



## **HUPO 2018**

**17th Annual World Congress of the Human  
Proteome Organization**

**September 30 to October 3, 2018**

Loews Royal Pacific • Orlando, FL • USA

**HPP Post-Congress Day, October 4**

# **FINAL PROGRAM**



# MOLECULAR & CELLULAR PROTEOMICS

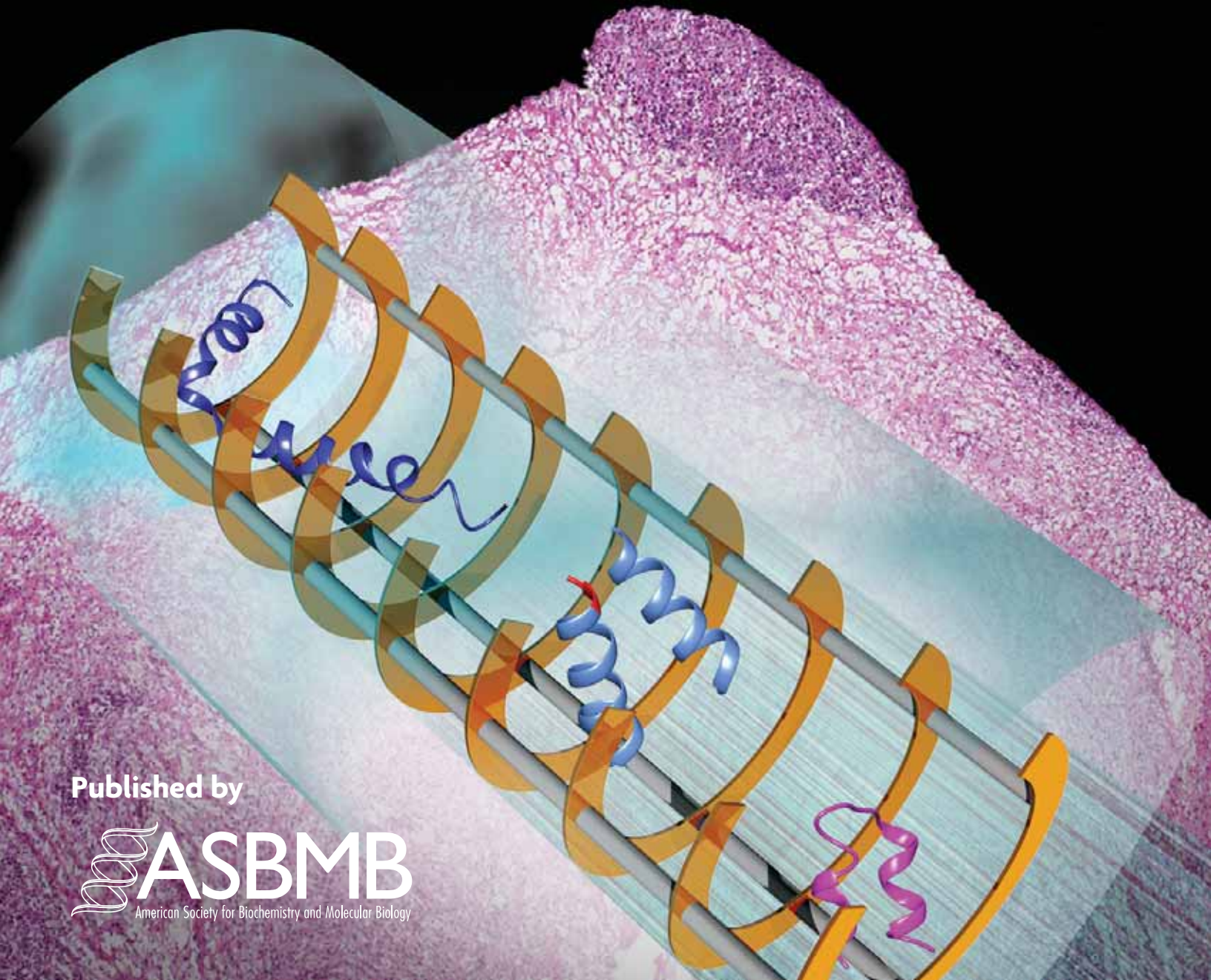
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**17th Human Proteome Organization World Congress**  
**Orlando, Florida • September 30 - October 3, 2018**  
 HPP Post-Congress Day, October 4

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*The HUPO 17<sup>th</sup> Annual World Congress is co-sponsored by US HUPO and HUPO.*



## WELCOME

Welcome to the 17th World Congress of the Human Proteome Organization (HUPO) in Orlando, Florida. We are honored to serve as your hosts for the 2018 HUPO meeting that we hope will excite, enlighten, intrigue, stimulate, motivate and encourage you.

Since the beginning of HUPO, we have all witnessed the evolution of proteomics, certainly the most interesting (and arguably the most challenging) field of biological research, which takes as its mission the study and understanding of all proteins, the machines of life, and strives to lead these discoveries towards One-Health, the integrated notion of wellbeing (the interface of nutrition, environment and humans). In the early years, our field focused primarily on developing the best technologies and chemistries to measure proteins. Its focus then evolved into a discipline establishing methods and standards that were robust and reproducible from laboratory to laboratory. Now, proteomics is everywhere, developing into routine tools in mainstream biology, that cover biology's depth and breadth, from agriculture to cancer and from specialty to high impact journals.

As we developed this program, we have sought to embrace two complementary notions. First, we view this annual gathering as our opportunity to update one another about the latest and most advanced findings in our field. To that end, we have identified 30 different sessions of the most exciting topics where proteomics is advancing our understanding. These sessions include invited speakers representing the most proficient cutting-edge leaders in our discipline, as well as speakers working on provocative projects selected from the submitted abstracts. The sessions cover a wide range of topics, including various diseases, technology advances, and novel scientific and analytical approaches. Our goal to establish a program of outstanding speakers who make it difficult for you to choose which talk to attend is a success. We are proud of the results and we hope you also leave with the same pride.

Second, we recognized that this was an opportune moment to begin to look beyond where proteomics is now, and think about where we should go in the future. We therefore reached outside our community and brought in brilliant scientists, working at exciting interfaces of biology not typically associated with HUPO, and whom we feel will stimulate new associations with proteomics. We want to hear about these new disciplines and begin the discussion about how proteomics could enhance them.

Of course, the most important part of this Congress is the interpersonal meetings in both formal and informal settings, to share the latest data, build new collaborations, discover new products and catch up with friends. To enhance this element, we selected this venue to provide everything under one roof – housing, meetings, posters and vendor booths. Please join us in the social activities, designed for having fun, including a visit to Universal Studios and a themed banquet.

Finally, it takes a village to plan such a meeting. We are deeply indebted to the many people who helped us plan this event, recommend and encourage new exciting speakers, reviewing abstracts, contacting vendors, managing logistics, attending numerous meetings and writing countless emails. We all thank them for their time and commitment.

As your hosts, we wish you a magical and productive 17th Annual HUPO congress.

### HUPO 2018 CONGRESS CHAIRS



**Joshua LaBaer**  
*Biodesign Institute,  
University of Arizona*



**Ileana Cristea**  
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**Robert Moritz**  
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## GENERAL INFORMATION

Welcome to the HUPO 17<sup>th</sup> Annual World Congress. All Congress activities (talks and posters) and exhibit booths are in the Loews Royal Pacific Hotel.

- **PLENARY AND PARALLEL ORAL SESSIONS** are located in the Oceana Ballroom complex
- **CORPORATE LUNCH SEMINARS AND EVENTS** may be found in the Exhibitor listing, pages 12 – 18.
- **POSTERS AND EXHIBIT BOOTHS** are in the Pacifica Ballroom.
- **INNOVATION STAGE** is located in the Pacifica Ballroom with the exhibits and posters.
- **REGISTRATION** is open 8:00 am - 7:00 pm Sunday, 7:30 am - 5:00 pm Monday - Wednesday.

**Free wifi** is available look for signs at the congress with sign-in details.

**PROGRAM CODES** for talks are as follows:

M, T, W = Day

O = Oral

A, B, C, D, E = Session

Time

For example, MOA am 9:10 = Monday Oral Session A at 9:10 am.

For posters 'Poster' is followed by board space.

**SPEAKERS** must load presentations **one day before** your talk in the Speaker Ready Room (Oceana 11). In the interest of preserving the time schedule and as a courtesy to other speakers, **no presentations may be loaded in the session rooms**. The Speaker Ready Room is open with a technician at these times:

Sunday, 1:00 - 6:00 pm

Monday - Tuesday, 8:00 am - 5:00 pm

Wednesday, 8:00 am - 2:00 pm

**POSTER PRESENTERS** must set up by 9:15 am Monday morning and remove posters at 10:30 am on Wednesday. All posters will be displayed Monday - Wednesday. **Refer to the poster numbers in this final program for board assignments**. Push pins are available at the Poster Supplies counter inside the Pacifica Ballroom.

All poster authors are encouraged to be present for the daily poster session 9:15 - 10:30 am and 3:50 - 5:00 pm according to this schedule:

- Odd-numbered boards present on Monday
- Even-numbered boards present on Tuesday
- All presenters on Wednesday morning.

**BIOINFORMATICS HUB** will have programs Sunday through Wednesday noon in Pacifica Ballroom. For details visit <https://github.com/CompMS/Overview/wiki/HUPO-2018>



**OPENING RECEPTION, SUNDAY, 7:15 - 9:00 PM** is in the Pacific Ballroom. Congress name badge is required.

**LUAU SOCIAL EVENT, 6:30 - 8:30 PM, TUESDAY.** Advanced ticket purchase required. Tickets may be purchased at registration through 12:00 pm, Monday.

**HALLOWEEN HORROR @ UNIVERSAL, 6:30 PM, WEDNESDAY.** Universal park is open for special Halloween Horror night on Wednesday. We have \$60 tickets available for purchase at registration counter. Meet up 6:30 pm on Wednesday to go to the park as a group. Halloween Horror night will have 'frights' throughout the park with some special haunted house. Harry Potter attractions within Universal will be fully functional and are non-fright! Join us to blow off some post-congress steam.

### CONGRESS REGULATIONS

- **Name badges** are required for all conference sessions, including the exhibit hall.
- **No smoking** is permitted inside the hotel.
- **Cell phones** must be **turned off** in oral sessions.
- **No photography or recording** in any session, including posters.
- **The placement of advertising** in the meeting area is prohibited.

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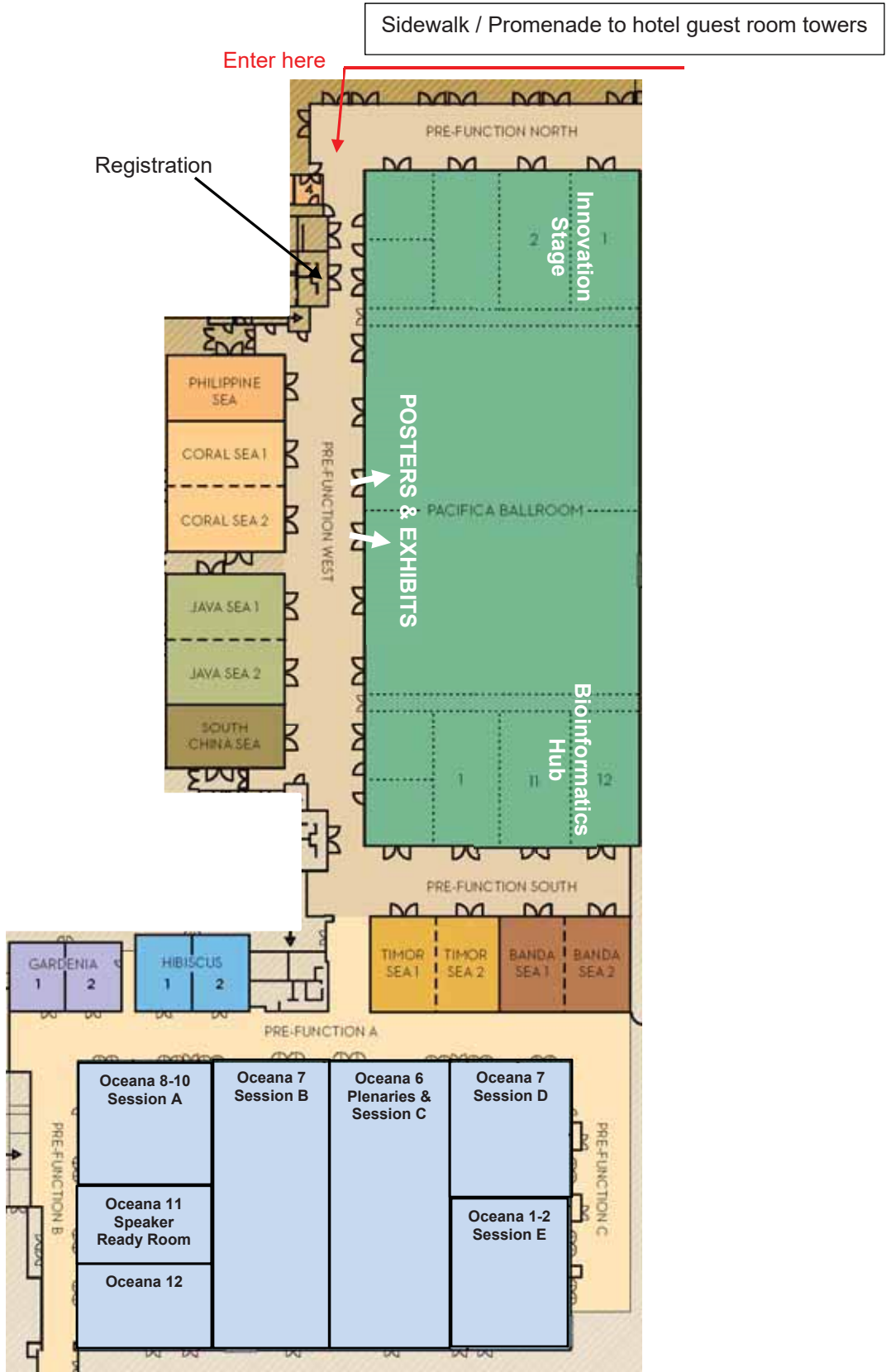
- register at <http://med.stanford.edu/hpop.html>
- email us at [sahadi@stanford.edu](mailto:sahadi@stanford.edu)
- find us at the hPOP table in the Posters-Exhibits

Participants can be one-time only  
(no commitment needed for future sampling).

We also welcome and encourage  
return participants.

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## HUPO AWARDS

### DISTINGUISHED ACHIEVEMENT IN PROTEOMIC SCIENCES

Sponsored by Journal of Proteome Research (ACS Publications)



#### Kathryn Lilley

*University of Cambridge, UK*



Dr. Lilley provides leading efforts in developing technologies to enable the measurement of proteome dynamics in a high throughput manner in space and time during cellular processes. These include methods for defining dynamic protein and protein-complex sub-cellular location called "Localization of Organelle Proteins using Isotope Tagging" (LOPIT) and more recently, hyperLOPIT. Other methods developed by Dr. Lilley encompass mapping the location of translation of proteins and their control; Multi protein complexes with novel protein-protein interaction methods using Parallel Affinity Capture (iPAC); and Selective Proteomic Proximity Labeling using Tyramide (SPPLAT), a method that enables the identification of proteins in the immediate vicinity of a target membrane protein. These methods are underpinned by robust statistical and computational data analysis tools developed by Dr. Lilley in an open-source software environment. Applying these tools to model systems of drosophila, systematic analysis of how chemicals poison bacterial cells, and studying the RNA binding proteome, provide solid demonstrations of the methods developed by Dr. Lilley. Few proteomics researchers have contributed comprehensive methods to study proteomes as has Dr. Lilley, and we are proud of her distinguished achievements to date for the 2018 Distinguished achievement award.

### SCIENCE AND TECHNOLOGY AWARD

Sponsored by HUPO Industrial Advisory Board

#### John Syka, Jae Schwartz, Lee Earley and Christopher Mullen

*Thermo-Fisher Scientific*



Drs. Syka, Schwartz, Earley and Mullen have all played a significant role in the development and commercialization of Electron Transfer Dissociation on state-of-the-art mass spectrometers from its early development at the U. Virginia by Dr. Coon and Dr. Hunt. ETD is a nonergodic type of dissociation that enables sequencing of labile post translational modifications on proteins and peptides. The ease of use of the commercial ETD device developed by the team at Thermo-Fisher has enable many researchers from around the world to access this technology in a technically simple device and straightforward application simultaneously, or concurrently, with standard CID fragmentation. ETD continues to play significant roles in advancement of research in the detection of post-translational modifications such as phosphorylation and glycosylation. Additionally, ETD is a crucial fragmentation technology that enables rich fragment and subsequent sequence information in Middle down and Top-down proteomics. This significant innovation in the proteomics field continues to have wide acceptance and also provides a springboard for proteomics technology to impact the field of translational clinical proteomics. The HUPO members applaud the efforts of the industrial team from Thermo-Fisher of Drs. Syka, Schwartz, Earley and Mullen resulting in readily available novel ETD fragmentation technology for the 2018 HUPO award in Science and Technology.

## HUPO AWARDS

### DISCOVERY IN PROTEOMIC SCIENCES

(shared by two recipients)

Sponsored by Journal of Proteomics (Elsevier BV)



**Ulrike Kusebauch**

*Institute for Systems Biology, USA*

Dr. Kusebauch, a native of Germany, has performed pioneering efforts for the establishment of the complete Human SRMAtlas. This achievement, led by Dr. Kusebauch and encompassing a large collaborative team from around the globe, provides a highly curated database of over 160,000 proteotypic peptide fragmentation spectra, their performance characteristics, selected reaction monitoring (SRM) assays, and a data-rich web based resource for the entire human proteome. This resource, generated on multiple mass spectrometry instruments, includes multiple fragmentation parameters and chromatographic traces for each peptide to enable any researcher to establish targeted identification and quantification of any accessible protein of the human proteome. This proteome-centric database is extensively integrated with bioinformatics knowledge bases and resources to provide an information rich resource for human proteome research. The SRMAtlas resource has extensive evidence of accessible assays for the human proteome covering proteins from all chromosomes with high quality quantitative SRM profiles to underpin HUPO HPP efforts in determining evidence for all proteins in the Human Proteome. This significant achievement provides a component of the MS Pillar substantiating the HPP efforts of HUPO and cuts across all other programs within HUPO (i.e., C-HPP, B/D-HPP etc.) making this resource also an achievement for HUPO. In addition, many biomarker evaluation projects worldwide have been accelerated with the use of this resource by providing readily available SRM assays, visual inspection of each assay, and assay parameter instrument files that are deployed in discovery and verification studies, ultimately fast tracking this process. She published this work in the journal *Cell* in 2016, and the SRMAtlas resource is freely available via the web. Dr. Kusebauch is a long-term HUPO member and active in proteomics for many years and these efforts by a young, talented female researcher being awarded is a demonstration of the talent in human proteome research that HUPO strives to support and recognize. The Human Proteome Organization congratulates Dr. Kusebauch, a worthy co-winner of the 2018 award for Discovery in Proteomics.



**Joshua Coon**

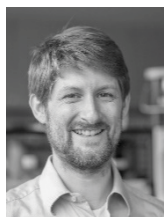
*University of Wisconsin, USA*

Dr. Coon has been at the forefront of discovery research and the tools he has created have gained wide use around the world. His research program is to facilitate, expedite, and comprehensively analyze proteins and metabolites by innovating new mass spectrometric technologies and apply these techniques to advance biomedical research. Dr. Coon has contributed heavily to aspects of proteomics and metabolomics research developing next generation instrumentation and instrument methods, proteomics workflows, novel isotopic labeling quantitative approaches, and associated software development that has influenced many labs around the globe. Specifically, these include; the development electron transfer dissociation (ETD), which is a complementary nature of ion fragmentation, where ETD provides new avenues for analyzing intact proteins, post-translational modifications, and structural aspects of the proteome. Dr. Coon started his independent program in 2005, shortly after co-developing ETD with Prof. Don Hunt (U. Virginia) and John Syka (Thermo-Fisher) and went on to apply the ETD technique to a number of biological questions with great success. Dr. Coon was also awarded the American Society for Mass Spectrometry's Biemann Medal in 2012 for this work. Other innovations supporting discovery research include: describing Parallel Reaction Monitoring (PRM), a method for a high resolution/accurate mass analyzer to permit the parallel detection of all target product ions in one, concerted high resolution mass analysis extending the specificity of SRM analysis; the GC-Orbitrap construction for identification of unknown metabolites from high-resolution, high mass accuracy discovery GC-MS experiments; novel proteome quantification methodology using metabolic labeling called NeuCode; and multi-omic data analysis and visualization to understand metabolism. We applaud the efforts of new HUPO member Dr. Coon, a well-deserved co-winner of the Discovery in Proteomics award for 2018.

## HUPO AWARDS

### CLINICAL AND TRANSLATIONAL PROTEOMICS (shared by two recipients)

Sponsored by Clinical Proteomics (BioMed Central)



**Bernd Bodenmiller**  
*University of Zurich, Switzerland*

Dr. Bernd Bodenmiller performs pioneering development work in Clinical and Translational Proteomics for single cell proteomic analyses, one of the transforming technologies in the emerging field of "Precision Medicine". Precision Medicine is one of the most disruptive and exciting developments in the field of clinical and translational proteomics and, in fact, in clinical research as a whole. Without innovative technologies to provide a sound experimental approach, Precision Medicine is just an empty term. Therefore, this new field of research necessitates approaches that enable a comprehensive analysis and visualization of the diseased tissue, ranging from molecules to networks and phenotypes, to cell-to cell interactions and to tissue morphology. Such analytical approaches will also support understanding of tissue biology, to define biomarkers and to identify novel routes of therapy. Dr. Bodenmiller and his group have pioneered the quantification of panels of proteins at the single cell level in cells in solution and in tissue based on mass cytometry, a technique that uses metals and inductively coupled plasma mass spectrometry to measure markers simultaneously with subcellular resolution. With the development of the most powerful targeted tissue imaging method in existence, called imaging mass cytometry (IMC), IMC currently enables to image 52 and soon >100 selected proteins, protein modifications and transcripts simultaneously with subcellular resolution in tissues. In other studies, the group of Dr. Bodenmiller showed how protein overexpression, a known driver of cancer initiation and progression, impacts single cell signaling networks. This work suggests why deregulation of the MAPK pathway renders it (unexpectedly) resistant to targeted inhibition. We applaud the efforts of Dr. Bodenmiller for his translational proteomic efforts to benefit human health, a well-deserved co-winner of the Clinical and Translational Proteomics award for 2018.



**Peipei Ping**  
*University of California Los Angeles, USA*

Dr. Peipei Ping is an extraordinary scientist who has made seminal contributions in the field of Clinical and Translational proteomics science. Dr. Ping is currently a Professor in the Departments of Physiology, Medicine, and Bioinformatics at UCLA. She was the recipient of HUPO Distinguished Service Award in 2013. Dr. Ping is a visionary leader who has made a profound impact in both proteomics and cardiac physiology communities with her keen leadership and devotion. She is internationally recognized for her expertise and achievements in proteomics, mitochondrial proteome biology, systems biology, and data science. Dr. Ping has a highly innovative research program and her lab has pioneered several major advances that have shifted paradigms in mitochondrial and protein degradation research. Notably, Dr. Ping has devoted her past decades to translational research, which has advanced the field of cardiology, her major focus. With the development of techniques and data resources specifically to tackle clinical aspects of cardiology research, Dr. Ping has advanced proteomics clinically in this translational field. Her work across many aspects of translational research has provided new critical mechanistic insights in cardiac hypertrophy, a clear advancement of the field of cardiology. We congratulate Dr. Ping for her outstanding achievements culminating in co-sharing the Clinical and Translational Proteomics award for 2018.

## OTHER AWARDS



**2018 Recipient of the MCP Lectureship: Carol V. Robinson**

*Molecular & Cellular Proteomics*, an official publication of the American Society for Biochemistry and Molecular Biology, introduced its sponsored lectureship series as part of its 10th anniversary celebration in 2011. Each lecturer is a leader in the field of proteomics who presents his or her particular research and interests, with the intent to expand on proteomics' potential to ask (and answer) increasingly complex questions associated with health, energy, food supply and the environment. The lectureships are given at germane meetings and symposia throughout the year, and the lecturers are chosen by the organizers of those meetings. Each lecturer is presented with a crystal plaque to commemorate the occasion.

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## EXHIBITORS

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 <h1 style="margin-left: 20px;">Agilent</h1>	<p><b>Booth 301</b>          Agilent Technologies is a worldwide provider of GC, LC, MS and Spectroscopy instruments, technologies, related consumables, support, services, and workflow solutions that enable labs to analyze, confirm and quantify substances of interest with confidence while maintaining the most stringent laboratory practices, from sample preparation to final report.</p> <p><b>AGILENT MONDAY LUNCH SEMINAR</b>, 12:20 - 1:45 pm, <i>South China Sea Room</i>          Pre-register online or at booth.</p> <p><b>Agilent Innovations for Proteomics Research</b>; Christine Miller, <i>Omics Market Manager</i>, and Randy Bolger, <i>Workflow Solutions Manager</i>          Join us for a two-part presentation on proteomics research. Our first presentation will describe proof-of-principle results demonstrating the value of combining metabolomics and proteomics analysis for better biological understanding. We will highlight the tools used in this approach. Our second presentation will discuss automation to improve sample preparation of complex proteomics samples while improving reproducibility and consistency across longitudinal studies.</p>
 <p style="text-align: center;">NEXT GENERATION PROTEOMICS</p>	<p><b>Booth 407</b>          Biognosys is dedicated to developing superior proteomics solutions and offers these solutions to researchers as products and services. These solutions rely on mass spectrometry with parallel data-independent acquisition (DIA), which allow simultaneous quantification of thousands of proteins in a single experiment. Our next generation technology provides multiplexed protein quantification with high precision and depth.</p> <p><b>BIOGNOSYS-EVOSEP TUESDAY LUNCH SEMINAR</b>, 12:20 - 1:45 pm, <i>Banda Sea Room</i>          Pre-register online or at booth.</p> <ul style="list-style-type: none"> <li>• <b>Robust Clinical Proteomics Workflow for Plasma and Cancer Tissue Analysis</b>; Matthias Mann, <i>Max Planck Institute for Biochemistry</i>,</li> <li>• <b>Clinical Proteomics for Biomarker Qualification in Fabry Disease</b>; Petra Oliva, <i>Sanofi Genzyme</i></li> </ul>
	<p><b>Booth 302</b>          BSI is well known for its PEAKS software suite and service platform for proteomics discovery and therapeutic protein characterization by LC-MS. The benchmark de novo sequencing capabilities offers advanced solutions for proteomic and therapeutic protein discovery as provided through peptide/protein identification &amp; quantification, peptide mapping, PTMs and sequence variants.</p>

## EXHIBITORS

	<p><b>Booth 400</b>          For more than 55 years, Bruker has enabled scientists to make breakthrough discoveries and develop new applications that improve the quality of human life. Bruker's high-performance scientific instruments and high-value analytical and diagnostic solutions enable scientists to explore life and materials at molecular, cellular and microscopic levels.</p> <p><b>BRUKER EXCEED SUNDAY (USER MEETING)</b>, 3:30-5:45 pm, <i>Timor Sea Room</i>          Pre-register online.</p> <p><b>BRUKER MONDAY LUNCH SEMINAR</b>, 12:20 – 1:45 pm, <i>Timor Sea Room</i>          Pre-register online or at booth.</p> <ul style="list-style-type: none"> <li>▪ <b>Introduction to the timsTOF Pro powered by PASEF</b>; Oliver Raether, <i>QTOF R&amp;D Manager, Bruker Daltonik</i></li> <li>▪ <b>Data-independent Parallel Accumulation – Serial Fragmentation (diaPASEF) on the tims – TOF Pro</b>; Matthias Mann, <i>Max Planck Institute of Biochemistry</i></li> </ul> <p><b>BRUKER TUESDAY LUNCH SEMINAR</b>, 12:20 – 1:45 pm, <i>Timor Sea Room</i>          Pre-register online or at booth.</p> <ul style="list-style-type: none"> <li>▪ <b>Our time with PASEF on the timsTOF Pro</b>; Tharan Srikumar, <i>Princeton University</i></li> <li>▪ <b>Urine Biomarker Discovery by Proteomics and Peptidomics: Towards "All-in-One Urine Test"</b>; Tadashi Yamamoto, <i>Niigata University</i></li> </ul>
 <p>Cambridge Isotope Laboratories, Inc.  <b>isotope.com</b></p>	<p><b>Booth 502</b>          Cambridge Isotope Laboratories, Inc. (CIL) is the world leader in the manufacture of stable isotope-labeled compounds (<sup>13</sup>C, D, <sup>15</sup>N, <sup>17</sup>O, <sup>18</sup>O) used for qualitative and quantitative, MS-based 'omics applications. Key innovative products include ProteusQC™ (for QC and quantitation), Mouse Express® mouse feed and mouse tissue, 99% enriched amino acids, SILAC and SILAM reagents, Fmoc protected amino acids and preloaded resins (for peptide synthesis), dimethyl labeling reagents, <sup>18</sup>O water (for enzymatic labeling), heavy-labeled proteins, INLIGHT™ (for glycan tagging), heavy labeled glycans, and PeptiQuant™ Plus MRM Assay Kits.</p>
 <p><b>CLINICAL          PROTEOMICS</b></p>	<p><b>Tabletop</b>          Placing an emphasis on the application of proteomic technology to all aspects of clinical research and molecular medicine, <i>Clinical Proteomics</i> provides a scholarly forum for novel scientific research across the broad spectrum of clinical and translational proteomics. The journal is committed to rapid scientific review and the timely publication of manuscripts.</p>
	<p><b>Booth 606</b>          Covaris is the recognized industry leader in NGS, utilizing its patented Adaptive Focused Acoustics® (AFA®) technology for DNA fragmentation. AFA-energetics™ is also used for a wide range of sample preparation applications including FFPE and cfDNA extraction, chromatin shearing, proteomics, epigenomics, cell lysis, and compound management.</p>
	<p><b>Tabletop</b>  <i>Journal of Proteomics</i> is published by Elsevier. Empowering Knowledge: Elsevier is a global information analytics business that helps institutions and professionals advance healthcare, open science and improve performance for the benefit of humanity.</p>

## EXHIBITORS

 <p>ESI Source Solutions</p>	<p><b>Booth 406</b> ESI Source Solutions provides unique products for researchers using capillary chromatography and nanospray interfaces to mass spectrometers in the field of proteomics, metabolomics and others. Located in Woburn, MA, we offer products like the ABIRD and CPS designed to optimize your current systems to improve overall reliability and sensitivity.</p>
	<p><b>Booth 503</b> Evosep aims to improve quality of life and patient care by radically innovating protein based clinical diagnostics. Making sample preparation and separation before MS analysis 10 times faster and 100 times more robust will enable truly large cohort studies for biomarker validation and provide the foundation for precision medicine.</p> <p><b>EVOSEP-BIOGNOSYS TUESDAY LUNCH SEMINAR, 12:20 - 1:45 pm, Banda Sea Room.</b> Pre-register online or at booth.</p> <ul style="list-style-type: none"> <li>• <b>Robust Clinical Proteomics Workflow for Plasma and Cancer Tissue Analysis;</b> Matthias Mann, <i>Max Planck Institute for Biochemistry</i>,</li> <li>• <b>Clinical Proteomics for Biomarker Qualification in Fabry Disease;</b> Petra Oliva, <i>Sanofi Genzyme</i></li> </ul>
	<p><b>Booth 206</b> HUPO19 is on the other side of the world and hosted by the Australasian Proteomics Society in Adelaide. Come and meet some of the team at our stand and we will help you organise your participation next year.</p>
	<p><b>Booth 306</b> INTAVIS Bioanalytical Instruments focuses on automation of complex scientific protocols. We concentrate on applications in proteomics, and functional genomics. The DigestPro MSi, is a dedicated platform for protein digestion and sample preparation in proteome research. It automates both in-gel and solution digest procedures for protein analysis by mass spectrometry.</p>
	<p><b>Tabletop</b> <i>Journal of Proteome Research</i> publishes all aspects of global protein analysis and function, including the dynamic aspects of genomics, spatio-temporal proteomics, metabolomics, clinical and agricultural proteomics, and methodological advances including bioinformatics. The emphasis is on a multidisciplinary approach to the life sciences through the synergy between the various "omics".</p>
	<p><b>Booth 504</b> JPT is an ISO certified provider of innovative peptide products and services focusing on proteomics, novel immune therapeutics &amp; diagnostics and protein profiling. JPT has a large portfolio of proprietary products such as SpikeMix™ Peptide Pools &amp; SpikeTides™ that provide access to economical stable isotope labeled and/or quantified peptide standards.</p>
	<p><b>Booth 203</b> Korea Basic Science Institute (KBSI) performs a role as a national institute with the world-class basic research infrastructure leading the innovation of analytical platforms and equipments. Now, we are introducing an automated software platform (IQ-GPA) developed for identification and quantification of intact N- and O-linked glycopeptides from mass spectrometry data.</p>
	<p><b>Booth 307</b> Take the guesswork out of protein identification with Mascot Server, the benchmark for database search. Get closer to your raw data with Mascot Distiller, giving direct access to all popular files formats for peak picking, de novo sequencing, quantitation, and more.</p>



## EXHIBITORS

	<p><b>Tabletop</b> The mission of <i>Molecular &amp; Cellular Proteomics</i> is to foster the development and applications of proteomics in basic and translational research. MCP publishes studies reporting significant biological or clinical discoveries underpinned by proteomic observations across all kingdoms of life. MCP also emphasizes articles describing innovative computational methods and technological advancements that enable future discoveries.</p>
	<p><b>Booth 201</b> Metabolon is the world's leading health technology company advancing metabolomics for precision medicine and every area of life sciences research. Our Precision Metabolomics™ is a powerful technology for assessing health and is delivering biomarker discoveries, innovative diagnostic tests, and ground-breaking data in genomics and population health initiatives.</p>
	<p><b>Booth 205</b> <i>Molecular Omics</i> is a new high quality journal for the –omics sciences, published by the Royal Society of Chemistry (a not-for-profit Publisher). It focuses on molecular level experimental and bioinformatics research in the –omics sciences. We especially welcome multidisciplinary papers that present studies combining different types of –omics, or the interface of –omics and other fields.</p>
	<p><b>Booth 200</b> MRM Proteomics Inc. specializes in precision proteomics technology, with the mission of delivering the highest quality services and products. We offer services for protein quantitation, structural characterization, tissue imaging, metabolomics, and clinical diagnostics, plus a line of easy-to-use MRM-MS kits. We also offer unique custom-tailored solutions for specific research projects.</p>
	<p><b>Booth 602</b> Olink Proteomics provides innovative solutions for targeted human protein biomarker discovery. Proseek® Multiplex enables rapid, high-throughput immunoassay analysis, exceptional data quality, and minimal sample consumption. Disease-focused panels allow simultaneous analysis of 92 biomarkers with 1 µl of sample, and are available as kits or via our Analysis Service.</p>
	<p><b>Booth 507</b> PharmaFluidics is a disruptive player in the field of micro-Chip chromatography for biomarker discovery and the analytical development of biopharmaceuticals. Our expertise is the lithographic design and surface treatment of silicon wafers for the development of analytical chromatography columns (µPAC™) with a separation bed of highly ordered and free-standing pillars.</p>
	<p><b>Booth 304</b> PreOmics is as a spin-off company from the proteomics research group of pioneer Matthias Mann at the Max-Planck Institute of Biochemistry. We at PreOmics develop and provide innovative technologies for mass spectrometry-based proteomics to make your research simply better. Our mission is to set the standard for MS-based protein analysis.</p>
	<p><b>Booth 505</b> Pressure BioSciences, Inc. (OTCQB: PBIO) is a leader in the development and sale pressure-based platform solutions for the life sciences industry. Our products and services are based on the unique properties of our patented, pressure-enhanced platforms: Pressure Cycling Technology (“PCT”) and Pressure Enabled Protein Manufacturing Technology (“PreEMT”). The PCT Platform is used in biomarker and target discovery, soil &amp; plant biology, anti-bioterror, and forensics. The PreEMT Platform employs high pressure for the disaggregation and controlled refolding of proteins.</p>

## EXHIBITORS

	<p><b>Booth 305</b>          Swiss company Prolab Instruments has been perfecting Micro- and Nano-UHPLC technology for over 20 years. Their Zirconium UHPLC pumps sets offer highest precision gradient control, huge dynamic flow ranges, fast equilibration (meaning shortest cycle times), and a host of flexible system integration options, catering both OEM and end users.</p> <p><b>PROLAB TUESDAY TALK @ INNOVATION STAGE, 9:45 – 10:00 am</b></p>
	<p><b>Booth 207</b>          Proteome Software sets the standard with our intuitive, high-quality proteomics, metabolomics and small molecule analytic software. From customer-focused design and development to specific, clear technical support and documentation, our dedicated team strives to create a satisfying user experience. Researchers and core labs around the world rely on Proteome Software.</p>
	<p><b>Booth 506</b>          Resyn Biosciences. Advanced magnetic microsphere reagents and solutions for automation of routine mass spectrometry workflows, increasing reproducibility for improved data quality.</p>
	<p><b>Booth 500</b>          SCIEX delivers advanced analytical technologies and software that contribute to the understanding and research of human disease. Innovative LC-MS, LC-MS/MS and CE solutions enable deeper analysis of complex biological systems by providing comprehensive quantitation and characterization required across proteomics, lipidomics and metabolomics — leading to advances in systems biology and biomarker discovery.</p> <p><b>SCIEX MONDAY LUNCH SEMINAR, 12:20 - 1:45 pm, Coral Sea Room</b>          Pre-register online or at booth.  <b>Quantitative Proteomics Profiling – SWATH® Acquisition as a Tool for the Research and Service Core Lab</b>          Yansheng Liu, <i>Yale University</i>, and Birgit Schilling, <i>Buck Institute for Research on Aging</i></p> <p><b>SCIEX INNOVATION SOCIAL HOUR, 6:00 – 8:00 pm, Coral Sea Room</b>          Pre-register online or at booth.          Reception, Alternative Fragmentation Technology for Proteomics Applications Presentation, and Interactive Poster Session, <i>Featuring Yves LeBlanc, SCIEX R&amp;D</i></p> <p><b>SCIEX TUESDAY LUNCH SEMINAR, 12:20 - 1:45 pm, Coral Sea Room</b>          Pre-register online or at booth.  <b>The TripleTOF® System in the Lab – Practical Applications and Exciting Innovations, Featuring Christie Hunter, SCIEX</b></p>
	<p><b>Booth 300</b>          Shimadzu is the leading provider of analytical measurement and testing instrumentation for a broad range of applications in science and industry. With the release of Shimadzu's LCMS-9030 Q-TOF and microflow HPLC, Mikros, along with automated Perfinity Workstation for proteomic workflows, Shimadzu brings unparalleled solutions for proteomic and systems biology research.</p>

EXHIBITORS

	<p><b>Booth 600</b> Synpeptide is the leading provider of custom peptide synthesis services and peptide modifications, especially for stable isotope labeled peptides, glycopeptides and glycosylated peptides, phosphorylated peptides, peptide libraries and so on. Pleased to be your partner.</p>
	<p><b>Booth 401</b> Thermo Fisher Scientific is the world leader in serving science. Our mission is to enable our customers to make the world healthier, cleaner and safer. Through our Thermo Scientific and Invitrogen brands, we help customers accelerate innovation and enhance productivity.</p> <p><b>THERMO MONDAY LUNCH SEMINAR, 12:20 – 1:45 pm, Java Sea Room</b> Pre-register online or at booth. <b>Latest Advances in Multiplexing Technology:</b></p> <ul style="list-style-type: none"> <li>• Establishing a Roadmap for Brain-based Protein Biomarkers in Alzheimer's Disease; Nicholas Seyfried, <i>Emory School of Medicine</i></li> <li>• Drug Effects on Protein Homeostasis; Marcus Bantscheff, <i>Cellzome, a GSK Company</i></li> </ul> <p><b>THERMO TUESDAY LUNCH SEMINAR, 12:20 – 1:45 pm, Java Sea Room</b> Pre-register online or at booth. <b>Precision Medicine:</b> Advancing Mass Spectrometry-based Large-cohort Proteomics for Precision Medicine – An International Cancer Moonshot Multi-site Study; Thomas Conrads, <i>Inova Schar Cancer Institute</i>; Yue Xuan, <i>Thermo Fisher Scientific</i></p> <p><b>THERMO WEDNESDAY LUNCH SEMINARS, 12:20 – 1:45 pm, Java Sea Room and South China Sea Room</b> Pre-register online or at booth.</p> <p><b>MS Toolbox for Systems Biology:</b> Understanding Interactions in Membrane Proteins – New Opportunities for Drug Discovery; Carol Robinson, <i>University of Oxford</i></p> <p><b>MS Targeted Assays:</b> New Quantitative Proteomic Assays for Cancer Signaling Pathways Using Multiplex IP and Targeted Mass Spectrometry; Jonathan Krieger, <i>The Hospital for Sick Children</i>, and Bhavin Patel, <i>Thermo Fisher Scientific</i></p>
	<p><b>Booth 404</b> PeakInvestigator® software revolutionizes MS analyses, delivering unprecedented quality and revealing features hidden in your raw profile data from ion trap, TOF, Orbitrap or FTICR-based instruments. PeakInvestigator's advanced signal processing algorithms dynamically differentiate peaks from noise, increase spectral resolution (up to 5-6x), and provide statistical confidence intervals on every peak.</p>

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## Booth 501

Waters Corporation creates business advantages for laboratory-dependent organizations by delivering scientific innovation to enable customers to make significant advancements. Waters helps customers make profound discoveries, optimize laboratory operations, deliver product performance, and ensure regulatory compliance with a connected portfolio of separations and analytical science, laboratory informatics, mass spectrometry, as well as thermal analysis.

### WATERS CORP. TUESDAY ACTIVITIES

#### Q&A Session | Progenesis QI for Proteomics Users, 9:30-10:15 am, *South China Sea Room*

Pre-register online or at booth.

Come join fellow users of the latest version of Progenesis QI for proteomics to gain insight into how Progenesis can help you analyze proteins and characterize peptides. We'll discuss some of the new features in Progenesis Version 4.1 that help you identify the peptides and proteins of interest and learn more about the use of Spectral Libraries.

#### Lunch Seminar, 12:20 – 1:45 pm, *South China Sea Room*

Pre-register online or at booth.

Join us for lunch and see how Waters is addressing the challenges of throughput in clinical proteomic studies with precision and accuracy. SONAR™ on Waters bench top Xevo G2-XS® Qtof brings the speed and selectivity for accurate quantitative analysis of the plasma proteome. High throughput SONAR methods can be combined with quantitative kits such as the Biognosys PQ500™ for accurate absolute quantitation of human biological fluids.

In this lunch seminar we will discuss the multi-omic application of SONAR for a respiratory medicine study.

- **Introduction of Absolute Quantification of Plasma Samples Based on Biognosys' PQ500 Reference Peptides**

*Presented by: Dr. Roland Bruderer, Research and Development, Biognosys*

- **Application of SONAR for enhanced throughput analysis with an exploratory targeted reagent strategy in clinical proteomics research**

*Presented by: Dr. Lee Gethings, Senior Manager Biomedical Research, Waters Corp.*

#### Overview | Progenesis QI for Proteomics, 4:00 - 4:45 pm, *South China Sea Room*

Pre-register online or at booth.

We invite you to get a better understanding of the concepts behind Progenesis QI for proteomics, one of the world-leading proteomics data analysis applications.

- Learn how missing values affects your experiment
- Understand the benefits of Co-Detection and why it is beneficial
- See how Progenesis can investigate peptides
- Learn how to create your own Spectral Libraries

# Clinical Proteomics

Published by BMC, and affiliated with the Human Proteome Organization (HUPO), *Clinical Proteomics* provides a scholarly forum for novel scientific research across the broad spectrum of clinical and translational proteomics.

