6th CCPBioSim/CCP5 Multiscale Modelling Conference

Programme

Monday Afternoon – https://ukri.zoom.us/j/99655158845
Tuesday Morning – https://ukri.zoom.us/j/91045594839
Tuesday Afternoon – https://ukri.zoom.us/j/92992230968
Wednesday Morning – https://ukri.zoom.us/j/92212955259

Day 1 - Monday 31 March 2025

12:00 – 13:20	Registration and Lunch		
13:20 - 13:30	Welcome / Introduction (Session Chair: Marc van der Kamp)		
13:30 - 14:05	Alessandro Troisi		
	High-throughput screening of polymeric semiconductors		
14:05 - 14:25	Kirill Zinovjev		
	QM/MM without QM: electrostatic machine learning embedding for		
	enzymatic catalysis, ligand binding and more		
14:25 - 14:45	Anna Bui		
	Learning classical density functionals for ionic fluids		
14:45 - 15:05	Edoardo Donadoni		
	The impact of polymer coating on nanoparticles interaction with lipid		
	membranes explored by coarse-grained molecular dynamics		
	simulations		
15:05 – 15:35	Coffee		
15:35 - 16:10	Chantal Valeriani		
	From active polymers' assembly to mechanical properties of a bacterial		
	biofilm		
16:10 - 16:30	Carlo Fonte		
	From Electrons to Reactors: ab initio multi-scale simulation of methane		
	oxidation over palladium oxide		
16:30 – 17:00	Flash Talks A		
17:00 - 18:30	Poster Session A - Odd Numbers		

Day 2 - Tuesday 1 April 2025

08:30 - 09:00	Coffee (Session Chair: Charlie Laughton)	
09:00 - 09:35	Matthias Schmidt	
	Neural functionals and unexpected symmetries in statistical mechanics	
09:35 - 09:55	Marko Hanzevacki	
	Reaction mechanism of choline degradation by choline trimethylamine-	
	lyase (CutC) revealed with QM/MM modelling	
09:55 - 10:30	Lorna Dougan	
	Current challenges and opportunities in translating the mechanical	
	properties of biomolecules across lengthscales	
10:30 - 11:05	Coffee	
11:05 - 11:40	Katarzyna Swiderek	
	Deciphering the role of protein electrostatic field in enzymatic catalysis:	
	A case study of three distinct heterogeneous Proteasome 🛚 -subunits	

11:40 - 12:00	Tom Headen		
	Wide Q-range Total Neutron Scattering: Simulation Based Data-		
	Refinement for Multi-Scale Systems		
12:00 - 13:30	Lunch (Session Chair: Paola Carbone)		
13:30 - 14:05	Valentina Erastova		
	Let's talk about dirt: from environmental sorptive materials of Earth to		
	biosignatures in Space		
14:05 - 14:25	Maria Grazia De Angelis		
	Multiscale and machine learning-assisted models for the design of		
	materials supporting net zero, energy transition, and circular economy		
14:25 – 14:45	Mark Miller		
	Superselectivity and Nucleation in Biomolecular Condensates		
14:45 – 15:15	Coffee		
15:15 – 15:50	Rebecca Wade		
	Multi-resolution molecular simulations to investigate the interplay		
	between cytochrome P450 interactions, conformational variability and		
	function		
15:50 – 16:10	Alberto Pérez de Alba Ortíz		
	Novel Bayesian approaches to biomolecular free-energy calculations:		
	from ion-responsive polysaccharide rigidities to nanoplastic-induced		
	protein dissociation		
16:10 – 17:00	Flash Talks B		
17:00 – 18:30	Poster Session B – Even Numbers		
19:00	Conference Dinner		

Day 3 - Wednesday 2 April 2025

08:30 - 09:00	Coffee (Session Chair: Stephen Cox)
09:00 - 09:35	Benedetta Mennucci
	Multiscale strategies to investigate light-responsive proteins
09:35 - 09:55	Thomas Sayer
	Small polaron transport in very large lattices with accurate numerics
09:55 - 10:15	You Lu
	Py-ChemShell's Biomolecular Simulation Workflow and Recent
	Advances
10:15 - 10:35	Jean-Marc Lude
	Molecular Density Functional Theory with Atomistic Dipolar Solvent to
	Study Pressure Effect on a Diels-Alder Reaction
10:35 - 11:05	Coffee
11:05 - 11:25	Shakir Ali Siddiqui
	Local Electric Field-Guided Redox Potential Tuning in de novo Proteins
11:25 - 11:45	Dmitry Nerukh
	Upscaling Simulations: Integrating Molecular Dynamics with
	Hydrodynamics
11:45 - 12:20	Jochen Blumberger
	Simulating Electronic Quantum Dynamics on the Nanoscale (10-
	100nm)
12:20 - 14:00	Lunch and Close

