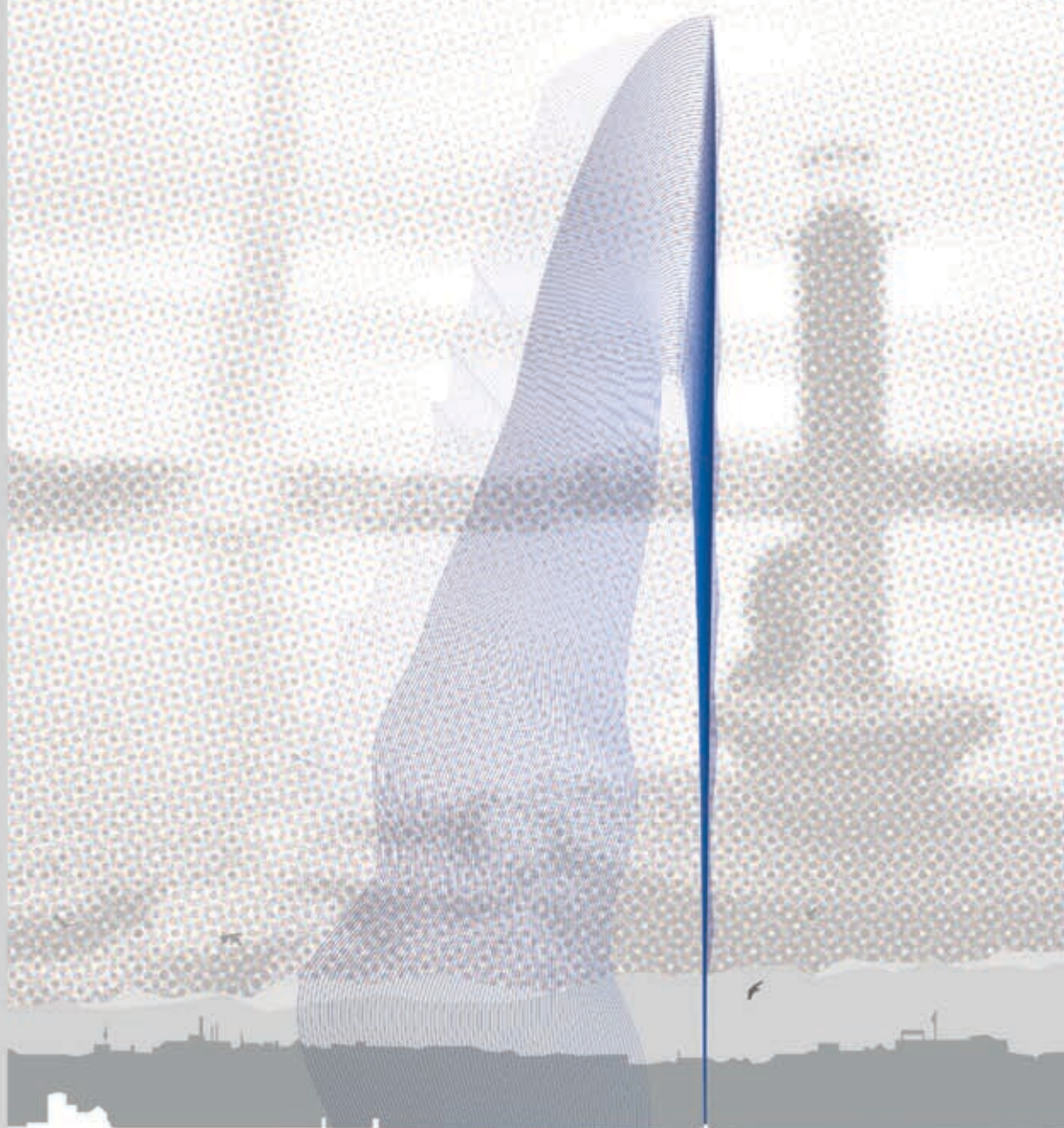


Final Programme

Translational Proteomics





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Acknowledgements

The HUP0 2011 Organising Committee would like to thank the following companies for their active participation and support:

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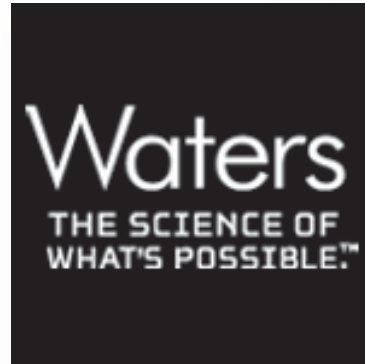


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Welcome message

Dear Colleagues,

On behalf of the Organizing committee, we are delighted to welcome you to Geneva, Switzerland, for the combined HUP0 10th Annual World Congress, 5th EuPA Annual Scientific Meeting and the 8th SPS scientific meeting.

The Scientific Program will focus on Translational Proteomics. We are confident that you will enjoy the Scientific Program of this International World Congress, in this small, yet international and sophisticated place; Geneva.

Just enjoy it!

Yours sincerely,

Jean- Charles Sanchez



Denis Hochstrasser



Luca Bini



Committees

CONGRESS CO-CHAIRS AND LOC MEMBERS

JEAN-CHARLES SANCHEZ

Department of Human Protein Sciences, Geneva University, Switzerland

DENIS HOCHSTRASSER

Department of Genetics and Laboratory Medicine & Department of Human Protein Sciences, University Hospitals & Geneva University, Switzerland

LUCA BINI

Department of Biotechnologies, Siena University, Italy

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Alex Scherl

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Natacha Turck

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CHAIRMEN OF HUPO EDUCATION DAY, HUPO CLINICAL DAY, EUPA DAY, HUPO & EUPA AWARDS AND HUPO & EUPA INITIATIVES

CLINICAL DAY CHAIRS: Pierre Burkhard & Daniel Chan

EDUCATION DAY CHAIRS: Thierry Rabilloud & Peter James

EUPA DAY CHAIRS: Jesus Jorrin Novo & Jean-Jacques Diaz

HUPO & EUPA AWARDS CHAIRS: Concha Gil & Maxey Chung

HUPO & EUPA INITIATIVES CHAIRS: Juan-Pablo Albar & Gil Omenn

Human Proteome Organisation

WHAT IS HUP0?

The Human Proteome Organisation (HUP0) is an international scientific organization representing and promoting proteomics through international cooperation and collaborations by fostering the development of new technologies, techniques and training.

HUP0 EXECUTIVE COMMITTEE

	2011 Makeup
President	Catherine Costello (2011-2012)
Past President	Young-Ki Paik (2011-2012)
Senior Vice President	Pierre Legrain (2011-2012)
Vice President	Bill Hancock (2011-2012)
Treasurer	Roman Zubarev (2011-2013)
Secretary General	Richard Simpson (2011-2012)
Member at Large	Denis Hochstrasser (2010-2011)
Member at Large	Gil Omenn (2011-2012)

HUP0 COUNCIL (BOARD OF DIRECTORS)

The HUP0 Council is the decision-making body. The Council is elected by the HUP0 members.

European Proteomics Association (EuPA)

EuPA

The European Proteomics Association (EuPA) is the federation of European national proteomics societies. It has been established to coordinate and integrate national initiatives within the proteomics field. The main objectives are to strengthen and promote fundamental research and applications, as well as education and training in all areas of proteomics throughout Europe. EuPA is a non-profit organisation.

CURRENT EXECUTIVE COMMITTEE

PRESIDENT: Gyorgy Marko-Varga

VICE-PRESIDENT: Jean-Charles Sanchez

CONFERENCE & COMMUNICATION COMMITTEE: Concha Gil

EDUCATION COMMITTEE: Garry Corthals

Eu-HUPO COMMITTEE: Thierry Rabilloud

FUNDING COMMITTEE: Guenter Thesseling

CURRENT EUPA COUNTRY MEMBERS

Austria, Czech Republic, Denmark, Finland, France, Germany, Greece, Hungary, Italy, Netherlands, Norway, Portugal, Russia, Spain, Sweden, Switzerland and UK

Swiss Proteomics Society (SPS)

The Swiss Proteomics Society (SPS) is a scientific society established for the public benefit to advance research, development and education in the proteomic sciences created in 2000. The aim of the SPS is to stimulate and coordinate proteomic activities in Switzerland as well as in other countries, with two major objectives:

- To hold and arrange courses and meetings on matters connected to proteomics;
- To promote the diffusion and exchange of information among people interested in proteomics.

SPS EXECUTIVE COMMITTEE

PRESIDENT: Dr Hans Voshol

CURRENT MEMBERS (2010-2013)

Dr Michael Affolter, Nestlé Research Centre, Lausanne

Dr Bruno Domon, Institute of Molecular Systems Biology, Zurich

Dr Niels Lion, Laboratoire d'Electrochimie Physique et Analytique, Lausanne

Dr Markus Muller, Swiss Institute of Bioinformatics, Geneva

Dr Alexander Scherl, Proteomics Core Facility, University of Geneva, Geneva

Dr Alexander Schmidt, Proteomics Core Facility, Biozentrum, University of Basel, Basel

Dr Hans Voshol, Novartis Institutes for Biomedical Research, Basel

Programme overview

Sunday 4th September 2011

	HALL A	HALL B	HALL C	ROOM K	ROOM L
09:00 - 10:30	<p>TRANSLATION OF PROTEOMICS DISCOVERY INTO CLINICAL PRACTICE: CRITICAL ISSUES</p> <p>CLINICAL DAY 1</p> <p>Chairs: Pierre Burkhard & Daniel Chan Speakers: Daniel W. Chan, Denis Hochstrasser & David Ransohoff</p>	<p>UP-DATE ON PROTEOMICS TOOLS</p> <p>EDUCATION DAY 1</p> <p>Chairs: Thierry Rabilloud & Peter James Speaker: Kathryn Lilley, Thierry Rabilloud</p>	<p>MICROORGANISM, PLANT, AND ANIMAL PROTEOMICS</p> <p>EUPA DAY 1</p> <p>Chairs: Jesus Jorriñ Novo & Jean-Jacques Diaz Speakers: Christian H. Ahrens, Peter Jungblut, Concha Gil</p>	<p>SYMPOSIUM ON NEW TECHNOLOGIES AND STANDARDIZATION IN PROTEOMICS</p> <p>Chairs: Bruno Dornon & Christoph Borchers</p> <p>Speakers: Susan Abbatiello, Roumen Bogoev, Eric Deutsch, Will Dracup, Chris Kinsinger, James Langridge, Alexander Makarov, Eugene Nikolaev, Richard Smith, Keith Waddell</p>	<p>HPP CONSORTIUM</p>
10:30 - 11:00	Coffee break				
11:00 - 12:30	<p>CLINICAL DAY 2</p> <p>Chairs: Pierre Burkhard & Daniel Chan Speakers: Steve Binder, Steve Skates, Ruedi Aebersold</p>	<p>EDUCATION DAY 2</p> <p>Chairs: Thierry Rabilloud & Peter James Speakers: Loic Dayon, Mike Mac Coss</p>	<p>EUPA DAY 2</p> <p>Chairs: Jesus Jorriñ Novo & Jean-Jacques Diaz Speakers: Stefanie Wienkoop, Dominique Job, Albert Heck, Fernando Corrales</p>		
12:30 - 14:00	Lunch				
13:30 - 16:30	<p>CLINICAL DAY 3</p> <p>Chairs: Pierre Burkhard & Daniel Chan Speakers: Burcu Ayoglu, Laura Chiecchi Natalia Tiberti, Xavier Robin</p>	<p>EDUCATION DAY 3</p> <p>Chairs: Thierry Rabilloud & Peter James Speaker: Susan Abbatiello, Pierre-Alain Binz, Tine Thingholm</p>	<p>EUPA DAY 3</p> <p>Chairs: Jesus Jorriñ Novo & Jean-Jacques Diaz Speakers: Niranjana Chakraborty, Adelina Rogowska-Wrzzesinska, Pao-Chi Liao, Amandine Boeuf, Jianke Li, Brian Tate Weinert</p>		<p>HUPO COUNCIL</p>
16:30 - 17:00	Coffee break				
17:00 - 17:30	OPENING CEREMONY				
17:30 - 19:00	<p>PLENERARY LECTURE 1: Marc Abrahams PLENERARY LECTURE 2: Michael Sternberg</p>				<p>EXHIBITION</p>
19:00 - 22:00					<p>WELCOME COCKTAIL</p>

Programme overview

Monday 5 th September 2011

	HALL A	HALL B	HALL C	ROOM K	ROOM E	ROOM F	ROOM G	ROOM L
7:00 - 8:30				HUMAN LIVER PROTEOME	C-HPP			
8:30 - 9:15	PLENERARY LECTURE 3 : Stylianos Antonarakis							
9:15 - 10:00	PLENERARY LECTURE 4: Ueli Schibler							
10:00 - 10:30	Coffee break (10:00 -10:30)							
10:30 - 12:30	S1 INFECTIOUS DISEASES Invited speaker S1: Jaques Schrenzel (30') Johan Malmström: (20') Jean-Philippe Charrier: (20') Marianne Danielsen: (20') Michael Snyder: (20')	S2 ORGANELLAR PROTEOMICS Invited speaker S2: Angus Lamond (30') Emma Lundberg: (20') Jens S Andersen: (20') Christine Carapito: (20') Domitille Schwartz: (20')	S3 MS QUANTITATION Invited speaker S3: Ruedi Aebersold (30') Leigh Anderson: (20') Matthias Selbach: (20') Ivo Fierro-Monti: (20') H.Christian Eberl: (20')	EXHIBITION	BRAIN PROTEOME PROJECT			
12:30 - 13:15	Lunch							
13:15 - 14:00	EUPA Present and Future			POSTER 1 (ODD #)	CELL BIOSCIENCES	BIORAD	BECKMAN COULTER	AB SCIEX
14:30 - 15:00	Coffee break (14:30-15:00)							
15:00 - 17:00	S4 ONCOLOGY Invited speaker S4: Giulio Superti Furga (30') Rodrigo Banderas: (20') Silvia Surinova: (20') Yue Fan: (20') Tamar Geiger: (20')	S5 POST-TRANSLATIONAL MODIFICATIONS Invited speaker S5: Ole Jensen (30') Marc Wilkins: (20') Petra Beli: (20') Christian Toft Madsen: (20') Al Burlingame: (20')	S6 PROTEIN ARRAYS Invited speaker S6: Dolores Cahit (30') Jochen M Schwenk: (20') Peter Nilsson: (20') Jörg D. Hoheisel: (20') Christer Wingren: (20')	EXHIBITION			PROTEOMIC STANDARDS INITIATIVE	
17:00 - 18:30	Human Glycomics / Proteome Initiative	HUPO Cardiovascular Initiative	Proteome Biology of Stem Cells Initiative	AGILENT	GE HEALTHCARE	AB SCIEX	PROTEOMICS MASTER	ETD
19:00 - 22:00	Speaker Dinner (upon invitation only)							

Programme overview

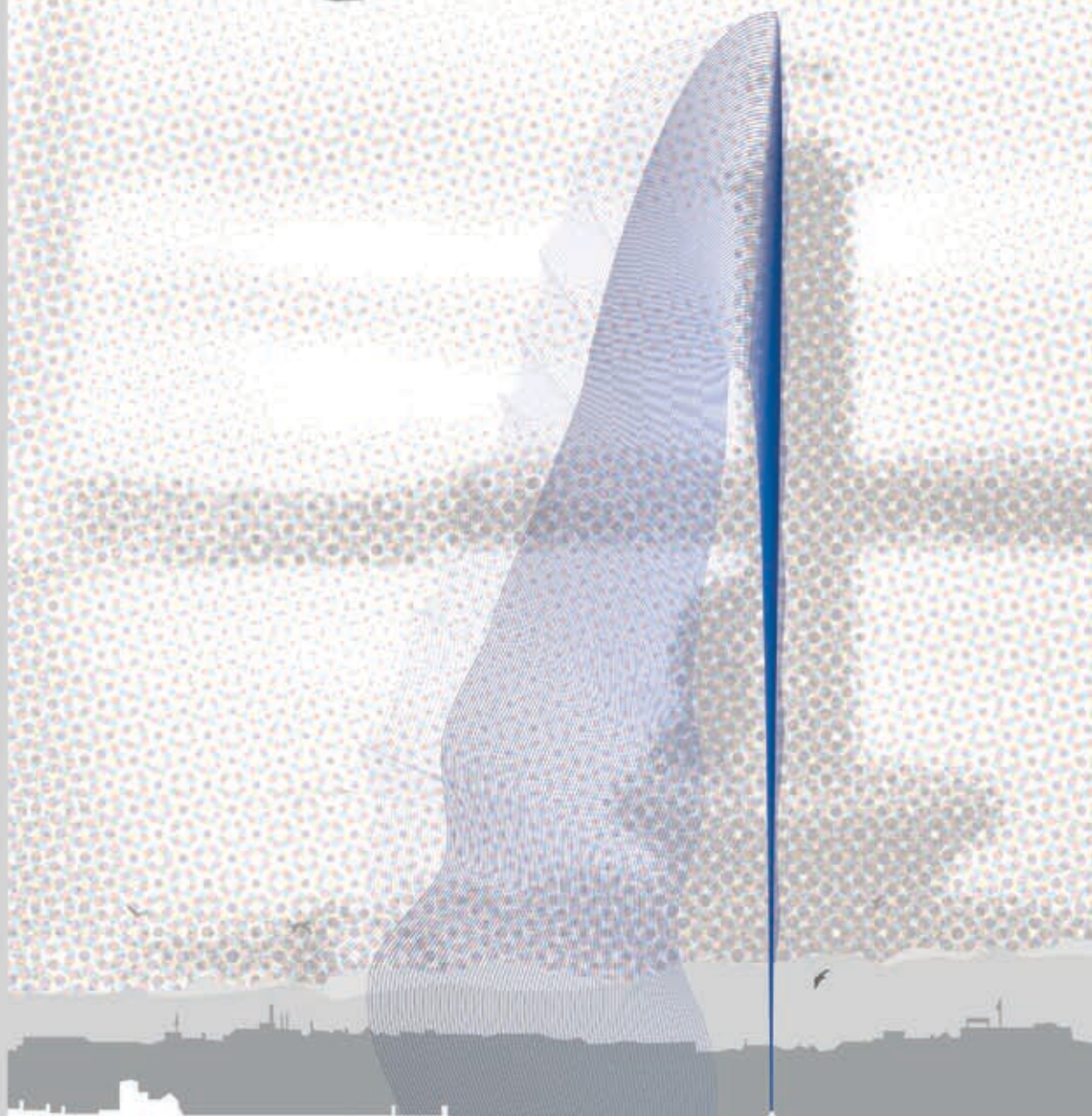
Tuesday 6 th September 2011

	HALL A	HALL B	HALL C	ROOM K	ROOM E	ROOM F	ROOM L
7:00 - 8:30				KIDNEY AND URINE INITIATIVE		AB SCIEX	
8:30 - 9:15	PLENERAY LECTURES 5: Martin Schwab						
9:15 - 10:00	PLENERAY LECTURES 6: Amos Bairoch						
10:00 - 10:30	Coffee break (10:00 - 10:30)						
10:30 - 12:30	S7 CNS Invited speaker S7: Joan Montaner (30') Uwe Schulte: (20') Angela Bachi: (20') Thomas Bock: (20') Ali Fathi: (20')	S8 MEMBRANES PROTEINS Invited speaker S8: Jerome Garin (30') Arie Admon: (20') Nobuaki Takemori: (20') Bernd Wollscheid: (20') Rebekah L. Gundry: (20')	S9 DATA AND KNOWLEDGE Chair S9: Amos Bairoch Nicki Packer: (20') Ruth Hüttenhain: (20') Joe Slagel: (20') Abdullah Kahraman: (20') Henning Hermjakob: (20')	EXHIBITION	HUMAN ANTIBODY INITIATIVE		
12:30 - 13:15	Lunch						
13:15 - 14:00							
14:30 - 15:00	Coffee break (14:30 - 15:00)	EUPA YOUNG INVESTIGATOR PRIZE	Coffee break (14:30 - 15:00)	POSTER 2 (EVEN #)	C-HPP (from 12.45)	LUMINEX	THERMO
15:00 - 17:00	S10 METABOLIC DISORDERS Invited speaker S10: Nelson Randall (30') Michael Przybylski: (20') Stefan Lehr: (20') Federica Taurino: (20') Feliciano Priego-Capote: (20')	S11 FUNCTIONAL PROTEOMICS Invited speaker S11: Mathias Mann (30') Joachim Klose: (20') Markku Varjosalo: (20') Bernd Bodenmiller: (20') Brett Larsen: (20')	S12 BIOMARKER DISCOVERY AND VALIDATION Invited speaker S12: C. Pineau (30') Thomas D Schreiber: (20') Christophe Masselon: (20') Annarita Farina: (20') Andrei P. Drabovich: (20')	EXHIBITION	EUPA COMPANY AFTERNOON	MODEL ORGANISMS	
17:00 - 18:30	HUPO General Assembly			BRUKER	PROTEOMICS MASTER CLASSES	DATA REPOSITORY	ETD WORKSHOP
19:00 - 22:00	Circus Congress night						



