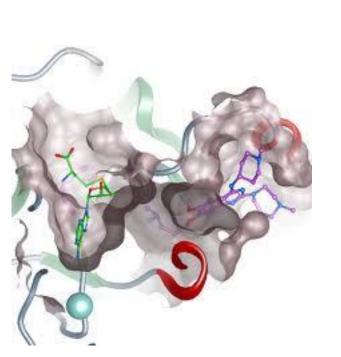


HPC in Drug Discovery



Ashutosh Tripathi, Ph.D. Bankaitis Lab Department of Molecular and Cellular Medicine TAMHSC

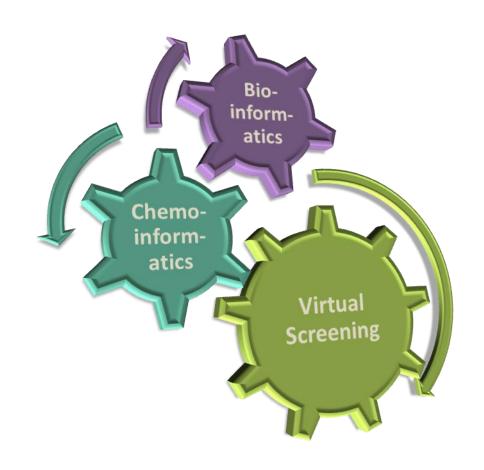


Technologies in Drug Discovery





- Expensive
- Protracted
- Difficult

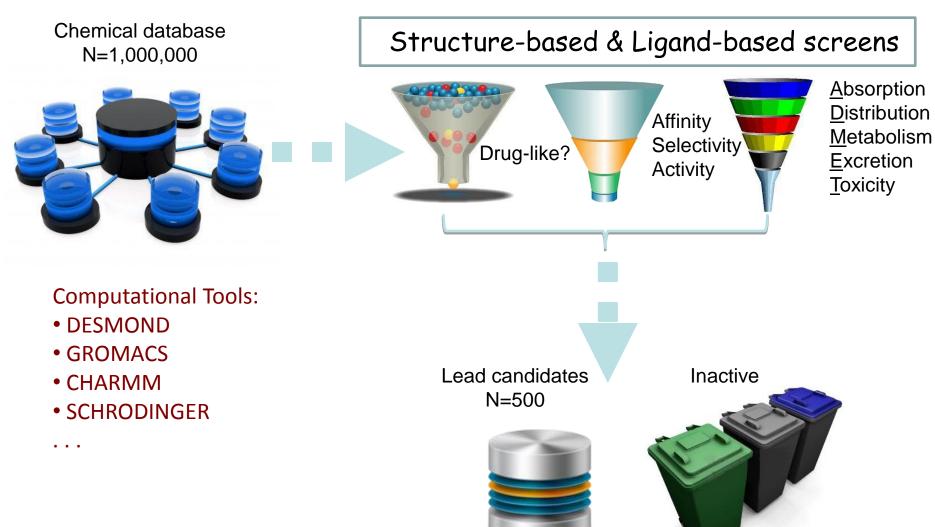


HPC MD-based Simulations

- Inexpensive
- Very fast
- Useful for early stages



In silico Screening Complements High-throughput Screening





Drug Discovery: Driven by Computation and Experiments



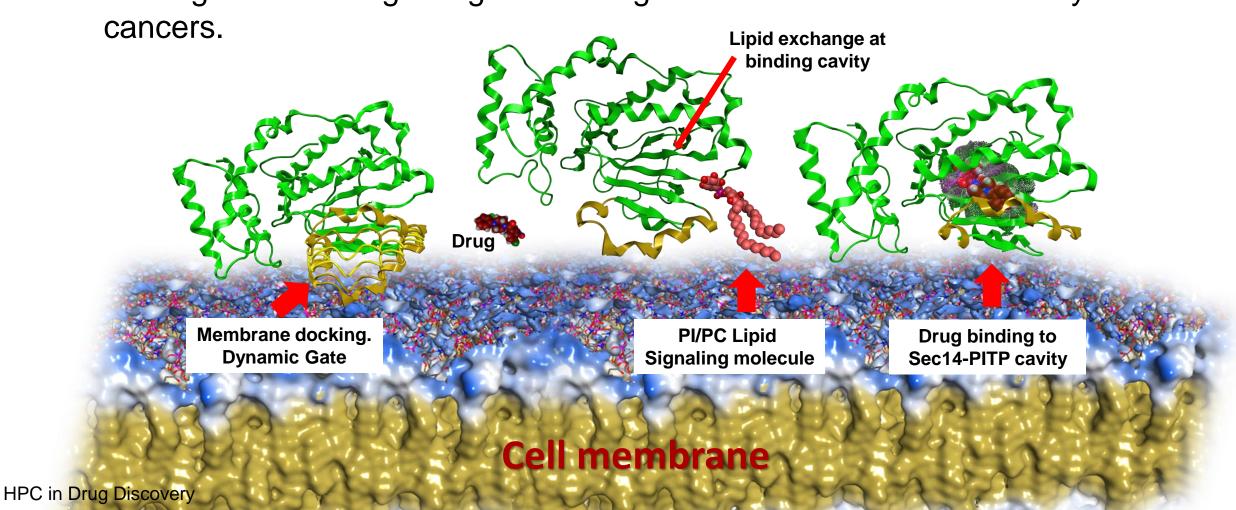
