

# Orbitrap LC-MS/MS User Meeting

Odense, Denmark | 4 December 2019

BMB seminar room entrance O (currently blocked),  
University of Southern Denmark, Campusvej 55 5230, Odense

The aim of the meeting is to share best-practice and meet colleagues and specialists in this field. We will be present with chemists and engineers working with Orbitrap™ Systems. The meeting is free but registration is required.

Time	Topic
09:00	Registration and coffee
09:30	Welcome and Introduction Willy Björklund, Thermo Fisher Scientific
09:40	New horizons of Orbitrap mass spectrometry Alexander Makarov, Thermo Fisher Scientific
10:15	Using Thermo Scientific™ SureQuant™ for Targeted Mass Spec Analysis Shannon Eliuk, Thermo Fisher Scientific
10:45	Coffee break
11:15	The best of two worlds - DIA & Thermo Scientific™ FAIMS Pro™ Interface on the Thermo Scientific™ Orbitrap Exploris™ 480 mass spectrometer Jesper Olsen, Novo Nordisk Foundation Center for Protein Research Proteomics Program
11:50	Exploring Orbitrap Technology for Single Cell Proteomics Simonas Savickas, DTU Bioengineering
12:30	Lunch
13:30	Going Beyond 11: Increasing quantitative proteomic sample multiplexing using next generation mass tags Ryan Bomgarden, R&D, Thermo Fisher Scientific
14:00	Experiences from the European Cancer Moonshot Lund Center Lazaro Betancourt, Lund University
14:30	Native protein characterization: A Biggie Small Story Vlad Zabrouskov, Thermo Fisher Scientific
15:00	General discussion and questions - followed by Snacks and Drinks

Register before November 28, 2019 at  
[thermofisher.com/nordic/events](https://thermofisher.com/nordic/events)