



Membrane transporters and receptors in Brain – Nissen laboratory

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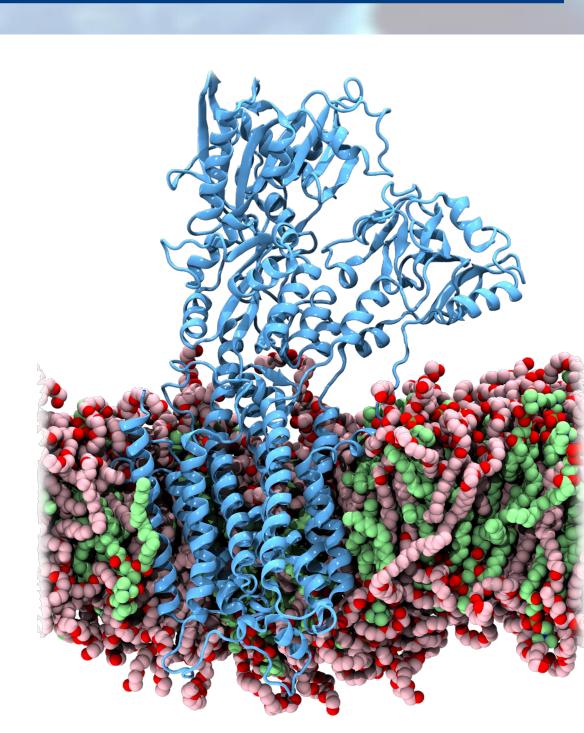
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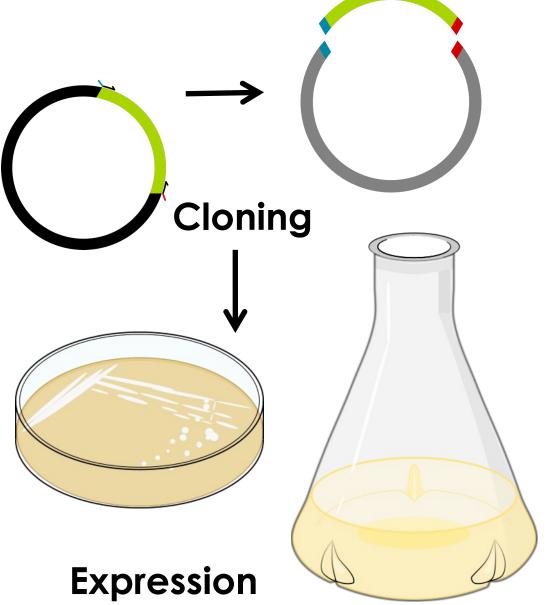
BACKGROUND/OVERVIEW The Nissen group investigates molecular mechanisms of membrane transport processes and biomembrane structure of importance in neuroscience.

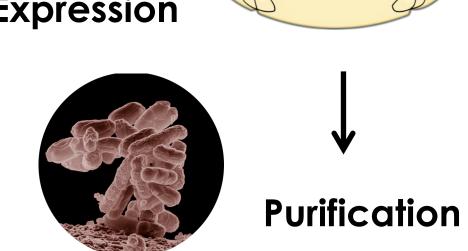
Methods: mainly focused on cryo-electron microscopy (Cryo-EM), biochemistry and biophysics of protein interactions and dynamics, and molecular cell biology using fluorescence microscopy and cryo-electron tomography. Key topics: membrane transporters and receptors in brain that are affected in neurological and psychiatric disorders, including P-type ATPases, Nat dependent neurotransmitter and chloride transporters (SLCs), insulin and sortilin receptors. We study structure, function and mechanisms of regulation, effects of disease causing mutations, and aim also for structure based drug discovery and protein engineering. We also engage in mathematical modelling of molecular networks in brain.

Research Areas: Structural neurobiology, Membrane transport and signalling, Cryo-EM, biochemistrybiophysics, drug discovery, network modelling.