HPC in Drug Discovery



Phosphatidylinositol Transfer Proteins (PITPs) Important in Cell Function

Important in Lipid-mediated cell signaling and metabolism.

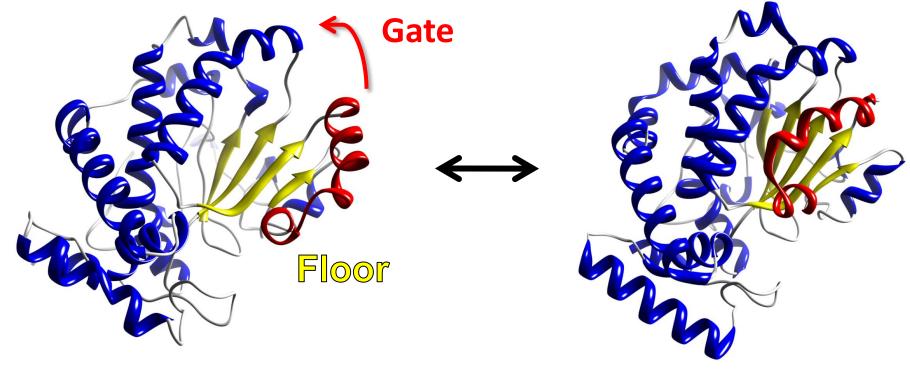
Derangement in Signaling: Neurodegenerative diseases and many forms of





Phosphatidylinositol Transfer Proteins (PITPs) Critical in Cell Function

- Important in Lipid-mediated cell signaling and metabolism.
- Derangement in Signaling: Neurodegenerative diseases and many forms of cancers.
- Undergoes Conformational Changes: Helical Gate Mediates Lipid Access/Exchange



