Structural biologists face several challenges when trying to solve the structure of large and dynamic complexes. However, the combination of different techniques as Mass Spectrometry and Cryo-EM, an approach known as integrative structural biology, is revolutionizing the understanding of protein structure, function and dynamics.

Program:

9.30 am – 10.00 am: Check-in and Coffee

10.00 am – 10.45 am: Natalia de Val PhD, Thermo Fisher
Introduction to Cryo-EM and Its Integration with Mass Spectrometry, Application Results

10.45 am – 11.00 am: Q&A

11.00 am – 11.30 am: Rosa Viner PhD, Thermo Fisher
Introduction to Mass Spectrometry and Its Integration with Electron Microscopy, Application Results

11.30 am – 11.45 am: Q&A

11.45 am – 12.45 pm: Lunch

12.45 pm – 1.15 pm: Vadim Cherezov PhD, USC
Structural Studies of G Protein-Coupled Receptors

1.15 pm – 1.45 pm: Iain D. G. Campuzano PhD FRSC, Amgen
Hydrophobic Protein Analysis in Industry: From Native-MS to High Throughput-MS Therapeutic Project Support

1.45 pm – 2.15 pm: Rachel Loo PhD, UCLA
Complementary MS and CryoEM: Macromolecular Complexes and Post-Translational Modifications

2.15 pm – 2.45 pm: Coffee Break

2.45 pm – 3.15 pm: Cornelius Gati PhD, USC
Structural Basis of GABA Reuptake Inhibition

3.15 pm – 3.45 pm: Lan Huang PhD, UCIrvine
Structural Analysis of Protein Complexes Using Cross-linking Mass Spectrometry

3.45 pm – 4.00 pm: Wrap-up