## Nuclear Quantum Effects

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## Introduction to Imaginary Time Path Integrals Methods

Qijing Zheng

Department of Physics

University of Science and Technology of China



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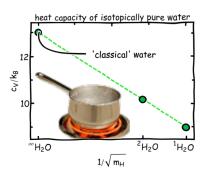
#### Outline

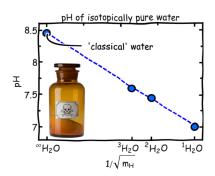
Introduction

2 Basics of Imaginary-time Path Integrals



#### Isotope Effects in Water





Proton sharing and delocalization



Strengthens hydrogen bonding, increases structure and slows the dynamics.

Hydrogen bond bending and distortion



Weakens hydrogen bonding, de-structures the liquid and speeds up dynamics

#### Experimentally Observed Isotope Effects in Water

Table 1. Experimentally Observed Isotope Effects in Water<sup>a</sup>

property	H <sub>2</sub> O	$D_2O$	$T_2O$	H <sub>2</sub> <sup>18</sup> O	ref(s)
bond energies (kJ/mol) (gas phase)	458.9	466.4 (1.6%)			49
dipole moment (D) (gas phase)	1.855	1.855			50
vibrational frequency (cm <sup>-1</sup> ) (gas phase)	3657.1	2671.6	2237.2	3649.7	51, 52
	1594.7	1178.4	995.4	1588.3	
	3755.9	2787.7	2366.6	3741.6	
water dimer dissociation energy (kJ/mol) (gas phase) 10 K	13.22	14.88 (12.7%)			53-55
melting point, T <sub>m</sub> (K) (at 1 atm)	273.15	276.97 (1.40%)	277.64 (1.64%)	273.46 (0.11%)	52, 56, 57
temperature of maximum density (TMD) (K)	277.13	284.34 (2.60%)	286.55 (3.40%)	277.36 (0.08%)	58, 59
critical temperature (K)	647.10	643.85 (-0.50%)	641.66 (-0.84%)		52, 56
molar density (mol/L)	55.35	55.14 (-0.38%)	55.08 (-0.49%)	55.42 (0.13%)	56, 60
molar density at the TMD (mol/L)	55.52	55.22 (-0.53%)	55.17 (-0.63%)	55.59 (0.13%)	56, 59
liquid/vapor surface tension (N/m)	0.07198	0.07187 (-0.15%)			56
specific heat capacity, C <sub>v</sub> [J/(K mol)]	74.54	84.42 (13.2%)			56, 61
translational diffusion (Å <sup>2</sup> /ps)	0.230	0.177 (-23.0%)		0.219 (-6.1%)	62-64
rotational diffusion (rad <sup>2</sup> /ps)	0.104	0.086 (-17%)			65
dielectric relaxation time at 20 °C (ps)	9.55	12.3 (29%)			66
viscosity (mPa s)	0.8904	1.0966 (23.2%)		0.9381 (5.4%)	56, 59
acidity pH/pD	7.00	7.43	7.6		56, 57

<sup>&</sup>quot;Values given are at 25 °C unless otherwise stated, and values in parentheses are percentage shifts relative to H2O.

M. Ceriotti, et al. Chem. Rev., 116, 7529 (2016)

#### Nuclear Quantum Effects

NQEs include tunneling and zero-point motion.



other nuclear quantum effects: exchange effects, quantum coherence...

• The thermal de Broglie wavelength of a particle:

$$\Lambda = \frac{h}{\sqrt{2\pi m k_B T}}$$

for a hydrogen atom at 300 K,  $\Lambda \approx 1.0 \,\text{Å}$ .

