

## **ISPPP 2013**

# 33rd International Symposium and Exhibit on the

# Separation and Characterization of Biologically Important Molecules

**ISPPP 2013 Scientific Program** (as of 6-19-2013)

# Wednesday, July 17, 2013

**Symposium Registration Open** 7:30 AM 10:00 AM Exhibition Opens in Harbor Ballroom II & III **Welcome and Opening Remarks in Burroughs Room** 8:10 AM Wednesday Session 1. Analytical Separations Location: Burroughs Room 8:20 AM L-101 **Orthogonal Bioseparations.** Mark Schure, Kroungold Analytical, Blue Bell, PA, USA 8:40 AM L-102 New Advances in Hydrophilic Interaction Chromatography of Peptides: Comparison of HILIC/SALT to HILIC and RPC. Colin Mant, Robert Hodges, Department of Biochemistry and Molecular Genetics, University of Colorado Denver, Aurora, CO, USA 9:00 AM L-103 Highly Efficient LC-MS of Intact Proteins using Sub-0.5µm Particles with Slip Flow. Zhen Wu<sup>1</sup>, Bingchuan Wei<sup>2</sup>, Ximo Zhang<sup>1</sup>, Mary Wirth<sup>1</sup>, <sup>1</sup>Purdue University, West Lafayette, IN, USA; <sup>2</sup>University of Washington, Seattle, WA, USA 9:20 AM L-104 Capillary-channeled Polymer (C-CP) Fibers for High Throughput Analytical Separations and Desalting of Proteins Prior to MALDI and ESI-MS. R. Kenneth Marcus<sup>1</sup>, Abby Schadock-Hewitt<sup>1</sup>, Benjamin T. Manard<sup>1</sup>, Carolyn Q. Burdette<sup>2</sup>, Marissa Pierson<sup>1</sup>, <sup>1</sup>Clemson University, Clemson, SC, USA; <sup>2</sup>National Institute of Standards and Technology, Gaithersburg, MD, USA 9:40 AM L-105 Increasing the Peak Capacity of Peptide Separations using Long Microcapillary Columns and Sub 2 µm Particles at 30,000+ psi. Kaitlin Fague, Justin Godhino, Edward Franklin, Jordan Stobaugh, University of North Carolina, Chapel Hill, NC. USA 10:00 AM **Break in Exhibit and Poster Session Hall** (Location: Harbor Ballroom II & III)

# Wednesday Session 2. Proteomics and Protein Measurements Location: Burroughs Room 10:40 AM L-106 Application of a Robust and Modular Cartridge based NanoLC System for **Proteomics.** Michael Bereman<sup>1</sup>, Tom Corso<sup>2</sup>, Colleen Van Pelt<sup>2</sup>, Michael MacCoss<sup>1</sup>, <sup>1</sup>University of Washington, Seattle, WA, USA; <sup>2</sup>CorSolutions, Ithaca, NY, USA 11:00 AM L-107 Protein Markers for Cancer Stem Cells in Pancreatic Cancer. David Lubman, Jianhui Zhu, University of Michigan, Ann Arbor, MI, USA 11:20 AM L-108 Target-based Multiplex MRM Assays. Christopher M. Colangelo, Lisa Chung, Shifman Mark, Abbott Thomas, Fumika Sakaue, Angus Nairn, Willaims Kenneth, Yale University, New Haven, CT, USA 11:40 AM L-109 Pretreatment of Human Fluid Samples for Trace Protein Analysis. Frank Jahnke, Sonata Biosciences, Inc., Auburn, CA, USA 12:00 PM Lunch on own 12:00-1:30 PM Free Vendor Seminar Sponsored by Waters Corporation Location: Carlton Room Must register with Waters in order to attend as seating is limited 1:30 - 3:00 PM Poster Session I in Exhibit and Poster Session Hall (Location: Harbor Ballroom II & III) 3:00 - 3:45 PM Break/Social in Exhibit and Poster Session Hall (Location: Harbor Ballroom II & III) Wednesday Session 3. Protein Therapeutics Location: Burroughs Room L-110 IgG Asparagine-linked Oligosaccharide Profiling by High-performance 3:45 PM Anion-exchange Chromatography with Pulsed Amperometric Detection. Jeffrey Rohrer<sup>1</sup>, Deanna Hurum<sup>2</sup>, Lipika Basumallick<sup>1</sup>, <sup>1</sup>Thermo Fisher Scientific, Sunnyvale, CA, USA; <sup>2</sup>Palo Alto, CA, USA 4:05 PM L-111 Size-exclusion Chromatography using Multi-angle Light Scattering (SEC-MALS) for the Characterization of Polypeptide Mixtures. Joseph Glajch, Ying Li, Momenta Pharmaceuticals, Cambridge, MA, USA 4:25 PM L-112 Developing an Automated Workflow for Disulfide Linkages Analysis of Biotherapeutics by High-Resolution LCMS. Asish Chakraborty<sup>1</sup>, Stephane Houel<sup>1</sup>, Henry Shion<sup>1</sup>, Scott Berger<sup>1</sup>, Weibin Chen<sup>1</sup>, Anurag Rathore<sup>2</sup>, <sup>1</sup>Waters Corporation, Milford, MA, USA; <sup>2</sup>Indian Institute of Technology, New Delhi, Delhi, INDIA

