

Quantitative Studies of Membrane Transport in the Dawn Light of Exascale Computing

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Acknowledging

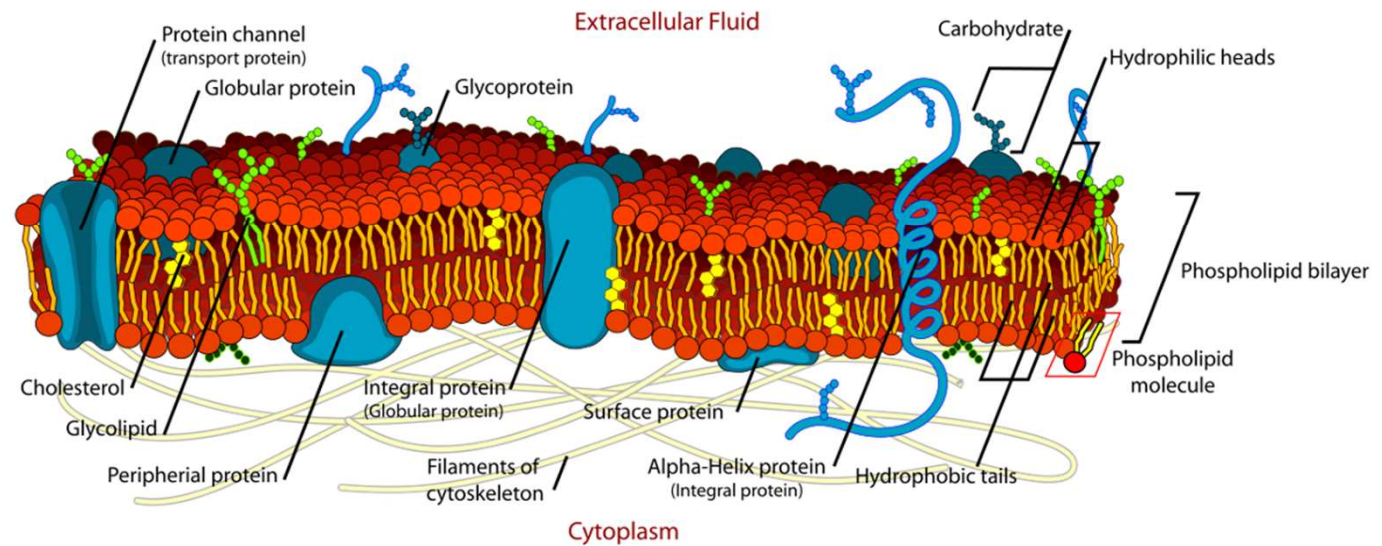


- Goal: Predict biological function from jiggling-and-wiggling of atoms

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- Obstacle 1: Time scale
- Obstacle 2: Intrinsic and artifactitious fluctuations
- Ways to overcome or circumvent
- Free or biased:
 - brute force vs biasing for enhanced sampling
 - down to the root: what drives the processes

~3 trillion water molecules

Cell: separation but not isolation
Membrane with machineries



[https://commons.wikimedia.org/wiki/File:Cell membrane detailed diagram en.svg](https://commons.wikimedia.org/wiki/File:Cell_membrane_detailed_diagram_en.svg)

