

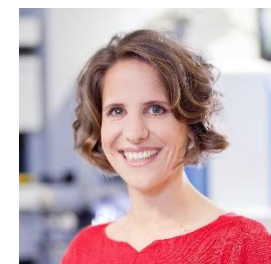
BayBioMS – Advanced mass spectrometry seminar

When: Thursday **10.04.2025, 3 pm**
Where: iGZW, conference room 3rd floor
Gregor-Mendel-Strasse 4

Doors open at 2:45 pm
Free drinks and snacks will be served!!

BayBioMS

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Dr. Christina Ludwig
(Proteomics)



Dr. Karin Kleigrew
(Metabolomics)



Amirhossein Sakhteman
(Bioinformatics)

BayBioMS proudly presents our special guest speaker:

Brendan MacLean

University of
Washington



“Skyline - A software solution for quantitative mass spectrometry studies in proteomics and metabolomics”

Skyline is a freely-available and open source software tool for analyzing proteomic as well as metabolomic mass spectrometry data in a targeted fashion. Over the last 15 years, Skyline has emerged as the most popular analysis software in the field of targeted proteomics. Brendan MacLean is the principal investigator of Skyline and responsible for all aspects of design, development and support in creating the Skyline Environment and its growing worldwide user community.