L-113 Analytical HIC for mAb Aggregate Analysis; How Does the Salt Ion Type Influence the Selectivity? Hannah Brueck, Judith Vajda, Werner Conze, Egbert Mueller,

5:05 - 7:00 PM Welcome Mixer in Exhibit and Poster Session Hall

Tosoh Bioscience GmbH, Stuttgart, GERMANY

(Location: Harbor Ballroom II & III)

4:45 PM

# Thursday, July 18, 2013

7:45 AM	Symposium Registration Open
10:00 AM	Exhibition Open in Harbor Ballroom II & III
	Thursday Session 4. Affinity and Ion Exchange Methods Location: Harbor Ballroom I
8:20 AM	L-114 <b>Discovery and Development of Universal Fc Binders for Antibody Purification.</b> Marc Arnold, Holger Bittermann, Thomas Neumann, Graffinity Pharmaceuticals GmbH, Heidelberg, GERMANY
8:40 AM	L-115 <b>Development of a Sialic Acid-specific Affinity Chromatography for the Purification and Separation of Glycoprotein Isoforms.</b> Matthias Meininger <sup>1</sup> , Francisco Vito Santos da Silva <sup>1</sup> , Samanta Cajic <sup>1</sup> , René Hennig <sup>1</sup> , Erdmann Rapp <sup>1</sup> , Frank Zwanziger <sup>2</sup> , Stefan Laufer <sup>2</sup> , Karl-Heinz Wiesmueller <sup>3</sup> , Heinz Rotering <sup>4</sup> , Udo Reichl <sup>1</sup> , Michael Wolff <sup>1</sup> , <sup>1</sup> Max Planck Institut for Dynamics of Complex Technical Systems, Magdeburg, GERMANY; <sup>2</sup> Eberhard-Karls-University, Tuebingen, GERMANY; <sup>3</sup> EMC Microcollections GmbH, Tuebingen, GERMANY; <sup>4</sup> Merckle Biotec GmbH, Ulm, GERMANY
9:00 AM	L-116 Reversible Cyclic and Polycyclic Peptides for the Discovery of Affinity Ligands. Stefano Menegatti, Robert Blackburn, Kevin Ward, Ruben Carbonell, North Carolina State University, Raleigh, NC, USA
9:20 AM	L-117 <b>Understanding Ion Exchange Adsorption Mechanism.</b> Francisco Marques <sup>1</sup> , Patricia Aguilar <sup>1</sup> , Marvin Thrash Jr. <sup>2</sup> , <u>Cristina Dias-Cabral<sup>1</sup></u> , <sup>1</sup> University of Beira Interior, Covilhã, PORTUGAL; <sup>2</sup> Central State University, Wilberforce, OH, USA
9:40 AM	L-118 Purification of a Potential Malaria Vaccine by Multimodal Ion Exchange Chromatography. Sonja Jensen, Jessica Paul, Gesine Cornelissen, Hamburg University of Applied Sciences, Hamburg, GERMANY
10:00 AM	Break in Exhibit and Poster Session Hall (Location: Harbor Ballroom II & III)