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## PROGRAM OVERVIEW

<p style="text-align: center;"><b>SUNDAY, SEPT 30</b></p> <p style="text-align: center;">9:00 am - 3:30 pm, <b>HPP Investigators' Day</b> (separate registration required, lunch included)            9:00 am - 3:30 pm, <b>ECR Mentoring Day</b> (separate registration required, lunch included)            9:00 am - 5:45 pm, Bioinformatics Hub (FREE), <i>Pacifica 11-12</i>            3:00 – 5:00 pm, <b>Core Facilities Workshop</b> (FREE), <i>South China Sea Room</i>            3:45 – 5:45 pm, <b>HUPO Council Meeting</b> (by invitation), <i>Hibiscus Room</i>            - CONGRESS BEGINS -            6:00 – 7:15 pm, <b>Plenary Session</b>, Carol Robinson and Beth Anderson, <i>Oceana 6</i>            7:15 – 9:00 pm, <b>Welcome Reception with exhibitors</b>, <i>Pacifica</i></p>		
<p style="text-align: center;"><b>MONDAY, OCT 1</b></p>	<p style="text-align: center;"><b>TUESDAY, OCT 2</b></p>	<p style="text-align: center;"><b>WEDNESDAY, OCT 3</b></p>
<p>8:30-9:15 am, <b>Plenary Session</b>, Stephen Quake, <i>Oceana 6</i></p> <p>9:15-10:30 am, <b>Poster Session</b>, Odd-numbers present, <i>Pacifica</i></p> <p>9:30 am - 5:00 pm, Bioinformatics Hub, <i>Pacifica</i></p> <p>10:30 am-12:20 pm, <b>Parallel Sessions</b></p> <ul style="list-style-type: none"> <li>• Affinity/Proximity (MOA am), <i>Oceana 8-10</i></li> <li>• Infectious Disease (MOB am), <i>Oceana 7</i></li> <li>• Cancer (MOC am), <i>Oceana 6</i></li> <li>• HPP: Partnering w-Pathology Toward Precision Medicine (MOD am), <i>Oceana 3-5</i></li> <li>• Statistics in Experimental Design (MOE am), <i>Oceana 1-2</i></li> </ul> <p>12:20-1:45 pm, <b>Corporate Lunch Seminars</b> (RSVP) or Lunch-on-your-own</p> <ul style="list-style-type: none"> <li>• Agilent, <i>South China Sea</i></li> <li>• Bruker, <i>Timor Sea</i></li> <li>• SCIEX, <i>Coral Sea</i></li> <li>• Thermo, <i>Java Sea</i></li> </ul> <p>2:00-3:50 pm, <b>Parallel Sessions</b></p> <ul style="list-style-type: none"> <li>• Aging (MOA pm), <i>Oceana 8-10</i></li> <li>• Nutrition &amp; Food (MOB pm), <i>Oceana 7</i></li> <li>• PTMs (MOC pm), <i>Oceana 6</i></li> <li>• HPP: Targeting the Proteome in Women's Health (MOD pm), <i>Oceana 3-5</i></li> <li>• Proteogenomics (MOE pm), <i>Oceana 1-2</i></li> </ul> <p>3:50-5:00 pm, <b>Poster Session</b>, Odd-numbers present, <i>Pacifica</i></p> <p>4:30-5:00 pm, <b>Ph.D. Poster Competition @ Innovation Stage</b>, <i>Pacifica</i></p> <p>5:00-5:45 pm, <b>Plenary Session</b>, Joel Dudley, <i>Pacifica</i></p>	<p>8:30-9:15 am, <b>Plenary Session</b>, Karolin Luger, <i>Oceana 6</i></p> <p>9:15-10:30 am, <b>Poster Session</b>, Even-numbers present, <i>Pacifica</i></p> <p>9:30 am - 5:00 pm, Bioinformatics Hub, <i>Pacifica</i></p> <p>10:30 am-12:20 pm, <b>Parallel Sessions</b></p> <ul style="list-style-type: none"> <li>• Systems Biology (TOA am), <i>Oceana 8-10</i></li> <li>• Immunology (TOB am), <i>Oceana 7</i></li> <li>• Structural Proteomics (TOC am), <i>Oceana 6</i></li> <li>• HPP: Metabolic Remodeling and Human Disease (TOD am), <i>Oceana 3-5</i></li> <li>• Rare Diseases (TOE am), <i>Oceana 1-2</i></li> </ul> <p>12:20-1:45 pm, <b>Corporate Lunch Seminars</b> (RSVP) or Lunch-on-your-own</p> <ul style="list-style-type: none"> <li>• Biognosys-Evosep, <i>Banda Sea</i></li> <li>• Bruker, <i>Timor Sea</i></li> <li>• SCIEX, <i>Coral Sea</i></li> <li>• Thermo, <i>Java Sea</i></li> <li>• Waters Corp., <i>South China Sea</i></li> </ul> <p>2:00-3:50 pm, <b>Parallel Sessions</b></p> <ul style="list-style-type: none"> <li>• Cardiology (TOA pm), <i>Oceana 8-10</i></li> <li>• Computational Advances (TOB pm), <i>Oceana 7</i></li> <li>• New Technological Advances in Proteomics (TOC pm), <i>Oceana 6</i></li> <li>• HPP: Harnessing the Immune System to Fight Disease (TOD pm), <i>Oceana 3-5</i></li> <li>• Microbiome &amp; Pathogen Infection (TOE pm), <i>Oceana 1-2</i></li> </ul> <p>3:50-5:00 pm, <b>Poster Session</b>, Odd-numbers present, <i>Pacifica</i></p> <p>5:00-5:30 pm, <b>HUPO General Assembly @ Innovation Stage</b>, <i>Pacifica</i></p> <p>6:30-8:30 pm, <b>LUAU Social Event</b>, Advance Purchase Ticket Req'd.</p>	<p>8:30-9:15 am, <b>Plenary Session</b>, Mary Higby Schweitzer, <i>Oceana 6</i></p> <p>9:15-10:30 am, <b>Poster Session</b>, All posters present, <i>Pacifica</i></p> <p>9:30 am - 12:00 pm, Bioinformatics Hub, <i>Pacifica</i></p> <p>10:30 am-12:20 pm, <b>Parallel Sessions</b></p> <ul style="list-style-type: none"> <li>• Activity/Chemical Proteomics (WOA am), <i>Oceana 8-10</i></li> <li>• Biomarkers, non-cancer (WOB am), <i>Oceana 7</i></li> <li>• Single-Cell Proteomics (WOC am), <i>Oceana 6</i></li> <li>• HPP: Human Chemosensation (WOD am), <i>Oceana 3-5</i></li> <li>• Neurodegenerative Diseases (WOE am), <i>Oceana 1-2</i></li> </ul> <p>12:20-1:45 pm, <b>Corporate Lunch Seminars</b> (RSVP) or Lunch-on-your-own</p> <ul style="list-style-type: none"> <li>• Thermo, <i>Java Sea and South China Sea</i></li> </ul> <p>2:00-3:50 pm, <b>Parallel Sessions</b></p> <ul style="list-style-type: none"> <li>• Drug Discovery (WOA pm), <i>Oceana 8-10</i></li> <li>• New Mass Spec Technologies (WOB pm), <i>Oceana 7</i></li> <li>• Metabolism (WOC pm), <i>Oceana 6</i></li> <li>• HPP: Unravelling Tissue Pathology through Cell Sampling(WOD pm), <i>Oceana 3-5</i></li> <li>• Personalized Wellness (WOE pm), <i>Oceana 1-2</i></li> </ul> <p>4:00-5:00 pm, <b>HUPO Award Session</b>, <i>Oceana 6</i></p> <p>5:00-5:30 pm, <b>Plenary Session</b>, Matthias Mann, <i>Oceana 6</i></p> <p>6:30 pm, optional event (buy tix at Reg), Halloween Horror @ Universal</p>
<p style="text-align: center;"><b>THURSDAY, OCT 4, 9:00 am – 3:30 pm, HPP Post-Congress Day</b> (separate registration required, lunch included)</p>		

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SUNDAY, SEPTEMBER 30 AND MONDAY, OCTOBER 1

6:00 - 7:15 pm  
**SUNDAY PLENARY**  
Session Chairs: Joshua LaBaer  
Oceana 6

8:30 - 9:15 am Monday  
**MONDAY MORNING PLENARY**  
Session Chair: Joshua LaBaer  
Oceana 6



6:00 - 6:30 pm  
**Carol V. Robinson**  
*University of Oxford*

MCP Lectureship Awardee  
**MCP** MOLECULAR & CELLULAR  
PROTEOMICS  
**Understanding Critical Protein  
Lipid Interactions Through  
Mass Spectrometry**



8:30 - 9:00 am  
**Stephen Quake**  
*Stanford University*

**Human Cell Atlas**

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9:00 - 9:15 am, **Human Proteome Project (HPP) Update**, Gil Omenn and Mark Baker



6:30 - 7:00 pm  
**Beth Anderson**  
*Arkitek Scientific*

**Data Are Beautiful**

9:15 - 10:30 am  
**POSTER SESSION**  
Pacifica

Odd-numbered posters present.

7:15 - 9:00 pm  
**WELCOME RECEPTION**  
Pacifica Ballroom  
Join your colleagues and friends for a welcome  
reception in the exhibit hall.

MONDAY, OCTOBER 1

10:30 am - 12:20 pm Monday  
**AFFINITY, PROXIMITY, & SPATIAL PROTEOMICS**  
 Session Chairs: Anne-Claude Gingras and  
 Michael Washburn  
 Oceana 8-10

- MOA am 10:30 **Capturing the RNA Binding Proteome in Time and Space**; Kathryn Lilley; *University of Cambridge, Cambridge, United Kingdom*
- MOA am 10:50 **The *in silico* Human Surfaceome & Technologies for the Elucidation of the Surfaceome Nanoscale Organization**; Bernd Wollscheid; *ETH Zurich; Zurich, Switzerland*
- MOA am 11:10 **Proximity Assays to Annotate Oncogenic Signaling-Associated Complexes: A Path towards Clinical Implementation**; Matthew Smith<sup>1</sup>; Brian Kelly<sup>2</sup>; Nathan Polaske<sup>2</sup>; Yuri Belosludstev<sup>2</sup>; Theresa Boyle<sup>1</sup>; Y. Ann Chen<sup>1</sup>; Eric Haura<sup>1</sup>; <sup>1</sup>*Moffitt Cancer Center, Tampa, Florida*; <sup>2</sup>*TRED, Roche Tissue Diagnostics, Tuscon, AZ*
- MOA am 11:22 **Interactomic Analysis of VAV1, a Key Signaling Molecules of the TCR Signaling Pathway in Primary T Cells**; Guillaume Gaud<sup>2</sup>; Romain Roncagalli<sup>3</sup>; Karima Chaoui<sup>1</sup>; Céline Colacios<sup>2</sup>; Sahar Kassem<sup>2</sup>; Bernard Monsarrat<sup>1</sup>; Odile Bulet-Schiltz<sup>1</sup>; Anne Gonzalez De Peredo<sup>1</sup>; Bernard Malissen<sup>3</sup>; Abdelhadi Saoudi<sup>2</sup>; <sup>1</sup>*Institute of Pharmacology and Structural Biology, Toulouse, France*; <sup>2</sup>*Centre de Physiopathologie de Toulouse Purpan, Toulouse, France*; <sup>3</sup>*Centre d'Immunologie de Marseille-Luminy, Marseille, France*
- MOA am 11:34 **Complex Complexes: Diversification of Sin3 HDAC Complexes via Protein Paralogs and Isoforms.**; Mark Adams<sup>1</sup>; Charles Banks<sup>1</sup>; Mihaela Sardi<sup>1</sup>; Janet Thornton<sup>1</sup>; Cassandra Eubanks<sup>1</sup>; Md Sayem Miah<sup>1</sup>; Laurence Florens<sup>1</sup>; Michael Washburn<sup>1,2</sup>; <sup>1</sup>*Stowers Institute for Medical Research, Kansas City, MO*; <sup>2</sup>*University of Kansas Medical Center, Kansas City, KS*
- MOA am 11:46 **An Enhanced Proximity Biotinylation Method for Characterisation of Large Protein Complexes**; Vincent Geoghegan; *University of York, York, UK*
- MOA am 11:58 **Antibodypedia – An Open Access Resource of Validated Antibodies**; Cristina Al-Khalili Szigyarto; Lukas Persson; Kalle von Feilitzen; Mathias Uhlen; *KTH-Royal Institute of Technology, Stockholm, Sweden*

10:30 am - 12:20 pm Monday  
**INFECTIOUS DISEASE**  
 Session Chair: Doug Sheeley  
 Oceana 7

- MOB am 10:30 **Antibiotic Resistance and Pathogenicity: Mechanistic Insights from the**

**Interactome**; James Bruce; Xuefei Zhong; Juan Chavez; Jared Mohr; Andrew Keller; Martin Mathay; Devin Schweppe; Xia Wu; *University of Washington, Seattle, WA*

- MOB am 10:50 **Peroxisome Maintenance En Route to Virus Replication: Integrated Proteomics, Lipidomics, Microscopy and Mathematical Modeling**; Ileana Cristea; *Princeton University, Princeton, NJ*
- MOB am 11:10 **Remodeling of the Glyco-phenotype of T Cell Surface Proteins with Antisense RNA of Human Immunodeficiency Virus**; Weiming Yang<sup>1</sup>; Minghui Ao<sup>1</sup>; Fabio Romero<sup>2</sup>; Hui Zhang<sup>1</sup>; <sup>1</sup>*Johns Hopkins University, Baltimore, MD*; <sup>2</sup>*University of Maryland, Baltimore, MD*
- MOB am 11:22 **HBx: Hepatitis B Virus Swiss Army Knife Enabling Successful Infection**; Emanuela Milani<sup>1</sup>; Bingqian Qu<sup>2</sup>; Stephan Urban<sup>2</sup>; Bernd Wollscheid<sup>1</sup>; <sup>1</sup>*Dep. of Health Sciences and Technology, ETH Zürich, Zürich, Switzerland*; <sup>2</sup>*University Clinic Heidelberg, Heidelberg, Deutschland*
- MOB am 11:34 **Proteolytic Events Regulate Virulence Processes of the Human Pathogen Salmonella as Shown by Quantitative Degradomics.**; Jeremy Clair; Ryan Sontag; Joshua Hansen; Aaron Wright; Joshua Adkins; *Pacific Northwest National Laboratory, Richland, WA*
- MOB am 11:46 **Systematic Identification of Mycobacterium Tuberculosis Effectors Reveals that BfrB Suppresses Innate Immunity**; Xiang He; Sheng-Ce Tao; *Shanghai Jiao Tong University, Shanghai, China*

10:30 am - 12:20 pm Monday  
**CANCER**  
 Session Chairs: Yu-Ju Chen and Sudhir Srivastava  
 Oceana 6

- MOC am 10:30 **Nickolas Papadopoulos**; *Johns Hopkins School of Medicine, Baltimore, MD*
- MOC am 10:50 **The sTRA Glycan Is Complementary to CA19-9 as a Serological Biomarker of Pancreatic Cancer**; Ben Staal<sup>1</sup>; Daniel Barnett<sup>1</sup>; Zonglin He<sup>2</sup>; Ying Huang<sup>2</sup>; Katie Partyka<sup>1</sup>; Aatur Singhi<sup>3</sup>; Richard Drake<sup>4</sup>; Anirban Maitra<sup>5</sup>; Randall Brand<sup>3</sup>; Brian Haab<sup>1</sup>; <sup>1</sup>*Van Andel Research Institute, Grand Rapids, MI*; <sup>2</sup>*Fred Hutchinson Cancer Research Center, Seattle, WA*; <sup>3</sup>*University of Pittsburgh Medical Center, Pittsburgh, PA*; <sup>4</sup>*Medical University of South Carolina, Charleston, SC*; <sup>5</sup>*MD Anderson Cancer Center, Houston, TX*
- MOC am 11:10 **A Proteomic Contexture of Exosomes Isolated from Viable Renal Cell Carcinoma Tissues, toward Development of Cancer Liquid Biopsy Diagnostics**; Atsushi Ikeda<sup>1</sup>; Kentaro Jingushi<sup>2</sup>; Naomi

MONDAY, OCTOBER 1

Ohnishi<sup>1</sup>; Motohide Uemura<sup>2</sup>; Kazutake Tsujikawa<sup>3</sup>; Koji Ueda<sup>1</sup>; <sup>1</sup>*Japanese Found. for Can. Res., Koto-Ku, Japan*; <sup>2</sup>*Dep. Therap. Uro. Onco, Osaka Univ, Osaka, Japan*; <sup>3</sup>*Lab. Mol. Cell. Phys, Osaka Univ, Osaka, Japan*

Toronto, Canada; <sup>16</sup>*Clinical Cooperation Unit Neuropathology, DKFZ, Heidelberg, Germany*; <sup>17</sup>*Inst of Neuropathology, Medical Faculty DKTK, Dusseldorf, Germany*; <sup>18</sup>*Div of Neurosurgery, Sickkids Hospital, Toronto, Canada*

MOC am 11:22 **Multi-omic Characterization of Pathway Abnormalities in High Grade Serous Ovarian Cancer**; Osama Arshad<sup>1</sup>; Jason McDermott<sup>1</sup>; Vladislav Petyuk<sup>1</sup>; Samuel Payne<sup>1</sup>; Marina Gritsenko<sup>1</sup>; Therese Clauss<sup>1</sup>; Ronald Moore<sup>1</sup>; Matthew Monroe<sup>1</sup>; Mathangi Thiagarajan<sup>2</sup>; Christopher Kinsinger<sup>3</sup>; Henry Rodriguez<sup>3</sup>; Richard Smith<sup>1</sup>; Tao Liu<sup>1</sup>; Karin Rodland<sup>1</sup>; <sup>1</sup>*Pacific Northwest National Laboratory, Richland, WA*; <sup>2</sup>*Frederick National Laboratory for Cancer Research, Frederick, MD*; <sup>3</sup>*National Cancer Institute, Bethesda, MD*

MOC am 11:46 **Proteomic-based Machine Learning Computational Analysis Discovered Biomarkers of Aberrant Vesicle-Exosomal Trafficking to Determine Chemotherapeutic Responses in the FFPE-Human Breast Cancer Sample**; Han Suk Ryu; Dohyun Han; Kyung-min Lee; Kwangsoo Kim; *Seoul National University Hospital, Seoul, South Korea*

MOC am 11:34 **Proteogenomic Landscape of Medulloblastoma Subgroups**; Antoine Forget<sup>1, 2</sup>; Loredana Martignetti<sup>3, 4</sup>; Stephanie Puget<sup>5</sup>; Laurence Calzone<sup>3, 4</sup>; Sebastian Brabetz<sup>6, 7</sup>; Daniel Picard<sup>8, 9</sup>; Arnau Montagud<sup>3, 4</sup>; Stephane Liva<sup>3, 4</sup>; Alexandre Sta<sup>3, 4</sup>; Florent Dingli<sup>10</sup>; Guillaume Arras<sup>10</sup>; Jaime Rivera<sup>10</sup>; Damarys Loew<sup>10</sup>; Aurore Besnard<sup>11</sup>; Joëlle Lacombe<sup>11</sup>; Mélanie Pagès<sup>11</sup>; Pascale Varlet<sup>11</sup>; Christelle Dufour<sup>12</sup>; Hua Yu<sup>1, 2</sup>; Audrey Mercier<sup>1, 2</sup>; Sophie Leboucher<sup>1, 13</sup>; Laura Sieber<sup>6, 7</sup>; Kevin Beccaria<sup>5</sup>; Michael Gombert<sup>9</sup>; Frauke Meyer<sup>8, 9</sup>; Nan Qin<sup>8, 9</sup>; Jasmin Bartl<sup>8, 9</sup>; Lukas Chavez<sup>6, 7</sup>; Konstantin Okonechnikov<sup>6, 7</sup>; Tanvi Sharma<sup>6, 7</sup>; Venu Thatikonda<sup>6, 7</sup>; Franck Bourdeaut<sup>14</sup>; Celio Pouponnot<sup>1, 2</sup>; Vijay Ramaswamy<sup>15</sup>; Andrey Korshunov<sup>16</sup>; Arndt Borkhardt<sup>9</sup>; Guido Reifenberger<sup>17</sup>; Patrick Pouillet<sup>3, 4</sup>; Michael D. Taylor<sup>18</sup>; Marcel Kool<sup>6, 7</sup>; Stefan M. Pfister<sup>6, 7</sup>; Daisuke Kawachi<sup>6, 7</sup>; Emmanuel Barillot<sup>3, 4</sup>; Marc Remke<sup>8, 9</sup>; Olivier Ayrault<sup>1, 2</sup>; <sup>1</sup>*Institut Curie, PSL Research Univ, CNRS UMR, Orsay, France*; <sup>2</sup>*Université Paris Sud, Univ Paris-Saclay, Orsay, France*; <sup>3</sup>*Institut Curie, Paris, France*; <sup>4</sup>*Inserm, Paris, France*; <sup>5</sup>*Dept of Pediatric Neurosurgery, Necker Hosp., Paris, France*; <sup>6</sup>*Hopp Children's Cancer Center at the NCT, Heidelberg, Germany*; <sup>7</sup>*Div of Pediatric Neurooncology-DKFZ-DKTK, Heidelberg, Germany*; <sup>8</sup>*Dept of Pediatric Neuro-Oncogenomics, DKTK, Dusseldorf, Germany*; <sup>9</sup>*Dept of Pediatric Oncology, DKTK, Dusseldorf, Germany*; <sup>10</sup>*Proteomics and Mass Spec Lab, Inst. Curie, Paris, France*; <sup>11</sup>*Dept of Neuropathology, Sainte-Anne Hospital, Paris, France*; <sup>12</sup>*Pediatric & Adolescent Oncol. Dpt, Gustave Roussy, Villejuif, France*; <sup>13</sup>*Institut Curie, Plateforme d'Histologie, Orsay, France*; <sup>14</sup>*PSL Univ., Institut Curie Research Center, SiRIC, Paris, France*; <sup>15</sup>*Div. of Haematology/Oncology, Sickkids Hosp.,*

MOC am 11:58 **Proteomics in PreCancer Atlas: an Evidence Based Biomarkers Discovery**; Sudhir Srivastava; *National Cancer Institute, NIH; Bethesda, MD*

**10:30 am - 12:20 pm Monday**  
**HPP: PARTNERING WITH PATHOLOGY TOWARDS PRECISION MEDICINE**  
**Session Chairs: Daniel W. Chan and Ed Nice**  
**Oceana 3-5**

MOD am 10:30 **Can Proteomic Pathology Deliver Solutions for Unmet Clinical Needs?**; Stephen Pennington<sup>1, 2</sup>; <sup>1</sup>*UCD Conway Institute, School of Medicine, Dublin, Ireland*; <sup>2</sup>*University College Dublin, Ireland*

MOD am 10:50 **Adoption of Proteomics into Clinical Practice- Barriers and Opportunities**; Peter Stewart<sup>1, 2</sup>; <sup>1</sup>*Royal Prince Alfred and Liverpool Hospitals, Sydney, Australia*; <sup>2</sup>*University of Sydney, Australia*

MOD am 11:10 **Immuno-Proteomics of Colon Cancer**; Michael Roehrl; *MSKCC, New York, NY*

MOD am 11:22 **Comparison of Fractionated versus Unfractionated Plasma Lipoproteome in the Context of Vascular Contributions to Alzheimer's Disease**; Danni Li; *University of Minnesota, Minneapolis, MN*

MOD am 11:34 **Proteomics of Kidney Biopsy Tissue and Urine Liquid Biopsy towards Precision Medicine**; Tadashi Yamamoto<sup>1, 2</sup>; Keiko Yamamoto<sup>1</sup>; Bo Xu<sup>1</sup>; Amr Elguoshy<sup>1</sup>; Yoshitoshi Hirao<sup>1</sup>; <sup>1</sup>*Biofluid Biomarker Center, Niigata University, Niigata, Japan*; <sup>2</sup>*Shinrakuen Hospital, Niigata, Japan*

MOD am 11:46 **An Approach to Spatiotemporally Resolve Protein Interaction Networks in Living Cells**; Ruth Hüttenhain<sup>1, 2</sup>; Braden T. Lobingier<sup>1</sup>; Kelsie Eichel<sup>1</sup>; Alice Y. Ting<sup>3</sup>; Brian Shoichet<sup>1</sup>; Mark von Zastrow<sup>1</sup>; Nevan J. Krogan<sup>1, 2</sup>; <sup>1</sup>*UCSF, San Francisco, CA*; <sup>2</sup>*Gladstone Institutes, San Francisco, CA*; <sup>3</sup>*Stanford University, Stanford, CA*

MOD am 11:58 **Improved Survival Prognostication of Node-Positive Malignant Melanoma Patients – A Proteogenomics Study Guided by Histopathological**

**Characterization**; Lazaro Betancourt; Krzysztof Pawlowski; Jonatan Eriksson; Marcell Szasz; Shamik Mitra; Indira Pla Parada; Charlotte Welinder; Henrik Ekedahl; Per Broberg; Roger Appelqvist; Maria Yakovleva; Yutaka Sugihara; Kenichi Miharada; Christian Ingvar; Lotta Lundgre; Bo Baldetorp; Håkan Olsson; Melinda Rezeli; Elisabet Wieslander; Peter Horvatovich; Johan Malm; Göran Jönsson; György Marko-Varga; *Lund University, Lund, Sweden*

**10:30 am - 12:20 pm Monday**  
**STATISTICS IN EXPERIMENTAL DESIGN**  
**Session Chairs: Olga Vitek and Pei Wang**  
**Oceana 1-2**

- MOE am 10:30 **Experimental Design and Data-Analysis in Label-Free Quantitative MS-based Proteomics**; Lieven Clement; *Ghent University, Ghent, Belgium*
- MOE am 10:50 **Components of Reproducible Quantitative Mass Spectrometry-Based Research: A Statistician's Perspective**; Olga Vitek; *Northeastern University, Boston, MA*
- MOE am 11:10 **Empirical Peptide-Level Statistics Allow Robust and Sensitive Differential Expression Detection in MS-Proteomics Data**; Constantin Ammar; Gergely Csaba; Markus Gruber; Ralf Zimmer; *LMU Munich, Munich, Germany*
- MOE am 11:22 **A Statistical Framework for Relative Quantification of Post-Translational Modifications in Global Proteomics Experiments**; Tsung-Heng Tsai<sup>1</sup>; Lillian Phu<sup>2</sup>; Yi Zeng<sup>2</sup>; Donald Kirkpatrick<sup>2</sup>; Erik Verschueren<sup>2</sup>; Olga Vitek<sup>1</sup>; <sup>1</sup>*Northeastern University, Boston, MA*; <sup>2</sup>*Genentech, Inc., South San Francisco, CA*
- MOE am 11:34 **Bayesian Confidence Intervals for Multiplexed Proteomics Integrate Ion Statistics with Peptide Quantification Concordance**; Leonid Peshkin<sup>2</sup>; Meera Gupta<sup>1</sup>; Lillia Ryazanova<sup>1</sup>; Martin Wühr<sup>1</sup>; <sup>1</sup>*Princeton University, Princeton, NJ*; <sup>2</sup>*Harvard Medical School, Boston, MA*
- MOE am 11:46 **MSstatsTMT: Statistical Detection of Differentially Abundant Proteins in Mass Spectrometry Experiments with Isobaric Labeling**; Ting Huang<sup>1</sup>; Meena Choi<sup>1</sup>; Manuel Tzouros<sup>2</sup>; Nikhil Pandya<sup>2</sup>; Balazs Banfai<sup>2</sup>; Tom Dunkley<sup>2</sup>; Olga Vitek<sup>1</sup>; <sup>1</sup>*Northeastern University, Boston, MA*; <sup>2</sup>*Hoffmann-La Roche Ltd, Basel, Switzerland*
- MOE am 11:58 **Cancer Proteogenomics**; David Fenyö; *New York University School of Medicine, New York, NY*

**12:20 – 1:45 pm Monday**  
**CORPORATE LUNCH SEMINARS or Lunch-on-your-own**

RSVP Required for Corporate Lunch Seminars.



*South China Sea Room*

**Agilent Innovations for Proteomics Research**  
 Christine Miller, *Omics Market Manager*  
 Randy Bolger, *Workflow Solutions Manager*



*Timor Sea Room*

**Introduction to the timsTOF Pro powered by PASEF**  
 Oliver Raether, *QTOF R&D Manager, Bruker Daltonik*

**Data-independent Parallel Accumulation – Serial Fragmentation (diaPASEF) on the tims – TOF Pro**  
 Matthias Mann, *Max Planck Institute of Biochemistry*



*Coral Sea Room*

**Quantitative Proteomics Profiling: SWATH® Acquisition as a Tool for the Research & Service Core Lab**

Yansheng Liu, *Yale University*  
 Birgit Schilling, *Buck Institute for Research on Aging*



*Java Sea Room*

**Latest Advances in Multiplexing Technology:**

- Establishing a Roadmap for Brain-based Protein Biomarkers in Alzheimer's Disease; Nicholas Seyfried, *Emory School of Medicine*
- Drug Effects on Protein Homeostasis; Marcus Bantscheff, *Cellzome, a GSK Company*

MONDAY, OCTOBER 1

2:00 - 3:50 pm Monday  
**AGING**  
 Session Chairs: Birgit Schilling and Paola Sebastiani  
 Oceana 8-10

2:00 - 3:50 pm Monday  
**NUTRITION AND FOOD**  
 Session Chairs: Subhra Chakraborty and Paola Roncada  
 Oceana 7

- MOA pm 2:00 **Proteomics Analysis of Skeletal Muscle in Healthy Human**; Ceereena Ubaida-Mohien; Alexey Lyashkov; Ravi Tharakan; Marta Gonzalez-Freire; Michelle Shardell; Ruin Moaddel; Chee Chia; Luigi Ferrucci; *National Institute on Aging, Baltimore, MD*
- MOA pm 2:20 **May Mitochondrial Dysfunction Predispose for Cancer?**; Christopher Gerner; *University of Vienna*
- MOA pm 2:40 **Mouse and human models of longevity indicate altered response to Diet Induced Free Radical Damage**; Nadia Ashrafi<sup>1, 2</sup>; Wendy E. Heywood<sup>1</sup>; Marie-Stéphanie Clerget-Froidevaux<sup>2</sup>; Barbara Demeneix<sup>2</sup>; Diana van Heemst<sup>3</sup>; Raymond Noordam<sup>3</sup>; Kevin Mills<sup>1</sup>; <sup>1</sup>*University College London, London, United Kingdom*; <sup>2</sup>*Evolution of Endocrine Regulations, Paris, France*; <sup>3</sup>*Leiden University Medical Center, Leiden, Netherland*
- MOA pm 2:52 **Representation of Neonatal and Paediatric Proteins in Current Plasma Proteome Databases**; Conor McCafferty<sup>1</sup>; Jochen M. Schwenk<sup>2</sup>; Vera Ignjatovic<sup>1, 3</sup>; <sup>1</sup>*Murdoch Children's Research Institute, Parkville, Australia*; <sup>2</sup>*KTH Royal Institute of Technology, Stockholm, Sweden*; <sup>3</sup>*The University of Melbourne, Parkville, Australia*
- MOA pm 3:04 **Identification and Quantification of Biomarkers for Senescent Cells Using Mass Spectrometry**; Nathan Basisty; Abhijit Kale; Herbert Kasler; Eric Verdin; Judith Campisi; Birgit Schilling; *The Buck Institute for Research on Aging, Novato, CA*
- MOA pm 3:16 **Establishing a Roadmap for Brain-based Protein Biomarkers in Alzheimer's Disease**; Maotian Zhou; Duc Duong; Lingyan Ping; Eric Dammer; Marla Gearing; James Lah; Allan Levey; Nicholas Seyfried; *Emory School of Medicine, Atlanta, GA*
- MOA pm 3:28 **Investigating the Relationship between Protein Glycation and Stability in Cells and Tissues**; Simone Di Sanzo; Joanna Kirkpatrick; Nicolas Huber; Alessandro Ori; *Leibniz Institute on Aging, Jena, Germany*

- MOB pm 2:00 **MPK4 Interacting Proteins in Crop Immunity Response**; Sixue Chen; *University of Florida, Gainesville, FL*
- MOB pm 2:20 **Omics Approaches to Uncover Tolerant Mechanism in Soybean at the Initial Flooding Stress**; Setsuko Komatsu; *Fukui University of Technology, Fukui, Japan*
- MOB pm 2:40 **Nutraceuticals, Deciphering Mechanism of Actions of Novel Bioactive Compounds by Thermal Proteome Profiling**; Ana M Carrasco del Amor<sup>1</sup>; Olatz Fresnedo<sup>2</sup>; Siegfried Ussar<sup>4</sup>; Ralph Urbatzka<sup>3</sup>; Susana Cristobal<sup>1</sup>; <sup>1</sup>*Linköping University, Linköping, Sweden*; <sup>2</sup>*University of the Basque Country, Leioa, ES*; <sup>3</sup>*CIIMAR - Interdisciplinary Center of Marine and En, Porto, PT*; <sup>4</sup>*Hemholtz Center Munich, Munich, DE*
- MOB pm 2:52 **Single Shot DIA Profiling of >1500 Plasma Proteomes of the Weight Loss and Maintenance Study DiOGenes**; Roland Bruderer<sup>1</sup>; Jan Muntel<sup>1</sup>; Sebastian Müller<sup>1</sup>; Oliver M. Bernhardt<sup>1</sup>; Tejas Gandhi<sup>1</sup>; Polina Mironova<sup>2</sup>; Ondine Walter<sup>2</sup>; Jérôme Carayol<sup>2</sup>; Arne Astrup<sup>3</sup>; Wim H.M. Saris<sup>4</sup>; Jörg Hager<sup>2</sup>; Armand Valsesia<sup>2</sup>; Loic Dayon<sup>2</sup>; Lukas Reiter<sup>1</sup>; <sup>1</sup>*Biognosys AG, Schlieren, Switzerland*; <sup>2</sup>*Nestle, Lausanne, Switzerland*; <sup>3</sup>*University of Copenhagen, Copenhagen, Denmark*; <sup>4</sup>*University of Maastricht, Maastricht, Netherlands*
- MOB pm 3:04 **Comparative Proteomics of Low Oxalate Tomatoes during Post-harvest Storage**; Sudip Ghosh; Kanika Narula; Pooja Aggarwal; Niranjana Chakraborty; Subhra Chakraborty; *National Institute of Plant Genome Research, New Delhi, India*
- MOB pm 3:16 **Identification of Functional Peptides with Tolerogenic Potential in a Partially Hydrolysed Infant Formula**; Joost Gouw<sup>1</sup>; Juandy Jo<sup>2, 3</sup>; Laura Meulenbroek<sup>1, 3</sup>; Sam Heijjer<sup>1, 3</sup>; Erica Kremer<sup>1</sup>; Elena Sandalova<sup>2, 3</sup>; André Knulst<sup>4</sup>; Sergio Oliviera<sup>1</sup>; Jan Knol<sup>1, 5</sup>; Johan Garssen<sup>1, 3</sup>; Anneke Rijnierse<sup>1</sup>; Léon Knippels<sup>1, 3</sup>; <sup>1</sup>*Danone Nutricia Research, Utrecht, Netherlands*; <sup>2</sup>*Danone Nutricia Research, Singapore*; <sup>3</sup>*UIPS, Utrecht, Netherlands*; <sup>4</sup>*Department of Dermatology and Allergology, Utrecht, Netherlands*; <sup>5</sup>*Laboratory of Microbiology, Wageningen, Netherlands*
- MOB pm 3:28 **Culture Independent Label Free Metho for Milk Metaproteome and Resistome Analysis**; Cristian Piras<sup>1</sup>; Alessio Soggiu<sup>1</sup>; Viviana Greco<sup>2</sup>; Luigi Bonizzi<sup>1</sup>; Alfonso Zecconi<sup>1</sup>; Andrea Urbani<sup>3</sup>; Claudia Gusmara<sup>1</sup>; Domenico Britti<sup>4</sup>; Paola

MONDAY, OCTOBER 1

Roncada<sup>4</sup>; <sup>1</sup>DIMEVET - University of Milan, Milano, Italy; <sup>2</sup>Fondazione Santa Lucia, Rome, Italy; <sup>3</sup>Catholic University of Sacred Heart, Rome, Italy; <sup>4</sup>Università Magna Græcia, Catanzaro, Italy

Xinlei Sheng; Ileana Cristea; Princeton University, Princeton, NJ

2:00 - 3:50 pm Monday  
**POSTTRANSLATIONAL MODIFICATIONS (PTMs)**  
 Session Chair: Karolin Luger and Yingming Zhao  
 Oceana 6

- MOC pm 2:00 **Quantitative Proteomics for Understanding Cancer Epigenetics;** Benjamin Garcia; *University of Pennsylvania School of Medicine, Philadelphia, PA*
- MOC pm 2:20 **Linear Ubiquitin Control Identified by Positional Proteomics Leading to Discovery of a Molecular Corrector to Rescue NFκB Activation in Immunodeficiency;** Christopher Overall; *University of British Columbia, Vancouver, BC*
- MOC pm 2:40 **Lysine Benzoylation Is a New Type of Histone Mark Regulated by SIRT2;** He Huang; Di Zhang; Mathew Perez-Neut; Yingming Zhao; *The University of Chicago, Chicago, IL*
- MOC pm 2:52 **Reanalysis of Global Proteomic and Phosphoproteomic Data Identified a Large Number of Glycopeptides;** Yingwei Hu; Punit Shah; David J. Clark; Minghui Ao; Hui Zhang; *Johns Hopkins University, Baltimore, Maryland*
- MOC pm 3:04 **Phosphoproteomic Landscapes of Cancer Cell Lines Predict Drug Response;** Martin Frejno<sup>1</sup>; Benjamin Ruprecht<sup>1, 2</sup>; Chen Meng<sup>1</sup>; Alexander Högberg<sup>1, 3</sup>; Jana Zecha<sup>1, 4</sup>; Dominic Helm<sup>1, 5</sup>; Thomas Oellerich<sup>6, 7</sup>; Sebastian Scheich<sup>7</sup>; Hans-Michael Kvasnicka<sup>7</sup>; Enken Drecoll<sup>1</sup>; Wilko Weichert<sup>1</sup>; Bernhard Kuster<sup>1, 8</sup>; <sup>1</sup>Technical University of Munich, Freising, Germany; <sup>2</sup>Merck & Co., Boston, MA; <sup>3</sup>University of Copenhagen, Copenhagen, Denmark; <sup>4</sup>German Cancer Consortium (DKTK), Munich, Germany; <sup>5</sup>EMBL, Heidelberg, Germany; <sup>6</sup>Cambridge University, Cambridge, United Kingdom; <sup>7</sup>Goethe University, Frankfurt, Germany; <sup>8</sup>Center for Integrated Protein Science (CIPSM), Munich, Germany
- MOC pm 3:16 **A Tandem Affinity Enrichment Method Identifies MacroH2A1 as a BRCA1/BARD1 E3 Ligase Substrate;** Beom-Jun Kim<sup>1</sup>; Doug Chan<sup>1</sup>; Sung Jung<sup>1</sup>; Yue Chen<sup>1</sup>; Jun Qin<sup>1, 2</sup>; Yi Wang<sup>1, 2</sup>; <sup>1</sup>Baylor College of Medicine, Houston, TX; <sup>2</sup>National Center for Protein Sciences, Beijing, China
- MOC pm 3:28 **Temporal-Spatial Orchestration of Protein Acetylation In Antiviral Response and Immunity;** Laura Murray;

2:00 - 3:50 pm Monday  
**HPP: TARGETING THE PROTEOME IN WOMEN'S HEALTH**  
 Session Chair: Jennifer Van Eyk  
 Oceana 3-5

- MOD pm 2:00 **Tony Whetton;** *University of Manchester, Manchester, UK*
- MOD pm 2:30 **Nur Yucer;** *Cedars Sinai Medical Center, Los Angeles, CA*
- MOD pm 3:00 **Location Is Everything: Protein Translocation as a Virus Replication Strategy;** Katelyn C Cook; Pierre M Jean Beltran; Michelle A Kennedy; Ileana M Cristea; *Princeton University, Princeton, NJ*
- MOD pm 3:15 **Improvement of Detection Sensitivity of nLC-MS towards the Single-Cell Proteomics in the Era of Precision Medicine;** Qing Kay Li<sup>1</sup>; Chuanzi Ouyang<sup>2</sup>; Hui Zhang<sup>2</sup>; <sup>1</sup>Johns Hopkins Hospitals, Baltimore, <Not Specified>; <sup>2</sup>Johns Hopkins University, Baltimore, Maryland
- MOD pm 3:30 **Evaluating Mitra® Microsampling Devices for Remote Monitoring of Apolipoproteins in Patients at Risk for Cardiac Events;** Kelly Mouapi<sup>1, 2</sup>; Irene van den Broek<sup>1, 2</sup>; Mitra Mastali<sup>1, 2</sup>; Qin Fu<sup>1, 2</sup>; Garth Fuller<sup>4</sup>; Sandy Young<sup>1, 3</sup>; Shivani Dhawan<sup>1, 3</sup>; Mayra Lopez<sup>4</sup>; Chrisandra Shufelt<sup>1, 3</sup>; Brennan Spiegel<sup>4</sup>; Noel Bairey Merz<sup>1, 3</sup>; Jennifer Van Eyk<sup>1, 2</sup>; <sup>1</sup>Smidt Heart Institute, Cedars-Sinai Medical Center, Los Angeles, CA; <sup>2</sup>Advanced Clinical Biosystems Research Institute, Los Angeles, CA; <sup>3</sup>Barbra Streisand Women's Heart Center, Smidt Heart, Los Angeles, CA; <sup>4</sup>Cedars-Sinai Center for Outcomes Research(CS-CORE), Los Angeles, CA

2:00 - 3:50 pm Monday  
**PROTEOGENOMICS**  
 Session Chair: Henry Rodriguez  
 Oceana 1-2

- MOE pm 2:00 **Antibiotics Discovery: From Peptidogenomics to Genome Mining to Spectral Networks;** Pavel A. Pevzner; *University of California, San Diego, San Diego, CA*
- MOE pm 2:20 **Karin Rodland;** *Pacific Northwest National Laboratory, Richland, WA*
- MOE pm 2:40 **Impact of Alternative Splicing on Proteoisoforms and Epigenetic Regulation during T cell Stimulation;** Laura Agosto<sup>1, 2</sup>; Michael J. Mallory<sup>1</sup>; Simone Sidoli<sup>1, 2</sup>; Amber K. Weiner<sup>1, 2</sup>; Kristen W. Lynch<sup>1</sup>; Benjamin A. Garcia<sup>1, 2</sup>; <sup>1</sup>University of Pennsylvania School of Medicine, Philadelphia, PA; <sup>2</sup>Penn Epigenetics Institute, Philadelphia, PA

MONDAY, OCTOBER 1

MOE pm 2:52 **Proteogenomic Subtypes of Squamous Cell Lung Cancer**; Paul Stewart; Eric Welsh; Robbert Slebos; Bin Fang; Victoria Izumi; Matthew Chambers; Guolin Zhang; Ling Cen; Fredrik Pettersson; Yonghong Zhang; Zhihua Chen; Chia-Ho Cheng; Katherine Fellows; Jewel Francis; Tania Mesa; Chaomei Zhang; Sean Yoder; Gina DeNicola; Amer Beg; Theresa Boyle; Jamie Teer; Ann Chen; John Koomen; Steven Eschrich; Eric Haura; *Moffitt Cancer Center, Tampa, FL*

MOE pm 3:04 **Proteogenomics Discovery of Human Coding Regions and Cancer Neoantigens**; Janne Lehti<sup>1,2</sup>; Yafeng Zhu<sup>1</sup>; Husen Umer<sup>1</sup>; Rui Branca<sup>1</sup>; <sup>1</sup>*Karolinska Institutet, Stockholm, Sweden*; <sup>2</sup>*SciLifeLab, Stockholm, Sweden*

MOE pm 3:16 **Proteogenomic Integration for Systematic Identification and Prioritization of Tumor Neoantigens**; Bo Wen; Yun Zhang; Noel Namai; Yongchao Dou; Bing Zhang; *Baylor College of Medicine, Houston, TX*

MOE pm 3:28 **The Proteome Data Commons in the Context of the NCI Cancer Research Data Commons**; Christopher R Kinsinger<sup>1</sup>; Izumi Hinkson<sup>1</sup>; Ratna R Thangudu<sup>2</sup>; Michael Holck<sup>2</sup>; Deepak Singhal<sup>2</sup>; Karen Ketchum<sup>2</sup>; Paul A Rudnick<sup>3</sup>; Nathan J Edwards<sup>4</sup>; Michael J MacCoss<sup>5</sup>; Anand Basu<sup>2</sup>; <sup>1</sup>*National Cancer Institute, Bethesda, Maryland*; <sup>2</sup>*ESAC, Rockville, MD*; <sup>3</sup>*Spectragen-Informatics, Bainbridge Island, WA*; <sup>4</sup>*Georgetown University Medical Center, Washington, DC*; <sup>5</sup>*University of Washington Genome Sciences, Seattle, WA*

5:00 - 5:45 pm Monday  
MONDAY AFTERNOON PLENARY  
Session Chair: Joshua LaBaer  
Oceana 6



Joel Dudley  
*Mount Sinai Medical Center*

Big Data

3:50 – 5:00 pm  
POSTER SESSION  
Pacifica

Odd-numbers present.

Don't miss the Ph.D. Poster Competition Finalists on the Innovation Stage, 4:30-5:00 pm

8:30 - 9:15 pm Tuesday  
**TUESDAY MORNING PLENARY**  
 Session Chair: Robert Moritz  
 Oceana 6



8:30 - 9:00 am  
**Karolin Luger**  
*Howard Hughes Medical  
 Institute and University of  
 Colorado at Boulder*

**Off to the Races –  
 Quantitating the  
 Recruitment of Proteins to  
 Sites of DNA Damage**

9:00 - 9:15 am, **Special International Initiatives Update,**  
 Fuchu He

9:15 - 10:30 am  
**POSTER SESSION**  
 Pacifica

Even-numbers present.