• NQEs are important for any vibrational mode for which  $\hbar\omega/k_BT>1$ . If  $T=300\,{\rm K},$   $\omega\approx208\,{\rm cm}^{-1}.$ 

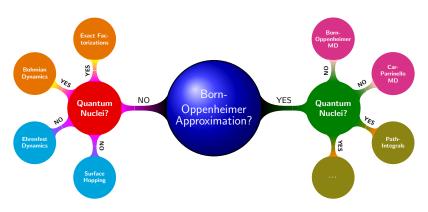
$$K_Q = \frac{\hbar\omega}{4} \coth \frac{\hbar\omega}{2k_BT}$$
 
$$\frac{\frac{\hbar\omega}{k_BT} \ll 1}{2}$$
  $K_C = \frac{k_BT}{2}$ 

### How to model the dynamics of electrons and nuclei from ab initio?

The time-dependent Schrödinger equation:

$$i\hbar \frac{\partial \Psi(\mathbf{r}, \mathbf{R}, t)}{\partial t} = \hat{\mathcal{H}}(\mathbf{r}, \mathbf{R}) \Psi(\mathbf{r}, \mathbf{R}, t)$$

In practice, approximations have to be made! 1



<sup>&</sup>lt;sup>1</sup> "Ab initio molecular dynamics", Mariana Rossi, DFT Workshop 2017

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#### Outline

Introduction

2 Basics of Imaginary-time Path Integrals

### Path Integral Isomorphism I

For a single particle moving in one spatial dimension potential  $^{2}$ 

$$\hat{\mathcal{H}} = \frac{\hat{p}^2}{2m} + \hat{V}(\hat{x}) \equiv \hat{T} + \hat{V}$$

The quantum partition function

$$\begin{split} Z &= \text{tr} \left[ e^{-\beta \hat{\mathcal{H}}} \right] = \text{tr} \left[ (e^{-\frac{\beta}{P} \hat{\mathcal{H}}})^P \right] = \text{tr} \left[ (e^{-\beta_P \hat{\mathcal{H}}})^P \right] \\ &= \int \mathrm{d}x_1 \, \left\langle x_1 | (e^{-\beta_P \hat{\mathcal{H}}})^P | x_1 \right\rangle \\ &= \int \mathrm{d}x_1 \dots \int \mathrm{d}x_P \, \left\langle x_1 | e^{-\beta_P \hat{\mathcal{H}}} | x_2 \right\rangle \dots \left\langle x_P | e^{-\beta_P \hat{\mathcal{H}}} | x_1 \right\rangle \end{split}$$

Note the connection between the quantum propagator and the density matrix,

$$\langle x'|e^{-i\hat{\mathcal{H}}t/\hbar}|x\rangle$$
  $t \to \tau = i\beta\hbar$   $\langle x'|e^{-\beta\hat{\mathcal{H}}}|x\rangle$ 

Hence the name imaginary-time path integral. By using the Trotter splitting<sup>3</sup>

$$\begin{split} e^{-\beta_P \hat{\mathcal{H}}} &= e^{-\beta_P \hat{V}/2} e^{-\beta_P \hat{T}} e^{-\beta_P \hat{V}/2} + \mathcal{O}(\beta_P^3) \\ \Rightarrow & \langle x_i | e^{-\beta_P \hat{\mathcal{H}}} | x_j \rangle \approx \left( \frac{m}{2\pi\beta_P \hbar^2} \right)^{1/2} e^{-\beta_P \left[ \frac{m}{2(\beta_P \hbar)^2} (x_i - x_j)^2 + \frac{1}{2} \left( V(x_i) + V(x_j) \right) \right]} \end{split}$$

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#### Path Integral Isomorphism II

$$\begin{split} Z &= \operatorname{tr} \left[ e^{-\beta \hat{\mathcal{H}}} \right] \\ &= \lim_{P \to \infty} \int \mathrm{d} x_1 \dots \int \mathrm{d} x_P \, \left( \frac{m}{2\pi \beta_P \hbar^2} \right)^{P/2} \exp \left\{ -\beta_P \underbrace{\sum_{i=1}^P \left[ \frac{m}{2(\beta_P \hbar)^2} (x_{i+1} - x_i)^2 + V(x_i) \right]}_{V_{\text{eff}}(x_1, \dots, x_P)} \right\} \bigg|_{x_{P+1} = x_1} \end{split}$$



- Maps the quantum partition function to the configuration integral of classical ring-polymers.
- **EXACT** when  $P \to \infty$ . Reduce to classical partition function when P = 1.
- The integral can be sampled with Monte Carlo method (PIMC, not covered in this talk), but for MD we need momenta!

