



Macromolecular Crystallography, Part B, Volume 277

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Macromolecular Crystallography, Part B is the "linch-pin" of the ongoing revolution in structural biology. In the past decade, fundamental transformations of nearly every technical aspect of structure determination by X-ray diffraction have taken place. The contributions of this volume and its companion **Volume 276** emphasize the revision and extension of statistical tools that underly all phases of structure determination, the practical and conceptual impact of synchrotron radiation, horizon phase-determination methods, and computerized automation.

Key Features

- * Horizon methods for phase determination
- * Structure building, refinement, and analysis
- * Analysis of static diffraction patterns and multiple diffraction patterns recorded as a function of time
- * Accessory software for manipulating, archiving, analyzing, and presenting structures
- * Documentation of integrated software packages containing the tools needed for structure solution

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