HUPO 2011
10TH WORLD CONGRESS
4-7 SEPTEMBER 2011
Geneva

Pre-Congress Programme



Pre-Congress Programme

SUNDAY 4 SEPTEMBER 2011

09:00-16:30

CLINICAL DAY

HALL A

Co-chairs: Daniel Chan, Pierre Burkhard

09:00 – 09:30	Introduction and overview of translational clinical proteomics Speaker: Daniel Chan
09:30 – 10:00	Clinical unmet needs: Opportunity for proteomic biomarkers Speaker: Denis Hochstrasser
10:00 – 10:30	The “Intended Use”: Study design for proteomic diagnostics Speaker: David Ransohoff
10:30 -11:00	Coffee Break
11:00 – 11:30	Development of clinical diagnostics: The process Speaker: Steve Binder
11:30 – 12:00	Validation of promising biomarkers: Critical Issues Speaker : Steve Skates
12:00 – 12:30	Creation of a biomarker pipeline: Critical issues Speaker: Ruedi Aebersold
12:30 – 13:30	Lunch
13:30 – 15:30	Four Selected from abstracts: Biomarker development studies with potentials for the translation into clinical diagnostics. (Presentation: 20 minutes. Questions from the judges: 10 minutes)
13:30 -14:00	B. Ayoglu
14:00 -14:30	L. Chiecchi
14:30 -15:00	N. Tiberti
15:00 -15:30	X. Robin
15:30 – 16:30	The verdict: A panel of judges (Chairs and Speakers)

Pre-Congress Programme

SUNDAY 4 SEPTEMBER 2011

09:00-16:30

EDUCATION DAY

HALL B

Co-chairs: Thierry Rabilloud, Peter James

09:00 – 09:15	Introduction from the co-chairs
09:15 – 10:00	Experimental and statistical planning Speaker: Kathryn Lilley
10:00 – 10:45	2D gel electrophoresis Speaker: Thierry Rabilloud
10:45 – 11:15	Coffee Break
11:15 – 12:00	Peptide IEF Speaker: Loic Dayon
12:00 – 12:45	Shotgun proteomics Speaker: Mike Mac Coss
12:45 – 14:00	Lunch
14:00 – 14:45	MRM Speaker: Susan Abbatiello
14:45 – 15:30	Bioinformatics and databases Speaker: Pierre-Alain Binz
15:30 – 16:15	Phosphorylations and glycosylations Speaker: Tine Thingholm
16:15 – 16:30	Conclusions from the co-chairs

Pre-Congress Programme

SUNDAY 4 SEPTEMBER 2011**09:00-16:30**

EuPA DAY

HALL C

Co-chairs: *Jésus Jorin Novo, Jean-Jacques Diaz*

- 09:00 – 09:30 **RNA-Seq and directed shotgun proteomics: towards complete proteome discovery in prokaryotes**
Speaker: Christian Ahrens
- 09:30 – 10:00 **Microorganisms**
Speaker: Peter Jungblut
- 10:00 – 10:30 **Candida albicans - host interaction: insights from proteomics**
Speaker: Concha Gil
- 10:30 – 11:00 Coffee Break
- 11:00 – 11:30 **Plants**
Speaker: Stepanie Wienkoop
- 11:30 – 12:00 **Plant proteomics: past, present and future. The launching of INPPO: the International Plant Proteomics Organization**
Speaker: Dominique Job
- 12:00 - 12:30 **What are animals? a proteomics view**
Speaker: Albert Heck
- 12:30 – 13:00 **Prohibitin 1 as a regulator of proliferation and cell death in liver cells. Mechanisms and implications in liver injury**
Speaker: Fernando Corrales
- 13:00 – 14:00 Lunch
- 14:00 – 15:30 **Six selected oral presentations (15 min each) from submitted contributions: 2 microorganisms, 2 plants, 2 animals**
Speakers: Niranjan Chakraborty, Adelina Rogowska - Wrzesienska, Pao - Chi Lao, Amandine Boeuf, Jianke Li, Brian Tate Weinert
- 15:30 – 16:30 **Discussion / Concluding remarks**
Jean-Jacques Diaz, Jésus Jorin Novo

Pre-Congress Programme

SUNDAY 4 SEPTEMBER 2011

09:00-16:30**SYMPOSIUM ON NEW TECHNOLOGIES AND STANDARDIZATION IN PROTEOMICS****ROOM K**

Topics covered New Technologies, Clinical Proteomics, Novel MS
Instrumentation and Standardization

Organized by HUP0 New Technology and Resources Committee

Chairs Christoph Borchers & Bruno Domon

Speakers Susan Abbatiello,
Roumen Bogoev,
Eric Deutsch,
Will Dracup,
Chris Kinsinger,
James Langridge,
Alexander Makarov,
Eugene Nikolaev,
Richard Smith,
Keith Waddell

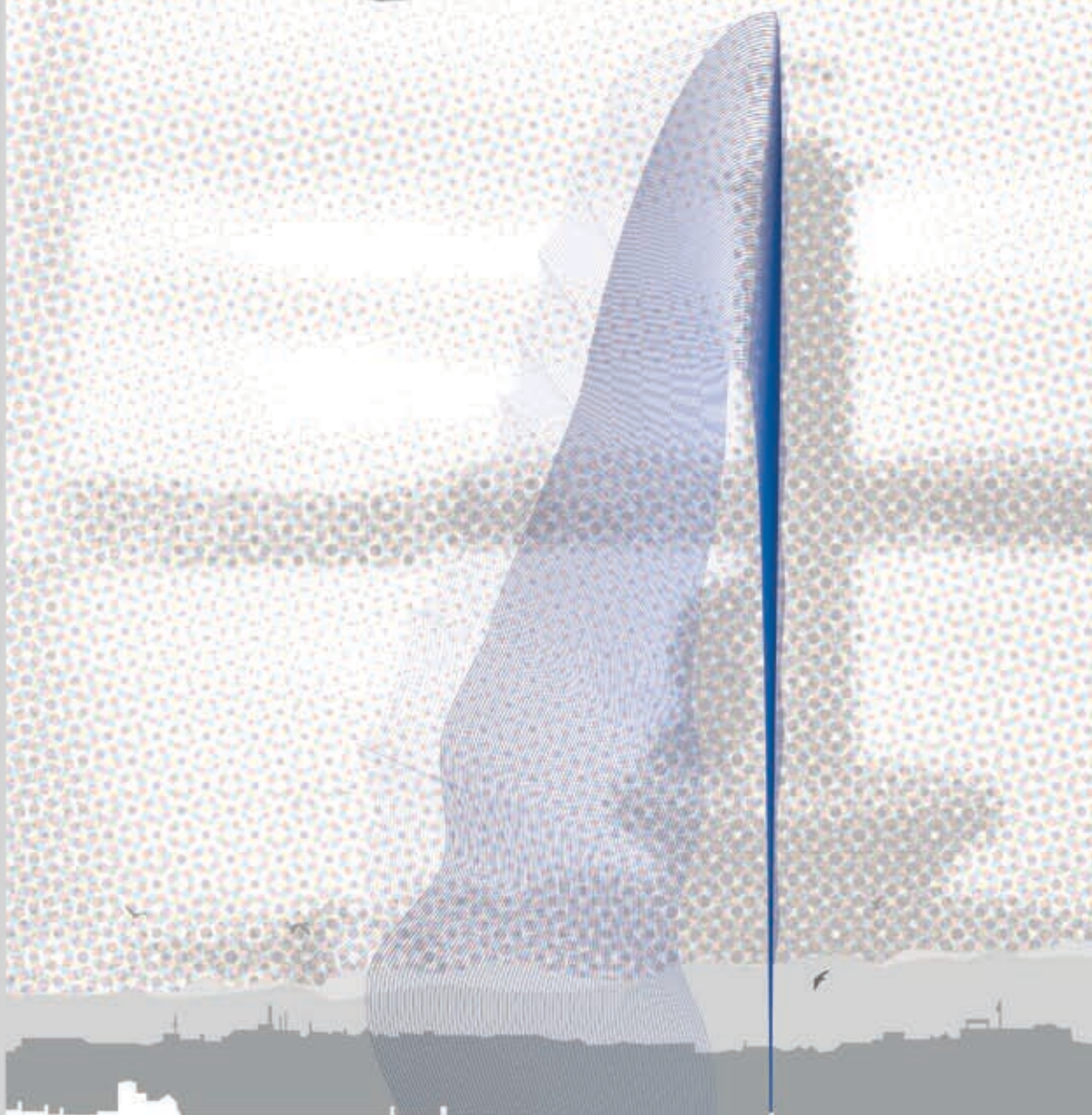
Free to HUP0 members and conference attendees.

13:30 - 16:30 Hupo Council

ROOM L

HUPO 2011
10TH WORLD CONGRESS
4-7 SEPTEMBER 2011
Geneva

Scientific Programme



Scientific Programme

SUNDAY 4 SEPTEMBER 2011

17:00-17:30	Opening ceremony Chairs: Patricia Palagi Speaker: Jean-Charles Sanchez, Henri Bounameaux & Catherine Costello	HALL A, HALL B
17:30-18:15	Plenary Lecture 1 Chair: Denis Hochstrasser The IG nobel prizes Speaker: Marc Abrahams	HALL A, HALL B
18:15-19:00	Plenary Lecture 2 Chair: Denis Hochstrasser Protein modelling and drug discovery Speaker: Michael Sternberg	HALL A, HALL B
19:00-22:00	Welcome Reception	EXHIBITION AREA

MONDAY 5 SEPTEMBER 2011

08:30-09:15	Plenary Lecture 3 Chair: Denis Hochstrasser The Connectivity of Genomic Elements Speaker: Stylianos Antonorakis <i>With the kind support of ASBMB</i>	HALL A, HALL B
09:15-10:00	Plenary Lecture 4 Chair: Denis Hochstrasser Proteins And Genes Of The Circadian Mammalian Timing System Speaker: Ueli Schibler	HALL A, HALL B
10:30-12:30	S1 Infectious diseases Chair: Jacques Schrenzel	HALL A
10:30-11:00	Mass Spectrometry In Clinical Microbiology: Fascination, Facts And Future Speaker: Jacques Schrenzel	
11:00-11:20	OP012 - PROTEOME-WIDE SELECTED REACTION MONITORING ANALYSIS OF COMPLETE MICROBIAL PROTEOMES – A CASE STUDY IN STREPTOCOCCUS PYOGENES VIRULENCE MECHANISMS Johan Malmström	

Scientific Programme

11:20-11:40	OP013 - MICROORGANISM CHARACTERIZATION FOR THE CLINICS USING SRM Jean-Philippe Charrier	
11:40-12:00	OP014 - IMPACT OF GUT MICROBIOTA ON HOST BIOLOGY - WHO IS DOING WHAT? Marianne Danielsen	
12:00-12:20	OP015 - FULL OMICS PROFILING REVEAL EXTENSIVE DYNAMIC CHANGES DURING NORMAL AND DISEASE STATES Michael Snyder	
10:30-12:30	S2 Organellar proteomics Chair: Angus Lamond	HALL B
10:30-11:00	Super Experiments: Studying Cell Biology Using Quantitative Proteomics Speaker: Angus Lamond	
11:00-11:20	OP016 - MAPPING THE SUBCELLULAR PROTEIN DISTRIBUTION IN THREE HUMAN CELL LINES – A SUBCELLULAR PROTEIN ATLAS Emma Lundberg	
11:20-11:40	OP017 - DISSECTING THE SPATIO-TEMPORAL ORGANIZATION OF THE HUMAN CENTROSOME PROTEOME Jens S Andersen	
11:40-12:00	OP018 - N-TERMINOMICS AND TARGETED PROTEOMICS FOR A BETTER CHARACTERIZATION OF THE HUMAN MITOCHONDRIAL PROTEOME Christine Carapito	
12:00-12:20	OP019 - MODULATION OF NEURONAL PENTRAXIN 1 EXPRESSION IN RAT PANCREATIC BETA-CELLS SUBMITTED TO GLUCOTOXIC STRESS Domitille Schvartz	
10:30-12:30	S3 MS Quantitation Chair: Ruedi Aebersold	HALL C
10:30-11:00	The Organization Of The Proteome In Space And Time Speaker: Ruedi Aebersold	
11:00-11:20	OP020 - AUTOMATION OF A SISCAPA MAGNETIC BEAD WORKFLOW FOR PROTEIN BIOMARKER QUANTITATION BY MASS SPECTROMETRY Leigh Anderson	
11:20-11:40	OP021 - GENOME-WIDE PARALLEL QUANTIFICATION OF MRNA AND PROTEIN LEVELS AND TURNOVER IN MAMMALIAN CELLS Matthias Selbach	

Scientific Programme

11:40-12:00	<p>OP022 - A DOUBLE SILAC STRATEGY FOR SIMULTANEOUS MEASUREMENT OF DIFFERENTIAL CHANGES IN PROTEIN DEGRADATION AND SYNTHESIS: APPLICATION TO STUDY THE EFFECTS OF INHIBITION OF THE HSP90 CHAPERONE MACHINE.</p> <p>Ivo Fierro-Monti</p>	
12:00-12:20	<p>OP023 - LABEL FREE QUANTITATIVE PROTEOMICS FOR THE DISCOVERY OF NOVEL MODIFICATION DEPENDENT HISTONE BINDING FACTORS</p> <p>H.Christian Eberl</p>	
15:00-17:00	<p>S4 Oncology</p> <p>Chair: Giulio Superti Furga</p>	HALL A
15:00-15:30	<p>Molecular Networks In Leukemia And Innate Immunity</p> <p>Speaker: Giulio Superti Furga</p>	
15:30-15:50	<p>OP024 - IDENTIFICATION OF A HIGH-SPECIFICITY AND SENSITIVITY AUTOANTIBODY-BASED COLORECTAL CANCER PREDICTOR PANEL BY COMBINING DIFFERENT PROTEIN MICROARRAY APPROACHES</p> <p>Rodrigo Barderas</p>	
15:50-16:10	<p>OP025 - RATIONAL DESIGN OF PLASMA PROTEIN SIGNATURES FOR EARLY DETECTION OF COLORECTAL CANCER</p> <p>Silvia Surinova</p>	
16:10-16:30	<p>OP026 - LABEL-FREE LC-MS/MS AND MRM DEVELOPMENT FOR DISCOVERY AND VERIFICATION OF BIOMARKERS FOR PROSTATE CANCER</p> <p>Yue Fan</p>	
16:30-16:50	<p>OP027 - ACCURATE QUANTIFICATION OF TUMOR PROTEOMES WITH SUPER-SILAC AND HIGH RESOLUTION MASS SPECTROMETRY</p> <p>Tamar Geiger</p>	
15:00-17:00	<p>S5 Post-translational modification</p> <p>Chair: Ole Jensen</p>	HALL B
15:00-15:30	<p>Posttranslational modifications. The eternal Frontier of proteomics?</p> <p>Speaker: Ole Jensen</p>	
15:30-15:50	<p>OP028 - BUILDING THE METHYLPROTEOME NETWORK</p> <p>Marc Wilkins</p>	
15:50-16:10	<p>OP029 - GLOBAL PROFILING OF DNA DAMAGE-REGULATED PHOSPHOPROTEOME AND ACETYLOME</p> <p>Petra Beli</p>	
16:10-16:30	<p>OP030 - PROTEOME-WIDE ANALYSIS OF THE UBIQUITIN-LIKE MODIFIER URM1 BY QUANTITATIVE MASS SPECTROMETRY</p> <p>Christian Toft Madsen</p>	

Scientific Programme

16:30-16:50	OP031 - O-GLCNACYLATION AND PHOSPHORYLATION AT THE SYNAPSE: IS THERE EVIDENCE FOR PTM CROSS-TALK? Al Burlingame	
15:00-17:00	S6 Protein Arrays Chair: Dolores Cahill	HALL C
15:00-15:30	Clinical and translational applications of candidate biomarkers identified using Protein arrays Speaker: Dolores Cahill	
15:30-15:50	OP032 - PLASMA BIOMARKER DISCOVERY BY NEXT GENERATION AFFINITY ARRAYS Jochen M Schwenk	
15:50-16:10	OP033 - PROTEOMIC PROFILING OF AUTOIMMUNITY ON ANTIGEN MICROARRAYS Peter Nilsson	
16:10-16:30	OP034 - AFFINITY-BASED PROTEOMIC ANALYSIS OF PANCREATIC CANCER Jörg D. Hoheisel	
16:30-16:50	OP035 - MOLECULAR DESIGN OF RECOMBINANT ANTIBODIES FOR MICROARRAY-BASED APPLICATIONS Christer Wingren	