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<sup>&</sup>lt;sup>2</sup> "Statistical Mechanics: Theory and Molecular Simulation", Mark E. Tuckerman

<sup>&</sup>lt;sup>3</sup>There are higher-order splitting techniques.

### Expectation Values from Path Integral

For a Hermitian operator  $\hat{A}$ , the expectation value follows,

$$\langle \hat{A} \rangle = \frac{1}{Z} \operatorname{tr} \left[ \hat{A} e^{-\beta \hat{\mathcal{H}}} \right] = \frac{1}{Z} \operatorname{tr} \left[ \hat{A} (e^{-\beta_P \hat{\mathcal{H}}})^P \right]$$

if  $\hat{A}$  is purely a function of position operator  $\hat{x}$ , i.e.  $\hat{A}(\hat{x})|x\rangle = a(x)|x\rangle$ , then <sup>4</sup>

$$\begin{split} \langle \hat{A} \rangle &= \frac{1}{Z} \lim_{P \to \infty} \int \mathrm{d}x_1 \dots \int \mathrm{d}x_P \, \left( \frac{m}{2\pi \beta_P \hbar^2} \right)^{P/2} \\ &\qquad \left( \frac{1}{P} \sum_{i=1}^P \mathsf{a}(x_i) \right) \exp \left\{ -\beta_P \sum_{i=1}^P \left[ \frac{m}{2(\beta_P \hbar)^2} (x_{i+1} - x_i)^2 + V(x_i) \right] \right\} \bigg|_{x_{P+1} = x} \end{split}$$

- $a(x_1, ..., x_P) = \frac{1}{P} \sum_{i=1}^{P} a(x_i)$  is referred to as the *estimator* of  $\langle \hat{A} \rangle$ .
- If  $\hat{A}$  is a function of momentum operator  $\hat{p}$ , then the cyclic-path condition  $x_{P+1} = x_1$  is released, in which case one should resort to the so-called open-path path integral method.
- The thermodynamic functions, which may depend on both  $\hat{x}$  and  $\hat{p}$ , are exceptional because thermodynamic relations can be used. For example

$$E = \langle \hat{\mathcal{H}} \rangle = \langle \frac{\hat{p}^2}{2m} + V(\hat{x}) \rangle = -\frac{\partial}{\partial \beta} \ln Z = \frac{1}{Z} \frac{\partial Z}{\partial \beta}$$

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<sup>&</sup>lt;sup>4</sup> "Statistical Mechanics: Theory and Molecular Simulation", Mark E. Tuckerman 🕨 🗅 🕨 🗸 📑 🕨 🔻 🚆 🕨

## Energy and Kinetic Energy Estimator

• The *primitive* energy estimator  $\varepsilon_{prim}(x_1, \ldots, x_P)$ 

$$\varepsilon_{\mathsf{prim}}(x_1, \dots, x_P) = \frac{P}{2\beta} - \sum_{i=1}^P \frac{mP}{2\beta^2 h^2} (x_{i+1} - x_i)^2 + \frac{1}{P} \sum_{i=1}^P V(x_i)$$
$$= \mathcal{K}_{\mathsf{prim}} + \frac{1}{P} \sum_{i=1}^P V(x_i)$$

