AGENDA

10:00  Part I: Characterizing the dynamics of alpha-synuclein oligomers using HDX-MS
Part II: Overcoming gas phase H/D-scrambling
Thomas JD Jørgensen, SDU

10:30  The power in combining HX-MS with other biophysical techniques
Simon Mysling, Finsen lab, Rigshospitalet

11:00  Coffee break

11:15  Fibrillation of proteins
Sara Snogerup Linse, Lund University

11:30  Intact protein analysis using the Orbitrap mass analyzer
Kai Scheffler, Thermo Fisher Scientific

12:00  Lunch Break

13:00  HDX-MS solution to probe protein conformation/conformation dynamics
Terry Zhang, Thermo Fisher Scientific

13:30  Taking the pain out of HDX-MS – options for automation
Peter Smith, LEAP Technology

14:00  Lab Session

15:00  End of meeting, Q&A Session

Registration

VENUE
Room: I1308
BMC (Biomedicinskt centrum)
Lund University, Sölvegatan 19, 221 84 Lund