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Rigid quantum Monte Carlo simulations of condensed molecular matter: Water clusters in the $n=2\rightarrow 8$ range

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The numerical advantage of quantum Monte Carlo simulations of rigid bodies relative to the flexible simulations is investigated for some simple systems. The results show that if high frequency modes in molecular condensed matter are predominantly in the ground state, the convergence of path integral simulations becomes nonuniform. Rigid body quantum parallel tempering simulations are necessary to accurately capture thermodynamic phenomena in the temperature range where the dynamics are influenced by intermolecular degrees of freedom; the stereographic projection path integral adapted for quantum simulations of asymmetric tops is a significantly more efficient strategy compared with Cartesian coordinate simulations for molecular condensed matter under these conditions. The reweighted random series approach for stereographic path integral Monte Carlo is refined and implemented for the quantum simulation of water clusters treated as an assembly of rigid asymmetric tops. © 2007 American Institute of Physics.

I. INTRODUCTION

A number of recent investigations have contributed to the development, testing, and applications of path integral Monte Carlo methods\(^4\)–\(^{25}\) in non-Euclidean manifolds\(^{26,28}\) arising in molecular condensed matter theory, whenever holonomic constraints are employed. Among the earliest attempts reported in the literature, one finds the simulation of bulk rigid water carried out in quaternion space with the fixed axis approximation.\(^{29}\) When there is a small number of linear rotors present in the system simulated, it is possible to split the kinetic energy operator in the density matrix expression. The angular momentum terms are isolated and then treated by vector space methods.\(^{30,34}\) Such an approach has been used to handle the simulation of a linear covalent molecule on surfaces,\(^{30,31}\) a linear molecule solvated by noble gas atoms,\(^{32,34}\) and para-D\(_2\), ortho-D\(_2\) clusters.\(^{35}\) Other authors have reported path integral Monte Carlo (PIMC) methods in torsional space using time slicing and angular variables.\(^{36,37}\) In a series of articles\(^{38,42}\) we have introduced and refined methodologies to carry out PIMC methods in rotational and torsional manifolds based on the DeWitt formula.\(^2\) We have established that torsional spaces and rotational spaces can be mapped using a general strategy; the maps are bijections from the manifold to an equidimensional Euclidean space. The stereographic projection coordinates extend from negative to positive infinity; therefore, once the metric of the manifold is transformed, it becomes possible to simulate the random walk using these “Cartesian-type” coordinates and to map the diffusion process back into the manifold. The one-to-one nature of the map between the angular variables and the stereographic projections is the key for the faithful representation of random paths in the manifold. In this manner, the notoriously difficult problem of evaluating the path integral inside open sets and the problem of imposing boundary conditions on the time evolution operator are avoided. Furthermore, using stereographic projections, it is possible to expand paths using a random series after generalizing the Feynman-Kaç formula in non-Euclidean spaces. This latter advance allows one to develop a fast converging algorithm for the PIMC method, without the need to evaluate the gradient or the Hessian of the potential. Additionally, one can develop efficient estimators based on numerical derivatives of the action; this development has been recently reported for path integral simulations with Cartesian coordinates by one of our groups.\(^{22,23}\) The extension of these techniques to non-Euclidean manifolds mapped with stereographic projections has been tested recently.\(^42\) Rigid hydrogen fluoride (HF) clusters from the dimer to the octamer have been simulated.\(^45\) Rigid diatomic molecules are linear rotors; however, these calculations would be intractable if the angular terms were treated with vector spaces.

