

Preparation and characterization of macromolecular complexes

November 19th to 23rd 2018

IGBMC, Strasbourg-Illkirch, France



INSTRUCT-FRISBI course



Preliminary program

Monday, Nov. 19th

Morning: Lectures

- 9h00-9h30 Course introduction
- 9h30-10h00 Participants will present their projects and aims
- 10h30-12h30 Methods for isolation of macromolecular complexes:
- Engineering cells by genome editing to facilitate purification of complexes
 - Baculovirus production of recombinant multiprotein complexes: General principles

Afternoon: Practicals

- 13h30-15h30 Engineering cells by genome editing: design of gene tagging experiments
- 16h00-18h00 Baculovirus production of recombinant multiprotein complexes: manipulation of insect cells and infection with baculovirus expressing exogenous complexes

Tuesday, Nov. 20th

Morning: Practicals

- 9h00-10h30 Biophysical characterization of complexes:
NanoDSF for stability optimization of protein samples: presentation of the method and data acquisition
- 11h00-12h30 EM Grid preparation and analysis of sample homogeneity

Afternoon: Practicals

- 13h30-15h30 Engineering cells by genome editing: CRISPR/Cas9 reagents and gene tagging
- 16h00-18h00 Baculovirus production of recombinant multiprotein complexes: transfection of constructs for virus production

Wednesday, Nov. 21st

Morning: Practicals

- 9h00-10h00 Baculovirus production of recombinant multiprotein complexes: construct design
- 10h00-11h00 Biophysical characterization of complexes: NanoDSF data analysis
- 11h00-12h00 Biophysical characterization of complexes: SEC MALLS (Multiple Angle Laser Light Scattering coupled to Size Exclusion Chromatography) for protein sample quality control: presentation of the method and data acquisition

Afternoon: Mini-Symposium

“State-of-art strategies technologies for preparation and biophysical characterization of macromolecular complexes in view of biochemical and structural studies”

- 13h30–14h00 Identification & reconstitution of multiprotein complexes: the Exon Junction Complex as model system – **H. le Hir**
- 14h15–14h45 New opportunities offered by genome editing strategies - **J.P. Concordet**
- 15h00–15h30 Strategies for determining the 3D structure of macromolecular complexes at atomic or pseudo-atomic resolution - **B. Klaholz**
- 15h45–16h15 X-ray Free Electron Laser: Opportunities for drug discovery – **M. Hennig**
- 16h15–16h45 *Coffee Break*
- 16h45–17h15 Nanobodies as tools for structural proteomics and beyond - **J. Steyaert**

Thursday, Nov. 22nd

Morning: Practicals

- 9h00-10h00 Biophysical characterization of complexes:
SV-AUC (Sedimentation Velocity Analytical Ultracentrifugation) for protein sample quality control: presentation of the method and sample preparation
- 10h00-11h00 Biophysical characterization of complexes: SEC MALLS (Multiple Angle Laser Light Scattering coupled to Size Exclusion Chromatography)
- 11h00-12h00 Lecture on co-expression in *E. coli* for the reconstitution of multiprotein complexes (Christophe Romier)

Afternoon: Practicals

- 13h30-15h30 Engineering cells by genome editing: Testing integration with a luciferase assay
- 16h00-18h00 Baculovirus production of recombinant multiprotein complexes: collect and analyze culture for expression of exogenous complexes

Friday, Nov. 23rd

Morning: Practicals and lecture/discussion

- 9h00-10h00 Biophysical characterization of complexes:
SV-AUC data analysis
- 10h00-11h00 Biophysical characterization of complexes: SEC MALLS and NanoDSF data analysis
- 11h00-12h00 Lecture on the vaccinia virus expression system for the reconstitution of multiprotein complexes (Marc Ruff & Robert Drillien)

Afternoon: Wrap-up session

- 13h30-14h00 Summary of experiments
- 14h00-15h30 General discussion