Sec14-OPEN

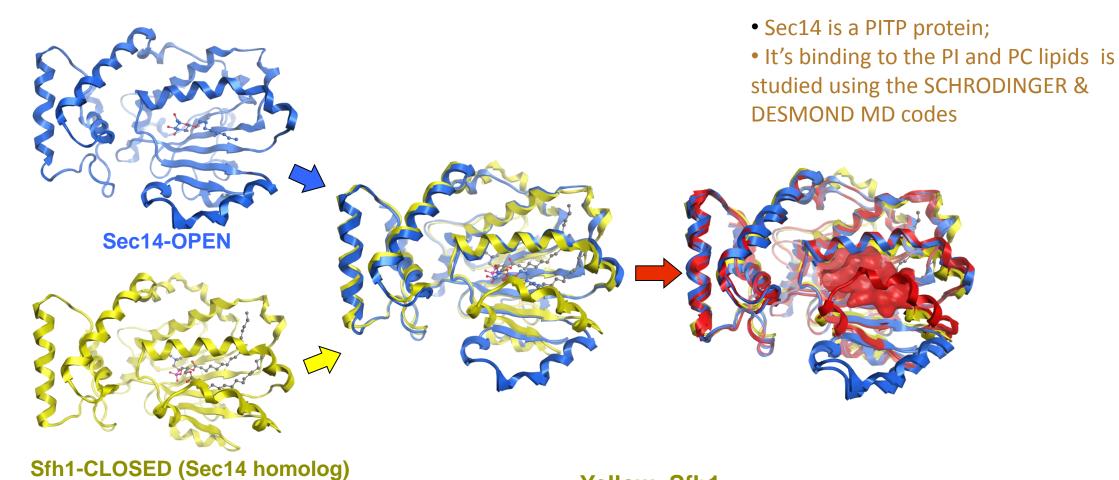
'Transition state'?

CLOSED

Schaaf et. al. Mol. Cell. (2008)



Homology Modeling of Sec14 (Closed Conformation)