The present article has two main purposes. Firstly, we investigate the efficiency gained in PIMC simulations of molecular condensed matter when holonomic constraints are used. Secondly, we make use of new improvements in the stereographic projection path integral algorithms applied to Cartesian products of inertia ellipsoid spaces to simulate wa-
ter clusters of several sizes in the 100–500 K range. For
the investigation of the efficiency gain, in particular, we are
interested to learn how rigid body simulations are more effi-
cient than flexible ones in quantum simulations. It is well
known that rigid body simulations improve the efficiency of
thermodynamic estimators in Metropolis simulations. How-
ever, the efficiency gained by treating high frequency modes
rigidly has not been investigated carefully for quantum par-
allel tempering simulations.\textsuperscript{43–46} The gain in efficiency in
classical Metropolis simulations of rigid bodies is usually
quantifiable as a decrease in the statistical fluctuation at all
the temperatures.\textsuperscript{17–49} We expect the same to hold for parallel
tempering simulations. However, we anticipate that in quan-
tum simulations the efficiency of the rigid body algorithms
will have two contributions; the first should be a similar
decrease in statistical fluctuations of the estimated thermody-
namics properties, and the second, and probably more impor-
tant contribution is the convergence with respect to the num-
ber of path variables of the properties, at temperatures where
the high frequency modes are predominantly in the ground
state. In this case, one expects the path integral Monte Carlo
approach based on Cartesian coordinates to experience non-
uniform convergence. Nonuniform convergence can be expe-
rienced even when highly efficient accelerating schemes,
such as the reweighted random series (RRS) method, are
employed in spaces mapped by Cartesian coordinates. In or-
der to verify these statements, we carry out a number of nu-
merical tests. We choose a coupled, harmonic, many-
particle one-dimensional system (a harmonic chain) to inves-
tigate the convergence issue with path integral simulations.
The knowledge of the eigenvalues of the Hessian for the full
system and its adiabatic limit allow us to use analytical
formulas for the heat capacity as a function of $T$ and the
Trotter number. Both primitive convergence and cubic con-
vergence can be inspected analytically for the full system
and its adiabatic limit. One can learn a great deal about the
convergence behavior of the four estimators by inspecting
graphs of the heat capacity as a function of the two variables.
The harmonic chain is a reasonable yet simple model for
condensed molecular matter, where the complications that
require differential manifolds are avoided by restricting
the motion to translations in one dimension only.

We carry out several rigid body quantum simulations for
water clusters in the dimer through the octamer range.\textsuperscript{50} For
this purpose, we revisit the RRS method recently developed
for non-Euclidean manifolds, and we find a simple but im-
portant improvement for its numerical implementation. We
simulate a particle in a ring and a rigid water monomer, with
its rotations hindered by an external potential field to test the
new improvements.

The remainder of this article is organized as follows: The
improved RRS stereographic projection path integral for
simulations in non-Euclidean spaces is discussed in Sec. II.
All the numerical tests are presented in Sec. III. Section IV
contains our conclusions.

II. METHODS

The partial averaging method and RRS approach have
been extended to non-Euclidean spaces in a series of previ-
ous works.\textsuperscript{38–42} We revisit the basic theory in the present
manuscript since a computationally more efficient approach
is developed. We begin with the elements of the density ma-
trix in non-Euclidean manifolds,

$$\rho(q,q',\beta) = \int Dq \exp \left\{ -\frac{1}{\hbar} \int_0^{\beta \hbar} d\tau \left[ \frac{1}{2} g_{\mu\nu} q^\mu q^\nu + V \right] \right\},$$

(1)

where $A_s$ contains the natural logarithm of the square root of
the determinant of the metric tensor evaluated at every point
along the path. To develop the equivalent of the Feynman-
Kaç formula in non-Euclidean spaces, we introduce a conve-
nient reference space,

$$\frac{\rho(q,q',\beta)}{\rho_0(q,q',\beta)} = \frac{\int Dq \exp\left\{ -(1/\hbar) \int_0^{\beta \hbar} d\tau [\mathcal{L}^{(0)} + \Delta \mathcal{L} + V] \right\}}{\int Dq \exp\left\{ -(1/\hbar) \int_0^{\beta \hbar} d\tau \mathcal{L}^{(0)} \right\}}.$$  

(2)

The symbol $\mathcal{L}^{(0)}$ represents a potential-free Lagrangian for a
system of particles constructed in a convenient reference
space of equal dimensionality with a constant (i.e., configu-
ration independent) metric tensor,

$$\mathcal{L}^{(0)} = \frac{1}{2} g^{(0)}_{\mu\nu} q^\mu q^\nu.$$  

(3)

The Lagrangian $\mathcal{L}^{(0)}$ is used to handle the tail portion of the
path truncated by the primitive algorithm to improve con-
vergence. The geometric interaction is defined as

$$\Delta \mathcal{L} = -\frac{N}{2\beta} \ln(\det(g_{\mu\nu})) + \frac{1}{2} \left( g_{\mu\nu} - g^{(0)}_{\mu\nu} \right) q^\mu q^\nu.$$  

(4)