- K<sub>prim</sub> is the estimator for the quantum kinetic energy, which scale linearly with the number of beads P.
- The virial theorem:

$$\frac{P}{2\beta} - \left\langle \sum_{i=1}^{P} \frac{mP}{2\beta^2 \hbar^2} (x_{i+1} - x_i)^2 \right\rangle = \left\langle \frac{1}{2P} \sum_{i=1}^{P} x_i \frac{\partial V}{\partial x_i} \right\rangle$$

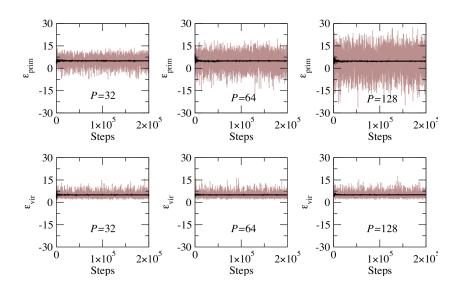
• The *virial* energy estimator  $\varepsilon_{\text{vir}}(x_1,\ldots,x_P)$  and virial kinetic energy estimator  $\mathcal{K}_{\text{vir}}$  5

$$\varepsilon_{\text{vir}}(x_1, \dots, x_P) = \frac{1}{P} \sum_{i=1}^P \frac{1}{2} x_i \frac{\partial V}{\partial x_i} + \frac{1}{P} \sum_{i=1}^P V(x_i)$$
$$= \frac{1}{2\beta} + \frac{1}{P} \sum_{i=1}^P \frac{1}{2} (x_i - x_c) \frac{\partial V}{\partial x_i} + \frac{1}{P} \sum_{i=1}^P V(x_i)$$

<sup>5</sup> "Statistical Mechanics: Theory and Molecular Simulation", Mark E. Tuckerman « ロ » « 🗗 » « 🛢 » « 🛢 » 🥞 🥏 🥏 🤄

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#### Comparison of Primitive and Virial Energy Estimator



<sup>5 &</sup>quot;Statistical Mechanics: Theory and Molecular Simulation", Mark E. Tuckerman « □ » « 🗗 » « 🛢 » « 🛢 » 🗦 🥏 🔗 ९ 🧇

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#### PIMD — Introducing Momenta

Insert 
$$P$$
 Gaussian integral into  $Z$ :  $\left(\frac{\beta}{2\pi\tilde{m}}\right)^{1/2}\int\mathrm{d}p\,e^{-\beta\frac{p^2}{2\tilde{m}}}=1$ 

$$Z = \left(\frac{1}{2\pi\hbar}\right)^P \prod_{i=1}^P \sqrt{\frac{m}{\tilde{m}_i}} \lim_{P \to \infty} \int \mathrm{d}x_1 \dots \int \mathrm{d}x_P \int \mathrm{d}p_1 \dots \int \mathrm{d}p_P$$

$$= \exp\left\{-\beta_P \sum_{i=1}^P \left[\frac{p_i^2}{2\tilde{m}_i} + \frac{m}{2(\beta_P \hbar)^2} (x_{i+1} - x_i)^2 + V(x_i)\right]\right\}$$

$$= \mathcal{C}\lim_{P \to \infty} \int \mathrm{d}\mathbf{x}^P \int \mathrm{d}\mathbf{p}^P \exp\left[-\beta_P H_P(\mathbf{x}, \mathbf{p})\right]$$

$$= \lim_{P \to \infty} \int \mathrm{d}\mathbf{x}^P \int \mathrm{d}\mathbf{p}^P \exp\left[-\beta_P H_P(\mathbf{x}, \mathbf{p})\right]$$

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• PIMD trajectories are obtained by integrating Hamilton's classical equaiton of motion

$$\frac{\mathrm{d}\boldsymbol{p}_i}{\mathrm{d}\boldsymbol{t}} = + \frac{\partial H_P(\mathbf{x}, \mathbf{p})}{\partial x_i}; \qquad \frac{\mathrm{d}x_i}{\mathrm{d}\boldsymbol{t}} = - \frac{\partial H_P(\mathbf{x}, \mathbf{p})}{\partial \boldsymbol{p}_i}$$

- $\beta_P H_P(\mathbf{x}, \mathbf{p}) \Rightarrow \beta (H_P(\mathbf{x}, \mathbf{p})/P)$ , which affects the dynamics but not the statistics.
- Different mass matrices give the same static average, only dynamics will be changed.
- Mathematically, the main difference between PIMD, PACMD and RPMD is the choice of the mass matrix. Physically, they differ dramatically!