TUESDAY 6 SEPTEMBER 2011

08:30-09:15	Plenary Lecture 5 Chair: Jean Charles Sanchez	HALL A, HALL B
	Growth Inhibitors For Fibre Tracts And Circuits In The Brain And Spinal Cord Limit Repair After Injury Speaker: Martin Schwab	
09:15-10:00	Plenary Lecture 6 Chair: Jean Charles Sanchez	HALL A, HALL B
	Next Prot: Exploring the universe of human proteins Speaker: Amos Bairoch	
10:30-12:30	S7 CNS Chair: Joan Montaner	HALL A
10:30-11:00	Plasma biomarkers in ischemic stroke Speaker: Joan Montaner	

Scientific Programme

11:00-11:20	OP036 - FUNCTIONAL PROTEOMICS OF NATIVE AMPA RECEPTOR COMPLEXES REVEALS AN UNEXPECTED MOLECULAR DIVERSITY Uwe Schulte	
11:20-11:40	OP037 - QUANTITATIVE PROTEOMICS TO UNRAVEL NEURAL PROGENITOR/STEM CELLS (NPCS) INTERCELLULAR COMMUNICATION Angela Bachi	
11:40-12:00	OP038 - A PROTEIN SURFACEOME MAP OF GLIOBLASTOMA BRAIN CANCER CELLS FOR CLINICAL DIAGNOSTIC APPLICATIONS Thomas Bock	
12:00-12:20	OP039 - 2D-DIGE PROFILING OF HUMAN EMBRYONIC STEM CELLS DURING NEURAL DIFFERENTIATION Ali Fathi	
<hr/>		
10:30-12:30	S8 Membranes proteins Chair: Jerome Garin	HALL B
10:30-11:00	Exploring membrane proteomes : Some emerging solutions to many challenges Speaker: Jerome Garin	
11:00-11:20	OP040 - ABUNDANCE OF CANCER BIOMARKERS AMONG THE PLASMA SOLUBLE HLA PEPTIDOME Arie Admon	
11:20-11:40	OP041 - AN ABSOLUTE QUANTIFICATION METHOD OF MOUSE TRANSMEMBRANE PROTEOME USING A COMBINATION OF WHEAT CELL-FREE SYNTHESIS AND MRM MASS SPECTROMETRY Nobuaki Takemori	
11:40-12:00	OP042 - THE CELL SURFACE PROTEIN ATLAS (CSPA) Bernd Wollscheid	
12:00-12:20	OP043 - CELL SURFACE CAPTURING REVEALS NEW MARKERS FOR IMMUNOPHENOTYPING CARDIAC PROGENITORS Rebekah L. Gundry	
<hr/>		
10:30-12:30	S9 Data and knowledge Chair: Amos Bairoch	HALL C
10:30-10:50	OP044 - UNICARB-KB: A NEW GLYCOINFORMATICS INITIATIVE Nicki Packer	
10:50-11:10	OP045 - A MASS SPECTROMETRIC MAP FOR CANCER-ASSOCIATED PROTEINS APPLIED TO HUMAN PLASMA AND URINE Ruth Hüttenhain	

Scientific Programme

11:10-11:30	OP046 - ANALYSIS OF LARGE LC-MS/MS DATASETS USING THE AMAZON WEB SERVICE CAPABILITIES OF THE TRANS-PROTEOMIC PIPELINE Joe Slagel	
11:30-11:50	OP047 - STRUCTURAL SYSTEMS BIOLOGY USING CROSS-LINK DATA AND COMPUTATIONAL MODELLING Abdullah Kahraman	
11:50-12:10	OP048 - PROTEOMEXCHANGE - STANDARDS AND INFRASTRUCTURE FOR PROTEOMICS Henning Hermjakob	
13:15-15:00	EUPA Young Investigator Price Chairs: Luca Bini & Garry Corthals	ROOM K
13:15- 13:35	OP078 - MOUSE MODEL DRIVEN DISCOVERY OF PROTEIN BIOMARKERS FOR OVARIAN CANCER AND THEIR VERIFICATION IN HUMAN PLASMA BY SRM Ruth Hüttenhain	
13:35- 13:55	OC079 - FROM HERO TO ZERO - INADEQUATE CELLULAR STRESS RESPONSES AFTER EXPOSURE TO PERITONEAL DIALYSIS FLUIDS AND POTENTIAL TARGETS FOR CYTOPROTECTION Klaus Kratochwill	
13:55- 14:15	OC080 - MALDI MSI FOR THE SCREENING OF OVARIAN CANCER BIOMARKERS: NEW INSIGHTS OF THE TECHNOLOGY FOR NEW ISSUES IN PHARMACOLOGY. Rémi Longuespée	
14:15- 14:35	OC081 - GLYCAN ISOFORMS-MODULATION OF THE METASTATIC PROCESS IN BREAST CANCER Giuseppe Palmisano	
14:35- 14:55	OC082 - CEREBROSPINAL FLUID BIOMARKERS TO DIAGNOSE THE MENINGO-ENCEPHALITIC STAGE OF SLEEPING SICKNESS Natalia Tiberti	
15:00-17:00	S10 Metabolic disorders Chair: Randal Nelson	HALL A
15:00-15:30	Building Multidimensional Biomarker Views Of Type 2 Diabetes And Comorbidities Based On Protein Microheterogeneity Speaker: Randal Nelson	

Scientific Programme

15:30-15:50	OP049 - CLINICAL DIAGNOSTICS OF LYSOSOMAL STORAGE DISEASES: DRY BLOOD SPOT ANALYSIS BY MULTIPLE REACTION MONITORING MASS SPECTROMETRY AND AFFINITY-PROTEOMICS OF LYSOSOMAL TARGET PROTEINS Michael Przybylski	
15:50-16:10	OP050 - IDENTIFICATION OF A NOVEL ADIPOKINE: DIPEPTIDYL PEPTIDASE 4 A POTENTIAL LINK BETWEEN OBESITY AND THE METABOLIC SYNDROME Stefan Lehr	
16:10-16:30	OP051 - STREPTOZOTOCIN-INDUCED DIABETES AFFECTS COMPLEX I ACTIVITY BY LOWERING NDUFS3 LEVELS IN RAT BRAIN MITOCHONDRIA Federica Taurino	
16:30-16:50	OP052 - PROTEIN GLYCATION IN HUMAN BODY FLUIDS AND BLOOD CELLS Feliciano Priego-Capote	
15:00-17:00	S11 Functional proteomics Chair: Mathias Mann	HALL B
15:00-15:30	In depth and rapid proteome coverage for functional proteomics Speaker: Mathias Mann	
15:30-15:50	OP053 - MALES AND FEMALES AND THE COMPLEXITY OF THE PROTEOME Joachim Klose	
15:50-16:10	OP054 - CHARTING THE HUMAN KINOME INTERACTOME Markku Varjosalo	
16:10-16:30	OP055 - SINGLE CELL MASS SPECTROMETRY TO COMPREHENSIVELY STUDY IMMUNE- AND CANCER CELL SIGNALING NETWORKS Bernd Bodenmiller	
16:30-16:50	OP056 - TRACKING AND FILTERING INTERACTION PROTEOMICS DATA Brett Larsen	
15:00-17:00	S12 Biomarker discovery tools Chair: Charles Pineau	HALL C
15:00-15:30	Fertichip: A Predictive Assay To Assess The Presence Of Spermatozoa In The Testes Of Infertile Men Speaker: Charles Pineau	
15:30-15:50	OP057 - COMBINING ULTRACENTRIFUGATION AND PEPTIDE TERMINI GROUP-SPECIFIC ENRICHMENT FOR MULTIPLEX ANALYSIS OF PLASMA PROTEINS Thomas D Schreiber	
15:50-16:10	OP058 - BLADDER CANCER CANDIDATE BIOMARKERS DISCOVERY BY AMT TAG PROTEOMICS ON A LARGE MULTI-SITE COHORT Christophe Masselon	

Scientific Programme

16:10-16:30	OP059 - PROTEOMIC ANALYSIS OF BILE AND PANCREATIC CYST FLUID REVEALS PROMISING SOURCES OF CANCER MARKERS Annarita Farina	
16:30-16:50	OP060 - DIFFERENTIAL DIAGNOSIS OF AZOOSPERMIA WITH PROTEOMIC BIOMARKERS QUANTIFIED IN SEMINAL PLASMA BY MULTIPLEX SELECTED REACTION MONITORING ASSAY Andrei P Drabovich	
17:00-18:30	HUP0 General assembly	HALL A
19:00-20:00	Reception at the Swiss National Circus	
20:00-23:00	Congress Circus Event	

WEDNESDAY 7 SEPTEMBER 2011

08:30-09:15	Plenary Lecture 7 Chair: Luca Bini	HALL A, HALL B
	Cell Therapy In Parkinson Disease Speaker: Patrick Brundin <i>With the kind support of Roche</i>	
09:15-10:00	Plenary Lecture 8 Chair: Luca Bini	HALL A, HALL B
	Cern, Particle Accelerators And The LHC Collider Speaker: Steve Myers	
10:30-12:30	S13 Cardiovascular Chair: Dario Neri	HALL A
10:30-11:00	Vascular Tumor Targeting: From Target Identification To Clinical Trials Speaker: Dario Neri	
11:00-11:20	OP061 - PLASMA PROTEOMIC BIOMARKERS OF ACUTE HEART AND KIDNEY REJECTION: FROM AN UNBIASED ITRAQ DISCOVERY TO A TARGETED MULTIPLE REACTION MONITORING VALIDATION W. Robert McMaster	
11:20-11:40	OP062 - CARDIAC ISCHEMIA BIOMARKER DISCOVERY AND VALIDATION Jennifer E Van Eyk	
11:40-12:00	OP063 - THE PROTEOME OF HUMAN LP(A) POINTS TO A ROLE IN WOUND HEALING Anne Von Zychlinski	

Scientific Programme

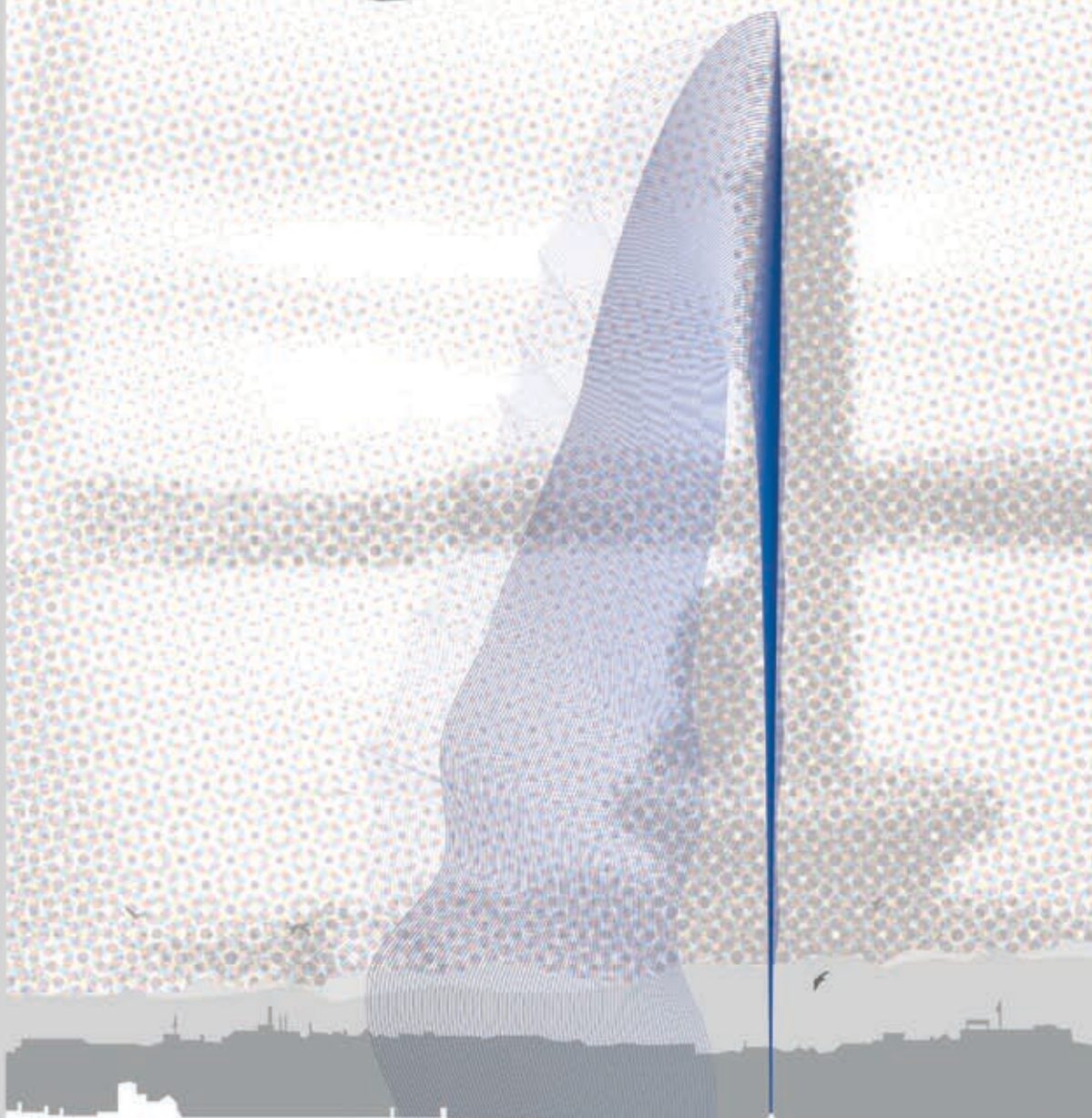
12:00-12:20	OP064 - IGFBP7 CONTRIBUTES TO VASCULAR REMOLDING DURING HYPERTENSION DEVELOPMENT IN SPONTANEOUS HYPERTENSIVE RATS REVEALED BY PROTEOMIC STUDY Jie Qiu	
10:30-12:30	S14 Pharmacotoxicology Chair: Gyorgy Marko-Varga	HALL B
10:30-11:00	Clinical Studies With Personalized Medicines And Biomarker Discovery Speaker: Gyorgy Marko-Varga	
11:00-11:20	OP065 - MONITORING DRUG METABOLIZING ENZYMES – ABSOLUTE QUANTIFICATION OF CYTOCHROME P450 3A4, 3A5, 3A7 AND MDR1 USING A SINGLE PEPTIDE TERMINUS-SPECIFIC ANTIBODY Oliver Poetz	
11:20-11:40	OP066 - CLOSTRIDIAL GLYCOSYLATING TOXINS REORGANIZE THE PROTEOME OF COLON CELLS Andreas Pich	
11:40-12:00	OP067 - IN SILICO DISEASE MODELS TO COMPARE KINASE INHIBITORS ACTION IN CANCER TREATMENTS Jacques Colinge	
12:00-12:20	OP068 - EXPLORING TIME-DEPENDENT DRUG PERTURBATIONS OF THE HER PHOSPHOTYROSINE SIGNALING NETWORK Manuel Tzouros	
10:30-12:30	S15 Imaging Chair: Ron Heeren	HALL C
10:30-11:00	Imaging Mass Spectrometry For Proteome Studies On Biological Surfaces Speaker: Ron Heeren	
11:00-11:20	OP069 - MALDI IMAGING IDENTIFIES PROGNOSTIC SEVEN-PROTEIN SIGNATURE OF NOVEL TISSUE MARKERS IN GASTRIC CANCER Axel Walch	
11:20-11:40	OP070 - THE HYBRID TECHNIQUE OF MALDI-TOF IMAGING AND MR IMAGING TO IMPROVE THE CORRELATION OF ANATOMICAL FEATURES AND MOLECULAR INFORMATION Herbert Thiele	
11:40-12:00	OP071 - DISTINGUISHING MORPHOLOGICALLY IDENTICAL TUMORS USING CORROBORATED MS-BASED MOLECULAR HISTOLOGY Liam Mcdonnell	

Scientific Programme

12:00-12:20	OP072 - MALDI IMAGING AT 20μM RESOLUTION OFFERS NEW POSSIBILITIES TO STUDY RAT SPERMATOGENESIS IN SITU Mélanie Lagarrigue	
13:15-15:00	Human Proteome Project Co-chairs: Juan Pablo Albar, Pierre Legrain, Gilbert Omenn, Young-Ki Paik	HALL A, HALL B
15:00-17:00	HUPO & EuPA Awards Chairs: Maxey Chung, Concha Gil EuPA Young Investigator Award Mike Dunn: EuPA Senior Scientist Award in Proteomics Steven Carr: HUP0 Discovery Award Amos Bairoch: HUP0 Distinguished Discovery Achievement Award Funchu He: HUP0 Distinguished Service Award Alexander Makarov: HUP0 Science Technology Award	HALL A, HALL B
17:00-18:30	Closing Ceremony Chair: Patricia Palagi	HALL A, HALL B
17:00 - 17:30	The Swiss Chocolate Speaker: Watzke Heribert	
17:30 - 18:00	Concluding Remarks Speaker: Jean-Charles Sanchez	
18:00 - 18:05	EUPA 2012: Glasgow	
18:00 - 18:10	HUP0 2012: Boston	
18:10 - 18:45	Farewell cocktail	

HUPO 2011
10TH WORLD CONGRESS
4-7 SEPTEMBER 2011
Geneva

Social Programme



Social Programme

OPENING CEREMONY (17:00-17:30) AND WELCOME RECEPTION (19:00 - 22:00)

SUNDAY 4 SEPTEMBER 2011 - PALEXPO

The Welcome Reception will take place inside the Exhibition Hall after the Plenary Lectures, from 19:00 – 21:00. The Welcome Reception is included in the Registration Fee.

CONGRESS RECEPTION

TUESDAY 6 SEPTEMBER 2011, 19:00 – 20:00

A buffet dinner is preceding the circus show.

- Meeting point : Circus Entrance, Plaine de Plainpalais
- Bus stop: Palexpo. **Bus n°5** direction Hôpital. Stop at Nations. Take **tram N°15** direction Palettes. Stop at Plainpalais.
- A reservation fee of CHF 54 is applicable.

CONGRESS CIRCUS EVENT

TUESDAY 6 SEPTEMBER 2011, 20:00 – 23:00

Come and enjoy a fantastic show in the Swiss National Circus (Circus Knie), a show with renowned international artists.

- Meeting point : Circus Entrance, Plaine de Plainpalais.
- Bus stop: Palexpo. **Bus n°5** direction Hôpital. Stop at Nations. Take **tram N°15** direction Palettes. Stop at Plainpalais.
- The cost for the circus show is included in the congress registration Fee.