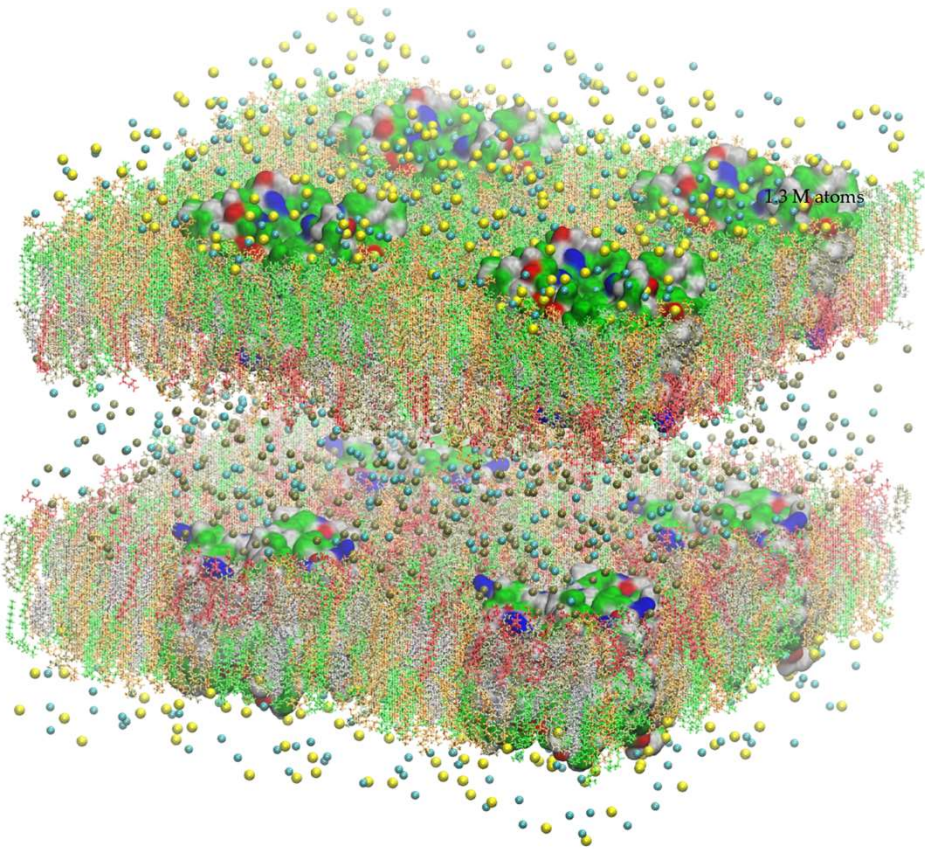
Study biological processes in a predictive manner?



From simplicity to simplicity via complex dynamics

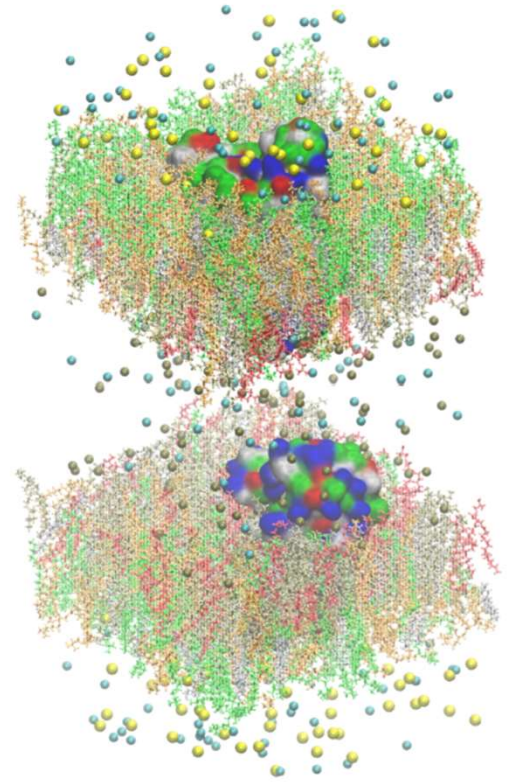
- Brute force molecular dynamics
 - Milliseconds by steps of femtoseconds?
 - Accuracy of force fields
 - Millions of degrees of freedom

