



Introduction to Protein Mass Spectrometry

Pradip Kumar Ghosh

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Introduction to Protein Mass Spectrometry provides a comprehensive overview of this increasingly important, yet complex, analytical technique. Unlike many other methods which automatically yield an absolutely unique protein name as output, protein mass spectrometry generally requires a deduction of protein identity from determination of peptide fragmentation products.

This book enables readers to both understand, and appreciate, how determinations about protein identity from mass spectrometric data are made. Coverage begins with the technical basics, including preparations, instruments, and spectrometric analysis of peptides and proteins, before exploring applied use in biological applications, bioinformatics, database, and software resources.

Citing the most recent and relevant work in the field of biological mass spectrometry, the book is written for researchers and scientists new to the field, but is also an ideal resource for those hoping to hone their analytical abilities.

- Offers introductory information for scientists and researchers new to the field, as well as advanced insight into the critical assessment of computer-analyzed mass spectrometric results and their current limitations
- Provides examples of commonly-used MS instruments from Bruker, Applied Biosystems, JEOL, Thermo Scientific/Thermo Fisher Scientific, IU, and Waters
- Includes biological applications and exploration of analytical tools and databases for bioinformatics

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