	Thursday Session 5. Monoclonal Antibody Separations Location: Harbor Ballroom I
10:40 AM	L-119 New Tools to Achieve Enhanced Process Productivity in the Purification of Monoclonal Antibodies and Other Recombinant Proteins. Milton Hearn, Monash University, Clayton, Victoria, AUSTRALIA
11:00 AM	L-120 Investigation of Monoclonal Antibody Purification by Aqueous Two-phase Extraction. Jan Mündges, Shuai Shi, Tim Zeiner, TU Dortmund University, Dortmund, GERMANY
11:20 AM	L-121 Fujifilm Diosynth's and Chromatan's Investigation of Countercurrent Tangential Chromatography for Purification of Monoclonal Antibodies. Oleg Shinkazh, Chromatan, State College, PA, USA
11:40 AM	L-122 <b>2D-DIGE for Host Cell Protein Analysis and Antibody Process Development.</b> Alois Jungbauer, BOKU, Vienna, AUSTRIA
12:00 PM	Lunch on own
12:00-1:30 PM	Free Vendor Seminar Sponsored by Shimadzu "Turning Mountains into Molehills: The New Landscape of Protein Analysis" Location: Griffin Room Must register with Shimadzu in order to attend as seating is limited SIGN UP BY CONTACTING <a href="https://www.surveymonkey.com/s/K5R8NVL">https://www.surveymonkey.com/s/K5R8NVL</a>
12:00-1:30 PM	Free Vendor Seminar Sponsored by Knauer "Contichrom®: A Versatile Purification Platform for Batch/CaptureSMB/MCSGP Biochromatography" Location: Carlton Room Must register with Knauer in order to attend as seating is limited SIGN UP BY CONTACTING nickel@knauer.net
1:30 - 3:00 PM	Poster Session II in Exhibit and Poster Session Hall (Location: Harbor Ballroom II & III)
3:00 - 3:30 PM	Break/Social in Exhibit and Poster Session Hall (Location: Harbor Ballroom II & III)

	Thursday Session 6. Protein Targets Location: Harbor Ballroom I
3:30 PM	L-123 Integration and Intensification of Downstream Bioprocessing based in Aqueous Two-phase Systems. Ana Azevedo, Raquel Aires-Barros, Instituto Superior Tecnico, Lisbon, PORTUGAL
3:50 PM	L-124 Multi-stage Enzyme Extraction using Aqueous Two-phase Systems – Experiment and Modeling. Axel Prinz, Katharina Koch, Tim Zeiner, TU Dortmund University, Dortmund, GERMANY
4:10 PM	L-125 Purifying and Concentrating Recovery Process Samples for Recombinant Protein Quantification. <u>Tanja Buch</u> , Ian Marison, DCU, Dublin, IRELAND
4:30 PM	L-126 Predicting Protein Solubility and Crystallization Behavior based on the Second Osmotic Virial Coefficient. Marcel Herhut, Christoph Brandenbusch, Gabriele Sadowski, Department of Biochemical and Chemical Engineering, Dortmund, GERMANY
4:50 PM	L-127 <b>Historical Retrospective of ISPPP Conferences.</b> Milton Hearn, Monash University, Clayton, Victoria, AUSTRALIA
5:10 PM	Pause
6:30 - 8:30 PM	Symposium Banquet - please purchase \$30 ticket online by July 1 Location: Burroughs room

# Friday, July 19, 2013

8:15 AM Symposium Registration Op	pen
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	Friday Session 7. Bio-Therapeutics Location: Harbor Ballroom I
8:45 AM	L-128 Improved Identification and Quantitation of Host Cell Proteins in Protein Therapeutics using 2D-LC and Ion Mobility. Weibin Chen, Catalin Doneanu, Keith Fadgen, Martha Stapels, Waters, Milford, MA, USA
9:05 AM	L-129 Additional Structural Insights on Therapeutic Glucocerebroside Variants and Impurities by SEC. John Thomas, Shire, Lexington, MA, USA
9:25 AM	L-130 TBD
9:45 AM	L-131 Impact of Plasmid Size on the Purification of Model pDNA Vaccines by HIC on Phenyl Membrane Adsorbers. Luis Raiado-Pereira, Jonathan de la Vega, D. Miguel F. Prazeres, Marilia Mateus, Instituto Superior Tecnico, Lisbon, PORTUGAL
10:05 AM	L-132 <b>Protein Biomarker Assays for Drug Safety Assessments.</b> Jennifer Colangelo, Pfizer, Groton, CT, USA
10:25 AM	Break