On the right, it is assumed that a trapezoid quadrature with $N$
points is used to evaluate the integral in $u = \tau/\beta \hbar$ so that

$$-\frac{1}{\hbar} \int_0^{\beta \hbar} d\tau \left[ \frac{N}{2\beta} \ln(\det(g_{\mu\nu})) \right]$$

$$= \frac{N}{2} \int_0^1 du (\ln(\det(g_{\mu\nu}))) = \frac{1}{2} \sum_{i=1}^{N} \ln(\det(g_{\mu\nu})) = A_s.$$  

(5)

To develop a random series algorithm, the infinite di-

dimensional Riemann integral is recast in terms of path vari-

ables $a_k^\mu$ defined by

$$q^\mu(u) = q^\mu_{\mu} + \hbar \beta^{1/2} \sum_{k=1}^{\infty} a_k^\mu \Lambda_k(u),$$  

(6)

where the functions $\Lambda_k(u)$ have the following property in the

\[ <.> \text{ denotes an ensemble average.} \]
and constructs the functions $\tilde{\Lambda}_k(u)$ so that the partial averaging expansion about the core path derived from the above is equal to the same derived with the infinite series. In Eq. (14) we define the volume element for the RRS path integral $\tilde{d}[a]=da_1\cdots da_{k_m}$, One repeats the steps starting from Eq. (8) through Eq. (11), but replaces the infinite tail series $\Sigma_{k=k_{m+1}}^\infty a_k \Lambda_k(u)$ with $\Sigma_{k=k_{m+1}}^{k_m'} a_k \tilde{\Lambda}_k(u)$ everywhere to find the functions $\tilde{\Lambda}_k(u)$ that best reproduce the convergence of the partial averaging method. The requirement for $\tilde{\Lambda}_k(u)$ is easily derived,}

$$\sum_{k=k_{m+1}}^{k_m'} \tilde{\Lambda}_k(u) = u(1-u) - \sum_{k=1}^{k_m} \Lambda_k(u).$$

So far, the theoretical treatment is identical to that employed in Euclidean spaces; for non-Euclidean spaces there
and we have used the following convenient notation:

\[ J = \left( \frac{1}{2} \pi \right)^{N/2} \beta^{-N/2} \int \ldots \int \, d[a], \]

Therefore, for an \( N \) point quadrature, the quantity \( J \) associated with the change of variables, \( dq \rightarrow da \), is \( J = (h^2 \beta)^{N/2} J_A \). Upon combining all these results, \( \rho^{\text{RW}} \) becomes,

\[ \rho^{\text{RW}}(q, q', \beta) = \left( \frac{1}{2} \pi \right)^{N/2} \beta^{-N/2} J_A \int d[a] \exp \left\{ - \beta \int_0^1 du U(q^\mu(u)) \right\}. \]
\[ \langle E \rangle_\beta = \frac{D}{2\beta} + \left\langle \frac{\partial}{\partial \beta} \left[ \beta \int_0^1 du U(q^\mu(u)) \right] \right\rangle. \] (28)

\[ \frac{C_V}{k_B} = \frac{D}{2} + \frac{D^2}{4} + D\beta^2 \left\langle \left[ \beta \int_0^1 du U(q^\mu(u)) \right]^2 \right\rangle \]

\[ + \beta^2 \left\langle \left[ \beta \int_0^1 du U(q^\mu(u)) \right] \right\rangle^2 \]

\[ - \beta^2 \left\langle \frac{\partial^2}{\partial \beta^2} \left[ \beta \int_0^1 du U(q^\mu(u)) \right] \right\rangle \]

\[ - \left\langle -\frac{D}{2} - \beta \left[ \beta \int_0^1 du U(q^\mu(u)) \right] \right\rangle^2. \] (29)

where the derivatives with respect to \( \beta \) are evaluated numerically. However, we do not divide by the effective masses when we rescale the path coefficients primarily because in non-Euclidean spaces effective masses (such as moments of inertia) are dependent on the configuration. Consequently, one is forced to decide on the precise form for the reference metric tensor \( g^{(0)}_{\mu \nu} \). The tensor \( g^{(0)}_{\mu \nu} \) is the “best constant” representation of \( g_{\mu \nu} \) on the manifold. We prefer to leave mass dimensions on the path coefficients to yield a “cleaner” and more general formalism for differential manifolds that is completely independent of the choice of \( g^{(0)}_{\mu \nu} \).

