



Recent Advances of Proteomics Applied to Cancer and Human Diseases

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Message from the Guest Editors

Cancer is a complex disease with dysregulation of normal cellular signaling networks that control cell behaviors, such as proliferation and apoptosis, caused by genetic, genomic, and epigenetic alterations at the cellular or tissue levels. It includes several modifications at the proteome level, such as modifications of protein expression level or localizations; abnormal protein expressions; modifications of post-translational patterns; and altered signalization, metabolic pathways, and immune system. In this Special Issue, we would like to gather contributions reporting very recent advances in the development of proteomics in cancer. We invite you to participate in this Special Issue, and we expect original works addressing the biological models (2D and 3D cellular models, biological fluids, and tissues from biobanks) suited for proteomics analysis, the discovery of new protein biomarkers and targets, the determination of post-translational modifications, their profiles, cross-talks and implications in signal transduction, cellular reprogramming, and immune response during tumoral progression.





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Message from the Editor-in-Chief

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