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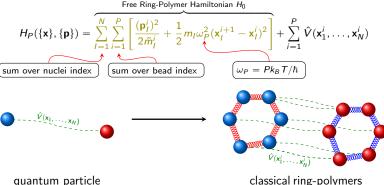
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### Multiparticle Generalization

For a system of N distinguishable nuclei, 6

$$\hat{\mathcal{H}}_{P} = \sum_{I}^{N} \frac{\mathbf{p}_{I}^{2}}{2m_{I}} + \frac{\hat{V}(\mathbf{x}_{1}, \dots, \mathbf{x}_{N})}{\text{Potential energy: could be from emperical potential or from abinitio calculations. } \frac{\mathbf{p}_{I}}{\mathbf{p}_{I}}$$

The PIMD Hamiltonian is



<sup>6</sup>We are neglecting the exchange effect, which is generally fine for nuclei (unless treating 4 K Helium) but not for electrons.

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4 D > 4 A > 4 B > 4 B >

#### Terminologies and Properties of the Ring-Polymer

• Radius of Gyration – the spread in imaginary time. For a free particle the root mean square radius of gyration is:

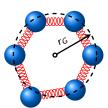
$$\left\langle r_G^2(T) \right\rangle^{1/2} = \frac{\Lambda(T)}{\sqrt{8\pi}} \qquad \qquad \Lambda(T) = \frac{h}{\sqrt{2\pi m k_B T}}$$

$$\Lambda(T) = \frac{h}{\sqrt{2\pi m k_B T}}$$

 Bead to bead distance – For a free particle the average is:

$$\sqrt{\frac{\beta\hbar^2}{Pm}}$$

Note the distance decreases as P increases.



 Centroid – the center of the Ring-Polymer.

$$x_c = \frac{1}{P} \sum_{i=1}^P x_i$$

Bead spring constants <sup>7</sup> – determined by mω<sup>2</sup><sub>P</sub>

$$\omega_P = \frac{1}{\beta_P \hbar} = P k_B T / \hbar$$

The overall object is referred to as an Imaginary Time Path or a Ring-Polymer.

<sup>7</sup>In some textbooks,  $\omega_P = \sqrt{P} k_B T / \hbar$  if the Hamiltonian is  $H_P(\mathbf{x}, \mathbf{p}) / P$ 

### Ring-Polymer Normal Modes

• Normal Modes: – the ring-polymer potential can be diagonalized

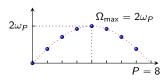
$$\sum_{i=1}^P \frac{1}{2} m \omega_P^2 (x_{i+1} - x_i)^2 \quad \Longleftrightarrow \quad \begin{aligned} \tilde{\mathbf{x}} &= \mathbf{U} \cdot \mathbf{x} \\ \mathbf{x} &= \mathbf{U}^T \cdot \tilde{\mathbf{x}} \end{aligned} \qquad \Longleftrightarrow \quad \sum_{j=1}^P \frac{1}{2} m \Omega_j^2 \tilde{x}_j^2;$$

where x is the bead Cartesian coordinate and  $\tilde{x}$  is the normal mode coordinate.

