

Tools for Analysis of Membrane Proteins

Brussels, 20 April 2017

Scientific committee

Nico Callewaert (VIB-UGent Center for Medical Biotechnology Center)
Rouslan Efremov (VIB-VUB Center for Structural Biology)
Mark Veugelers (VIB Technology Watch team)

o9h3o-o9h35 Welcome

SESSION 1: BIOCHEMICAL AND BIOPHYSICAL APPROACHES TO STUDY MEMBRANE PROTEINS

ogh35-ogh50 **Trainer 1**: Nico Callewaert or Rouslan Efremov

Suggested title: Introduction to membrane proteins

ogh50-10h20 Trainer 2: Tim Dafforn, School of Biosciences - University of Birmingham, GB

Detergent free production of membrane proteins

10h20-10h50 Trainer 3: Anass Jawhari, Calixar, FR

Stabilization of native and functional membrane proteins for drug discovery

10h50-11h15 Coffee

Session 2: Membrane Protein Interactions

11h15-11h50 Trainer 4: Cédric Govaerts, Université Libre de Bruxelles, BE

Suggested title: Biophysics and biochemistry of membrane proteins

11h50-12h25 Trainer 5: Arnold Driessen, University of Groningen, NL

Suggested title: Biophysics tools for characterizing protein-protein and protein-ligand

interactions

12h25-13h30 Lunch



Session 3: Signaling/Cell Biology across membranes and drug development

13h30-14h05 **Trainer 6**: Robert Tampé, Institute of Biochemistry, Goethe-University Frankfurt, DE Suggested title: Imaging membrane proteins

14ho5-14h40 Trainer 7: Chris Tate, MRC Laboratory of Molecular Biology, UK

Stabilization of membrane proteins in specific conformations for structural studies and drug screening

14h4o-15h15 **Trainer 8:** Graeme Fraser, Euroscreen, BE

Suggested title: Drug screening assays targeting signal-transducing membrane proteins

15h15-15h45 Coffee

Session 4: Structural Biology of Membrane Proteins

15h45-16h20 Trainer 9: Bernadette Byrne, Imperial College London, GB

Suggested title: Expression and purification of functionally active Eukaryotic membrane proteins

16h2o-16h55 Trainer 10: Werner Kühlbrandt, Max-Planck-Institute of Biophysics, DE

Suggested title: Structural studies of membrane proteins: X-ray crystallography and cryo-EM