Yellow: Sfh1

Blue: Sec14-OPEN

Red: Sec14 Homology Model-CLOSED

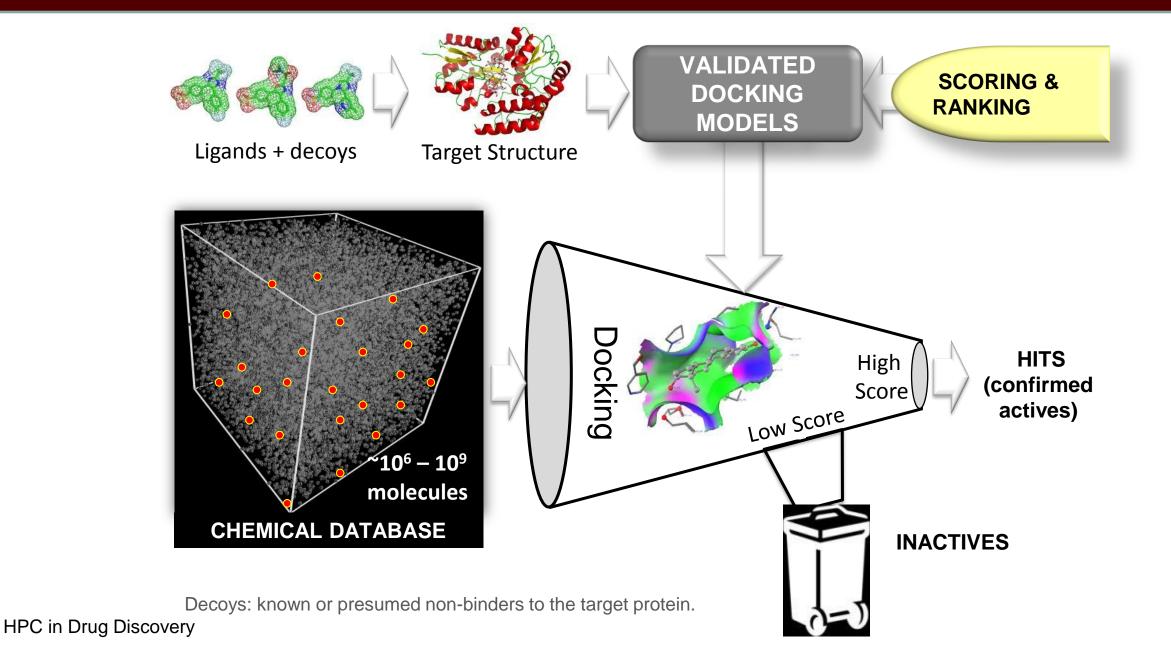
Identity: 62 %

Similarity: 78 %

Homology: 77 %

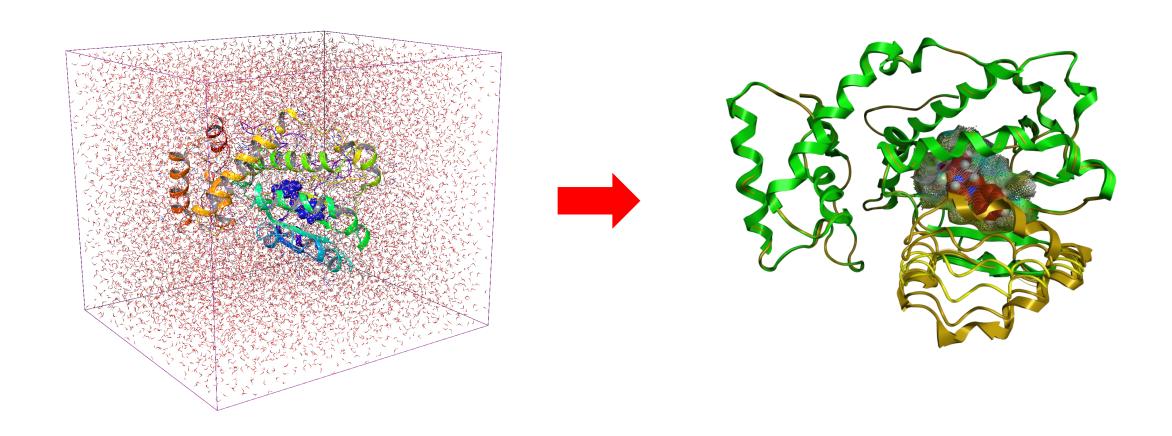


Docking-Based Virtual Screening





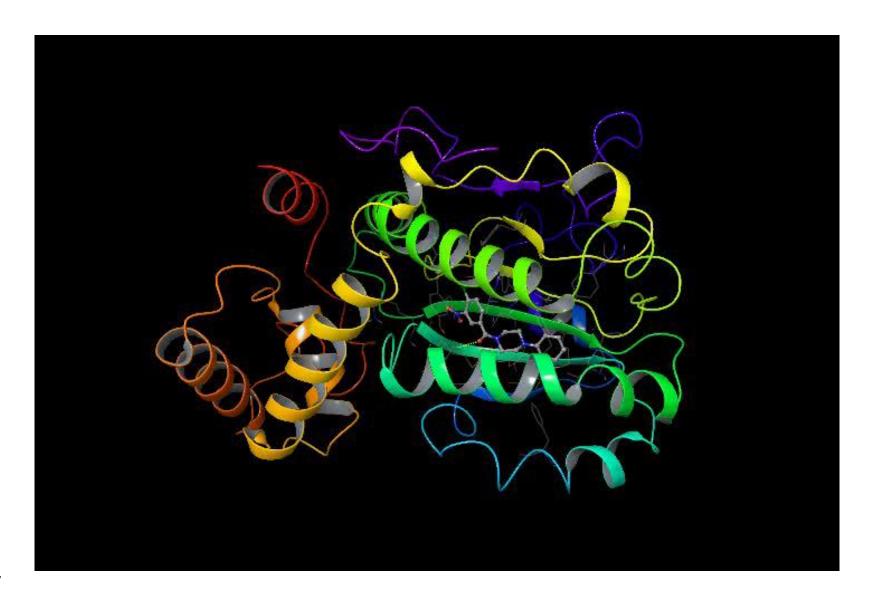
Sec14 Homology Model: All-Atom MD Simulation



