Quantum mechanical free energy calculations using path integral molecular dynamics

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Background

- 1. Free energy calculations
 - ▶ Free energy calculations are an integral part of chemistry
 - ▶ Provide insight into equilibrium structures, reaction rates, etc.
 - ▶ Helmholtz free energy directly related to partition function:

$$A = -k_B T \ln(Z)$$

- \blacktriangleright In principle, easy to calculate. In practice, Z is very difficult
- 2. Nuclear quantum effects
 - ▶ Important in molecular dynamics simulations¹
 - ▶ Use Feynman path integrals

What if we want to do free energy calculations on systems with non-negligible nuclear quantum effects?

¹M. Ceriotti, et al., Phys. Rev. Lett. **103**, 030603 (2009), M. Ceriotti, et al., Proc. Natl. Acad. Sci. **110**, 15591–15596 (2013).

Umbrella sampling with post-quantization restraints (PQR)

Classical Umbrella sampling¹ and WHAM²

• Umbrella sampling introduces a biasing potential to the system

$$V_{\text{total}}(q) = V(q) + \frac{1}{2}k(q - q_{eq})^2$$

- Bias is required to sample desired configurations
 - ► Water dimer would evaporate at high temperature
- Unbias simulation windows using the Weighted Histogram Analysis Method (WHAM)



¹G. M. Torrie, and J. P. Valleau, Chem. Phys. Lett. 28, 578–581 (1974), G. Torrie, and J. Valleau, J. Comp. Phys. 23, 187–199 (1977).

²S. Kumar, et al., J. Comp. Chem. 13, 1011–1021 (1992).

Umbrella sampling for path integrals

Unbiased path integral definition

$$Z = \operatorname{Tr}\left[e^{-\beta\left(\hat{K}+\hat{V}\right)}\right]$$
$$= \lim_{P \to \infty} \int \mathrm{d}q_1 \dots \mathrm{d}q_P \prod_{i=1}^P \rho_\tau(q_i, q_{i+1})$$

where
$$\rho_{\tau}(q_i, q_{i+1}) = e^{-\frac{\tau}{2}V(q_i)}\rho_0(q_i, q_{i+1})e^{-\frac{\tau}{2}V(q_{i+1})}$$

 \blacksquare Simple solution

$$Z_{\text{bias}} = \text{Tr} \left[e^{-\beta \left(\hat{K} + \hat{V} + \hat{V}_{\text{bias}} \right)} \right]$$
$$= \lim_{P \to \infty} \int dq_1 \dots dq_P \prod_{i=1}^P \rho_{\tau}^{\text{bias}}(q_i, q_{i+1})$$



ring polymer

2 atoms with P=4,R=4

where

$$\rho_{\tau}^{\text{bias}}(q_i, q_{i+1}) = e^{-\frac{\tau}{2}(V(q_i) + V_{\text{bias}}(q_i))} \rho_0(q_i, q_{i+1}) e^{-\frac{\tau}{2}(V(q_{i+1}) + V_{\text{bias}}(q_{i+1}))}$$
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Umbrella sampling for path integrals

Unbiased path integral definition

$$Z = \operatorname{Tr}\left[e^{-\beta\left(\hat{K}+\hat{V}\right)}\right]$$
$$= \lim_{P \to \infty} \int \mathrm{d}q_1 \dots \mathrm{d}q_P \prod_{i=1}^P \rho_\tau(q_i, q_{i+1})$$

where
$$\rho_{\tau}(q_i, q_{i+1}) = e^{-\frac{\tau}{2}V(q_i)}\rho_0(q_i, q_{i+1})e^{-\frac{\tau}{2}V(q_{i+1})}$$

■ Small change to simple solution

$$Z_{PQR} = \operatorname{Tr} \left[e^{-\beta(\hat{K} + \hat{V})} e^{-\beta V_{\text{bias}}} \right]$$
$$= \lim_{P \to \infty} \int dq_1 \dots dq_P e^{-\beta V_{\text{bias}}(q_1)} \prod_{i=1}^P \rho_\tau(q_i, q_{i+1})$$



ring polymer

2 atoms with P=4,R=1

 \blacktriangleright Only requires 1 $V_{\rm bias}$ term

Test Method - Ar_2 and Ne_2^1



¹K. P. Bishop, and P.-N. Roy, J. Chem. Phys. **148**, 102303 (2018).