**Don't miss the ProLabs Instruments presentation on the  
 Innovation Stage, 9:45 - 10:00 am.**

10:30 am - 12:20 pm Tuesday  
**SYSTEMS BIOLOGY**  
 Session Chair: Robert Moritz  
 Oceana 8-10

TOA am 10:30 **Pathology from the Molecular Scale on  
 Up;** Garry P. Nolan; *Stanford University  
 School of Medicine, Stanford, CA*

TOA am 10:50 **Systems Genetics Approaches to Map  
 the Functional Organization of the  
 Proteome;** Jasmin Coulombe-Huntington;  
 Thierry Bertomeu; Linnea Olofsson;  
 Caroline Huard; Daniel St-Cyr; Lily Zhang;  
 Andrew Chatr-ayramontri; Mike Tyers;  
*Université de Montréal, Montréal, Canada*

TOA am 11:10 **Towards the Human Co-Receptome: A  
 Systematic Exploration of the  
 Immunoglobulin Superfamily  
 Interactome;** Nadia Martinez; *Genentech,  
 South San Francisco, CA*

TOA am 11:22 **Systems Pharmacology Dissection of  
 Cell-Specific Cholesterol Regulation  
 Mechanisms Reveals Large  
 Pharmacodynamic Variability;** Peter  
 Blattmann<sup>1</sup>; David Henriques<sup>2</sup>; Michael  
 Zimmermann<sup>1</sup>; Fabian Frommelt<sup>1</sup>; Uwe  
 Sauer<sup>1</sup>; Julio Saez-Rodriguez<sup>3,4</sup>; Ruedi  
 Aebersold<sup>1,5</sup>; <sup>1</sup>*ETH Zurich, Zurich,  
 Switzerland*; <sup>2</sup>*Spanish Council for Scientific  
 Research, Vigo, Spain*; <sup>3</sup>*RWTH-Aachen*

*University, Aachen, Germany*; <sup>4</sup>*European  
 Molecular Biology Laboratory - European B,  
 Hinxton, United Kingdom*; <sup>5</sup>*University of  
 Zurich, Zurich, Switzerland*

TOA am 11:34 **Building a Global Map of Human Protein  
 Complexes: Synthesis of >15k Mass  
 Spectrometry Experiments;** Kevin Drew;  
 John B. Wallingford; Edward Marcotte;  
*University of Texas, Austin, TX*

TOA am 11:46 **Charting of Endothelin B Receptor  
 Signaling using Phosphoproteomics  
 Discovers Critical Kinases for  
 Endothelin Induced Cell Migration;**  
Alexander Schaefer<sup>1</sup>; Richard W. D.  
 Welford<sup>2</sup>; Imke Renz<sup>2</sup>; Francois Lehembre<sup>2</sup>;  
 Peter M. A. Groenen<sup>2</sup>; Enio Gjerga<sup>3</sup>; Julio  
 Saez-Rodriguez<sup>3,4</sup>; Ruedi Aebersold<sup>1,5</sup>;  
 Matthias Gstaiger<sup>1</sup>; <sup>1</sup>*ETH Zurich, Zurich,  
 Switzerland*; <sup>2</sup>*Idorsia Pharmaceuticals,  
 Allschwil, Switzerland*; <sup>3</sup>*RWTH Aachen  
 University, Aachen, Germany*; <sup>4</sup>*European  
 Molecular Biology Laboratory, Hinxton, UK*;  
<sup>5</sup>*Faculty of Science, University of Zürich,  
 Zurich, Switzerland*

TOA am 11:58 **A High-Resolution Organellar  
 Proteomics Approach to Study  
 Subcellular Distribution and Dynamics of  
 Proteins in Cells;** Johannes Jordan<sup>1</sup>;  
 Wolfgang Bildl<sup>1</sup>; Clara Steinbrueck<sup>1</sup>;  
 Alexander Haupt<sup>1</sup>; Maciej Kocylowski<sup>1</sup>;  
 Astrid Kollwe<sup>1</sup>; Bernd Fakler<sup>1,2</sup>; Uwe  
 Schulte<sup>1,3</sup>; <sup>1</sup>*University of Freiburg,  
 Germany, Freiburg, Germany*; <sup>2</sup>*Center for  
 Biological Signaling Studies (BIOSS),  
 Freiburg, Germany*; <sup>3</sup>*Logopharm GmbH,  
 March-Buchheim, Germany*

10:30 am - 12:20 pm Tuesday  
**IMMUNOLOGY**  
 Session Chair: Pierre Thibault  
 Oceana 7

TOB am 10:30 **Advanced Immunopeptidomics based  
 Discovery Engine for the Development of  
 Personalized Cancer Immunotherapy;**  
 Michal Bassani-Sternberg; *Unil-Chuv*

TOB am 10:50 **Allegedly Non-Coding Regions Are the  
 Main Source of Tumor-Specific Antigens;**  
 Claude Perreault; *IRIC-Université de  
 Montréal, Montreal, Canada*

TOB am 11:10 **Systematic Profiling of HLA Class I  
 Peptide Epitopes by LC-MS/MS in Mono-  
 Allelic Cells Improves Neoantigen  
 Binding Prediction Algorithms;** Susan  
 Klaeger<sup>1</sup>; Derin Keskin<sup>2,3</sup>; Siranush  
 Sarkizova<sup>4</sup>; Karl R Clauser<sup>1</sup>; Oliver Spiro<sup>1</sup>;  
 Hasmik Keshishian<sup>1</sup>; Christina Hartigan<sup>1</sup>;  
 Nir Hacohen<sup>1,5</sup>; Catherine J Wu<sup>1,2</sup>; Steven  
 A Carr<sup>1</sup>; <sup>1</sup>*The Broad Institute, Cambridge,  
 MA*; <sup>2</sup>*Dana-Farber Cancer Institute, Boston,  
 MA*; <sup>3</sup>*Brigham and Women's Hospital,  
 Boston, MA*; <sup>4</sup>*Harvard Medical School,  
 Boston, MA*; <sup>5</sup>*Massachusetts General  
 Hospital, Boston, MA*



TUESDAY, OCTOBER 2

- TOB am 11:22 **Integrated Proteome and HLA Peptidome Quantitation with Tandem Mass Tags;** Patrick Murphy<sup>1</sup>; Prathyusha Konda<sup>1</sup>; Joao Paulo<sup>3</sup>; Heiko Schuster<sup>2</sup>; Daniel Kowalewski<sup>2</sup>; Youra Kim<sup>1</sup>; Derek Clements<sup>1</sup>; Michael Giacomantonio<sup>1</sup>; Stefan Stevanovic<sup>2</sup>; Steve Gygi<sup>3</sup>; Shashi Gujar<sup>1</sup>; <sup>1</sup>Dalhousie University, Halifax, Canada; <sup>2</sup>Tubingen University, Tuebingen, Germany; <sup>3</sup>Harvard Medical School, Boston, MA
- TOB am 11:34 **Is Hybrid Peptide Formation a New Post-Translational Modification that Drives Autoimmunity?** Timothy Wiles; Anita Hohenstein; Thomas Delong; University of Colorado Skaggs School of Pharmacy, Aurora, CO
- TOB am 11:46 **Citrullinated Glucose-regulated Protein 78: An Autoantigen in Human Type 1 Diabetes;** Lut Overbergh<sup>1</sup>; Inne Crèvecoeur<sup>1</sup>; Fernanda MC Sodr <sup>1</sup>; Aisha Callebaut<sup>1</sup>; Gabriele B fagan<sup>2</sup>; Mei-Ling Yang<sup>3</sup>; David Arribas-Layton<sup>2</sup>; Meghan Marr <sup>4</sup>; Dana P Cook<sup>1</sup>; Etienne Waelkens<sup>1</sup>; Roberto Mallone<sup>5</sup>; Jon Piganelli<sup>4</sup>; Rita Derua<sup>1</sup>; Mark Mamula<sup>3</sup>; Eddie A James<sup>2</sup>; Chantal Mathieu<sup>1</sup>; Mijke Buitinga<sup>1</sup>; <sup>1</sup>KU Leuven, Leuven, Belgium, Leuven, Belgium; <sup>2</sup>Benaroya Research Institute, Seattle, WA, USA; <sup>3</sup>Yale University School of Medicine, New Haven, CT, USA; <sup>4</sup>University of Pittsburgh, Division of Pediatric S, Pittsburgh, PA, USA; <sup>5</sup>INSERM, Paris, France
- TOB am 11:58 **Cytoplasmic Interactions of the Pathogen Recognition Receptor cGAS Modulate Type I IFN induction during Herpesvirus Infection;** Krystal Lum; Bokai Song; Joel Federspiel; Benjamin Diner; Ileana Cristea; Princeton University, Princeton, NJ
- Benjamin A. Garcia<sup>1,2</sup>; <sup>1</sup>University of Pennsylvania, Epigenetics Institute, Philadelphia, PA; <sup>2</sup>University of Pennsylvania School of Medicine, Philadelphia, PA
- TOC am 11:22 **Cell-deep Structural Biology Insights from Surface-exposed Biotins;** David-Paul Minde; Manasa Ramakrishna; Kathryn Lilley; University of Cambridge, Cambridge, United Kingdom
- TOC am 11:34 **Structure and Protein Interaction-based Gene Ontology Annotations Reveal Likely Functions of Uncharacterized Proteins on Human Chromosome 17;** Chengxin Zhang; Gilbert Omenn; Yang Zhang; University of Michigan, Ann Arbor, MI
- TOC am 11:46 **Novel Strategies for Enrichment of Membrane Proteins and Structural Characterization by Top-down Mass Spectrometry with Ultra-Violet Photodissociation (UVPD);** Julian Whitelegge; Semel Institute, UCLA, Los Angeles, CA
- TOC am 11:58 **Conformational Switching of the MLKL Pseudokinase Domain Promotes MLKL Tetramerization and Cell Death by Necroptosis;** Jarro Sandow<sup>1</sup>; Emma Petrie<sup>1</sup>; Annette Jacobson<sup>1</sup>; Brian Smith<sup>3</sup>; Michael Griffin<sup>2</sup>; Isabelle Lucet<sup>1</sup>; Weiwen Dai<sup>1</sup>; Samuel Young<sup>1</sup>; Maria Tanzer<sup>1</sup>; Ahmad Wardak<sup>1</sup>; Lung-Yu Liang<sup>1</sup>; Angus Cowan<sup>1</sup>; Joanne Hildebrand<sup>1</sup>; Wilhelmus Kersten<sup>1</sup>; Guillaume Lessene<sup>1</sup>; John Silke<sup>1</sup>; Peter Czabotar<sup>1</sup>; Andrew Webb<sup>1</sup>; James Murphy<sup>1</sup>; <sup>1</sup>Walter & Eliza Hall Institute, Parkville, Australia; <sup>2</sup>The University of Melbourne, Parkville, Australia; <sup>3</sup>LaTrobe University, Bundoora, Australia

10:30 am - 12:20 pm Tuesday  
**STRUCTURAL PROTEOMICS**  
 Session Chairs: Patrick Griffin and Carol Robinson  
 Oceana 6

- TOC am 10:30 **Different Means to Solubilize Membrane Proteins for MS Analysis: Going More Native;** Nina Morgner; Oliver Peetz; Nils Hellwig; Goethe University Frankfurt/Main, Frankfurt/Main, Germany
- TOC am 10:50 **Photo-Crosslinking Mass Spectrometry and Integrative Modeling Enables Rapid Screening of Antigen Interactions Involving Bacterial Transferrin Receptors;** Daniel S. Ziemianowicz; Dixon Ng; Anthony B. Schryvers; David Schriemer; University of Calgary, Calgary, Canada
- TOC am 11:10 **Deciphering the Role of Histone H2A Proteolysis during Stem Cell Differentiation and its Consequence in Nucleosome Stability;** Marie Coradin<sup>1</sup>; Kelly Karch<sup>1,2</sup>; Enrique Lin-Shiao<sup>1,2</sup>; Simone Sidoli<sup>1,2</sup>; Shelley Berger<sup>1,2</sup>;

10:30 am - 12:20 pm Tuesday  
**HPP: METABOLIC REMODELING AND HUMAN DISEASE**  
 Session Chairs: Fernando Corrales and Ferdinando Cerchiello  
 Oceana 3-5

- TOD am 10:30 **Nonalcoholic Fatty Liver Disease Diversity: Learning from Mouse Models;** Cristina Alonso; OWL Metabolomics, Derio, Spain
- TOD am 10:50 **Technology for Clinical Proteomics and Its Application to Liver Disease;** Matthias Mann; Max Planck Institute of Biochemistry, Martinsried, Germany
- TOD am 11:10 **Quantitative Targeted Proteomic Analysis of One-Carbon Metabolism Proteins in Human Liver Cancer;** Fernando Corrales<sup>1</sup>; Alberto Paradela<sup>4</sup>; Ver nica Ambao<sup>2</sup>; Ignacio Granero<sup>4</sup>; Bruno Sangro<sup>3</sup>; <sup>1</sup>Centro Nacional de Biotecnolog a, CSIC; CIBEREHD, Madrid, Spain; <sup>2</sup>Centro de Investigaciones Endocrinol gicas CONICET, Buenos Aires,

TUESDAY, OCTOBER 2

- Argentina; <sup>3</sup>Clínica Universidad de Navarra-  
IDISNA and CIBEREHD, Pamplona, Spain;  
<sup>4</sup>Centro Nacional de Biotecnología, CSIC,  
Madrid, Spain
- TOD am 11:22 **Systems Analysis Reveals  
Phosphatidylcholine Metabolism  
Changes in Relapsed Multiple Myeloma;**  
Ahmed Mohamed<sup>1</sup>; Joel Collins<sup>3, 4</sup>; Hui  
Jiang<sup>2</sup>; Jeffrey Molendijk<sup>2</sup>; Thomas Stoll<sup>2</sup>;  
Kate Markey<sup>1, 5</sup>; Michelle Hill<sup>1, 2</sup>; <sup>1</sup>QIMR  
Berghofer Medical Research Institute, and  
The, Brisbane, Australia; <sup>2</sup>UQ Diamantina  
Institute, University of Queensland,  
Brisbane, Australia; <sup>3</sup>Princess Alexandra  
Hospital, Brisbane, Australia; <sup>4</sup>Toowoomba  
Hospital, Toowoomba, Australia; <sup>5</sup>Royal  
Brisbane and Women's Hospital, Brisbane,  
Australia
- TOD am 11:34 **NEDDylated Proteome in Non Alcoholic  
Fatty Liver Disease;** Marina Serrano-  
Maciá<sup>1</sup>; Mikel Azkargorta<sup>1, 4</sup>; Jorge Simón<sup>1</sup>;  
Naroa Goikoetxea-Usandizaga<sup>1</sup>; Teresa  
Cardoso<sup>1</sup>; David Fernandez-Ramos<sup>1</sup>;  
Virginia Gutierrez de Juan<sup>1</sup>; Marta Varela-  
Rey<sup>1, 5</sup>; Pablo Fernandez-Tussy<sup>1</sup>; Fernando  
Lopitz-Otsoa<sup>1</sup>; Patricia Aspichueta<sup>2</sup>; Paula  
Iruzubieta<sup>2</sup>; Javier Crespo<sup>2</sup>; Selly C Lu<sup>3</sup>;  
José M. Mato<sup>1</sup>; Felix Elortza<sup>1, 4</sup>; María Luz  
Martinez-Chantar<sup>1, 5</sup>; <sup>1</sup>CIC bioGUNE, Derio,  
Spain; <sup>2</sup>UPV/EHU, Leioa, Spain; <sup>3</sup>Cedars-  
Sinai Medical Center, LA, CA; <sup>4</sup>ProteoRed-  
ISCIII, Derio, Spain; <sup>5</sup>CIBERehd, Derio,  
Spain
- TOD am 11:46 **Comparative Proteomic and Lipidomic  
Profiling Reveals Broad Dysregulation of  
Lipid Metabolism in Triple-Negative  
Breast Cancer Development;** Ling Lin<sup>1</sup>;  
Songping Lin<sup>2</sup>; Huali Shen<sup>1</sup>; Pengyuan  
Yang<sup>1</sup>; <sup>1</sup>Fudan University, Shanghai, China;  
<sup>2</sup>Affiliated Union Hospital of Fujian Medical  
Univer, Fujian, China
- TOD am 11:58 **A Molecular Portrait of Ground State  
Pluripotency;** Ana Martinez-Val<sup>1</sup>; Cian  
Lynch<sup>2, 3</sup>; Manuel Serrano<sup>2, 3</sup>; Javier  
Muñoz<sup>1</sup>; <sup>1</sup>Proteomics Unit, CNIO, Madrid,  
Spain; <sup>2</sup>Cellular Plasticity and Disease  
Group, IRB, Barcelona, Spain; <sup>3</sup>Tumour  
Suppression Group, CNIO, Madrid, Spain
- 10:30 am - 12:20 pm Tuesday  
RARE DISEASES  
Session Chair: Nicholas J. Schork  
Oceana 1-2
- TOE am 10:30 **Integrated Approaches to Patient-  
Specific Research;** Nicholas J. Schork<sup>1, 2</sup>;  
<sup>1</sup>Translational Genomics Research Institute  
(TGen), Phoenix, AZ; <sup>2</sup>City of Hope/TGen  
IMPACT Center, Duarte, CA
- TOE am 10:50 **Pediatric Cancer: A Genomics-based  
Study of a Rare Disease;** Elaine R. Mardis;  
Nationwide Children's Hospital, Columbus,  
OH
- TOE am 11:10 **Quantitative Proteomic Analyses of  
Uterine Leiomyomas from Hereditary  
Leiomyomatosis and Renal Cell Cancer  
Patients.;** Thomas Conrads<sup>1</sup>; Christopher  
Tarney<sup>2</sup>; Nicholas Bateman<sup>2</sup>; Niyati Parikh<sup>2</sup>;  
Ming Zhou<sup>1</sup>; Kelly Conrads<sup>2</sup>; James  
Segars<sup>3</sup>; Paul Driggers<sup>3</sup>; Yovanni  
Casablanca<sup>2</sup>; Chad Hamilton<sup>2</sup>; G. Larry  
Maxwell<sup>4</sup>; <sup>1</sup>Inova Schar Cancer Institute,  
Falls Church, VA; <sup>2</sup>Gynecologic Cancer  
Center of Excellence, Annandale, VA;  
<sup>3</sup>Johns Hopkins School of Medicine,  
Baltimore, MD; <sup>4</sup>Obstetrics and Gynecology,  
Inova Fairfax Hospital, Falls Church, VA
- TOE am 11:22 **Plasma Proteomic Profiling to Identify  
Potential Biomarkers for Early Diagnosis  
of Multiple Myeloma in Premalignant  
Disease;** Yurany Moreno<sup>1</sup>; Nicola J  
Weston-Bell<sup>1</sup>; Kate Vandylke<sup>3</sup>; Duncan  
Hewett<sup>3</sup>; Andrew Zannettino<sup>3</sup>; Spiros  
Garbis<sup>2</sup>; Surinder S Sahota<sup>1</sup>; <sup>1</sup>Tumour  
Immunogenetics Group, University of  
Southampton, United Kingdom; <sup>2</sup>Proteomics  
Group, University of Southampton, United  
Kingdom; <sup>3</sup>Myeloma Research Laboratory,  
The University of Adelaide, Australia
- TOE am 11:34 **Angiotensin II Signature Proteins as  
Non-Invasive Markers of Fibrosis in  
Kidney Transplant Recipients;** Zahraa  
Mohammed-Ali<sup>1</sup>; Tomas Tokar<sup>1</sup>; Ihor  
Batruch<sup>2</sup>; Shelby Reid<sup>3</sup>; Alexandre Tavares-  
Brum<sup>4</sup>; Paul Yip<sup>1</sup>; Héloïse Cardinal<sup>4</sup>;  
Marie-Josée Hébert<sup>4</sup>; Yanhong Li<sup>1</sup>; S.  
Joseph Kim<sup>1, 3</sup>; Igor Jurisica<sup>1, 3</sup>; Rohan  
John<sup>1</sup>; Ana Konvalinka<sup>1</sup>; <sup>1</sup>University Health  
Network, Toronto, ON, Canada; <sup>2</sup>Lunenfeld-  
Tanenbaum Research Institute, Toronto,  
ON, Canada; <sup>3</sup>University of Toronto,  
Toronto, ON, Canada; <sup>4</sup>Centre Hospitalier  
de l'Université de Montréal, Montréal, QC,  
Canada
- TOE am 11:46 **Aptamer-based Proteomics Identifies  
Potential Predictive Biomarkers of  
Doxorubicin-induced Cardiotoxicity;** Li-  
Rong Yu<sup>1</sup>; Jaclyn Daniels<sup>1</sup>; Zhijun Cao<sup>1</sup>;  
Richard Beger<sup>1</sup>; Issam Makhoul<sup>2</sup>; Angela  
Pennisi<sup>2</sup>; Jeanne Wei<sup>2</sup>; Jane Bai<sup>3</sup>; Julia  
Lathrop<sup>4</sup>; Jinong Li<sup>5</sup>; Valentina Todorova<sup>2</sup>;  
<sup>1</sup>National Center for Toxicological  
Research, FDA, Jefferson, AR; <sup>2</sup>University  
of Arkansas for Medical Sciences, Little  
Rock, AR; <sup>3</sup>Center for Drug Evaluation and  
Research, FDA, Silver Spring, MD; <sup>4</sup>Center  
for Biologics Evaluation and Research,  
FDA, Silver Spring, MD; <sup>5</sup>Center for Devices  
and Radiological Health, FDA, Silver  
Spring, MD
- TOE am 11:58 **Targeted Proteomic Analysis of  
Formalin-Fixed Paraffin Embedded  
Prostate Biopsies with Outcomes Data to  
Identify Candidate Biomarkers for  
Aggressive Prostate Cancer;** Yuqian  
Gao<sup>1</sup>; Hui Wang<sup>1</sup>; Denise Young<sup>2</sup>; Jennifer  
Cullen<sup>2, 3</sup>; Yingjie Song<sup>2</sup>; Yongmei Chen<sup>2</sup>;

TUESDAY, OCTOBER 2

Athena Schepmoes<sup>1</sup>; Gyorgy Petrovics<sup>2, 3</sup>;  
Thomas Fillmore<sup>1</sup>; Tujin Shi<sup>1</sup>; Wei-Jun  
Qian<sup>1</sup>; Richard Smith<sup>1</sup>; Sudhir Srivastava<sup>4</sup>;  
Jacob Kagan<sup>4</sup>; Albert Dobi<sup>2, 3</sup>; Inger  
Rosner<sup>2</sup>; Karin Rodland<sup>1</sup>; Isabell  
Sesterhenn<sup>5</sup>; Shiv Srivastava<sup>2</sup>; Tao Liu<sup>1</sup>;  
<sup>1</sup>Pacific Northwest National Laboratory,

Richland, WA; <sup>2</sup>Center for Prostate Disease  
Research, Bethesda, MD; <sup>3</sup>John P. Murtha  
Cancer Center, Bethesda, MD; <sup>4</sup>National  
Cancer Institute, Bethesda, MD; <sup>5</sup>Joint  
Pathology Center, Silver Springs, MD

12:20 – 1:45 pm Tuesday  
CORPORATE LUNCH SEMINARS or Lunch-on-your-own

RSVP Required for Corporate Lunch Seminars.



Banda Sea Room

**Robust Clinical Proteomics Workflow for Plasma and  
Cancer Tissue Analysis**; Matthias Mann, *Max Planck  
Institute for Biochemistry*; **Clinical Proteomics for  
Biomarker Qualification in Fabry Disease**; Petra Oliva,  
*Sanofi Genzyme*



Java Sea Room

**Precision Medicine**: Advancing Mass Spectrometry-based  
Large-cohort Proteomics for Precision Medicine – An  
International Cancer Moonshot Multi-site Study; Thomas  
Conrads, *Inova Schar Cancer Institute*; Yue Xuan, *Thermo  
Fisher Scientific*



Timor Sea Room

**Our time with PASEF on the timsTOF Pro**; Tharan  
Srikumar, *Princeton University*; **Urine Biomarker Discovery  
by Proteomics and Peptidomics: Towards "All-in-One  
Urine Test"**; Tadashi Yamamoto, *Niigata University*

Waters

THE SCIENCE OF WHAT'S POSSIBLE.®

South China Sea Room

**Precision, Accuracy, and Throughput in Clinical  
Proteomics**

Introduction of Absolute Quantification of Plasma Samples  
Based on Biognosys' PQ500 Reference Peptides; Roland  
Bruderer (Biognosys)

Application of SONAR for enhanced throughput analysis with  
an exploratory targeted reagent strategy in clinical proteomics  
research; Lee Gethings, (Waters Corp.)



Coral Sea Room

**The TripleTOF® System in the Lab – Practical  
Applications and Exciting Innovations**; Christie Hunter,  
*SCIEX*

2:00 - 3:50 pm Tuesday CARDIOLOGY Session Chairs: Peipei Ping and Pothur Srinivas Oceana 8-10		2:00 - 3:50 pm Tuesday COMPUTATIONAL ADVANCES Session Chairs: Nuno Bandeira and David Fenyo Oceana 7	
TOA pm 2:00	<b>Top-Down Proteomics in Cardiac Disease and Regeneration;</b> Ying Ge; <i>University of Wisconsin, Madison, WI</i>	TOB pm 2:00	<b>Imputing Missing Values in Proteomics Data from Mass Spectrometry based Experiments;</b> Pei Wang; <i>Icahn School of Medicine at Mount Sinai, New York, NY</i>
TOA pm 2:20	<b>Oxidative Stress Post-translational Modification Landscape in Cardiac Hypertrophy Revealed By Machine Learning Approaches;</b> Peipei Ping; <i>UCLA, Los Angeles, CA</i>	TOB pm 2:20	<b>Low Abundance Peptide Sequencing by Deep Learning;</b> Lei Xin <sup>1</sup> ; Hieu Tran <sup>2</sup> ; Xin Chen <sup>1</sup> ; Rui Qiao <sup>2</sup> ; CY Liu <sup>2</sup> ; <u>Baozhen Shan</u> <sup>1</sup> ; Ming Li <sup>2</sup> ; <sup>1</sup> <i>Bioinformatics Solutions Inc, Waterloo, Canada</i> ; <sup>2</sup> <i>University of Waterloo, Waterloo, Canada</i>
TOA pm 2:40	<b>Proteomic Mapping Reveals Differences in the Bioenergetics of the Heart;</b> Wendy Heywood <sup>1</sup> ; Richard Collis <sup>2, 4</sup> ; Jonathan Searle <sup>1</sup> ; Ivan Doykov <sup>1</sup> ; Caroline Coats <sup>2</sup> ; Michael Ashworth <sup>3</sup> ; Perry Elliott <sup>2, 4</sup> ; <u>Kevin Mills</u> <sup>1</sup> ; <sup>1</sup> <i>UCL Great Ormond Street Institute of Child Health, London, United Kingdom</i> ; <sup>2</sup> <i>UCL Institute of Cardiovascular Science, London, UK</i> ; <sup>3</sup> <i>Great Ormond Street Hospital, London, UK</i> ; <sup>4</sup> <i>Barts Hospital, London, UK</i>	TOB pm 2:40	<b>A Combined Identification and Quantification Error Model of Label-Free Protein Quantification;</b> Matthew The; <u>Lukas Käll</u> ; <i>KTH - Royal Inst of Technology, Stockholm, Sweden</i>
TOA pm 2:52	<b>Circulating Concentrations of MYDGF in Healthy Individuals and Patients with Myocardial Infarction as Assessed by a New, Quantitative LC-MS Assay;</b> <u>Andreas Pich</u> <sup>1</sup> ; Felix Polten <sup>1</sup> ; Marc Reboll <sup>1</sup> ; Kerstin Bethmann <sup>1</sup> ; Christian Widera <sup>1</sup> ; Evangelos Giannitsis <sup>2</sup> ; Jochen Tillmanns <sup>1</sup> ; Tibor Kempf <sup>1</sup> ; Johann Bauersachs <sup>1</sup> ; Kai Wollert <sup>1</sup> ; <sup>1</sup> <i>Hannover Medical School, Hannover, Germany</i> ; <sup>2</sup> <i>University of Heidelberg, Heidelberg, Germany</i>	TOB pm 2:52	<b>See Deeper with iSwathX: An Effortless Approach to Combine DDA Data Libraries for DIA Data Analysis;</b> <u>Zainab Noor</u> <sup>1</sup> ; Abidali Mohamedali <sup>1</sup> ; Mark S. Baker <sup>2</sup> ; Shoba Ranganathan <sup>1</sup> ; <sup>1</sup> <i>Department of Molecular Sciences, Macquarie Univer, Sydney, Australia</i> ; <sup>2</sup> <i>Department of Biomedical Sciences, Macquarie University, Sydney, Australia</i>
TOA pm 3:04	<b>Getting to the Heart of the Matter: Multispecies Heart Tissue Proteome Characterization;</b> <u>Joel Federspiel</u> <sup>1</sup> ; Caralynn Wilczewski <sup>2</sup> ; Laura Herring <sup>2</sup> ; Samvida Venkatesh <sup>1</sup> ; Lauren Wasson <sup>2</sup> ; Frank Conlon <sup>2</sup> ; Ileana Cristea <sup>1</sup> ; <sup>1</sup> <i>Princeton University, Princeton, NJ</i> ; <sup>2</sup> <i>University of North Carolina at Chapel Hill, Chapel Hill, NC</i>	TOB pm 3:04	<b>Topological Scoring of Protein Interaction Networks;</b> <u>Mihaela Sardu</u> <sup>1</sup> ; Joshua Gilmore <sup>2</sup> ; Brad Groppe <sup>3</sup> ; Arnob Dutta <sup>4</sup> ; Laurence Florens <sup>1</sup> ; Michael Washburn <sup>1, 5</sup> ; <sup>1</sup> <i>Stowers Institute for Medical Research, Kansas City, MO</i> ; <sup>2</sup> <i>Boehringer Ingelheim Vetmedica, St. Joseph, MO</i> ; <sup>3</sup> <i>Thermo Fisher Scientific, Waltham, MA</i> ; <sup>4</sup> <i>University of Rhode Island, Kingston, RI</i> ; <sup>5</sup> <i>The University of Kansas Medical Center, Kansas City, MO</i>
TOA pm 3:16	<b>Phosphoproteomic Profiling Reveals Perturbed Cardiac Signaling in Dilated Cardiomyopathy Patients;</b> <u>Sina Hadipour-Lakmehsari</u> <sup>1</sup> ; Uros Kuzmanov <sup>1</sup> ; Andrew Emili <sup>1</sup> ; Gavin Oudit <sup>2</sup> ; Anthony Gramolini <sup>1</sup> ; <sup>1</sup> <i>University of Toronto, Toronto, Canada</i> ; <sup>2</sup> <i>University of Alberta, Edmonton, Canada</i>	TOB pm 3:16	<b>The Functional Human Phospho-Proteome;</b> David Ochoa; Andrew Jarnuczak; Pedro Beltrao; <u>Juan Antonio Vizcaino</u> ; <i>EMBL-European Bioinformatics Institute (EMBL-EBI), Hinxton, United Kingdom</i>
TOA pm 3:28	<b>Global Proteomic and Transcriptomic Analyses Identify a Profile that Distinguishes Advanced Heart Failure Patients Capable of Cardiac Recovery Following LVAD-Unloading;</b> Christopher Tracy; Aman Makaju; Rachit Badolia; Sutip Navankasattusas; Dinesh Ramadurai; Anna Bakhtani; Lauren McCreath; Nikolaos Diakos; Craig Selzman; Stavros Drakos; <u>Sarah Franklin</u> ; <i>University of Utah, Salt Lake City, UT</i>	TOB pm 3:28	<b>Functional 5' UTR Motif Discovery with LESMoN: Local Enrichment of Sequence Motifs in Biological Networks;</b> <u>Mathieu Lavallée-Adam</u> <sup>1, 2</sup> ; Philippe Cloutier <sup>3</sup> ; Benoit Coulombe <sup>3, 4</sup> ; Mathieu Blanchette <sup>1</sup> ; <sup>1</sup> <i>McGill University, Montreal, Canada</i> ; <sup>2</sup> <i>University of Ottawa, Ottawa, Canada</i> ; <sup>3</sup> <i>Institut de recherches cliniques de Montréal, Montreal, Canada</i> ; <sup>4</sup> <i>Université de Montréal, Montreal, Canada</i>

**2:00 - 3:50 pm Tuesday**  
**NEW TECHNOLOGICAL ADVANCEMENTS**  
**IN PROTEOMICS**  
**Session Chairs: Kimberly Lee and Julian Saba**  
**Oceana 6**

- TOC pm 2:00 **Scratching the Surface: Ligand-Based Receptor Capture Methodologies to Explore the Surfaceome of Living Cells and Its Interactors**; Maria Pavlou; Laura A. Lopez-Garcia; Sandra Marder; Levent Demiray; Paul Helbling; *Dualsystems Biotech AG, Schlieren, Switzerland*
- TOC pm 2:20 **Integrating Cryo Electron Microscopy and LC-MS/MS for In-Depth Characterization of Viral Glycoproteins**; Joost Snijder<sup>1</sup>; David Veesler<sup>2</sup>; <sup>1</sup>*Snijder Bioscience, Arnhem, The Netherlands*; <sup>2</sup>*University of Washington, Seattle, WA*
- TOC pm 2:40 **Evaluation of a Novel LC System that Embeds Analytes in Pre-Formed Gradients for Rapid, Ultra-Robust Proteomics**; Nicolai Bache<sup>1</sup>; Philipp E. Geyer<sup>2</sup>; Dorte B. Bekker-Jensen<sup>3</sup>; Ole Hoerning<sup>1</sup>; Lasse Falkenby<sup>1</sup>; Peter V. Treit<sup>2</sup>; Sophia Doll<sup>2</sup>; Igor Paron<sup>2</sup>; Florian Meier<sup>2</sup>; Jesper V. Olsen<sup>3</sup>; Ole Vorm<sup>1</sup>; Matthias Mann<sup>2</sup>; <sup>1</sup>*EVOSEP Biosystems, Odense, Denmark*; <sup>2</sup>*Max Planck Institute of Biochemistry, Martinsried, Germany*; <sup>3</sup>*University of Copenhagen, Copenhagen, Denmark*
- TOC pm 2:52 **Efficient Top-Down Characterization of Native Proteoforms by Electron Capture Dissociation at Chromatographic Speeds with Q-ToF and Orbitrap Instruments**; Joseph S Beckman<sup>1</sup>; Valery G. Voinov<sup>2</sup>; Yury Vasil'ev<sup>2</sup>; Blaine Roberts<sup>3</sup>; Jared Shaw<sup>4</sup>; <sup>1</sup>*Oregon State University, Corvallis, OR*; <sup>2</sup>*e-MSion, Inc, Corvallis, OR*; <sup>3</sup>*Florey Institute, Melbourne, AU*; <sup>4</sup>*PNNL, Richland, WA*
- TOC pm 3:04 **Mass Spectrometry Imaging as Tool in Cancer Diagnostics and Cancer Drug Testing Platform**; Peter Hoffmann<sup>1</sup>; Parul Mittal<sup>2</sup>; Mitchell Acland<sup>2</sup>; Georgia Arentz<sup>1</sup>; Gurjeet Kaur<sup>3</sup>; Martin K. Oehler<sup>4</sup>; <sup>1</sup>*Future Industries Institute, UniSA, Mawson Lakes, Australia*; <sup>2</sup>*Adelaide Proteomics Centre, Adelaide, Australia*; <sup>3</sup>*Research in Molecular Medicine (INFORMM), Penang, Malaysia*; <sup>4</sup>*Department of Gynaecological Oncology, RAH, Adelaide, Australia*
- TOC pm 3:16 **Digging Deeper into the Plasma Proteome, a Novel Nanoflow LCMS Approach using Micro Pillar Array Columns (µPAC™)**; Jeff Op De Beeck<sup>1</sup>; Paul Jacobs<sup>1</sup>; Natalie Van Landuyt<sup>1</sup>; Wim De Malsche<sup>2</sup>; Gert Desmet<sup>2</sup>; Jarne Pauwels<sup>3</sup>; An Staes<sup>3</sup>; Francis Impens<sup>3</sup>; Kris Gevaert<sup>3</sup>; <sup>1</sup>*PharmaFluidics, Zwijnaarde, Belgium*; <sup>2</sup>*Vrije Universiteit Brussel, Brussel,*

*Belgium*; <sup>3</sup>*VIB Proteomics Expertise Center, UGent, Gent, Belgium*

TOC pm 3:28

**Comprehensive Single-Shot Proteomics Experiments with LC-FAIMS-MS/MS**; Alexander S. Hebert<sup>1</sup>; Satendra Prasad<sup>2</sup>; Michael W. Belford<sup>2</sup>; Derek J. Bailey<sup>2</sup>; Susan E. Abbatiello<sup>2</sup>; Romain Huguet<sup>2</sup>; Graeme C. McAlister<sup>2</sup>; Eloy R. Wouters<sup>2</sup>; Jean-Jacques Dunyach<sup>2</sup>; Dain Brademan<sup>1</sup>; Michael S Westphall<sup>1</sup>; Joshua J. Coon<sup>1, 3</sup>; <sup>1</sup>*University of Wisconsin - Madison, Madison, WI*; <sup>2</sup>*Thermo Fisher Scientific, San Jose, CA*; <sup>3</sup>*Morgridge Institute for Research, Madison, WI*

**2:00 - 3:50 pm Tuesday**  
**HPP: HARNESSING THE IMMUNE SYSTEM TO**  
**FIGHT DISEASE**  
**Session Chairs: Arie Admon and Ileana Cristea**  
**Oceana 3-5**

- TOD pm 2:00 **A Tissue-based Draft Map of the Murine MHC class I Immunopeptidome**; Heiko Schuster<sup>2</sup>; Wenguang Shao<sup>1</sup>; Tobias Weiss<sup>3</sup>; Patrick Pedrioli<sup>1</sup>; Patrick Roth<sup>3</sup>; Michael Weller<sup>3</sup>; David Campbell<sup>4</sup>; Eric Deutsch<sup>4</sup>; Robert Moritz<sup>4</sup>; Oliver Planz<sup>2</sup>; Hans-Georg Rammensee<sup>2</sup>; Ruedi Aebersold<sup>1</sup>; Etienne Caron<sup>1</sup>; <sup>1</sup>*ETH Zurich, Zurich, Switzerland*; <sup>2</sup>*University of Tübingen, Tübingen, Germany*; <sup>3</sup>*University Hospital Zürich, Zürich, Switzerland*; <sup>4</sup>*Institute for Systems Biology, Seattle, WA*
- TOD pm 2:20 **Targeted Proteomics-Driven Computational Modeling of Macrophage Microbial Sensing Pathways**; Nathan Manes; Jessica Mann; Pauline Kaplan; Martin Meier-Schellersheim; Iain Fraser; Ronald Germain; Aleksandra Nita-Lazar; *National Institutes of Health, Bethesda, MD*
- TOD pm 2:40 **Comparison of the HLA Peptidome of Primary and PDX Human Tumors Enables Identification of Neoepitopes of Potential for Personalized Immunotherapy**; Nataly Nataly Rijensky<sup>1</sup>; Netta Shraga<sup>2</sup>; Eilon Barnea<sup>1</sup>; Eitan Rubin<sup>3</sup>; Yitzhak Haviv<sup>2</sup>; Arie Admon<sup>1</sup>; <sup>1</sup>*Technion-Israel Inst. of Tech, Haifa, Israel*; <sup>2</sup>*Bar Ilan University, Zfat, Israel*; <sup>3</sup>*Ben-Gurion University of the Negev, Beer-Sheva, Israel*
- TOD pm 2:52 **Microscale Cell Surface Capture Technology for Discovery of Cell Surface N-Glycoproteins and its Application to Human Primary Cardiomyocytes**; Rachel A Jones Lipinski; Matthew Waas; Ranjuna Weerasekera; Theodore R. Keppel; Christopher Ashwood; Rebekah L. Gundry; *Medical College of Wisconsin, Milwaukee, WI*
- TOD pm 3:04 **Gas Phase Ion Fractionation Provides Unparalleled Sensitivity for Proteomic and Immunopeptidomic Analyses**; Sibylle Pfammatter<sup>1</sup>; Eric Bonneil<sup>1</sup>; Joel Lanoix<sup>1</sup>; Marie-Pierre Hardy<sup>1</sup>; Satendra Prasad<sup>2</sup>;

TUESDAY, OCTOBER 2

Michael Belford<sup>2</sup>; Jean-Jacques Dunyach<sup>2</sup>;  
Claude Perreault<sup>1</sup>; Pierre Thibault<sup>1</sup>;  
<sup>1</sup>Universite de Montreal, Montreal, Canada;  
<sup>2</sup>Thermo Fisher Scientific, San Jose, CA

TOD pm 3:16 **Longitudinal Omics and Biomolecular Network Analysis Dissect Host-specific Immune Dynamics in Multihost Fungal disease**; Pooja Aggarwal; Kanika Narula; Sudip Ghosh; Rajul Tayal; Nirranjan Chakraborty; Subhra Chakraborty; *National Institute of Plant Genome Research, New Delhi, India*

TOD pm 3:28 **Computational Immunomics Reveals Differential Human Immune Recognition of the *Candida albicans* Cell Surface Proteome during Dimorphic Transition in Invasive Candidiasis**; Aida Pitarch; César Nombela; Concha Gil; *Complutense University, Madrid, Spain*

**2:00 - 3:50 pm Tuesday**  
**MICROBIOME AND PATHOGEN INFECTIONS**  
**Session Chair: Frank Schmidt and Concha Gil**  
**Oceana 1-2**

TOE pm 2:00 **Ionbot: A Novel, Fully Data-Driven Search Engine for Open Modification and Mutation Searches with Applications in Quantitative (Meta-)Proteomics**; Sven Degroeve; Lennart Martens; *VIB-UGent Center for Medical Biotechnology, Ghent, Belgium*

TOE pm 2:20 **Pathogen-Specific Monoclonal Antibodies Maintain Short-Chain Fatty Acids (SCFA) And The Intestinal Microbiome**; Sonja Hess; *Medimmune, Gaithersburg, MD*

TOE pm 2:40 **Multi-Omics Comparative Analysis Reveals Host Signaling Pathways Affected by the Gut Microbiota**; Nathan Manes<sup>1</sup>; Natalia Shulzhenko<sup>2</sup>; Arthur Nuccio<sup>1</sup>; Sara Azeem<sup>1</sup>; Andrey Morgun<sup>2</sup>; Aleksandra Nita-Lazar<sup>1</sup>; <sup>1</sup>*National Institutes of Health, Bethesda, MD*; <sup>2</sup>*Oregon State University, Corvallis, OR*

TOE pm 2:52 **Quantifying Functional Microbiomes using MetaQuant: An Integrated, Quantitative Metaproteomics Tool**

**Reveals Connections between Taxa, Function and Protein Expression in Microbiomes.**; Pratik Jagtap<sup>1</sup>; Caleb Easterly<sup>1</sup>; Nadia Szeinbaum<sup>2</sup>; Andrea Argentini<sup>3</sup>; Subina Mehta<sup>1</sup>; Ray Sajulga<sup>1</sup>; Bart Mesuere<sup>4</sup>; James Johnson<sup>1</sup>; Carolin Kolmeder<sup>5</sup>; Praveen Kumar<sup>1</sup>; Jennifer Glass<sup>2</sup>; Joel Rudney<sup>1</sup>; Lennart Martens<sup>3</sup>; Brook Nunn<sup>6</sup>; Timothy Griffin<sup>1</sup>; <sup>1</sup>*University of Minnesota, Minneapolis, Minnesota*; <sup>2</sup>*Georgia Tech, Atlanta, GA*; <sup>3</sup>*VIB-UGent Center for Medical Biotechnology, Ghent, Belgium*; <sup>4</sup>*University of Ghent, Ghent, Belgium*; <sup>5</sup>*University of Helsinki, Helsinki, Finland*; <sup>6</sup>*University of Washington, Seattle, WA*

TOE pm 3:04 **Strength in Numbers: Impact of Oligomerization of Antiviral Proteins in Immune Response**; Timothy Howard; Krystal Lum; Catherina Pan; Ileana Cristea; *Princeton University, Princeton, NJ*

TOE pm 3:16 **Exploring the Anti-Staphylococcal Antibody Response using a Bead-Based Array Approach**; Tanja Meyer; Stephan Michalik; Barbara Bröker; Uwe Völker; *University Medicine Greifswald, Greifswald, Germany*

TOE pm 3:28 **Bacterial Proteotyping using Machine Learning Defined Peptide Signatures and Validation on Q-Exactive HF-X Coupled to Capillary Flow Liquid Chromatography**; Florence Roux-Dalvai<sup>1</sup>; Clarisse Gotti-Barban<sup>1</sup>; Mickaël Leclercq<sup>1</sup>; Frédéric Fournier<sup>1</sup>; Marie-Claude Hélie<sup>2</sup>; Judith Marcoux<sup>1</sup>; Isabelle Kelly<sup>1</sup>; Tabiwang N. Arrey<sup>4</sup>; Julie Bestman-Smith<sup>3</sup>; Claire Daully<sup>5</sup>; Maurice Boissinot<sup>2</sup>; Michel G. Bergeron<sup>2</sup>; Arnaud Droit<sup>1</sup>; <sup>1</sup>*Proteomics Platform - CHU Quebec Laval University, Quebec, Canada*; <sup>2</sup>*Infectiology CHU Quebec Laval University, Quebec, Canada*; <sup>3</sup>*Enfant Jesus Hospital CHU Quebec ULaval, Quebec, Canada*; <sup>4</sup>*Thermo Fisher Scientific, Bremen, Germany*; <sup>5</sup>*Thermo Fisher Scientific, Paris, France*

TUESDAY, OCTOBER 2

3:50 - 5:00 pm  
POSTER SESSION  
Pacifica  
Even-numbers present.