Workflow





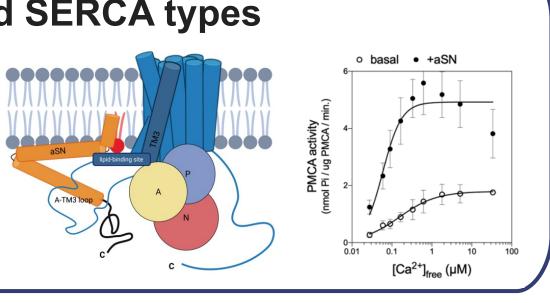
Bacteria

Mammalian cells

Projects

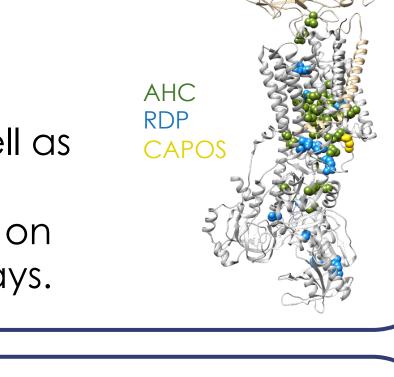
Ca²⁺-ATPases in neurons – PMCA and SERCA types

- Characterization of isoforms.
- Regulation by lipids, autoinhibition a-synuclein.



Na⁺,K⁺-ATPase

- Study of disease variants causing rare neurological disorders (AHC, CAPOS) associated with encephalopathies as well as cognitive and motor disabilities.
- Characterization of isoforms with a focus on specific functions and regulatory pathways.



Phospholipid flippases

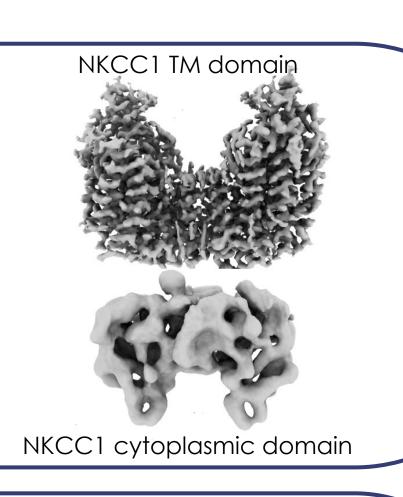
- Structural and functional characterization of yeast and mammalian "lipid-pumps".
 - Substrate identification
 - Mechanism of autoinhibition by lipids and termini.
 - Physiological function.

Drug discovery

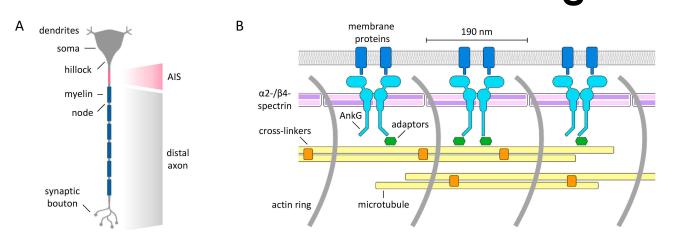
 Compound screening for the inhibition of Cu-ATPase and its disease mutants for the potential treatment of Wilson's disease. A genetic disorder characterized by accumulation of copper in the liver and brain.

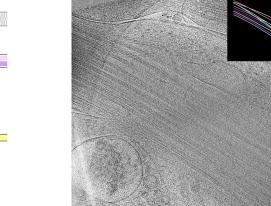
Solute carriers (SLC)

 Characterization of secondary active transporters carrying ions, neurotransmitters, amino acids, polyamines and phosphate.