- $\tilde{x}_1$  corresponds to the centroid motion and  $\Omega_1=0.$
- The transformation  $\tilde{\mathbf{x}} \Leftrightarrow \mathbf{x}$  can be done with FFT.<sup>8</sup>
- The normal mode frequencies <sup>9</sup>

$$\Omega_j = 2\omega_P \sin\left(\frac{(j-1)\pi}{P}\right)$$

e.g. 
$$P = 8$$



• The matrix element  $U_{kj}$  of the unitary transformation matrix **U** for even P

$$\begin{cases} \sqrt{1/P}, & k = 1 \\ \sqrt{2/P}\cos\left(\frac{2\pi}{P}(j-1)(k-1)\right), & 2 \leqslant k \leqslant P/2 \\ \sqrt{1/P}(-1)^j, & k = P/2 + 1 \\ \sqrt{2/P}\sin\left(\frac{2\pi}{P}(j-1)(k-1)\right), & P/2 + 2 \leqslant k \leqslant P \end{cases}$$

and 
$$\tilde{x}_k = \sum_{j=1}^P U_{kj} x_j$$
.

• Note if we choose  $\beta_P H_P \Rightarrow \beta(H_P/P)$ , then

$$\tilde{\mathbf{x}} = \frac{1}{\sqrt{P}} \mathbf{U} \cdot \mathbf{x}$$
$$\mathbf{x} = \sqrt{P} \mathbf{U}^T \cdot \tilde{\mathbf{x}}$$



<sup>&</sup>lt;sup>8</sup> J. Chem. Phys., 104, 2028(1996).

<sup>&</sup>lt;sup>9</sup>This reminds me of the phonon dispersion of 1-D atomic chain Q.J. Zheng (D.P. USTC)

### Convergence of Standard Path Integral MD

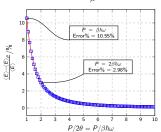
For the 1D harmonic oscillator:  $V(x) = \frac{1}{2}m\omega x^2$ , the RP potential in the normal-mode coordinates

$$\sum_{i=1}^{P} \left[ \frac{m}{2(\beta_{P}\hbar)^{2}} (x_{i+1} - x_{i})^{2} + V(x_{i}) \right] \quad \Rightarrow \quad \sum_{i=1}^{P} \frac{1}{2} m(\Omega_{i}^{2} + \omega^{2}) \tilde{x}_{i}^{2}$$

then  $\langle \tilde{x}_i^2 \rangle = 1/\beta_P m(\Omega_i^2 + \omega^2)$ , the energy of the harmonic oscillator  $\langle E \rangle_P$ 

$$\begin{split} \langle E \rangle_P &= \langle T \rangle_P + \langle V \rangle_P = 2 \langle V \rangle_P = 2 \cdot \frac{1}{2} m \omega^2 \langle x^2 \rangle = \frac{m \omega^2}{P} \sum_{i=1}^P \langle x_i^2 \rangle = \frac{m \omega^2}{P} \sum_{i=1}^P \langle \tilde{x}_i^2 \rangle \\ &= \frac{\theta}{\beta} \sum_{i=1}^P \frac{\theta}{P^2 \sin^2 \left( \frac{(i-1)\pi}{P} \right) + \theta^2} \qquad (\theta = \beta \hbar \omega/2) \end{split}$$

the exact solution is  $\langle E \rangle = \frac{\theta}{\beta} \coth(\theta)$ 



- Commonly used convergence criteria:  $P > \hbar \omega_{\rm max}/k_B T$
- Standard PIMD converges as  $1/P^2$ .
- Some properties converges faster, e.g. RDF converges fast but heat capacity very slowly.

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## Reducing the Number of Beads

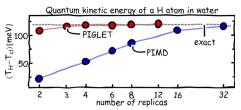
• Higher-order Trotter's expansion and better estimator.

Jang et al. J. Chem. Phys., 115, 7832 (2001)

Ring-polymer Contraction.

Markland et al. J. Chem. Phys., 464, 256 (2008)

 Smart thermostating of internal modes, e.g., generalized Langevin equation based colored noise thermostats.