5:00 - 5:30 pm  
HUPO GENERAL ASSEMBLY MEETING @  
INNOVATION STAGE

Flow into the Innovation Stage at the end of Tuesday afternoon posters.  
Grab a free beer, wine, or soda! Hear a brief update on HUPO happenings.



6:30 – 8:30 pm  
LUAU SOCIAL EVENT

Advance ticket required. Available for purchase at registration through 12:00 pm on Monday.

Polynesian-themed Luau cook-out dinner featuring entertainment (think fire-dancing).



Sponsored in part by



8:30 - 9:15 pm Wednesday  
**WEDNESDAY MORNING PLENARY**  
 Session Chair: John Yates  
 Oceana 6



**Mary Higby Schweitzer**  
 North Carolina State  
 University  
 ProteoPaleontology

WOA am 11:34 **Chemical Proteomics Unravel Protein Targets and Binding Sites of Sphingolipid like Small Molecules with Anticancer Properties;** Peter Kubiniok<sup>1</sup>; Alison McCracken<sup>2</sup>; Brendan Finicle<sup>2</sup>; Lorenzo Sernissi<sup>1</sup>; Stephen Hanessian<sup>1</sup>; Aimee Edinger<sup>2</sup>; Pierre Thibault<sup>1</sup>; <sup>1</sup>University of Montreal, Montreal, Canada; <sup>2</sup>University of California, Irvine, CA

WOA am 11:46 **Inside-out: Targeting Matrix Metalloproteinases and the Surrounding Proteome in the Healing Skin Wound by Hybrid Degradomics;** Simonas Savickas<sup>1,3</sup>; Tobias Kockmann<sup>2</sup>; Ulrich auf dem Keller<sup>1,3</sup>; <sup>1</sup>Technical University of Denmark, Lyngby, Denmark; <sup>2</sup>Functional Genomics Center Zurich, Zurich, Switzerland; <sup>3</sup>Institute of Molecular Health Science, Zurich, Switzerland

WOA am 11:58 **Ubiquitinome Dynamics upon Proteasome Modulation;** Lennart Van der Wal; Karel Bezstarosti; Karen Sap; Dick Dekkers; Erikjan Rijkers; Jeroen A.A. Demmers; Erasmus University Medical Center Rotterdam, Rotterdam, Netherlands

9:15 - 10:30 am  
**POSTER SESSION**  
 Pacifica

ALL posters present.  
 Please remove posters at 10:30 am.

10:30 am - 12:20 pm Wednesday  
**ACTIVITY / CHEMICAL PROTEOMICS**  
 Session Chairs: Marcus Bantscheff and Christopher Overall  
 Oceana 8-10

10:30 am - 12:20 pm Wednesday  
**BIOMARKERS, NON-CANCER**  
 Session Chairs: Karin Rodland and Hui Zhang  
 Oceana 7

WOA am 10:30 **Expanding the Cancer Cell DUBome using Advanced Chemoproteomics;** Adan Pinto-Fernandez<sup>1</sup>; Abigail Schofield<sup>1</sup>; Eidarus Salah<sup>1</sup>; Sebastian Mathea<sup>2</sup>; Simon Davis<sup>1</sup>; Philip Charles<sup>1</sup>; Roman Fischer<sup>1</sup>; Benedikt M. Kessler<sup>1</sup>; <sup>1</sup>University of Oxford, Oxford, UK; <sup>2</sup>Goethe-Universität, Frankfurt, Germany

WOA am 10:50 **Deciphering the Cell-Specific Redoxome by Chemical Proteomics;** Jing Yang; National Center for Protein Sciences, Beijing, China

WOA am 11:10 **Proteome Analysis using Label-free DARTS and LC-MS/MS Method Reveals a Target Protein of Small Molecule Inhibitor of Autophagy;** Hui-Yun Hwang<sup>1</sup>; Yoon Sun Cho<sup>1</sup>; Jin Young Kim<sup>2</sup>; Ki Na Yun<sup>2</sup>; Jong Shin Yoo<sup>2</sup>; György Marko-Varga<sup>1,3</sup>; Ho Jeong Kwon<sup>1</sup>; <sup>1</sup>Yonsei University, Seoul, South Korea; <sup>2</sup>Korea Basic Science Institute, Chungbuk, South Korea; <sup>3</sup>Lund University, Lund, Sweden

WOA am 11:22 **Photoaffinity Probes and Quantitative Proteomics Enable Assessment of Target Engagement and Compound Potency in Live Cells;** H.Christian Eberl; Johanna Vappiani; Anne J. Wagner; Stephanie Lehmann; Marcel Muelbaier; Marcus Bantscheff; Cellzome, a GSK company, Heidelberg, Germany

WOB am 10:30 **Cardiovascular Disease: Moving to Precision Medicine and Health;** Jennifer Van Eyk; Cedars Sinai Medical Center; Los Angeles, CA

WOB am 10:50 **Plasma Protein Biomarker Discovery and Down Selection for Subsequent Validation in The Environmental Determinants of Diabetes in the Young Cohort;** Bobbie-Jo Webb-Robertson<sup>1</sup>; Ernesto Nakayasu<sup>1</sup>; Charles Ansong<sup>1</sup>; Lisa Bramer<sup>1</sup>; Marina Gritsenko<sup>1</sup>; Therese Clauss<sup>1</sup>; Paul Piehowski<sup>1</sup>; Athena Schepmoes<sup>1</sup>; Bryan Stanfill<sup>1</sup>; Daniel Orton<sup>1</sup>; Ronald Moore<sup>1</sup>; Brigitte Frohnert<sup>2</sup>; Marian Rewers<sup>2</sup>; Richard Smith<sup>1</sup>; Jeffrey Krischer<sup>3</sup>; Thomas Metz<sup>1</sup>; <sup>1</sup>Pacific Northwest National Laboratory, Richland, WA; <sup>2</sup>University of Colorado School of Medicine, Denver, CO; <sup>3</sup>University of South Florida, Tampa, FL

WOB am 11:10 **Mitochondrial Proteins as Parkinson's Disease Circulatory Biomarkers – A Translational Study;** Sandra I. Anjo<sup>1,2</sup>; Patrícia Valério dos Santos<sup>3</sup>; Maria Luiza Constante Rosado<sup>4,5</sup>; Graça Baltazar<sup>4</sup>; Mário Grãos<sup>1,6</sup>; Bruno Manadas<sup>1</sup>; <sup>1</sup>Center for Neuroscience and Cell Biology, UC, Coimbra, Portugal; <sup>2</sup>Faculty of Medicine, UC, Coimbra, Portugal; <sup>3</sup>Centro Hospitalar de Setúbal, Setúbal, Portugal; <sup>4</sup>Faculty of Health Sciences, UBI, Covilhã, Portugal; <sup>5</sup>Centro Hospitalar Cova da Beira, E.P.E.,



Covilhã, Portugal; <sup>6</sup>Biocant, Biotechnology Transfer Association, Cantanhede, Portugal

WOB am 11:22 **Development a of Novel Targeted Proteomic Plasma Biomarker Panel for Hypertrophic Cardiomyopathy;** Wendy Heywood<sup>4</sup>; Gabriella Captur<sup>2</sup>; Caroline Coats<sup>2</sup>; Stefania Rosmini<sup>2</sup>; Vimal Patel<sup>3</sup>; Richard Collis<sup>2</sup>; Nina Patel<sup>4</sup>; Petros Syrris<sup>3</sup>; Paul Bassett<sup>1</sup>; Ben O'Brian<sup>2</sup>; James Moon<sup>2</sup>; Perry Elliott<sup>3</sup>; Kevin Mills<sup>4</sup>; <sup>1</sup>UCL, London, United Kingdom; <sup>2</sup>Barts Heart Centre, St Bartholemews Hospital, London, UK; <sup>3</sup>UCL Institute of Cardiovascular Science, London, UK; <sup>4</sup>UCL Great Ormond Street Institute of Child Health, London, UK

WOB am 11:34 **A Novel and Robust Method for Urinary Proteome Profiling Applied in an Exploratory Case-Control Study;** Ireshyn Govender<sup>1,2</sup>; Stoyan Stoychev<sup>1</sup>; Demetra Mavri-Damelin<sup>2</sup>; Ebrahim Variava<sup>2</sup>; Faheem Seedat<sup>2</sup>; Neil Martinson<sup>2</sup>; Dalu Mancama<sup>1</sup>; <sup>1</sup>Council for Scientific and Industrial Research, Pretoria, South Africa; <sup>2</sup>University of the Witwatersrand, Johannesburg, South Africa

WOB am 11:46 **Astroglial Injury-Defined Biomarkers for Assessment of Traumatic Brain Injury;** Ina-Beate Wanner; Julia Halford; Sean Shen; Joseph Loo; University of California, Los Angeles, Los Angeles, CA

WOB am 11:58 **Remodeling of the Glyco-phenotype of T Cell Surface Proteins with Antisense RNA of Human Immunodeficiency Virus;** Weiming Yang<sup>1</sup>; Minghui Ao<sup>1</sup>; Fabio Romerio<sup>2</sup>; Hui Zhang<sup>1</sup>; <sup>1</sup>Johns Hopkins University, Baltimore, MD; <sup>2</sup>University of Maryland, Baltimore, MD

10:30 am - 12:20 pm Wednesday  
SINGLE CELL PROTEOMICS  
Session Chair: Nikolai Slavov  
Oceana 6

This session is generously supported by



WOC am 10:30 **Dissecting the Spatiotemporal Subcellular Distribution of the Human Proteome;** Emma Lundberg; KTH Royal Institute of Technology, Stockholm, Sweden

WOC am 10:50 **Single-Molecule Protein Sequencing;** Jagannath Swaminathan; Alexander A. Boulgakov; Erik T. Hernandez; Angela M. Bardo; James L. Bachman; Joseph Marotta; Amber M. Johnson; Eric V. Anslyn; Edward M. Marcotte; University of Texas at Austin, Austin, TX

WOC am 11:10 **Automated Sample Preparation for High-Throughput Single-Cell Proteomics;** Harrison Specht<sup>1</sup>; Guillaume Harmange<sup>1</sup>; David Perlman<sup>1</sup>; Ed Emmott<sup>1</sup>; Zach Niziolek<sup>2</sup>; Bogdan Budnik<sup>2</sup>; Nikolai Slavov<sup>1</sup>; <sup>1</sup>Northeastern University, Boston, MA; <sup>2</sup>Harvard University, Cambridge, MA

WOC am 11:22 **Comparison of Novel Quantitative Methods for Single Cell Proteomics;** Akos Vegvari<sup>1</sup>; Christian Beusch<sup>1</sup>; Alexandra Alexandridou<sup>1</sup>; Amirata S Dibavar<sup>1</sup>; Jaakko S Teppo<sup>2</sup>; Roman A Zubarev<sup>1</sup>; <sup>1</sup>Karolinska Institutet, Stockholm, Sweden; <sup>2</sup>University of Helsinki, Helsinki, Finland

WOC am 11:34 **Cellular and Subcellular Heterogeneity of Metabolites, Lipids, and Peptides in Selected and Perturbed Cells Explored by Single-Cell Mass Spectrometry;** Linwen Zhang; Nikkita Khattar; Akos Vertes; The George Washington University, Washington, DC

WOC am 11:46 **Cell-Type Resolved Proteome and Lipidome of Human Lung at Population and Single Cell Level.;** Jeremy Clair; Jennifer Kyle; Ying Zhu; Lisa Bramer; Paul Piehowski; Ryan Kelly; Charles Ansong; PNNL, Richland, WA

WOC am 11:58 **PASEF for Ultra-Sensitive Shotgun Proteomics;** Romano Hebel<sup>1</sup>; Heiner Koch<sup>1</sup>; Christopher M. Adams<sup>2</sup>; Scarlet Koch<sup>1</sup>; Markus Lubeck<sup>1</sup>; Florian Meier<sup>3</sup>; Andreas Brunner<sup>3</sup>; Matthias Mann<sup>3</sup>; <sup>1</sup>Bruker Daltonik GmbH, Bremen, Germany; <sup>2</sup>Bruker Daltonics Inc., San Jose, USA; <sup>3</sup>Max Planck Institute of Biochemistry, Martinsried, Germany

**10:30 am - 12:20 pm Wednesday**  
**HPP: HUMAN CHEMOSENSATION:**  
**OLFACTION AND TASTE**  
**Session Chairs: Mark Baker and Jong Shin Yoo**  
**Oceana 3-5**

- WOD am 10:30 **Odorant Receptors and the Perception of Odors**; Andreas Keller; *Rockefeller University, New York, NY*
- WOD am 10:50 **Bitter Sweet Taste Proteins**; Danielle Reed; *Monell Chemical Senses Center, Philadelphia, PA*
- WOD am 11:10 **Comprehensive Proteomic Approaches for Membrane Proteins Analysis**; Jing Gao; Yanting Zhou; Hongwen Zhu; Zhenyun Zhu; Hu Zhou; *Shanghai Institute of Materia Medica*
- WOD am 11:22 **Changes in Proteome of Age-Related Maculopathy Susceptibility Protein 2 (ARMS2) Edited Retinal Pigment Epithelial Cells in Response to Oxidative Stress**; Meleha Ahmad<sup>1</sup>; Todd Greco<sup>2</sup>; Marisol Cano<sup>1</sup>; James Handa<sup>1</sup>; Karl Wahlin<sup>1</sup>; Donald Zack<sup>1</sup>; Srinivas Sripathi<sup>1</sup>; Ileana Cristea<sup>2</sup>; Richard Semba<sup>1</sup>; <sup>1</sup>*Wilmer Eye Institute, Baltimore, MD*; <sup>2</sup>*Princeton University, Princeton, NJ*
- WOD am 11:34 **Identification of Missing Proteins and Alternative Splicing Variants from Human Olfactory Epithelial Tissue**; Heeyoun Hwang<sup>1</sup>; Ji Eun Jeong<sup>1,2</sup>; Hyun Kyoung Lee<sup>1,2</sup>; Ki Na Yun<sup>1</sup>; Hyun Joo An<sup>2</sup>; Bonghee Lee<sup>3</sup>; Young-Ki Paik<sup>4</sup>; Gi Taek Yee<sup>5</sup>; Jin Young Kim<sup>1</sup>; Jong Shin Yoo<sup>1,2</sup>; <sup>1</sup>*KBSI, Chungju-Si, South Korea*; <sup>2</sup>*Chungnam National Univ, Daejeon, South Korea*; <sup>3</sup>*Gachon University, Incheon, South Korea*; <sup>4</sup>*Yonsei Univ, Seoul, South Korea*; <sup>5</sup>*Gil Medical Center, Incheon, South Korea*
- WOD am 11:46 **Identifiable Human Olfactory Receptor Proteome using High-Stringency Mass Spectrometry**; Subash Adhikari; Samridhi Sharma; Seong Beom Ahn; Mark S. Baker; *Macquarie University, Sydney, Australia*
- WOD am 11:58 **Speaker Roundtable**

**10:30 am - 12:20 pm Wednesday**  
**NEURODEGENERATIVE DISEASES**  
**Session Chair: Nicholas Seyfried**  
**Oceana 1-2**

- WOE am 10:30 Victor Faundez; *Emory University, Atlanta, GA*
- WOE am 10:50 Wilfried Rossoll; *Mayo Clinic, Jacksonville, FL*
- WOE am 11:10 **Large-scale Proteomic Analysis of Alternative Open Reading Frames Reveals Novel Proteins in Neurodegenerative Diseases such as Amyotrophic Lateral Sclerosis**; Marie Brunet; Jennifer Raisch; Mylene Brunelle; Jean-Francois Lucier; Jean-Fancois

Jacques; Nathalie Rivard; Xavier Roucou; *University of Sherbrooke, Sherbrooke, Canada*

- WOE am 11:22 **Hydralazine Induces Stress Resistance and Extends C. Elegans Lifespan by Activating the NRF2/SKN-1 Signalling Pathway**; Hamid Mirzaei; *UT Southwestern, Dallas, TX*
- WOE am 11:34 **Urine Proteomics for Biomarker Discovery in Neurodegenerative Disease**; Jenny Hällqvist<sup>1</sup>; Ross W. Paterson<sup>2</sup>; Robert Clayton<sup>1</sup>; Nick Fox<sup>2</sup>; Jonathan M. Schott<sup>2</sup>; Henrik Zetterberg<sup>3,5</sup>; Amanda Heslegrave<sup>3,4</sup>; Wendy Heywood<sup>1</sup>; Kevin Mills<sup>1</sup>; <sup>1</sup>*GOS Institute of Child Health, UCL, London, UK*; <sup>2</sup>*Dementia Research Centre, UCL IoN, London, UK*; <sup>3</sup>*Department of Molecular Neuroscience, UCL IoN, London, UK*; <sup>4</sup>*UK Dementia Research Institute at UCL, London, UK*; <sup>5</sup>*Clinical Neurochemistry Laboratory, SUH, Mölndal, Sweden*
- WOE am 11:46 **Autoantigen Anoctamin 2 Associates with Increased Risk for Multiple Sclerosis**; Peter Nilsson; *KTH - Royal Institute of Technology, Solna, Sweden*
- WOE am 11:58 **An update on the HUPO Brain Protein Project (HBPP)**; Klaus Oliver Schubert<sup>1,2</sup>; Katrin Marcus<sup>4</sup>; Robert McCollumsmith<sup>5</sup>; Daniel Martins-De-Souza<sup>6</sup>; Charlotte Teunissen<sup>7</sup>; Peter Nilsson<sup>3</sup>; <sup>1</sup>*Univ of Adelaide, Adelaide, Australia*; <sup>2</sup>*Northern Adelaide Mental Health Service, Salisbury, Australia*; <sup>3</sup>*KTH Royal Inst of Technology, Stockholm, Sweden*; <sup>4</sup>*Ruhr Univ, Bochum, Germany*; <sup>5</sup>*Univ of Toledo, Toledo, OH*; <sup>6</sup>*Univ of Campinas, Campinas, Brazil*; <sup>7</sup>*VU Univ, Amsterdam, Netherlands*

**12:20 – 1:45 pm Wednesday**  
**CORPORATE LUNCH SEMINARS or Lunch-on-your-own**  
 RSVP Required for Corporate Lunch Seminars.



Java Sea Room and South China Sea Room

**MS Toolbox for Systems Biology:**

- Understanding Interactions in Membrane Proteins – New Opportunities for Drug Discovery; Carol Robinson, *University of Oxford*

**MS Targeted Assays:**

- New Quantitative Proteomic Assays for Cancer Signaling Pathways Using Multiplex IP and Targeted Mass Spectrometry; Jonathan Krieger, *The Hospital for Sick Children*, and Bhavin Patel, *Thermo Fisher Scientific*

2:00 - 3:50 pm Wednesday  
**DRUG DISCOVERY**  
 Session Chair: Mike Tyers  
 Oceana 8-10

- WOA pm 2:00 **Protein Acylation Is a General Regulatory Mechanism in Biosynthetic Pathway of Acyl-CoA-Derived Bioactive Natural Products**; Minjia Tan; *Shanghai Institute of Materia Medica,, Shanghai, China*
- WOA pm 2:20 **Interactome Rewiring following Pharmacological Targeting of BET Bromodomains**; Jean-Philippe Lambert<sup>1</sup>; Sarah Picaud<sup>2</sup>; Takao Fujisawa<sup>3</sup>; Huayun Hou<sup>4</sup>; Pavel Savitsky<sup>2</sup>; Liis Uusküla-Reimand<sup>4</sup>; Gagan D. Gupta<sup>1</sup>; Hala Abdouni<sup>1</sup>; Zhen-Yuan Lin<sup>1</sup>; Monika Tucholska<sup>1</sup>; James D.R. Knight<sup>1</sup>; Beatriz Gonzalez-Badillo<sup>1</sup>; Nicole St-Denis<sup>1</sup>; Joseph A. Newman<sup>2</sup>; Manuel Stucki<sup>5</sup>; Laurence Pelletier<sup>1,4</sup>; Nuno Bandeira<sup>6</sup>; Michael D. Wilson<sup>4,4</sup>; Panagis Filippakopoulos<sup>2,3</sup>; Anne-Claude Gingras<sup>1,4</sup>; <sup>1</sup>*Lunenfeld-Tanenbaum Research Inst at Mt Sinai Hosp, Toronto, Canada*; <sup>2</sup>*Structural Genomics Consortium, Oxford, UK*; <sup>3</sup>*Ludwig Institute for Cancer Research, Oxford, UK*; <sup>4</sup>*University of Toronto, Toronto, Canada*; <sup>5</sup>*University of Zurich, Schlieren, Switzerland*; <sup>6</sup>*University of California, San Diego, San Diego, CA*
- WOA pm 2:40 **Multipronged Quantitative Proteomics Analyses Reveal Alterations in Kinase Activities as a Fundamental Mechanism of Action of Circadian Period Altering Drugs**; Sandipan Ray<sup>1,2</sup>; Akhilesh B. Reddy<sup>1</sup>; <sup>1</sup>*The Francis Crick Institute, 1 Midland Road, London NW1 1AT, United Kingdom*; <sup>2</sup>*Institute of Neurology, University College London, Queen Square, London WC1N 3BG, United Kingdom*
- WOA pm 2:52 **Limited Proteolysis as a Target Deconvolution Strategy in Mammals**; Nigel Beaton<sup>1</sup>; Roland Bruderer<sup>1</sup>; Ilaria Piazza<sup>2</sup>; Paola Picotti<sup>2</sup>; Lukas Reiter<sup>1</sup>; <sup>1</sup>*Biognosys, Schlieren, Switzerland*; <sup>2</sup>*ETH Zurich, Zurich, Switzerland*
- WOA pm 3:04 **Drug Effects on Protein Homeostasis**; Nico Zinn<sup>1</sup>; Maria Faelth-Savitiski<sup>1</sup>; Mikhail Savitski<sup>2</sup>; Giovanna Bergamini<sup>1</sup>; Marcus Bantscheff<sup>1</sup>; <sup>1</sup>*Cellzome, A GSK company, Heidelberg, Germany*; <sup>2</sup>*EMBL, Heidelberg, Germany*
- WOA pm 3:16 **Building a Toolkit for Studying the Orphan Kinome**; Laurie Parker<sup>1</sup>; Minervo Perez<sup>1,2</sup>; John Blankenhorn<sup>1</sup>; Lindsay Breidenbach<sup>1</sup>; Hannah Peterson<sup>1</sup>; <sup>1</sup>*University of Minnesota, Minneapolis*; <sup>2</sup>*Purdue University, West Lafayette, IN*

2:00 - 3:50 pm Wednesday  
**NEW MASS SPEC TECHNOLOGIES**  
 Session Chairs: Lissa Anderson and John Yates  
 Oceana 7

Livia Eberlin's invited talk is generously supported by



- WOB pm 2:00 **Development of the MasSpec Pen for Tissue Analysis and Cancer Diagnosis**; Livia Eberlin; *University of Texas, Austin, TX*
- WOB pm 2:20 **The Important Role of Proteoforms in Human Health and Disease: The Significance and Relevance of the Cell-Based Human Proteome Project**; Neil Kelleher; *Northwestern University, Evanston, IL*
- WOB pm 2:40 **Trapped Ion Mobility Mass Spectrometry for Improved Sensitivity and Fastest Proteomics**; Oliver Raether<sup>1</sup>; Markus Lubeck<sup>1</sup>; Heiner Koch<sup>1</sup>; Scarlet Koch<sup>1</sup>; Florian Meier<sup>2</sup>; Andreas Brunner<sup>2</sup>; Juergen Cox<sup>2</sup>; Matthias Mann<sup>2</sup>; <sup>1</sup>*Bruker Daltonik GmbH, Bremen, Germany*; <sup>2</sup>*Max Planck Institute of Biochemistry, Martinsried, Germany*
- WOB pm 2:52 **EASI-tag Enables Accurate Multiplexed and Interference-free MS2-based Proteome Quantification**; Sebastian Virreira Winter; Florian Meier; Christoph Wichmann; Juergen Cox; Matthias Mann; Felix Meissner; *Max Planck Institute of Biochemistry, Martinsried/Munich, Germany*
- WOB pm 3:04 **Molecularly Imprinted Polymers (MIPs) for the Detection of Low Abundance Proteins as Biomarkers for Lung Cancer**; Rachel Norman; Sergey Piletsky; Leong Ng; Antonio Guerreiro; Francesco Canfarotta; Donald Jones; *University of Leicester, Leicester, United Kingdom*
- WOB pm 3:16 **Protein Analysis using Sub-Nanopore Sensors**; Mikhail Kolmogorov<sup>1</sup>; Gregory Timp<sup>2</sup>; Pavel Pevzner<sup>1</sup>; <sup>1</sup>*UC San Diego, La Jolla, California*; <sup>2</sup>*University of Notre Dame, Notre Dame, IN*
- WOB pm 3:28 **MaxQuant Software for Ion Mobility Enhanced Shotgun Proteomics**; Nikita Prianichnikov<sup>1</sup>; Christoph Wichmann<sup>1</sup>; Scarlet Beck<sup>2</sup>; Heiner Koch<sup>2</sup>; Markus Lubeck<sup>2</sup>; Romano Hebel<sup>2</sup>; Juergen Cox<sup>1</sup>; <sup>1</sup>*Max Planck Institute of Biochemistry, Martinsried, Germany*; <sup>2</sup>*Bruker Daltonik GmbH, Bremen, Germany*

**2:00 - 3:50 pm Wednesday  
METABOLISM  
Session Chair: Robert Moritz  
Oceana 6**

- WOC pm 2:00 **Lipids: Why Bother? From R&D to Clinical Applications;** Anne K. Bendt; Tze Ping Loh; Markus R. Wenk; *National University of Singapore, Singapore*
- WOC pm 2:20 **Healthy Lipidome Is a Resource for Understanding Metabolic Diseases;** Andrej Shevchenko; *MPI of Molecular Cell Biology and Genetics, Dresden, Germany*
- WOC pm 2:40 **Plasma Biomarkers of Cardiovascular Disease Associated with Type 1 Diabetes in Children;** Chantal Attard<sup>1</sup>; Stefan Bjelosevic<sup>2</sup>; Jasmine Wong<sup>1</sup>; Fergus Cameron<sup>1, 3</sup>; Paul Monagle<sup>2, 3</sup>; Vera Ignjatovic<sup>1, 2</sup>; <sup>1</sup>*Murdoch Children's Research Institute, Parkville, Australia;* <sup>2</sup>*The University of Melbourne, Parkville, Australia;* <sup>3</sup>*Royal Children's Hospital, Parkville, Australia*
- WOC pm 2:52 **Multi-omic Profiling of the Liver in a Rat Model of Type 2 Diabetes;** Desmond Li<sup>1</sup>; Lauren Smith<sup>1</sup>; Yen Chin Koay<sup>1, 2</sup>; Holly McEwen<sup>1, 3</sup>; Anthony Don<sup>1, 3</sup>; John O'Sullivan<sup>1, 2</sup>; Stuart Cordwell<sup>1</sup>; Melanie White<sup>1</sup>; <sup>1</sup>*University of Sydney, Sydney, Australia;* <sup>2</sup>*Heart Research Institute, Sydney, Australia;* <sup>3</sup>*ACRF Centenary Cancer Research Centre, Sydney, Australia*
- WOC pm 3:04 **Urinary Peptidomic Analysis Reveals Bioactive Uromodulin Peptides in Early Type 1 Diabetes;** Julie Van<sup>1</sup>; Sergi Clotet Freixas<sup>2</sup>; Xiaohua Zhou<sup>1</sup>; Ihor Batruch<sup>3</sup>; Etienne Sochett<sup>1</sup>; Farid Mahmud<sup>1</sup>; Eleftherios Diamandis<sup>1, 3</sup>; James Scholey<sup>1, 2</sup>; Ana Konvalinka<sup>1, 2</sup>; <sup>1</sup>*University of Toronto, Toronto, Canada;* <sup>2</sup>*University Health Network, Toronto, Canada;* <sup>3</sup>*Mount Sinai Hospital, Toronto, Canada*
- WOC pm 3:16 **Elucidating Changes in Plasma Protein Profiles related to Bariatric Surgery: An IMI DIRECT Study;** Ragna Häussler<sup>1</sup>; Matilda Dale<sup>1</sup>; Mun-Gwan Hong<sup>1</sup>; Cecilia Thomas<sup>1, 2</sup>; Johann Gassenhuber<sup>3</sup>; Violeta Raverdy<sup>4</sup>; Francois Pattou<sup>4</sup>; Jochen Schwenk<sup>1</sup>; <sup>1</sup>*Science for Life Laboratory, Solna, Sweden;* <sup>2</sup>*The Novo Nordisk Foundation Center, Copenhagen, Denmark;* <sup>3</sup>*Sanofi-Aventis Deutschland GmbH, Frankfurt am Main, Germany;* <sup>4</sup>*Inserm U1190, Lille, France*
- WOC pm 3:28 **A Systematic Map of Protein-Metabolite Interactions Reveals Principles of Chemical Communication;** Ilaria Piazza; Paola Picotti; *ETH Zurich, Zürich, Switzerland*

**2:00 - 3:50 pm Wednesday  
HPP: UNRAVELLING TISSUE PATHOLOGY THROUGH CELL MAPPING  
Session Chairs: Justyna Fert-Bober and Emma Lundberg  
Oceana 3-5**

- WOD pm 2:00 **Highly Multiplexed Imaging of Tissues with Subcellular Resolution by Imaging Mass Cytometry.;** Bernd Bodenmiller; *University of Zurich, Zurich, Switzerland*
- WOD pm 2:30 **Integrated omics for Spatial Mapping of the Human Proteome - Understanding the Molecular Repertoire in Health and Disease;** Cecilia Lindskog; *Uppsala University, Uppsala, Sweden*
- WOD pm 3:00 **Mass Spectrometry Imaging of the Metabolome and Lipidome Through Development of an Innovative Ionization Method;** David Muddiman; *North Carolina State University, Raleigh, NC*
- WOD pm 3:20 **A High-Precision Tissue-based Mouse Proteome with BoxCar and Data-Independent Acquisition;** Florian Meier<sup>1</sup>; Oliver Bernhardt<sup>2</sup>; Marta Murgia<sup>1</sup>; Catherine Vasilopoulou<sup>1</sup>; Michael Wierer<sup>1</sup>; Lynn Verbeke<sup>2</sup>; Tejas Gandhi<sup>2</sup>; Lukas Reiter<sup>2</sup>; Matthias Mann<sup>1</sup>; <sup>1</sup>*MPI of Biochemistry, Martinsried, Germany;* <sup>2</sup>*Biognosys AG, Schlieren, CH*

**2:00 - 3:50 pm Wednesday  
PERSONALIZED WELLNESS  
Session Chairs: Sanjeeva Srivastava and Mike Snyder  
Oceana 1-2**

- WOE pm 2:00 **The Value of Imputing -Omics Data into Biobanks Linked to Electronic Health Records;** Nancy J. Cox; *Vanderbilt University, Nashville, TN*
- WOE pm 2:20 **Proteomics Analysis in Context of Personal, Dense, Dynamic Data Clouds from Thousands of People;** Nathan Price; *Institute for Systems Biology, Seattle, WA*
- WOE pm 2:40 **Integrative Proteomics and Transcriptomics for Personalized Wellness: Using Saliva and Blood to Monitor Immune Response in Individuals;** George Mias; *Michigan State University, East Lansing, MI*
- WOE pm 2:52 **Longitudinal Multi-omics Profiling in Insulin Resistant and Sensitive Prediabetic Population;** Sara Ahadi; Wenyu Zhou; Reza Sailani; Kevin Contrepois; Mike Snyder; *Stanford University, Palo Alto, CA*

WOE pm 3:04

**Advancing Mass Spectrometry-based Large-Cohort Proteomics for Precision Medicine: An International Cancer Moonshot Multi- Site Study;** Yue Xuan<sup>1,2</sup>; Nicholas W. Bateman<sup>3</sup>; Sebastien Gallien<sup>2,14</sup>; Yue Zhou<sup>15</sup>; Niyati Parikh<sup>3</sup>; Mo Hu<sup>15</sup>; Pedro Navarro<sup>1</sup>; Yuju Chen<sup>4</sup>; Albert Sickmann<sup>5</sup>; Bernd Wollscheid<sup>6</sup>; Connie R. Jimenez<sup>7</sup>; Martin R. Larsen<sup>8</sup>; Hu Zhou<sup>9</sup>; Siqi Liu<sup>10</sup>; Zhinan Chen<sup>11</sup>; Thomas Kislinger<sup>12</sup>; Ben Crossett<sup>13</sup>; Brian Hood<sup>3</sup>; Reta Birhanu Kitata<sup>4</sup>; Christin Lorenz<sup>5</sup>; Christina Loosse<sup>5</sup>; Sandra Goetze<sup>6</sup>; Sander Piersma<sup>7</sup>; Davide Chiasserini<sup>7</sup>; Muhammad Tahir<sup>8</sup>; Hongwen Zhu<sup>9</sup>; Guixue Hou<sup>10</sup>; Xiuxuan Sun<sup>11</sup>; Andrew Macklin<sup>12</sup>; Amanda Khoo<sup>12</sup>; Benjamin L. Parker<sup>16</sup>; Stuart J. Cordwell<sup>16</sup>; Thomas P. Conrads<sup>3,17</sup>; <sup>1</sup>Thermo Fisher Scientific (Bremen) GmbH, Bremen, Germany; <sup>2</sup>Thermo Fisher Precision Medicine Science Center, Cambridge, MA; <sup>3</sup>Gynecologic Cancer Center of Excellence, HJF, Bethesda, MD; <sup>4</sup>Institute of Chemistry, Academia Sinica, Taipei, Taiwan; <sup>5</sup>Leibniz-Institut für Analytische Wissenschaften, Dortmund, Germany; <sup>6</sup>Institute of Molecular Systems Biology (IMSB), ETH, Zurich, Switzerland; <sup>7</sup>Dept. Medical Oncolog, VU University Medical Center, Amsterdam, Netherlands; <sup>8</sup>Dept. Biochemistry and Molecular Biology, SDU, Odense, Denmark; <sup>9</sup>Shanghai Institute of Materia Medica, Shanghai, China; <sup>10</sup>BGI-SHENZHEN, Shenzhen, China; <sup>11</sup>The Fourth Military Medical University, Xi'an, China; <sup>12</sup>Princess Margaret Cancer Centre, Toronto, Canada; <sup>13</sup>Sydney Mass Spectrometry, The University of Sydney, Sydney, Australia; <sup>14</sup>Thermo Fisher Scientific, Paris, France; <sup>15</sup>Thermo Fisher Scientific (China) Co. Ltd, Shanghai, China; <sup>16</sup>School of Life and Environ. Sci., Univ. Sydney, Sydney, Australia; <sup>17</sup>The Inova Schar Cancer Institute, Annandale, VA

WOE pm 3:16

**Mike Snyder;** *Stanford University, Stanford, CA*

WOE pm 3:28

**Speaker Roundtable**

4:00 - 5:00 pm  
**HUPO AWARDS SESSION**  
 Presentations for all HUPO Awards



5:00 - 5:30 pm Wednesday  
**WEDNESDAY AFTERNOON PLENARY**  
 Session Chairs: Ileana Cristea  
 Oceana 6



**Matthias Mann**  
*Max-Planck Institute for Biochemistry*

# Journal of Proteomics



An official journal of



**Editor-in-Chief: Juan J. Calvete**

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## POSTER LIST

All posters displayed Monday - Wednesday. Odd-numbered posters present Monday.

Even-numbered posters present Tuesday.

All posters present Wednesday morning.

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## POSTERS

All posters displayed Monday - Wednesday. Odd-numbered posters present Monday.  
Even-numbered posters present Tuesday. All posters present Wednesday morning.

### B/D HPP Posters 001-005

- Poster 001 **Subcellular Proteome Analysis of the Pancreatic Beta Cell lines: INS1e and MIN6;** Kyle McClary<sup>1</sup>; Sanraj Mittal<sup>1</sup>; John Yates, PhD<sup>2</sup>; Raymond C. Stevens, PhD<sup>1</sup>; <sup>1</sup>Univ of Southern California, Los Angeles, CA; <sup>2</sup>The Scripps Research Institute, La Jolla, CA
- Poster 002 **What are the 'Popular Proteins' in Rheumatic and Autoimmune Diseases?** Cristina Ruiz-Romero<sup>1</sup>; Maggie P. Y. Lam<sup>2</sup>; Peter Nilsson<sup>3</sup>; Patrik Önerfjord<sup>4</sup>; Paul J. Utz<sup>5</sup>; Jennifer E. Van Eyk<sup>6</sup>; Vidya Venkatraman<sup>6</sup>; Justyna Fert-Bober<sup>6</sup>; Fiona E. Watt<sup>7</sup>; Francisco Javier Blanco García<sup>1</sup>; <sup>1</sup>Proteomics Group GIR-ProteoRed/ISCIII INIBIC-CHUAC, A Coruña, Spain; <sup>2</sup>NIH BD2K Center of Excellence at UCLA, Los Angeles, United States; <sup>3</sup>Affinity Proteomics, SciLifeLab, KTH, Stockholm, Sweden; <sup>4</sup>Dpt. of Clinical Sciences, Section Rheumatology, Lund, Sweden; <sup>5</sup>Div. of Immunology and Rheumatology, Stanford Univ, Palo Alto, United States; <sup>6</sup>Dpt. Medicine and The Heart Institute, Cedars-Sinai, Los Angeles, United States; <sup>7</sup>Kennedy Institute of Rheumatology, Oxford Univ, Oxford, United Kingdom
- Poster 003 **Targeting Endothelial Erk1/2-Akt Axis as a Regeneration Strategy to Bypass Fibrosis during Chronic Liver Injury;** Ying Jiang; Yuanxiang Lao; Li Yanyan; He Fuchu; Beijing Institute of Lifeomics, Beijing, China
- Poster 004 **An Update on the Human Plasma Proteome;** Jochen Schwenk<sup>1</sup>; Eric Deutsch<sup>2</sup>; <sup>1</sup>Science for Life Laboratory, Solna, Sweden; <sup>2</sup>Institute for Systems Biology, Seattle, WA
- Poster 005 **SWATH-MS with Internal Landmarks for Quantitative Proteomics of Liver Cirrhosis Patient Urine;** Bo Xu<sup>1</sup>; Yoshitoshi Hirao<sup>1</sup>; Masaaki Takamura<sup>2</sup>; Keiko Yamamoto<sup>1</sup>; Amr Elguoshy<sup>1</sup>; Tadashi Yamamoto<sup>1</sup>; <sup>1</sup>BBC, Niigata University, Niigata, Japan; <sup>2</sup>Div. of Gast. and Hepat., Niigata University, Niigata, Japan

### BIOMARKERS Posters 006 - 035

- Poster 006 **Immunoproteomics Profiling of Citrullinated AAgeome Reveals Next-Generation Biomarkers for Rheumatoid Arthritis;** Wei Yu<sup>1</sup>; Hongye Wang<sup>1</sup>; Xiaolong Guan<sup>2</sup>; Haiyong Wang<sup>2</sup>; Fei Wang<sup>2</sup>; Lei Song<sup>1</sup>; Mingwei Liu<sup>1</sup>; Haoyu Wang<sup>3</sup>; Lihui Yang<sup>1</sup>; Jiayu Dai<sup>1</sup>; Te Liang<sup>1</sup>; Hu Duan<sup>1</sup>; Dong Li<sup>1</sup>; Yuan Liu<sup>1</sup>; Zhonglin Fu<sup>1</sup>; Xiaoling Yan<sup>2</sup>; Guorui Liu<sup>2</sup>; Linghui Li<sup>2</sup>; Andrea Throop<sup>4</sup>; Joshua LaBaer<sup>4</sup>; Xiao-Jun Li<sup>2</sup>; Jun Qin<sup>1</sup>; Bei Zhen<sup>1</sup>; Xiaobo Yu<sup>1</sup>; <sup>1</sup>Beijing Proteome Research Center, PHOENIX Center, Beijing, China; <sup>2</sup>Jinling Hospital, Nanjing University, Nanjing, China; <sup>3</sup>Department of Biostatistics, Columbia University, New York,

NY; <sup>4</sup>Biodesign Institute, Arizona State University, Tempe, AZ

- Poster 007 **Early Candidate Urine Biomarkers for Detecting Alzheimer's Disease before Amyloid-β Plaque Deposition in an APP (swe)/PSEN1dE9 Transgenic Mouse Model;** Fanshuang Zhang<sup>1</sup>; Jing Wei<sup>2</sup>; Xundou Li<sup>1</sup>; Chao Ma<sup>1</sup>; Youhe Gao<sup>2</sup>; <sup>1</sup>Basic Medicine Peking Union Medical College, Beijing, China; <sup>2</sup>Beijing Normal University, Beijing, China
- Poster 008 **Characterization of Human Multipotent Stromal Cells Secretome in Response to *in vitro* Passaging;** Ramavati Pal; Food and Drug Administration, Silver Spring, MD
- Poster 009 **A Spectral-Library Based Quantitative Study of Protein Signatures to Predict Response of Pancreatic Cancer Patients Receiving Chemotherapy;** Hong Peng<sup>1</sup>; Ru Chen<sup>2</sup>; Teresa Brentnall<sup>2</sup>; Vincent Picozzi<sup>3</sup>; Sheng Pan<sup>1</sup>; <sup>1</sup>The University of Texas Health Science Center, Houston, TX; <sup>2</sup>University of Washington, Seattle, WA; <sup>3</sup>Virginia Mason Medical Center, Seattle, WA
- Poster 010 **MIF Induce Th17-Related Cytokines Secretion in PBMC from Rheumatoid Arthritis Patients: Analysis through the Heat Map Method;** Luis Alexis Hernández-Palma<sup>1</sup>; Samuel García-Arellano<sup>1</sup>; Richard Bucala<sup>2</sup>; Mara Anaís Llamas-Covarrubias<sup>1</sup>; Ulises de la Cruz-Mosso<sup>1</sup>; Sergio Cerpa-Cruz<sup>3</sup>; José Francisco Muñoz-Valle<sup>1</sup>; <sup>1</sup>Universidad de Guadalajara, Guadalajara, Jalisco, México; <sup>2</sup>Yale University School of Medicine, New Haven, Connecticut, USA; <sup>3</sup>Hospital Civil de Guadalajara Fray Antonio Alcalde, Guadalajara, Jalisco, México
- Poster 011 **Preliminary Steps towards the Generation of a Rat Plasma Spectral Library;** Janet Kelsall<sup>1</sup>; Rachael Eineman<sup>1,2</sup>; Dave Lee<sup>1</sup>; Laura Cove-Smith<sup>3</sup>; Alison Backen<sup>3</sup>; John Radford<sup>4</sup>; Howard Mellor<sup>5</sup>; Kevin Hickling<sup>5</sup>; Marie South<sup>5</sup>; Jason Kirk<sup>5</sup>; Ivona Baricevic-Jones<sup>1</sup>; Julie Brazzatti<sup>1</sup>; Anthony Whetton<sup>1</sup>; Kim Linton<sup>4</sup>; <sup>1</sup>Stoller Biomarker Discovery Centre, Manchester, United Kingdom; <sup>2</sup>Manchester Molecular Pathology Innovation Centre, Manchester, United Kingdom; <sup>3</sup>Medical Oncology, Christie NHS Foundation Trust, Manchester, United Kingdom; <sup>4</sup>Manchester Cancer Research Centre, Wilmslow Road, Manchester, United Kingdom, M20 4QL; <sup>5</sup>AstraZeneca, Alderley Park, Macclesfield, SK10 4TG, United Kingdom
- Poster 012 **Differences of Saliva Composition in Relation to Tooth Decay and Gender;** Lucie Kulhová<sup>1</sup>; Adam Eckhardt<sup>2</sup>; Ivan Mikšík<sup>2</sup>; <sup>1</sup>Faculty of Science, Charles University, Prague, Czech Republic; <sup>2</sup>Institute of Physiology, Prague, Czech Republic
- Poster 013 **Developing Serum Multi-marker Panels for Diagnosing Hepatocellular Carcinoma Using Multiple Reaction Monitoring-Mass**



## POSTERS

All posters displayed Monday - Wednesday. Odd-numbered posters present Monday.  
Even-numbered posters present Tuesday. All posters present Wednesday morning.