Ultrastructure of the axon initial segment (AIS)

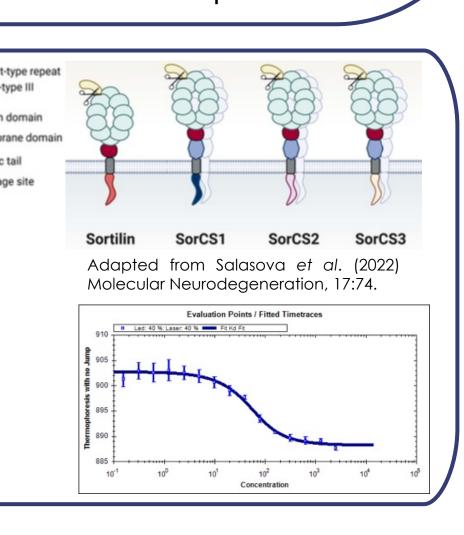




- Cultivation of primary neurons.
- Cryo-Correlative Light and Electron microscopy & Cryo-Electron Tomography for the investigation of AIS molecular structure and identification of receptors and transporters.

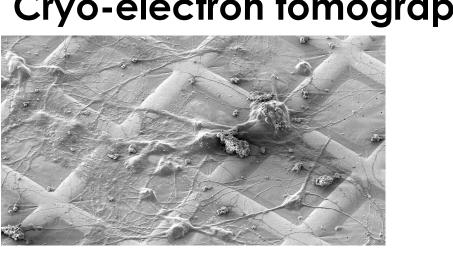
Sortilin receptors

 Structural and biophysical characterization of interactions with neuronal receptors essential for synaptic plasticity (TrkB, p75^{NTR}, NMDA), and other disease-related proteins (ApoE).

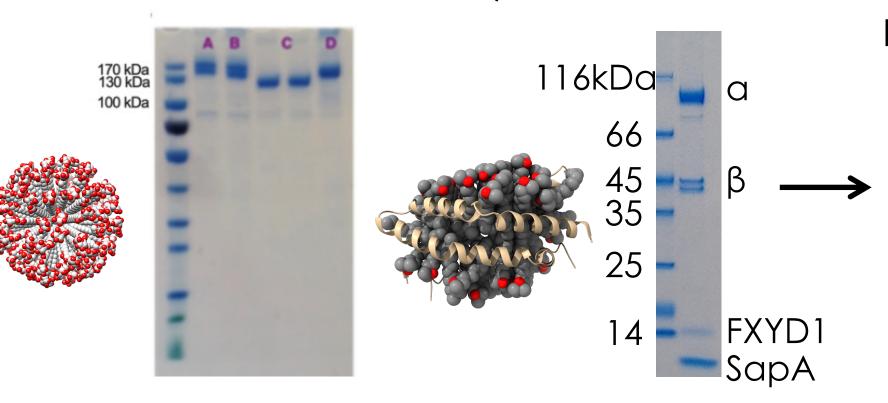


3D reconstruction

Expansion microscopy Cryo-electron tomography



Use of membrane mimetics, nanodiscs



 Detergents, e.g. PMCA variants Nanodiscs, e.g. Na+,K+-ATPase

Functional characterization

- Activity assays
- Fluorescence assays
- Proteoliposome assays Uptake assays
- Protein interactions

dynamics/stability

Protein

Cryo-EM

Data processing

Projects

BSc projects Course projects MSc projects

Educational tracks

Molecular Biology Molecular Medicine Nanoscience Medicinal Chemistry Biology Engineering

REFERENCES

- 1. Monomeric a-Synuclein activates the Plasma Membrane Calcium Pump. EMBO J. 2023, Dec 1;42(23):e111122
- 2. Cryo-EM structure of the human NKCC1 transporter reveals mechanisms of ion coupling and specificity. EMBO J. 2022, Dec 1;41(23):e110169.
- 3. Activation and substrate specificity of the human P4-ATPase ATP8B1. Nat Commun. 2023, 14(1):7492
- 4. Structural Investigations of Full-Length Insulin Receptor Dynamics and Signalling. JMB 2022, 434(5):167458
- Structural insights into the inhibition of glycine reuptake. Nature 2021, 591 (7851):677-681

