HPC in Drug Discovery

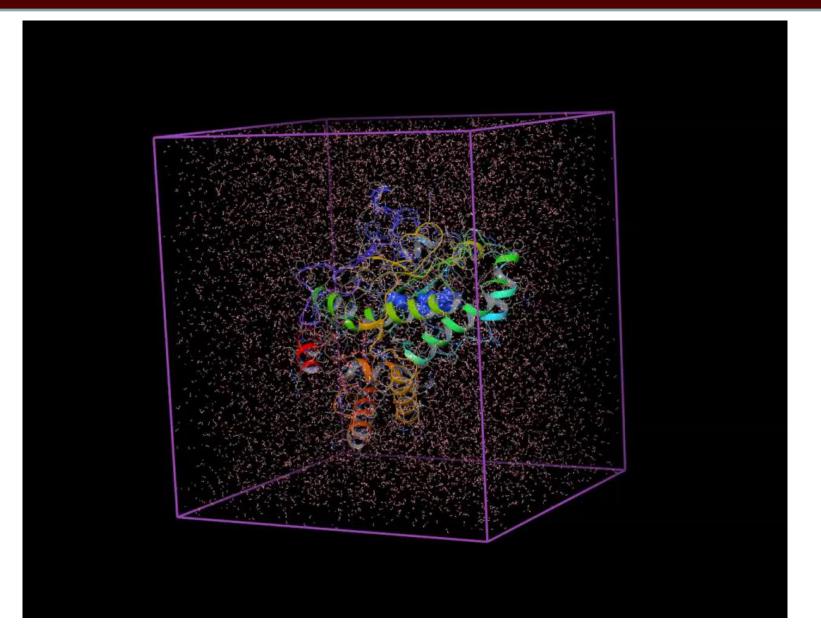


Simulation of Drug Binding to Protein Using DESMOND MD Code



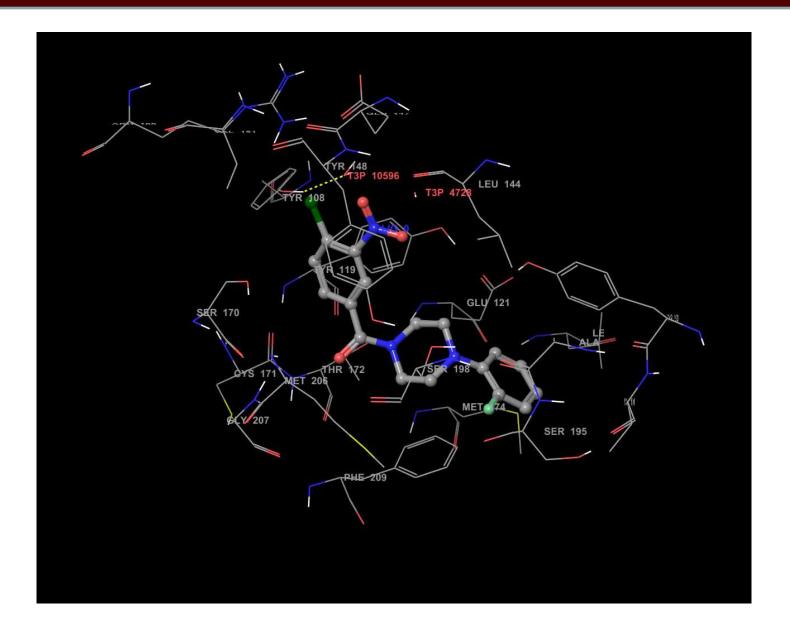


All-Atom Simulation of Sec14 in Explicit Water Molecules (used DESMOND MD Code)





Small molecule (drug) vibrating in Sec14 binding pocket (Used DESMOND MD Code)





Conclusion



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- ■Bachelor of Pharmacy, Institute of Engineering and Technology, M.J.P. Rohilkhand University, India.



Research Interest

- Computer-aided drug design.
- Algorithm and software development for designing new drugs.
- Cancer therapeutics.
- Clinical informatics.
- ADME/Tox QSAR modeling.