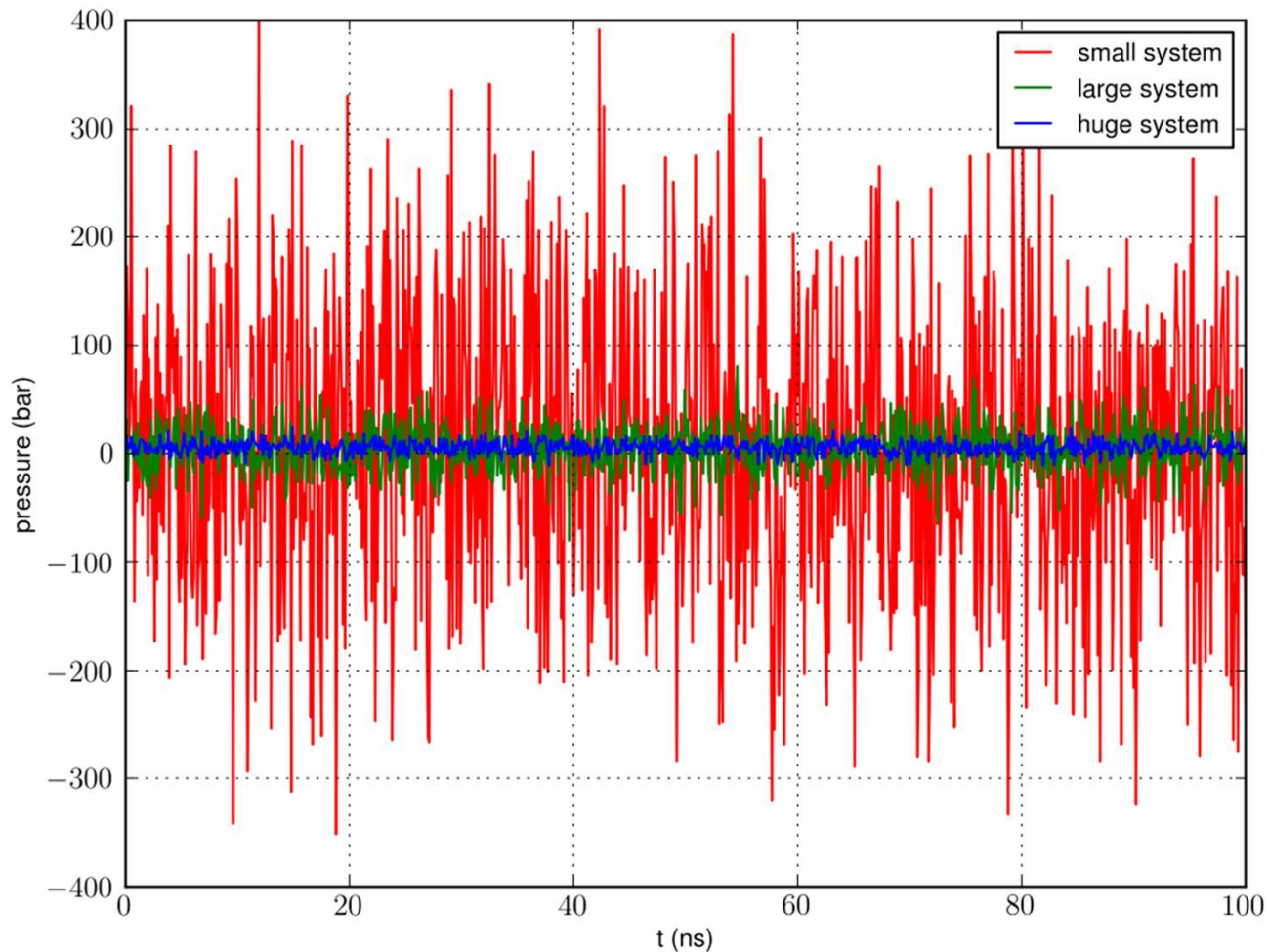
- The driving “force”
 - Free energy gradient
 - “Mechanical” forces
 - Entropic “forces”

