



# Acceleration and Improvement of Protein Identification by Mass Spectrometry

*Willy Vincent (Ed.) Bienvenut*

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At present where protein identification and characterisation using mass spectrometry is a method of choice, this book is presenting a review of basic proteomic techniques. The second part of the book is related to the novel high throughput protein identification technique called the 'molecular scanner'. Several protein identification techniques are described, especially the peptide mass fingerprint with MALDI-MS based method. E.g. ionisation process, matrix available, signal reproducibility and suppression effect, as well as data treatment for protein identification using bioinformatics tools.

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