

Introduction to mass spetrometry-based proteomics

The CF proteomics is happy to announce that also this year it will give an introduction to mass spectrometry-based proteomics for everybody that is interested.

The course will give an insight into how to submit samples to our facility, what happens with your samples once they are submitted and what data you will obtain from us. It is designed to give you a better understanding on the limitations and possibilities of mass spectrometry.

The course will be split into three parts:

- Wednesday, 10th April 2024, 10 11:30: "An overview and introduction to mass spectrometry-based proteomics"
- Thursday, 11th April 2024, 13:30 15:00: "Introduction in sample preparation and common contamination sources"
- Friday, 12th March 2024, 13:30 15:00: "Proteomics data analysis and statistics"

As the course will be in the Golgi, we will be limited by space (up to 15 persons) and would therefore ask you to register for the course via this <u>form</u>. If not working, please write to proteomics@leibniz-fli.de.

For all PhD candidates, this course will be part of LGSA training. A certificate of attendance will be provided to all participants.