



Low-Abundance Proteome Discovery: State of the Art and Protocols

Egisto Boschetti, Pier Giorgio Righetti

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Low-Abundance Proteome Discovery addresses the most critical challenge in biomarker discovery and progress: the identification of low-abundance proteins. The book describes an original strategy developed by the authors that permits the detection of protein species typically found in very low abundance and that may yield valuable clues to future discoveries. Known as combinatorial peptide ligand libraries, these new methodologies are one of the hottest topics related to the study of proteomics and have applications in medical diagnostics, food quality, and plant analysis. The book is written for university and industry scientists starting proteomic studies of complex matrices (e.g., biological fluids, biopsies, recalcitrant plant tissues, foodstuff, and beverage analysis), researchers doing wet chemistry, and graduate-level students in the areas of analytical and biochemistry, biology, and genetics.

- Covers methodologies for enhancing the visibility of low-abundance proteins which, until now, has been the biggest challenge in biomarker progress
- Includes detailed protocols that address real-life needs in laboratory practice
- Addresses all applications, including human disease, food and beverage safety, and the discovery of new proteins/peptides of importance in nutraceuticals
- Compiles the research and analytic protocols of the two scientists who are credited with the discovery of these landmark methodologies, also known as combinatorial peptide ligand libraries, for the identification of low-abundance proteins

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