



Advanced Applications of Mass Spectrometry for Proteomics Analysis

Guest Editor:

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Message from the Guest Editor

Dear Colleagues,

Mass spectrometry is firmly established as an essential analytical technique in almost every branch of the life sciences, and in particular biomedical research. It is at the heart of proteomics, the term given to large-scale and/or unbiased protein identification, quantitation, and characterization. Advances in proteomics-based research are underpinned by new technologies and methods, and this Special Edition aims to highlight recent developments including but not limited to sample preparation, applications of data-independent acquisition, micro-scale analysis, and characterization of post-translational modifications.

Dr. Richard Wilson
Guest Editor





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Message from the Editor-in-Chief

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