Unbiased, unaccelerated MD
Accelerated MD: REMD, *etc.*



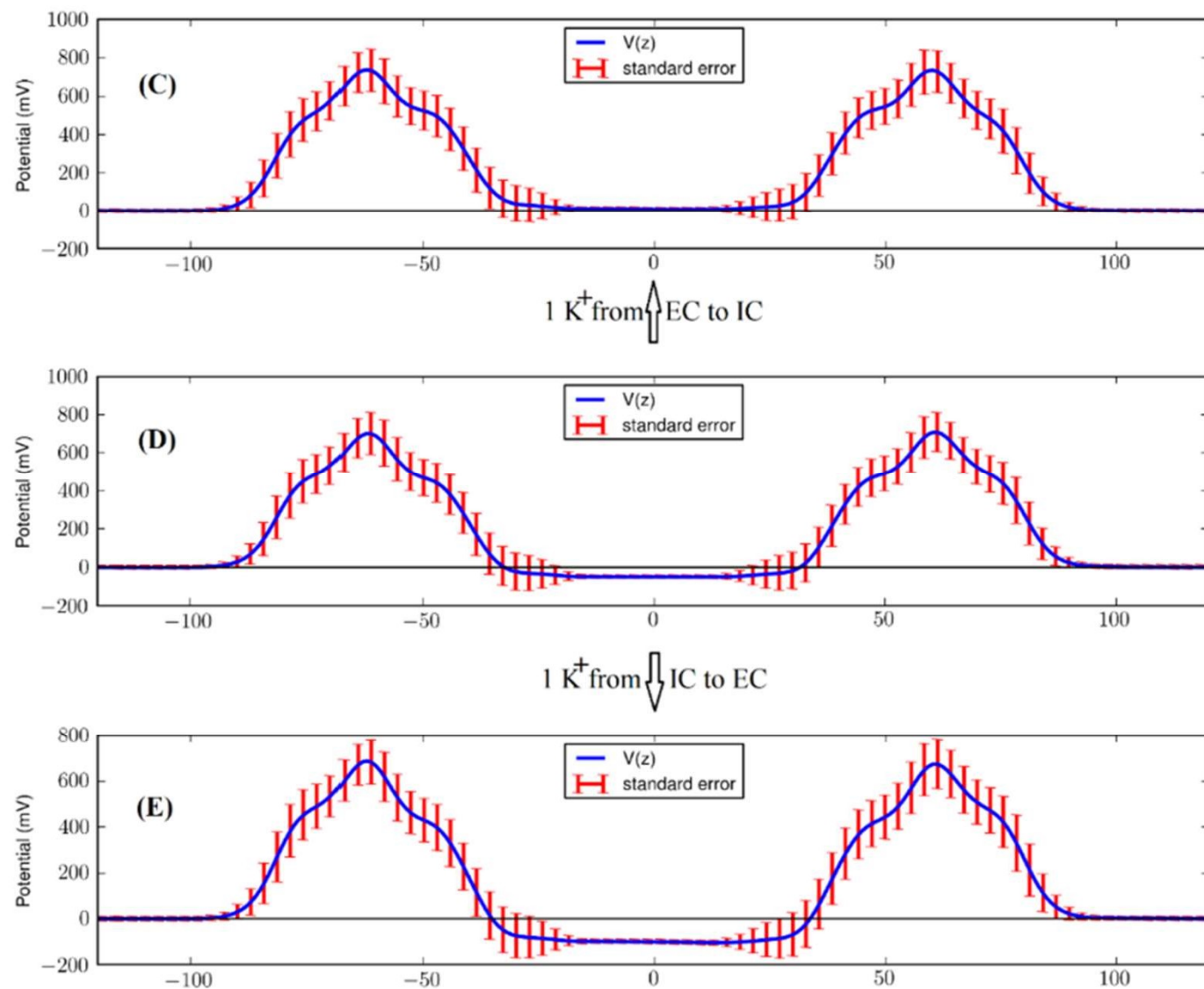
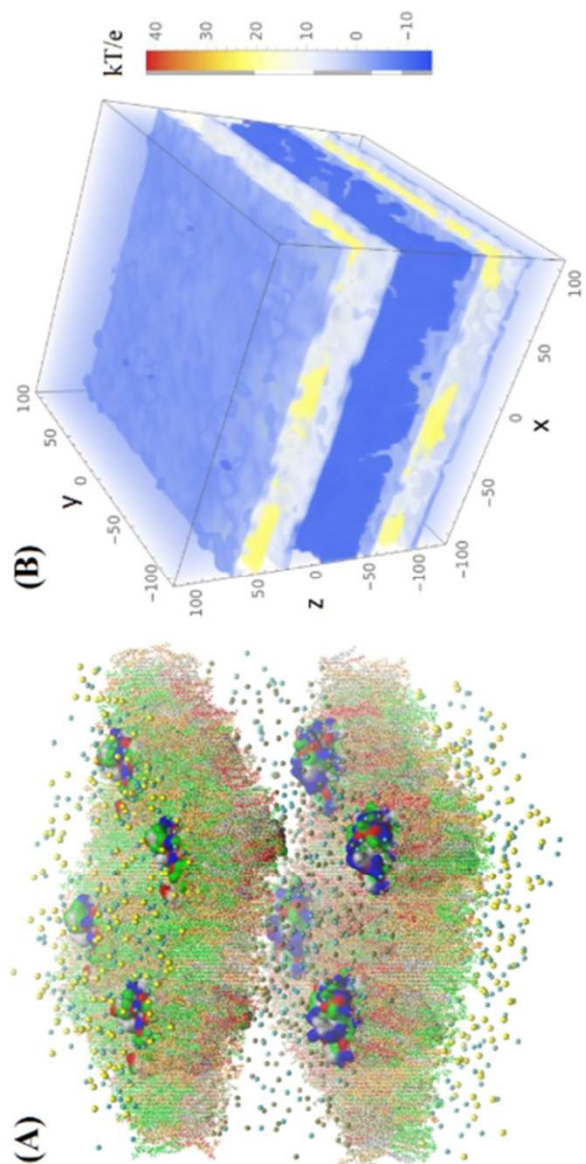

