

Posters

Kathrin Bach	Unraveling the role of the human Sorting and Assembly Machinery (SAM) complex in premature aging
Anika Baier	Cryo-EM Analyses of the Conformational Spectrum of a Nucleotide-Bound ABC Transporter in Various Membrane Mimics
Florestan Bilsing	Structural basis of Protein O-mannosylation by Pmt4
Claudia Bohg	From High-Throughput Screening to Structural Insight in Lipid Bilayers
Andreas Boland	Structure of ATTRv-F64S fibrils isolated from skin tissue of a living patient
Alexandra Deaconescu	Autoinhibition Is Not a Universal Feature of Transcription-Repair Coupling Factors
Eddy Javier Figuerola Marquez	Mechanisms of Wnt protein secretion
Tom Goetze	Investigation of Cellular Ultrastructure in Cyanobacteria using in situ Cryogenic Electron Tomography
Monika Gunkel	Illuminating Cryo EM in Cologne
Zeyaul Islam	Characterizing the higher oligomeric form of SOX5: A transcription factor involved in chondrogenesis
Kevin Janson	Novel sample delivery for small nanoparticles and biomolecules for cryo electron microscopy
Athithya Jeyarasaa	Structural Analysis of SLC9C1 – A VHH-Driven Approach
Michael Kamel	Conformational plasticity of P-glycoprotein and its modulation by verapamil.

Posters

Alexej Kedrov	Substrate-induced assembly and functional mechanism of the bacterial membrane protein insertase SecYEG-YidC
Sagar Khavnekar	Towards Understanding RNA Biology with Cryo-Electron Tomography: Moving from Breathtaking Renderings to Quantitative, Context-Aware RNA Cartography
Uljana Kravcenko	Structural insights into the molecular architecture of synaptic vesicles
Anuj Kumar	Cryo-EM structure of the ATP driven Methyl-coenzyme M reductase activation complex
Sunil Kumar	Engineered Biocompatible Virus-like Particles: A Promising Nanocarrier for Targeted Anti-cancer Drug Delivery in Cervical Cancer Treatment
Mariya Lazarova	SLC9C1 at activating hyperpolarizing conditions – the endeavor of cryo-EM in proteoliposomes
Yuan E Lee	Improved Cryo-EM Resolution of Enterococcus hirae V-ATPase: Structural Insights into the Full Complex
Nicolas Lichti	Investigation of PE/PPE proteins and MmpL/S transporters of Mycobacterium tuberculosis
Mauro Maiorca	Uncovering Hidden Functional States in Cryo-EM Datasets with EMPROVE
Daniel Mann	Near atomic resolution and chemical contrast of ice-embedded proteins from scanning transmission electron microscopy
Vasilii Mikirtumov	Activation of RyR1 in native membranes
Agnes Moe	Molecular architecture and lipid transfer between the two mitochondrial membranes

Posters

Tomas Pascoa	Mechanism and in situ architecture of a class II benzoyl-CoA reductase
Martina Peter	Nutrient import in mycobacteria: Towards understanding the mycobacterial outer membrane
Maximilian Ruettermann	Structural and functional analysis of GTPase regulators
Jakob Ruickoldt	CryoCrane - An open source GUI for the visualisation of cryo-EM screening data sets
Daniel Schaefer	Combining phase-contrast STEM imaging with electron energy loss spectroscopy
Jan Hannes Schaefer	Dissecting the crosstalk between transthyretin amyloids and the extracellular matrix
Klara Theresa Scholtissek	Unleashing the activity of PC flipping ATP10A - a driver of endocytosis
Jan Michael Schuller	Dissecting the Wood-Ljungdahl Pathway - Molecular Basis of Acetogenesis
Heesang Song	Structural insights into phosphoglucomutase 1 as a glycogen metabolism regulator
Muratha Sottatipreedawong	Structure and function of mitochondrial membrane folding complex
Irene Vercellino	Cryo-EM structures of naturally occurring dimeric photosystem II complexes lacking the Mn_4CaO_5 cluster
Carl von Hallerstein	Atomic-level membrane protein structure determination in vacuum using ES-IBD and cryo-EM
Shutian Wang	Structural analysis of the insulin receptor in native vesicles by cryo-ET subtomogram averaging

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Céline Zheng-Gérard **Structural investigation of the 60 kDa potassium channel TASK-1 using cryo-EM**