

**28th Sep 2022**

08:30-08:45	<b>Welcome and Introductory Comments</b> Co-chair: Da Ren, <i>Amgen Inc.</i>
08:45-09:45	<b>Keynote I</b> <b>In Quest of SPARQLING Lights in the Dark Human Proteome</b> <u>Lydie Lane</u> <i>Swiss Institute of Bioinformatics</i>
09:45-10:15	<b>Networking Break</b>
10:15-11:50	<b>Session I - Contemporary Bioinformatics Applications for Mass Spectrometry</b> Session Chairs: Da Ren, <i>Amgen Inc.</i> and Richard Rogers, <i>Umoja BioPharma, Inc.</i>  <b>10:20-10:50</b> <b>Development of Automated Workflows for Analysis and Reporting of Product-Specific Peptide Mapping Data</b> <u>Michael Pettit</u> <i>Janssen Research &amp; Development, LLC, Malvern, PA, USA</i>  <b>10:50-11:20</b> <b>Chemometric Approaches to Evaluate and Predict the Performance of Biopharmaceutical Manufacturing Processes</b> <u>Noemi Dorival-Garcia,</u> <i>The National Institute for Bioprocessing Research and Training (NIBRT), Dublin, Ireland</i>  <b>11:20-11:50</b> <b>Fast, Large-Scale Analysis of Mass Spectrometry Data Repositories to Find the Needle in the Haystack</b> <u>Simon Letarte</u> <i>Gilead Sciences, Oceanside, CA, USA.</i>
12:05-12:35	<b>Technical Seminar Sponsored by SCIEX</b>  <b>12:05-12:35</b> <b>A Disruptive Approach to the Characterization of mAb Charge Variants During Process Development</b> <u>Greg Adams</u> <i>FUJIFILM Diosynth Biotechnologies, Morrisville, North Carolina, USA</i>
12:40-13:10	<b>Technical Seminar Sponsored by Agilent Technologies, Inc.</b>  <b>Recent Advances in Oligo Purity and Sequence Determinations by LCMS: Maximizing Throughput, Confidence, and Coverage.</b> <u>Peter Rye</u> <i>Agilent Technologies, Lexington, MA, USA</i>
13:10-14:20	<b>Poster Session I</b>

<p>14:20-15:25</p>	<p><b>Session II - Molecular Design, Developability and Biotransformation</b>  Session Chairs: Anders Lund, <i>Synlogic</i> and Andrew Mahan, <i>Janssen Research &amp; Development, LLC</i></p> <p><b>14:25-14:55</b>  <b>LC-MS-based Target Engagement Assays, Enabling Decision Making in Drug Development</b>  <u>Michelle Robinson</u>  <i>Merck &amp; Co., Inc., West Point, PA, USA</i></p> <p><b>14:55-15:25</b>  <b>Two-Dimensional Liquid Chromatography-Mass Spectrometry (2D-LC-MS) for Simultaneous Multi-Attribute Characterization of Adeno-Associated Viruses</b>  <u>Zhijie Wu</u>,  <i>Regeneron Pharmaceuticals, Inc., Tarrytown, NY, USA</i></p>
<p>15:25-17:00</p>	<p><b>Session III - Emerging Mass Spectrometry Technologies for HOS</b>  Session Chairs: John Valliere-Douglass, <i>Seagen Inc.</i> and Christopher Yu, <i>Genentech, a Member of the Roche Group</i></p> <p><b>15:30-16:00</b>  <b>Variable Temperature Native Mass Spectrometry for Studying Thermodynamics of Protein-Ligand Interactions</b>  <u>Thomas Walker</u>  <i>Texas A&amp;M University, College Station, TX, USA</i></p> <p><b>16:00-16:30</b>  <b>Study of Monoclonal Antibody Aggregation by Size Exclusion Chromatography Coupled With Native Mass Spectrometry</b>  <u>Jing Xu</u>  <i>Biogen, Cambridge, MA, USA</i></p> <p><b>16:30-17:00</b>  <b>Evaluation of Monoclonal Antibody HOS Change Upon Forced Degradation by Biophysical and Mass Spectrometry-Based Technologies</b>  <u>Yanchun Lin</u>  <i>Washington University in St Louis, St Louis, MO, USA.</i></p>
<p>17:00-18:00</p>	<p><b>Welcome and New Member Reception</b></p>

