

## Program

### Thursday, June 20<sup>th</sup>, 2024

<b>12:00-13:00</b>	<b>Registration opens. Welcome coffee and small snacks.</b>
13:00-13:05	<b>Welcome</b> – Michal Bassani (CHUV) & LS <sup>2</sup> Proteomics Board
<b><u>13:05-14:35</u></b>	<b><u>Oral Session I</u></b>
<b>13:05-13:50</b>	<b><u>Keynote 1:</u></b>  <b>Kirti Sharma</b> (Kymera Therapeutics, Boston) “Leveraging mass spectrometry proteomics to advance TPD based degraders from early discovery to clinic”.
13:50-14:35	<b>Oral and Flash presentations - PTMs and interactions</b>
<b>14:35-15:10</b>	<b>Coffee Break</b>
<b><u>15:10-16:25</u></b>	<b><u>Oral Session II</u></b>
15:10-16:25	<b>Oral and Flash presentations - Methods</b>
<b>16:25-16:55</b>	<b>Coffee Break</b>
<b><u>16:55-17:40</u></b>	<b><u>Oral Session III</u></b>
16:55-17:05	<b>Sponsored talk - Jean-Baptiste Vincendet (Sciex)</b>
17:06-17:40	<b>Oral and Flash presentations - Basic Biology</b>
<b><u>17:40-18:20</u></b>	<b><u>Career Development – panel discussion</u></b>
<b>18:35-18:45</b>	<b>Group picture</b>
<b>18:45-20:00</b>	<b>Apéro (upon registration only!)</b>
<b>20:00-22:30</b>	<b>Dinner in the conference hotel (upon registration only!)</b>

Friday, June 21<sup>st</sup>, 2024

08:00-09:30

Workshop (*upon registration only*):  
"How can proteomics drive drug development?".

09:30-10:30

Oral Session IV

09:30-10:20

**Oral and Flash presentations - Clinical and Spatial**

09:30-09:35

**Sponsor talk - Torsten Müller** (Bruker Daltonics)

10:21-10:26

**Sponsor talk - Dominic G. Hoch** (Thermo Fisher Scientific)

**10:27-11:00**

**Coffee Break**

11:00-12:45

Oral Session V

11:00-12:00

**Oral and Flash presentations - Clinical and Spatial**

11:46-11:51

**Sponsor talk - Tinkara Vozel** (Olink)

12:00-12:45

**Keynote 2:**

**Dr. Fabian Coscia** (Max Delbrück Center for Molecular  
Medicine, DE)  
"Spatial tissue proteomics to assess health and disease".

12:45-13:00

Awards and closing remarks.

12:45-12:55

**Best oral presentation prize & best flash talk prize**

12:55-13:00

**Closing remarks** – Michal Bassani (CHUV) & LS<sup>2</sup> Proteomics  
Board

**13:00-14:30**

**Farewell Lunch** (*upon registration only!*)

14:30-15:30

LS<sup>2</sup> Proteomics Board Meeting

## Detailed Program

### Thursday, June 20<sup>th</sup>, 2024

- 12:00-13:00**      **Registration opens. Welcome coffee and small snacks.**
- 13:00-13:05      **Welcome** – Michal Bassani (CHUV) & LS<sup>2</sup> Proteomics Board
- 13:05-14:35**      **Oral Session I**
- 13:05-13:50**      **Keynote 1:**
- Kirti Sharma** (Kymera Therapeutics, Boston)  
“Leveraging mass spectrometry proteomics to advance TPD based degraders from early discovery to clinic”.
- 13:50-14:35**      **Oral and Flash presentations - PTMs and interactions**
- 13:50-14:02      **Melanie Brunner** (University of Fribourg)  
Proteome-wide in vitro kinase and phosphatase assays to determine enzyme-substrate interactions
- 14:03-14:15      **Viviane Reber** (ETH Zurich)  
More than inhibition: Identification of a Pseudoactive Kinase Conformation and Altered Protein-Protein Interactions induced by ATP-Competitive Inhibitors
- 14:16-14:20      **Veronique Laforte** (Biognosys AG)  
Improved PTM proteomics analyses in Spectronaut®
- 14:21-14:25      **Benjamin Steinmetz** (ETH Zurich)  
Two-Step Enrichment of RNA-Peptide Adducts from Human Cell Lysates
- 14:26-14:30      **Ibrahim Bayramoglu** (University of Fribourg)  
Using Phosphoproteomics to Explore Autophagy Relevant TBK1 Signal Transduction
- 14:31-14:35      **Luca Riermeier** (ETH Zurich)  
Characterization of the organization of protein-RNA complexes using structural proteomics methods

14:35-15:10

**Coffee Break**

15:10-16:25

Oral Session II

15:10-16:25

**Oral and Flash presentations - Methods**

15:10-15:22

**Christopher Below** (Biognosys AG)

Synchronizing the TIMS ramp with the quadrupole scan demonstrates a significant qualitative and quantitative improvement in DIA analysis

15:23-15:35

**Til Schlotter** (ETH Zurich)

Interface Nanopores for Single-Molecule Protein Analysis

15:36-15:48

**Liliana Malinovska** (Biognosys AG)

High-throughput and automatable drug target identification using limited proteolysis coupled to mass spectrometry (LiP-MS) suitable for phenotypic screening applications

15:49-15:53

**Paolo Nanni** (University of Zurich)

A transfer-free proteomics approach for the analysis of few dispensed cells: a core facility perspective

15:54-15:58

**Romain Hamelin** (EPF Lausanne)

Wide-Window Acquisition (WWA) DDA for label-free quantitative analysis of low protein yielding systems using an Orbitrap