Free energy profiles for MB-pol¹



¹K. P. Bishop, and P.-N. Roy, J. Chem. Phys. **148**, 102303 (2018).

Free energy differences for MB-pol²

• $\Delta A = -12.90 \pm 0.05 \text{ kJ/mol}$, Expt. value¹ = $-13.2 \pm 0.12 \text{ kJ/mol}$



- ¹B. E. Rocher-Casterline, et al., J. Chem. Phys. 134, 211101 (2011).
- ²K. P. Bishop, and P.-N. Roy, J. Chem. Phys. **148**, 102303 (2018).

Quantum mechanical free energy calculations using constrained path integral molecular dynamics

Formalism

 \blacksquare Free energy for a specific value of a reaction coordinate, ξ^*

$$A(\xi^*) = -k_B T \ln \rho(\xi^*) = -k_B T \ln \langle \xi^* | \operatorname{Tr}_{\mathbf{X}} e^{-\beta \hat{H}} | \xi^* \rangle$$

■ The derivative of the free energy is thus

$$A'(\xi^*) = \frac{\partial}{\partial \xi} A(\xi^*) = \frac{\partial}{\partial \xi} (-k_B T \ln \rho(\xi^*)) = -k_B T \frac{\rho'(\xi^*)}{\rho(\xi^*)}$$

where $\rho(\xi^*) = \int \mathrm{d}\mathbf{q} \,\delta(\xi(\mathbf{q}_1) - \xi^*) \,\langle \mathbf{q} | e^{-\beta \hat{H}} | \mathbf{q} \rangle$

- The final component to evaluate is $\rho'(\xi^*)$ and Dmitri Iouchtchenko has formally derived 2 estimators:
 - 1. Evaluate derivative in operator representation and then discretize
 - 2. Discretize and evaluate derivative in path integral representation

Estimator 1

The derivative of the density is given by

$$\rho'(\xi^*) \approx \int \mathrm{d}\mathbf{q}_1 \cdots \int \mathrm{d}\mathbf{q}_P \, \widetilde{\delta(\xi(\mathbf{q}_1) - \xi^*)} \pi(\mathbf{q}_1, \dots, \mathbf{q}_P) \times \left[\frac{\partial}{\partial \xi} \ln |J(\mathbf{q}_1)| - \frac{\beta}{P} \sum_{i=1}^f \frac{\partial q_i(\mathbf{q}_1)}{\partial \xi} \sum_{j=1}^P \frac{\partial V(\mathbf{q}_j)}{\partial q_i} \right]$$

where $\pi(\mathbf{q}_1, \ldots, \mathbf{q}_P) = \langle \mathbf{q}_1 | e^{-\tau \hat{H}} | \mathbf{q}_2 \rangle \langle \mathbf{q}_2 | e^{-\tau \hat{H}} | \mathbf{q}_3 \rangle \ldots \langle \mathbf{q}_P | e^{-\tau \hat{H}} | \mathbf{q}_1 \rangle$ and $J(\mathbf{q}_1)$ is the Jacobian transformation between coordinates. The derivative of the free energy may then be evaluated using

$$A'_{\text{Est1}}(\xi^*) = -k_B T \left\langle \frac{\partial}{\partial \xi} \ln |J(\mathbf{q}_1)| - \frac{\beta}{P} \sum_{i=1}^f \frac{\partial q_i(\mathbf{q}_1)}{\partial \xi} \sum_{j=1}^P \frac{\partial V(\mathbf{q}_j)}{\partial q_i} \right\rangle_{\xi^*, P}$$