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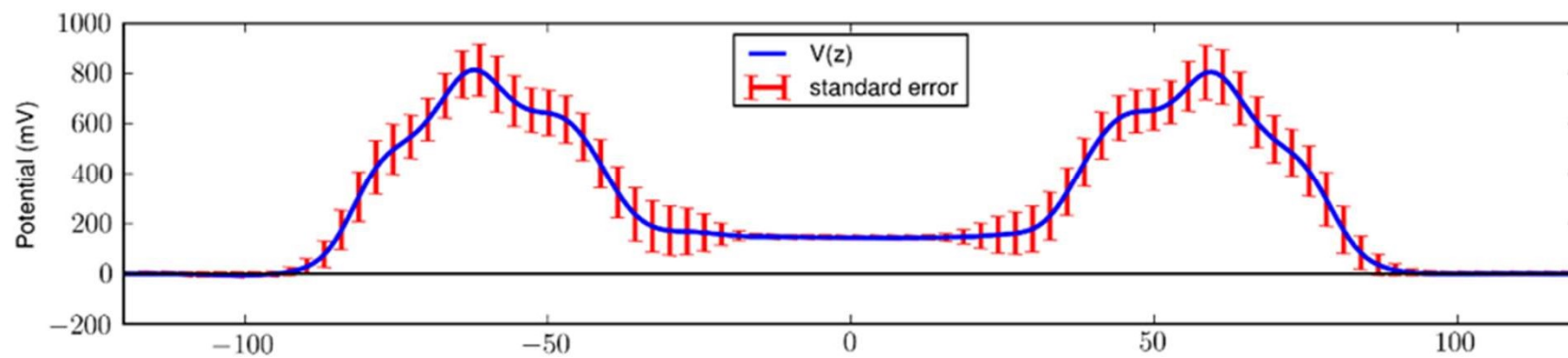
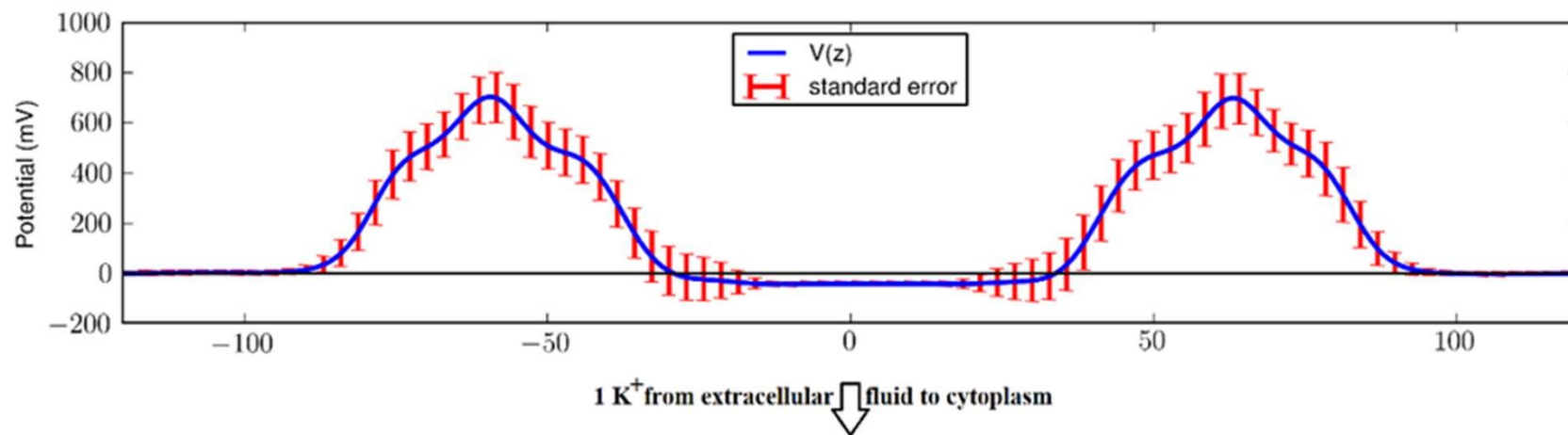




NPT
ensemble
sampling:
N constant,
T constant,
p constant

Pressure fluctuations in an NPT simulation of a small system consisting of 183,508 atoms vs. that of a large system consisting of 1,447,456 atoms vs. that of a huge system consisting of 23,186,176 atoms.