	Friday Session 8. Glycans and Complex Carbohydrates  Location: Harbor Ballroom I
11:00 AM	L-133 <b>Separations of Intact Glycoproteins by HILIC.</b> <u>Barry Boyes<sup>1</sup>, Ron Orlando<sup>2</sup>, Joseph DeStefano<sup>1</sup>, <sup>1</sup>Advanced Materials Technologies, Inc., Wilmington, DE, USA; <sup>2</sup>University of Georgia, Athens, GA, USA</u>
11:20 AM	L-134 <b>Progress Towards Automated Sequencing of Heparin/Heparan Sulfate.</b> <u>Joshua S. Sharp</u> , Complex Carbohydrate Research Center, University of Georgia, Athens, GA, USA
11:40 AM	L-135 <b>HILIC-MS of Glycans and Glycopeptides.</b> <u>Joseph Zaia</u> , Center for Biomedical Mass Spectrometry, Boston University, Boston, MA, USA
12:00 PM	L-136 <b>Hypothesis Driven Glycomics.</b> Shujuan Tao <sup>1</sup> , Yining Huang <sup>1</sup> , Barry Boyes <sup>2</sup> , Ron Orlando <sup>1</sup> , <sup>1</sup> Complex Carbohydrate Research Center, University of Georgia, Athens, GA, USA; <sup>2</sup> Advanced Materials Technologies, Inc., Wilmington, DE, USA
12:20 PM	Closing Remarks
12:30 PM	Adjourn

See the list of poster presentations on the following pages

# **Preliminary List of Poster Presentations**

Posters are located in the Exhibit Hall in Harbor Ballroom II & III

Wednesday & Thursday Poster Session Times: 1:30 – 3:00 PM

All posters must be put up onto the poster boards on Wednesday morning and left on the poster boards for two days. Do not remove the poster until the end of Poster Session II on Thursday afternoon. The size of each poster board is 3.6 feet wide (110cm) by 3.6 feet high (110cm). *Please see "Poster Information"* at <a href="https://www.ISPPP.org">www.ISPPP.org</a>.

- Epitope Mapping and the Selection of MAbs used in the Diagnostic Immunoassays. Cheng Zhao, Bryan Tieman, Bailin Tu, Robert Ziemann, Jeffrey Fishpaugh, Carol Ramsay, Abbott Laboratories. Abbott Park, IL, USA
- An Alternative Capture Step for Monoclonal Antibodies: Phenyl Boronate as a New Multi-modal Ligand. Ana M. Azevedo, Raquel dos Santos, Sara A.S.L. Rosa, M. Raquel Aires-Barros, Instituto Superior Tecnico, Lisbon, PORTUGAL
- Phenyl Boronic Acid as Ligand for a Multimodal Chromatography: Adsorption Behavior Comparison between Control Pore Glass and Agarose Matrixes. Rimenys Jr. Carvalho<sup>1</sup>, James Woo<sup>2</sup>, Karim A. Nakamura<sup>2</sup>, Maria Raquel Aires-Barros<sup>1</sup>, <u>Ana M. Azevedo<sup>1</sup></u>, Steven M. Cramer<sup>2</sup>, <sup>1</sup>Instituto Superior Tecnico, Lisbon, PORTUGAL; <sup>2</sup>Rensselaer Polytechnic Institute, Troy, NY, USA
- Aggregate Removal with Cation Exchange Chromatography (NuviaTM HR-S). Paul K. Ng, Mark A. Snyder, Bio-Rad Laboratories, Hercules, CA, USA
- Optimized One-day Assay for Quantitation of Monosaccharide Content in Proteins by 2-Picoline-Borane Reductive Amination. Nicholas Woon, Genentech, South San Francisco, CA, USA
- Chiral Separation of D,L-Phenylglycine using an Enantioselective Membrane Formed by Polycondensation of Bovine Serum Albumin with 1,6-Diisocyanatohexane on a Polysulfone Membrane. <u>Li-Ming Yuan</u>, Guang-Yong Zeng, Department of Chemistry, Yunnan Normal University, Kunming, Yunnan, P.R. CHINA
- Methodology of Monoclonal Antibody Charge-variant Analysis by Ion-exchange Chromatography. Hillel Brandes, Roy Eksteen, Sigma Aldrich / Supelco, Bellefonte, PA, USA
- Analysis of Associated Forms of Insulins. <u>Sara Fexby Garmer</u>, Martin Mårtensson, Dorte Bjerre Steensgaard, Aage Hvass, Svend Havelund, Novo Nordisk A/S, Måløv, DENMARK
- 2D Separation and Immunoreactive Coverage of CHO and E.coli Host Cell Proteins (HCP) by Polyclonal Antibodies – Use of Fluorescent Labeling (Cyanine Dyes) in 2D Western Blotting. Harbhajan Dhillon, Mark Abbott, Jonathan Basch, Mark Panek, Bristol-Myers Squibb, East Syracuse, NY, USA
- Lipidomics using Ion Mobility Mass Spectrometry with Transomics Informatics. Giuseppe Astarita<sup>1</sup>, Roy Martin<sup>1</sup>, Giorgis Isaac<sup>1</sup>, James Langridge<sup>2</sup>, Weibin Chen<sup>1</sup>, Waters Corporation, Milford, MA, USA; Waters Corporation, Manchester, UK
- N-linked Glycan Profile Comparison Between the Innovator and a Biosimilar Etanercept. Ying Qing Yu, Weibin Chen, Waters Corporation, Milford, MA, USA
- Characterization of Two Novel Analytical Chromatographic Columns for Orthogonal Analysis
  of Monoclonal Antibody and Protein Aggregates and their Isoforms. <u>Justin Steve</u>, Atis
  Chakrabarti, Tosoh Bioscience LLC, King of Prussia, PA, USA

