

Welcome to Integrated Structural Biology SciLifeLab 4th users meeting

December 10, 2024



Location: Air and Fire, Tomtebodavägen 23a, Solna

December 10, 9-16

9.00-9.10	Welcome and introduction
9.10-9.30	Oscar Jungholm, Karolinska Institute Crystals, KRAS, and Core Collaborations: My Journey Through Protein Science
9.30-9.50	John Cowgill, Stockholm University Structural basis for activation and potentiation in a human $\alpha 5\beta 3$ GABAA receptor
9.50-10.10	Nicholas Croy, Uppsala University Investigating the catalytic mechanism of Rubisco with high-resolution cryo-EM
10.10-10.40	Fika
10.40-11.00	Julia Kinsolving, Karolinska Institute Structural and functional characterization of Frizzled-7 for drug discovery
11.00-11.20	Josy ter Beek, Umeå University A first glimpse of a Gram-positive Type 4 Secretion system that facilitates the spread of antibiotic resistance
11.20-11.50	Marten Chaillet, Utrecht University 3D template matching allows automated picking in cryo-electron tomography
12.00-13.00	Lunch
13.00-14.30	Flash talks from infrastructures Protein Production Sweden, Tomas Nyman, Karolinska Institute ProLinC, Dean Derbyshire, Linköping University Cryo-EM, Tim Schulte, Stockholm University Swedish NMR Centre, Cecilia Persson, University of Gothenburg Structural Proteomics, Simon Ekström, Lund University PReSTO, Martin Moche, Karolinska Institute
14.30-15.00	Fika
15.00-15.20	Arvid Eskilson, Linköping University Functional mutations of USP14 reveal dynamic networks connecting the active site and proteasome interaction surface.
15.20-15.40	Maximilian Kahle, Stockholm University CryoEM structure of the NorQD metallo-chaperone complex
15.40-16.00	Wrap-up