Student Travel Award Winners

Sergi Clotet, Canada
- **MP09-001**: Silac-Based Proteomics of Human Kidney Cells Reveals a Novel Link between Male Sex Hormones and Impaired Energy Metabolism in Diabetic Kidney Disease

Victoria Dardov, United States
- **WO08-004**: Proteomic Analysis of Motor Neurons from Induced Pluripotent Stem Cells: ALS

Fei Fang, China
- **MO02-004**: All MS/MS Ions Monitoring Acquired by Data-Dependent Acquisition without Dynamic Exclusion: A New Concept for In-Depth Protein Quantification

Humberto Gonczarowska-Jorge, Germany
- **MO08-003**: Subtilisin for Large Scale (Phospho) Proteomics – The Beginning of a Wonderful Love Story?

Honggang Huang, Denmark
- **MP09-002**: Quantitative Simultaneous Multiple Ptmomics Characterization of Arteries from Patients with Atherosclerosis and Type 2 Diabetes

Jiangming Huang, China
- **TP02-012**: Simplified Cell Strategy for Large Scale Identification of Mucin-type O-glycoproteins

Emila Kurbasic, Sweden
- **TP01-041**: Changes in Protein Expression between Primary Breast Tumour and Lymph Node or Distant Metastases

Hiromi Koh, Singapore
- **MP06-025**: EBprotV2: Statistical Analysis of Labeling-based Quantitative Proteomics Data with Applications to Clinical Data

Hiroshi Kusamoto, Japan
- **WP03-003**: Thio-tag Tip Method by Using zinc (II)–cyclen-attached Agarose Beads for Enrichment of Cysteine-Containing Biomolecules

Chongyang Li, Canada
· **TP03-004**: PIAS1-mediated SUMOylation of BAF57 Is a Critical Regulator of Cell Growth and Drug Sensitivity in Ovarian Cancer Cells

Yang Li, China
· **TO06-003: A High-Content Functional Mycobacterium Tuberculosis Proteome Microarray and Its Applications**

Parul Mittal, Australia
· **WP16-003**: Lymph Node Metastasis of Primary Endometrial Cancers: Associated Proteins Revealed by Maldi Imaging

Dan Bi Park, South Korea
· **TP02-026**: Glycomic Approach for Design of Humanized Mouse Model via Nano-LC/MS and LC/MS/MS

Gun Wook Park, South Korea
· **TP02-025**: Integrated GlycoProteome Analyzer (I-GPA) for Automatic Identification and Quantitation of Site-Specific N-Glycosylation in Human Plasma

Cristian Piras, Italy
· **TO10-004**: Adipose Tissue Pathways in Obesity: Iberian Pig As Large Animal Model of Metabolic Disorders

Priya Sivadasan, India
· **TP01-073**: Salivary Proteins from Pre-Malignant and Malignant Lesions of the Oral Cavity and Their Translational Potential for Early Diagnosis

Zhiduan Su, Australia
· **WO03-004**: Proteomic and Redox Proteomic Analyses Reveal a Dual ROS-regulation of Glucose Uptake in Adipocytes

Chisato Takahashi, Japan
· **MP05-020: Phosphoproteomics-based Prediction of Cellular Protein Kinome Profiles**

Mathias Walzer, Germany
· **TO10-003: The HUPO-PSI Quality Control Working Group: Making QC More Accessible for Biological Mass Spectrometry**

Xin Wang, Japan

· **WP12-010**: **Proteomic Analysis to Reveal the Calcium Function on Protein Glycosylation in Endoplasmic Reticulum of Soybean under Flooding and Drought Stresses**

Churat Weeraphan, Thailand

· **TP01-088**: **Phosphoproteome Profiling of Isogenic Cholangiocarcinoma Exosomes Reveal Differential Expression of a Key Metastatic Factor**

Ting Wu, Australia

· **MP07-022: Quantitative Shifts in the Influenza Immunopeptidome Reveal the Relative Contributions of Direct and Cross-Presentation to T cell Mediated Immunity**

Jeonghun Yeom, South Korea

· **WP01-011: Identification for Protein-level Evidence of Genomic Variants in Cancer Cells Using New Proteogenomic Approach**

Kun-Hsing Yu, United States

· **TP01-096**: **Predicting Ovarian Cancer Patients' Clinical Response to Platinum-based Chemotherapy by Their Tumor Proteomic Signatures**

Bo Zhang, Sweden

· **MP06-051**: **Diffacto: A Robust and Accurate Quantification-Centered Proteomics Method for Large-scale Differential Analysis**

Yinghua Zhao, China

· **WP15-016: Application of High Throughput Urinary Proteomic Strategy in the Diagnosis of Acute Appendicitis with Confusable Acute Abdomens**