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Autor: Muñoz Victor

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Protein folding is the process by which newly synthesized proteins fold into the specific three-dimensional structures defining their biologically active states while avoiding aggregation into pathogenic assemblies. It has always been a major focus of research in biochemistry and has often been seen as the unsolved second part of the genetic code. Protein Folding, Misfolding and Aggregation: Classical Themes and Novel Approaches includes chapters in the areas that have witnessed major developments and are written by top experts in the field. The book is unique in its scope and in its coverage of all the new developments in this area by filling a much needed gap in the current literature." The book is essential reading for graduate and postdoctoral students actively involved in protein folding research, other scientists interested in the recent progress of the field, and instructors revamping the protein folding section of their biochemistry and biophysics courses.