

Conference on Biomolecule Analysis

Registration & Accommodation

REGISTRATION

Please register online under

<http://campus.uni-muenster.de/konf.html>

Registration fee:

Regular: 50 €

Students: 10 €

On-site: 60 €

VENUE



Max-Planck-Institute
for Molecular Biomedicine

Röntgenstraße 20
D-48149 Münster

ACCOMMODATION

Close to the conference site:

Haus Niemann
Horstmarer Landweg 126
48149 Münster

Tel: +49 (0) 251 / 82828

Fax: +49 (0) 251 / 88800

(5-10 min walking distance)

Hotel Jellentrup

Hüfferstraße 52

48149 Münster

Tel: +49 (0) 251 / 98105-0

Fax: +49 (0) 251 / 98105-40

(20 min walking distance)

Münster Tourist Information

Tel: +49 (0) 251 / 492 2710

Fax: +49 (0) 251 / 492 7743

E-mail: tourismus@stadt-muenster.de

<http://www.muenster.de/tourismus>

CONFERENCE OFFICE & ORGANISATION

IZKF Scientific Office

Albert-Schweitzer-Campus 1, Building D3, 48149 Münster

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E-mail: sekretariat.cu-proteomics@uni-muenster.de



Münster Conference on Biomolecule Analysis

December 2- 3, 2015

Core Unit Proteomics
Interdisciplinary Centre for Clinical Research
(IZKF) Münster

Medizinische Fakultät
der Westfälischen
Wilhelms-Universität Münster
Dekanat
Domagkstraße 3
48149 Münster

Adresse des IZKF:
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48149 Münster

medizinische Fakultät
Hohe Schule der
münster Medizin.

medizinische
Fakultät
Westfälische
Wilhelms-Universität Münster

WESTFÄLISCHE
WILHELMS-UNIVERSITÄT
MÜNSTER

Conference on Biomolecule Analysis

Welcome Address

The 2nd Münster Conference on Biomolecule Analysis, focusing on advances in the field of proteomics will be held on December 3, 2015 at the Max-Planck Institute for Molecular Biomedicine in Münster. The conference will be jointly hosted by the Interdisciplinary Centre for Clinical Research (IZKF) Münster and Serva Electrophoresis GmbH.

The organizers are proud to present eminent speakers who have influenced proteomics research on many fronts. The keynote lecture will be delivered by Ruedi Aebersold, the Co-Founder of the Institute for Systems Biology in Seattle, who is now heading a group of quantitative mass spectrometry at ETH Zurich. Another highlight in this top line-up of speakers will be an application example of high-end mass spectrometry involving a mummy given by Anna and Andrej Shevchenko of MPI Dresden.

We will also hold three pre-conference workshops given by the technology leaders in proteomic analysis on 02. December. Due to high demand some of the workshops will be held again on 03. December after the conference. Sponsoring companies will have their products on display on the day of the conference.

Graduate students are especially encouraged to present their work in poster form. The two best posters will be awarded a book voucher at a value of 150 EUR each.

We are looking forward to welcoming you in Münster to a day of stimulating discussions with fellow academicians and industrial partners.

Programm

Wednesday, 2. December

WORKSHOPS	
09:00 - 11:30	Reiner Westermeier Serva Electrophoresis GmbH Proteomics sample preparation and gel electrophoresis. Fluorescence in electrophoresis - new methods in protein visualisation
11:30	Lunch
12:00 - 14:00	Robert van Ling ThermoFisher Scientific Nano liquid chromatography: Introduction and live demo
14:00 - 15:00	Marc Kipping Waters Corporation Data Dependent (DDA) and Data Independent (DIA) LC-MS Acquisition Strategies in Quantitative Proteomics
	Martin Zeller ThermoFisher Scientific Mass spectrometry in proteomics
16:00	PRE-CONFERENCE SOCIAL LWL Museum für Kunst und Kultur Guided Tour of the Special Exhibition „Wilhelm Morgner and Modern Art“

Thursday, 3. December

8:30	CONFERENCE REGISTRATION
9:00	OPENING & WELCOME Simone König, Günter Theßeling
9:10	KEYNOTE LECTURE Ruedi Aebersold Quantitative measurement of proteins and proteomes by targeted mass spectrometry: Reproducibility matters

Programm

9:55	SESSION I Boris Mačák Dissecting the cell cycle control in fission yeast by quantitative phosphoproteomics
10:20	Martin Eisenacher A Day in the Life of a 'Bioinformatics for Proteomics' Researcher
10:45	COFFEE BREAK & POSTER SESSION
11:15	SESSION II Dörte Becher Quantitative proteomics at a global scale – methods and applications in bacteria
11:40	Emöke Bendixen Host genetics influence on E.Coli virulence
12:05	LUNCH & POSTER SESSION
13:15	SESSION III Eberhard Krause MS analysis of 'difficult' protein modifications
13:40	Andrej Shevchenko Ancient cheese found with mummies
14:30	POSTER AWARDS & CLOSING REMARKS
14:45	END OF CONFERENCE
15:00 - 17:00	WORKSHOPS Robert van Ling ThermoFisher Scientific
17:00 - 17:30	Marc Kipping Waters Corporation
17:30 - 18:00	Martin Zeller ThermoFisher Scientific