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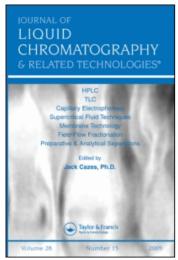
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Journal of Liquid Chromatography & Related Technologies

Publication details, including instructions for authors and subscription information: http://www.informaworld.com/smpp/title~content=t713597273

The Book Corner

To cite this Article (2006) 'The Book Corner', Journal of Liquid Chromatography & Related Technologies, 29: 11, 1675 — 1682

To link to this Article: DOI: 10.1080/10826070600678399 URL: http://dx.doi.org/10.1080/10826070600678399

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Journal of Liquid Chromatography & Related Technologies®, 29: 1675–1682, 2006

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DOI: 10.1080/10826070600678399

The Book Corner

Separation Methods in Microanalytical Systems, Edited by Jörg P. Kutter and Yolanda Fintschenko, 2006, CRC Taylor & Francis Group, New York, NY, 575 pages. Price: \$159.95

Separation Methods in Microanalytical Systems is a well designed, organized, and written book which deals with a timely topic.

In the last couple of decades scientists started talking about miniaturization of analytical instrumentation and lab-on-a-chip. This book concerns itself with certain aspects of microfluidics—the behavior of fluids in confined spaces and the manipulation of these fluids—namely, the possibility to perform chemical analyses, biochemical assays, and similar processes. The products of this kind of research are often dubbed micro-total analysis systems (μ -TAS) or, more generally, lab-on-a-chip (LOC) devices. As it is intended for a wide audience, it was also written by contributors from many of the disciplines that constitute the backbone of the LOC community.

Of course, this book cannot attempt to cover the entire field of LOC. Instead it focuses on what has been one of the main driving forces behind the development of LOC for the last 15 years: miniaturized separation systems. Separation units are still at the heart of many micro-TAS and LOC devices, and modern separation techniques are indispensable tools for analytical chemists. This book gives an overview of separation techniques on micro-fabricated devices: theoretical background information, design and understanding, fabrication and material issues, implementations, and separation systems in relation to other parts of LOC applications (sample preparation, detection, etc.). It is intended as a one-stop shopping guide for questions concerning separation techniques in microanalytical devices. It is, however, not so much meant only as a quick reference guide, but rather as a place to linger and browse. It is very likely that the information is provided in several locations within the book. A multiauthor volume gives the reader different styles, different approaches, and different opinions. Many topics are so common that they reappear in different chapters, showing different angles to approach a given problem, reflecting the different backgrounds from which researchers attach the same issues.

This excellent volume makes a good reference for all those interested in microfludics and can be a text for a graduate course.

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Haleem J. Issaq, Ph.D. Editor, The Book Corner

Using Mass Spectrometry Drug Metabolism Studies, Edited by Walter A. Korfmacher, 2005, CRC Press, New York, NY, 370 pages. Price: \$159.95

Mass spectrometry (MS) is, without a doubt, the most important and developed analytical technique for the analysis not only of small organic compounds but large biomolecules ranging from amino acids to peptides to proteins. The introduction of time-of-flight and electrospray ionization revived mass spectrometry and elevated it to a very high level where MS is a must technique for the analytical laboratory.

The current book is a welcome addition to books dealing with MS. According to the editor the book was designed to be a resource book for professionals in both mass spectrometry and drug metabolism areas, but will also be helpful to medicinal chemists interested in learning more about

drug metabolism issues in new drug discovery. The text features current knowledge in stand-alone chapters that address specific topics thoroughly enough to be read independently, with notes and references to other chapters for further reading. The first eight chapters discuss current topics regarding the use of MS for analyzing various types of *in vitro* and *in vivo* drug metabolism samples, and the final four chapters describe the latest MS technology and its uses. Throughout, expert authors demonstrate how to apply MS to determine drug metabolism parameters. They also detail the different drug metabolism concepts and their significance.

The book's twelve chapters deal with different topics that at times seem to be remotely connected. The mass spectrometer topics are scattered toward the end of the book; Chapter 7 deals with MS while Chapter 8 deals with special requirements for metabolite characterization, and Chapter 9 is a return to MS-APPI, Chapter 12 deals with electrospray ionization. Although the topics of the chapters are not well organized, each chapter is well written, illustrated and referenced.