Posters With Prizes Sponsored by the RSC Journals







Poster	Presenter Name	Poster Title
Number		
1	Daymahayan Fiana	Duadiation anatomic about discussion was this
1	Baumhauer, Fiona	Predicting azeotropic phase diagrams using machine
		learned classical density functional theory
2	Brukhno, Andrey	Shapespyer: a Python driven toolchain for soft matter
		simulations
2	Challandle - Dairechi	De also assistante de la forma field forma a field forma a field forma field formation formation for a field for a fie
3	Chattopadhyay, Rajorshi	Development of polarizable force field for La3+ in Cl-
		bearing hydrothermal fluids
4	Doveiko, Daniel	Binding Energy Calculations of Anthracene and
		Rhodamine 6G H-type Dimers: A Comparative Study of
		DFT and SMD Methods
5	Epstein, Alexander	Understanding Hydrophobicity in Complex Geometries
		via Classical Density Functional Theory
6	Fairchild, Connie	A Minimal Model for the Frequency-Dependent
		Conductivity of Ionic Liquids
7	Findlay, Elliot	Modelling Soil-release Polymers on Fabric Surfaces:
		Multiscale Insights from Dissipative Particle Dynamics
		and Enhanced Sampling MD Simulations
8	Fitkin, Arielle	A computational investigation of metal-organofluorine
		interactions and their role in selective metal deposition
9	Freeman, Colin	Mineral Control over Molecular Binding and Decay
10	Galappaththi Guruge,	Computational insights into structural changes in
	Amali	PEODT:PSS interfaced with water
	7 111011	1 200 m oo meenassa man mater
11	Gopalakrishna Rao, Aditya	Multiscale simulations of the mechanism of a de novo
		designed Diels Alderase photoenzyme catalysing [2+2]
		cycloadditions: implications for photobiocatalyst design
12	Hayton, John	Silver Iodide in Water – A Machine Learned Interatomic
		Potential with Electrostatics
13	Hodala, Aydin	The effect of aggregation on the pKa of oleic acid
1.4	lia Henriian	Discovering Blood Brain Bornier Borneschle Comment
14	Jia, Hengjian	Discovering Blood Brain Barrier Permeable Compounds

		with Cliques Descriptors
15	Igaev, Maxim	Microtubule dynamics are defined by conformations and stability of clustered protofilaments
16	Kanagarajan, Ajeeth	Negative Chemotaxis of polymeric vesicles against lactic acid concentration gradient
17	Kazmierczak, Magdalena	Machine learning potentials for accurate and efficient multiscale enzymatic Diels – Alder reaction modelling
18	Lei, Li	Coarse-grained investigation of the adsorption behaviour of antibodies at water–hexadecane interfaces
19	Liu, Yufeng	Prediction of rhamnolipid partitioning into lipid bilayers using coarse-grained molecular dynamics with the Martini 3 force field
20	Meadows, James	Controlling Polymorph Crystallisation Using Structured Ternary Fluids
21	Meulemans, Arne	Organic Thermoelectric Polymers: atomistic modelling of semicrystalline P3HT
22	Morado, Joao	Enhancing Electrostatic Embedding for ML/MM Free Energy Simulations
23	Morgan, Liam	Understanding the Behaviour of Hole States for Copper Substitutionals in MgO
24	Novi Inverardi, Giovanni	Structural and dynamical assessment of the adsorption of silica oligomers on biomolecules via Molecular Dynamics
25	Ramos, Carlos	Capturing complexity in enzyme catalysis: Multidimensional Free Energy Landscape Explorations Using Multiscale Methods to Fight Infection Diseases.
26	Sahnoune, Meriem	Coarse-grained modeling of insulin adsorption on plasticized polyvinyl chloride surfaces
27	Thomas, Dominic	Understanding electrochemical interfaces with neural functional theory
28	Tomlinson, Emma	Understanding DNA-Protein Interactions in Eukaryotic Topoisomerases through atomistic molecular
29	Underhill, Juno	Spectral tuning in a de novo riboflavin protein

30	Yang, Chao-Yu	FibrilGen: a program to template cross-β
		nanostructures at the atomic level
31	Ymeraj, Matilda	Structure-based virtual screening targeting APSR:
		exploring covalent inhibitors against Pseudomonas
32	Zhang, Xibei	Transforming Tuberculosis Care: Rapid Accurate and
		Reliable Computer-Based Prediction of Drug Resistance
		to Guide Targeted Treatments
33	Zhou, Fulu	Applicability of the Thermodynamic and Mechanical
		Route to the Young Equation for Rigid and Soft Solids: A
		Molecular Dynamics Simulations Study of a Lennard-
		Jones System Model
34	Donadoni, Edoardo	Characterizing graphene biointeractions: coarse-
		grained modeling of ion binding, small molecule
		adsorption and protein corona formation
35	Güven, Jasmin	Protocols for free energy predictions for beta-
		lactamases: insights from cross-class inhibitors
36	Kearney, Eoin	Coarse-grained molecular dynamics simulations of
		surfactant membrane-water partitioning
37	Kolli, Hima Bindu	Multiscale Simulation Scattering Intensity Calculator
		(MuSSIC):Validation and Application to Soft-Matter
		Systems
38	Petho, Andras	Investigating the conductivity of multi-heme
		cytochromes