

Tuesday, 10.05.2017, 5 pm

Location: IGZW

Conference Room 3rd floor
Gregor-Mendel-Strasse 4

Doors open at 4:45 pm

Refreshments will be served

Methods covered

- ❖ Bioinformatics special seminar
by Jürgen Behr



Dr. Jürgen Behr
BayBioMS

Experimental design and data analysis in
proteomics and metabolomics



Dr. Karin Kleigrew
(Metabolomics)



Dr. Jürgen Behr
(Bioinformatics)

Presentation of Research Projects



Dr. Peng Yu
Chair of Proteomics
and Bioanalytics

Trinity P1 mixed mode chromatography enables
mass spectrometry compatible orthogonal
peptide separation to reversed phase



Stephanie Heinzlmeir
(Proteomics)



Maria Reinecke
Chair of Proteomics
and Bioanalytics

Selectivity profiling of small molecule kinase
inhibitors using chemical proteomics



Dr. Christina Ludwig
(Proteomics)