- Spectrometry; Injoon Yeo<sup>1</sup>; Hyunsoo Kim<sup>2</sup>; Areum Sohn<sup>2</sup>; Gi-Ae Kim<sup>3</sup>; Young-Suk Lim<sup>4</sup>; Youngsoo Kim<sup>1</sup>; <sup>1</sup>Departments of Biomedical Engineering, Seoul National University College of Medicine, Seoul, Korea; <sup>2</sup>Departments of Biomedical Sciences, Seoul National University College of Medicine, Seoul, Korea; <sup>3</sup>Health Screening and Promotion Center, Asan Medical Center, Seoul, Korea; <sup>4</sup>Department of Gastroenterology, University of Ulsan College of Medicine, Seoul, Korea**
- Poster 014 **Discovery of Serum Biomarkers for Pancreatic Cancer by Lectin Affinity Capture Coupled with iTRAQ-Based Quantitative Glycoproteomics Approach; Chia-Chun Wu<sup>1</sup>; Yu-Ting Lu<sup>1</sup>; Yun-Hsin Chan<sup>2</sup>; Ta-Sen Yeh<sup>2</sup>; Jau-Song Yu<sup>1</sup>; <sup>1</sup>Chang Gung University, Taoyuan, Taiwan; <sup>2</sup>Chang Gung Memorial Hospital, Taoyuan, Taiwan**
- Poster 015 **Discovery of Prognostic Biomarkers for Hepatocellular Carcinoma by Mass Spectrometry-Based Phosphoproteomics Approach; Ye-Hsuan Sun<sup>2</sup>; Yu-Tsun Lin<sup>1</sup>; Kun-Yi Chien<sup>2,4</sup>; Chau-Ting Yeh<sup>2,3</sup>; Jau-Song Yu<sup>1,4</sup>; <sup>1</sup>Department of Biochemistry & Molecular Biology, Chang Gung University, Taoyuan, Taiwan; <sup>2</sup>Graduate Institute Of Biomedical Sciences, Chang Gung University, Taoyuan, Taiwan; <sup>3</sup>Liver Research Center, Chang Gung Memorial Hospital, Taoyuan, Taiwan; <sup>4</sup>Molecular Medicine Research Center, Chang Gung University, Taoyuan, Taiwan**
- Poster 016 **Identification of Potential Serum Protein Biomarkers for Recurrence in Gastric Cancer Patients Using a Quantitative Multiple Reaction Monitoring Approach; Byoung-Kyu Cho; Min Jueng Kang; Eugene C. Yi; Seoul National University, Seoul, South Korea**
- Poster 017 **PRM and MRM Methods to Identify Prognostic Biomarkers of Tocilizumab; Jinwoo Jung<sup>1</sup>; Byoung Kyu Cho<sup>1</sup>; Yeong Wook Song<sup>1,2</sup>; Eugene C. Yi<sup>1</sup>; <sup>1</sup>Seoul National University, Seoul, South Korea; <sup>2</sup>Division of Rheumatology, Seoul National University, Seoul, South Korea**
- Poster 018 **Quantitative Proteomic Analysis of Pancreatic Cyst Fluid Proteins Associated with Malignancy in Intraductal Papillary Mucinous Neoplasms; Misol Do<sup>1</sup>; Dohyun Han<sup>4</sup>; Joseph Injae Wang<sup>2</sup>; Hyunsoo Kim<sup>2</sup>; Wooil Kwon<sup>3</sup>; Youngmin Han<sup>3</sup>; Jin-Young Jang<sup>3</sup>; Youngsoo Kim<sup>1,2</sup>; <sup>1</sup>Department of Biomedical Sciences, Seoul National University College of Medicine, 103 Daehak-ro, Seoul, South Korea; <sup>2</sup>Department of Biomedical Engineering, Seoul National University College of Medicine, 103 Daehak-ro, Seoul, South Korea; <sup>3</sup>Department of Surgery, Seoul National University College of Medicine, 103 Daehak-ro, Seoul, South Korea; <sup>4</sup>Proteomics Core Facility, Seoul National University Hospital, 101 Daehak-ro, South Korea**
- Poster 019 **Variability Assessment of 90 Salivary Proteins in Intraday and Interday Samples from Healthy Donors by Multiple Reaction Monitoring-Mass Spectrometry; Lichieh Julie Chu<sup>1</sup>; Yung-Chin Hsiao<sup>1</sup>; Wei-Fan Chiang<sup>2</sup>; Yao-Ning Chuang<sup>1</sup>; Yu-Sun Chang<sup>1</sup>; Jau-Song Yu<sup>1</sup>; <sup>1</sup>Chang Gung University, Taoyuan, Taiwan; <sup>2</sup>Chi-Mei Medical Center, Liouying, Taiwan**
- Poster 020 **Development of an Automated Immuno-MALDI Mass Spectrometry Assay for Detection of Interstitial Collagenase in Dried Saliva Spot Sample.; Yung-Chin Hsiao<sup>1</sup>; Kun-Yi Chien<sup>1</sup>; Lang-Ming Chi<sup>2</sup>; Shih-Yu Lin<sup>1</sup>; Wei-Fang Chiang<sup>3</sup>; Yu-Sun Chang<sup>1</sup>; Jau-Song Yu<sup>1</sup>; <sup>1</sup>Chang Gung University, Tao-Yuan, Taiwan; <sup>2</sup>Chang Gung Memorial Hospital, Tao-Yuan, Taiwan; <sup>3</sup>Chi-Mei Medical Center, Liouying, Taiwan**
- Poster 021 **Comprehensive Proteomic Profiling of Serum Exosomes Identifies Novel Biomarkers for Early Detection of Gastric Cancer; Naomi Ohnishi; Japanese Foundation for Cancer Research, Tokyo, Japan**
- Poster 022 **Proteomic Study on Advanced Glycation End-Products Treatment in Kidney of Mice; Eun Hee Han; Young-Ho Chung; Korea Basic Science Institute (KBSI), Cheongju-Si, South Korea**
- Poster 023 **Proline-Rich Protein 4 (PRR4) as a Potential Tear Biomarker for Personalized Diagnosis of Dry Eye Disease and Glaucoma; Natarajan Perumal; Caroline Manicam; Alexandra Tschäbunin; Aline Ratcliffe; Laura Gronbach; Maya Scieranski; Norbert Pfeiffer; Franz Grus; University Medical Centre Mainz, Mainz, Germany**
- Poster 024 **A Fast, Simple and Robust Sample Preparation Workflow Enables High-Throughput Plasma Protein Profiling; Mo Hu; Yue Zhou; Jing Li; Thermo Fisher Scientific, Shanghai, China**
- Poster 025 **Distinguishing Pancreatic Cancer from Benign Diseases and Healthy Individuals by Mass Spectrometry-Based Metabolomic Pipeline; Xiaohui Liu; Yueting Xiong; Pengyuan Yang; Fudan University, Shanghai, China**
- Poster 026 **Plasma Proteomics in Children Diagnosed with Acute Lymphoblastic Leukemia: A Pilot Study; Sandra Calderon-Rodriguez<sup>1</sup>; Carolina Sanabria-Salas<sup>1,2</sup>; Adriana Umana-Perez<sup>1</sup>; <sup>1</sup>National University Colombia, Bogota, Colombia; <sup>2</sup>Instituto Nacional de Cancerologia, Bogota, Cundinamarca**
- Poster 027 **Quantitative Proteomic Signature of First-Episode Psychosis Patients' PBMCs – Preliminary Results; Cátia Santa<sup>1,2</sup>; Manuel Coroa<sup>3,4</sup>; Sofia Morais<sup>3,4</sup>; Sandra I. Anjo<sup>1,3</sup>; Inês Baldeiras<sup>1,3</sup>; Nuno Madeira<sup>3,4</sup>; António Macedo<sup>3,4</sup>; Bruno Manadas<sup>1</sup>; <sup>1</sup>Center for Neuroscience**

## POSTERS

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and Cell Biology, UC, Coimbra, Portugal;  
<sup>2</sup>Institute for Interdisciplinary Research, UC,  
Coimbra, Portugal; <sup>3</sup>Faculty of Medicine, UC,  
Coimbra, Portugal; <sup>4</sup>Psychiatry Department,  
CHUC, Coimbra, Portugal

- Poster 028 **Searching for New Blood Biomarkers of Wilson's Disease Using Translational Proteomics.**; Maud Lacombe; CEA/DRF/BIG/BGE/EDyP, Grenoble, France
- Poster 029 **Identification of Diagnostic Biomarkers for Lung Cancer by Quantitative Proteomic Analysis;** Yan Ping Zhu; Binzhou Medical University, Yantai, China
- Poster 030 **Identification of Differential Expression Proteins in Esophageal Cancer by Label-Free Quantitative Proteomics Analysis;** Xiaoying Qi; Binzhou Medical University, Yantai, China
- Poster 031 **Investigating the Proteases/Peptidases Implicated in the Urinary Normal Peptidome Generation; towards New Trends in Biomarker Discovery.**; Amr Elguoshy<sup>1,2</sup>; Yoshitoshi Hirao<sup>1</sup>; Keiko Yamamoto<sup>1</sup>; Bo Xu<sup>1</sup>; Toshiaki mitsu<sup>2</sup>; Tadashi Yamamoto<sup>1</sup>; <sup>1</sup>Biofluid Biomarker Center, Niigata University, Niigata, Japan; <sup>2</sup>Graduate School of Science and Technology, Niigata university, Japan
- Poster 032 **Strategy to Establish New Clinical Biomarkers: From Proteomics Selection of Biomarker Candidates to Validation for Clinical Use;** Keiko Yamamoto; Yoshitoshi Hirao; Amr Elguoshy; Xu Bo; Tadashi Yamamoto; Biofluid Biomarker Center, Niigata University, Niigata, Japan
- Poster 033 **Early Candidate Biomarkers in Urine of Walker-256 Lung Metastasis Rat Model;** Jing Wei<sup>1</sup>; Na Ni<sup>2</sup>; Linpei Zhang<sup>1</sup>; Youhe Gao<sup>1</sup>; <sup>1</sup>Beijing Normal University, Beijing, China; <sup>2</sup>Chongqing Medical University, Chongqing, China
- Poster 034 **A Diagnostic Panel of Urine Protein Biomarkers Predicts Lung Cancer from Healthy Controls and Other Tumors;** Pei Zhen<sup>1,2</sup>; Chunchao Zhang<sup>2</sup>; Changqing Sun<sup>2</sup>; Yi Wang<sup>1</sup>; Guangshun Wang<sup>2</sup>; Jun Qin<sup>1,3</sup>; <sup>1</sup>The PHOENIX Center, Beijing, China; <sup>2</sup>Joint Center for Translational Medicine, Tianjin, China; <sup>3</sup>Baylor College of Medicine, Houston, USA
- Poster 035 **Analysis of Toxicologically Relevant Proteins in Pesticide-Treated HepaRG Cells by MS-Based Immunoassays;** Felix Schmidt<sup>1</sup>; Andreas Steinhilber<sup>1</sup>; Helen Hammer<sup>2</sup>; Almut Mentz<sup>3</sup>; Jörn Kalinowski<sup>3</sup>; Dajana Lichtenstein<sup>4</sup>; Albert Braeuning<sup>4</sup>; Philip Marx-Stoelting<sup>4</sup>; Alfonso Lampen<sup>4</sup>; Thomas Joos<sup>1</sup>; Oliver Pötz<sup>1,2</sup>; <sup>1</sup>NMI Reutlingen, Reutlingen, Germany; <sup>2</sup>SIGNATOPE GmbH, Reutlingen, Germany; <sup>3</sup>Bielefeld University, Bielefeld, Germany; <sup>4</sup>German Federal Institute for Risk Assessment, Berlin, Germany

## CANCER Posters 036 - 071

- Poster 036 **Identification of Tumor Specific Peptides Using HLA Peptidome;** Sunny Heo; Asan Medical Center, Seoul, South Korea
- Poster 037 **Proteomic Profiling of Proteolytic Processing Events in Plasma Samples from Melanoma Patients;** Francine Braga<sup>1</sup>; Alexandre Tashima<sup>2</sup>; Eduardo Kitano<sup>3</sup>; Ana Maria Chudzinski-Tavassi<sup>3</sup>; Roger Chammas<sup>4</sup>; André Zelanis<sup>1</sup>; <sup>1</sup>Laboratório de Proteômica Funcional, UNIFESP, São José Dos Campos, SP, Brazil; <sup>2</sup>Department of Biochemistry, UNIFESP, São Paulo, SP, Brazil; <sup>3</sup>Centre of Excellence in New Target Discovery, São Paulo, SP, Brazil; <sup>4</sup>Instituto do Câncer do Estado de São Paulo, São Paulo, SP, Brazil
- Poster 038 **Quantitative Shotgun Proteomics Unveils Candidate Novel Cervical Cancer-Specific Proteins;** Alberto Ramírez Torres<sup>1</sup>; Jeovanis Gil<sup>1</sup>; Sandra Contreras<sup>1</sup>; Graciela Ramírez<sup>2,3</sup>; Heriberto Valencia<sup>2,3</sup>; Alejandro García Carranca<sup>2,3</sup>; Sergio Encarnacion-Guevara<sup>1</sup>; <sup>1</sup>CCG-Universidad Nacional Autónoma de México, Cuernavaca, Morelos, México; <sup>2</sup>IIB-Universidad Nacional Autónoma de México, Ciudad de México, México; <sup>3</sup>Instituto Nacional de Cancerología, Ciudad de México, México
- Poster 039 **The Proteome of OSCC-Derived Extracellular Vesicles Reflects Tumor Aggressiveness and Clinical Staging;** Ana Kariana de Oliveira<sup>1,2</sup>; Ariane Busso-Lopes<sup>1</sup>; Jamile Sá<sup>1,2</sup>; César Rivera<sup>1,3</sup>; Alan Santos-Silva<sup>2</sup>; Márcio Lopes<sup>2</sup>; Adriana Paes Leme<sup>1</sup>; <sup>1</sup>Mass Spectrometry Laboratory, LNBio, CNPEM, Campinas, Brazil; <sup>2</sup>Department of Oral Diagnosis, UNICAMP, Campinas, Brazil; <sup>3</sup>Department of Basic Biomedical Sciences, UTALCA, Talca, Chile
- Poster 040 **Unravelling the Impact of ASPP-PP1 Interactions Using Phosphoproteomics Studies in KI Mouse and Cancer Cell Lines;** Kundan Sharma; Elizabeth Slee; Hokfung Chan; Xin Lu; Ludwig Cancer Research, University of Oxford, Oxford, United Kingdom
- Poster 041 **Bioenergetic Reprogramming Profoundly Changes the Mitochondrial Proteome Resulting in Marked Changes to Morphology and Susceptibility to Induced Cell Death;** Rebekah Jukes-Jones<sup>1</sup>; Gareth J Miles<sup>3</sup>; Kelvin Cain<sup>1</sup>; Claudia Langlais<sup>2</sup>; <sup>1</sup>MRC Toxicology Unit, Leicester, United Kingdom; <sup>2</sup>Immunocore, Abingdon, Ox; <sup>3</sup>Leicester University, Leicester, LE
- Poster 042 **Characterization of Malignant Pleural Mesothelioma by Comprehensive Proteomics Study;** Jongmin Choi; Hyun-Sung Lee; Bryan M. Burt; Sung Jung; Baylor College of Medicine, Houston, TX

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- Poster 043 **Investigating Novel Mutant p53 Interacting Proteins in Cancer Cells;** Maríel Mendoza; Katherine Alexander; Enrique Lin Shiao; Charly Ryan Good; Benjamin A. Garcia; Shelley L. Berger; *University of Pennsylvania, Philadelphia, PA*
- Poster 044 **Mass Spectrometry-Based Proteomic Profiling of Pediatric Neuroblastic Tumors;** Rebecca C Poulos<sup>1</sup>; Qing Zhong<sup>1</sup>; Brett Tully<sup>1</sup>; Sumanth Nagabushan<sup>2, 3</sup>; Priya Duggal<sup>1</sup>; Sadia Mahboob<sup>1</sup>; Belinda Serafin<sup>1</sup>; Peter G Hains<sup>1</sup>; Phillip J Robinson<sup>1</sup>; Roger Reddel<sup>1</sup>; Rosemary Balleine<sup>1</sup>; <sup>1</sup>*Children's Medical Research Institute, University of Sydney, NSW, Australia*; <sup>2</sup>*Sydney Children's Hospital Network, Westmead, NSW, Australia*; <sup>3</sup>*University of Sydney, NSW, Australia*
- Poster 045 **Generation of the CanPath Prototype - a Platform for Predictive Cancer Pathway Modelling;** Magdalena Bober-Andres<sup>1</sup>; Daniel Heinzmann<sup>1</sup>; Monika Banko-Bielecka<sup>1</sup>; Oliver Rinner<sup>1</sup>; Christoph Wierling<sup>2</sup>; Thomas Kessler<sup>2</sup>; Artur Muradyan<sup>2</sup>; Louisa Krützfeldt<sup>2</sup>; Moritz Schütte<sup>2</sup>; Felix Dreher<sup>2</sup>; Bodo Lange<sup>2</sup>; <sup>1</sup>*Biognosys AG, Schlieren, Switzerland*; <sup>2</sup>*Alacris Theranostics GmbH, Berlin, Germany*
- Poster 046 **Pitchfork Approach for Membrane Proteome Profiling of Human Pheochromocytoma and Paraganglioma;** Ondrej Vit<sup>1</sup>; Karel Pacak<sup>2</sup>; Jiri Petrak<sup>1</sup>; <sup>1</sup>*First Faculty of Medicine, Charles University, Vestec, Czech Republic*; <sup>2</sup>*NICHD-NIH, Bethesda, MD*
- Poster 047 **Proteomic Analysis of Nasopharyngeal Carcinoma Cells with Activated NLRP3 Inflammatory Specks by iTRAQ Technology;** I-Che Chung; Chih-Ching Wu; Yu-Sun Chang; *Chang Gung University, Kweishan, Taiwan*
- Poster 048 **Quantitative Phosphoproteomics Indicates Altered Cell Migration in Prolonged Cabozantinib-Treated Renal Cell Carcinoma Cell Lines;** Yu-Heng Hsieh<sup>1</sup>; Shao-Kuan Chen<sup>1, 2</sup>; Yen-Chieh Wang<sup>1</sup>; Teh-Sheng Hsieh<sup>3</sup>; Chih-Jung Huang<sup>1, 3</sup>; Wei-Chi Ku<sup>1</sup>; <sup>1</sup>*Fu Jen Catholic University, New Taipei, Taiwan*; <sup>2</sup>*Sijhih Cathay General Hospital, New Taipei, Taiwan*; <sup>3</sup>*Cathay General Hospital, Taipei, Taiwan*
- Poster 049 **Biomarker Discovery in Triple Negative Breast Cancer Using iTRAQ-Based Quantitative Proteomic Analysis;** Songping Lin; Yuxiang Lin; Fangmeng Fu; Chuan Wang; *Union Hospital of Fujian, Fuzhou, China*
- Poster 050 **Comprehensive Proteomic Mapping of Chronic Myelogenous Leukemia;** Sameh Magdeldin; Aya Osama; *CCHE 57357, Cairo, Egypt*
- Poster 051 **Systems-Wide Profiling of Proteolytic Events in Murine Melanoma Secretome Using Terminal Amine Isotopic Labeling of Substrates;** Tarcísio Liberato<sup>1, 2</sup>; Isabella Fukushima<sup>1, 2</sup>; Dayelle Pessotti<sup>1</sup>; Débora Andrade-Silva<sup>3</sup>; Eduardo S. Kitano<sup>4</sup>; Solange M.T. Serrano<sup>3</sup>; André Zelanis<sup>1, 2</sup>; <sup>1</sup>*UNIFESP - Brazil, São José Dos Campos, Brazil*; <sup>2</sup>*UNIFESP - Functional Proteomics Laboratory, São José dos Campos, Brazil, São Paulo*; <sup>3</sup>*Lab. Especial de Toxinologia Aplicada, I. Butantan, São Paulo, SP, Brazil*; <sup>4</sup>*Centre of Excellence in New Target Discovery, I. Butantan, Sao Paulo, Brazil*
- Poster 052 **Proteomic Profiles of Glioma Subtypes and Glioblastoma Stem Cells Reveal Conserved Profiles According to IDH Mutation Status;** Ugljesa Djuric<sup>1</sup>; Jennifer Kao<sup>1</sup>; Ithor Batruch<sup>2</sup>; Stefan Jevtic<sup>1</sup>; Ken Aldape<sup>1</sup>; Phedias Diamandis<sup>1</sup>; <sup>1</sup>*University Health Network, Toronto, Canada*; <sup>2</sup>*Mount Sinai Hospital, Toronto, Canada*
- Poster 053 **Proteomics Confirms Lower Cancer Cell-Surface uPAR Superimposed on KRAS Mutation Carrying Cells Can Negate Many of the Hallmarks of Cancer;** Seong Beom Ahn<sup>1</sup>; Abidali Mohamedali<sup>2</sup>; Dana Pascovici<sup>3</sup>; Subash Adhikari<sup>1</sup>; Mark Baker<sup>1</sup>; <sup>1</sup>*Biomedical Sciences, Macquarie University, Sydney, Australia*; <sup>2</sup>*Molecular Sciences, Macquarie University, Sydney, Australia*; <sup>3</sup>*APAF, Macquarie University, Sydney, Australia*
- Poster 054 **A Quantitative Analysis of Colon Adenocarcinoma Using MS-Based Proteomics;** Sanjeeva Srivastava; *IIT Bombay, Mumbai, India*
- Poster 055 **Proteomic and Integrated Omic Analyses Reveal Drivers in a Subset of Aggressive Primary Lung Cancers;** Shideh Mirhadi; Michael Moran; *University of Toronto, Toronto, Canada*
- Poster 056 **Integrated Proteogenomic Data Analysis Pipeline and Its Applications to the Analysis of CPTAC Ovarian Cancer Data;** Yingwei Hu; Jianbo Pan; David J. Clark; Punit Shah; Minghui Ao; Michale Schnaubelt; Lijun Chen; Jiang Qian; Zhen Zhang; Daniel W. Chan; Hui Zhang; *Johns Hopkins University, Baltimore, MD*
- Poster 057 **Integrative Mass Spectrometry and RNA-Sequencing Identifies Candidate Immunotherapeutic Targets in Neuroblastoma;** Amber K. Weiner<sup>1, 2</sup>; Alexander B. Radaoui<sup>2</sup>; Nathan M. Kendersky<sup>1, 2</sup>; Simone Sidoli<sup>1</sup>; Karina L. Conkrite<sup>2</sup>; Jo Lynne Harenza<sup>2</sup>; Zalman Vaksman<sup>2</sup>; Komal S. Rathi<sup>2</sup>; Pichai Ramen<sup>2</sup>; Daniel Martinez<sup>2</sup>; Tricia Bhatti<sup>2</sup>; Matthew Tsang<sup>2</sup>; Bruce Pawel<sup>2</sup>; Benjamin A. Garcia<sup>1</sup>; John M. Maris<sup>2</sup>; Sharon J. Diskin<sup>2</sup>; <sup>1</sup>*University of Pennsylvania, Philadelphia, PA*; <sup>2</sup>*Children's Hospital of Philadelphia, Philadelphia, PA*
- Poster 058 **Bothrops Jararaca Snake Venom Increases Level of Several Cancer-Related Proteins on Different Tumor Cell Lines;** Carolina Yukiko Kisaki; Ismael Feitosa Lima; Hugo Aguirre

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- Armelin; Leo Kei Iwai; *Butantan Institute, São Paulo, Brazil*
- Poster 059 **Neoadjuvant Chemotherapy-Treated Ovarian Cancer Patients Have Unique Tumor Proteome Alterations Associated with Volume of Residual Disease;** Nicholas Bateman<sup>1</sup>; Emily Penick<sup>1</sup>; Kelly Conrads<sup>1</sup>; Ming Zhou<sup>3</sup>; Guisong Wang<sup>1</sup>; Niyati Parikh<sup>1</sup>; Kathleen Darcy<sup>1</sup>; Yovanni Casablanca<sup>1</sup>; Paulette Mhaweche-Fauceglia<sup>2</sup>; Thomas Conrads<sup>3</sup>; G. Larry Maxwell<sup>4</sup>; <sup>1</sup>*Gynecologic Cancer Center of Excellence, Annandale, VA;* <sup>2</sup>*Keck School of Medicine, University of Southern Cal, Los Angeles, CA;* <sup>3</sup>*Inova Schar Cancer Institute, Falls Church, VA;* <sup>4</sup>*Obstetrics and Gynecology, Inova Fairfax Hospital, Falls Church, VA*
- Poster 060 **Inhibition of Cell Proliferation and Altered Proteome Analysis after GnRH Agonist Treatment in Glioblastoma Cell Line;** Priyanka Harishchandra Tripathi<sup>1, 2</sup>; Jyoti Arora<sup>1</sup>; Ravindra Varma Polisetty<sup>3</sup>; Ravindra Kumar Saran<sup>4</sup>; Fouzia Siraj<sup>1</sup>; Neetu Mishra<sup>2</sup>; Ravi Sirdeshmukh<sup>5</sup>; Poonam Gautam<sup>1</sup>; <sup>1</sup>*ICMR-National Institute of Pathology, New Delhi, India;* <sup>2</sup>*Symbiosis School of Biological Sciences, Pune, India;* <sup>3</sup>*SriVenkateshwar College, Delhi University, New Delhi, India;* <sup>4</sup>*Govind Ballabh Pant Hospital, New Delhi, India;* <sup>5</sup>*Institute of Bioinformatics, Bangalore, India*
- Poster 061 **HER2 Proteomic Signature in Gastric Cancer;** Jeong-Won Kang; Hark Kim; *National Cancer Center, Goyang, South Korea*
- Poster 062 **Post-Translational Crosstalk Networks Identify Strategies to Overcome EMT-Mediated Resistance to EGFR Inhibitors;** Guolin Zhang<sup>1</sup>; Karen Ross<sup>2</sup>; Bin Fang<sup>1</sup>; Jun-Min Zhou<sup>1</sup>; Paul A Stewart<sup>1</sup>; Emma Adhikari<sup>1</sup>; Eric A Welsh<sup>1</sup>; Xuefeng Wang<sup>1</sup>; John M Koomen<sup>1</sup>; Cathy H Wu<sup>2, 3</sup>; Eric B Haura<sup>1</sup>; <sup>1</sup>*H.Lee Moffitt Cancer Center & Research Institute, Tampa, FL;* <sup>2</sup>*Georgetown University Medical Center, Washington, DC;* <sup>3</sup>*University of Delaware, Newark, DE*
- Poster 063 **Breast Cancer Quantitative Proteome and Proteogenomic Landscape;** Henrik J Johansson<sup>1</sup>; Fabio Socciarelli<sup>1</sup>; Nathaniel Vacanti<sup>1</sup>; Mads H. Haugen<sup>2</sup>; Yafeng Zhu<sup>1</sup>; Ioannis Siavelis<sup>1</sup>; Alejandro Fernandez<sup>1</sup>; Miriam R. Aure<sup>2</sup>; Bengt Sennblad<sup>3</sup>; Mattias Vesterlund<sup>1</sup>; Rui M. Branca<sup>1</sup>; Lukas M. Orre<sup>1</sup>; Mikael Huss<sup>3</sup>; Erik Fredlund<sup>1</sup>; Elsa Beraki<sup>2</sup>; Øystein Garred<sup>2</sup>; Jorrit Boekel<sup>1</sup>; Torill Sauer<sup>4</sup>; Wei Zhao<sup>5</sup>; Silje Nord<sup>2</sup>; Elen K. Högländer<sup>2</sup>; Daniel C. Jans<sup>6</sup>; Hjalmar Brismar<sup>6</sup>; Tonje H. Haukaas<sup>8</sup>; Ellen Schlichting<sup>2</sup>; Bjørn Naume<sup>2</sup>; OSBREAC OSBREAC<sup>7</sup>; Elin Borgen<sup>2</sup>; Vessela N. Kristensen<sup>2</sup>; Hege G. Russnes<sup>2</sup>; Ole Christian Lingjærde<sup>2</sup>; Gordon B. Mills<sup>5</sup>; Kristine K. Sahlberg<sup>2</sup>; Anne-Lise Børresen-Dale<sup>2</sup>; Janne Lehtiö<sup>1</sup>; <sup>1</sup>*Karolinska Institutet, Stockholm, Sweden;* <sup>2</sup>*Oslo University Hospital, Oslo,*
- Norway;* <sup>3</sup>*Stockholm University, Solna, Sweden;* <sup>4</sup>*Akershus University Hospital, Lørenskog, Norway;* <sup>5</sup>*The University of Texas MD Anderson Cancer Center, Houston, USA;* <sup>6</sup>*KTH Royal Institute of Technology, Stockholm, Sweden;* <sup>7</sup>*www.osbreac.no, Oslo, Norway;* <sup>8</sup>*The Norwegian University of Science and Technology, Trondheim, Norway*
- Poster 064 **Polycomb Loss Enhances Oncogenesis but Leads to Therapeutic Vulnerabilities in Malignant Peripheral Nerve Sheath Tumors;** John Wojcik<sup>1</sup>; Dylan Marchione<sup>1</sup>; Simone Sidoli<sup>1</sup>; Benjamin Garcia<sup>1, 2</sup>; <sup>1</sup>*University of Pennsylvania, Philadelphia, PA;* <sup>2</sup>*University of Pennsylvania School of Medicine, N/A, N/A*
- Poster 065 **Epigenetic Dysregulation Drives Altered Chromatin-Reader Interactions in Diffuse Intrinsic Pontine Glioma;** Dylan Marchione; John Wojcik; Benjamin A. Garcia; *University of Pennsylvania, Philadelphia, PA*
- Poster 066 **Rapid Plasma Biomarker Validation by Reverse Phase Protein Array;** Tesshi Yamada; *National Cancer Center Research Institute, Chuo-Ku, Japan*
- Poster 067 **Analysis of the Epidermal Growth Factor-Induced Phosphorylation of Actinin-4 involved in Cancer Metastasis;** Nami Miura<sup>1</sup>; Kaoru Onidani<sup>1</sup>; Kazufumi Honda<sup>1, 2</sup>; <sup>1</sup>*National Cancer Center, Chuo-Ku, Tokyo, Japan;* <sup>2</sup>*Japan Agency for Medical Research and Development, Tokyo, Japan*
- Poster 068 **BRK Mediated Phosphorylation Regulates SMAD4 Control of the TGF- $\beta$ /SMAD4 Signaling Pathway to Control SLUG, SNAIL and Metastatic Potential.;** Md Sayem Miah<sup>1, 2</sup>; Charles Banks<sup>1</sup>; Yetunde Ogunbolude<sup>2</sup>; Edward Bagu<sup>2</sup>; Anita Saraf<sup>1</sup>; Gaye Hattem<sup>2</sup>; Cassandra Eubanks<sup>1</sup>; Mihaela Sardi<sup>1</sup>; Laurence Florens<sup>1</sup>; Kiven Lukong<sup>2</sup>; Michael Washburn<sup>1, 3</sup>; <sup>1</sup>*Stowers Institute, Kansas City, MISSOURI;* <sup>2</sup>*University of Saskatchewan, Saskatoon, Canada;* <sup>3</sup>*University of Kansas Medical Centre, Kansas City, KS*
- Poster 069 **Screening of Novel Molecular Therapeutic Targets for Tongue Cancer Using a Kinase Antibody Library;** Kaoru Onidani<sup>1, 2</sup>; Yukio Watabe<sup>2</sup>; Nami Miura<sup>1</sup>; Takahiko Shibahara<sup>2</sup>; Kazufumi Honda<sup>1, 3</sup>; <sup>1</sup>*National Cancer Center Research Institute, Tokyo, Japan;* <sup>2</sup>*Tokyo Dental College, Tokyo, Japan;* <sup>3</sup>*AMED CREST, Tokyo, Japan*
- Poster 070 **Identification of Aggressive Prostate Cancers: In-depth Proteomics of Tissues and post-DRE urines;** Thomas Kislinger<sup>1</sup>; Andrew Macklin<sup>1</sup>; Amanda Khoo<sup>2</sup>; Katharina Fritsch<sup>2</sup>; Yunee Kim<sup>2</sup>; Ankit Sinha<sup>2</sup>; Vincent Huang<sup>3</sup>; Julie Livingstone<sup>3</sup>; Vladimir Ignatchenko<sup>1</sup>; Theodoros van der Kwast<sup>1</sup>; Rob Bristow<sup>1</sup>; Stanley Liu<sup>5</sup>; Julius Nyalwidhe<sup>4</sup>; Jouhyun Jouhyun<sup>3</sup>; John Semmes<sup>4</sup>; Paul Boutros<sup>3</sup>; <sup>1</sup>*Princess Margaret Cancer Centre, Toronto, Canada;* <sup>2</sup>*University of*

## POSTERS

All posters displayed Monday - Wednesday. Odd-numbered posters present Monday.  
Even-numbered posters present Tuesday. All posters present Wednesday morning.

Toronto, Toronto, Canada; <sup>3</sup>Ontario Institute for Cancer Research, Toronto, Canada; <sup>4</sup>Eastern Virginia Medical School, Norfolk, USA; <sup>5</sup>Sunnybrook Health Sciences Centre, Toronto, Canada

Germany; <sup>2</sup>Institute for Clinical Chemistry, UMC, Goettingen, Germany; <sup>3</sup>Bioanalytical Mass Spectrometry, MPI, Goettingen, Germany; <sup>4</sup>Neuromuscular and Cardiovascular Cell Biology, MDC, Berlin, Germany

Poster 071 **High Resolution Protein Mapping of ROS1-Rearranged NSCLC Cell Lines: Defining Mechanisms of Acquired Crizotinib Resistance;** Sarah Hayes<sup>1,2</sup>; Christoph Krisp<sup>3</sup>; Amanda Hudson<sup>1,2</sup>; Stephen Clarke<sup>4</sup>; Nick Pavlakis<sup>4</sup>; Mark Molloy<sup>3</sup>; Viive Howell<sup>1,2</sup>; <sup>1</sup>Kolling Institute of Medical Research, St Leonards, Australia; <sup>2</sup>Sydney Medical School, University of Sydney, Sydney, Australia; <sup>3</sup>Australian Proteome Analysis Facility, Sydney, Australia; <sup>4</sup>Department of Medical Oncology, RNSH, Sydney, Australia

Poster 076 **Profiling the Proteomic and Lipidomic Dysregulations of Mouse Aorta during Atherosclerotic Plaque Formation;** Juanjuan Xie<sup>1</sup>; Xiangdong Yang<sup>3</sup>; Huali Shen<sup>1</sup>; Pengyuan Yang<sup>1,2</sup>; <sup>1</sup>Institute of Biomedical Science, Fudan University, Shanghai, China; <sup>2</sup>Department of Chemistry, Fudan University, Shanghai, China; <sup>3</sup>Department of Cardiology, Zhongshan Hospital, Shanghai, China

Poster 077 **Global Human and Mouse Phosphoproteomic Profiling of Signaling Pathway Aberrations in Hypertrophic Cardiomyopathy;** Uros Kuzmanov<sup>1</sup>; Rachel Vanderlaan<sup>1</sup>; Hongbo Guo<sup>1</sup>; Sina Hadipour-Lakmehsari<sup>1</sup>; Parveen Sharma<sup>2</sup>; Phyllis Billia<sup>3</sup>; Andrew Emili<sup>1</sup>; Anthony Gramolini<sup>1</sup>; <sup>1</sup>University of Toronto, Toronto, Canada; <sup>2</sup>University of Liverpool, Liverpool, United Kingdom; <sup>3</sup>University Health Network, Toronto, Canada

### CARDIOVASCULAR Posters 072 - 077

Poster 072 **Differences in Plasma Fibrin Clot Composition in Patients with Thrombotic Antiphospholipid Syndrome Compared with Venous Thromboembolism;** Aneta Stachowicz<sup>1,2</sup>; Michał Ząbczyk<sup>1</sup>; Joanna Natorska<sup>1</sup>; Maciej Suski<sup>1</sup>; Rafał Olszanecki<sup>1</sup>; Ryszard Korbut<sup>1</sup>; Jacek Wiśniewski<sup>2</sup>; Anetta Undas<sup>1</sup>; <sup>1</sup>Jagiellonian University Medical College, Krakow, Poland; <sup>2</sup>Max Planck Institute of Biochemistry, Martinsried, Germany

Poster 073 **Integrated Dissection of the Cysteine Oxidative Modification Proteome During Cardiac Hypertrophy;** Jie Wang<sup>1,3</sup>; Howard Choi<sup>2,3</sup>; Neo Chung<sup>3</sup>; Quan Cao<sup>1,3</sup>; Dominic Ng<sup>1,3</sup>; Bilal Mirza<sup>1,3</sup>; Sarah Sruggs<sup>1,3</sup>; Ding Wang<sup>1,3</sup>; Anders Garlid<sup>1,3</sup>; Peipei Ping<sup>1,3</sup>; <sup>1</sup>Departments of Physiology and Medicine, UCLA, Los Angeles, CA; <sup>2</sup>Department of Bioinformatics, UCLA, Los Angeles, CA; <sup>3</sup>NIH BD2K Center of Excellence, UCLA, Los Angeles, CA

Poster 074 **Phosphopeptide Enrichment and Analysis of Human Ischemic Cardiomyopathic Tissues Reveal Infarct Versus Non-Infarct Unique Signaling Pathways;** Da Hye (Julia) Kim<sup>1,2</sup>; Uros Kuzmanov<sup>2,3</sup>; Sina Hadipour-Lakmehsari<sup>1,2</sup>; Andrew Emili<sup>3,4</sup>; Gavin Oudit<sup>5,6</sup>; Anthony Gramolini<sup>1,2</sup>; <sup>1</sup>Physiology, University of Toronto, Toronto, Canada; <sup>2</sup>Ted Rogers Centre for Heart Research, Toronto, Canada; <sup>3</sup>Donnelly Centre for Cellular and Biomolecular Res, Toronto, Canada; <sup>4</sup>Molecular Genetics, University of Toronto, Toronto, Canada; <sup>5</sup>Medicine, University of Alberta, Edmonton, Canada; <sup>6</sup>Mazankowski Alberta Heart Institute, Edmonton, Canada

Poster 075 **Phospholamban Targeted Proximity Labeling for Proteomic Mapping of sarcoendoplasmic Reticulum Subdomains;** Daniel Kownatzki-Danger<sup>1</sup>; Christof Lenz<sup>2,3</sup>; Henning Urlaub<sup>2,3</sup>; Michael Gotthardt<sup>4</sup>; Stephan E. Lehnart<sup>1</sup>; <sup>1</sup>Cardiology and Pneumology, UMC, Goettingen,

### CHEMICAL PROTEOMICS Posters 078 - 080

Poster 078 **Development of the Human Proteome Peptide Catalog – A comprehensive Repository of Reference Peptides for the Human Proteome;** Karsten Schnatbaum<sup>1</sup>; Daniel P. Zolg<sup>2</sup>; Mathias Wilhelm<sup>2</sup>; Tobias Knaute<sup>1</sup>; Johannes Zerweck<sup>1</sup>; Holger Wenschuh<sup>1</sup>; Bernhard Kuster<sup>2,3</sup>; Ulf Reimer<sup>1</sup>; <sup>1</sup>JPT Peptide Technologies GmbH, Berlin, Germany; <sup>2</sup>Chair of Proteomics and Bioanalytics, TU Munich, Freising, Germany; <sup>3</sup>Center for Integrated Protein Science Munich, Freising, Germany

Poster 079 **Analysis of Receptor Tyrosine Kinase Inhibitor Distribution in Tumor Xenograft Mouse Using MALDI Mass Spectrometry Imaging;** Tae Young Kim<sup>1</sup>; Seung Hyun Pan<sup>1</sup>; Yonghyo Kim<sup>1</sup>; Yutaka Sugihara<sup>2</sup>; Melinda Rezeli<sup>2</sup>; Marcell Szasz<sup>3</sup>; Gyorgy Marko-Varga<sup>2</sup>; Ho Jeong Kwon<sup>1</sup>; <sup>1</sup>Yonsei university, Seoul, South Korea; <sup>2</sup>Lund university, Lund, Sweden; <sup>3</sup>National Koranyi Institute, Budapest, Hungary

Poster 080 **Elucidating Off-Target Proteins of RTKi by Combinatory Method of Label-Free DARTS and LC-MS/MS;** Seung Hyun Pan<sup>1</sup>; Tae Young Kim<sup>1</sup>; EunSun Ji<sup>2</sup>; Jin Young Kim<sup>2</sup>; Jong Shin Yoo<sup>2</sup>; Ho Jeong Kwon<sup>1</sup>; <sup>1</sup>Yonsei University, Seoul, South Korea; <sup>2</sup>KBSI, Ochang, South Korea

### C-HPP Posters 081 - 090

Poster 081 **Identification of the Missing Protein Hyaluronan Synthase 1 in Mesenchymal Stem Cells Derived from Adipose Tissue or**

## POSTERS

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- Umbilical Cord**; Miguel Marcilla<sup>1</sup>; Luis Felipe Clemente<sup>2</sup>; Maria Luisa Hernaez<sup>2</sup>; Antoni Ramos-Fernandez<sup>3</sup>; Gertrudis Ligeró<sup>4</sup>; Concha Gil<sup>2</sup>; [Fernando Corrales](#)<sup>1</sup>; <sup>1</sup>*Centro Nacional de Biotecnología, CSIC, Madrid, Spain*; <sup>2</sup>*Faculty of Pharmacy, University Complutense, Madrid, Spain*; <sup>3</sup>*Proteobotics, Madrid, Spain*; <sup>4</sup>*Andalusian Public Health System Biobank, Granada, Spain*
- Poster 082 **Missing Protein Detection in Human Embryonic Tissues Using a Proteogenomics Approach**; José González-Gomariz<sup>1, 2</sup>; Guillermo Serrano<sup>1</sup>; Alba Garin-Muga<sup>3</sup>; [Fernando J. Corrales](#)<sup>4</sup>; Elizabeth Guruceaga<sup>1, 2</sup>; Victor Segura<sup>1, 2</sup>; <sup>1</sup>*Bioinformatics Platform, CIMA, Pamplona, Spain*; <sup>2</sup>*IdiSNA, Pamplona, Spain*; <sup>3</sup>*Vicomtech, San Sebastian, Spain*; <sup>4</sup>*Proteomics Unit, CNB, Madrid, Spain*
- Poster 083 **Chromosome 19: Progress in Hunting Missing Proteins and Future Strategies.**; Jeovanis Gil<sup>1</sup>; Ramiro Alonso Bastida<sup>1</sup>; Ariadna Ortega Lozano<sup>1</sup>; Leopoldo Gómez Caudillo<sup>1</sup>; Magdalena Hernández Ortiz<sup>1</sup>; Alejandro García Carranca<sup>2, 3</sup>; [Sergio Encarnacion-Guevara](#)<sup>1</sup>; <sup>1</sup>*CCG-Universidad Nacional Autónoma de México, Cuernavaca, Morelos, México*; <sup>2</sup>*IIB-Universidad Nacional Autónoma de México, Ciudad de México, México*; <sup>3</sup>*Instituto Nacional de Cancerología, Ciudad de México, México*
- Poster 084 **Creating a Complete Human Full-Length Plasmid Collection for C-HPP and Proteomics Studies**; [Jin Park](#); Vel Murugan; Joseph Miceli; Mitch Magee; Joshua LaBaer; *Arizona State University, Tempe, Arizona*
- Poster 085 **Chromosome 17 Missing Proteins: Recent Progress and Future Directions as Part of the Next-50MP Challenge**; [Hongjiu Zhang](#); Omer Siddiqui; Yuanfang Guan; Gilbert Omenn; *University of Michigan, Ann Arbor, MI*
- Poster 086 **Multi-Omics Data Analysis Pipeline for Identifying, Functional Annotating of Single Amino-Acid Variants**; [SooYoun Lee](#)<sup>1</sup>; Heeyoun Hwang<sup>1, 2</sup>; Young Mook Kang<sup>1</sup>; Ji Eun Jeong<sup>1</sup>; Jin Young Kim<sup>1</sup>; Jong Shin Yoo<sup>1</sup>; <sup>1</sup>*Biomedical Omics Research Center, Korea Basic Science Institute, Ochang, 28119, Republic of Korea*; <sup>2</sup>*Graduated School of Analytical, Science and Technology, Chungnam National, University, Daejeon, 34134 Republic of Korea*
- Poster 087 **Congenital Zika Syndrome (Microcephaly) Signature in Amniotic Fluid**; [Gilberto B Domont](#)<sup>1</sup>; Fabio CS Nogueira<sup>1</sup>; Rafael Melani<sup>1</sup>; Adriana SO Melo<sup>2</sup>; <sup>1</sup>*Fed Univ of Rio de Janeiro, Rio De Janeiro, Brazil*; <sup>2</sup>*IPESQ, Campina Grande, Brazil*
- Poster 088 **ASV-ID, a Proteogenomic Analysis Method for Identifying Alternative Splice Variants of the Human Proteome**; [Seul-Ki Jeong](#); Chae-Yeon Kim; Young-Ki Paik; *Yonsei Proteome Research Center, Seoul, South Korea*
- Poster 089 **Deciphering the Dark Proteome: Use of the Testis and Characterization of Two Dark Proteins**; Nathalie Melaine<sup>1</sup>; Emmanuelle Com<sup>1</sup>; Pascale Bellaud<sup>2</sup>; Laetitia Guillot<sup>1</sup>; Mélanie Lagarrigue<sup>1</sup>; Nick A. Morrice<sup>3</sup>; Blandine Guével<sup>1</sup>; Régis Lavigne<sup>1</sup>; Juan-Felipe Velez de la Calle<sup>4</sup>; Jörg Dojahn<sup>5</sup>; [Charles Pineau](#)<sup>1</sup>; <sup>1</sup>*PROTIM - Irset - Inserm U1085, Rennes, France*; <sup>2</sup>*H2P2 Core Facility, UMS BioSit, Univ Rennes, Rennes, France*; <sup>3</sup>*Sciex, Phoenix House Lakeside Drive Centre Park, Warrington, UK*; <sup>4</sup>*Unité FIV, Clinique Pasteur, Brest, France*; <sup>5</sup>*Sciex, Landwehrstr. 54, Darmstadt, Germany*
- Poster 090 **Improvement of Peptide Separation for Exploring the Missing Proteins Localized on Membranes**; [Zhilong Lin](#)<sup>1, 2</sup>; Yuanliang Zhang<sup>1, 2</sup>; Piliang Hao<sup>3</sup>; Kexia Hou<sup>3</sup>; Yuanyuan Sui<sup>3</sup>; Keren Zhang<sup>1, 2</sup>; Yanbin He<sup>1, 2</sup>; Hong Li<sup>4</sup>; Huanming Yang<sup>1, 5</sup>; Siqi Liu<sup>1, 2</sup>; Yan Ren<sup>1, 2</sup>; <sup>1</sup>*BGI-Shenzhen, Shenzhen, China*; <sup>2</sup>*China National GeneBank, Shenzhen, China*; <sup>3</sup>*ShanghaiTech University, Shanghai, China*; <sup>4</sup>*Shenzhen Seventh People's Hospital, Shenzhen, China*; <sup>5</sup>*James D. Watson Institute of Genome Sciences, Hangzhou, China*