ACCOMPANYING PERSONS' EXCURSION

CITY TOUR OF GENEVA

SUNDAY, 4 SEPTEMBER, MONDAY, 5 SEPTEMBER

3 HOURS - 9:00 TO 12:00 & 14:00 TO 17:00

Nobody should leave Geneva without seeing its monuments and historical sites. We will take you to the most important ones in three hours, including the symbol of the city, the famous "water jet"; this impressive 460ft (140m) high lake water fountain, the first of its kind, was completed in 1891. The flower clock in the English garden is another symbol of Geneva, known all over the world. The monument of Duke of Brunswick, who left his entire fortune to the City of Geneva, is located right on the opposite side of Lake Geneva. The historical Old Town will be visited on foot to discover its carefully preserved charm. The Cathedral Saint-Pierre constructed in the 12th and 13th centuries dominates the Old Town. Right next to it is the Auditorium, where Jean Calvin taught. Calvin arrived in 1536 in Geneva. Leading figure of the protestant Reformation, he deeply influenced religion, ethics and social life in the western world. The oldest privately constructed building is the Maison Tavel, now transformed into a museum of old Geneva. With its parks and gardens, its monuments and museums, Geneva is today truly international and cosmopolitan, not only in population but also in its whole atmosphere.

HUPO 2011
10TH WORLD CONGRESS
4-7 SEPTEMBER 2011
Geneva

HUPO Initiative Programmes



HUP0 Initiative Programmes

SUNDAY 4 SEPTEMBER 2011

09:00-12:00

HPP CONSORTIUM

ROOM L

Organized by Pierre Legrain, Gil Omenn and Young-Ki Paik

Theme Next steps in the development of HPP to become a top scientific initiative.

Goals To establish a formal HPP consortium (Technology, C-HPP and B/D-HPP).

Chair: Gil Omenn/Pierre Legrain

09:00 - 09:10	Introduction by Gil Omenn
09:10 - 09:25	Update on the HPP Working Group Activity (2010/2011) by Pierre Legrain, Manager of HPP WG, CEA, France (PC31)
09:25 - 11:25	Discussion: "Formation of the HPP Consortium" <ol style="list-style-type: none"> 1. Proposal for the Formation of the Technology Consortium of HPP with IAB inputs: The Three Pillars Moderator: Bill Hancock, Co-Chair of IAB and of C-HPP Consortium Technology & Bioinformatics Update, Ruedi Aebersold, Mathias Uhlen, Amos Bairoch, Eric Deutsch 2. Update on the Chromosome-Centric HPP (C-HPP) Working Group Activity by Young-Ki Paik, Chair of C-HPP, YPRC, Korea (PC032/OC074) 3. Presentation of a Proposal for the Formation of Biology/Disease-HPP Consortium by Gil Omenn & Ruedi Aebersold 4. Leadership: Nomination of Chairs of Technology, B/D-HPP and HPP consortium Moderator: Pierre Legrain 5. Introduction of Senior Scientific Advisory Board for HPP Moderator: Young-Ki Paik
11:25 - 11:35:	Conclusions and Action Plans
11:35 - 12:00	General Discussion

HUP0 Initiative Programmes

MONDAY 5 SEPTEMBER

07:00 – 08:30
HUMAN LIVER PROTEOME
ROOM K

- 07:00 - 08:30 Chairs: José M. Mato, Laura Beretta & Fuchu He
- 07:00 - 07:15 **OC001**
**SEARCHING FOR NON-INVASIVE BIOMARKERS FOR LIVER INJURY:
 EXOSOME AND PEPTIDOME APPROACHES**
 Felix Elortza
- 07:15 - 07:30 **OC002**
**PROTEOMIC ANALYSIS OF HEPATOCELLULAR CARCINOMA
 IDENTIFICATION OF TARGET PROTEINS AND FUNCTIONAL IMPLICATIONS**
 Fernando Corrales
- 07:30 - 07:45 **OC003**
A METHOD FOR PROFILING TRANSCRIPTION FACTORS
 Jun Qin
- 07:45 - 08:00 **OC004**
A WORKFLOW FOR IN DEPTH COVERAGE OF THE MOUSE LIVER PROTEOME
 Jun Qin
- 08:00 - 08:15 **OC005**
**THE NOVEL ROLE OF AN EXOSOME SECRETION RELATED PROTEIN
 IN THE METASTASIS OF HEPATOCELLULAR CARCINOMA**
 Ying Jiang
- 08:15 - 08:30 **Concluding remarks**

MONDAY 5 SEPTEMBER 2011

10:30-12:30
HBPP SESSION - HUP0 2011
ROOM K
“Launching Human Brain Proteome Atlas Project”

- 10:30 - 10:50 **OC021**
Strategies for creating Human Brain Proteome Atlas
 Young Mok Park
- 10:50 - 11:10 **OC022**
**Strategies of tissue and cell sampling in early stages of AD in order to
 assess and to compensate confounding ante- and post-mortem factors**
 Helmut Heinsen
- 11:10 - 11:30 **OC024**
Maldi imaging For neuropathologies investigation.
 Michel Salzet (or Isabelle Fournier)

HUP0 Initiative Programmes

- 11:30 - 11:50 **OC025**
Biomarker for Neurodegenerative Diseases - Brain Regions of Interest
 Helmut Meyer
- 11:50 - 12:10 **OC023**
The level of DNA hydroxymethylation in Alzheimer's disease (AD) and normally aged brain.
 Harry Steinbusch
- 12:10 - 12:30 **OC026**
Characterization and quantification of α -synuclein – a potential biomarker for Parkinson's disease
 Katrin Marcus

MONDAY 5 SEPTEMBER

EuPA PRESENT AND FUTURE

Chair: Gyorgy Marko-Vargas

13:15 – 15:00

HALL A

OC027

SHAPING THE FUTURE OF THE EUROPEAN PROTEOMICS ASSOCIATION EUPA

Gyorgy Marko-Vargas

MONDAY 5 SEPTEMBER 2011

HUP0 PROTEOMICS STANDARDS INITIATIVE

15:00-17:00

ROOM G

The HUP0 Proteomics Standards Initiative (PSI) defines community standards for data representation in proteomics to facilitate data comparison, exchange and verification. In this session we discuss current status and future plans, and invite feedback from the community and fellow HUP0 Initiatives.

Agenda

- 15:00 – 15:10 **Welcome**
 PSI overview
 Pierre-Alain Binz, SIB.
- 15:10 – 15:25 **Molecular Interactions**
 As a key result of the EU PSIMEx grant, the PSICQUIC common query interface for currently 15 interaction data resources has been published
 Sandra Orchard, EBI

HUP0 Initiative Programmes

15:25 – 15:40	<p>Mass Spectrometry</p> <p>The Transitions Markup Language (TraML) is a new format being developed for encoding selected reactions monitoring transitions</p> <p>Eric Deutsch, ISB</p>
15:40 – 15:55	<p>Protein Identifications</p> <p>mzIdentML (released) and mzQuantML (under development) provide a full-featured representation for MS-based protein identification and quantification with full evidence track</p> <p>Martin Eisenacher, MPC Bochum</p>
15:55 – 16:05	<p>MIAPE Quant</p> <p>MIAPE guidelines for Mass Spectrometry Quantification are now being defined. These guidelines will help to standardize the reporting for the quantitative data and will allow a better understanding and critical evaluation of these approaches</p> <p>Salvador Martinez-Bartolomé, ProteoRed</p>
16:05 – 16:15	<p>mzTab</p> <p>The mzTab format provides a concise overview of protein identifications and quantitations in a standardized format</p> <p>Juan Antonio Vizcaino, EBI</p>
16:15 – 16:25	<p>2D case study</p> <p>A DIGE study on the effects of salbutamol on the rat muscle proteome - an exemplar of best practice for data sharing in proteomics</p> <p>Juan Pablo Albar, ProteoRed</p>
16:25 – 16:35	<p>ProteomeXchange</p> <p>The ProteomeXchange consortium aims to provide a single point of submission to a network of collaborating proteomics repositories</p> <p>Henning Hermjakob, EBI</p>
16:35 - 17:00	<p>Open Discussion</p>

MONDAY 5 SEPTEMBER 2011

HUMAN DISEASE GLYCOMICS / PROTEOME INITIATIVE

Chairs: Hisashi Narimatsu, Naoyuki Taniguchi, Anne Dell

17:00-19:00

HALL A

17:00 - 17:20	<p>OC046</p> <p>FUCOSYLATED ALPHA-FETOPROTEIN(AFP): STRUCTURAL BASIS AND DEVELOPMENT OF ANTIBODY-LECTIN ENZYME IMMUNOASSAY</p> <p>Naoyuki Taniguchi</p>
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HUP0 Initiative Programmes

- 17:20 - 17:40 **OC047**
SUGARS IN THE REAL WORLD: ARE CULTURED CELLS A GOOD MODEL
SYSTEM FOR STUDYING PROTEIN GLYCOSYLATION?
Nicki Packer
- 17:40 - 18:00 **OC048**
GLYCOPROTEOMICS APPROACH TOWARD DISCOVERY OF GLYCOBIOMARKERS
Hisashi Narimatsu
- 18:00 - 18:20 **OC049**
ABERRANT O-GLYCOSYLATION OF IGA1 AND PATHOGENESIS
OF IGA NEPHROPATHY
Jan Novak
- 18:20 - 18:40 **OC050**
MASS SPECTROMETRIC STRATEGIES FOR GLYCOMICS AND GLYCOPROTEOMICS
Stuart Michael Haslam
- 18:40 - 19:00 **Open Discussion**

MONDAY 5 SEPTEMBER 2011

HUP0 CARDIOVASCULAR INITIATIVE

Chair: Peipei Ping

17:00-19:00

HALL B

- 17:00 -17:30 **Proteome Biology of Cardiovascular Medicine: On the Boat of
Translational Research**
Peipei Ping
- 17:30 -18:00 **Reorganization of cyclic nucleotide signaling pathways in dilated
cardiomyopathic hearts**
Arjen Scholten
- 18:00 -18:30 **Human Protein Knowledgebase for the Heart and Soul**
Henning Hermjakob
- 18:30 - 19:00 **Open Discussion**

HUP0 Initiative Programmes

MONDAY 5 SEPTEMBER 2011

17:00-19:00

PROTEOME BIOLOGY OF STEM CELLS INITIATIVE

HALL C

Chairs: J. Krijgsveld , A. Heck

- | | |
|---------------|--|
| 17:00 - 17:20 | OC051
USING MASS SPECTROMETRY BASED PROTEOMICS TO UNRAVEL KEY MOLECULAR COMPONENTS IN STEM CELL BIOLOGY
A. Heck |
| 17:20 - 17:40 | OC052
PROTEOME ANALYSIS OF HAEMATOPOIETIC STEM CELLS AND MULTIPOTENT PROGENITOR CELLS
D. Klimmeck |
| 17:40 - 18:00 | OC061
QUANTITATIVE PROTEOMICS TO UNRAVEL NEURAL PROGENITOR/STEM CELLS (NPCS) INTERCELLULAR COMMUNICATION
A. Bachi |
| 18:00 - 18:20 | OP086
GLYCOBIOLOGY AND GLYCOMICS OF EMBRYONIC STEM CELLS
S. Dalton |
| 18:20 - 18:40 | OC053
IN DEPTH QUANTITATIVE PROTEOMICS ON HUMAN EMBRYONIC STEM CELL DIFFERENTIATION
B. Blagoev |
| 18:40 - 19:00 | Open Discussion |

MONDAY 5 SEPTEMBER 2011

17:00-19:00

PROTEOMICS MASTER CLASSES

ROOM G

Chair: Thierry Rabilloud

DETERMINATION AND STUDY OF PHOSPHOSITES IN PROTEINS

Speakers : Martin Larsen, Garry Corthals, Tine Thingholm

The HUP0 master classes aim at giving an advanced knowledge on hot topics to researchers who are not beginners in proteomics but are not necessarily experts in these precise topics. To this purpose, experts in the field will be present for the duration of the workshop to answer the practical and fundamental questions of the audience in a workshop/round table style."

HUP0 Initiative Programmes

MONDAY 5 SEPTEMBER 2011

17:00-19:00

MANUAL INTERPRETATION OF ELECTRON TRANSFER
DISSOCIATION (ETD) MASS SPECTRA OF PEPTIDES

ROOM L

Speaker: Donald F. Hunt,

Electron transfer dissociation mass spectrometry is a break-through technology for sequencing post-translationally modified peptides. In this technique, radical anions of polyaromatic hydrocarbons (fluoranthene and azulene) are employed to transfer electrons to the carbonyl group in the polyamide backbone of multiply charged peptides generated by electrospray ionization. Capture of an electron into the peptide backbone reduces the positive charge on the ion by one, and forms a carbonyl radical anion that then abstracts a proton from a nearby protonated amino group. The resulting carbonyl radical triggers cleavage of the adjacent nitrogen-carbon bond to produce fragments of type c' and z•.

The purpose of the workshop is to provide instruction on how to manually interpret peptide ETD mass spectra. Following a lengthy tutorial about ion structures, fragmentation pathways, predictable changes in fragment ion isotope patterns, etc., we will outline a general approach for the manual interpretation of peptide ETD spectra, solve the sequence of several post-translationally modified peptides, assign homework spectra, and reconvene on the second day of the workshop to go over the homework problems and to explore additional applications of this powerful technology. A packet of lecture notes and handouts will be provided. Attendees should bring an inexpensive calculator, pad of paper, and a small ruler.

TUESDAY 6 SEPTEMBER 2011

07:00-08:30

HKUPP INITIATIVE (KIDNEY AND URINE INITIATIVE)

ROOM K

Chairs: Tadashi Yamamoto, Harold Mischak, Roby Langham

07:00 - 07:15

OC054

PARTICIPATION OF HUMAN KIDNEY AND URINE PROTEOME PROJECT (HKUPP)
INITIATIVE IN CROSSINITIATIVE ANALYSIS AND HUMAN PROTEOME PROJECT
T. Yamamoto

07:15 - 07:30

OP055

COMPARISON OF ON-SITE PROTEOMICS STUDY ON LASER MICRODISSECTED FFPE
AND FROZEN HUMAN GLOMERULUS TISSUES
B. Xu

HUP0 Initiative Programmes

07:30 - 07:45	OP056 L-FABP AND H-FABP AS MARKERS FOR ACUTE RENAL INJURY: A CLINICAL PROTEOMICS STUDY H. Dihazi
07:45 - 08:00	OP057 Mass spectrometric characterization of human urine proteome changes induced by space flight Kononokhin A.S
08:00 - 08:15	OP058 Extensive profiling of human urine proteome: construction of SRM transitions for quantification of urinary biomarkers for acute and chronic kidney diseases. Y. Yoshida
08:15 - 08:30	Open Discussion

TUESDAY 6 SEPTEMBER 2011

HAI SESSION (HUMAN ANTIBODY INITIATIVE)

Chairs: Michael Snyder and Mathias Uhlen

10:30-12:30

ROOM K

10:30 - 10:40	Welcome and short background to HAI Michael Snyder and Mathias Uhlen
10:40 - 10:55	NIH Protein capture reagents Salvatore Sechi
10:55 - 11:20	Recombinant antibodies with phage ESCape Michael Snyder
11:20 - 11:45	The Human Protein Atlas Mathias Uhlen
11:45 - 12:10	Antibodypedia - a portal for validated antibodies Kalle Jonasson
12:10 - 12:30	HAI contribution to the HPP - discussion Michael Snyder and Mathias Uhlen

HUP0 Initiative Programmes

TUESDAY 6 SEPTEMBER 2011

12:45-15:00

C-HPP CONSORTIUM

ROOM E

Organized by Young-Ki Paik, Bill Hancock and György Marko-Varga

Theme Setting up short-term and long-term plans coupled to the C-HPP Guidelines
Goals To execute C-HPP guidelines and organization of the C-HPP sub-committees
Co-Chairs Young-Ki Paik, Bill Hancock and György Marko-Varga

- 12:45 - 12:55 **Introduction-Key Issues for the Execution of C-HPP Guidelines**
 by Young-Ki Paik, Chair of C-HPP Consortium
- 12:55 - 13:20 **Major Technological Issues for the C-HPP: Web portal, data submission, samples, data integration, etc.**
 by Bill Hancock, co-chair of C-HPP consortium
- C-HPP Web Portal** by Amos Bairoch
- Data Submission & Integration** by Henning Hermjakob (EBI) & Eric Deutsch (ISB)
- Geomics / Transcripomics** by Michael Snyder
- Samples and Affinity Reagents** by Mathias Uhlen
- 13:20 - 14:10 **Introduction of New C-HPP Teams Since Sydney Congress**
 Chair: György Marko-Varga
 5 minutes for each presentation
- | | |
|---------------|--|
| Chromosome 1 | (China) by Fuchu He |
| Chromosome 2 | (Switzerland) by Pierre-Alain Binz (OC077) |
| Chromosome 3 | (Japan) by Toshihide N.Shimura |
| Chromosome 6 | (Canada) by Paul Keown |
| Chromosome 7 | (Australia) by Mark Baker |
| Chromosome 11 | (Korea) by Jong Shin Yoo |
| Chromosome 14 | (France) by Jerome Garin & Charles Pineau |
- 14:10 - 14:35 **Update on the ongoing C-HPP**
 Chair: William Hancock
- Chromosome 18 (Russia)** by Alex Archakov (OP081/OP077) & A.V Lisirsa (OP078 / OP080)
- Chromosome 19 (Sweden / others)** by Juan Pablo Albar & György Marko-Varga (OP076)
- Others (Until time is permitted)**
- 14:30 - 15:00 **Setting Up the Action Plans, Guidelines, and Governance: General discussion**
 Chair: Young-ki Paik

HUP0 Initiative Programmes

TUESDAY 6 SEPTEMBER 2011

15:00-17:00

EUPA COMPANY AFTERNOON

ROOM E

TUESDAY 6 SEPTEMBER 2011

15:00-17:00

INITIATIVE ON MODEL ORGANISM PROTEOMES (iMOP) (OC098)

ROOM F

Chairs: S. Schrimpf, M. Hengartner

15.00 – 15.10	Introduction Michael Hengartner
15.10 – 15.35	PaxDB Christian von Mering
15.35 – 16.00	Farm animal proteomics (OC099) Emøke Bendixen
16.00 – 16.25	Plant proteomics (OC100) Joshua Heazlewood
16.25 – 16.50	Host-pathogen interaction Dirk Bumann
16.50 – 17.00	General discussion about iMOP

TUESDAY 6 SEPTEMBER

17:00 – 18:30

PROTEOMICS MASTER CLASSES

ROOM E

Chair: Thierry Babilloud

HOW TO BUILD EFFICIENT SRM ASSAYS

Speakers: Ruedi Aebersold, Steve Carr, Susan Abbatiello

The HUP0 master classes aim at giving an advanced knowledge on hot topics to researchers who are not beginners in proteomics but are not necessarily experts in these precise topics. To this purpose, experts in the field will be present for the duration of the workshop to answer the practical and fundamental questions of the audience in a workshop/round table style."

HUP0 Initiative Programmes

TUESDAY 6 SEPTEMBER 2011

17:00-18:30

PROTEOMICS REPOSITORIES AND JOURNALS – A PARTNERSHIP MADE IN H...?