- Greater Loading Capacity and Resolution for Improved Process-scale Peptide Purification.

  Jochen Saar<sup>1</sup>, Reno Nguyen<sup>2</sup>, Chitra Sundarajan<sup>3</sup>, Scott Anderson<sup>4</sup>, Dennis McCreary<sup>5</sup>, <u>Janine Sinck<sup>6</sup></u>, <sup>1</sup>Worms, GERMANY; <sup>2</sup>Hesperia, CA, USA; <sup>3</sup>Hyderabad, Andrapradesh, INDIA; <sup>4</sup>Deerfield, IL, USA; <sup>5</sup>Columbia, MD, USA; <sup>6</sup>Allentown, PA, USA
- Charge Surface Modified C18 Columns for Increasing Peak Capacity in LC-MS Peptide Separations with Formic Acid Mobile Phases. <u>Matthew Lauber</u>, Stephan Koza, Kenneth Fountain, Waters Corporation, Milford, MA, USA
- Purification of a Potential Malaria Vaccine by Multimodal Ion Exchange Chromatography.
   Sonja Jensen, Jessica Paul, Gesine Cornelissen, Hamburg University of Applied Sciences, Hamburg, GERMANY
- A Quality by Design Approach: Systematic Optimization of Malaria Vaccine Purification with IMAC. <u>Jessica Paul</u>, Arthur Dukart, Jasmin Zuehlke, Gesine Cornelissen, Hamburg University of Applied Sciences, Hamburg, GERMANY
- Characterization of Two Novel High Capacity Strong Ion Exchange Resins. Chinlun Huang, J. Kevin O'Donnell, Tosoh Bioscience, King of Prussia, PA, USA
- Automating Workflows for Developing Separation and Reporting Methods for Size Exclusion Chromatography. <u>Thomas E. Wheat</u>, Aparna Chavali, Patricia McConville, Waters Corporation, Milford, MA, USA
- Aggregates and Particle Characterization to Support Biomanufacturing Process Development.
   Yogesh Mudaliar, Rong-Rong Zhu, Tim Hanley, EMD Millipore, Bedford, MA, USA
- Enthalpic and Entropic Contributions in Lysozyme Adsorption onto a Cation-exchange Support. Francisco Marques<sup>1</sup>, Goncalo Silva<sup>1</sup>, Marvin Thrash, Jr.<sup>2</sup>, Cristina Dias-Cabral<sup>1</sup>, <sup>1</sup>CICS-UBI-Health Sciences Research Centre, University of Beira Interior, Covilha, PORTUGAL; <sup>2</sup>Department of Water Resources and Environmental Engineering, College of Science and Engineering, Central State University, Wilberforce, OH, USA
- Superficially Porous Particles for Peptide and Protein Analysis. <u>Barry Boyes</u>, Joseph Kirkland, Stephanie Schuster, Brian Wagner, Joseph DeStefano, Advanced Materials Technologies, Inc., Wilmington, DE, USA
- Malaria Vaccine Purification via Expanded Bed Adsorption Chromatography Combined with an Ultrafiltration. <u>Sarah Schreiber</u>, Sven Oliver Borchert, Jessica Paul, Gesine Cornelissen, University of Applied Sciences, Hamburg, GERMANY
- Microcalorimetric Study of Linear Plasmid DNA Adsorption onto an Ion Exchange Support.
   Patricia Aguilar<sup>1</sup>, Filipa Pires<sup>1</sup>, Marvin Thrash, Jr.<sup>2</sup>, Cristina Dias-Cabral<sup>1</sup>, <sup>1</sup>CICS-UBI-Health Sciences
   Research Centre, University of Beira Interior, Covilha, PORTUGAL; <sup>2</sup>Department of Water Resources
   and Environmental Engineering, College of Science and Engineering, Central State University,
   Wilberforce, OH, USA
- Continuous-batch Protein Chromatography Continuous Capture Coupled with Two-step Automated Batch Polish. <u>Peter Tiainen</u>, Jais Rose Bjelke, Ditte Skibstrup, Haleh Ahmadian, Novo Nordisk A/S, Malov, DENMARK
- Separation of DMB-Labeled Sialic Acids for the Comparison of Biosimilars to Reference Materials with an Improved Chromatographic Method. Xiaoning Lu, <u>Hillel Brandes</u>, Dave Bell, Roy Eksteen, Sigma-Aldrich, Bellefonte, PA, USA