III. NUMERICAL TESTS

A. The harmonic chain

In using the adiabatic approximation, one assumes that there are at least two sets of degrees of freedom. One set is characterized by very low frequencies associated with intramolecular degrees of freedom, while the other set is characterized by relatively elevated frequencies associated with intermolecular degrees of freedom. If the molecules in the sample of condensed matter are not too large, there is a relatively large gap in the frequencies between these two sets. To study how the convergence of PIMC simulations is affected by releasing intramolecular constraints, let us consider the following example which represents a simplified model of a relatively large sample of condensed molecular matter. Let us consider a chain of \( n \) point masses, of mass \( m \), connected by ideal springs (Fig. 1). There are two sets containing \([n/2]\) springs. The springs in the first set share the same value for the force constant \( k_1 \). Those in the second set share the same value, \( k_2 \). The spring constants alternate between these two values along the chain. Each point mass is only allowed to translate backward and forward along the direction of the chain. In Fig. 1, \( k_1 \) represents the force constant for the intramolecular degrees of freedom, and \( k_2 \) the same for the intermolecular ones. The mass-weighted Hessian for the flexible case is the \( n \times n \) matrix \( H \).

\[ \begin{pmatrix} k_1 & -k_1 & 0 & 0 \\ -k_1 & k_1 + k_2 & -k_2 & 0 \\ 0 & -k_2 & k_1 + k_2 & -k_1 \\ 0 & 0 & -k_1 & k_1 + k_2 \end{pmatrix}. \] (30)

The adiabatic approximation for the model is sketched at the bottom half of Fig. 1. Each pair of point masses connected by a spring with constant \( k_1 \) is now replaced by a 2m point mass; each of these is connected to its neighbor(s) by an ideal spring with constant \( k_2 \). The mass-weighted Hessian is now \([n/2] \times [n/2]\) in dimension.

\[ H^{(a)} = \frac{1}{2m} \begin{pmatrix} k_2 & -k_2 & 0 & 0 \\ -k_2 & 2k_2 & -k_2 & 0 \\ 0 & -k_2 & 2k_2 & -k_2 \\ 0 & 0 & -k_2 & 2k_2 \end{pmatrix}. \] (31)

Figure 2 contains the eigenvalues of the flexible harmonic chain with \( n=1000 \) point masses for \( k_1=0.1 \) a.u. and \( k_2=0.001 \) a.u. The eigenfrequencies \( \omega_i \) are sorted in ascending order, scaled by the square root of the mass, and plotted against the sorting index. The eigenfrequencies \( \omega_i \) of the “intramolecular” degrees of freedom (\( i \leq 500 \)) are on the left hand side of the graph in Fig. 2; those for the intermolecular degrees of freedom cases are on the right. The value of \( \omega_i \) are discrete because the size \( n \) is finite. However, there is a gap much larger than usual between \( \omega_{501} \) and \( \omega_{500} \). With close inspection of Fig. 2, one can easily verify that \( \omega_{i+1} \approx \sqrt{2k_1/m} \), for \( 500 < i \leq 1000 \), whereas \( \omega_i \approx \omega' = \sqrt{2k_2/m} \) as \( i \to 500 \). The adiabatic limit produces a graph in the \( 1 \leq i \leq 500 \) range (white squares) that is indistinguishable from the flexible case in the same index range on the scale of Fig. 2.
1. The O(N) convergence of the quantum heat capacity

Once the frequencies are computed from the eigenvalues of the Hessian, one can study the convergence of the path integral estimators using analytical formulas. Since the problem is reduced to a set of \( n \) independent oscillators, it is simple to generalize well known analytical expressions for the thermodynamic properties at finite \( N \) of a single harmonic oscillator. \( N \) is the Trotter number or the number of random series coefficients. There are several questions that can be explored quickly with the simple harmonic chain. The first three of the following four questions, in particular, are important for the present discussion: How do the thermodynamic properties converge with the adiabatic approximation compared to the unconstrained counterpart? How much numerical advantage does the adiabatic approximation yield at cold temperatures? How does the numerical advantage gained by using constraints change for algorithms with improved convergence relative to the linear (or primitive) one? Does the size of the chain impact the convergence properties? To answer these questions we use the frequency sets \( \{ \omega_i \}_{i=1}^n \) for the flexible harmonic chain and \( \{ \omega_i^m \}_{i=1}^{m^2} \) for the adiabatic approximation to calculate the heat capacity of the chain at several temperatures. To simulate linear convergence on \( C_V/k_B \), we use the following expression derived from the discretized path integral with \( N \) slices,