### CHROMATIN DYNAMICS

#### Poster 091

- Poster 091 **Measuring Histone Protein Dynamics by Hydrogen-Deuterium Exchange Mass Spectrometry**; [Geoffrey Dann](#); Kelly Karch; Abigail Lemmon; Benjamin Garcia; *University of Pennsylvania, Philadelphia, PA*

### CLINICAL PROTEOMICS

#### Posters 092 - 107

- Poster 092 **Clinical Proteomics Analysis Using Data Independent Acquisition (DIA) Identified Classifiers for Molecular Characterization of Lymphoma**; Haikuo Li<sup>1</sup>; Jinghan Wang<sup>2</sup>; Wenjuan Yu<sup>2</sup>; Fang Yu<sup>2</sup>; Zhongqi Li<sup>2</sup>; Xin Ku<sup>1</sup>; Jie Jin<sup>2</sup>; [Wei Yan](#)<sup>1</sup>; <sup>1</sup>*Shanghai Jiao Tong University, Shanghai, China*; <sup>2</sup>*The First Affiliated Hospital, Zhejiang University, Hangzhou, China*
- Poster 093 **Development of Robust and Reproducible Renal Proteome Assays from Kidney Biopsies**; Wouter Knoi<sup>1</sup>; Petra Jansen<sup>1</sup>; Jesper Kers<sup>1, 2</sup>; [Garry Corthals](#)<sup>1</sup>; <sup>1</sup>*University of Amsterdam, Amsterdam, The Netherlands*; <sup>2</sup>*Amsterdam University Medical Centre, Amsterdam, The Netherlands*
- Poster 095 **Biomarker Candidate Discovery in Blood and Cerebral Spinal Fluid from patients with Neurodegenerative Diseases**; [Shaochun Zhu](#)<sup>1, 2</sup>; Gunnar Wingsle<sup>3</sup>; Lars Forsgren<sup>1</sup>; Miles Trupp<sup>1</sup>; <sup>1</sup>*Pharmacology and Clinical Neuroscience, Umeå Unive, Umea, Sweden*; <sup>2</sup>*Umeå Biotech Incubator, Umea, Sweden*; <sup>3</sup>*Swedish University of Agricultural Sciences SLU, umea, sweden*

## POSTERS

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- Poster 096 **Investigation of Targetable Biomarkers for Non-Small-Cell Lung Carcinoma (NSCLC) in Human Blood Plasma**; Barbara Helm<sup>1</sup>; Magdalena Szczygiel<sup>1, 3</sup>; Alexander Gorol<sup>1</sup>; Marvin Wäsch<sup>1</sup>; Marc Schneider<sup>2, 3</sup>; Thomas Muley<sup>2, 3</sup>; Ursula Klingmüller<sup>1, 3</sup>; <sup>1</sup>German Cancer Research Center (DKFZ), Heidelberg, Germany; <sup>2</sup>Thoraxklinik at Heidelberg University, Heidelberg, Germany; <sup>3</sup>Translational Lung Research Center, Member of DZL, Heidelberg, Germany
- Poster 097 **Differential Plasma Proteomic Analysis of B-thalassemia Patients in Response to Hydroxyurea Treatment**; Muhammad Zohaib<sup>1</sup>; Saqib Ansari<sup>2</sup>; Tahir shamsi<sup>2</sup>; Roman Zubarev<sup>3</sup>; Shamshad Zarina<sup>1</sup>; <sup>1</sup>NCP, University of Karachi, Karachi, PK; <sup>2</sup>NIBD, Karachi, PK; <sup>3</sup>Karolinska Institute, Stockholm, Se
- Poster 098 **Peripheral Immunophenotypes in Non-Small-Cell Lung Cancer Patients Carrying Different EGFR Genotypes**; Yu-Teng Jheng<sup>1</sup>; Po-Hao Feng<sup>1, 2</sup>; Zi-Ming Huang<sup>2</sup>; San-Yuan Wang<sup>1</sup>; Kang-Yun Lee<sup>1</sup>; Ching-Shan Luo<sup>2</sup>; Sheng-Ming Wu<sup>1</sup>; Chia-Li Han<sup>1</sup>; <sup>1</sup>Taipei Medical University, Taipei, Taiwan; <sup>2</sup>Shuang Ho Hospital, New Taipei City, Taiwan
- Poster 099 **Immunodepletion Using MARS-14 Column Enables Higher Coverage of CSF Proteome than "Equalization" by ProteoMiner Hexapeptide Ligand Library.**; Eliska Doktorova; Marek Svitek; Karel Holada; Jiri Petrak; First Faculty of Medicine, Charles University, Prague, Czech Republic
- Poster 100 **Identification and Overexpression of Proteins during Human Corneal Epithelial Wound Healing *in vitro* Model**; Shamim Mushtaq<sup>1</sup>; Meraj Zehra<sup>1</sup>; Nikhat Ahmed<sup>2</sup>; <sup>1</sup>Ziauddin University, Karachi, Pakistan; <sup>2</sup>Barrett Hodgson University, Karachi, Please Select
- Poster 101 **Distinction between Molecular Subtypes Group 3 and Group 4 of Medulloblastoma Using Quantitative Proteomics**; Lenka Hernychova<sup>1</sup>; Marta Nekulova<sup>1</sup>; Marta Jezova<sup>2</sup>; Lenka Dosedelova<sup>1</sup>; Michaela Scigelova<sup>3</sup>; Borivoj Vojtesek<sup>1</sup>; Karel Zitterbart<sup>4</sup>; <sup>1</sup>Masaryk Memorial Cancer Institute, Brno, CZ; <sup>2</sup>Department of Pathology, University Hospital, Brno, CZ; <sup>3</sup>Thermo Fisher Scientific, Bremen, D; <sup>4</sup>Department of Pediatric Oncology, University Hosp., Brno, CZ
- Poster 102 **Differential Analysis of the Proteome of Ovarian Endometriosis**; Urja Jaiswal<sup>1</sup>; Raj Kumar Yadav<sup>1</sup>; Alka Kriplani<sup>2</sup>; Kallol Kumar Roy<sup>2</sup>; <sup>1</sup>Dept. of Physiology, AIIMS, New Delhi, India; <sup>2</sup>Dept. of Obs & Gynae, AIIMS, New Delhi, India
- Poster 103 **Translating Serologic Response to the *Candida albicans* Cell Wall-Associated Proteome during Dimorphic Transition into a Molecular Discriminator for Invasive Candidiasis**; Aida Pitarch; César Nombela; Concha Gil; Complutense University, Madrid, Spain
- Poster 104 **Proteomics of Laser-captured Microdissected Glomeruli and Tubulointerstitium Reveals Compartment-Specific Altered Extracellular Matrix of Kidney Allografts with Antibody-Mediated Rejection**; Sergi Clotet<sup>1</sup>; Caitriona McEvoy<sup>1</sup>; Ihor Batruch<sup>2</sup>; Max Kotlyar<sup>3, 4</sup>; Chiara Pastrello<sup>3, 4</sup>; Julie Van<sup>1</sup>; Andrea Bozovic<sup>5</sup>; Vathany Kulasingam<sup>5</sup>; PeiXuen Chen<sup>6</sup>; Eleftherios P Diamandis<sup>2</sup>; Igor Jurisica<sup>3, 4</sup>; Andrzej Chruscinski<sup>7</sup>; Rohan John<sup>1, 5</sup>; Ana Konvalinka<sup>1, 7</sup>; <sup>1</sup>Toronto General Hospital Research Institute, UHN, Toronto, Canada; <sup>2</sup>Lunenfeld-Tanenbaum Research Institute, MSH, UoT, Toronto, Canada; <sup>3</sup>Kremlil Research Institute, UHN, Toronto, Canada; <sup>4</sup>Dpt of Medical Biophysics & Computer Science, UoT, Toronto, Canada; <sup>5</sup>Laboratory Medicine and Pathobiology, UHN, Toronto, Canada; <sup>6</sup>Dpt of Medicine & Institute of Medical Science, UoT, Toronto, Canada; <sup>7</sup>Dpt of Medicine, Division of Nephrology, UHN, Toronto, Canada
- Poster 105 **The Investigation of Drug Resistance of *Acinetobacter baumannii* Based on DIA Quantitative Proteomics Approach**; Ming Ke<sup>1</sup>; Naikei Wong<sup>2</sup>; Yan Ren<sup>1</sup>; Siqi Liu<sup>1</sup>; <sup>1</sup>BGI, Shenzhen, China; <sup>2</sup>The third people's hospital of Shenzhen, Shenzhen, China
- Poster 106 **Cross-Omics Analysis of Proteome and Transcriptome Dynamics During Clinical Peritoneal Dialysis Therapy**; Klaus Kratochwill; Rebecca Herzog; Andreas Vychytil; Christoph Aufricht; Medical University of Vienna, Vienna, Austria
- Poster 107 **Diagnosis of Malignant Pleural Mesothelioma Cancer Relying on Targeted Proteomics in Blood**; Ferdinando Cerciello<sup>1</sup>; Meena Choi<sup>2</sup>; Sara L Sinicropi-Yao<sup>1</sup>; Katie Lomeo<sup>1</sup>; Joseph M. Amann<sup>1</sup>; Emanuela Felley-Bosco<sup>3</sup>; Rolf A. Stahel<sup>3</sup>; Bruce Robinson<sup>4</sup>; Jenette Creaney<sup>4</sup>; Harvey I. Pass<sup>5</sup>; Olga Vitek<sup>2</sup>; David P. Carbone<sup>1</sup>; <sup>1</sup>James Thoracic Center, The Ohio State University, Columbus, OH; <sup>2</sup>Northeastern University, Boston, MA; <sup>3</sup>University Hospital Zürich, Zürich, Switzerland; <sup>4</sup>University of Western Australia, Nedlands, Western Australia; <sup>5</sup>New York University, New York, NY

## COMPUTATION, INFORMATICS AND BIG DATA Posters 108 - 133

- Poster 108 **Automated Workflow Composition in Mass Spectrometry-Based Proteomics**; Magnus Palmblad<sup>1</sup>; Anna-Lena Lamprecht<sup>2</sup>; Jon Ison<sup>3</sup>; Veit Schwämmle<sup>4</sup>; <sup>1</sup>Leiden University, Leiden, Netherlands; <sup>2</sup>Utrecht University, Utrecht, Netherlands; <sup>3</sup>Technical University of Denmark, Kongens Lyngby, Denmark; <sup>4</sup>University of Southern Denmark, Odense, Denmark

## POSTERS

All posters displayed Monday - Wednesday. Odd-numbered posters present Monday.  
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- Poster 109 **Comprehensive Computational Pipeline for Conventional MS/MS Data Processing and Open Search-Based PTM Characterization;** Felipe da Veiga Leprevost; Andy Kong; Dmitry Avtonomov; Hui-Yin Chang; Guo Ci Teo; Daniel Geiszler; Alexey Nesvizhskii; *University of Michigan, Ann Arbor, MI*
- Poster 110 **Functional Analysis of Protein Lists Using Popular Proteins Across Human Diseases;** Edward Lau<sup>1</sup>; Maggie Pui Yu Lam<sup>2</sup>; <sup>1</sup>*Stanford University, Palo Alto, CA*; <sup>2</sup>*University of Colorado Anschutz Medical Campus, Aurora, CO*
- Poster 111 **Charge Deconvolution of Crowded Spectra;** Andrew Nichols; Elizabeth Yang; Yong J. Kil; Marshall Bern; *Protein Metrics Inc., San Carlos, CA*
- Poster 112 **A Novel Computational Strategy for Top Down Proteomics, Based on All Ion Fragmentation and Capillary Electrophoresis;** Andrew Collins<sup>1</sup>; Ranjeet Bhamber<sup>2</sup>; Andrew Dowsey<sup>2</sup>; Matthias Vonderach<sup>1</sup>; Claire Evers<sup>1</sup>; Andrew Jones<sup>1</sup>; <sup>1</sup>*University of Liverpool, Liverpool, United Kingdom*; <sup>2</sup>*University of Bristol, Bristol, UK*
- Poster 113 **An Online Service for Proteomics Data Mining Using Clustered Spectra;** Mingze Bai<sup>1,2</sup>; Johannes Griss<sup>4</sup>; Yasset Perez-Riverol<sup>3</sup>; Weimin Zhu<sup>1</sup>; Juan Antonio Vizcaino<sup>3</sup>; Henning Hermjakob<sup>1,3</sup>; <sup>1</sup>*National Center for Protein Sciences, Beijing, China*; <sup>2</sup>*Chongqing Key Lab on Big Data for Bio Intelligence, Chongqing, China*; <sup>3</sup>*European Bioinformatics Institute, Hinxton, United Kingdom*; <sup>4</sup>*Medical University of Vienna, Vienna, Austria*
- Poster 114 **OpenProt Unveils Yet Unseen Depths of Eukaryotic Proteomes;** Sebastien Leblanc; *Sherbrooke University, Sherbrooke, Canada*
- Poster 115 **Using Sub-Ranked Database Matching Scores for Improving the Peptide and Protein Identification Performance;** Ying-Lan Chen; Wei-Hung Chang; Yet-Ran Chen; *Academia Sinica, Taipei, Taiwan*
- Poster 116 **TACO, a Database Integrating Transcriptome Alterations, Pathway and Prognosis in Cancers;** Tingwen Chen; Po-Hao Chou; Jau-Song Yu; *Chang Gung University, Kwei-Shan, Taiwan*
- Poster 117 **Development and Validation of The multi-marker panel for Diagnosis of Pancreatic Cancer Using Deep Learning Algorithm;** Yoseop Kim<sup>1</sup>; Hyunsoo Kim<sup>2,3</sup>; Jin-Young Jang<sup>4</sup>; Youngsoo Kim<sup>1,2</sup>; <sup>1</sup>*Bioengineering, Seoul National University, Seoul, South Korea*; <sup>2</sup>*Biomedical Sciences, Seoul National University, Seoul, South Korea*; <sup>3</sup>*Biomedical Engineering, Seoul National University, Seoul, South Korea*; <sup>4</sup>*Surgery, Seoul National University, Seoul, South Korea*
- Poster 118 **Impact of Different Quantitation and Normalization Algorithms on Proteomics-based Biomarker Discovery – A Case Study on Lung Cancer Tissue Data;** Ching-Tai Chen<sup>1</sup>; Jen-Hung Wang<sup>1</sup>; Yi-Ju Chen<sup>2</sup>; Yu-Ju Chen<sup>2</sup>; Ting-Yi Sung<sup>1</sup>; <sup>1</sup>*Institute of Information Science, Academia Sinica, Taipei, Taiwan*; <sup>2</sup>*Institute of Chemistry, Academia Sinica, Taipei, Taiwan*
- Poster 119 **Improved Survival Prognostication of Node-Positive Malignant Melanoma Patients Utilizing Shotgun Proteomics Guided by Histopathological Characterization and Genomic Data;** Jonatan Eriksson<sup>1</sup>; Krzysztof Pawlowski<sup>1,3</sup>; Peter Horvatovich<sup>2</sup>; Gyorgy Marko-Varga<sup>1</sup>; <sup>1</sup>*Lund University, Lund, Sweden*; <sup>2</sup>*University of Groningen, Groningen, Netherlands*; <sup>3</sup>*Warsaw University of Life Sciences, Warsaw, PL*
- Poster 120 **Improved Peptide Identification in Shotgun Proteomics Data Using an Efficient Open Search Engine;** Hao Chi<sup>1</sup>; Chao Liu<sup>1</sup>; Hao Yang<sup>1</sup>; Wen-Feng Zeng<sup>1</sup>; Wen-Jing Zhou<sup>1</sup>; Yue-He Ding<sup>2</sup>; Yao Zhang<sup>3</sup>; Zhen-Lin Chen<sup>1</sup>; Rui-Xiang Sun<sup>2</sup>; Tao Liu<sup>1</sup>; Guang-Ming Tan<sup>1</sup>; Meng-Qiu Dong<sup>2</sup>; Ping Xu<sup>3</sup>; Pei-Heng Zhang<sup>1</sup>; Si-Min He<sup>1</sup>; <sup>1</sup>*Institute of Computing Technology, CAS, Beijing, China*; <sup>2</sup>*National Institute of Biological Sciences, Beijing, Beijing, China*; <sup>3</sup>*Beijing Institute of Lifeomics, Beijing, China*
- Poster 121 **ProteinExplorer: a Repository-Scale Resource for Exploration of Protein Detection in Public Mass Spectrometry Datasets;** Benjamin Pullman<sup>1</sup>; Julie Wertz<sup>1</sup>; Jeremy Carver<sup>1</sup>; Nuno Bandeira<sup>1,2</sup>; <sup>1</sup>*Computer Science and Engineering, UC San Diego, La Jolla, CA*; <sup>2</sup>*Skaggs School of Pharmacy UC San Diego, La Jolla, CA*
- Poster 122 **Fast and Efficient Mapping of Peptide Sequences and their Variants to Proteome Databases Using Full Inverted Indices;** Luis Mendoza; Eric Deutsch; Robert Moritz; *Institute for Systems Biology, Seattle, WA*
- Poster 123 **ProDiGy<sup>KDS</sup>: Towards the Omics Datasets Analyses of Precision Medicine Based on the PMap;** Dong Li; *Beijing Institute of Life Omics, Beijing, China*
- Poster 124 **Evaluation of Protein-Protein Interaction Detection Methods as a source of Capturing Domain-Motif Interactions;** Sobia Idrees; Richard Edwards; *University of New South Wales, Sydney, Australia*
- Poster 125 **Reactome Multi-Scale Pathway Visualisation;** Antonio Fabregat Mundo<sup>1</sup>; Kostas Sidiropoulos<sup>1</sup>; Guilherme Viteri<sup>1</sup>; Cristoffer Sevilla<sup>1</sup>; Henning Hermjakob<sup>1,2</sup>; <sup>1</sup>*European Bioinformatics Institute (EMBL-EBI), Cambridge, United Kingdom*; <sup>2</sup>*National Center for Protein Sciences, Beijing, China*



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Even-numbered posters present Tuesday. All posters present Wednesday morning.

Poster 126 **Proteostasis Network in NAFLD Mice from Heavy Water Metabolic Labeling and LC-MS;** [Rovshan Sadygov](#)<sup>1</sup>; Kwangwon Lee<sup>2</sup>; Sergei Ilichenko<sup>2</sup>; Takhar Kasumov<sup>2</sup>; Ahmad Borzou<sup>1</sup>; <sup>1</sup>University of Texas Medical Branch, Galveston, TX; <sup>2</sup>North East Ohio Medical University, Rootstown, OH

Poster 127 **The Human Proteome as of 2018, from the HUPO Human Proteome Project;** [Gilbert Omenn](#)<sup>1</sup>; Lydie Lane<sup>2</sup>; Eric W. Deutsch<sup>3</sup>; Jochen Schwenk<sup>4</sup>; Christopher Overall<sup>5</sup>; Fernando J. Corrales<sup>6</sup>; Jennifer Van Eyk<sup>7</sup>; Mark Baker<sup>8</sup>; Michael P. Snyder<sup>9</sup>; Young-Ki Paik<sup>10</sup>; <sup>1</sup>University of Michigan, Ann Arbor, MI; <sup>2</sup>Swiss Institute of Bioinformatics, Geneva, Switzerland; <sup>3</sup>Institute for Systems Biology, Seattle, WA; <sup>4</sup>SciLifeLab, Stockholm, Sweden; <sup>5</sup>University of British Columbia, N/A, N/A; <sup>6</sup>Centro Nacional de Biotecnología (CSIC), Madrid, Spain; <sup>7</sup>Cedars Sinai Medical Center, N/A, N/A; <sup>8</sup>Macquarie University, Sydney, Australia; <sup>9</sup>Stanford University, Palo Alto, CA; <sup>10</sup>Yonsei University, Seoul, Korea

Poster 128 **Clustering Tandem Mass Spectra Using Locality Sensitive Hashing;** [Lei Wang](#); Sujun Li; Haixu Tang; Indiana University, Bloomington, IN

Poster 129 **The UniProt Knowledgebase – from Genome to Proteome to Function;** Sandra Orchard; EMBL-EBI, Cambridge, United Kingdom

Poster 130 **Global Detection and Quantification of Modified Ribonucleosides from RNA Using a HPLC Coupled Mass Spectrometry Approach;** [Selene Swanson](#)<sup>1</sup>; Michael Washburn<sup>1,2</sup>; Laurence Florens<sup>1</sup>; <sup>1</sup>Stowers Institute for Medical Research, Kansas City, MO; <sup>2</sup>University of Kansas Medical Center, Kansas City, KS

Poster 131 **Batch Effects in Large-Scale Proteomic Studies: Diagnostics and Correction;** [Jelena Ćuklina](#)<sup>1,2</sup>; Chloe Lee<sup>1</sup>; Evan G. Williams<sup>1</sup>; Tatjana Sajic<sup>1</sup>; Ben C Collins<sup>1</sup>; María Rodríguez Martínez<sup>2</sup>; Patrick Pedrioli<sup>1</sup>; Ruedi Aebersold<sup>1</sup>; <sup>1</sup>ETH Zürich, Zurich, Switzerland; <sup>2</sup>IBM Zurich Research Laboratory, Rüschlikon, Switzerland

Poster 132 **ProteomeTools: Update of the World's Largest Synthetic Peptide and Data Resource for Human Proteome Research;** [Daniel Paul Zolg](#)<sup>1</sup>; Mathias Wilhelm<sup>1</sup>; Siegfried Gessulat<sup>1,3</sup>; Tobias Schmidt<sup>1</sup>; Patroklos Samaras<sup>1</sup>; Karsten Schnatbaum<sup>2</sup>; Johannes Zerweck<sup>2</sup>; Ulf Reimer<sup>2</sup>; Hans-Christian Ehrlich<sup>3</sup>; Pedro Navarro<sup>4</sup>; Bernard Delanghe<sup>4</sup>; Andreas Huhmer<sup>5</sup>; Bernhard Kuster<sup>1,6</sup>; <sup>1</sup>Technical University of Munich, Freising, Germany; <sup>2</sup>JPT Peptide Technologies GmbH, Berlin, Germany; <sup>3</sup>SAP SE, Potsdam, Germany; <sup>4</sup>Thermo Fisher Scientific, Bremen, Germany; <sup>5</sup>Thermo Fisher Scientific, San Jose, CA; <sup>6</sup>Bavarian Center for Biomolecular Mass Spectrometry, Freising, Germany

Poster 133 **Mutated Nucleotide and Amino-acid sequence Generator (MuNAGE): Novel Proteogenomics Software to Generate Sample-Specific Database;** [Emi Hattori](#); Kumiko Shiozawa; Tadashi Kondo; Division of Rare Cancer Research, NCC, Tokyo, Japan

### CROSS-LINKING / MOLECULAR PAINTING Posters 134 - 135

Poster 134 **Identification of Cross-Linked Peptides and Oxidation Products in CRP Exposed to UV and Rose Bengal-Mediated Oxidation;** Michele Mariotti; University Of Copenhagen, Copenhagen, Denmark

Poster 135 **Molecular Architecture of the Antiophidic Protein Dm64 and Its Complex with Myotoxin II from *Bothrops asper* Venom;** Barbara S. Soares<sup>1</sup>; Surza L. G. Rocha<sup>1</sup>; Diogo B. Lima<sup>2</sup>; Fabio C. Gozzo<sup>3</sup>; Borries Demeler<sup>4</sup>; Tayler Williams<sup>4</sup>; Janelle Arnold<sup>4</sup>; Tatiana A. C. B. Souza<sup>5</sup>; Jonas Perales<sup>1</sup>; Richard H. Valente<sup>1</sup>; Bruno Lomonte<sup>6</sup>; Francisco Gomes-Neto<sup>1</sup>; [Ana Gisele C. Neves-Ferreira](#)<sup>1</sup>; <sup>1</sup>Oswaldo Cruz Institute, Fiocruz, Rio de Janeiro, Brazil; <sup>2</sup>Pasteur Institute, Paris, France; <sup>3</sup>University of Campinas, Campinas, Brazil; <sup>4</sup>University of Texas Health Science Center, San Antonio, USA; <sup>5</sup>Carlos Chagas Institute, Fiocruz, Curitiba, Brazil; <sup>6</sup>Clodomiro Picado Institute, San José, Costa Rica

### DATA-INDEPENDENT ACQUISITION (DIA) Posters 136 - 147

Poster 136 **Comprehensive Phosphoproteomics Analysis of Drug-Treated Cardiomyocytes Using Versatile DIA Workflows;** [Nathalie Selevsek](#)<sup>1</sup>; Laura Kunz<sup>1</sup>; Carla Pluess<sup>2</sup>; Adrian Roth<sup>2</sup>; Ralph Schlapbach<sup>1</sup>; <sup>1</sup>FGCZ, ETH Zurich, Zurich, Switzerland; <sup>2</sup>Roche Innovation Center Basel, Basel, Switzerland

Poster 137 **DISCO: Exploration of DIA Data Using Data-Driven Analysis;** [David Shteynberg](#); Mukul Midha; Michael Hoopmann; Samuel Bader; Luis Mendoza; Eric Deutsch; Robert Moritz; Institute for Systems Biology, Seattle, WA

Poster 138 **Strategies and Challenges for Big Clinical SWATH-MS Dataset Analysis;** [Mukul Midha](#); David Campbell; Michael R. Hoopmann; David Shteynberg; Ulrike Kusebauch; Christopher L. Moss; Robert L. Moritz; Institute for Systems Biology, Seattle, WA

Poster 139 **Accelerating DIA Studies to Extend Workflow Utility, Using Fast Microflow LC Gradients;** [Christie Hunter](#)<sup>1</sup>; Nick Morrice<sup>2</sup>; Zuzana Demianova<sup>3</sup>; <sup>1</sup>SCIEX, Redwood City, CA; <sup>2</sup>SCIEX, Warrington, UK; <sup>3</sup>SCIEX, Darmstadt, Germany

Poster 140 **Simplifying the Use of Ion Libraries during Data Processing of Data Independent Acquisition Proteomics Data;** [Christie Hunter](#)<sup>1</sup>;

## POSTERS

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Matt Huebsch<sup>2</sup>; Adam Lau<sup>2</sup>; Kathleen Lewis<sup>1</sup>;  
Sara Ahadi<sup>4</sup>; Nick Morrice<sup>5</sup>; Arianna Jones<sup>3</sup>;  
<sup>1</sup>SCIEX, Redwood City, CA; <sup>2</sup>SCIEX, Concord,  
Canada; <sup>3</sup>SCIEX, Framingham, MA; <sup>4</sup>Stanford,  
Palo Alto, CA; <sup>5</sup>SCIEX, Warrington, United  
Kingdom

Poster 141 **Accurate Benchmarking of Acquisition Parameters and Processing Softwares for Data Independent Acquisition Analyses of Proteomic Samples**; Clarisse Gotti-Barban; Florence Roux-Dalvai; Claudine Lamothe; Frédéric Fournier; Arnaud Droit; *CHU de Quebec - Laval University, Quebec, QC*

Poster 142 **MS2 Chromatograms Alignment for Improved Protein Quantification in Large-Scale Targeted Proteomics Studies**; Shubham Gupta<sup>1</sup>; Sara Ahadi<sup>2</sup>; Hannes Röst<sup>1</sup>; *1*University of Toronto, Toronto, Canada; *2*Stanford University School of Medicine, Stanford, CA

Poster 143 **Identification, Quantification and Monitoring of Low-Abundance Host Cell Proteins during Monoclonal Antibody Purification**; Catalin Doneanu<sup>1</sup>; Alex Xenopoulos<sup>2</sup>; Romas Skudas<sup>2</sup>; Mark Bennett<sup>1</sup>; Ying Qing Yu<sup>1</sup>; Asish Chakraborty<sup>1</sup>; Weibin Chen<sup>1</sup>; *1*Waters, Milford, Massachusetts; *2*EMD Millipore, Bedford, MA

Poster 144 **Comparison of Quantitative Reproducibility between DDA Precursor and DIA Fragment Quantification Techniques**; Seth Just<sup>1</sup>; Susan Weintraub<sup>2</sup>; Sammy Pardo<sup>2</sup>; Jacob Lippincott<sup>1</sup>; Susan Ludwigsen<sup>2</sup>; Brian Searle<sup>1</sup>; *1*Proteome Software, Portland, OR; *2*University of Texas HSC, San Antonio, TX

Poster 145 **Pros and Cons of Isobaric Labelling Quantification and Label Free Single Shot DIA**; Jan Muntel<sup>1</sup>; Roland Bruderer<sup>1</sup>; Joanna Kirkpatrick<sup>2</sup>; Oliver Bernhardt<sup>1</sup>; Lynn Verbeke<sup>1</sup>; Tejas Gandhi<sup>1</sup>; Ting Huang<sup>3</sup>; Olga Vitek<sup>3</sup>; Alessandro Ori<sup>2</sup>; Lukas Reiter<sup>1</sup>; *1*Biognosys AG, Schlieren, Switzerland; *2*Leibniz Institute on Aging, Jena, Germany; *3*Northeastern University, Boston, MA

Poster 146 **Machine Learning of 1566 Prostate Proteomes Generated by PCT-SWATH Uncovers a Protein Signature Predicting Survival that Outperforms Gleason Score**; Tiannan Guo<sup>1,2</sup>; Qing Zhong<sup>3,4</sup>; Tiansheng Zhu<sup>1</sup>; Rohan Shah<sup>4</sup>; Guobo Chen<sup>5</sup>; Rebecca Poulos<sup>4</sup>; Jelena Ljubicic<sup>3</sup>; Peter Hains<sup>4</sup>; Natasha Lucas<sup>4</sup>; Yi Zhu<sup>1,2</sup>; Rutishauser Dorothea<sup>3</sup>; Rui Sun<sup>1</sup>; Hannes Roest<sup>2</sup>; George Rosenberger<sup>2</sup>; Janis Neumann<sup>6</sup>; Konstantina Charmpi<sup>6</sup>; Matteo Manica<sup>7</sup>; Marija Buljan<sup>2</sup>; Wenguang Shao<sup>2</sup>; Guan Ruan<sup>1</sup>; Niels Rupp<sup>3</sup>; Daniel Schirmacher<sup>2</sup>; Pedrioli Patrick<sup>2</sup>; Maria Rodriguez<sup>7</sup>; Andreas Beyer<sup>6,8</sup>; Roger Reddel<sup>4</sup>; Phil Robinson<sup>4</sup>; Peter Wild<sup>3,9</sup>; Ruedi Aebersold<sup>2,10</sup>; *1*Westlake University, Hangzhou, China; *2*ETH Zurich, Zurich, Switzerland; *3*University Hospital Zurich, Zurich, Switzerland; *4*ProCan, Children's Medical Research Institute, USYD, Sydney, New South

Wales, Australia; <sup>5</sup>People's Hospital of Hangzhou Medical College, Hangzhou, China; <sup>6</sup>CECAD, University of Cologne, Cologne, Germany; <sup>7</sup>IBM, Zurich, Switzerland; <sup>8</sup>CMMC, University of Cologne, Cologne, Germany; <sup>9</sup>University Hospital Frankfurt, Frankfurt am Main, Germany; <sup>10</sup>University of Zurich, Zurich, Switzerland

Poster 147 **Investigating the Plasma Proteome Using a Novel Data Independent (DIA) Approach for Determining the Mechanistic Processes Involved in Respiratory Conditions**; Christopher Hughes; Lee Gethings; Robert S. Plumb; *Waters Corporation, Wilmslow, UK*

### ENVIRONMENTAL PROTEOMICS Poster 149

Poster 149 **Protein Profiling of Drought Tolerance and – Susceptible Soybean Cultivars Showed Major Shift in Antioxidant and Defense Proteins' Abundance**; Ramesh Katam<sup>1</sup>; Kambham R Reddy<sup>2</sup>; Mahya Bahmani<sup>1</sup>; *1*Florida A&M University, Tallahassee, FL; *2*Mississippi State University, Mississippi State, MS

### GLYCOPROTEOMICS AND GLYCOMICS Posters 150 - 170

Poster 150 **Differentiation of  $\alpha$ 2,3 and  $\alpha$ 2,6 Sialic Acid-Linked Glycan Isomers Using Differential Mobility Spectrometry**; Randy Arnold<sup>1</sup>; Catherine Lane<sup>2</sup>; Kirsty McManus<sup>3</sup>; Philip Widdowson<sup>3</sup>; Sarah Flowers<sup>4</sup>; Gerard Powell<sup>3</sup>; Ian Anderson<sup>3</sup>; J. Larry Campbell<sup>5</sup>; *1*SCIEX, Redwood City, CA; *2*SCIEX, Warrington, UK; *3*Allergan Biologics Limited, Liverpool, UK; *4*Georgetown University, Washington, DC; *5*SCIEX, Concord, CA

Poster 151 **Breast Cancer Tumor Necrosis Associated Peptide and Glycan Co-localizations in FFPE Tissues by MALDI Imaging Mass Spectrometry**; Danielle Scott; Laura Spruill; Pegggi Angel; Richard Drake; *Medical University of SC, Charleston, SC*

Poster 152 **Automatic Identification and Quantitation of Site-Specific N- and O-Glycoproteins in Human Serum with IQ-GPA and Database Search**; Gun Wook Park<sup>1</sup>; Young-Mook Kang<sup>1</sup>; Ju Yeon Lee<sup>1</sup>; Hyun Kyoung Lee<sup>1,2</sup>; Jin Young Kim<sup>1</sup>; Jong Shin Yoo<sup>1,2</sup>; *1*Korea Basic Science Institute, Cheongju-Si, South Korea; *2*Chungnam National University, Daejeon, South Korea

Poster 153 **Development of Biomarker for Biliary Tract Cancer and Cholelithiasis Using Serum Haptoglobin Glycan**; Miyako Nakano<sup>1</sup>; Taiki Sugiyama<sup>1</sup>; Shiro Takahashi<sup>1</sup>; Eiji Miyoshi<sup>2</sup>; *1*Hiroshima University, Higashi-Hiroshima, Japan; *2*Osaka University Graduate School of Medicine, Suita, Japan

Poster 154 **ABO Antigens on the Epithelial Cell Membrane Decrease in Cancerous Parts;**

## POSTERS

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- Poster 154 **Asaka Naya**<sup>1</sup>; Yoshimi Higashi<sup>1</sup>; Miyo Oda<sup>2</sup>; Koji Arihiro<sup>2</sup>; Miyako Nakano<sup>1</sup>; *<sup>1</sup>Hiroshima University, Higashi-Hiroshima, Japan; <sup>2</sup>Hiroshima University Hospital, Hiroshima, Japan*
- Poster 155 **Influence of Glucan on DNA Repair in Human Lymphocytes after Radiation Exposure**; Dr.Thulasi Pillai; *Model College, Mumbai, India*
- Poster 156 **Hydrophilic Multilayer Mesoporous Magnetic Probe for Endogenous Glycopeptides Analysis from Complex Biosample**; Yilin Li; *Fudan University, Shanghai, China*
- Poster 157 **Glycopeptide Fragmentation Optimization and Quantitation by Multi Collision Energy Ramp Scanning Quadrupole Data Independent Acquisition**; Lee Gethings<sup>1</sup>; Christopher Hughes<sup>1</sup>; YiJu Chen<sup>2</sup>; YuJu Chen<sup>2</sup>; Johannes Vissers<sup>1</sup>; *<sup>1</sup>Waters, Wilmslow, United Kingdom; <sup>2</sup>Academia Sinica, Taipei, Taiwan*
- Poster 158 **In-Depth Structural Characterization of Erythropoietin**; Minkyung So; Myung-Sin Lim; Byoung Joon Ko; *Division of drug screening and evaluation, New Dru, Osong medical innovation foundation Osong saengmyu, Heungdeok-gu, Cheongju-si, Chungbuk,*
- Poster 159 **Proteomics and Intact Glycoproteomics Analysis of Exosomes from Macrophage Cell Lines**; Jialin Liu; Pengyuan Yang; *Fudan, Shanghai, China*
- Poster 160 **The Nature of Phosphatidylinositol Mannosidases (PIMs) Interaction with the PPE68 Protein – Revealing Novel Insights in its Immunogenicity and Virulence**; Nagender Rao Rameshwaram<sup>1</sup>; Kristina Thomsson Hulthe<sup>2</sup>; Rahila Qureshi<sup>1</sup>; Chunsheng Jin<sup>2</sup>; Niclas G Karlsson<sup>2</sup>; Sangita Mukhopadhyay<sup>1</sup>; *<sup>1</sup>Centre For DNA Fingerprinting & Diagnostics, Hyderabad, India; <sup>2</sup>Medical Biochemistry, University of Gothenburg, Gothenburg, Sweden*
- Poster 161 **Glycoproteomics-Based Signatures for Tumor Subtyping and Clinical Outcome in Human High-Grade Serous Ovarian Cancer**; Jianbo Pan; Yingwei Hu; Shisheng Sun; Lijun Chen; Jianying Zhou; Michael Schnaubelt; Minghui Ao; Jiang Qian; Zhen Zhang; Daniel W. Chan; Hui Zhang; *Johns Hopkins University, Baltimore, MA*
- Poster 162 **GlycoStore: A Bioinformatics Platform for LC and CE Glycomics Data**; Matthew Campbell<sup>1</sup>; Sophie Zhao<sup>2</sup>; Jodie Abrahams<sup>1</sup>; Ian Walsh<sup>2</sup>; Louise Royle<sup>3</sup>; Pauline Rudd<sup>2</sup>; *<sup>1</sup>Institute for Glycomics, Gold Coast, Australia; <sup>2</sup>Bioprocessing Technology Institute, A\*STAR, Singapore, Singapore; <sup>3</sup>Ludger Ltd, Abingdon, United Kingdom*
- Poster 163 **Genomic, Proteomic and Glycoproteomic Characterization of Human High-Grade Serous Ovarian Carcinoma**; Yingwei Hu; Jianbo Pan; Punit Shah; Minghui Ao; Lijun Chen; Michael Schnaubelt; Jiang Qian; Zhen Zhang; Daniel W. Chan; Hui Zhang; *Johns Hopkins University, Baltimore, MD*
- Poster 164 **Exploring the Cell-, Protein- and Tumour-Grade-Specific N-Glycosylation Forming the Prostate Cancer Tumour-Microenvironment**; Rebeca Kawahara Sakuma<sup>1,2</sup>; Christopher Ashwood<sup>1</sup>; Hannes Hinneburg<sup>1</sup>; Saulo Recuero<sup>3</sup>; Miguel Srougi<sup>3</sup>; Katia R. M. Leite<sup>3</sup>; Nicole H. Packer<sup>1</sup>; Giuseppe Palmisano<sup>2</sup>; Morten Thaysen-Andersen<sup>1</sup>; *<sup>1</sup>Macquarie University, Sydney, Australia; <sup>2</sup>University of São Paulo, São Paulo, Brazil; <sup>3</sup>Faculdade de Medicina da USP, São Paulo, Brazil*
- Poster 165 **Automated Annotation of Glycoproteomics Mass Spectrometry Studies Enabled by the Integration of DrawGlycan with GlycoPAT**; Sriram Neelamegham; Kai Cheng; Alan Friedman; Jun Qu; *State Univ. of New York, Buffalo, NY*
- Poster 166 **mOGP 1.0-Making O-glycoproteomics More Convenient and Meaningful**; Weiqian Cao; Jiangming Huang; Mengxi Wu; Yang Zhang; Biyun Jiang; Pengyuan Yang; *Fudan University, Shanghai, China*
- Poster 167 **Towards Universal Glycoproteome Analysis Using pGlycoNovo: Intact N-Glycopeptide Profiling Across Seven Model Species**; Mingqi Liu<sup>1</sup>; Wenfeng Zeng<sup>2</sup>; Weiqian Cao<sup>1</sup>; Huali Shen<sup>1</sup>; Simin He<sup>2</sup>; Pengyuan Yang<sup>1</sup>; *<sup>1</sup>Fudan University, Shanghai, China; <sup>2</sup>Chinese Academy of Sciences, Beijing, China*
- Poster 168 **Protein Glycosylation in Pancreatic Ductal Adenocarcinoma and its Implication in Chemoresistance**; Sheng Pan<sup>1</sup>; Teresa Brentnall<sup>2</sup>; Ru Chen<sup>2</sup>; *<sup>1</sup>University of Texas Health Science Center, Houston, TX; <sup>2</sup>University of Washington, Seattle, WA*
- Poster 169 **1<sup>st</sup> Human Glycoproteomics Initiative (HGI) Study: Community Evaluation of Software for Automated Intact Glycopeptide Identification by Mass Spectrometry**; Morten Thaysen-Andersen<sup>1</sup>; Daniel Kolarich<sup>2</sup>; Rebeca Kawahara Sakuma<sup>1,8</sup>; Hannes Hinneburg<sup>1</sup>; Kai-Hooi Khoo<sup>3</sup>; Katalin Medzihradsky<sup>4</sup>; Joseph Zaia<sup>5</sup>; Goran Larsson<sup>6</sup>; Stuart Haslam<sup>7</sup>; Giuseppe Palmisano<sup>8</sup>; Jong Shin Yoo<sup>9</sup>; Nicole H. Packer<sup>1,2</sup>; *<sup>1</sup>Macquarie University, Sydney, Australia; <sup>2</sup>Griffith University, Southport, Australia; <sup>3</sup>Academia Sinica, Taipei, Taiwan; <sup>4</sup>The University of California, San Francisco, CA; <sup>5</sup>Boston University, Boston, MA; <sup>6</sup>Gothenburg University, Gothenburg, Sweden; <sup>7</sup>Imperial College, London, UK; <sup>8</sup>University of Sao Paulo, Sao Paulo, Brazil; <sup>9</sup>Korea Basic Science Institute, Daejeon, Korea*
- Poster 170 **Rapid Profiling of Prostate Cancer-Specific PSA Glycoforms as a Specificity-Enhanced Secondary Biomarker**; Yoshimi Haga<sup>1</sup>; Motohide Uemura<sup>2</sup>; Kentaro Inamura<sup>3</sup>; Kengo Takeuchi<sup>3,4</sup>; Norio Nonomura<sup>2</sup>; Koji Ueda<sup>1</sup>;

## POSTERS

All posters displayed Monday - Wednesday. Odd-numbered posters present Monday.  
Even-numbered posters present Tuesday. All posters present Wednesday morning.

