



Dear friends of OpenMS,

Welcome to the 9th OpenMS User Meeting.

During the meeting, we will present the capabilities of OpenMS, and discuss your needs and future directions in mass spectrometry.

This year's meeting will include an **optional OpenMS tutorial session on the first day (Wednesday, September 21)** with the **general conference starting on Thursday, September 22**.

Traditionally, the schedule of our meetings is very flexible, allowing discussions and contributions from the participants.

You can **bring your data**, an **OpenMS wish list**, **ideas for collaboration projects**, and also **give a talk about your project**. This may include suggestions in areas where currently available software solutions were unsatisfactory. We would be very happy to welcome you in Tübingen this year.

We hope you will enjoy our user meeting.

Wednesday, September 21st (Tutorial Session)

Wilhelm-Schickard-Institut , Sand 14

Time	Event	Location
09:00	Registration and Coffee	WSI Sand, A104
10:00	Welcome and Opening to Tutorials	WSI Sand, A104
10:15	Tutorial Session 1 <i>Track 1: Hands-on Tutorial</i> <i>Track 2: Bring Your Own Data (BYOD)</i>	
12:30	Lunch	
13:45	Tutorial Session 2 <i>Track 1: Hands-on Tutorial (continued)</i> <i>Track 2: Bring Your Own Data (BYOD)</i>	WSI Sand, A104
15:30	Coffee break	
16:00	Tutorial Session 3 <i>Track 1: Hands-on Tutorial (continued)</i> <i>Track 2: Bring Your Own Data (BYOD)</i>	
18:00	Dinner (individual) / Free evening	

Thursday, September 22st (Conference)

Konferenzzentrum (conference center) Schnarrenberg, Otfried-Müller-Straße 6

Time	Event	Location
09:00	Registration and Coffee	conference room 1
09:15	Welcome and Opening of the User Meeting	
09:30	Introduction to OpenMS 2.1 <i>Oliver Kohlbacher, University of Tübingen</i>	
10:00	Session 1: Developer talks Julianus Pfeuffer: "Protein inference using Bayesian methods" Timo Sachsenberg: "OpenMS tools for Protein-SIP and RNA-Protein cross-link identification"	
11:00	Coffee break	
11:15	Session 2: Developer talks Erhan Kenar: "Label-free quantification of Metabolites " Marc Rurik: "LC-MS-based Metabolomics in OpenMS"	
12:15	Lunch	
13:15	Meet the developers: Open discussion & Collaboration	conference room 1
14:45	Coffee break	
15:00	Meet the developers: Open discussion & Collaboration	conference room 1

15:30	Social Event: <i>Museum at the Hohentübingen Castle, Stocherkahnfahrt and Dinner</i>	city
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Friday, September 23st (Conference)

Konferenzzentrum (conference center) Schnarrenberg, Otfried-Müller-Straße 6

Time	Event	Location
09:30	Contributed talks 1: Tamara Scheidt: <i>“Comprehensive phosphoproteome analysis of activated hedgehog signaling in medulloblastoma cells”</i>	conference room 1
10:00	Coffee break	
10:30	Keynote talk: Metabolite Identification Prof. Dr. Sebastian Böcker – Friedrich-Schiller-Universität Jena, Lehrstuhl für Bioinformatik	
11:30	Tjeerd Dijkstra: <i>“Missing Data in Quantitative Proteomics”</i> Roger Olivella: <i>“Implementing a web-based QC system using OpenMS”</i>	conference room 1
12:30	Lunch	
13:30	Open discussion & Collaboration	conference room 1
14:30	Closing remarks	
14:45	Open discussion & Collaboration / Coffee	
15:30	End of meeting / Free evening	