All-atom model system of eight GLUT1 in asymmetric environments mimicking the membrane of a human erythrocyte. The system consists of 290,776 atoms. The membrane potential of this model system is -30 mV, which will increase to $+168$ mV when a single cation is moved across the membrane along the EC-to-IC direction.

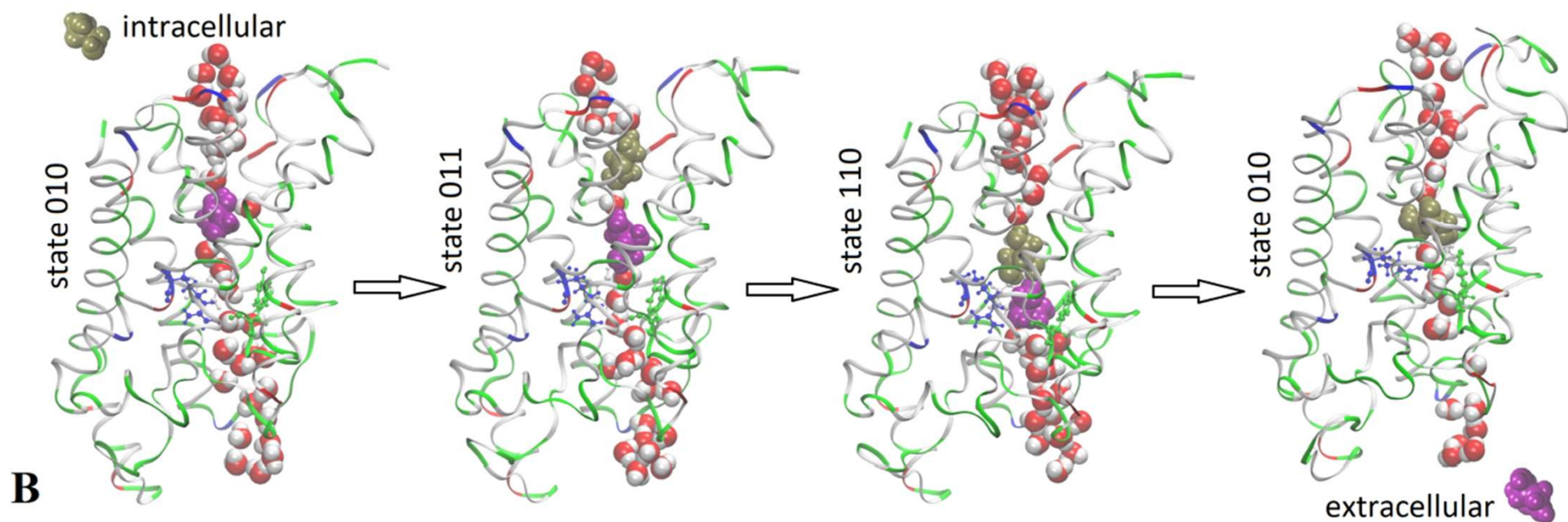


Fig. 3 A. Eight states of the glycerol-AQP3 complex and transitions between them. The three wells on a red curve indicate the binding sites of glycerol inside the AQP3 channel. Each green nut indicates the presence of a glycerol molecule at the binding site. **B.** An IC-to-EC transport event. The protein AQP3 monomer is shown in ribbons and the ar/R sf residues (Phe63, Tyr212, Arg218) in ball-and-sticks, all colored by residue types (hydrophobic, white; hydrophilic, green; negatively charged, red; positively charged, blue). Water molecules inside and near the AQP3 channel are shown in space-filling spheres colored by atoms (O, red; H, white). Two glycerol molecules are shown in space-filling spheres: GOL661 colored purple and GOL102 colored gold.

- Two fundamental problems in typical MD studies of "small system":
1. NPT ensemble but pressure is hardly constant
 2. Membrane potential fluctuates too wildly

Good news: By Going larger!

Frontera Benchmarks

