Computational Modeling Core Mini-symposium

Date: Wednesday May 2nd

Location: University of Chicago, Gordon Center for Integrative Science (GCIS), room W301

Schedule:

8:00 AM Breakfast and welcome

AIM 1

- 8:30 Lei Huang Webserver for automated optimization force field parameters
- 8:50 Christopher Mayne VMD interface and automated force field optimization
- 9:10 Wei Han Development of a multiscale CG model for protein folding, aggregation and dynamics in membrane
- 9:30 Emad Tajkhorshid Developing an atomistic, highly mobile membrane model (HMMM) and its wide applications
- 9:50 Fatemeh Khalili Asymmetric membrane potential and ionic concentration
- 10:10 Janamejaya Chowdhary polarizable force field for lipid membranes

AIM 2

- 10:30 Shahid Islam KcsA modeling with spin label distribution from ESR/DEER histogram distances
- 10:50 Fatemeh Khalili Na/K Pump probed by LRET
- 11:10 Elia Zomot The crystallized structure of the carnitine/ γ —butyrobetaine antiporter CaiT is a substrate-releasing conformation
- 11:30 Hang Yu Membrane Sculpting by F-BAR Domains studied by Molecular Dynamics Simulations

Break for lunch

AIM 3

- 12:30 PM Avisek Das Conformational transition pathways from coarse grained ANM normal modes
- 12:50 Mert Gur Transition pathways of dopamine transporters explored by combining molecular dynamics simulations and monte carlo sampling of collective modes
- 1:10 Mahmoud Moradi Optimizing Transition Pathway for Large—Scale Conformational Changes in ABC Transporters Using Non—Equilibrium Work
- 1:30 Giray Enkavi Chemomechanical Coupling in MFS Transporters
- 1:50 Wei Han Proton Transfer Pathway in CIC Antiporter

AIM 4

2:15 — Wonpil Im — restrained ensemble for solid state NMR (special talk of 30 min)