- Bioseparations with 3.5- and 5-Micron Wide-Pore Superficially Porous Particles. <u>Joseph DeStefano</u>, Joseph Kirkland, Stephanie Schuster, Bill Johnson, Advanced Materials Technologies, Inc., Wilmington, DE, USA
- Twin Column CaptureSMB: A Novel Cyclic Process to Increase the Capacity Utilization in Protein A Chromatography. Monica Angarita<sup>1</sup>, Baur Daniel<sup>1</sup>, Thomas Muller-Spath<sup>2</sup>, Roel Lievrouw<sup>3</sup>, Geert Lissens<sup>3</sup>, Guido Strohlein<sup>2</sup>, Massimo Morbidelli<sup>1</sup>, <sup>1</sup>ETH Zurich, Zurich, SWITZERLAND; <sup>2</sup>ChromaCon AG, Zurich, SWITZERLAND; <sup>3</sup>JSR Life Sciences, Leuven, BELGIUM
- Microfluidic CE-MS Applied to Protein, Peptide, and Small Molecule Characterization. Gregory <u>Roman</u><sup>1</sup>, Scott Mellors<sup>2</sup>, Martin Gilar<sup>1</sup>, James Murphy<sup>1</sup>, J. Michael Ramsey<sup>2</sup>, <sup>1</sup>Waters Corporation, Milford, MA, USA; <sup>2</sup>University of North Carolina, Chapel Hill, NC, USA
- Effects of Mobile Phase Optimization on Analyte Behaviour in Size Exclusion Chromatography of Biomolecules. <u>James Martosella</u>, Andrew Coffey<sup>2</sup>, <sup>1</sup>Agilent Technologies, Wilmington, DE, USA <sup>2</sup>Agilent Technologies, Church Stretton, UK
- Fast and Efficient Reversed-phase Liquid Chromatography/Mass Spectrometry
   Characterization of Glycosylation in the Fc Region of a Recombinant IgG(1) Therapeutic
   Monoclonal Antibody (mAb). <u>James Martosella</u>, Phu Duong, Alex Zhu, Agilent Technologies,
   Wilmington, DE, USA
- Peptide Mapping of a Therapeutic Monoclonal Antibody (mAb): Optimizations for Increasing Speed and Peptide Identifications. <u>James Martosella</u>, Ning Tang<sup>2</sup>, Alex Zhu<sup>1</sup>, <sup>1</sup>Agilent Technologies, Wilmington, DE, USA; <sup>2</sup>Agilent Technologies, Santa Clara, CA, USA
- Countercurrent Tangential Chromatography for Purification of Monoclonal Antibodies. Oleg Shinkazh, Chromatan, State College, PA, USA
- A Novel Automated Enrichment Process for the Isolation of Product-related Impurities from Active Pharmaceutical Ingredients. <u>Thomas Muller-Spath</u>, Nicole Ulmer, Lars Aumann, Guido Strohlein, Michael Bavand, ChromaCon AG, Zurich, SWITZERLAND
- Combining Small-scale Purification and Analysis of Monoclonal Antibodies on One Instrument. Sonja Schneider, Agilent Technologies, Waldbronn, GERMANY
- Amphoteric Ion-Exchange Separation of Biomolecules with Porous or Non-Porous Polymer-Based Resins. Ken Tseng<sup>1</sup>, Toshi Ono<sup>1</sup>, Tsunehisa Hirose<sup>2</sup>, Kazuhiro Kimata<sup>2</sup>, <sup>1</sup>Nacalai USA Inc., San Diego, CA, USA; <sup>2</sup>Nacalai Tesque, Kyoto, JAPAN
- New Wide Pore Media Improves Loading Capacity and Productivity of Peptide and Protein Purification by Flash Chromatography. Bopanna NK, Chitra Sundararajan, Melissa Wilcox, <u>Janine Sinck</u>, Reno Nguyen, Grace Discovery Sciences, Deerfield, IL, USA
- Isomeric Separation of Procainamide Labeled N-glycans by Using Novel Superficially Porous Particle HILIC Column. Shujuan Tao<sup>1</sup>, Yining Huang<sup>1</sup>, Barry Boyes<sup>2</sup>, Ron Orlando<sup>1</sup>, <sup>1</sup>CCRC, University of Georgia, Athens, GA, USA; <sup>2</sup>AMT, Wilmington, DE, USA
- Two Dimensional Separations: Which Dimension Plays the Most Important Role in Protein Identification Efficiency? Darryl Johnson<sup>1</sup>, Barry Boyes<sup>2</sup>, Ron Orlando<sup>1</sup>, <sup>1</sup>CCRC, University of Georgia, Athens, GA, USA; <sup>2</sup>AMT, Wilmington, DE, USA