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Haleem J. Issaq, Ph.D. Editor, The Book Corner

Knowledge Discovery in Proteomics, by Igor Jurisica and Dennis Wigle, 2006, Chapman & Hall/CCR Taylor & Francis Group, New York, NY, 318 pages. Price: \$89.95

Knowledge Discovery in Proteomics is another recent addition in books published dealing with proteomics. The book is made up of seven chapters with five specific domains. The authors state, "We felt this book was a timely discussion of some of the key issues in the field. In subsequent chapters we discuss a number of examples from our own experience that represent some of the challenges of knowledge discovery in high-throughput proteomics. This discussion is by no means comprehensive, and does not attempt to highlight all relevant domains. However, we hope to provide the reader with an overview of what we envision as an important and emerging field in its own right by discussing the challenges and potential solutions to the problems presented. We have selected five specific domains to discuss: (1) Mass spectrometry based protein analysis; (2) Protein-protein interaction network analysis; (3) Systematic high-throughput protein crystallization; (4) A systematic and integrated analysis of multiple data repositories using a diverse set of algorithms and tools; and (5) Systems biology. In each of these areas, we describe the challenges created by the type of data produced, and potential solutions to the problem of data mining within the domain. We hope this stimulates even more discussion, and newer and better ways to deal with the problems at hand."

The book is very well written and illustrated. However, the areas of proteome fractionation and separation are rather limited and weak. The references are placed at the end of the book and not at the end of each chapter. This volume of the Mathematical Biology and Medicine series is reasonably priced.

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Haleem J. Issaq, Ph.D. Editor, The Book Corner

Volume 44 Advances in Chromatography, Edited by Eli Grushka and Nelu Grinberg, 2006, CRC Taylor & Francis Group, New York, NY, 337 pages. Price: \$94.95

Advances in Chromatography, a compilation of timely topics, is a very successful series which was first published forty years ago. The recently published Volume 44 is a welcome addition to the chromatographer's library. The current volume introduces Dr. N. Grinberg who replaces Professor Phyllis Brown and joins Professor Eli Grushka as co-editor. We wish Professor Brown the very best and we will miss her insight.

In the Foreword of this volume Professor Grushka tells the reader a short history of this series and the scientists who made it a must reference. He writes: "With Volume 44 of the Advances in Chromatography series I say goodbye, reluctantly, to Phyllis R. Brown, who served as one of the editors of the series. Advances in Chromatography was started by J. Calvin Giddings and Roy A. Keller in 1965. Very quickly, the series became an important tool for quick dissemination of new developments in the field. In Volume 12 of the series (1975) Jack Cazes and I joined Giddings and Keller as editors. In 1976 (Volume 14) Phyllis replaced Roy Keller. Jack Cazes left the series in the mid 1980s and Cal Giddings stepped down from his position as executive editor in 1993. Over the last 10 years or so Phyllis and I were the sole editors of the Advances in Chromatography series. Several years ago Phyllis retired from her position as a chemistry professor at the University of Rhode Island, and now she has decided that the time has come to hand over her editorial responsibilities to the younger generation. Nelu Grinberg was selected to replace Phyllis, and with this volume he makes his debut as an editor. Phyllis served as an editor for almost 30 years. In many respects she was the primary moving force of the series. The fact that Advances in Chromatography is still going strong after 40 years is due to her foresight, determination, diligence, and perseverance. Phyllis has a very good eye for recognizing the latest developments in chromatography and related techniques. Her purpose was always to "bring to our readers the latest developments and advances at the forefront of the field." Using her many acquaintances and contacts, we succeeded in staying on top of the field and providing our readers with up-to-date reviews in all areas of chromatography and other elution techniques. Based on her own research, Phyllis recognized very early the usefulness of HPLC in life sciences. She saw to it that the Advances in Chromatography series educates its readers about the power of the technique as applied to the biological field."

This volume is made up of twelve chapters dealing with widely different but timely topics. This volume conveys the enormous potential, versatility, and challenges of the multi-channel microchip concept and its applications in cellular biology and genomic sequencing, describes multiple-channel microchip designs, substrates, sample loading, detection methods, and applications, discusses the increasingly popular use of temperature in

chromatographic separation, details the use of different separation techniques for determining the lipophilicity of analytes, covers HPLC, TLC, counter-current chromatography, explains how high-performance liquid chromatography (HPLC) can be used in controlling the quality of the pharmaceutical product, presents polysaccharide CSPs in terms of their mechanisms of action and interaction with enantiomeric analytes, illustrates the effects of chaotropic agents in reversed-phase HOLC, and highlights the challenges presented by the separation of water-insoluble proteins.

Each chapter is written by an expert in his area of research. The chapters, overall, are well written and illustrated.

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