ROOM F

Chairs: Henning Hermjakob and Juan Pablo Albar

Proteomics repositories and journals are the two disseminators of proteomics data and results, so clearly this is a partnership made in heaven, or is it?

In the standard reference example of genomics, the world used to be easy: You generate a sequence, you submit it to a repository, and you publish on it. In proteomics, the situation slowly developed in this direction, but this development is far from complete, and even the “standard” genomics model is starting to creak under the pressure of high throughput sequencing.

In this workshop we will discuss past, present, and future of the interplay between journals and repositories, how well the current situation is serving the community, and which directions future developments should/could take.

Agenda:

- | | |
|---------------|--|
| 17:00 – 17:10 | Welcome and ProteomeXchange concept
Chair: Henning Hermjakob, European Bioinformatics Institute |
| 17:10 – 17:40 | A journal, or a database?
Scott Edmunds, BGI Shenzhen; Editor in chief, GigaScience journal |
| 17:40 – 18:00 | Raising the standards
Robert Chalkley, UCSF; Editor, Molecular and Cellular Proteomics journal &
Henning Hermjakob, European Bioinformatics Institute |
| 18:00 – 18:30 | Podium discussion
Chair: Juan Pablo Albar,
Participants: Scott Edmunds, Robert Chalkley, Eric Deutsch
Christopher Kinsinger, Juan J. Calvete |

This workshop is sponsored by the EU ProteomeXchange grant.

HUP0 Initiative Programmes

WEDNESDAY 7 SEPTEMBER 2011

07:00-08:30

IAB WORKSHOP AND AWARD

ROOM E

Chairs: A. Paulus, Bill Hanckok

- 07:00 – 07:25 **The Orbitrap story: from dream to mainstream**
Alexander Makarov - Thermo Fisher Scientific, Bremen, Germany
Winner - HUP0 Science and Technology Award
- 07:25 – 07:35 **Advanced mass spectrometry for speed independent proteomics**
Jim Landridge, Director, Proteomics - Waters
- 07:35 – 07:45 **New approaches for improving assay reproducibility for quantitation of peptides
and new approaches for improving protein identification**
Keith Waddell, LC/MS Application Solutions Manager - Agilent
- 07:45 – 07:55 **Integrated Mass Spectrometry-based Solutions: Around the Proteome in 10 Minutes**
Andreas Hühmer, Director of Proteomics Marketing - Thermo Fisher
- 07:55 – 08:05 **PTMScan Technology: Innovative Tools for Pathway Profiling, Biomarker Discovery
and Target Identification**
John-David Herlihy, Pharma Service and Technical Application Specialist -
Cell Signaling Technology
- 08:05 – 08:15 **Extending the reach of proteomics - tools and trends**
Ron Bonner, Principal Scientist - ABSciex
- 08:15 – 08:25 **Advancements in low abundant proteins detection and consistency
of proteome analysis**
Roumen Bogoev, Senior Product Manager - BioRad

WEDNESDAY 7 SEPTEMBER 2011

10:30-12:30

HUMAN PLASMA PROTEOME INITIATIVE

ROOM K

Chairs: Gil Omenn, Ruedi Aebersold, Mark Baker

- 10:30 – 11:00 **OC117**
THE HUPO HUMAN PLASMA PROTEOME PROJECT: 2011 UPDATE
Gil Omenn
- 11:00 – 11:30 **OC118**
**ENABLING COMPREHENSIVE PROTEOMICS MEASUREMENTS WITH
THE DEVELOPMENT OF THE COMPLETE HUMAN PEPTIDE- AND SRM-ATLAS**
Robert Moritz

HUP0 Initiative Programmes

11:30-12:00	OC119 CEDAR: A MULTI-TIERED PROTEIN IDENTIFICATION SCHEME FOR SHOTGUN PROTEOMICS, APPLIED TO THE HUMAN PLASMA PEPTIDEATLAS Eric Deutsch
12:00-12:30	PC033 THE HUMAN PLASMA PROTEOME IN PEPTIDEATLAS Eric Deutsch

WEDNESDAY 7 SEPTEMBER 2011

13:15-15:00

HPP PLENARY SESSION

Organized by Gil Omenn, Pierre Legrain and Young-Ki Paik

Theme	Future Direction of HPP
Goals	To establish the global commitment and engagement for the HPP
Moderator	Young-Ki Paik,

Topic - I : Technology and Biology/Disease-Driven Human Proteome Project (B/D-HPP)

13:15 - 13:30	Network-Centric Human Proteome Project Ruedi Aebersold
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13:30 - 13:45	Human Antibody Resources for HPP Mathias Uhlen
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Topic-II: Chromosome-Centric Human Proteome Project (C-HPP)

13:45 - 14:00	Encode the Human Parts List Michael Snyder
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14:00 - 14:15	Studies on Chromosome Biology and Cancer Proteomics within C-HPP (OC075) Bill Hancock
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Topic-III: Clinical Implications of HPP and Concluding Remarks

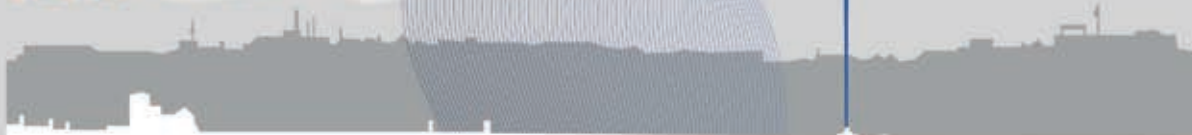
14:15 - 14:30	Clinical proteomics: are we missing opportunities?" Denis Hochstrasser
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14:30 - 14:45	HPP Goals and Vision Cathy Costello
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14:45 - 15:00	General Discussion
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HUPO 2011
10TH WORLD CONGRESS
4-7 SEPTEMBER 2011
Geneva

Industry Workshop Programmes



Industry Workshop Programmes

MONDAY 5 SEPTEMBER 2011

13:15 – 14:45

THERMO SCIENTIFIC

ROOM K

More Capabilities, New Possibilities in Proteomics Confirmation

13:15 - 14:00 **Applications of Recent Developments in Orbitrap Instrumentation to Quantitative Proteomics**
Speaker: Bruno Domon (Luxembourg)

14:00 - 14:45 **Decoding the Spatial and Temporal Regulation of Protein Interactions using Isobaric Tags**
Speaker: Ileana Cristea (USA)

For advanced registration for the Thermo Scientific workshop and to be kept informed of the latest Thermo Scientific plans for HUP0 2011 please access the following link: www.thermoscientific.com/hupo2011

PROTEINSIMPLE

ROOM E

Innovative Protein Characterization Using an Automated Platform for Both Size- and Charge-Based Detection

13:15 - 13:45 **A new view of proteins: Using a novel automated platform for detection and characterization of proteins**
Speaker: Peter Fung (USA)

13:45 - 14:15 **Analysis of RAF/MEK/ERK signaling following inhibition of the HSP90 molecular chaperone**
Speaker: Paul Andrew Clarke (UK)

14:15 - 14:45 **Fully automated size-based separation and Western blot analysis of proteins in capillaries based on NanoPro technology**
Speaker: Annegret Boge (USA)

BIO-RAD LABS

ROOM F

2-D Electrophoresis for Consistent, Meaningful results

13:15 - 13:45 **Experimental Planning for Consistent 2-D Results**
Speaker: Kathryn Lilley (UK)

13:45 - 14:15 **High reproducible 2-DE: Still a valuable tool for comparative protein profiling**
Speaker: Stefan Lehr (Germany)

14:15 - 14:45 **How 2-D Electrophoresis is beneficial to most Cell Protein Characterization in Bio pharmaceuticals**
Speaker: Martha Jackson (USA)

Industry Workshop Programmes

MONDAY 5 SEPTEMBER 2011

13:15 – 14:45

BECKMAN COULTER

ROOM G

Digging Deeper into the Proteome: Unleashing the Power of your Mass Spectrometer with CESI-MS

13:15 - 14:00 **Revealing Hidden Information in Complex Mixtures of Peptides through CESI-MS**
Speaker: Jim Thorn (UK)

14:00 - 14:45 **CESI-MS, A Promising Technique For The Identification Of Modified Peptides: Comparison With NanoLC-MS**
Speaker: Herbert Lindner (Austria)

To register, please email: CESI-EUROPE@BECKMAN.COM

AB SCIEX

ROOM L

Furthering the Human Proteome Project (HPP)

13:15 – 13:40 **An Introduction to the HPP and its Analytical Challenges**
Speaker: Pierre Legrain (France)

13:45 – 14:10 **Evaluation of new nanoLC/MS technologies for Targeted Proteomics**
Speaker: Sue Abatiello (USA)

14:15 – 14:40 **The HPP/Chromosome 19 Initiative**
Speaker: Juan Pablo Albar (Spain)

You are cordially invited to attend the AB Sciex workshops at HUP0 10th world Conference in Geneva, Switzerland.

Register now on <http://www.absciex/events/conferences/HUP02011>

After the workshops, you are invited to join us on Stand #1020 to learn more about our latest innovations, meet our experts or participate to one of our activities.

MONDAY 5 SEPTEMBER 2011

17:00 – 18:30

AGILENT TECHNOLOGIES

ROOM K

Meeting of the Minds

Agilent recognises the next generation of scientists is key to proteomics research and invites you to attend “**Meeting of the Minds**” on Monday evening, providing early career scientists the opportunity to meet and mix with today’s proteomics leaders. This is a rare opportunity for you to mingle and talk with many of the HUP0 keynote speakers and dignitaries in a relaxed and informal manner. Registration is essential to attend this event as numbers are limited, so be sure to [register](#) upfront today!

17:00 – 17:10 **Opening and Welcome Address**
Speaker: Gus Salem

Industry Workshop Programmes

17:10 – 18:30	Meeting of the Minds Meet the Proteomics Experts Rudolf Aebersold (Switzerland) Leigh Anderson (USA) Garry Corthals (Finland) Catherine Costello (USA) Ron Heeren (The Netherlands) Kathryn Lilley (UK) Ole Norregaard Jensen (Danmark) Gilbert Omenn (USA) Jesus V.Jorrin-Novo (Spain)
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MONDAY 5 SEPTEMBER 2011

GE HEALTHCARE

Accelerate your cancer research

17:00 – 18:30

ROOM E

Speaker: Jos Buijs (Sweden)

Speaker: Peter Gimeson (Sweden)

Speaker: Rita Marouga (Sweden)

AB SCIEX

New Technologies and Workflows in Peptide Quantitation

ROOM F

17:00 – 17:15	Development Challenges and Future Considerations for QTRAP Technology Speaker: Jim Hager (Canada)
17:20 – 17:45	A Quantitative Mass Spectrometry Assay for Detecting Tumor Specific Mutant Proteins Speaker: Keith Ashman (Spain)
17:50 – 18:05	Further Enhancing Peptide Quantitation Using Ultra Low flow CESI Speaker: Jeff Chapman (USA)
18:10 – 18:30	cHiPLC Workflows for MRM Quantitation in Proteomics Speaker: Lorne Taylor (Canada)

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Industry Workshop Programmes

TUESDAY 6 SEPTEMBER 2011**07:30 – 08:30**

AB SCIEX

ROOM F

MS/MSALL with SWATH Acquisition, a New Paradigm in Quantitative Proteomics

07:30 – 07:50 **Accelerating Biomarker and Proteomics Research with the TripleTOF 5600 System**
Speaker: Christie Hunter (USA)

07:55 – 08:25 **SWATH Acquisition and its Application to Quantitative Proteomics**
Speaker: Ruedi Aebersold (Switzerland)

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TUESDAY 6 SEPTEMBER 2011**13:15 – 14:45**

WATERS

ROOM K

Advanced in Mass Spectrometry for Quantitative Proteomics

Speakers: Kostas Petritis (USA)
Robert Tonge (UK)

TUESDAY 6 SEPTEMBER 2011**13:15 – 14:45**

LUMINEX

ROOM F

The next dimensions in multiplexing

13:15 – 13:35 **From Discovery to Validation: xMAP® Technology implementing Novel Biomarkers**
Speaker: F. Topin (The Netherlands)

13:35 – 13:55 **Suspension Bead Arrays as Tools in Proteomics**
Speaker: J. Schwenk (Sweden)

13:55 – 14:15 **Immunoassays in Multiplex for Biomarker Discovery and Validation**
Speaker: T. Joos (Germany)

14:15 – 14:35 **Autoantibody Biomarker Discovery And Validation: From Functional Protein Microarrays To Xmap Technology**
Speaker: Lisa Freeman-Cook (USA)

Industry Workshop Programmes

THERMO SCIENTIFIC

ROOM L

More Capabilities, New Possibilities in Comprehensive Proteomics

- 13:15 – 14:00 **Proteomics with a Quadrupole Orbitrap Hybrid Instrument**
Speaker: Annette Michalski (Germany)
- 14:00 – 14:45 **Towards Targeted Analysis of Kinase Signaling Networks**
Speaker: Claus Jorgensen (UK)

For advanced registration for the Thermo Scientific workshop and to be kept informed of the latest Thermo Scientific plans for HUP0 2011 please access the following link: www.thermoscientific.com/hupo2011

TUESDAY 6 SEPTEMBER 2011

17:00 – 18:30

BRUKER DALTONIK

ROOM K

Beyond Traditional Concepts - Proteomics goes Clinical

- 17:00 – 17:20 **Clinical Applications of MALDI Imaging**
Speaker: A. Walch (Germany)
- 17:20 – 17:40 **Gaining New Insights with Protein Glycosylation Studies**
Speaker: M. Wuhrer (The Netherlands)
- 17:40 – 18:00 **Next Generation Proteomics**
Speaker: Arnd Ingedoh (Germany)
- 18:00 – 18:20 **UHPLC and CaptiveSpray: The Total Proteomics Solution**
Speaker: Kerry Nugent

Poster Programme

Odd numbers: Monday 5 2011

Even numbers: Tuesday 6 2011

Odd and Even numbers: Wednesday 7 2011

Posters

MS QUANTITATION

P1000 - A SYSTEMATIC APPROACH TO ASSESS AMINO ACID CONVERSIONS IN SILAC EXPERIMENTS

Alberto Paradela

P1001 - ABSOLUTE QUANTIFICATION OF ALPHA-SYNUCLEIN PROTEIN IN A RAT MODEL OF PARKINSON'S DISEASE USING MRM

Adrien W Schmid

P1002 - INVERSE REGULATION IN THE METABOLIC GENES PCKA AND METE REVEALED BY PROTEOMIC ANALYSIS OF THE SALMONELLA RCSCDB REGULON

Alberto Paradela

P1004 - THREE BOTTLENECKS IN HPP IMPLEMENTATION

Alexander I Archakov

P1005 - IDENTIFICATION AND ABSOLUTE QUANTIFICATION OF A POTENTIAL THERMAL DEGRADATION MARKER OF PLASMA TRANSFUSION

Alexia Ortiz

P1006 - LOADQUANT: A PLATFORM FOR HANDLING QUANTITATIVE PROTEOMIC DATA OUTPUT FROM MATRIX SCIENCE'S MASCOT DISTILLER™

Allan Stensballe

P1007 - SI TRACEABLE QUANTIFICATION OF MYOGLOBIN FOR THE CHARACTERISATION OF A MYOGLOBIN CANDIDATE REFERENCE MATERIAL

Amalia Muñoz

P1008 - IDENTIFYING QUALITATIVE AND QUANTITATIVE DIFFERENCES IN RECOMBINANTS AND ANTIBODY ENRICHMENTS FOR IGF1 AS PROVIDED BY DIFFERENT VENDORS AND DIFFERENT PREPARATIONS USING HIGH RESOLUTION MASS SPECTROMETRY

Amol Prakash

P1009 - QUANTITATIVE ANALYSIS OF SKIN SUCTION BLISTER FLUID

Andre Mueller

P1010 - QUANTITATIVE ANALYSIS OF ENERGY METABOLIC PATHWAYS IN BREAST CANCER CELLS UNDER DIFFERENT GROWTH CONDITIONS BY SELECTED REACTION MONITORING

Andrei P Drabovich

P1011 - EVALUATION OF A LABEL-FREE QUANTITATIVE PROTEOMIC METHOD TOGETHER WITH SAMPLE FRACTIONATION FOR THE LARGE-SCALE ANALYSIS OF INFLAMMATORY ENDOTHELIAL CELLS

Anne Gonzalez De Peredo

P1012 - COMBINATION OF IRREVERSIBLE BINDING AND MRM FOR LOW- AND ULTRA-LOW COPIED PROTEINS DETECTION AND QUANTITATION

Artur Kopylov

Posters

- P1013 - **MS/MS-BASED QUANTITATIVE PROTEOMICS BY ISOBARIC PEPTIDE TERMINI LABELING**
Bernd Thiede
- P1014 - **CHARACTERIZATION OF CELLULAR PROTEASOME COMPLEXES DIVERSITY BY PROTEOMIC APPROACHES**
Bertrand Fabre
- P1015 - **PEPTIDE QUANTIFICATION BASED ON A NOVEL QUADRUPOLE-ORBITRAP MASS SPECTROMETER**
Bruno Domon
- P1016 - **OBVIATING TRANSITION OPTIMIZATION IN SRM ASSAYS TO DETECT AND QUANTIFY ANY PROTEIN OF THE HUMAN PROTEOME**
Caroline S. Chu
- P1017 - **INFLUENCE OF DIETARY COMPONENTS ON THE LEVELS OF REDUCTASES IN HUMAN COLORECTAL CANCER CELLS**
Cédric Bovet
- P1018 - **PROTEOME PROFILING AND QUANTIFICATION OF MOUSE LIPID DROPLET BY USING ¹³C LABELLED QUANTITATIVE PEPTIDE CONCATAMER**
Chen Ding
- P1019 - **QUANTITATION OF TYPE 1 DIABETES MHC PEPTIDE EPITOPES**
Chor Teck Tan
- P1020 - **ACCELERATING DIGESTION OF PROTEOMES USING TRYPSIN-IMMOBILIZED MAGNETIC NANOPARTICLES**
Christie Hunter
- P1021 - **HIGH SENSITIVITY PHOSHOPEPTIDE QUANTITATION USING NANOFLOW LC ON A TRIPLE QUADRUPOLE MASS SPECTROMETER**
Christine Miller
- P1022 - **MUDPIT-SRM – A HIGHLY MULTIPLEXED STRATEGY FOR LOW ABUNDANT PROTEIN BIOMARKER IDENTIFICATION**
Christoph Krisp
- P1023 - **MRMAID2: MINING PRIDE FOR EVIDENCE-BASED SRM TRANSITIONS**
Conrad Bessant
- P1024 - **MULTIPLEXED ABSOLUTE QUANTIFICATION OF PROTEINS DURING EARLY EMBRYOGENESIS BY SELECTED REACTION MONITORING**
Daniela Deutsch
- P1025 - **G-PROTEIN COUPLED RECEPTOR QUANTIFICATION USING PEPTIDE-GROUP SPECIFIC ENRICHMENT COMBINED WITH INTERNAL PEPTIDE STANDARD ITRAQ REPORTER CALIBRATION**
David Eisen