,

Estimator 2

An alternative for the derivative of the density is given by

$$\rho'(\xi^*) \approx \int d\mathbf{q}_1 \cdots \int d\mathbf{q}_P \underbrace{\delta(\xi(\mathbf{q}_1) - \xi^*) \pi(\mathbf{q}_1, \dots, \mathbf{q}_P)}_{-\frac{\beta}{P} \frac{\partial V(\mathbf{q}_1)}{\partial \xi} - \sum_{i=1}^f \frac{m_i P}{\hbar^2 \beta} \frac{\partial q_i(\mathbf{q}_1)}{\partial \xi} \underbrace{(2q_i(\mathbf{q}_1) - q_i(\mathbf{q}_2) - q_i(\mathbf{q}_P))}_{(2q_i(\mathbf{q}_1) - q_i(\mathbf{q}_2) - q_i(\mathbf{q}_P))}$$

The derivative of the free energy may then be evaluated using

$$A'_{\text{Est2}}(\xi^*) = -k_B T \left\langle \frac{\partial}{\partial \xi} \ln |J(\mathbf{q}_1)| - \frac{\beta}{P} \frac{\partial V(\mathbf{q}_1)}{\partial \xi} - \sum_{i=1}^f \frac{m_i P}{\hbar^2 \beta} \frac{\partial q_i(\mathbf{q}_1)}{\partial \xi} \left(2q_i(\mathbf{q}_1) - q_i(\mathbf{q}_2) - q_i(\mathbf{q}_P) \right) \right\rangle_{\xi^*, P}$$

Implementing the estimators within PIMD

- \blacksquare The reaction coordinate for the first bead needs to be fixed at ξ^*
 - ▶ Requires a *constraint*
 - \blacktriangleright Our implementation uses the CCMA 1 constraint of OpenMM 2
- Implementation is initially benchmarked against Lennard-Jones dimers
 - ▶ Constraint between Lennard-Jones atoms
- Estimators are then used to evaluate free energy profile of water dimer and compared to PQR results
 - ▶ Constraint between oxygen atoms
 - ▶ q-SPC/Fw³, q-TIP4P/F⁴ and MB-pol⁵

¹P. Eastman, and V. S. Pande, J. Chem. Theory Comput. 6, 434-437 (2010).

²P. Eastman, et al., PLOS Comput. Biol. 13, 1–17 (2017).

³F. Paesani, et al., J. Chem. Phys. **125**, 184507 (2006).

⁴S. Habershon, et al., J. Chem. Phys. **131**, 024501 (2009).

⁵V. Babin, et al., J. Chem. Theory Comput. 9, 5395-5403 (2013), V. Babin, et al., J. Chem. Theory Comput. 10, 1599-1607 (2014).

Constrained PIMD simulations of Ar_2 at 10K



Density / arb. unit

Free energy derivatives of Ar_2 and Ne_2



Free energy profiles of Ar_2 and Ne_2



Free energy derivatives of MB-pol



Free energy profiles of MB-pol



ΔA for MB-pol

• $\Delta A = -13.03 \pm 0.14 \text{ kJ/mol}$, Expt. value¹ = $-13.2 \pm 0.12 \text{ kJ/mol}$



¹B. E. Rocher-Casterline, et al., J. Chem. Phys. 134, 211101 (2011).

Conclusions and future work

 \blacksquare PQR methodology works well down to 25 K

- ▶ Optimization of Langevin friction
- $\Delta A = -12.90 \pm 0.05 \text{ kJ/mol}$
- Constraint methodology also works well down to 25 K
 - ▶ Currently only supports atom-atom distance constraints
 - ► Estimator 3 is the most stable estimator for the water dimer over a broad temperature range
 - $\Delta A = -13.03 \pm 0.14 \text{ kJ/mol}$
- GPU implementation within OpenMM for constrained PIMD
- Systematic smoothing of free energy derivatives prior to integration
- Extensions to adaptive biasing methods such as metadynamics

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