chief

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