

3rd Annual Danish Workshop on Advances in Molecular Simulation 9-10 Dec 2025, AIAS, Aarhus University

Day 1	
8:30-9:00	Registration, coffee, poster set-up
9:00-9:15	Introduction
9:15-10:00	Chair: Weria Pezeshkian, KU Mohsen Sadeghi , Zuse Institut, Berlin. <i>Computational biology at the mesoscale: From membrane mechanics to cellular dynamics</i>
10:00-10:20	Beatrice Geiger , NBI, KU. <i>Membrane necks' split personality: the nuclear envelope under osmotic pressure</i>
10:20-10:40	Adria Bravo Vidal , NBI, KU. <i>Multi-body fluctuation-induced forces between membrane proteins: insights from mesoscale simulations</i>
10:40-11:10	Coffee break
11:10-11:55	Chair: Anna Duncan, AU Weria Pezeshkian , NBI, KU <i>Multiscale simulation of biomembranes: Bridging the gap between simple models and complex reality</i>
11:55-12:15	Amanda Dyrholm Stange , AU. <i>Toward a design framework for chemically modified RNA: Integrating cryo-EM, functional assays, and molecular dynamics</i>
12:15-12:35	Darian Yang , KU. <i>Driving MD sampling with dynamic experimental data for active refinement of conformational ensembles</i>
12:35-13:15	Lunch
13:15-14:30	Poster session 1
14:30-15:00	Coffee
15:00-15:45	Chair: Vincent Nieto, AU Matteo Degiacomi , University of Edinburgh. <i>Learning (from) protein dynamics</i>
15:45- 16:05	Lorena Zuzic , AU. <i>CHARMM and Amber force fields produce different outcomes in MD simulations of multicomponent systems</i>
16:05-16:25	Isabell Lindahl , NBI, KU. <i>Who induces what? Exploring membrane curvature generation and sensing of mitochondrial membrane proteins using coarse-grained MD</i>
17:00	Walk to dinner
18:15	Conference dinner at <i>Den Gamle By</i>
Day 2	
8:30-9:00	Coffee, arrival
9:00-9:20	Chair: Mikkel Dahl Andreasen, AU Lasse Messell Desdorf , AU. <i>Binding mechanism of NBCn2</i>
9:20-9:40	Luise Jacobsen , SDU. <i>A novel model for proton transport mediated by Uncoupling Protein 1</i>
9:40-10:25	Lucie Delemotte , SciLifeLab, Stockholm. <i>Voltage-gated ion channel activation and gating: insights from a family-wide molecular dynamics investigation</i>
10:25 – 10:55	Coffee break
10:55-11:40	Chair: Leonhard Starke, AU Ilpo Vattulainen , University of Helsinki. <i>Cell membrane structures as barriers and operating environments for biological functions</i>
11:40-12:00	King Ifashe , University of Oxford. <i>Molecular dynamics reveal conformational restriction eliminates cross-reactivity in engineered therapeutic T-Cell receptors</i>
12:00-12:20	Ali Asghar Hakami Zanjani , SDU. <i>Quantifying membrane geometry in molecular dynamics simulations</i>
12:20-13:00	Lunch
13:00-14:00	Poster session 2
14:00-14:20	Chair: Tommy Rosendahl, AU Daria Gusew , KU. <i>Modeling chemical exchange dynamics in proteins using molecular dynamics simulations and NMR relaxation</i>
14:20-14:40	Fabian Schuhmann , NBI, KU. <i>TS2CG 2.0 — the membrane builder</i>
14:40-15:00	Hassan Ghermezcheshme , SDU. <i>Realistic modeling of PET nanoplastics for interactions with a model lipid membrane</i>
15:00-16:00	Coffee and concluding remarks