

Modern Molecular Simulations using Extended-Lagrangian Approaches

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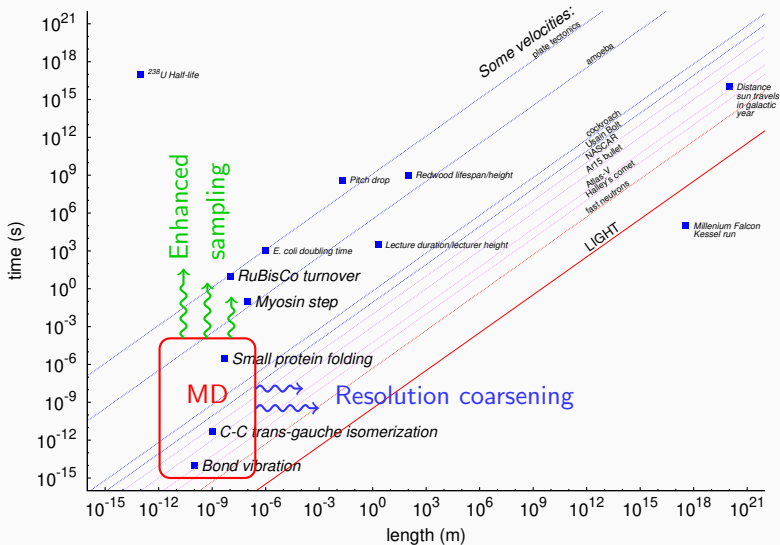
March 25, 2024

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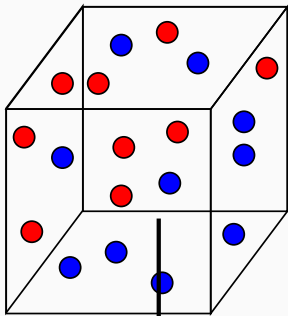
1. Introduction and Motivation
2. Summary

Introduction and Motivation

Lengthening the Reach of Molecular Dynamics Simulations



The Essential Problem: Poor Sampling of Feature Space



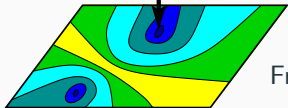
Configuration space $\mathbf{x}^{3N} \equiv (x_1, y_1, z_1, \dots)$

MD $m_i \ddot{x}_i = -\nabla_i V(\mathbf{x}) + \text{ensemble forces}$

Samples $\Rightarrow \mathbf{x}(t_1), \mathbf{x}(t_2), \dots$

Ergodicity $\langle X \rangle \approx \frac{1}{n_\tau} \sum_{i=1}^{n_\tau} X[\mathbf{x}(t_i)]$

Mapping function $\theta(\mathbf{x}) \mapsto \mathbf{z}$



Feature space $\mathbf{z}^M \equiv (z_1, z_2, \dots)$

Free energy $F(\mathbf{z}) = -k_B T \ln \langle \delta [\theta(\mathbf{x}) - \mathbf{z}] \rangle$

Summary

Summary

- OTFP combines enhanced sampling and free-energy-profile generation to provide deeper understanding of biomolecular mechanisms
- We recapitulate water-mediated “knock-on” transport of multiple K^+ ions in Kv1.2, but also predict reduced barriers for dry transport
- Demonstrated Markovian milestoning along minimum free-energy pathways to estimate entry and exit rates of O_2 in MSOX
- MSOX: Simulations agree with experiment that modified ping-pong mechanism is more likely than ping-pong because substrate influences O_2 entry and exit kinetics
- MSOX: Substrate-induced gatekeeper closing and active-site desolvation likely underly the substrates effect on O_2 .

Acknowledgments

- **OTFP:** Dr. S. Alexis PAZ (N. U. Córdoba, Argentina)
- **Milestoning:** Dr. Anthony BUCCI (West Pharma)
- **Other current projects:**
 - Steven GOSSERT (2021): HIV-1 Entry Inhibitor Design
 - Natasha VERGARA (2022): HIV-1 Entry Inhibitor Design
 - Ming HUANG (2023): Molecular modeling of thermosets
 - Dr. Salman ZARRINI: Molecular modeling of thermoset/glass sizing fluid
 - Dr. Gourav SHRIVASTAV: Milestoning and advanced methods
 - Dr. Mohammadjavad MOHAMMADI: HIV-1 Entry Inhibitor Design
- **Collaborators**
 - Prof. Eric VANDEN-EIJNDEN (NYU; Mathematics)
 - Prof. Irwin CHAIKEN (DUCOM; Biochemistry)
 - Prof. Amos SMITH III (UPenn; Organic Chemistry)
- **Funding (this talk):** NIH R01GM100472, NSF DMR1207389
- **Resources:** www.github.com/cameronabrams/cfacv,
www.github.com/cameronabrams/psfgen