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1.3.4 What Are Non-Native Protein Conformations?: Random Coils, Molten Globules, and Folding Intermediates; 1.3.5 Protein Folding Pathways; 1.4 Protein Energy Landscapes and the Folding Problem; 1.4.1 Protein Conformational Ensembles and Energy Landscapes: Enthalpic and Entropic Considerations
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3.1.2 Macromolecular Mass: Terms and Definitions

Sommario/riassunto

The definitive guide to mass spectrometry techniques in biology and biophysics. The use of mass spectrometry (MS) to study the architecture and dynamics of proteins is increasingly common within the biophysical community, and *Mass Spectrometry in Structural Biology and Biophysics: Architecture, Dynamics, and Interaction of Biomolecules, Second Edition* provides readers with detailed, systematic coverage of the current state of the art. Offering an unrivalled overview of modern MS-based armamentarium that can be used to solve the most challenging problems in biophysics, structural biology, and beyond.