## FREE VENDOR SEMINARS

Free Vendor Seminar sponsored by WATERS - Wednesday, July 17 @ 12:00-1:30PM

Free Vendor Seminar sponsored by SHIMADZU - Thursday, July 18 @ 12:00-1:30PM Must register with Shimadzu in order to attend as seating is limited SIGN UP BY CONTACTING https://www.surveymonkev.com/s/K5R8NVL

## Turning Mountains into Molehills: The New Landscape of Protein Analysis

Turn your 24 hour tryptic digests into 1 minute digests. Turn your web browser into a world class bioinformatics platform. Turn your MegaDalton sized protein complexes into measurable ions. This workshop will provide your organization with the facts and critical lessons you need to know for the successful adoption and integration of ultra-fast protein digestions, ultra-fast mass spectrometry, powerful new cloud-based informatics in your laboratory and ultra-high mass detection. Please join us for this interactive workshop where you will discover new Perfinity integrated Digestion Platform (iDP) technologies that enable automated, reproducible protein digests on a sub-minute timescale, greatly improving the applicability of peptide based SRM assays. Cloud-based informatics solutions that provide a unique opportunity to improve not only laboratory data storage, but also enable on-cloud workflow execution and project sharing will also be presented. Finally, the additional challenge of analyzing high molecular weight protein complexes and aggregates with the AXIMA MegaTOF will be discussed. These tools will save your laboratory time, money and resources and will greatly improve data sharing with collaborators.

Free Vendor Seminar sponsored by KNAUER - Thursday, July 18 @ 12:00-1:30PM Must register with Knauer in order to attend as seating is limited SIGN UP BY CONTACTING <a href="mailto:nickel@knauer.net">nickel@knauer.net</a>

# Contichrom®: A Versatile Purification Platform for Batch/CaptureSMB/MCSGP Biochromatography

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# List of Exhibiting Organizations

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