Posters

- P1026 - **MZDB: AN OPTIMIZED FILE FORMAT FOR THE EFFICIENT ANALYSIS OF LC-MS DATASETS**
David Bouyssié
- P1027 - **QUANTITATIVE ANALYSIS OF LARGE PEPTIDES USING PSEUDO-SELECTED REACTION MONITORING WITH ELECTRON TRANSFER DISSOCIATION**
Didia Coelho Graça
- P1028 - **COMPLETING THE RHODOPSIN MACHINE - CO-IMMUNOPRECIPITATION AND ICPL-TECHNOLOGY AS PROMISING TOOLS**
Eva-Maria Keidel
- P1029 - **EASYPROT: A PLATFORM FOR MS1/MS2 PROTEIN IDENTIFICATION AND LABEL-FREE QUANTIFICATION**
Florent Gluck
- P1030 - **GENERAL ANALYSIS OF PROTEIN RELATIVE EXPRESSION DATA FROM ISOBARIC TAGGING EXPERIMENTS**
Florian P Breitwieser
- P1031 - **MULTI-SITE ASSESSMENT OF PROTEOED PLASMA REFERENCE SAMPLE FOR BENCHMARKING LC-MS PLATFORM PERFORMANCE**
Francesc Canals
- P1032 - **BIOMARKER VALIDATION FOR COLORECTAL CANCER SCREENING USING SRM AND MRM3 AND CORRELATION WITH ELISA TEST**
Genevieve Choquet
- P1033 - **DETECTING AND AVOIDING REDUNDANCY IN SELECTED REACTION MONITORING (SRM) - A COMPUTATIONAL APPROACH TO DESIGN SELECTIVE SRM ASSAYS**
Hannes Luc Röst
- P1034 - **NEXT GENERATION QUANTITATIVE PROTEOMICS: OVERCOMING THE NEED FOR NON-OVERLAPPING ISOTOPE PATTERNS AND ALLOWING FOR NOVEL MULTIPLEXED LABELING STRATEGIES**
Hanno Steen
- P1035 - **A ROBUST QUANTITATION PLATFORM FOR HIGH THROUGHPUT LABEL-FREE SHOTGUN PROTEOMICS TECHNIQUES BY LC-MS/MS ANALYSIS**
Hee-Jung Jung
- P1036 - **TARGETED LABEL-FREE QUANTIFICATION IN A 3D ION TRAP INSTRUMENT**
Huisong Pak
- P1037 - **MULTIPLE REACTION MONITORING OF PHOSHOPEPTIDE LABELED WITH 6-PLEX TANDEM MASS TAGS(TMT) FOR QUANTITATIVE PHOSPHOPROTEOMICS**
Hye Kyeong Min

Posters

- P1038 - **PROTEOMIC ANALYSIS OF THE CAENORHABDITIS ELEGANS SIR-2.1 PROTEIN DEACETYLASE DEFICIENT STRAIN**
Hyoung Joo Lee
- P1039 - **INVESTIGATING THE RELATIONSHIP BETWEEN SAMPLE COMPLEXITY AND ACCURATE ASSESSMENT OF QUANTITATIVE CHANGE - HOW UNIQUELY CAN AN ION BE MEASURED?**
James Langridge
- P1040 - **HIGH RESOLUTION PEPTIDE ISOELECTRIC FOCUSING PREFRACTIONATION (HRIEF-LC-MS/MS) ENABLES COMPREHENSIVE HUMAN PROTEOMICS**
Janne Lehtiö
- P1043 - **DIFFERENTIAL ANALYSIS OF MEMBRANE PROTEOMES FROM CLOSTRIDIUM ACETOBUTYLICUM AT ACIDOGENESIS AND SOLVENTOGENESIS USING ITRAQ**
Jingjing Zhao
- P1044 - **ESTABLISHMENT OF QUALIFICATION PROCESS FOR 158 BIOMARKER CANDIDATES IN CYST FLUID USING HIGH THROUGHPUT MS-BASED PROTEOMIC APPROACH**
Jisook Park
- P1045 - **ANALYSIS OF DISEASE CAUSING MUTATIONS BY MASS SPECTROMETRY**
Johan Palmfeldt
- P1046 - **ACCURATE QUANTIFICATION OF PREDICTED MICRORNA TARGETS IN MOUSE HEART USING HIGH-RESOLUTION TARGETED PROTEOMICS**
Johannes A Hewel
- P1047 - **THE USE OF A PEPTIDE RETENTION-TIME CALIBRATION MIXTURE AS A TOOL IN SCHEDULED SELECTED REACTION MONITORING METHOD CONSTRUCTION**
John C. Rogers
- P1048 - **UNDERSTANDING THE ROLE OF TRK RECEPTORS IN NEUROBLASTOMA USING QUANTITATIVE PHOSPHOPROTEOMICS APPROACHES**
John C. Rogers
- P1049 - **LABEL-FREE QUANTITATION OF A UNIQUE PROSTATE CANCER PROGRESSION MODEL USING LC-FTICR-MS**
John Lapek
- P1050 - **QUANTIFYING THE EFFECTS OF IONIZING RADIATION AND RADIOPROTECTORS ON IMMUNE CELLS USING OFFGELC-MSE**
Jonathan Erde
- P1051 - **ABSOLUTE QUANTIFICATION OF A LARGE SET OF TRANSCRIPTION FACTORS INVOLVED IN THE ADIPOGENIC GENE REGULATORY NETWORK VIA A MASS SPECTROMETRY-BASED APPROACH**
Jovan Simicevic

Posters

P1052 - **IMPACT OF HIGH RESOLUTION MRM ANALYSIS ON COMPLEX PROTEOMES**

Kathleen Lewis Torpey

P1053 - **ANALYSIS/QUANTIFICATION OF DIFFERENT FORMS OF C-REACTIVE PROTEIN (CRP) IN REFERENCE MATERIALS**

Katja Hanisch

P1054 - **SELECTIVE QUANTIFICATION OF NEWLY SYNTHESIZED PROTEINS IN LPS-STIMULATED MACROPHAGES**

Katrin Schuschke

P1055 - **PRECISE MEASUREMENTS OF PROTEIN ABUNDANCE IN SACCHAROMYCES CERREVISIAE; COMPARING LC-MRM TO LABEL-FREE LC-MS**

Kevin T Collins

P1056 - **IN VIVO QUANTIFICATION OF LIVER SIGNALING AT THE SYSTEMS LEVEL**

Kirti Sharma

P1057 - **SIMULTANEOUS MEASUREMENT OF PLASMA APOLIPOPROTEINS BY MRM AND MTRAQ® LABELING FOR CANCER DIAGNOSTICS**

Kwang Youl Kim

P1058 - **SIMULTANEOUS MEASUREMENT OF PLASMA APOLIPOPROTEINS BY MRM AND MTRAQ LABELING FOR CANCER DIAGNOSTICS**

Kwang Youl Kim

P1059 - **SERUM ANALYSIS BASED ON MULTIPLE REACTION MONITORING USING LC/MS/MS FOR THE DETECTION OF PANCREATIC CANCER**

Kyoung-Jin Park

P1060 - **TOWARDS QUANTIFYING PROTEIN-PROTEIN INTERACTIONS USING SYNCHROTRON-BASED OXIDATIVE FOOTPRINTING AND SELECTED REACTION MONITORING**

Lars Malmström

P1061 - **AUTOMATIC AND ADAPTIVE PROCESSING ALGORITHM OF SRM TRACES FOR PROTEIN QUANTITATION**

Laurent Gerfault

P1062 - **DIFFERENTIAL ANALYSIS OF GLIOBLASTOMA MULTIFORME TISSUES IMPLICATES AN INTRIGUING SIGNALING PATHWAY**

Leroi V. Desouza

P1063 - **SWATH-MS: DATA INDEPENDENT CAPTURE OF MS/MS SEQUENCES FOR QUANTIFICATION OF PEPTIDE KINETICS IN A CELL SIGNALLING CASCADE**

Lorne Taylor

Posters

- P1064 - **MULTIPLEXING SIMS ON A NOVEL BENCH TOP ORBITRAP WITH A QUADRUPOLE MASS FILTER TO INCREASE SENSITIVITY FOR PEPTIDE QUANTITATION**
Louis Maljers
- P1065 - **SWATH MS: A METHOD FOR COMPLETE SAMPLE CONTENT ANALYSIS COMBINING A DATA INDEPENDENT ACQUISITION METHODOLOGY AND A TARGETED DATA ANALYSIS STRATEGY**
Ludovic CJ Gillet
- P1066 - **ISOBARIQ: SOFTWARE FOR ISOBARIC QUANTITATIVE PROTEOMICS USING IPTL, ITRAQ AND TMT**
Magnus Øverlie Arntzen
- P1067 - **A ROBUST LABEL-FREE ANALYSIS WORKFLOW FOR HIGH-RESOLUTION MASS SPECTROMETRY**
Marianne Sandin
- P1068 - **ANALYSIS OF IONIZING RADIATION-INDUCED MODULATION OF THE CELLULAR PROTEOME: USING QUANTITATIVE PROTEIN PROFILING BY SILAC AND SRM**
Mariette Matondo
- P1069 - **PREST-SILAC COMBINATION ENABLES ABSOLUTE QUANTIFICATION AND MULTIPLEXED DETERMINATION OF PROTEIN COPY NUMBERS IN CELL LINES**
Marlis Zeiler
- P1070 - **DEVELOPMENT OF AN MRM ASSAY PANEL FOR CARDIOVASCULAR DISEASES**
Melinda Rezeli
- P1071 - **OPTIMIZATION OF EXPERIMENTAL PARAMETERS FOR UNBIASED LABEL-FREE QUANTITATIVE MASS SPECTROMETRY OF PROTEINS**
Michael Groessel
- P1072 - **SPECIFICITY AND COMMONALITY OF THE PHOSPHOINOSITIDE-BINDING PROTEOME ANALYZED BY QUANTITATIVE MASS SPECTROMETRY**
Michael L. Nielsen
- P1073 - **ACCURATE QUANTIFICATION OF PROTEINS IN COMPLEX MATRICES FOR THE PRODUCTION OF MATRIX CERTIFIED REFERENCE MATERIALS**
Milena Quaglia
- P1074 - **QUANTITATIVE PROTEOMICS OF YEAST CELLS IN SINGLE LC MS/MS RUNS USING A BENCH-TOP QUADRUPOLE ORBITRAP**
Nagarjuna Nagaraj
- P1075 - **SYSTEMATIC QUANTIFICATION OF URINARY PROTEINS USING SELECTED REACTION MONITORING**
Nathalie Selevsek
- P1076 - **BIOINFORMATICS SUPPORT AT THE EARLY STAGES OF SRM ASSAY DEVELOPMENT**
Nikolaos Berntenis

Posters

P1077 - **EVALUATION OF PEPTIDE ALKYLATION SIDE-REACTIONS IN COMMON PROTEIN DIGESTION PROTOCOLS**

Núria Colomé

P1078 - **VALIDATION OF A MASS SPECTROMETRIC METHOD FOR SERUM HEPcidIN**

Outi Itkonen

P1079 - **FN1, TIMP1, AND SERPINA1 REGULATE THE MIGRATION, INVASION, AND PERICELLULAR FIBRONECTIN ASSEMBLY OF CANCER CELLS IN METASTASIS: A NOVEL BIOLOGICAL SIGNIFICANCE REVEALED BY QUANTITATIVE PROTEOMICS ANALYSIS OF NSCLC CELL SECRETOMES**

Pao-Chi Liao

P1080 - **HIGH-THROUGHPUT SRM-LIKE ANALYSIS OF DATA GENERATED UPON SWATH MS DATA INDEPENDENT ACQUISITION**

Pedro Navarro

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- P1852 - **PROTEOMIC ANALYSIS OF HUMAN HEPATOCELLULAR CARCINOMA IN VIETNAM**
Trinh Hong Thai
- P1853 - **A PILOT STUDY FOR THE USE OF RANDOM PEPTIDE ARRAYS FOR BIOMARKER DISCOVERY IN BREAST CANCER**
Ulf Reimer
- P1854 - **DISCOVERY AND VERIFICATION OF CNDP1 AS A PROSTATE CANCER BIOMARKER CANDIDATE IN PLASMA**
Ulrika Margareta Igel
- P1855 - **VERIFICATION OF BIOMARKERS FOR BREAST CANCER DIAGNOSIS BY MULTIPLE REACTION MONITORING**
Un-Beom Kang
- P1856 - **INVESTIGATION OF POTENTIAL BIOMARKERS OF SOCIALLY-SIGNIFICANT HUMAN DISEASES IN SERUM PEPTIDOME**
Vadim M Govorun
- P1857 - **SINGLE CELL MALDI-TOF MASS SPECTROMETRY ANALYSIS: MAMMALIAN OOCYTE PROFILING COULD REFLECT ITS QUALITY**
Valérie Labas
- P1858 - **QUANTITATIVE PROTEOMIC (ITRAQ) ANALYSIS OF 1ST TRIMESTER MATERNAL PLASMA SAMPLES IN PREGNANCIES AT RISK FOR PREECLAMPSIA**
Varaprasad Kolla
- P1859 - **TOWARDS EARLY MELANOMA BIOMARKERS: SYSTEMATIC SECRETOME PROFILING OF HUMAN TISSUE SAMPLES AND PRIMARY STROMA AND TUMOR CELLS**
Verena Paulitschke
- P1860 - **THE HUMAN INTER-INDIVIDUAL URINARY PROTEOME ANALYSIS DEFINING THE MINIMAL SAMPLE SIZE FOR CLINICAL PROTEOMIC RESEARCH**
Wei Sun
- P1861 - **IDENTIFICATION OF TBRG4 AS A NOVEL ONCOGENE IN BREAST CARCINOMA**
Wei Zhang
- P1862 - **POTENTIAL URINARY BIOMARKERS FOR PREDICTING THE CLINICAL OUTCOME OF REFLUX NEPHROPATHY IN CHILDREN**
Wei-Jung Chen

Posters

P1863 - IDENTIFICATION OF A NEW SET OF IMMUNOHISTOCHEMICAL BIOMARKERS FOR CLASSIFICATION OF DYSPLASTIC NODULES AND SMALL HEPATOCELLULAR CARCINOMA

Yan Li

P1864 - TOWARDS QUANTITATIVE ANALYSIS OF HUMAN SWEAT PROTEOME

Yohann Couté

P1865 - BIOMARKER DISCOVERY OF HIV-RELATED FATIGUE

Young Ah Goo

P1866 - DIFFERENTIAL PROTEOME PROFILING USING ITRAQ IN MICROALBUMINURIC AND NORMOALBUMINURIC TYPE 2 DIABETIC PATIENTS

Youngsoo Kim

P1867 - GLYCOPROTEINS ASSOCIATED WITH DIFFERENT HISTOLOGICAL SUBTYPES OF OVARIAN TUMORS

Yuan Tian

P1868 - HIGH-SPEED MEMBRANE PROTEOMIC SIGNATURE BY INFORMATICS-ASSISTED LABEL-FREE QUANTITATION FOR CANCER BIOMARKER DISCOVERY AND VERIFICATION

Yu-Ju Chen

P1869 - DEVELOPMENT OF THE PROTEASE IMMOBILIZED MICRO-FLOW CHIP FOR THE PROTEIN STRUCTURAL ANALYSES USING MASS SPECTROMETRY

Yukie Sasakura

P1870 - CIRCULATING BIOMARKER-IGM COMPLEXES IN THE SERUM OF PEDIATRIC PATIENTS WITH OSTEOSARCOMA AND EWING'S TUMOR

Yulia Savitskaya

CENTRAL NERVOUS SYSTEM

P1871 - PROTEOMICS OF HUMAN MYELIN AND AXOGLIAL-APPARATUS: IMPLICATION OF PERTURBED AXOGLIAL-APPARATUS IN PEDIATRIC ONSET MULTIPLE SCLEROSIS

Ajit Singh Dhaunchak

P1872 - IMMUNOPROTEOMIC PROFILING OF BRAIN STEROID 5 ALPHA REDUCTASES IN SLEEP-DEPRIVED RATS

Alessio Soggiu

P1873 - MRM QUANTITATION OF JC VIRUS IN BRAIN TISSUE FOR A CASE OF PROGRESSIVE MULTIFOCAL LEUKOENCEPHALOPATHY (PML) CAUSED BY SELECTIVE IMMUNE DEFICIENCY

Alexander Buko

P1874 - QUANTITATIVE ANALYSIS OF MYELIN BASIC PROTEIN DEGRADATION USING 18O-ISOTOPE LABELING

Alexey Kononikhin

Posters

- P1875 - IDENTIFICATION OF SYNUCLEINS AND OTHER SYNAPSE-ASSOCIATED PROTEINS IN ZEBRAFISH AND HUMAN BRAIN TISSUE
Ann Brinkmalm
- P1876 - INDUCTION OF NEURONAL DEATH BY MICROGLIAL AGE-ALBUMIN: IMPLICATIONS FOR ALZHEIMER'S DISEASE
Bonghee Lee
- P1877 - PROTEOME ANALYSIS OF POST-TRANSPLANTATION RECOVERY MECHANISMS OF EAE MODEL OF MULTIPLE SCLEROSIS TREATED WITH STEM CELL-DERIVED NEURAL PRECURSORS
Davood Bagher Nasrabadi
- P1878 - EVOLUTIONARY AND STRUCTURAL RELATIONSHIPS OF IONOTROPIC RECEPTORS NMDA, AMPA AND KAINATE IN FOUR SPECIES OF PRIMATES
Edgar Antonio Reyes-Montaña
- P1879 - ALTERED PROTEIN TRAFFICKING QUANTIFIED IN THE SYNAPSE OF SCHIZOPHRENIC POST-MORTEM BRAIN USING LC-SRM/MS AND A LABELED MOUSE BRAIN STANDARD
Eugene Ciccimaro
- P1880 - MASS SPECTROMETRIC CHARACTERIZATION OF AMYLOID ISOFORMS FROM CAT CEREBROSPINAL FLUID. DETERMINATION OF THE CAT AMYLOID SEQUENCE
Gunnar Brinkmalm
- P1881 - IDENTIFY THE PROTEINS ARE INVOLVED IN PARKINSON'S DISEASE USING PROTEOMICS
Jin Sup Sin
- P1882 - PROTEOMIC IDENTIFICATION OF DIFFERENTIALLY EXPRESSED GENES DURING THE DIFFERENTIATION FROM EMBRYONIC STEM CELLS OF CYNOMOLGUS MONKEY (MACACA FASCICULARIS) TO NEURAL STEM CELLS IN VITRO
Kuniko Akama
- P1883 - SYSTEMS BIOLOGY APPROACHES TOWARDS HUMAN BRAIN DISEASE: CATHEPSIN D DEFICIENCY
Maciej Maurycy Lalowski
- P1884 - EXERCISE EFFECTS ON THE HIPPOCAMPUS PROTEOME IN BRAIN AGING: PUTATIVE BIOMARKERS
Marilisa Sulpizio
- P1885 - DISTRIBUTION AND FUNCTION RESEARCH OF VITAMIN D BINDING PROTEIN IN EXPERIMENTAL AUTOIMMUNE ENCEPHALOMYELITIS
Shilian Liu
- P1886 - CNDP2 OVEREXPRESSION IN NIGRAL TISSUE OF PARKINSON'S DISEASE PATIENTS
Virginie Licker