<sup>1</sup>Cancer Proteomics Group, JFCR, Tokyo, Japan; <sup>2</sup>Department of Urology, Osaka University Graduate S, Osaka, Japan; <sup>3</sup>Division of Pathology, JFCR, Tokyo, Japan; <sup>4</sup>Pathology Project for Molecular Targets, JFCR, Tokyo, Japan

### HIGH RESOLUTION MASS SPECTROMETRY Posters 172 - 173

- Poster 172 **Quantitative Phosphoproteomics Reveals *in vivo* Boron Deficiency Induced Signaling Dynamics in Arabidopsis Roots**; Yanmei Chen; *China Agricultural University, Beijing, China*
- Poster 173 ***de novo* Sequencing of Tandem Mass Spectra Reveals Dark Matter of Cyclopeptidomics**; Bahar Behsaz<sup>1</sup>; Hosein Mohimani<sup>4</sup>; Alexey Gurevich<sup>2</sup>; Andrey Prjibelski<sup>2</sup>; Mark F. Fisher<sup>3</sup>; Larry Smarr<sup>1</sup>; Pieter C. Dorrestein<sup>1</sup>; Joshua S. Mylne<sup>3</sup>; Pavel A. Pevzner<sup>1</sup>; <sup>1</sup>UC San Diego, La Jolla, CA; <sup>2</sup>Saint Petersburg State University, Saint Petersburg, Russia; <sup>3</sup>The University of Western Australia, Crawley, Australia; <sup>4</sup>Carnegie Mellon University, Pittsburgh, PA

### IMAGING Posters 174 - 175

- Poster 174 **Mass Spectrometry Imaging of Synovium Reveals Molecular Profiles with Diagnostic Potential in Arthritis**; Beatriz Rocha<sup>1</sup>; Berta Cillero-Pastor<sup>2</sup>; Cristina Ruiz-Romero<sup>1</sup>; Andrea Cuervo<sup>3</sup>; Ron M A Heeren<sup>2</sup>; Juan D Cañete<sup>3</sup>; Francisco J Blanco<sup>1</sup>; <sup>1</sup>Proteomics Group-GIR-Proteored/ISCII INIBIC-CHUAC, A Coruña, Spain; <sup>2</sup>M4i Institute-IMS Division/Maastricht University, Maastricht, The Netherlands; <sup>3</sup>Arthritis Unit, Hospital Clinic/IDIBAPS, Barcelona, Spain
- Poster 175 **Optimised Desorption Electrospray Ionisation Mass Spectrometry Imaging (DESI-MSI) Method for the Analysis of Proteins/Peptides Directly from Tissue Sections**; Mark Towers<sup>1</sup>; James Hughes<sup>2</sup>; Rian Griffiths<sup>2</sup>; Patricia Lalor<sup>2</sup>; Helen Cooper<sup>2</sup>; Emmanuelle Claude<sup>1</sup>; <sup>1</sup>Waters, Wilmslow, United Kingdom; <sup>2</sup>University of Birmingham, Birmingham, UK

### IMMUPEPTIDOMICS Posters 176 - 181

- Poster 176 **A Chemical Derivatization Strategy for Extending the Identification of MHC Class I Immunopeptides**; Rui Chen; Francois Fauteux; Simon Foote; Jacek Stupak; Tammy-Lynn Trembley; Komal Gurnani; Kelly Fulton; Risini Weeratna; Susan Twine; Jianjun Li; *National Research Council Canada, Ottawa, Canada*
- Poster 177 **A New Algorithm for Identification of Immunopeptides from LC-MS Data with High Sensitivity**; Lin He; Lei Xin; Xin Chen; Baozhen

Shan; *Bioinformatics Solutions Inc., Waterloo, Canada*

- Poster 178 **Detection of Citrullinated Residues in Glucose-Regulated Protein 78 in Human Islets of Langerhans by LC-MS/MS Using Data Dependent Acquisition**; Aisha Callebaut<sup>1</sup>; Mijke Buitinga<sup>1</sup>; Marco Bugliani<sup>2</sup>; Etienne Waelkens<sup>1</sup>; Piero Marchetti<sup>2</sup>; Rita Derua<sup>1</sup>; Chantal Mathieu<sup>1</sup>; Lut Overbergh<sup>1</sup>; <sup>1</sup>KU Leuven, Leuven, Belgium; <sup>2</sup>Pisa University, Pisa, Italy
- Poster 179 **Immunoproteomic Approach for Identification of Allergenic Proteins in Pecan Nut and Read Oak Pollen Grains**; José Ángel Huerta Ocampo<sup>1</sup>; Alejandra Valenzuela Corral<sup>2</sup>; María Del Refugio Robles Burgueño<sup>2</sup>; Ana María Guzmán Partida<sup>2</sup>; Miguel Ángel Hernández Oñate<sup>1</sup>; Joel David Flores Rivas<sup>3</sup>; Luis Manuel Terán Juárez<sup>4</sup>; <sup>1</sup>CONACYT-Centro de Investigación en Alimentación y, Hermosillo, Mexico; <sup>2</sup>CIAD, A.C., Hermosillo, Sonora; <sup>3</sup>IPICYT, A.C, San Luis Potosí, México; <sup>4</sup>Instituto Nacional de Enfermedades Respiratorias, Mexico City, México
- Poster 180 **Characterisation of Flucloxacillin-Modified Proteins Leading to the Presentation of Flucloxacillin-Modified MHC Peptides and their Importance in iDILI**; James Waddington<sup>1</sup>; Xiaoli Meng<sup>1</sup>; Patricia Illing<sup>2</sup>; Arun Tailor<sup>1</sup>; Rosalind Jenkins<sup>1</sup>; Anthony Purcell<sup>2</sup>; Dean Naisbitt<sup>1</sup>; B. Kevin Park<sup>1</sup>; <sup>1</sup>University of Liverpool, Liverpool, United Kingdom; <sup>2</sup>Monash University, Melbourne, Australia
- Poster 181 **Constrained *de novo* Sequencing of neo-Epitope Peptides Using Tandem Mass Spectrometry**; Sujun Li; Alex DeCourcy; Haixu Tang; *Indiana University Bloomington, Bloomington, IN*

### METABOLOMICS Posters 182 - 186

- Poster 182 **CCSPredict: Using a Machine Learning Approach for Higher Confidence in Lipid Identification**; Lucy Woods; Sebastian Wegner; Heiko Neuweger; Ulrike Schweiger-Hufnagel; Sven Meyer; Aiko Barsch; Nikolas Kessler; *Bruker Daltonics GmbH, Bremen, Germany*
- Poster 183 **High Throughput Targeted Workflows for Metabolomics / Lipidomics Studies**; Christie Hunter<sup>1</sup>; Khatereh Motamedchaboki<sup>1</sup>; Mackenzie Pearson<sup>1</sup>; Santosh Kapil<sup>2</sup>; Paul Baker<sup>2</sup>; <sup>1</sup>SCIEX, Redwood City, CA; <sup>2</sup>SCIEX, Framingham, MA
- Poster 184 **Metabolomics Profiling of Parapneumonic Effusion Reveals Regulatory Roles of Dipeptides for Neutrophils**; Pei-Chun Hsueh; Chih-Ching Wu; *Chang Gung University, Taoyuan, Taiwan*
- Poster 185 **Analysis of RNA Mononucleosides by DIA nanoLC-MS/MS Can Reveal Inducible tRNA Wobble Position Modification Abundance Changes**; Kevin A. Janssen<sup>1</sup>; Marianne

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Even-numbered posters present Tuesday. All posters present Wednesday morning.

Kramer<sup>2</sup>; Ranran Wu<sup>1</sup>; Brian D. Gregory<sup>2</sup>;  
Benjamin A. Garcia<sup>1</sup>; <sup>1</sup>University of  
Pennsylvania School of Medicine, Philadelphia,  
PA; <sup>2</sup>University of Pennsylvania, Philadelphia,  
PA

Poster 186 **Developments of Liquid Chromatography-Mass Spectrometry Methods for Endogenous Metabolites in Glycolysis and Endogenous Nucleotides Used in Cancer Therapy;** Zhenyun Zhu; Jing Gao; Hongwen Zhu; Hu Zhou; Shanghai Institute of Materia Medica, Chinese Acad, Shanghai, China

### METAPROTEOMICS Posters 187 - 188

Poster 187 **Metaproteomic Analysis of the Infant Fecal Microbiome.;** Laetitia Cortes; Caprion Biosciences Inc., Montreal, Canada

Poster 188 **Prediction-Based Reduction of the Search Space Leads to Increased Identifications in Metaproteomics Without Impacting Sensitivity;** Tim Van Den Bossche<sup>1,2</sup>; <sup>1</sup>VIB - UGent Center for Medical Biotechnology, Ghent, Belgium; <sup>2</sup>Department of Biochemistry, Ghent University, Ghent, Belgium

### MICROBIOME ANALYSIS Poster 189

Poster 189 **Influence of the Gut Microbiota on Histone Acetylation through Butyrate Oxidation;** Peder Lund<sup>1</sup>; Sarah Smith<sup>1</sup>; Johayra Simithy<sup>1</sup>; Lillian Chau<sup>1</sup>; Elliot Friedmann<sup>1</sup>; Yedidya Saiman<sup>1</sup>; Sophie Trefely<sup>2</sup>; Mariana Lopes<sup>1</sup>; Zuo-Fei Yuan<sup>1</sup>; Kevin Janssen<sup>1</sup>; Yemin Lan<sup>1</sup>; Nathaniel Snyder<sup>2</sup>; Gary Wu<sup>1</sup>; Benjamin Garcia<sup>1</sup>; <sup>1</sup>University of Pennsylvania, Philadelphia, PA; <sup>2</sup>Drexel University, Philadelphia, PA

### MULTI-OMICS Posters 190 - 201

Poster 190 **Multi-Omic Profiling of the Liver in a Rat Model of Type 2 Diabetes;** Desmond Li<sup>1</sup>; Lauren Smith<sup>1</sup>; Yen Chin Koay<sup>1,2</sup>; Holly McEwen<sup>1,3</sup>; Anthony Don<sup>1,3</sup>; John O'Sullivan<sup>1,2</sup>; Stuart Cordwell<sup>1</sup>; Melanie White<sup>1</sup>; <sup>1</sup>University of Sydney, Sydney, Australia; <sup>2</sup>Heart Research Institute, Sydney, Australia; <sup>3</sup>ACRF Centenary Cancer Research Centre, Sydney, Australia

Poster 191 **An Integrated Analysis of Proteomics, Peptidomics, Metabolomics, and Inflammation Markers for Assessment of Pre-analytical Variability of Human Plasma;** Zhijun Cao<sup>1</sup>; Jaclyn Daniels<sup>1</sup>; Beate Kamlage<sup>2</sup>; Antje Wagner-Golbs<sup>2</sup>; Mackean Maisha<sup>1</sup>; Jinchun Sun<sup>1</sup>; Laura Schnackenberg<sup>1</sup>; Lisa Pence<sup>1</sup>; Thomas Schmitt<sup>1</sup>; Sarah Rogstad<sup>3</sup>; Richard Beger<sup>1</sup>; Li-Rong Yu<sup>1</sup>; <sup>1</sup>National Center for Toxicological Research, FDA, Jefferson, AR; <sup>2</sup>Metanomics Health GmbH, Berlin, Germany; <sup>3</sup>Center for Drug Evaluation and Research, FDA, Silver Spring, MD

Poster 192 **Proteogenomic Approach Reveals Translation from Untranslated Regions in Gastric Cancer;** Jinwon Lee; Seunghyuk Choi; Seungjin Na; Eunok Paek; Hanyang University, Seoul, South Korea

Poster 193 **The Mechanisms for the Formation of Proteome Complexity Revealed by Multi-Omic Analyses;** Dong Yang; Chao Gao; Pan Shen; Fuchu He; Beijing Institute of Lifeomics, Beijing, China

Poster 194 **A High Throughput Single Platform For High throughput Quantitative MultiOmic Studies;** Billy Molloy; Lee Gethings; Robert Plumb; Waters, Wilmslow, United Kingdom

Poster 195 **A Comprehensive Integrative Multiomics Investigation of Malaria and Dengue;** Sanjeeva Srivastava; IIT Bombay, Mumbai, India

Poster 196 **Proteogenomics Landscape of Dehydration-Afflicted Grasspea: New Insights into Stress Tolerance;** Divya Rathi; Akanksha Pareek; Subhra Chakraborty; Niranjana Chakraborty; NIPGR, New Delhi, India

Poster 197 **Proteomic and Proteogenomic Heterogeneity of HeLa Cells across Laboratories: Implications for Research Reproducibility;** Yansheng Liu<sup>1</sup>; Yang Mi<sup>2</sup>; Torsten Mueller<sup>3</sup>; Saskia Kreibich<sup>4</sup>; Evan Williams<sup>3</sup>; Audrey Van Drogen<sup>3</sup>; Christelle Borel<sup>5</sup>; Pierre-Luc Germain<sup>6</sup>; Max Frank<sup>3</sup>; Isabell Bludau<sup>3</sup>; Martin Mehnert<sup>3</sup>; Michael Seifert<sup>7</sup>; Mario Emmenlauer<sup>8</sup>; Isabel Sorg<sup>8</sup>; Fedor Bezrukov<sup>5</sup>; Frederique Sloan Bena<sup>9</sup>; Hu Zhou<sup>10</sup>; Christoph Dehio<sup>8</sup>; Giuseppe Testa<sup>6</sup>; Julio Saez-Rodriguez<sup>2</sup>; Stylianos Antonarakis<sup>5</sup>; Wolf-Dietrich Hardt<sup>4</sup>; Ruedi Aebersold<sup>3,11</sup>; <sup>1</sup>Yale University School of Medicine, West Haven, CT; <sup>2</sup>JRC-COMBINE, RWTH Aachen Uni, Aachen, Germany; <sup>3</sup>Institute of Molecular Systems Biology, ETH Zurich, Zurich, Switzerland; <sup>4</sup>Institute of Microbiology, ETH Zurich, Zurich, Switzerland; <sup>5</sup>University of Geneva Medical School, Geneva, Switzerland; <sup>6</sup>European Institute of Oncology, Milan, Italy; <sup>7</sup>TU Dresden, Dresden, Germany; <sup>8</sup>University of Basel, Basel, Switzerland; <sup>9</sup>University Hospitals of Geneva, Geneva, Switzerland; <sup>10</sup>Shanghai Institute of Materia Medica, CAS, Shanghai, China; <sup>11</sup>Faculty of Science, University of Zurich, Zurich, Switzerland

Poster 198 **Personal Proteogenomic Analysis Using Haploid Genome Assemblies;** James Wright<sup>1</sup>; Lu Yu<sup>1</sup>; Jonathan Mudge<sup>2</sup>; Carrie Davis<sup>3</sup>; Thomas Gingeras<sup>3</sup>; Adam Frankish<sup>2</sup>; Jyoti Choudhary<sup>1</sup>; <sup>1</sup>The Institute Cancer Research, London, United Kingdom; <sup>2</sup>European Bioinformatics Institute, Cambridge, United Kingdom; <sup>3</sup>Cold Spring Harbor Laboratory, Woodbury, NY

Poster 199 **An Integrated Atlas of Protein Expression in Human Cancer Derived from Publicly**

## POSTERS

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**Available Datasets;** Andrew Jarnuczak<sup>1</sup>; Hanna Najgebauer<sup>1</sup>; Mitra Barzine<sup>1</sup>; Fatemeh Ghavidel<sup>2</sup>; Yasset Perez-Riverol<sup>1</sup>; Alvis Brazma<sup>1</sup>; Juan Antonio Vizcaino<sup>1</sup>; <sup>1</sup>EMBL-European Bioinformatics Institute (EMBL-EBI), Hinxton, United Kingdom; <sup>2</sup>Department of Informatics, University of Bergen, Bergen, Norway

Poster 200 **Multi-Omic Characterisation of Bladder and Lung Carcinomas Using a Novel Scanning Quadrupole DIA Acquisition Method;** Lee Gethings; Adam King; Robert Plumb; *Waters, Wilmslow, United Kingdom*

Poster 201 **Phenome Study on the Tissues of Esophageal Squamous Cell Carcinoma;** Yan Ren<sup>1</sup>; Guixue Hou<sup>1</sup>; Shaohang Xu<sup>1</sup>; Xiaomin Lou<sup>2</sup>; Siqi Liu<sup>1</sup>; <sup>1</sup>BGI-Shenzhen, Shenzhen, China; <sup>2</sup>Beijing Institute of Genomics, Beijing, China

### NEUROLOGICAL DISEASES / NEUROPROTEOMICS Posters 202 - 214

Poster 202 **Proteomic Profile of the Hippocampus from Patients with Mesial Temporal Lobe Epilepsy Reveals the Molecular Mechanisms Related to Disease Progression;** Amanda Morato Do Canto<sup>1, 7</sup>; Alexandre Barcia Godoi<sup>1, 7</sup>; André Vieira<sup>1, 2</sup>; Fabio Rogerio<sup>1, 3</sup>; Clarissa Yasuda<sup>1, 4</sup>; Enrico Ghizoni<sup>4</sup>; Helder Tedeschi<sup>4</sup>; Albert Baskar Arul<sup>5, 6</sup>; Renã A S Robinson<sup>5, 6</sup>; Fernando Cendes<sup>1, 4</sup>; Iscia Lopes Cendes<sup>1</sup>; <sup>1</sup>BRAINN - UNICAMP, Campinas, Brazil; <sup>2</sup>Biology Institute, Campinas, SP, Brazil; <sup>3</sup>Department of Pathological Anatomy, Campinas, SP, Brazil; <sup>4</sup>Department of Neurology, Campinas, SP, Brazil; <sup>5</sup>Department of Chemistry, Nashville, TN; <sup>6</sup>Vanderbilt University, Nashville, TN; <sup>7</sup>Department of Medical Genetics, Campinas, SP, Brazil

Poster 203 **Development of the CSF-PR 2.0 Tool for Exploring Cerebrospinal Fluid Mass Spectrometry Biomarker Datasets - Updated with New Data;** Astrid Guldbrandsen; Yehia Farag; Ragnhild Reehorst Lereim; Frode Berven; Harald Barsnes; *University of Bergen, Bergen, Norway*

Poster 204 **Network Based Integration of Proteomic and Genomic Data Unravels New Key Astrocytic Players in ALS;** Iñigo Barrio-Hernandez<sup>1</sup>; Pedro Beltrao<sup>1</sup>; Andras Lakatos<sup>2</sup>; <sup>1</sup>EMBL-EBI, Saffron Walden, United Kingdom; <sup>2</sup>Dept. Clin. Neuro., Cambridge, UK

Poster 205 **The Role of Circulating Extracellular Vesicles in Systemic Response to Ischemic Stroke;** Livia Rosa-Fernandes; Maja Møller-Nielsen; Martin R Larsen; Bettina Clausen; Kate Lambertsen; *University of Southern Denmark, Odense, Denmark*

Poster 206 **Phosphoproteomic Analysis of the Dorsal Dentate Gyrus Laser-Microdissected from**

**the Hippocampus of an Animal Model of Mesial Temporal Lobe Epilepsy;** Amanda Morato Do Canto<sup>1, 2</sup>; Alexandre Hilario Berenguer Matos<sup>1</sup>; Beatriz Ayoama Bertelli<sup>3</sup>; Alexandre Barcia de Godoi<sup>1</sup>; Andre Schwambach Vieira<sup>3</sup>; Iscia Lopes-Cendes<sup>1</sup>; <sup>1</sup>FCM-UNICAMP, Campinas, Brazil; <sup>2</sup>BRAINN, Campinas, Brazil; <sup>3</sup>IB-UNICAMP, Campinas, Brazil

Poster 207 **Sex-Dependent Differences in Hippocampal Proteome from Organotypic Slice Cultures.;** Simone Nardin Weis<sup>1</sup>; Marina Firmino de Oliveira<sup>1</sup>; Jaques Miranda F. Souza<sup>1</sup>; Juliana Bender Hoppe<sup>2</sup>; Alan R. Mól<sup>1</sup>; Christianne G. Salbego<sup>2</sup>; Consuelo M. R. de Lima<sup>1</sup>; Carlos André O. Ricart<sup>1</sup>; Wagner Fontes<sup>1</sup>; Marcelo Valle de Sousa<sup>1</sup>; <sup>1</sup>Universidade de Brasília, Brasilia, Brazil; <sup>2</sup>Universidade Federal do Rio Grande do Sul, Porto Alegre, Brazil

Poster 208 **Early Changes in the Human Hippocampal Proteome at the Onset of Tau Burden;** Clarissa Ferolla Mendonça<sup>1</sup>; Magdalena Kuras<sup>2</sup>; Péter Döme<sup>3</sup>; Fabio Nogueira<sup>1</sup>; Gilberto B Domont<sup>1</sup>; Melinda Rezel<sup>2</sup>; Gyorgy Marko-Varga<sup>2</sup>; <sup>1</sup>Universidade Federal do Rio de Janeiro, Rio De Janeiro, Brazil; <sup>2</sup>Lund University, Lund, Sweden; <sup>3</sup>Semmelweis University, Budapest, Hungary

Poster 209 **Intron-Mediated Enhancement Boosts Rtn4 circRNA Expression: A Robust Method for Exploring circRNA Function;** Dingding Mo; Xinping Li; *Max Planck Institute for Biology of Ageing, Cologne, Germany*

Poster 210 **Histone  $\beta$ -hydroxybutyrylation in neuroblastoma cells;** Alexander Ma; Kaichen Chu; Di Zhang; Jun Ding; Yingming Zhao; *University of Chicago, Chicago, Illinois*

Poster 211 **Silver Nanoparticle-Induced Expression of Proteins Related to Oxidative Stress and Neurodegeneration in an *in vitro* Human Blood-Brain Barrier Model;** Asif Manzoor Khan; *University of Southern Denmark, Odense, Denmark*

Poster 212 **Proteomic Analysis of Rat Hippocampus Exposed to 10-Day Morphine Treatment and Subsequent 20-Day Drug Withdrawal;** Hana Ujčíková; Michal Jager; Lenka Roubalova; Petr Svoboda; *Institute of Physiology CAS, Prague 4, Czech Republic*

Poster 213 **Unveiling the Mice Cerebellum Proteomic Changes under the Effect of Rattle Snake *Crotalus durissus terrificus* Venom;** Fabio Montoni<sup>1</sup>; Diana Andreotti<sup>2</sup>; Rosangela Eichler<sup>3</sup>; Ismael Lima<sup>1</sup>; Emer Ferro<sup>3</sup>; Hugo Armelin<sup>1</sup>; Leo Iwai<sup>1</sup>; <sup>1</sup>LETA/CeTICS, Instituto Butantan, Brazil, Sao Paulo, Brazil; <sup>2</sup>Lab of Molecular Neuropharmacology, ICB, USP, Sao Paulo, Brazil; <sup>3</sup>Lab of Pharmacology of Intracell Peptides, ICB USP, Sao Paulo, Brazil

## POSTERS

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Poster 214 **A Label-free Quantification Approach to Identify Differentially Expressed Proteins between Wild Type and Transgenic Alzheimer Rat Brains**; Pritha Bagchi; Eric Dammer; Geng Wang; Robert Cohen; Nicholas Seyfried; *Emory School of Medicine, Atlanta, Georgia*

### NEW TECHNOLOGIES Posters 215 - 239

Poster 215 **Quantitation of Intact Proteins in Human Plasma Using Top-Down Parallel Reaction Monitoring-MS**; Daojing Wang; *Newomics Inc., Berkeley, CA*

Poster 216 **Micro Pillar Array Columns: A Novel Robust Chromatography Platform for Deep and Reproducible Proteome Coverage**; Robert Van Ling<sup>1, 2</sup>; Jeff Op De Beeck<sup>1, 2</sup>; Kurt van Mol<sup>1, 2</sup>; Bo Claerebout<sup>1, 2</sup>; Natalie Van Landuyt<sup>1, 2</sup>; Wim De Malsche<sup>3</sup>; Gert Desmet<sup>3</sup>; Paul Jacobs<sup>1, 2</sup>; <sup>1</sup>PharmaFluidics, Zwijnaarde, Belgium; <sup>2</sup>PharmaFluidics, Zwijnaarde, Belgium; <sup>3</sup>Vrije Universiteit Brussel, Brussels, Belgium

Poster 217 **Comprehensive Proteome Mapping of a Human Cancer Cell Line Using LC-FAIMS Pro-MS/MS**; Romain Huguet; Satendra Prasad; Joshua Silveira; Graeme McAlister; Philip Remeš; Derek Bailey; Qingyu Song; Michael Belford; Eloy Wouters; Jean-Jacques Dunyach; Vlad Zabrouskov; Susan Abbatiello; *Thermo Fisher Scientific, San Jose, California*

Poster 218 **Fast Microflow Chromatography for Accelerating Protein Identification Experiments**; Christie Hunter<sup>1</sup>; Nick Morrice<sup>2</sup>; Zuzana Demianova<sup>3</sup>; <sup>1</sup>SCIEX, Redwood City, CA; <sup>2</sup>SCIEX, Warrington, UK; <sup>3</sup>SCIEX, Darmstadt, Germany

Poster 219 **EASyPep - A New Simplified and Optimized Workflow for MS Sample Preparation**; Amarjeet Flora; Sergei Snovida; Ryan Bomgarden; John Rogers; *Thermo Fisher Scientific, Rockford, IL*

Poster 220 **Universal Sample Processing for Highly Reproducible Proteomic Sample Preparation of Diverse Sample Types**; John Wilson<sup>1, 3</sup>; Visa Meyyappan<sup>2</sup>; Domenic Nicholas Narducci<sup>2</sup>; Ben Neely<sup>4</sup>; Jim Laugharn<sup>2</sup>; Darryl Pappin<sup>1, 3</sup>; <sup>1</sup>ProtiFi, LLC, Huntington, NY; <sup>2</sup>Covaris, Inc., Woburn, MA; <sup>3</sup>Cold Spring Harbor Laboratory, Cold Spring Harbor, New York; <sup>4</sup>NIST, Charleston, SC

Poster 221 **Toward the Ideal Mass Analyzer with Data-Independent Acquisition and Parallel Accumulation – Serial Fragmentation (diaPASEF)**; Florian Meier<sup>1</sup>; Andreas-David Brunner<sup>1</sup>; Max Frank<sup>2</sup>; Eugenia Voytik<sup>1</sup>; Markus Lubeck<sup>3</sup>; Heiner Koch<sup>3</sup>; Scarlet Koch<sup>3</sup>; Oliver Räther<sup>3</sup>; Ben C. Collins<sup>4</sup>; Ruedi Aebersold<sup>4, 5</sup>; Hannes Röst<sup>2</sup>; Matthias Mann<sup>1, 6</sup>; <sup>1</sup>Max Planck Institute of Biochemistry, Martinsried, Germany;

<sup>2</sup>Donnelly Centre for Cellular and Biomol. Research, Toronto, Canada; <sup>3</sup>Bruker Daltonik GmbH, Bremen, Germany; <sup>4</sup>ETH Zurich, Zurich, Switzerland; <sup>5</sup>University of Zurich, Zurich, Switzerland; <sup>6</sup>NNF Center for Protein Research, Copenhagen, Denmark

Poster 222 **Magnetic HILIC Microparticles Enabling Automated Offi-Line MS Sample Preparation**; Stoyan Stoychev<sup>1</sup>; Previn Naicker<sup>1</sup>; Siphon Mamputha<sup>1</sup>; Justin Jordaan<sup>2</sup>; <sup>1</sup>CSIR, Pretoria, SA; <sup>2</sup>Resyn Biosciences, Pretoria, SA

Poster 223 **Short LC-Gradients for High Throughput and Deep Shotgun Proteomics Using PASEF on a TIMS Equipped QTOF**; Thomas Kosinski; Scarlet Koch; Thorsten Ledertheil; Christian Meier-Credo; Christoph Gebhardt; Heiner Koch; *Bruker Daltonik GmbH, Bremen, Germany*

Poster 224 **Cutting-Edge nanoLC Column Technology and its Capabilities in Advanced Mass Spectrometry Proteomics**; Yufeng Shen; *CoAnn Technologies, LLC, Richland, WA*

Poster 225 **Comparison of Two Solid-Phase Extraction (SPE) Methods for the Identification and Quantification of Porcine Retinal Protein Markers by LC-MS/MS**; Carsten Schmelter; Sebastian Funke; Jana Tremli; Anja Beschnitt; Natarajan Perumal; Norbert Pfeiffer; Franz H. Grus; *University Medical Center, Mainz, Germany*

Poster 226 **Maximum MS Utilization in High-Throughput and Deep Dive Low-Flow LC-MS Proteomics**; Alexander Boychenko<sup>1</sup>; Christopher Pynn<sup>1</sup>; Wim Decrop<sup>1</sup>; Martin Ruehl<sup>1</sup>; Bart van den Berg<sup>1</sup>; Mike Baynham<sup>2</sup>; Remco Swart<sup>1</sup>; <sup>1</sup>Thermo Fisher Scientific, Germering, Germany; <sup>2</sup>Thermo Fisher Scientific, Runcorn, UK

Poster 227 **Proteome Level de novo Sequencing with a Pair of Newly Developed Mirror Proteases of super-LysargiNase and Ac-Trypsin**; Ping Xu<sup>1</sup>; Hao Yang<sup>2</sup>; Yanchang Li<sup>3</sup>; Ming-Zhi Zhao<sup>1</sup>; Wei-di Xiao<sup>3</sup>; Yi-hao Wang<sup>3</sup>; Jun-Ling Zhang<sup>3</sup>; Christopher Overall<sup>4</sup>; Hao Chi<sup>2</sup>; Si-min He<sup>2</sup>; <sup>1</sup>Beijing Proteome Research Center, Changping District, China; <sup>2</sup>Institute of Computing Technology, Beijing, China; <sup>3</sup>National Center for Protein Sciences Beijing, Beijing, China; <sup>4</sup>Centre for Blood Research, University of British C, Vancouver, Canada

Poster 228 **Results As Soon As Possible (rASAP): 2 Hours from Lysis to Label Free Quantification of Cells and Tissues Using Subtilisin**; Humberto Gonczarowska-Jorge<sup>2</sup>; Stefan Lorocho<sup>2</sup>; Margherita Dell'Aica<sup>1</sup>; Albert Sickmann<sup>2</sup>; Christoph Borchers<sup>1, 3</sup>; Kristina Lorenz<sup>2</sup>; Andreas Roos<sup>2</sup>; Rene Zahedi<sup>1, 2</sup>; <sup>1</sup>Lady Davis Proteomics Centre, Montreal, Canada; <sup>2</sup>ISAS, Dortmund, Germany; <sup>3</sup>UVic-Genome BC Proteomics Centre, Victoria, BC

## POSTERS

All posters displayed Monday - Wednesday. Odd-numbered posters present Monday.  
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- Poster 229 **Snapshots of the EGFR Signaling Pathway Acquired with High Temporal Resolution Using a Microfluidic Device;** Margherita Dell'Aica<sup>1,2</sup>; Pedro Novo<sup>2</sup>; Denisa Hathazi<sup>2</sup>; Albert Sickmann<sup>2</sup>; Andreas Roos<sup>2</sup>; Dirk Janasek<sup>2</sup>; Rene Zahedi<sup>2,3</sup>; <sup>1</sup>Lady Davis Proteomics Centre, Montreal, Canada; <sup>2</sup>ISAS, Dortmund, Germany; <sup>3</sup>McGill University, Montreal, Canada
- Poster 230 **High Resolution Peptide Separation for Illuminating Human Proteome;** Yasushi Ishihama; Kosuke Ogata; Koshi Imami; Naoyuki Sugiyama; *Kyoto University, Kyoto, Japan*
- Poster 231 **Developing Tools to Facilitate Blood-Based Protein Biomarker Discovery in a Non-Model Organism;** Benjamin Neely<sup>1</sup>; Florian Marty<sup>2</sup>; Marion Neely<sup>3</sup>; Lori Schwacke<sup>4</sup>; <sup>1</sup>NIST, Charleston, SC; <sup>2</sup>Biognosys AG, Schlieren, Switzerland; <sup>3</sup>JHT, Inc. (NCCOS/NOS/NOAA), Charleston, SC; <sup>4</sup>National Marine Mammal Foundation, San Diego, CA
- Poster 232 **Screening and Epitope Mapping of Antibodies for Immuno-Mass Spectrometric Assays Using a Novel immuno-MALDI (iMALDI) Approach;** Huiyan Li<sup>1,2</sup>; Claudia Fredolini<sup>3</sup>; Jochen Schwenk<sup>3</sup>; Christoph Borchers<sup>1,2</sup>; <sup>1</sup>Jewish General Hospital Proteomics Centre, McGill, Montreal, Canada; <sup>2</sup>UVic-Genome BC Proteomics Centre, Victoria, Canada; <sup>3</sup>Science for Life Laboratory, KTH, Solna, Sweden
- Poster 233 **A Genetic Approach Toward Mass Spectrometry-Based Comprehensive and Sensitive Quantification of Yeast Proteome;** Keiji Kito; *Meiji University, Kawasaki, Japan*
- Poster 234 **Quantifying the Fetal Tissue Translatome Using a Novel Isotopic Labeling Approach Reveals Temporal and Tissue-Specific Regulatory Networks during Development;** Josue Baeza<sup>1</sup>; Barbara Coons<sup>2</sup>; William Peranteau<sup>2</sup>; Benjamin Garcia<sup>1</sup>; <sup>1</sup>University of Pennsylvania, Philadelphia, Pennsylvania; <sup>2</sup>Children's Hospital of Philadelphia, Philadelphia, Pennsylvania
- Poster 235 **Advance Analysis of Proteomics and Peptidomics by Ion Mobility Mass Spectrometry for Biomarker Discovery;** Yoshitoshi Hirao; Amr Elguoshy; Bo Xu; Keiko Yamamoto; Tadashi Yamamoto; *BBC, Niigata University, Niigata, Japan*
- Poster 236 **Development of a Functional Proteomics Technology for Biomarker and Drug target Discovery;** Xing Wang; *Array Bridge Inc., St. Louis, Missouri*
- Poster 237 **Improved iST Workflows for the Streamlined Analysis of Tissues and High-Throughput Preparation of Samples Using Isobaric Labelling;** Fabian Hosp; Garwin Pichler; Nils Kulak; *PreOmics, Martinsried, Germany*
- Poster 238 **Cell Lysate Microarray for Mapping the Network of Genetic Regulators for Histone Marks;** Li Cheng<sup>1</sup>; Junbiao Dai<sup>2</sup>; Sheng-Ce Tao<sup>1</sup>; <sup>1</sup>Shanghai Jiao Tong University, Shanghai, China; <sup>2</sup>Chinese Academy of Sciences, Shenzhen, China
- Poster 239 **Development of an Efficient Proteomics Sample Preparation Method for Human Gut Proteomics;** Rakesh Singh<sup>1</sup>; Om Prakash<sup>1,2</sup>; Roger Mercer<sup>1</sup>; <sup>1</sup>Florida State University, Tallahassee, FL; <sup>2</sup>National Centre for Microbial Resources, Pune, India

### PATHOGEN PROTEOMICS Posters 240 - 247

- Poster 240 **Plasma Proteome Signature of Sepsis: A Functionally Connected Protein Network;** Genaro Pimienta; *Sanford Burnham Prebys Medical Discovery Institute, La Jolla, CA*
- Poster 241 **Quantitative Proteomics and Phosphoproteomic Profiling of THP1 Cells After Dengue Infection;** Rosa Victoria Pando-Robles<sup>1</sup>; Angel Ambrocio<sup>1</sup>; Rosa del Angel<sup>2</sup>; Juan Osés-Prieto<sup>3</sup>; Cesar Batista<sup>4</sup>; Alma Burlingame<sup>3</sup>; <sup>1</sup>instituto Nacional de Salud Publica, Cuernavaca, Mexico; <sup>2</sup>CINVESTAV, Mexico city, Mexico; <sup>3</sup>University of California, San Francisco, USA; <sup>4</sup>UNAM, Cuernavaca, Mexico
- Poster 242 **Insights into the Human Pathodegradome of the *Staphylococcus aureus* V8 Protease;** Andrew Frey; Dale Chaput; Lindsey Shaw; *University of South Florida, Tampa, FL*
- Poster 243 **Systems-Wide Hijacking of Host Cells During Herpes Simplex Virus (HSV-1) Infection;** Katarzyna Kulej<sup>1,2</sup>; Ashley N. Della Fera<sup>2</sup>; Eui Tae Kim<sup>1,2</sup>; Matthew J. Charman<sup>1,2</sup>; Simone Sidoli<sup>1</sup>; Benjamin A. Garcia<sup>1</sup>; Matthew D. Weitzman<sup>1,2</sup>; <sup>1</sup>University of Pennsylvania, Philadelphia, PA; <sup>2</sup>Children's Hospital of Philadelphia, Philadelphia, PA
- Poster 244 **Tc-STAMS2: A Novel *Trypanosoma cruzi* Strain Typing Assay Using MS2 Peptide Spectral Libraries;** Gilberto Santos de Oliveira<sup>1</sup>; Rebeca Sakuma<sup>1</sup>; Livia Rosa-Fernandes<sup>2</sup>; Simon Ngao Mule<sup>1</sup>; Carla Cristi Avila<sup>1</sup>; Marta M.G. Teixeira<sup>1</sup>; Martin R. Larsen<sup>2</sup>; Giuseppe Palmisano<sup>1</sup>; <sup>1</sup>Department of Parasitology, Sao Paulo, Brazil; <sup>2</sup>Dartment of biochemistry and molecular biology, Odense, DK
- Poster 245 **The Impact of Mutations on Protein Expression Pattern in Intracellular Bacteria *Ehrlichia chaffeensis*;** Chandramouli Kondethimmanahalli; Roman Ganta; *College of Vet Medicine, Kansas State University, Manhattan, KS*
- Poster 246 **Culture Independent Label Free Method for Milk Metaproteome and Resistome Analysis;** Cristian Piras<sup>1</sup>; Alessio Soggiu<sup>1</sup>; Viviana Greco<sup>2</sup>;



## POSTERS

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Luigi Bonizzi<sup>1</sup>; Alfonso Zecconi<sup>1</sup>; Andrea Urbani<sup>3</sup>; Claudia Gusmara<sup>1</sup>; Domenico Britti<sup>4</sup>; Paola Roncada<sup>4</sup>; <sup>1</sup>DIMEVET - University of Milan, Milano, Italy; <sup>2</sup>Fondazione Santa Lucia, Rome, Italy; <sup>3</sup>Catholic University of Sacred Heart, Rome, Italy; <sup>4</sup>Università Magna Græcia, Catanzaro, Italy

Poster 247 **Identification of Serine 119 as an Effective Inhibitor Binding Site of *M. tuberculosis* Ubiquitin-Like Protein Ligase PafA**; He-Wei Jiang; *Shanghai Jiaotong University, Shanghai, China*

### POST-TRANSLATIONAL MODIFICATIONS Posters 248 - 279

Poster 248 **Ion Mobility High Resolution QTOF MS - Impact of PASEF on Detection of Post-Translational Modifications**; Allan Stensballe<sup>1</sup>; Kenneth kastaniegaard<sup>1</sup>; Thomas Bouet Guldbæk Poulsen<sup>1</sup>; Dres Damgaard<sup>2</sup>; Claus Henrik Nielsen<sup>2</sup>; <sup>1</sup>Aalborg University, Aalborg, Denmark; <sup>2</sup>Danish National Hospital, Copenhagen, Denmark

Poster 249 **Multiplexed Quantitative Analysis of APC/C-Specific Ubiquitin Substrates**; Lu Yu<sup>1</sup>; Theodoros Roumeliotis<sup>1</sup>; Gabor Bakos<sup>2</sup>; Igor Gak<sup>2</sup>; Jörg Mansfeld<sup>2</sup>; Jyoti Choudhary<sup>1</sup>; <sup>1</sup>Institute of Cancer Research, London, United Kingdom; <sup>2</sup>Technische Universität Dresden, Dresden, Germany

Poster 250 **Novel Antibody Reagents for the Characterization of Protein ADP-Ribosylation**; Matthew Fry<sup>1</sup>; Alvin Lu<sup>2</sup>; Rami Najjar<sup>1</sup>; Mario Niepel<sup>2</sup>; Matthew P Stokes<sup>1</sup>; <sup>1</sup>Cell Signaling Technology INC, Danvers, Massachusetts; <sup>2</sup>Ribon Therapeutics, Lexington, MA

Poster 251 **High Sensitivity Phosphoproteomics using PASEF on a TIMS-QTOF mass spectrometer**; Heiner Koch<sup>1</sup>; Kristina Desch<sup>2</sup>; Scarlet Koch<sup>1</sup>; Matt Willetts<sup>3</sup>; Thomas Kosinski<sup>1</sup>; Markus Lubeck<sup>1</sup>; Erin Schuman<sup>2</sup>; Julian Langer<sup>2</sup>; <sup>1</sup>Bruker Daltonik GmbH, Bremen, Germany; <sup>2</sup>Max-Planck-Institute for Brain Research, Frankfurt am Main, Germany; <sup>3</sup>Bruker Daltonics Inc., Billerica, Germany

Poster 252 **An Approach for the Site-Specific Quantitation of Protein Core Fucosylation in a Large Scale**; Yi Huang; Xianyuan Zhao; Zixiang Yu; Xiaohong Qian; Wantao Ying; *Beijing Institute of Lifeomics, Changping District, China*

Poster 253 **Quantitative Proteomics and Phosphoproteomics Analysis Revealed Different Regulatory Mechanisms of Halothane and Rendement Napole Genes in Porcine Muscle Metabolism**; Honggang Huang<sup>1</sup>; Martin Larsen<sup>2</sup>; Rene Lametsch<sup>3</sup>; <sup>1</sup>Arla Foods Ingredients, Videbæk, Denmark; <sup>2</sup>University of Southern Denmark, Odense,

*Denmark*; <sup>3</sup>University of Copenhagen, Copenhagen, Denmark

Poster 254 **Posttranslational Modifications and Data-Independent Acquisitions – Challenges and Opportunities**; Xueshu Xie<sup>1</sup>; Nathan Basisty<sup>1</sup>; Matthew Stokes<sup>2</sup>; Christie Hunter<sup>3</sup>; Kimberly Lee<sup>2</sup>; Birgit Schilling<sup>1</sup>; <sup>1</sup>The Buck Institute, Novato, CA; <sup>2</sup>Cell Signaling Technology, Inc., Danvers, MA; <sup>3</sup>SCIEX, Redwood City, CA

Poster 255 **Expanding the Citrullinome of Synovial Fibrinogen from Rheumatoid Arthritis using MS: Identification of Putative Sites of Pathogenic and Prognostic Relevance**; Mandvi Sharma<sup>1</sup>; Dres Damgaard<sup>2,3</sup>; Ladislav Senolt<sup>4</sup>; Birte Svensson<sup>1</sup>; Anne-Christine Bay Jensen<sup>5</sup>; Claus Henrik Nielsen<sup>2,3</sup>; Per Häggglund<sup>6</sup>; <sup>1</sup>Technical University of Denmark, Kgs. Lyngby, Copenhagen, Denmark; <sup>2</sup>University of Copenhagen, Copenhagen, Denmark; <sup>3</sup>Copenhagen University Hospital, Rigshospitalet, Copenhagen, Denmark; <sup>4</sup>Faculty of Medicine, Charles University, Prague, Czech Republic; <sup>5</sup>Nordic Biosciences, Herlev, Denmark; <sup>6</sup>Panum Institute, University of Copenhagen, Copenhagen, Denmark

Poster 256 **Radio-Sensitizing Effects of VE-821: Phosphoproteomic and Metabolomic Changes After ATR Inhibition in Irradiated MOLT-4 Cells**; Barbora Šalovská<sup>1,2</sup>; Hana Janečková<sup>3,4</sup>; Ivo Fabrik<sup>5</sup>; Radana Karlíková<sup>3,4</sup>; Lucie Čecháková<sup>1</sup>; Martin Ondrej<sup>1</sup>; Marek Link<sup>1</sup>; David Friedecký<sup>4</sup>; Aleš Tichý<sup>1,5</sup>; <sup>1</sup>University of Defence in Brno, Hradec Králové, Czech Republic; <sup>2</sup>Institute of Molecular Genetics of the ASCR, Prague, Czech Republic; <sup>3</sup>Palacký University Olomouc, Olomouc, Czech Republic; <sup>4</sup>University Hospital Olomouc, Olomouc, Czech Republic; <sup>5</sup>Biomedical Research Centre, University Hospital, Hradec Králové, Czech Republic

Poster 257 **Method Development for Phosphorylation and Glycosylation Detection using Orbitrap Fusion Lumos**; Susanne Breitkopf; Jeffrey A. Culver; Michelle F. Clasquin; Bei Betty Zhang; Mara Monetti; *Pfizer, Inc, Cambridge, MA*

Poster 258 **(Phospho)Proteomics Quantification strategies: A Systematic Comparison of SILAC, TMT and Label-Free Techniques to Study EGFR Signal Transduction Networks in CRC**; Markus Stepath<sup>1</sup>; Abdelouahid Maghnouj<sup>2</sup>; Birgit Zülch<sup>1</sup>; Karin Schork<sup>1</sup>; Michael Turewicz<sup>1</sup>; Martin Eisenacher<sup>1</sup>; Stephan Hahn<sup>2</sup>; Barbara Sitek<sup>1</sup>; Thilo Bracht<sup>1</sup>; <sup>1</sup>Ruhr-University Bochum - MPC, Bochum, Germany; <sup>2</sup>Ruhr-University Bochum - MGO, Bochum, Germany

Poster 259 **Adenoviral Proteins E1B55K and E4orf6 Use Non-Degradative Ubiquitination to regulate Viral Late Protein Expression**; Christin Herrmann<sup>1,2</sup>; Jennifer Liddle<sup>1,2</sup>; Joseph Dybas<sup>1,2</sup>; Benjamin Garcia<sup>2</sup>; Matthew Weitzman<sup>1,2</sup>; <sup>1</sup>Children's Hospital of Philadelphia,

## POSTERS

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*Philadelphia, PA; <sup>2</sup>University of Pennsylvania, Philadelphia, PA*

*of Pharmacy, Aurora, CO; <sup>2</sup>Barbara Davis Center for Childhood Diabetes, Aurora, CO*