Ceriotti et al. J. Chem. Phys., 133, 124104 (2010)

### Frequencies in a PIMD Simulation

For the 1D harmonic oscillator:  $V(x) = \frac{1}{2}m\omega x^2$ , the normal mode frequencies

$$ilde{\Omega}_i = \sqrt{\Omega_i^2 + \omega^2}; \qquad \Omega_i = 2\omega_P \sin\left(rac{(i-1)\pi}{P}
ight)$$

Hence the highest frequency is

$$\tilde{\Omega}_{\mathsf{max}} = \sqrt{\Omega_{\mathsf{max}}^2 + \omega^2} = \sqrt{4\omega_P^2 + \omega^2}$$

If we use the typical convergence criteria:  $P=\alpha\hbar\omega/k_BT$ , where  $\alpha>1$  determines how accurate the calculation is

$$\tilde{\Omega}_{\text{max}} = (4\alpha^2 + 1)^{\frac{1}{2}}\omega$$

- For  $\alpha=1$ , the highest frequency in PIMD is  $\sqrt{5}$  times larger than classical MD.
- For a *naive* implementation of PIMD, the time step should be  $\sqrt{5}$  times smaller than classical MD.
- Methods to allow bigger time-steps:
  - Scale the normal mode masses so they all oscillate at the same frequency.
     Multiple time-scale molecular dynamics:
  - use smaller time-steps for bead forces.
  - Use an integrator where the free ring polymer is evolved exactly.

#### PIMD, CMD and RPMD 10

#### **CMD**

- Approximate quantum dynamics
- Path centroid idea

$$m\ddot{x}_c = \left\langle \delta(x_0 - x_c) \frac{-1}{P} \sum_{i=1}^{P} \frac{\partial V(x_i)}{\partial x_i} \right\rangle$$

 Sample the whole configuration space available to the non-centroid modes at each given position of centroid  $x_0$ .

- PIMD with adiabatic decoupling of centroid and non-centroid motions.
- No thermostat for the centroid mode.
- Thermostatting for noncentroid mode as in PIMD.
- Must sample initial conditions.

#### **PIMD**

- Static equilibrium properties.
- Need to efficiently and ergodically sample the phase surface.
  - Thermostatting
  - Coordinate transformation and bead masses scaling

**NQEs** 

#### **RPMD**

- Approximate quantum dynamics
- PIMD with physical bead masses.
- Thermostats should not be used.
- Must sample initial conditions.



<sup>&</sup>lt;sup>10</sup> J. Chem. Phys., **130**, 194510(2009);

J. Chem. Phys., 129, 074501(2008); Q.J. Zheng (D.P. USTC)

## PIMD, (PA)CMD and RPMD — Fictitious Masses

The RP Hamiltonian in the normal-mode coordinates

$$\begin{split} H_{P}(\{\tilde{\mathbf{x}}\}, \{\tilde{\mathbf{p}}\}) &= \sum_{i=1}^{P} \left[ \frac{\tilde{p}_{i}^{2}}{2\tilde{m}_{i}} + \frac{1}{2} m \Omega_{i}^{2} \tilde{x}_{i}^{2} \right] + \sum_{i=1}^{P} V\left(x_{i}(\tilde{x}_{1}, \dots, \tilde{x}_{P})\right) \\ &\Rightarrow \quad \tilde{m}_{i} \ddot{\tilde{x}}_{i} = -m \Omega_{i}^{2} \tilde{x}_{i} - \sum_{j=1}^{P} \frac{\partial V\left(x_{j}(\tilde{x}_{1}, \dots, \tilde{x}_{P})\right)}{\partial \tilde{x}_{i}} \\ &\Rightarrow \quad \tilde{m}_{1} \ddot{\tilde{x}}_{1} = -\frac{1}{\sqrt{P}} \sum_{j=1}^{P} \frac{\partial V(x_{j})}{\partial x_{j}} \end{split}$$