29th Sep 2022

08:30-08:35	<b>Welcome and Introductory Comments</b> Speaker: Christopher Chumsae, <i>Bristol-Myers Squibb Company</i>
08:35-09:35	<b>Keynote II</b>  <b>08:35-09:35</b> <b>MAM: Multi-Artifacts Monitoring – Operators, Processes and Modalities</b> <u>Zhaohui (Sunny) Zhou</u> <i>Northeastern University, Boston, MA, USA</i>
09:35-10:05	<b>Networking Break</b>
10:05-11:40	<b>Session IV - Mass Spectrometry in Gene and Cell Therapy</b> Session Chairs: Jason Rouse, <i>Pfizer, Inc.</i> and Christopher Yu, <i>Genentech, a Member of the Roche Group</i>  <b>10:10-10:40</b> <b>Application of LC-MS for Analysis and Characterization of Pharmaceutical mRNA Products</b> <u>Tao Jiang</u> <i>Moderna Inc., Norwood, MA, USA</i>  <b>10:40-11:10</b> <b>Unlocking Analytics for AAV Gene Therapy Programs: Leveraging Standard Biotherapeutic Strategies to Transform New Modalities</b> <u>Thomas Powers</u> <i>Pfizer, Chesterfield, MO, USA</i>  <b>11:10-11:40</b> <b>Characterization of Host Cell Proteins for AAV Using Native Digestion and Quantitative Proteomics</b> <u>Jean Lodge</u> , Lihua Huang, Zhirui Lian, Yuwei Tian <i>Eli Lilly and Company, Indianapolis, IN, USA</i>
11:55-12:25	<b>Technical Seminar Sponsored by Thermo Fisher Scientific</b>  <b>11:55-12:25</b> <b>Profiling Complex Biotherapeutics With Charge Reduction or Charge Detection Native Mass Spectrometry</b> <u>Wendy Sandoval</u> <i>Genentech Inc., South San Francisco, CA, USA.</i>
12:30-13:00	<b>Technical Seminar Sponsored by Waters Corporation</b>  <b>12:30-13:00</b> <b>Advancing Mass Spectrometry Workflows: From the Revolutionary to the Routine</b> <u>Scott Berger</u> <i>Waters Corporation, Milford, MA, USA</i>

13:00-14:00	<b>Poster Session II</b>
14:00-15:05	<p><b>Session V - Next Generation Investigators</b>  Session Chairs: Jonathan (JJ) Josephs, <i>Genentech, a Member of the Roche Group</i> and Douglas Richardson, <i>Merck &amp; Co., Inc.</i></p> <p><b>14:05-14:25</b>  <b>Localization of O-Glycosylation in a CD24 Fc-Fusion Protein by Site-Directed Mutagenesis and Native Mass Spectrometry</b>  <u>Zac VanAernum</u>  <i>Merck &amp; Co., Inc., Rahway, NJ, USA</i></p> <p><b>14:25-14:45</b>  <b>Guiding the Cell Line Development of Bispecific Molecules Using a Newly Developed Native SECMS and Quantitative Chain Ratio Assay</b>  <u>Elsa Gorre</u>  <i>Janssen Pharmaceutical R&amp;D, LLC, Spring House, PA, USA</i></p> <p><b>14:45-15:05</b>  <b>Application of Free Cysteine Footprinting by Differential Alkylation in BioProcess Development</b>  <u>Roger Liu</u>  <i>Bristol-Myers Squibb Company, Devens, MA, USA. <sup>2</sup>Bristol-Myers Squibb Company, New Brunswick, NJ, USA</i></p>
15:05-15:30	<b>Networking Break</b>
15:30-16:30	Roundtable Discussions
16:30-17:30	<b>Exhibitor and Poster Reception</b>