15:59-16:03

**Emmanuelle Lezan** (Hoffmann-La Roche)

Comparison of two sample preparation methods for the analysis of human plasma and mouse serum using LC-MS/MS

16:04-16:08

**Sebastian Müller** (Biognosys AG)

Facilitating large-scale proteomics by combining automated sample preparation with short gradient LC-MS acquisitions

16:09-16:13

**Ane Laura Fineid Pedersen** (Nestle Research)

Assessing the performance of proteomic workflows with a two-proteome mixture: quantitative accuracy and precision of label-based and label-free approaches

16:14-16:24 **Sponsored talk**  
**Olivier Buetzberger** (Agilent Technologies)  
Leveraging Agilent's portfolio for high-throughput liquid handling, imaging, and detection methods in your proteomics workflow

**16:25-16:55** **Coffee Break**

**16:55-17:40** **Oral Session III**

16:55-17:05 **Sponsor talk**  
**Jean-Baptiste Vincendet** (Sciex)  
ZenoTOF 7600: current key applications in proteomics and perspectives

17:06-18:20 **Oral and Flash presentations - Basic Biology**

17:06-17:18 **Marc van Oostrum** (University of Basel)  
The proteomic landscape of synaptic diversity across brain regions and cell types

17:19-17:31 **Alexandre Leytens** (University of Fribourg)  
Targeted proteomics addresses selectivity and complexity of protein degradation by autophagy

17:32-17:36 **Ramon Pfaendler** (ETH Zurich)  
A systems view of cellular state heterogeneity in human pluripotent stem cells

17:36-17:40 **Sandra Schär** (Biognosys AG)  
Ultra-deep profiling of human and rodent tissues allows for deepest ever analysis of protein expression levels

**17:40-18:20** **Career Development – panel discussion**

**18:35-18:45** **Group picture**

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**20:00-22:30** **Dinner in the conference hotel** (*upon registration only!*)

Friday, June 21<sup>st</sup>, 2024

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Workshop (*upon registration only*):  
"How can proteomics drive drug development?".

*Confirmed panelists:*

**Andreas Hofmann** (Novartis)

**Debora Bonenfant** (Monte Rosa Therapeutics)

**Manuel Tzouros** (Roche)

**Kirti Sharma** (Kymera Therapeutics)

09:30-10:30

Oral Session IV

09:30-10:30

**Oral and Flash presentations - Clinical and Spatial**

09:30-09:35

**Sponsored talk**

**Torsten Müller** (Bruker Daltonics)

Innovations in 4D-Proteomic Analysis with timsTOF Mass Spectrometers

09:35-09:47

**Jianbo Fu** (ETH Zurich)

MultiOmicsAgent: Guided extreme gradient-boosted decision trees-based approaches for biomarker-candidate discovery in multi-omics data

09:48-10:00

**Neil Taylor** (University of Tuebingen)

Genetic code expansion-derived approach to detect oxidative stress-induced changes in the neuronal proteome

10:01-10:05

**Axel Martinelli** (BigOmics)

Exploring a Sarcoidosis Olink dataset with the Omics Playground platform

10:06-10:10

**Alex Soltermann** (Pathologie Länggasse)

Amyloid typing by mass spectrometry

- 10:11-10:15      **Arthur Viodé** (Biognosys AG)  
An efficient and deep plasma workflow applied discover biomarker candidates of response to chemo-immunotherapy in patients with thoracic malignancies
- 10:16-10:20      **Antonino Zito** (BigOmics Analytics)  
Imputation of Missing Values in Proteomics
- 10:21-10:26      **Sponsor talk**  
**Dominic G. Hoch** (Thermo Fisher Scientific)  
Accelerating translational omics – Discovery to validation at unprecedented scale
- 10:27-11:00**      **Coffee Break**
- 11:00-12:45**      **Oral Session V**
- 11:00-12:00**      **Oral and Flash presentations - Clinical and Spatial**
- 11:00-11:12      **Thibault Courtellemont** (EPF Lausanne)  
DIY Spatial Proteomics: Creative Solutions for an Affordable Pipeline
- 11:13-11:25      **Alexia Clavier** (University of Bern)  
Perfusate biomarkers of DCD cardiac graft quality identified with proteomics: Studies in an isolated rat heart model
- 11:26-11:30      **Sandrine Reymond** (University of Geneva)  
DIA-based investigation of plasma-derived extracellular vesicles for the discovery of biomarkers for acute stroke diagnosis
- 11:31-11:35      **Sibilla Sander** (University of Fribourg)  
Proteomic profiling of cancer-associated fibroblasts in squamous cell carcinoma
- 11:36-11:40      **Quentin Enjalbert** (PreOmics)  
Unique high-throughput workflow for deeper plasma/serum proteome coverage enables discovery of novel biomarkers

- 11:41-11:45      **Yong-Qi Gao** (EPF Lausanne)  
Dissecting neuron-specific roles of reactive lipid-derived electrophiles in *C. elegans* neurodegenerative disease models
- 11:46-11:51      **Sponsor talk**  
**Tinkara Vozel** (Olink)  
PEA Technology in Action: Case Studies from Clinical Research  
Frontiers
- 12:00-12:45      **Keynote 2:**  
  
**Fabian Coscia** (Max Delbrück Center for Molecular Medicine, DE)  
“Spatial tissue proteomics to assess health and disease”
- 12:45-13:00**      **Awards and closing remarks.**
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- 12:55-13:00      **Closing remarks** – Michal Bassani (CHUV) & LS<sup>2</sup> Proteomics Board
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