Posters

POSTERS SUBMITTED FOR HUP0 INITIATIVES

P1887 - PROTEOME CHARACTERISTICS OF HUMAN WHARTON'S JELLY MESENCHYMAL STEM CELLS BY SDS-PAGE COMBINED WITH NANOLC-MS/MS ANALYSIS

Chang-Chun Hsiao

P1888 - CRYOPRESERVATION EFFECTS ON WJCS PROTEOME

Fabrizio Di Giuseppe

P1889 - STATE OF BINDING SITES OF SERUM ALBUMIN PREGNANT WOMEN IN HEALTH AND DISEASE

Natallia Dmitrievna Lukouskaya

P1890 - POSITIONAL PROTEOMICS OF THE WING IMAGINAL DISC

Nikiana Simigdala

P1891 - PROTEOMIC PROFILING OF MOUSE LIVER CELL SYSTEM

Ying Jiang

POSTERS SUBMITTED FOR CLINICAL DAY

P1892 - SIMULTANEOUS DETECTION OF AUTOANTIBODIES AGAINST P53 AND A P53 PEPTIDE IN THE SERA OF WOMEN WITH OVARIAN CANCER

Steven Binder

POSTERS SUBMITTED FOR EUPA DAY

P1893 - PROTEOME ANALYSIS OF CANOLA SEEDLING ROOTS UNDER SALT STRESS

Ali Bandehagh

P1894 - DIFFERENTIAL PROTEOME ANALYSIS OF AN ALKALIPHILIC BACTERIUM, EXIGUOBACTERIUM SP. FOR STUDY OF CELLULAR RESPONSES TO ALKALINE CONDITIONS

Niha Mohan Kulshreshtha

OTHER POSTERS

P1895 - THE IMPACT OF INDIVIDUAL GENOME ON PROTEIN IDENTIFICATION WITH SHOTGUN MASS SPECTROMETRY DATA

Emøke Bendixen

P1896 - DISULFIDE MAPPING USING ACCURATE MASS ETD AND MSN TECHNIQUES

Iman Mohtashemi

Posters

P1897 - THE CHROMOSOME 19 CONSORTIUM- A GLOBAL RESEARCH INITIATIVE

Juan Pablo Albar

P1898 - PROTEORED MIAPE WEB TOOLKIT: SHARING PROTEOMICS EXPERIMENTS WITHIN THE COMING PROTEOMEXCHANGE CONSORTIUM

Salvador Martinez-Bartolome

General Information

Congress Registration Fees

Monday 5 September 2011			
	Up to 16 May 2011	After 16 May 2011	Onsite (as of 3 September 2011)
HUPO Member	CHF 825	CHF 975	CHF 1'175
Student Members*	CHF 320	CHF 390	CHF 470
HUPO Non-member	CHF 955	CHF 1'105	CHF 1'305
Student Non-member *	CHF 400	CHF 470	CHF 550
Pre-Congress	CHF 300	CHF 400	CHF 500
Pre-Congress Student *	CHF 110	CHF 120	CHF 140
Accompanying Person	CHF 225	CHF 225	CHF 225
Exhibitor	CHF 325	CHF 325	CHF 325

*For student registrations, an email from your mentor or the head of department is required including name of the student, name of the department head/mentor, affiliation, and registration number, must be submitted during the registration process.

HUPO MEMBERSHIP

If you are not yet a member of the HUPO, you can become a member by choosing the following options when registering:

- I am not a HUPO member but I wish to apply at no cost for 2011

You will pay the "Non-member" fee which will include your HUPO membership participation for 2011.

If your membership status is "expired" you may choose to renew it by selecting:

- My HUPO membership is expired and I wish to renew it now

You will pay the "Non-member" fee which will re-activate HUPO membership participation for 2011.

As an "active" member you will also be allowed to renew your membership for another year by choosing:

- My HUPO membership is active; renew for another year

You will pay the "Non-member" fee which will re-activate HUPO membership participation for 2012.

For all other questions related to HUPO Membership, please go to the HUPO website: www.hupo.com or visit HUPO at their stand N° XX

Delegates' registration fee includes:

- All congress sessions from Sunday 4 September to Wednesday 7 September 2011 (E&T sessions not included)
- Entrance to the Opening Ceremony on Sunday 4 September 2011
- Welcome Reception in the Exhibition on Sunday 4 September 2011
- Circus Show on Tuesday 6 September 2011
- Lunches from Monday 5 September to Wednesday 7 September 2011
- Morning and afternoon coffee breaks from Monday 5 September to Wednesday 7 September 2011
- Entrance to the Exhibition from Monday 5 September to Wednesday 7 September 2011
- Biodegradable congress bag
- Congress material including USB key with abstracts

General Information

Accompanying persons' registration fee includes:

- Entrance to the Opening Ceremony on Sunday 4 September 2011
- Welcome Reception in the Exhibition on Sunday 4 September 2011
- Circus Show on Tuesday 6 September 2011
- Lunches from Monday 5 September to Wednesday 7 September 2011
- Morning and afternoon coffee breaks from Monday 5 September to Wednesday 7 September 2011
- Entrance to the Exhibition from Monday 5 September to Wednesday 7 September 2011
- City Tour of Geneva
- Biodegradable congress bag
- Congress material including USB key with abstracts

Pre-Congress Day

Registration for the Pre-Congress Day was on a reservation basis and was subject to a fee:

	Up to 16 May 2011	After 16 May 2011	Onsite (as of 3 September 2011)
Pre-Congress	CHF 300	CHF 400	CHF 500
Pre-Congress Student *	CHF 110	CHF 120	CHF 140

Registered participants will receive a voucher for the session they have reserved and paid for to grant them access. The morning coffee break and lunch are included in the fees and will be served in the foyer of Rooms ABC.

Hotel and Registration Information

MCI Suisse SA has reserved a large number of rooms in different hotel categories and offers discounted room rates for the Congress participants. Only reservations made through MCI Suisse SA will benefit from these rates. All these hotels are located in the City Centre, or near the Congress Centre.

LIST AND CITY MAP

HOTELS IN GENEVA (SWISS FRANCS)

HOTELS 5*	LOCATION	DISTANCE TO PALEXPO
BEAU RIVAGE	1	8' driving distance
DE LA PAIX	2	10' driving distance
INTERCONTINENTAL	3	5' driving distance
MANDARIN ORIENTAL	4	10' driving distance
STARLING	5	3' walking distance

HOTELS 4* SUP	LOCATION	DISTANCE TO PALEXPO
BRISTOL	6	10' driving distance
EPSOM	7	10' driving distance
ROYAL	8	10' driving distance

HOTELS 4*	LOCATION	DISTANCE TO PALEXPO
AUTEUIL	9	10' driving distance
CORNAVIN	10	10' driving distance
LES NATIONS	11	4' driving distance
NH GENEVA AIRPORT	12	7' driving distance
NH REX	13	7' driving distance
NOVOTEL GENEVE CENTRE	14	10' driving distance
CROWNE PLAZA (former Ramada Park)	15	4' driving distance
WARWICK	16	8' driving distance

HOTELS 3* SUP	LOCATION	DISTANCE TO PALEXPO
EDELWEISS	17	10' driving distance
JADE	18	10' driving distance
KIPLING	19	10' driving distance

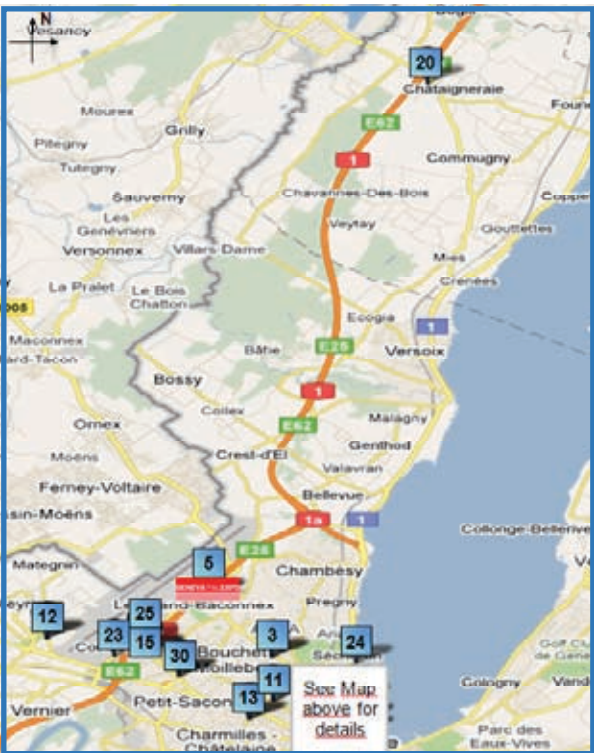
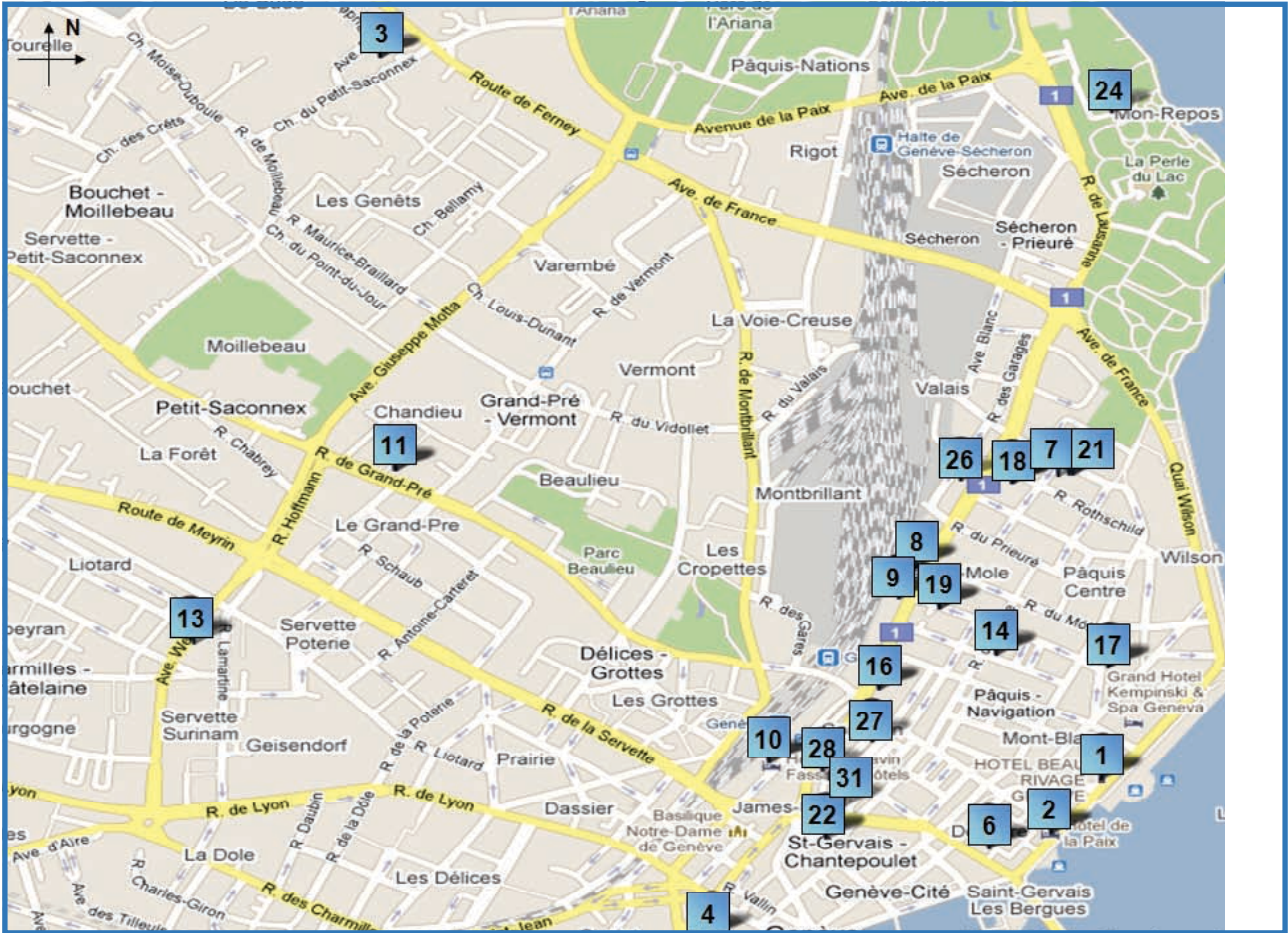
HOTELS 3*	LOCATION	DISTANCE TO PALEXPO
CHAVANNES DE BOGIS	20	20' driving distance
DRAKE & LONGCHAMP	21	8' driving distance
EXCELSIOR	22	10' driving distance
EXPRESS BY HOLIDAY INN	23	3' driving distance
MON REPOS	24	7' driving distance
NASH AIRPORT	25	4' driving distance
DESIGN HOTEL (Former Quality Hotel Geneva)	26	8' driving distance
STRASBOURG-UNIVERS	27	9' driving distance
SUISSE	28	8' driving distance

HOTELS 2* SUP	LOCATION	DISTANCE TO PALEXPO
ETAP **	30	5' driving distance
LIDO	31	9' driving distance

Hotel and Registration Information

LIST AND CITY MAP

HOTELS IN GENEVA (SWISS FRANCS)



Hotel and Registration Information



FOR ANY QUESTION, KINDLY CONTACT:

HUP0 2011

c/o MCI

Registration + Hotel Accommodation



Tel: + 41 22 33 99 578

Fax: + 41 22 33 99 621

mci

E-mail: hupo2011reg@mci-group.com

Website: www.hupo2011.com

RESERVATION AND PAYMENTS

Only guaranteed reservations will be accepted. To guarantee your reservation a deposit covering one night is required to be paid to MCI Suisse SA either by bank transfer or by credit card (deposits will be deducted from the final hotel bill when paid by the participant at the hotel). Please indicate the participant's name and "HUP0 2011" on ALL payments.

INQUIRIES

Any inquiries or requests for additional information, changes or cancellations of room reservations should be addressed directly to MCI by email or in writing.

GROUP RESERVATIONS

Group reservations (from 10 rooms) will be handled by MCI Suisse SA with separate contracts. Please contact MCI for further information.

CANCELLATION AND MODIFICATION CONDITIONS

Registration:

All cancellations must be sent to MCI Suisse S.A. in writing (fax, letter, or e-mail). For cancellations received before **July 6th, 2011**, fees will be refunded less **CHF 100.-** for administrative costs. After this date, no refund will be possible.

A handling fee of **CHF. 60.-** per registration will be charged for every registration modification received after **July 6th, 2011**.

SOCIAL EVENTS AND EXCURSIONS:

Cancellations can be made free of charge until August 3rd, 2011. After this date, no refund is possible.

MCI reserves the right to cancel or modify the excursion program if the minimum participation is not reached or due to factors beyond the control of the organizers.

HOTELS

To guarantee the reservation a deposit covering the first night of your stay is required. From **July 6th, 2011** only payments made by credit card will be accepted. All cancellations must be sent to MCI Suisse S.A. in writing (fax, letter, or e-mail). For cancellations received before July 6th, 2011 deposits will be refunded less **CHF 100.-** for administrative costs. After this date, no refund will be possible.

In case of late cancellation (after July 6th, 2011), rooms cancelled, cancelled nights or no-show, we reserve the right to charge the full stay if the hotel room cannot be re-sold.

A handling fee of **CHF. 60.-** per hotel room will be charged for every hotel modification received after **July 6th, 2011**.

Hotel and Registration Information

CHECK-IN/CHECK-OUT TIME

Hotel check-in time is 14:00 or later. If you are arriving after 18:00, please notify MCI Suisse SA in advance. Check-out time is noon.

DISCLAIMER

MCI Suisse SA as the Congress planner, HUP0 and committee members and officers of the 10th Congress of the HUP0 claim no liability for the act of any supplier to this meeting, nor liability for: personal injury, the safety of any attendee while in transit to or from this event, for any loss or damage, for delays in transport by air, sea, rail, road, weather, in case of strikes, sickness, war or other causes.

METHODS OF PAYMENT

Payment by personal checks cannot be accepted.

All payments are in Swiss Francs (CHF). For hotel bookings, a deposit payment for the first night must accompany the hotel accommodation form. Additional expenses such as mini-bar, laundry, etc. must be paid directly to the hotel when checking out.

From **06 July**, only payments made by credit card will be accepted.

General Congress Information

CONGRESS INFORMATION

ABSTRACTS ON USB KEY

The HUP0 2011 Abstracts will be available on a USB key. Participants will receive a voucher which will allow them to collect the key at the Agilent exhibition stand (stand N° 1340) during the exhibition opening hours.

BADGES

Name badges must be visible and used at all times, anywhere at the Congress Centre.

IDENTIFICATION:

No colour:	Participant badge (all access)
Gold colour:	Committees & Board
Blue colour:	Speakers & Chairs
Green colour:	Exhibitor
Red colour:	Accompanying Person

BANKS & AUTOMATIC TELLER MACHINES (ATM)

There is an ATM located at Palexpo. It is located near the entrance foyer of the Congress Centre. Please ask the Registration Desk for further information.

CERTIFICATE OF ATTENDANCE

Certificate of Attendance can be collected at the Registration Area (Self-Badge Printing desk) as of Tuesday, 6 September 2011, morning.

BIODEGRADABLE CONGRESS BAG

Participants will receive a congress bag voucher which will allow them to collect the bag with all the congress documents at the Registration Area (Congress Bags desk).

CONGRESS SECRETARIAT

HUP0 2011

c/o MCI

Rue de Lyon 75

CH-1211 Geneva 13

Switzerland

Tel: + 41 22 33 99 578

Fax: + 41 22 33 99 621

E-mail: hupo2011@mci-group.com

Website: www.hupo2011.com

General Congress Information

CONGRESS SPECIAL ISSUE

The Scientific Committee is pleased to announce that a Special Issue containing selected papers from the Congress entitled Translational Proteomics will be published in the Journal of Proteomics. All manuscripts will be refereed according to the scientific standards of the journal. Pierre Lescuyer, Laura Bianchi, Denis Hochstrasser, Luca Bini and Jean-Charles Sanchez will be acting as Guest Editors.