30th Sep 2022

08:30-08:35	<b>Welcome and Introductory Comments</b> Speaker: Da Ren, <i>Amgen Inc.</i>
08:35-09:35	<b>Keynote III</b> <b>Charge Detection Mass Spectrometry for the Analysis of Gene Therapy Vectors</b> Martin Jarrold <i>Indiana University</i>
09:35-10:00	<b>Networking Break</b>
10:00-12:05	<b>Session VI - Product and Process Characterization</b> Session Chairs: Christopher Chumsae, <i>Bristol-Myers Squibb Company</i> and Ingo Lindner, <i>Roche Diagnostics GmbH</i>  <b>10:05-10:35</b> <b>Identification of N-Linked Glycans From Bovine Alpha-Thrombin Using a Combination of Lc-MS Techniques</b> <u>Dennis Gessmann</u> <i>Pfizer, Inc., Andover, MA, USA.</i>  <b>10:35-11:05</b> <b>Probing Residual Host Cell Proteins in Monoclonal Antibody and Its High Molecular Weight Species: A Different Way to Study Difficult-to-Remove HCPs</b> <u>Yunli Hu</u> <i>Regeneron Pharmaceuticals, Inc., Tarrytown, NY, USA</i>  <b>11:05-11:35</b> <b>Identification of Hetero-Aggregates in Antibody Co-Formulations by Multi-Dimensional Liquid Chromatography Coupled to Mass Spectrometry</b> <u>Felix Kuhne</u> <i>Roche Diagnostics GmbH, Penzberg, Germany</i>  <b>11:35-12:05</b> <b>Performance Characteristics of Mass Spectrometry- Based Methods for Quantitation of Nitrosamines: Insight From an Inter-Laboratory Study</b> <u>Jingyue Yang</u> <i>CDER, FDA, Saint Louis, MO, USA</i>
12:20-12:50	<b>Technical Seminar Sponsored by MOBILion Systems, Inc.</b>  <b>12:20-12:50</b> <b>Tackling the Glycan Isomer Challenge for Your Biotherapeutics With MOBIE®</b> <u>Heidi Vitrac</u> <i>MOBILion Systems, Chadds Ford, PA, USA</i>
12:55-13:25	<b>Technical Seminar Sponsored by Genovis Inc.</b>  <b>12:55-13:25</b> <b>Enzymatic Strategies for Characterization of Biopharmaceuticals</b>

	<p><u>Laurent Rieux</u> <i>Genovis Inc., Cambridge, MA, USA</i></p>
<p>13:30-15:05</p>	<p><b>Session VII - New Technologies, Approaches and Methods</b> Session Chairs: Richard Rogers, <i>Umoja BioPharma, Inc.</i> and John Valliere-Douglass, <i>Seagen Inc.</i></p> <p><b>13:35-14:05</b> <b>Lab of the Future: Fully Integrated Automation for Protein Characterization and Biologics Development</b> <u>Hirsh Nanda</u> <i>Janssen Pharmaceutical Companies of Johnson &amp; Johnson, Springhouse, PA, USA.</i></p> <p><b>14:05-14:35</b> <b>The Application of Mass Spectrometry for Potency Readout of mRNA Vaccines From in Vitro Translation</b> <u>Alyssa Stiving</u> <i>Merck &amp; Co., Inc., West Point, PA, USA.</i></p> <p><b>14:35-15:05</b> <b>Peptide Mapping: Two Lc Methods Are More Informative (And Faster!) Than One</b> <u>James Arndt</u> <i>Teva Pharmaceuticals, West Chester, PA, USA</i></p>
<p>15:05-15:15</p>	<p><b>Poster Award Announcement</b></p>
<p>15:15-15:30</p>	<p><b>Closing Remarks</b> Co-chair: Christopher Chumsae, <i>Bristol-Myers Squibb Company</i></p>