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|--|---|
| <p>Poster 260 <b>A Specific Dual Functional Probe-Hydrophilic Mercaptosuccinic Acid Coupled Magnetic Mesoporous Titania for Simultaneous Capture of Glycopeptides and Phosphopeptides;</b> <u>Nianrong Sun</u>; Chunhui Deng; <i>Fudan University, Shanghai, China</i></p>   | <p>Poster 269 <b>oxSWATH: An Integrative Method for a Comprehensive Redox-Centered Analysis Combined with a Generic Differential Proteomics Screening;</b> <u>Sandra I. Anjo</u><sup>1, 2</sup>; Matilde M. Melo<sup>1</sup>; Liliana R. Loureiro<sup>1, 3</sup>; Lúcia Sabala<sup>1, 3</sup>; Pedro Castanheira<sup>4</sup>; Mário Grãos<sup>1, 4</sup>; Bruno Manadas<sup>1</sup>; <sup>1</sup><i>Center for Neuroscience and Cell Biology, UC, Coimbra, Portugal</i>; <sup>2</sup><i>Faculty of Sciences and Technology, UC, Coimbra, Portugal</i>; <sup>3</sup><i>Department of Chemistry, University of Aveiro, Aveiro, Portugal</i>; <sup>4</sup><i>Biocant, Biotechnology Transfer Association, Cantanhede, Portugal</i></p> |
| <p>Poster 261 <b>Quantitative Proteomic Approaches Identify Regulatory Enzymes and Substrates for Lysine 2-Hydroxyisobutyrylation Pathway;</b> <u>He Huang</u><sup>1</sup>; Zhouqing Luo<sup>2</sup>; Shankang Qi<sup>1</sup>; Jing Huang<sup>2</sup>; Lunzhi Dai<sup>3</sup>; Junbiao Dai<sup>4</sup>; Yingming Zhao<sup>1</sup>; <sup>1</sup><i>The University of Chicago, Chicago, 0</i>; <sup>2</sup><i>Tsinghua University, Beijing, China</i>; <sup>3</sup><i>Sichuan Univeristy, Sichuan, China</i>; <sup>4</sup><i>Shenzhen Institutes of Advanced Technology, Chines, Shenzhen, China</i></p> | <p>Poster 270 <b>Ub<sup>KEKS</sup> : A Novel ubiquitin Variant Expressed from a Pseudogene.;</b> <u>Marie-Line Dubois</u>; Patrick Delattre; Jean-François Jacques; Dominique Levesque; Vivian Delcourt; Maxime Beaudoin; Mylène Brunelle; Sondos Samandi; Marie Brunet; Pierre Lavigne; Xavier Roucou; François-Michel Boisvert; <i>University of Sherbrooke, Sherbrooke, Canada</i></p>   |
| <p>Poster 262 <b>Assessing an Automated Phosphopeptide Enrichment for High-Throughput Malignant Melanoma Phosphoproteomics from Patient Tissues;</b> <u>Jimmy Rodriguez</u>; Magdalena Kuras; Melinda Rezeli; Lazaro Betancourt; Gyorgy Marko-Varga; <i>Centre of Excellence in Biological and Medical Mas, Malmo, Sweden</i></p>  | <p>Poster 271 <b>Quantitative Proteomics Identifies Novel PIAS1 Protein Substrates Involved in Cell Migration and Motility;</b> <u>Chongyang Li</u><sup>1</sup>; Francis McManus<sup>1</sup>; Trent Nelson<sup>1</sup>; Mirela Cristina Pascariu<sup>1</sup>; Pierre Thibault<sup>1, 2</sup>; <sup>1</sup><i>Institute for Research in Immunology and Cancer, Montréal, Canada</i>; <sup>2</sup><i>Department of Chemistry, Université de Montréal, Montréal, Canada</i></p>  |
| <p>Poster 263 <b>FDR Estimation for Hybrid Mass Spectral Library Search Identifications in Bottom-up Proteomics;</b> <u>Meghan Burke</u>; Zheng Zhang; Yuri A. Mirokhin; Dmitrii V. Tchekhovskoi; Yuxue Liang; Stephen E. Stein; <i>NIST, Gaithersburg, &lt;Not Specified&gt;</i></p>  | <p>Poster 272 <b>Proteoform Atlas Of Extracellular Matrix Predicts Clade Specific Functionality Of Wall-Associated Signaling Components In Plant;</b> <u>Kanika Narula</u>; Pooja Choudhary; Arunima Sinha; Sudip Ghosh; Eman Elagamey; Niranjana Chakraborty; Subhra Chakraborty; <i>National Institute of Plant Genome Research, New Delhi, India</i></p>   |
| <p>Poster 264 <b>Proteomic-Scale Approaches for Quantifying Irreversible Cysteine Redox Post-Translational Modifications using Parallel Reaction Monitoring Mass Spectrometry in Myocardial Ischemia / Reperfusion;</b> <u>Alexander Rookyard</u>; <u>Stuart Cordwell</u>; <i>The University of Sydney, The University Of Sydney, Australia</i></p>  | <p>Poster 273 <b>Redox Regulation of Fetal and Adult Hematopoiesis;</b> <u>Kristyna Pimkova</u>; Maria Jassinskaja; Emil Johansson; Jenny Hansson; <i>Lund University, Lund, Sweden</i></p>   |
| <p>Poster 265 <b>Developing Workflow for Simultaneous Analyses of Phosphoproteomics and Glycoproteomics;</b> <u>Kyung-Cho Cho</u>; Lijun Chen; Yingwei Hu; Michael Schnaubelt; Hui Zhang; <i>Johns Hopkins, Baltimore, Maryland</i></p>  | <p>Poster 274 <b>MS Search Parameter Refinement Avoids ADPr-Acceptor Site Localization Bias and Identifies Tyrosine as Novel ADPr-Acceptor Site with Significant Functional Consequences;</b> <u>Deena Leslie Pedrioli</u><sup>1</sup>; Mario Leutert<sup>1, 3</sup>; Vera Bilan<sup>1</sup>; Kapila Gunasekera<sup>1</sup>; Kathrin Nowak<sup>1</sup>; Lars Malnström<sup>2</sup>; Michael Hottiger<sup>1</sup>; <sup>1</sup><i>University of Zürich, Zürich, Switzerland</i>; <sup>2</sup><i>3S3IT, University of Zurich, Zürich, Switzerland</i>; <sup>3</sup><i>Molecular Life Science PhD Program of the Life Sc, Zürich, Switzerland</i></p>  |
| <p>Poster 266 <b>Dehydration-Induced Alterations phosphorylation Status in the Nuclear Proteomic Landscape of Chickpea;</b> <u>Pragya Barua</u>; Dipak Gayen; Nilesh Vikram Lande; Subhra Chakraborty; Niranjana Chakraborty; <i>NIPGR, New Delhi, India</i></p>   | <p>Poster 275 <b>Global Age-Specific Changes in Protein Post Translational Modifications (PTMs) in Neonatal, Paediatric and Adult Plasma;</b> <u>Xiaomin Song</u><sup>1</sup>; Dana Pascovici<sup>1</sup>; Jemina Wu<sup>1</sup>; Paul Monagle<sup>2, 3</sup>; Mark Molloy<sup>4</sup>; <u>Vera Ignjatovic</u><sup>2, 5</sup>; <sup>1</sup><i>Australian Proteome Analysis Facility,</i></p>  |
| <p>Poster 267 <b>A Novel Method of Quantifying Protein Methylation Utilizing SWATH-MS;</b> <u>Aaron Robinson</u>; Shelly Lu; Jennifer Van Eyk; <i>Cedars Sinai Medical Center, Los Angeles, California</i></p>   |   |
| <p>Poster 268 <b>Hybrid Insulin Peptides (HIPs) are Detectable in Human Islets by Mass Spectrometry;</b> <u>Timothy Wiles</u><sup>1</sup>; Roger Powell<sup>1</sup>; Scott Beard<sup>2</sup>; Anita Hohenstein<sup>1</sup>; Cole Michel<sup>1</sup>; Thomas Delong<sup>1</sup>; <sup>1</sup><i>University of Colorado Skaggs School</i></p>  |   |

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Sydney, Australia; <sup>2</sup>The University of Melbourne, Parkville, Australia; <sup>3</sup>Royal Children's Hospital, Parkville, Australia; <sup>4</sup>Kolling Institute of Medical Research, St. Leonards, Australia; <sup>5</sup>Murdoch Children's Research Institute, Parkville, Australia

Poster 276 **An Automated and Reproducible Workflow for Human Cancer Cell Line Phosphopeptide Analysis;** Shuai Wu; Linfeng Wu; Agilent Technologies Inc., 384, <Not Specified>

Poster 277 **Do Not Dry TMT-Labeled Phosphopeptide Samples in Presence of NHS-Quenching Reagent;** Yumi Kwon; Shinyeong Ju; Cheolju Lee; Korea Institute of Science and Technology, Seoul, South Korea

Poster 278 **Effective Mass Spectrometry-Based Methods to Globally and Site-Specifically Analyze Glycoproteins;** Ronghu Wu; Georgia Institute of Technology, Atlanta, GA

Poster 279 **Revealing the Regulation of Growth and the Fatty Acid Metabolism of the Mandibular Gland of Honeybee Based on Phosphoproteomic Analysis;** Yue Hao; IAR, CAAS, Beijing, China

### PRECISION MEDICINE Posters 280 - 285

Poster 280 **Peripheral Blood Proteins Associated with Long-Term response to Lithium in Bipolar Disorder;** Klaus Oliver Schubert<sup>1,2</sup>; Georgia Arentz<sup>3</sup>; Bernhard Baune<sup>4</sup>; Peter Hoffmann<sup>3</sup>; <sup>1</sup>University of Adelaide, Adelaide, Australia; <sup>2</sup>Northern Adelaide Mental Health Service, Salisbury, Australia; <sup>3</sup>University of South Australia, Adelaide, Australia; <sup>4</sup>University of Melbourne, Melbourne, Australia

Poster 281 **Immunoproteomic Identification of IgE Binding Proteins from Ligustrum Pollen: Monosensitized- vs Polysensitized Allergic Patients;** Luis Manuel Teran<sup>1</sup>; Bessy Mani<sup>2</sup>; Jose Angel Huerta-Ocampo<sup>3</sup>; Fernando Gandhi Pavon Romero<sup>1</sup>; Ana Paulina Barba de la Rosa<sup>2</sup>; <sup>1</sup>INER, Distrito Federal, Mexico; <sup>2</sup>IPICYT, S.L.P., Mexico; <sup>3</sup>CIAD, Hermosillo, Mexico

Poster 283 **A Mass Spectrometric Proteome Profiling Workflow from Heart Tissue to Accelerate Cardiac Research and Diagnostics;** Christof Lenz<sup>1,2</sup>; Lisa Neuenroth<sup>1</sup>; Soeren Brandenburg<sup>3</sup>; Stephan E. Lehnart<sup>3,4</sup>; Henning Urlaub<sup>1,2</sup>; <sup>1</sup>Clinical Chemistry, UMC, Goettingen, DE; <sup>2</sup>MPI for Biophysical Chemistry, Goettingen, DE; <sup>3</sup>Cardiology and Pneumology, UMC, Goettingen, DE; <sup>4</sup>DZHK (German Centre for Cardiovascular Research), Goettingen, DE

Poster 284 **An Oncoproteogenomic Strategy for Multiplexed Screening of EGFR Mutations in Non-small-cell Lung Cancer;** Chi-Ting Lai<sup>1,2</sup>; Yi-Ju Chen<sup>1</sup>; Wai-Kok Choong<sup>3</sup>; Shr-Uen Lin<sup>4</sup>; Ya-Hsuan Chang<sup>5</sup>; Jie-Ning Zhang<sup>6</sup>; Sung-Liang Yu<sup>6</sup>; Ting-Yi Sung<sup>3</sup>; Hsuan-Yu Chen<sup>2,3</sup>; Chia-Li Han<sup>7</sup>; Yu-Ju Chen<sup>1,2</sup>; <sup>1</sup>Institute of Chemistry,

Academia Sinica, Taipei, Taiwan; <sup>2</sup>Genome and Systems Biology Degree Program, NTU, Taipei, Taiwan; <sup>3</sup>Institute of Information Science, Academia Sinica, Taipei, Taiwan; <sup>4</sup>Graduate institute of oncology, NTU, Taipei, Taiwan; <sup>5</sup>Institute of Statistical Science, Academia Sinica, Taipei, Taiwan; <sup>6</sup>CLSMB, College of Medicine, NTU, Taipei, Taiwan; <sup>7</sup>School of Pharmacy, TMU, Taipei, Taiwan

Poster 285 **Development of an Assay for Monitoring Plasma Protein Variants for Clinical Use to Assess Novel Therapies for Acute Liver Dysfunction;** Ivan Doykov<sup>1</sup>; Wendy Heywood<sup>1</sup>; Valeria Iansante<sup>2</sup>; Emer Fitzpatrick<sup>2</sup>; Anil Dhawan<sup>2</sup>; Celine Filippi<sup>2</sup>; Kevin Mills<sup>1</sup>; <sup>1</sup>University College London, London, United Kingdom; <sup>2</sup>Kings's College Hospital, London, UK

### PROTEIN COMPLEXES AND INTERACTOMICS Posters 286 - 294

Poster 286 **The Yeast Interactome Approaching Completeness – Combining a Robust High-Throughput Pull-Down Workflow with a Fast and Sensitive Evosep/timsTOF Setting;** André Clemens Michaelis<sup>1</sup>; Andreas-David Brunner<sup>1</sup>; Florian Meier<sup>1</sup>; Matthias Mann<sup>1,2</sup>; <sup>1</sup>Max Planck Institute of Biochemistry, Martinsried, Germany; <sup>2</sup>NNF Center for Protein Research, Copenhagen, Denmark

Poster 287 **EGFR Interactome Reveals Multiple Pathways and Regulatory Mechanism of Drug-resistance in Non-Small Cell Lung Cancer;** Pei-Shan Wu<sup>1,2</sup>; Miao-Hsia Lin<sup>2</sup>; Szu-Hua Pan<sup>3,4</sup>; Yu-Ju Chen<sup>2,5</sup>; <sup>1</sup>Genome and Systems Biology Degree Program, NTU, Taipei, Taiwan; <sup>2</sup>Institute of Chemistry, Academia Sinica, Taipei, Taiwan; <sup>3</sup>Institute of Medical Genomics and Proteomics, NTU, Taipei, Taiwan; <sup>4</sup>Degree Program of Translational Medicine, NTU, Taipei, Taiwan; <sup>5</sup>Department of Chemistry, NTU, Taipei, Taiwan

Poster 288 **Probing Novel Immunoglobulin Super Family Receptor Interactions on the Cell Surface;** Bushra Husain; Erik Verschueren; Nadia Martinez-Martin; Genentech Inc., South San Francisco, California

Poster 289 **A Sequential Affinity Purification and Mass Spectrometry Approach for Identifying Shared Interactions of Associated Protein Pairs;** Xingyu Liu<sup>1</sup>; Ying Zhang<sup>1</sup>; Jeffrey Lange<sup>1</sup>; Brian Slaughter<sup>1</sup>; Jay Unruh<sup>1</sup>; Tim Wen<sup>1</sup>; Laurence Florens<sup>1</sup>; Susan Abmayr<sup>1,2</sup>; Jerry Workman<sup>1</sup>; Michael Washburn<sup>1,2</sup>; <sup>1</sup>Stowers Institute for Medical Research, Kansas City, MO; <sup>2</sup>University of Kansas Medical Center, Kansas City, KS

Poster 290 **Proteomics Uncovers Lipid raft protein NTAL as a Regulator of Leukemia cells Proliferation and Death;** Carolina Thome<sup>1,2</sup>; Germano Ferreira<sup>1,2</sup>; Andreia Leopoldino<sup>1</sup>; Gustavo de Souza<sup>3</sup>; Eduardo Magalhães Rego<sup>1</sup>.

## POSTERS

All posters displayed Monday - Wednesday. Odd-numbered posters present Monday.  
Even-numbered posters present Tuesday. All posters present Wednesday morning.

<sup>2</sup>; Vitor Faça<sup>1,2</sup>; <sup>1</sup>University of São Paulo, Ribeirão Preto, Brazil; <sup>2</sup>Cell-Based Therapy Center, Ribeirão Preto, Brazil; <sup>3</sup>Federal University of Rio Grande do Norte, Natal, Brazil

Simone Sidoli<sup>1</sup>; Zuo-Fei Yuan<sup>1</sup>; Peder J. Lund<sup>1</sup>; Xiaolu Zhao<sup>2</sup>; Benjamin A. Garcia<sup>1</sup>; <sup>1</sup>University of Pennsylvania, Philadelphia, PA; <sup>2</sup>Wuhan University, Wuhan, China

Poster 291 **Proximity-Based Proteomic Profiling of DNA Double-Strand Break Repair Proteins Identifies Shieldin Complex as Novel Regulator of NHEJ**; Rajat Gupta<sup>1</sup>; Kumar Somyajit<sup>1</sup>; Takeo Narita<sup>1</sup>; Elina Maskey<sup>1</sup>; Andre Stanlie<sup>2</sup>; Magdalena Kremer<sup>3</sup>; Dimitris Typas<sup>1</sup>; Michael Lammers<sup>3</sup>; Niels Mailand<sup>1</sup>; Andre Nussenzweig<sup>2</sup>; Jiri Lukas<sup>1</sup>; Chunaram Choudhary<sup>1</sup>; <sup>1</sup>The NNF Center for Protein Research, Copenhagen, Denmark; <sup>2</sup>National Institutes of Health, Bethesda, USA; <sup>3</sup>Institute for Genetics and CECAD, Cologne, Germany

Poster 292 **Multilayered Proteomic Analysis of Cancer-Related Mutations in the Dyrk2 Kinase Complex**; Martin Mehnert; Rodolfo Ciuffa; Fabian Frommelt; Federico Uliana; Audrey van Drogen; Matthias Gstaiger; Ruedi Aebersold; ETH Zurich, Zurich, Switzerland

Poster 293 **Assembling Active Histone Deacetylases into Chromatin Remodelers: A Complex Task Illuminated Using Well-Placed Affinity Tags to Probe Protein Interaction Networks**; Charles Banks<sup>1</sup>; Sayem Miah<sup>1</sup>; Mark Adams<sup>1</sup>; Cassandra Eubanks<sup>1</sup>; Janet Thornton<sup>1</sup>; Laurence Florens<sup>1</sup>; Michael Washburn<sup>1,2</sup>; <sup>1</sup>Stowers Institute for Medical Research, Kansas City, MO; <sup>2</sup>University of Kansas Medical Center, Kansas City, KS

Poster 294 **Interactome Analysis of ER $\alpha$  and AP-2 $\gamma$  in breast cancer cells**; Edwin Cheung; University of Macau, Taipa, Macau

### PROTEIN QUALITY CONTROL Poster 295

Poster 295 **Systematic Analysis of the Mitochondrial Protein Synthesis Network**; Heaseung Sophia Chung; J. Wade Harper; Harvard Medical School, Boston, MA

### PROTEOFORM BIOLOGY Posters 296 - 298

Poster 296 **Evaluating the timsTOFPRO Bottom-Up Proteomics Platform Potential for Proteoform Profiling and Top-Down approaches**; Pierre-Olivier Schmit<sup>1</sup>; Kristina Marx<sup>2</sup>; Gary Kruppa<sup>3</sup>; <sup>1</sup>Bruker France S.A, Wissembourg, France; <sup>2</sup>Bruker Daltonik GmbH, Bremen, Germany; <sup>3</sup>Bruker Daltonics, Inc, Billerica, USA

Poster 297 **H2AV Lysine Crotonylation: An Epigenetic Switch During Human Myogenic Differentiation**; Natarajan Bhanu; Zuo-fei Yuan; Benjamin A Garcia; Epigenetics Institute, Perelman School Of Medicine, Philadelphia, PA

Poster 298 **Metabolic Labeling and Quantitative Proteomics for Interrogation of Proteome-Wide Acetylation Dynamics**; Yekaterina Kori<sup>1</sup>;

### PROTEOMICS IN AGEING AND AGE-RELATED DISEASES Posters 299 - 301

Poster 299 **Cellular Senescence; A Driver Of The Pro-aging Side Effects Of Antiretroviral Therapies**; Chisaka Kuehnemann<sup>1</sup>; Nathan Basisty<sup>1</sup>; Christopher Wiley<sup>1</sup>; Birgit Schilling<sup>1</sup>; Judith Campisi<sup>1,2</sup>; <sup>1</sup>Buck Institute, Novato, CA; <sup>2</sup>Lawrence Berkeley National Laboratory, Berkeley, CA

Poster 300 **Use of Nucleic Acid Programmable Protein Array (NAPPA) to Study Autoantibodies in Alzheimer's Disease**; Yanyang Tang; Biodesign Institute, Arizona State University, Tempe, Arizona

Poster 301 **Analysis of Differentially Expressed Hippocampal Proteins: Predicted role of Nuclear HIST4H4 and HIST1H2BB protein in AD pathology**; Nikhat Ahmed Siddiqui; Barrett Hodgson University, Karachi, Pakistan

### PROTEOMICS IN DRUG DEVELOPMENT Posters 302 - 304

Poster 302 **A Recombinant Asp-Specific Protease for Bottom-up and Multi-Enzyme LC-MS/MS Workflows**; Chris Hosfield; Michael Rosenblatt; Marjeta Urh; Promega Corporation, Madison, WI

Poster 303 **Clarification of the Signaling Network Affected by the TNIK Inhibitor, NCB-0846, Using Reverse-phase Protein Array**; Mari Masuda<sup>1</sup>; Takaomi Inoue<sup>2</sup>; Yuko Uno<sup>2</sup>; Naoko Goto<sup>1</sup>; Masaaki Sawa<sup>2</sup>; Tesshi Yamada<sup>1</sup>; <sup>1</sup>National Cancer Center Research Institute, Tokyo, Japan; <sup>2</sup>Carna Biosciences Inc, Kobe, Japan

Poster 304 **Assessment of Awareness About Immunization Among Parents in Population**; Hira Sabir Malik; Bahria university, Karachi, Pakistan

### PROTEOMICS IN MICROBIOLOGY Posters 305 - 312

Poster 305 **Intracellular Tandem Mass Tag (TMT) Proteomic Analyses of HIV-1 Infected Macrophages after Cocaine and Sig1R Antagonist (BD1047) Treatments**; Omar Vélez López<sup>1</sup>; Loyda Meléndez<sup>2</sup>; Abiel Roche Lima<sup>2</sup>; Kelvin Carrasquillo Carrión<sup>2</sup>; Carla Salgado Ramírez<sup>2</sup>; Yadira Cantres Rosario<sup>2</sup>; Eraysy Machín Martínez<sup>2</sup>; Manuel Alvarez Ríos<sup>3</sup>; <sup>1</sup>University of Puerto Rico MS- Microbiology, San Juan, Puerto Rico; <sup>2</sup>University of Puerto Rico MS, San Juan, Puerto Rico; <sup>3</sup>University of Puerto Rico RP, San Juan, PR

Poster 306 **Development of Data-Independent MS Platform to Quantify Phenol Soluble**

## POSTERS

All posters displayed Monday - Wednesday. Odd-numbered posters present Monday.  
Even-numbered posters present Tuesday. All posters present Wednesday morning.

- Modulins Isoforms in Culture Media of *Staphylococcus aureus* from Bacteremia Patients;** Jiyoung Yu<sup>1</sup>; Eun Sil Kim<sup>2, 3</sup>; Yumi Oh<sup>1, 2</sup>; Hwangkyo Jeong<sup>1, 2</sup>; Jeonghun Yeom<sup>1</sup>; Yong Pil Chong<sup>3</sup>; Yang Soo Kim<sup>3</sup>; Kyung-Kon Kim<sup>1, 2</sup>; <sup>1</sup>Asan Medical Center, Seoul, South Korea; <sup>2</sup>University of Ulsan, College of Medicine, Seoul, South Korea; <sup>3</sup>Department of Infectious Diseases, Asan Medical Ce, Seoul, South Korea
- Poster 307 **Comparative Proteome Analysis for Korean Specific *Staphylococcus aureus*;** Yumi Oh<sup>1, 2</sup>; Eun Sil Kim<sup>2, 3</sup>; Jiyoung Yu<sup>1</sup>; Hwangkyo Jeong<sup>1, 2</sup>; Jeonghun Yeom<sup>1</sup>; Chong Yong Pil<sup>3</sup>; Yang Soo Kim<sup>3</sup>; Kyung-Kon Kim<sup>1</sup>; <sup>1</sup>Asan Medical Center, Seoul, South Korea; <sup>2</sup>University of Ulsan, College of Medicine, Seoul, South Korea; <sup>3</sup>Department of Infectious Diseases, Asan Medical Ce, Seoul, South Korea
- Poster 308 **Proteomic Analysis of Bacterial Peptide Products from Stress-Modified mRNAs Using a Synthetic Biology Approach;** Randi Turner; Daniel Dwyer; University of Maryland College Park, College Park, MD
- Poster 309 **Comparative Proteomic Profiling Reveals New Insight between Different Growth Phase of Biofilm Extractomes from *Staphylococcus aureus* Using TMT-based Quantitative MS;** Md Arifur Rahman<sup>1</sup>; Ardeshir Amirkhani<sup>2</sup>; Durdana Chowdhury<sup>1</sup>; Mark Molloy<sup>2</sup>; Dana Pascovici<sup>2</sup>; Maria Mempo<sup>1</sup>; Mark Baker<sup>1</sup>; Honghua Hu<sup>1</sup>; Karen Vickery<sup>1</sup>; <sup>1</sup>Macquarie University, Sydney, Australia; <sup>2</sup>Australian Proteome Analysis Facility, Sydney, Australia
- Poster 310 **Systematic Analysis Revealed a Subset of Heat Shock Response Genes are Required for Optimal Growth of *Halobacterium salinarum*;** Ming-Lung Ho<sup>1</sup>; Shen-Lin Chen<sup>1</sup>; Yu-Mei Hsieh<sup>1</sup>; Minzhen Luo<sup>1</sup>; Rueyhung R Weng<sup>2</sup>; Wailap V Ng<sup>1</sup>; <sup>1</sup>National Yang Ming University, Taipei, Taiwan; <sup>2</sup>National Taiwan University, Taipei, Taiwan
- Poster 311 **Probing Protein State in Bacteria by Thermal Proteome Profiling;** Andre Mateus; Jacob Bobonis; Nils Kurzawa; Frank Stein; Dominic Helm; Johannes Hevler; Athanasios Typas; Mikhail Savitski; EMBL, Heidelberg, Germany
- Poster 312 **Multi-Omics Analysis of a Nutrient Transport Protein Required for Full Virulence in *Campylobacter jejuni*;** Lok Man; Stuart Cordwell; The University of Sydney, Sydney, Australia
- Poster 315 **Efficient Plasma Sample Preparation for MS-Based Quantitative Profiling;** Sergei Snovida; Amarjeet Flora; Ryan Bomgarden; John Rogers; Thermo Fisher Scientific, Rockford, IL
- Poster 316 **Evaluation of timsTOF Pro in Multiplexed Workflows;** Henry Shwe; Joel Federspiel; Xinlei Sheng; Ileana Cristea; Tharan Srikumar; Princeton University, Princeton, NJ
- Poster 317 **Development of a Quantitative Proteomic Standard for Tandem Mass Tags (TMT);** Jae Choi<sup>1</sup>; Aaron Robitaille<sup>2</sup>; Tabiwang Arrey<sup>3</sup>; Rosa Viner<sup>2</sup>; Andreas Huhmer<sup>2</sup>; John Rogers<sup>1</sup>; <sup>1</sup>Thermo Fisher Scientific, Rockford, IL; <sup>2</sup>Thermo Fisher Scientific, San Jose, CA; <sup>3</sup>Thermo Fisher Scientific, Bremen, Germany
- Poster 318 **Method Development for Quantification of Vitamin D-Binding Protein in Prenatal Serum using LC-MRM;** Lisa Kilpatrick<sup>1</sup>; Ashley Boggs<sup>2</sup>; Stephen Long<sup>2</sup>; Karen Phinney<sup>1</sup>; <sup>1</sup>NIST, Gaithersburg, MD; <sup>2</sup>NIST, Charleston, SC
- Poster 319 **Highly Reproducible and Accurate Label Free Quantification Using the PASEF Method on a TIMS-QTOF Mass Spectrometer;** Heiner Koch<sup>1</sup>; Gary Kruppa<sup>2</sup>; Scarlet Koch<sup>1</sup>; Thomas Kosinski<sup>1</sup>; Markus Lubeck<sup>1</sup>; Florian Meier<sup>3</sup>; Andreas Brunner<sup>3</sup>; Matthias Mann<sup>3</sup>; <sup>1</sup>Bruker Daltonik GmbH, Bremen, Germany; <sup>2</sup>Bruker Daltonics Inc., Billerica, US; <sup>3</sup>Max Planck Institute of Biochemistry, Martinsried, Germany
- Poster 320 **Landscape of Deubiquitinating Enzymes (DUBs) in KRAS Mutants Using Activity Based Protein Profiling (ABPP) and Bioinformatics Analysis;** Emma Adhikari; Moffitt Cancer Center, Tampa, <Not Specified>
- Poster 321 **In-Depth Secretome Analysis in stage-Specific Colon Cancer Cell Lines;** Jeyalakshmi Kandhavelu<sup>1</sup>; Stoyan Stoychev<sup>2</sup>; Kumar Subramanian<sup>1</sup>; Amber Khan<sup>1</sup>; Paul Ruff<sup>1</sup>; Clement Penny<sup>1</sup>; <sup>1</sup>University of the Wits, Johannesburg, South Africa; <sup>2</sup>Council for Scientific and Industrial Research, Pretoria, South Africa
- Poster 322 **Fully Validated SRM-MS-Based Method for Absolute Quantification of PIVKA-II in Human Serum: Clinical Applications for Patients with HCC;** Areum Sohn<sup>1</sup>; Hyunsoo Kim<sup>1</sup>; Injoon Yeo<sup>2</sup>; Yoseop Kim<sup>2</sup>; Minsoo Son<sup>2</sup>; Su Jong Yu<sup>3</sup>; Jung-Hwan Yoon<sup>3</sup>; Youngsoo Kim<sup>1, 2</sup>; <sup>1</sup>Seoul National University College of Medicine, Seoul, South Korea; <sup>2</sup>Seoul National University, Biomedical Engineering, Seoul, South Korea; <sup>3</sup>Medical Research Center, Liver Research Institute, Seoul, South Korea
- Poster 323 **Clinical Assay for AFP-L3 Using Multiple Reaction Monitoring-Mass Spectrometry for**

### QUANTITATIVE PROTEOMICS Posters 313 - 337

- Poster 313 **Automated TMT Labeling Using Solid Phase Micro Extraction Cartridges;** Greg Foster; Aaron Robitaille; Daniel Lopez-Ferrer; Thermo Fisher Scientific, San Jose, CA
- Poster 314 **Standardization of Sample Preparation for Proteomics Applications;** Aaron Robitaille<sup>1</sup>;

## POSTERS

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- Diagnosing Hepatocellular Carcinoma.;** Hyunsoo Kim<sup>1</sup>; Areum Sohn<sup>1</sup>; Injun Yeo<sup>2</sup>; Su Jong Yu<sup>3</sup>; Jung-Hwan Yoon<sup>3</sup>; Youngsoo Kim<sup>1, 2</sup>; <sup>1</sup>*Seoul National University College of Medicine, Seoul, South Korea*; <sup>2</sup>*Department of Biomedical Engineering, Seoul, Republic of Korea*; <sup>3</sup>*Department of Internal Medicine, Seoul, Republic of Korea*
- Poster 324 **A novel UHPLC-MRM-MS methodology for reproducible and fast quantification of histone PTMs;** Joseph Cesare; Simone Sidoli; Zuo-Fei Yuan; Hyoungjoo Lee; Benjamin Garcia; *University of Pennsylvania, Philadelphia, PA*
- Poster 325 **Development of a Multiplexed Assay for Oral Cancer Candidate Biomarkers Using Peptide Immunoaffinity Enrichment and Targeted Mass Spectrometry;** Yung-Chin Hsiao<sup>1</sup>; Lang-Ming Chi<sup>3</sup>; Kun-Yi Chien<sup>1</sup>; Wei-Fan Chiang<sup>2</sup>; Kai-Ping Chang<sup>3</sup>; Jau-Song Yu<sup>1</sup>; <sup>1</sup>*Chang Gung University, Tao-Yuan, Taiwan*; <sup>2</sup>*Chi-Mei Medical Center, Liouying, Taiwan*; <sup>3</sup>*Chang Gung Memorial Hospital, Taoyuan, Taiwan*
- Poster 326 **Proteomic Comparison of 11 Human Tissues using 11-plex Tandem Mass Tags and Synchronous Precursor Selection MS3 Analysis.;** Anna M. Vildhede<sup>2</sup>; Chuong Nguyen<sup>1</sup>; Emi Kimoto<sup>1</sup>; A. David Rodrigues<sup>1</sup>; Manthana V. Varma<sup>1</sup>; Robert A. Everley<sup>1</sup>; <sup>1</sup>*Pfizer R&D, Groton, CT*; <sup>2</sup>*AstraZeneca, Gothenburg, Sweden*
- Poster 327 **Targeted Proteomics Applied to the Research in Public Health: Analysis of Energy Metabolism of Isolated Monocytes from Sepsis Patients Plasma;** Pedro Mendes Azambuja Rodrigues<sup>1</sup>; Monique Ramos Oliveira Trugilho<sup>2, 3</sup>; Gabriel Reis Alves Carneiro<sup>4, 5</sup>; Fabio Cesar Souza Nogueira<sup>4, 5</sup>; Gilberto Barbosa Domont<sup>5</sup>; Richard Hemmi Valente<sup>2</sup>; Fernando Augusto Bozza<sup>1</sup>; Giselle Villa Flor Brunoro<sup>2</sup>; <sup>1</sup>*Evandro Chagas National Infectology Institute, Rio de Janeiro, Brazil*; <sup>2</sup>*Laboratory of Toxinology, Oswaldo Cruz Institute, Rio de Janeiro, Brazil*; <sup>3</sup>*Center of Technological Development in Health, Rio de Janeiro, Brazil*; <sup>4</sup>*Laboratory of Proteomics, UFRJ, Rio de Janeiro, Brazil*; <sup>5</sup>*Laboratory of Protein Chemistry, UFRJ, Rio de Janeiro, Brazil*
- Poster 328 **Discrepancy in Stably Expressed Proteins Within and Across Human Tissue Types;** Christine Wegler<sup>1, 2</sup>; Magnus Ölander<sup>1</sup>; Per Artursson<sup>1</sup>; <sup>1</sup>*Uppsala University, Uppsala, Sweden*; <sup>2</sup>*DMPK, AstraZeneca Gothenburg, Gothenburg, Sweden*
- Poster 329 **Analysis of Grape Berry Proteome to Determine their Relative Abundance During Berry Development and Ripening;** Amber Deets<sup>1, 2</sup>; Ramesh Katam<sup>1</sup>; <sup>1</sup>*Florida A&M University, Tallahassee, FL*; <sup>2</sup>*Stetson University, Deland, FL*
- Poster 330 **Rapid Qualitative and Absolute Quantification of Plasma based proteins using a Novel Scanning Quadrupole DIA Acquisition Method;** Christopher Hughes<sup>1</sup>; Lee Gethings<sup>1</sup>; Florian Marty<sup>2</sup>; Sebastian Müller<sup>2</sup>; Jose Castro-Perez<sup>3</sup>; Robert Plumb<sup>3</sup>; <sup>1</sup>*Waters Corporation, Wilmslow, United Kingdom*; <sup>2</sup>*Biognosys AG, Schlieren, Switzerland*; <sup>3</sup>*Waters Corporation, Milford, MA*
- Poster 331 **Absolute Quantification of Apolipoproteins Using a High Precision QPrEST-Based SRM Assay;** Andreas Hober<sup>1, 2</sup>; Jonas Malmqvist<sup>3</sup>; Maria Ryaboshapkina<sup>3</sup>; Björn Forsström<sup>1, 2</sup>; Mathias Uhlen<sup>1, 2</sup>; Fredrik Edfors<sup>1, 4</sup>; Tasso Milliotis<sup>3</sup>; <sup>1</sup>*Science for Life Laboratory, KTH, Solna, Sweden*; <sup>2</sup>*Department of Protein Science, KTH, Stockholm, Sweden*; <sup>3</sup>*AstraZeneca, Mölndal, Sweden*; <sup>4</sup>*Stanford, Department of Genetics, Stanford, CA*
- Poster 332 **The Proteomic and Biochemical Studies demonstrate that 300-Mediated Lysine 2-Hydroxyisobutyrylation Regulates Glycolysis;** He Huang<sup>1</sup>; Shuang Tang<sup>2</sup>; Ming Ji<sup>2</sup>; Xiaojing Liu<sup>3</sup>; Jason W. Locasale<sup>3</sup>; Xiaoling Li<sup>2</sup>; Yingming Zhao<sup>1</sup>; <sup>1</sup>*The University of Chicago, Chicago, <Not Specified>*; <sup>2</sup>*National Institute of Environmental Health Science, NC, 27709*; <sup>3</sup>*Duke University School of Medicine, Durham, NC*
- Poster 333 **Global Proteomic Profiling of Dehydration-Modulated Mitochondrial Dynamics and Defense Response in Rice;** Dipak Gayen; Pragya Barua; Nilesh Vikram Lande; Subhra Chakraborty; Niranjana Chakraborty; *NIPGR, New Delhi, India*
- Poster 334 **IonStar Enables High-Precision, Low-Missing-Data Proteomics Quantification in Large Biological Cohorts;** Jun Qu; *SUNY-Buffalo, Buffalo, NY*
- Poster 335 **Proteome-Wide Analysis of the NSm Protein Effect in Primary Macrophages Following Rift Valley Fever Virus Infection;** Magali Boissiere<sup>2</sup>; Carole Tamietti<sup>2</sup>; Dominique Simon<sup>3</sup>; Magalie Duchateau<sup>1</sup>; Natalia Pietrosevoli<sup>4</sup>; Quentin Gai Gianetto<sup>1, 4</sup>; Véronique Hourdel<sup>1</sup>; Félix Kreher<sup>5</sup>; Jean-Jacques Panthier<sup>3</sup>; Felix Rey<sup>2</sup>; Marie Flamand<sup>2</sup>; Mariette Matondo<sup>1</sup>; <sup>1</sup>*Proteomics Platforms, IP, MSBio unit, CNRS USR 2000, Paris, France*; <sup>2</sup>*Structural Virology, Institut Pasteur Paris, Paris, France*; <sup>3</sup>*Mouse Functional Genetics, Institut Pasteur Paris, Paris, France*; <sup>4</sup>*Bioinformatics and Biostatistics Hub, C3BI, USR 37, Paris, France*; <sup>5</sup>*Institute of Infection, Immunity and Inflammation, Glasgow, Scotland*
- Poster 336 **Accurate, Sensitive, and Precise Multiplexed Proteomics Using the Complement Reporter Ion Cluster;** Matthew Sonnett; Eyan Yeung; Martin Wühr; *Princeton University, Princeton, NJ*

## POSTERS

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Poster 337 **Quantitative Proteomics Uncovers a Novel USP9X substrate TTK for Tumorigenesis in Non-Small Cell Lung cancer;** Hu Zhou<sup>1</sup>; Xiangling Chen<sup>1</sup>; Chengli Yu<sup>1</sup>; Jing Gao<sup>1</sup>; Hongwen Zhu<sup>1</sup>; Han He<sup>1</sup>; Ruimin Huang<sup>1</sup>; Hua Xie<sup>1</sup>; Daming Gao<sup>2</sup>; <sup>1</sup>*Shanghai Institute of Materia Medica, Chinese Acad, Shanghai, China*; <sup>2</sup>*Institute of Biochemistry and Cell Biology, CAS, Shanghai, China*

### SIGNALING AND BIOCHEMICAL PATHWAY PROTEOMICS Posters 338 - 349

Poster 338 **Applications of SureQuant™ Pathway Panels for Quantitative Analysis of Cancer Signaling Proteins;** Penny Jensen<sup>1</sup>; Bhavin Patel<sup>1</sup>; Leigh Foster<sup>1</sup>; Renuka Sabinis<sup>1</sup>; Aaron Gajadhar<sup>1</sup>; Jonathan R. Krieger<sup>2</sup>; Jiefei Tong<sup>2</sup>; Michael F. Moran<sup>3</sup>; Rosa Viner<sup>4</sup>; Andreas Huhmer<sup>4</sup>; Kay Opperman<sup>1</sup>; John Rogers<sup>1</sup>; <sup>1</sup>*Thermo Fisher Scientific, Rockford, IL*; <sup>2</sup>*The Hospital for Sick Children, Toronto, Canada*; <sup>3</sup>*University of Toronto, Toronto, Canada*; <sup>4</sup>*Thermo Fisher Scientific, San Jose, CA*

Poster 339 **Targeted Mass Spectrometry Assay Kits for Absolute Quantitation of Signaling Pathway Proteins;** Bhavin Patel<sup>1</sup>; Penny Jensen<sup>1</sup>; Leigh Foster<sup>1</sup>; Renuka Sabnis<sup>1</sup>; Abid Haseeb<sup>1</sup>; Aaron Gajadhar<sup>2</sup>; Rosa Viner<sup>2</sup>; Sebastien Gallien<sup>3</sup>; Andreas Huhmer<sup>2</sup>; Kay Opperman<sup>1</sup>; John Rogers<sup>1</sup>; <sup>1</sup>*Thermo Fisher Scientific, Rockford, IL*; <sup>2</sup>*Thermo Fisher Scientific, San Jose, CA*; <sup>3</sup>*Thermo Fisher Scientific, PMSC, Cambridge, MA*

Poster 340 **Systems Toxicology Assessment of Potential Modified Risk Tobacco Products: Effects on Lung, Liver, and Heart in ApoE-/- Mice Using iTRAQ;** Catherine Nury<sup>1</sup>; Blaine Phillips<sup>2</sup>; Justyna Szostak<sup>1</sup>; Bjoern Titz<sup>1</sup>; Walter Schlage<sup>3</sup>; Emmanuel Guedj<sup>1</sup>; Patrice Leroy<sup>1</sup>; Gregory Vuillaume<sup>1</sup>; Florian Martin<sup>1</sup>; Ansgar Buettner<sup>4</sup>; Ashraf Elamin<sup>1</sup>; Alain Sewer<sup>1</sup>; Nicolas Sierro<sup>1</sup>; Mohamed Amin Choukrallah<sup>1</sup>; Thomas Schneider<sup>1</sup>; Nikolai Ivanov<sup>1</sup>; Patrick Vanscheeuwijck<sup>1</sup>; Manuel Peitsch<sup>1</sup>; Julia Hoeng<sup>1</sup>; <sup>1</sup>*PMI R&D, Philip Morris Products S.A., Neuchatel, Switzerland*; <sup>2</sup>*PMI R&D, Philip Morris International Research Lab, Singapore, Singapore*; <sup>3</sup>*Biology Consultant, Bergisch Gladbach, Germany*; <sup>4</sup>*Histovia GmbH, Overath, Germany*

Poster 341 **High Throughput Signaling Pathway Analysis Using Multiplex-Immunoprecipitation and Fast LC-PRM;** Sebastien Gallien<sup>1,2</sup>; Aaron Gajadhar<sup>3</sup>; Bhavin Patel<sup>4</sup>; Tabiwang Arrey<sup>5</sup>; Dave Sarracino<sup>2</sup>; Sarah Trusiak<sup>2</sup>; Yue Xuan<sup>2,5</sup>; Emily Chen<sup>2</sup>; <sup>1</sup>*Thermo Fisher Scientific, Paris, France*; <sup>2</sup>*Thermo Fisher Scientific, PMSC, Cambridge, MA*; <sup>3</sup>*Thermo Fisher Scientific, San Jose, CA*; <sup>4</sup>*Thermo Fisher Scientific, Rockford, IL*; <sup>5</sup>*Thermo Fisher Scientific, Bremen, Germany*

Poster 342 **Proteomic Analysis with Inguinal White Adipose Tissue of CXCL5 KO Mice Revealed Increased Energy Consumption Activity;** Dong Wook Kim; Da Bin Lee; Je-Yoel Cho; , *Seoul, South Korea*

Poster 343 **Profiling the Kinome and Phosphoproteome of mutant KRAS-driven Pancreatic Ductal Adenocarcinoma;** Lee Graves<sup>1</sup>; Laura Herring<sup>1</sup>; Thomas Gilbert<sup>1</sup>; Nely Dicheva<sup>1</sup>; Emily Werth<sup>1</sup>; Emily Wilkerson<sup>1</sup>; Angelina Vaseva<sup>2</sup>; Kirsten Bryant<sup>1</sup>; Devon Blake<sup>1</sup>; Nathaniel Diehl<sup>1</sup>; Naim Rashid<sup>1</sup>; Channing Der<sup>1</sup>; <sup>1</sup>*University of North Carolina at Chapel Hill, Chapel Hill, NC*; <sup>2</sup>*University of Texas, San Antonio, TX*

Poster 344 **Global Proteome Landscape During Antigen Dependent Differentiation Reveals Dynamic Cell Signaling Profiles Across Distinct B-Cells Subpopulations;** Manuel Fuentes; *University of Salamanca, Salamanca, Spain*

Poster 345 **Multi-Faceted Chemical and Genetic Knock-Down Approaches to Mapping Functional Protein Networks of DNA Methyltransferase I (DNMT1);** Rob Ewing; Emily Bowler; Paul Skipp; *University of Southampton, Southampton, United Kingdom*

Poster 346 **The Functional Role of Mitochondrial Sirtuin Signalling Underlying Acute Angiotensin II-Mediated Oxidative Stress in Mouse Ophthalmic Artery;** Natarajan Perumal; Lars Straßburger; Adrian Gericke; Franz Grus; Norbert Pfeiffer; Caroline Manicam; *University Medical Centre Mainz, Mainz, Germany*

Poster 347 **Assessment of MFG-E8 Protein Using Label Free and iTRAQ Proteomics for an Intrinsic Component of the Cell Growth Regulator;** Syed Azmal Ali; *ICAR-NDRI, Karnal, India*

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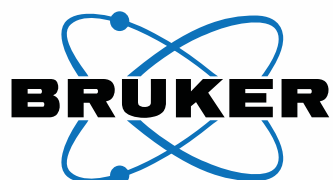


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