AB Applied Biosystems CIEX

Free Seminar

Advances in Mass Spectrometry Workflows for Proteomics:

Identification, Characterization, Biomarker Discovery and Validation

8:00 am to 3:00 pm January 31st, 2006

Cambridge Hyatt Regency, Massachusetts, 575 Memorial Drive, Cambridge, Massachusetts Tel: 617.492.1234 Fax: 617-491-6906

Join us for an educational seminar where you will hear from customer researchers as well as Applied Biosystems/MDS SCIEX experts. Below is the exciting Boston agenda for our January 31st meeting.

8:30	Welcome/Introduction
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- 8:45 The Role of Proteomics in Systems Pharmacology Steve Martin, Beyond Genomics Medicine
- 9:30 Advances in LC-MALDI Technology for Definitive Protein ID in Complex Samples Matt Willetts, Applied Biosystems
- 10:00 Break
- 10:15 Quantitative Proteomic Analysis of Cellular Signaling Networks Forest White, Biological Engineering, Massachusetts Institute of Technology
- 11:00 Mass Spectrometry: The Indispensable Tool in Discovery, Identification and Validation of Endometrial Cancer Markers Michael Siu, York University
- 11:45 Merging Brains and Brawn Introducing the New QSTAR® Elite LC/MS/MS System for Proteomics Research Dominic Gostick, Applied Biosystems
- 12:15 Lunch
- 1:15 Strategies for Biomarker Verification / Validation Using the MIDAS[™] Workflow on the 4000 QTRAP[®] LC/MS/MS System Jim Ferguson, Applied Biosystems
- 2:00 Protein expression profiling and biomarker discovery using stable isotopic amino acids in cell culture Invitrogen
- 2:30 Questions, adjourn

There is no charge for attendance, but space is limited. For registration details and the date and location of the free seminar in your area.

Visit – http://info.appliedbiosystems.com/MSRoadshow

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