

## Membrane Proteins — Topology, Dynamics & MD Setup — Hands-on

Learn how to prepare, embed and simulate membrane proteins in realistic lipid bilayers. This module connects topology prediction, bilayer construction, MD setup and analysis so you can study orientation, dynamics and interactions of channels, transporters and receptors in silico.

### Membrane Proteins — Topology, Dynamics & MD Setup

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#### Session 1

**Fee: Rs 23800** [Apply Now](#)

#### Membrane Protein Classes & Topology

Classes and architectures of membrane proteins

[single and multi pass helices](#) [beta barrels](#) [channels,](#)  
[transporters and GPCRs](#)

Topology prediction and orientation in membranes

[transmembrane segment prediction](#) [inside outside](#)  
[topology mapping](#) [signal peptides and tail anchors](#)

Lipid environment and biophysical context

**bilayer thickness and hydrophobic matching** **lipid composition and asymmetry ideas** **cholesterol and raft concepts**

### **Session 2**

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## **Bilayer Building & Embedding Workflows**

Building lipid bilayers for simulations

**choice of lipid types and mixtures** **bilayer size and area per lipid** **asymmetric leaflets and mixtures**

Embedding membrane proteins in bilayers

**alignment to membrane normal** **packing lipids around the protein** **removing overlaps and clashes**

Solvation, ions and box setup for MD

**water models and box geometry** **ion placement and neutralization** **checking overlaps and system sanity**

### **Session 3**

**Fee: Rs 31800** [Apply Now](#)

## **MD Protocols for Membrane Systems**

Force fields and parameters for membranes

**protein and lipid force field choices** **coarse grained vs all atom ideas** **mixing and compatibility issues**

Equilibration strategies for membrane systems

**restrained minimization and short MD** **releasing protein and lipid restraints** **monitoring area per lipid and thickness**

Production runs and trajectory management

**time step and ensemble choices** **long timescale**  
**planning** **checkpointing and storage practices**

#### **Session 4**

**Fee: Rs 34800** Apply Now

### Dynamics Analysis, Lipid Contacts & Reporting

Protein stability, orientation and dynamics readouts

**RMSD and RMSF in membranes** **helix tilt and rotation**  
**metrics** **pore radius and channel opening ideas**

Lipid interactions and local environment

**annular lipid contacts** **specific lipid binding sites**  
**cholesterol and headgroup preferences**

Deliverables: membrane MD report and figures

**summary plots for bilayer properties** **snapshots and**  
**movies of key events** **structure to mechanism**  
**narrative**