• PIMD — scale the mass to bring all the frequencies to the same value  $\omega_p$ .

$$\tilde{m}_1 = m, \qquad \tilde{m}_i = 4 \sin^2 \left( \frac{(i-1)\pi}{P} \right) m \quad (2 \leqslant i \leqslant P)$$

ullet (PA)CMD — adiabatic parameter  $\gamma>1$ , shifting the noncentroid frequencies  $\gamma\omega_P$  off the physical spectra  $\omega_{\text{max}}$ .

$$\tilde{m}_1 = m, \qquad \tilde{m}_i = 4 \sin^2 \left( \frac{(i-1)\pi}{P} \right) m/\gamma^2 \quad (2 \leqslant i \leqslant P)$$

RPMD — Use real masses for all the beads.

$$\tilde{m}_i = m, \quad (1 \leqslant i \leqslant P)$$

《中》《圖》《意》《意》。 意 Q.J. Zheng (D.P. USTC)

#### Initializing PIMD — Positions and Momenta

$$H_{P}(\{\mathbf{x}\}, \{\mathbf{p}\}) = \sum_{l=1}^{N} \sum_{i=1}^{P} \left[ \frac{(\mathbf{p}_{l}^{i})^{2}}{2\tilde{m}_{l}^{i}} + \frac{1}{2} m_{l} \omega_{P}^{2} (\mathbf{x}_{l}^{i+1} - \mathbf{x}_{l}^{i})^{2} \right] + \sum_{i=1}^{P} \hat{V}(\mathbf{x}_{1}^{i}, \dots, \mathbf{x}_{N}^{i})$$

• To initialize PIMD requires specification of 3NP positions and 3NP momenta.

#### Momenta

• We inserted *P* Gaussians when introducing momenta

$$\left(\frac{\beta}{2\pi\tilde{m}}\right)^{1/2}\int\mathrm{d}\boldsymbol{p}\,\mathrm{e}^{-\beta\boldsymbol{p}^2/2\tilde{m}}$$

 We can then sample the momentum of each bead from a Gaussian distribution with

$$\begin{split} &\bar{p}=0\\ &\sigma_{p}=\sqrt{\frac{\tilde{m}}{\beta_{P}}}=\sqrt{\tilde{m}k_{B}PT} \end{split}$$

#### Positions

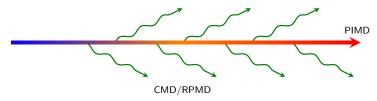
- Start all beads at the same positions and equilibrate.
  - RP will expand under NVE, T will drop.
  - Strong Thermostatting is need!
- Sample from the free ring-polymer distribution.
  - The only potential is the harmonic springs.
  - In normal mode coordinates

$$\sum_{j=1}^{P} \frac{1}{2} m \Omega_{j}^{2} \tilde{x}_{j}^{2} \quad \Rightarrow \quad \int \mathrm{d}\tilde{x}_{j} \, \mathrm{e}^{-\beta_{P} \left(\frac{1}{2} m \Omega_{j}^{2} \tilde{x}_{j}^{2}\right)}$$

- $\tilde{\tilde{x}}_j = 0$  and  $\sigma_{\tilde{x}_j} = \sqrt{\frac{1}{\beta_P m \Omega_i^2}}$ .
- In practice, RP slightly too extended.

### Initializing CMD/RPMD

- CMD and RPMD are intended to provide approximations to quantum dynamics and not to efficiently sample the quantum phase space.
- The ergodicity problem can be circumvented by launching trajectories from many different choice of configurations and momenta.



- Question Run a PIMD trajectory using an efficient thermostat scheme.
- Pick configurations and momenta from the thermostatted trajectory and launch RPMD or CMD trajectories from them.
- Ideally the separation in time between each should be determined by the correlation time of the properties in the system.

# Thank you!