Those wishing to submit a full-length article or technical reports for publication in the Special Issue, will have the option to do so online, here, as from **15 September 2011**. Please note that the deadline for submission of full-length articles for the Special Issue of the Journal of Proteomics is **15 November 2011**.

It is anticipated that the Special Issue will be published within 6 months from the date of the Congress.

INSTRUCTIONS FOR SUBMISSION:

- Manuscripts must be submitted electronically through the Journal of Proteomics submission website at: <http://ees.elsevier.com/jprot>
- The covering letter of the article should be addressed to the Guest Editors
- To ensure that all manuscripts are correctly identified for inclusion into the HUP0-2011 special issue, authors must select "SI: Translational Proteomics" when they reach the "Article Type" step in the submission process.
- Please consult all of the instructions in the Author Information section before beginning your submission.

DUPLICATION / RECORDING

Any form of duplication (recording, audiotaping, digital taping, photography and video) is prohibited in the sessions. Permission to do so should be sought from the Congress Organisers.



General Congress Information

EXHIBITION AREA INFORMATION

The HUP0 2011 Exhibition, featuring commercial displays of International Organisations, Pharmaceutical Companies, Media Publishers and Scientific Societies, will be located in Hall 1 of the Congress Centre. The Registration Desk, catering facilities and Poster Sessions will also be located in the Exhibition hall.

EXHIBITION AREA SCHEDULE

Sunday 04 September	16:00-21:00
Monday 05 September	08:00-17:00
Tuesday, 6 September	08:00-17:00
Wednesday, 7 September	08:00 -15:30

LUNCHES AND COFFEE BREAKS

Lunches and coffee breaks are included in the Registration Fee.

They will be served in the Exhibition Hall during the Exhibition opening hours from Monday 5 to Tuesday 6 September.

On Sunday, 4 September the morning coffee break and the lunch will be served in the Foyer of Rooms ABC and on Wednesday, 7 September, the morning coffee break and the lunch will be served in the Exhibition Hall.

A cash bar will be available inside the Exhibition during the Exhibition opening hours.

INTERNET

Free WIFI is provided in the Exhibition Hall and in front of rooms ABC. Please refer to the Registration Desk for more information. A Cyber Cafés is situated inside the Exhibition.

HUP0 OFFICE

The HUP0 Office is located in Office 3 on the mezzanine leading to rooms D-E-F-G.

FUTURE MEETINGS

A Future Meetings table is located in the Exhibition Area.

LANGUAGE

English is the official Congress language.

OFFICIAL CARRIER

Swiss International Air Lines is the official carrier for the HUP0 2011 Congress and is offering special Congress fares to all participants.



PARKING

There is a public parking at Palexpo, P12. It is open and accessible 24 h.

Cost: CHF 1.50 per hour, CHF 15 per day

General Congress Information

POSTER HELP DESK

There is a Poster Help Desk in the Poster Area. The Poster Help Desk has material to affix posters.

POSTER SESSIONS

Posters will be displayed for the duration of the poster session. Poster presenters have been scheduled to be by their poster according to the programme times. Odd numbers will be presented on Monday 5 September, even numbers will be presented on Tuesday 6 September. Wednesday 7 September remains for free presentations.

REGISTRATION DESK

The Registration Desk at Palexpo will be open as follows:

Saturday 3 September	15:00 - 18:00
Sunday 4 September	08:00 - 18:00
Monday 5 September	08:00 - 19:00
Tuesday 6 September	08:00 - 19:00
Wednesday 7 September	08:00 - 18:00

RULES

Smoking Policy – The HUP0 2011 Congress is a non-smoking event. Participants are kindly requested to refrain from smoking in the Congress venue, including the Exhibition Area.

Mobile Phones – Delegates are kindly requested to keep their mobile phones in the off position in the rooms where scientific and educational sessions are being held, as well as during poster sessions.

TRANSPORTATION

With any hotel reservation, a free transportation ticket on the Geneva public transportation network (TPG-Unireso) for the duration of the congress is provided. During HUP0 2011, the Congress badge will also grant participants free access to the Geneva public transport network.

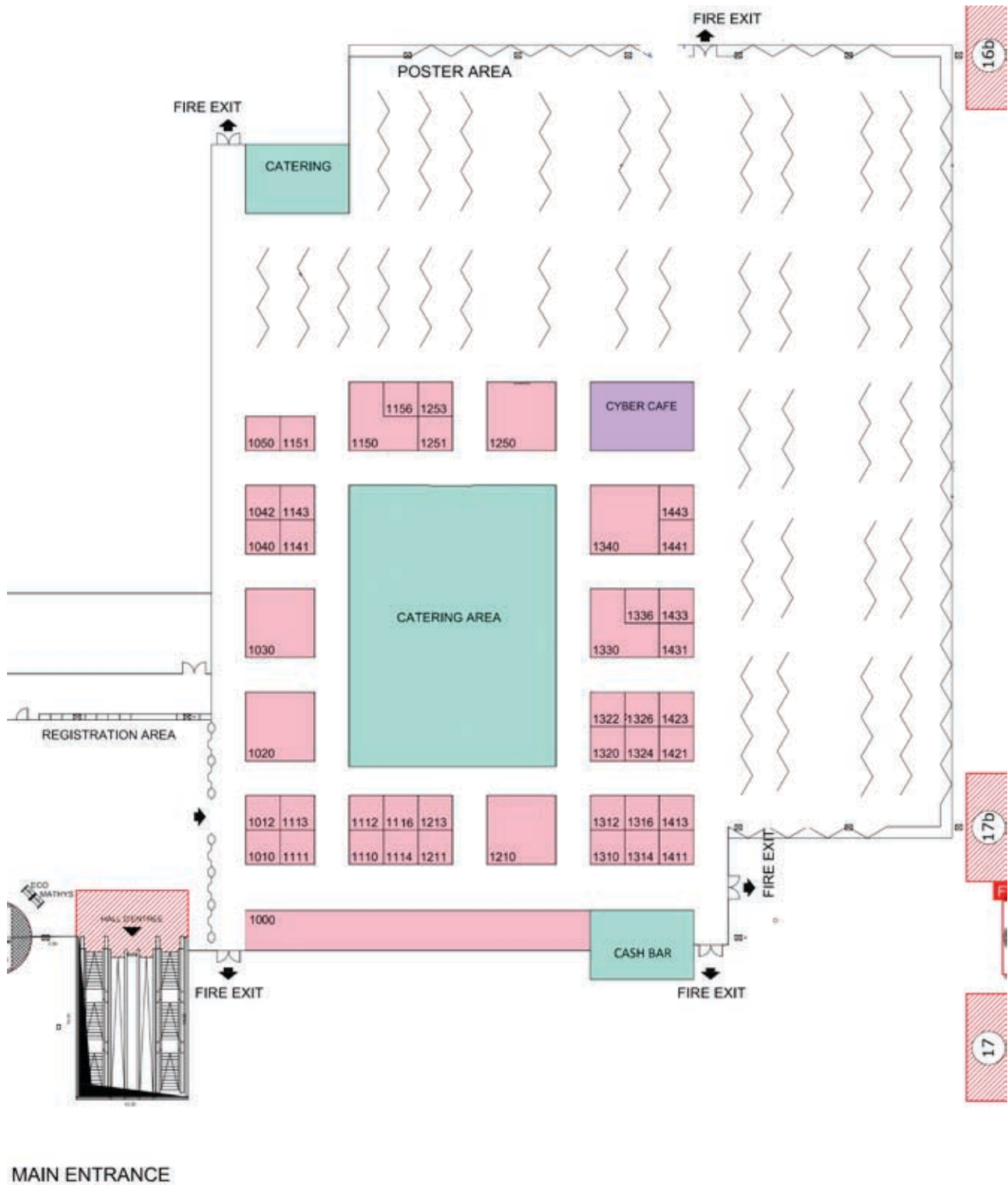
SPEAKER PREVIEW ROOM

Please ensure that you have visited the Speaker Preview Room to confirm your audiovisual requirements at least 2 hours prior to the start of your session. However, it is recommended speakers register with the Speaker Preview Room the day before their presentation. The Speaker Preview Room is located at the entrance of the Congress Centre.

Opening times are as follows:

Saturday 3 September	14:00 - 18:00
Sunday 4 September	08:00 - 20:00
Monday 5 September	07:00 - 19:00
Tuesday 6 September	07:00 - 19:00
Wednesday 7 September	07:00 - 17:30

Exhibition Floor Plan



Exhibitor List

COMPANY	STAND N°
ABCAM PLC	1314
AB SCIEX	1020
ACS PUBLICATIONS	1111
AGILENT TECHNOLOGIES	1340
BECKMAN COULTER, INC	1322
BIOGNOSYS AG	1151
BIOINFORMATIC SOLUTIONS	1421
BIO-RAD	1210
BRUKER DALTONIK GMBH	1150
BUCHER BIOTEC AG	1113
CAMBRIDGE ISOTOPE LABORATORIES, INC	1143
CAPROTEC BIOANALYTICS GMBH	1112
CELL SIGNALING TECHNOLOGY	1443
DECODON & DYE DIAGNOSTICS	1253
DENATOR AB	1431
DIONEX	1156
EKSIGENT	1020
ELSEVIER	1040
ENZO LIFE SCIENCES (ELS) AG	1110
EUPA	1040
EUROPEAN PHARMACEUTICAL REVIEW	1050
GE HEALTHCARE	1330
GENEBIO	1012
GENEDATA	1141
HUPO 2012	1326
JPT PEPTIDE TECHNOLOGIES GMBH	1326
LIFE TECHNOLOGIES GMBH	1423
LUMINEX COPORATION	1000
MATRIX SCIENCE	1336
MERCK MILLIPORE	1213
MILTENYI BIOTEC	1320
NONLINEAR DYNAMICS	1433
OLINK BIOSCIENCE	1310
PEAK SCIENTIFIC INSTRUMENTS LTD	1116
PROMEGA	1043
PROTEINSIMPLE (EX CELL BIOSCIENCES)	1114 & 1271
PROTEOME SOFTWARE	1324
RANDOX LABORATORIES LTD	1411
SERVA ELECTROPHORESIS GMBH	1251
SIGMA LIFE SCIENCE	1010
SILANTES GMBH	1441
THERMO FISCHER SCIENTIFIC	1250
WATERS	1030
WILEY - BLACKWELL	1312

Company Profiles

ABCAM PLC

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ORDERS@ABCAM.COM
www.abcam.com

BOOTH N°1314



Abcam is a producer and distributor of research-grade antibodies and associated proteomics research products. Headquartered in Cambridge (UK), Abcam has subsidiaries in Cambridge (US), Tokyo (Japan) and Hong Kong (China), serving a global customer base of over 75 countries.

Abcam's online catalog has over 70,000 products and includes a growing range of non-antibody products such as proteins, peptides, lysates, immunoassays and kits.

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AB SCIEX

110 MARSH DRIVE
FOSTER CITY, CA 94404
USA
www.absciex.com

BOOTH N°1020



AB SCIEX helps to improve the world we live in. AB SCIEX solutions for biomarker & omics research enable scientists to push the limits in discovery proteomics, targeted peptide quantitation, lipidomics, metabolomics, protein characterization, and tissue imaging. With over 20 years of proven innovation and world-renowned TripleTOF™, QTRAP®, TOF/TOF™, and Eksigent® systems, AB SCIEX excels by listening to and understanding the ever-evolving needs of its thousands of customers to develop reliable, sensitive and intuitive solutions that continue to redefine what is achievable. For more information about AB SCIEX biomarker & omics solutions, please go to www.absciex.com/omics.

ACS PUBLICATIONS

1155 SIXTEENTH STREET NW
ZIP CODE, CITY 20036
WASHINGTON, DC
USA
T_PRITCHETT@ACS.ORG
<http://pubs.acs.org>

BOOTH N°1111



The world's most trusted journal in the international field of proteomics, Journal of Proteome Research (JPR) celebrates its 10th year of publishing high-quality research in 2011. Led by Editor-in-Chief William S. Hancock, JPR publishes content encompassing all aspects of global protein analysis and function with emphasis on the synergy between physical and life sciences that result in a multidisciplinary approach to the understanding of biological processes. Visit 1111 to learn about our newest title ACS Synthetic Biology and a 2GB flash drive when you take our survey (while supplies last).

Company Profiles

AGILENT TECHNOLOGIES

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Bioinformatics Solutions Inc. develops software as:
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We hold in principle that it is not enough to develop solely on theory; we must develop with customer needs in mind. We believe the only solution is one that incorporates highly confident results and timely results, a satisfying product experience, customer support and two-way communication.

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caprotec bioanalytics GmbH is a biotechnology company based in Berlin, Germany. caprotec has developed a novel technology to reduce the complexity of protein mixtures in a targeted manner. The Capture Compound Mass Spectrometry (CCMS) technology is based on trifunctional Capture Compounds™ which enable capturing and isolating proteins based on their functionality. The CCMS technology is commercialized as caproKit™ reagents for research applications in the life science area. With the ImproMed™ application of CCMS caprotec is forming collaborations with pharmaceutical and biotechnology companies.

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DECODON GmbH is a bioinformatics company that develops innovative software for the modern Life Sciences. Its bioinformatics tools are helping scientists to generate knowledge from the massive amounts of data that are accumulated by recent methods in global functional genomics.

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One of the journals we have on display is the Journal of Proteomics which covers all areas of applied and basic research in Proteomics using multi-disciplinary approaches to unravel biological processes. Emphasis is placed on translational research and biomarker discovery in human, animal, micro-organism and plant systems.

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European Pharmaceutical Review is the leading bi-monthly technical journal for the pharmaceutical industry, and essential reading for anyone involved in drug discovery or manufacturing.

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JPT Peptide Technologies is the leading provider of low cost peptide libraries and microarrays worldwide for applications in multiplexed SRM/MRM assays, immune response and enzyme profiling. SpikeTidesTM is a unique approach to provide small scale unlabeled and isotopically labeled peptides for the development of high-throughput assays for quantification of biomarker expression levels. PepStarTM is a proprietary peptide microarray platform for testing humoral immune responses for many different antigens (including cancer, HCMV, HIV, malaria, mycobacterium tuberculosis, and others). This technology allows systematic profiling of antibody signatures, patient stratification and humoral immune response profiling using patient samples related to infectious and autoimmune diseases, allergies and cancer.

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Company Profiles

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3. Service for in vivo proteomics in mice.

Contacts:

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The logo for Thermo Scientific features the word "Thermo" in a large, bold, red, sans-serif font. Below it, the word "SCIENTIFIC" is written in a smaller, black, sans-serif font, with all letters in uppercase.

Company Profiles

Thermo Scientific, part of Thermo Fisher Scientific Inc., the world leader in analytical instrumentation, develops integrated solutions to help solve the analytical challenges in proteomics. Thermo Scientific solutions leverage the unmatched performance, speed, sensitivity and accuracy of their market-leading Orbitrap, ion trap and triple quadrupole mass spectrometers for definitive quantitation of protein mixtures. Join our workshops on 5th & 6th September and visit our two booths see example workflows and learn more.

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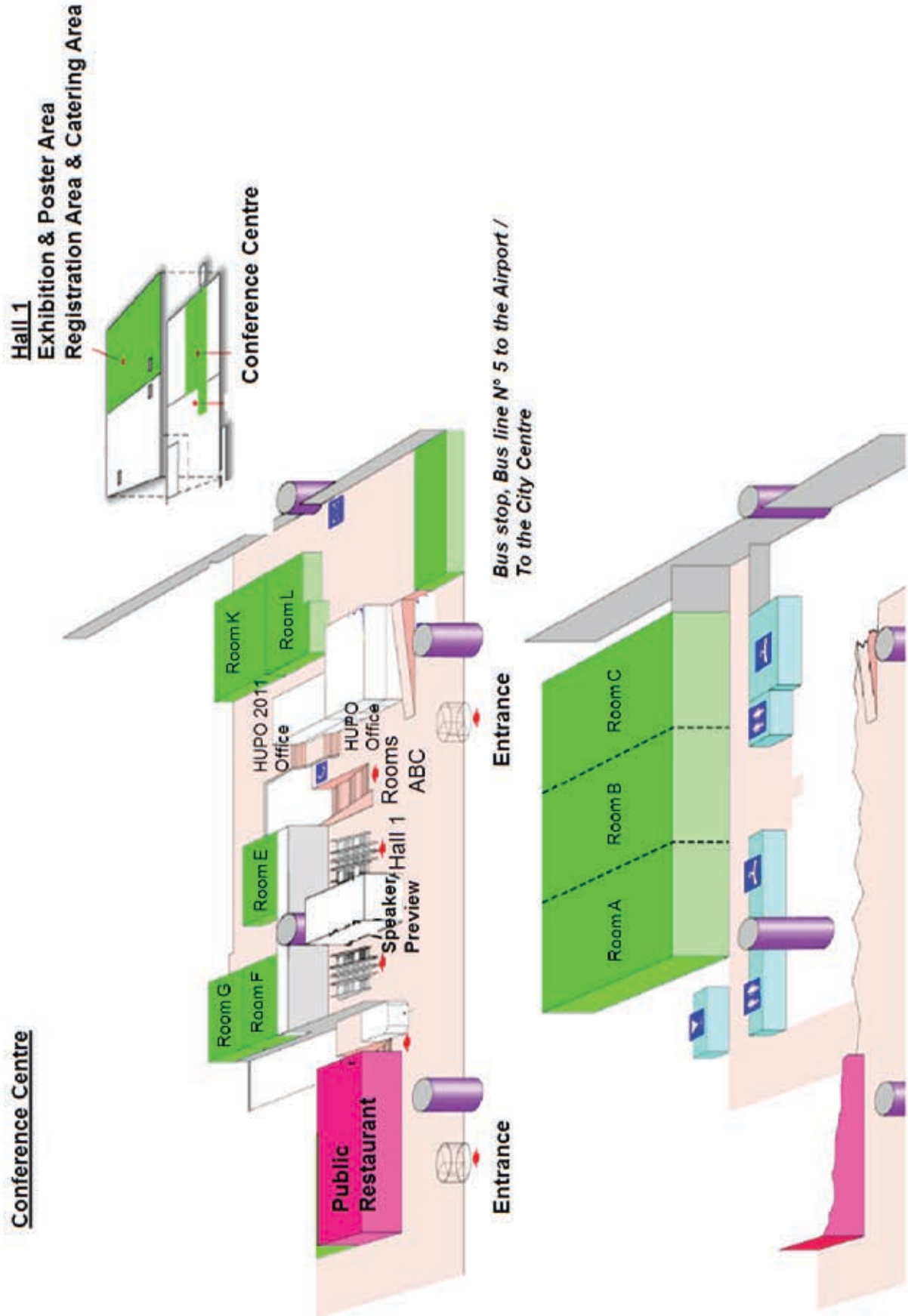
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