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