

Finding optimal reaction coordinates via kinetic rate minimization



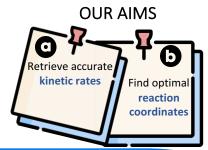
Line Mouaffac¹, Karen Palacio-Rodriguez², Fabio Pietrucci¹

¹Sorbonne Université, Musée National d'Histoire Naturelle, Institut de Minéralogie, de Physique des Matériaux et de Cosmochimie, Paris, France ²Department of Theoretical Biophysics, Max Planck Institute of Biophysics, Frankurt am Main, Germany



INTRODUCTION

While collective variables (CV) are ubiquitously used to model physico-chemical transformations, finding optimal reaction coordinates (RC) that yield accurate thermodynamic and kinetic properties is a fundamental challenge in the field of atomistic simulations. Projecting onto an accurate RC helps to accurately and efficiently estimate reaction rates and identify the common properties of transition states. Customarily, optimal RCs are defined as monotonic one-to-one functions of the committor [1]. We are proposing an original computationally affordable method to optimize reaction coordinates [2]. It is based on the fact that sub-optimal RCs yield kinetic rates higher than optimal ones.





THEORETICAL FRAMEWORK

In 2016, Zhang et. [3] investigated the repercussions on the reaction rate k of projecting the full dynamics onto a lowdimensional space defined by the CV





Effective dynamics $q(\{\overline{R_i}\})$

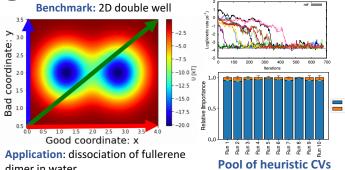
The rate of the full dynamics is less than or equal than the one computed using effective dynamics. The kinetic rate can be used as an $k^{true}_{A \to B} \le k^{model}$

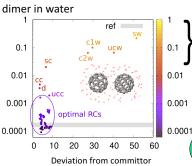
optimization criterion for CVs

The free energy F(q) and diffusion D(q) are obtained by training an overdamped Langevin model via likelihood maximization [4].

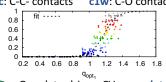
$$\dot{q} = -\beta D(q) \frac{\partial F(q)}{\partial q} + \frac{\partial D(q)}{\partial q} + \sqrt{2D(q)} \eta(t)$$
Free energy (thermodynamics) Diffusion profile (kinetics)

METHOD VALIDATION





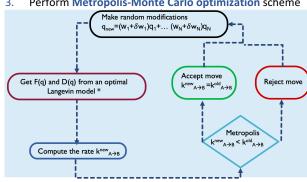
sw: entropy sc: carbon entropy ucw: VdW Energy. d: distance cc: C-C- contacts c1w: C-O contacts



Our data-driven CVs correlate well with the committor using only a fraction of the computational cost

COMPUTATIONAL METHODS

- Create a pool of collective variables
- Generate an initial random linear combination of CVs $q_{old} = w_1 q_1 + w_2 q_2 + ... w_N q_N$
- Perform Metropolis-Monte Carlo optimization scheme



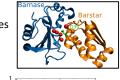
Get rate and optimal linear combination after convergence

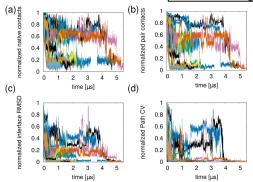
$$\frac{1}{k_{A \to B}} = \int_{q_0}^{b} dy \frac{e^{\beta F(y)}}{D(y)} \int_{a}^{y} dz e^{-\beta F(z)}$$

METHOD PERSPECTIVES

Application: predict dissociation rates of proteinprotein complexes in water

Brute force association trajectories provided by DE Shaw [5]





We developed an efficient & computationally affordable method that automates the optimization of RCs

Our method could be applied to find dissociation rates of proteins and find optimal RCs that describe the transition

REFERENCES

- [1] J. Lu and E. Vanden-Eijnden JCP, 2014.
- [2] L. Mouaffac, K. Palacio-Rodriguez, and F. Pietrucci JCTC, 2023.
- [3] W. Zhang, C. Hartmann, and C. Schütte Faraday discuss., 2016.
- [4] K. Palacio-Rodriguez and F. Pietrucci JCTC, 2022.
- [5] A. Pan, D. Jacobson, K. Yatsenko, D. Sritharan, T. Weinreich, D.Shaw *PNAS* (2019)