\[
\left( \frac{C_V}{k_B} \right)_N = \sum_i \frac{1}{4} \left( \frac{\hbar \omega_i}{2} \right)^2 \times \left\{ \frac{2}{\omega_i + \omega_i^{-1}} \right\}^2 + \frac{1}{N+1} \left[ \frac{\omega_i^N + \omega_i^{-N}}{\omega_i^{N+1} - \omega_i^{-N+1}} \right] \left( \frac{\omega_i - \omega_i^{-1}}{\omega_i + \omega_i^{-1}} \right)^2,
\]

where

\[
\bar{\omega}_i = \frac{1}{2N+1} + \sqrt{1 + \frac{1}{4N+1} \left( \frac{\hbar \omega_i}{2} \right)^2}.
\]

For the following discussion it is convenient to introduce two dimensionless temperature scales,

\[
T^* = \frac{1}{\beta \hbar \sqrt{2k_2/m}}, \quad T^* = \frac{1}{\beta \hbar \sqrt{2k_1/m}}.
\]

Let us consider the case for the harmonic chain with \( m=1, k_1=0.1 \) a.u., \( k_2=0.001 \) a.u. For this case, the reader should note that \( T^*=T^*/10 \). Figure 3 contains graphs of the function in Eq. (32) plotted versus \( T^* \) for several values of \( N \). The left graph contains several adiabatic cases for different values of \( N \), except for the set of points connected by the heavy dashed line; these correspond to the converged quantum result. The graph on the right contains plots for different values of \( N \) for the flexible chain. The results in Fig. 3 demonstrate that the adiabatic approximation yields a substantial numerical advantage while remaining very accurate in the \( 0.1 \leq T^* \leq 1 \) range; in this, the adiabatic approximation can compute \( C_V/k_B \) with \( N=16 \), whereas the flexible computation needs a much greater number of slices (or path variables) to achieve a comparable convergence.
2. The convergence of a higher order quantum heat capacity estimator

The results obtained with Eq. (32) are compared with those obtained with the Fourier path integral $T$-$H$ estimator since its analytical expression for the harmonic oscillator displays cubic convergence. The finite $N$, $T$-$H$ heat capacity estimator for the harmonic chain is

$$
\left( \frac{C_V}{k_B T/N} \right) = \sum_{i} \left\{ \frac{1}{2} - \frac{\beta \hbar \omega_i}{24} + \gamma_i + \frac{1}{2(1 - 8 \alpha_i)} - \frac{8(\alpha_i - \eta_i)}{(1 - 8 \alpha_i)^2} \right\},
$$

where $\alpha_i$, $\gamma_i$, and $\eta_i$ are the following sums over the random series coefficient index $k$:

$$\alpha_i = \sum_{k=1}^{N} \frac{(\beta \hbar \omega_i/k \pi)^2}{(k \pi)^2 + (\beta \hbar \omega_i)^2},$$

$$\gamma_i = \sum_{k=1}^{N} \frac{\beta \hbar \omega_i}{k \pi} \left( \frac{(\beta \hbar \omega_i)^2}{(k \pi)^2 + (\beta \hbar \omega_i)^2} \right),$$

$$\eta_i = \sum_{k=1}^{N} \frac{2}{(k \pi)^2 + (\beta \hbar \omega_i)^2}.$$

A set of curves similar to those in Fig. 3 is produced with Eq. (35) and plotted in Fig. 4. The superiority of the $T$-$H$ estimator is clear when one compares the curves in Fig. 4 with their counterparts in Fig. 3 for both the flexible and rigid cases. The numerical advantage of the adiabatic approximation in the $0.1 \leqslant T \leqslant 1$ range remains clear even when convergence is accelerated.

B. The particle in a ring and the hindered rotations of a rigid asymmetric top

A number of successful numerical tests have been carried out using the primitive Fourier path integral with projection coordinates in ellipsoids of inertia in earlier works. The testing of every part of an algorithm that performs path integration over a manifold composed of Cartesian products of many inertia ellipsoids with the RRS is a complex task. It is productive to brake down such algorithms by testing the theory gradually with systems of increasing complexity. We use the monodimensional non-Euclidean space $S^1$ to verify that the improved RRS method converges with better properties than the corresponding primitive algorithm. The particle in a unit radius ring with 207 a.u. of mass is subjected to a sinusoidal potential,

$$V = \frac{4 \xi}{\xi^2 + 4} = \cos \theta,$$

where $0 < \theta < 2 \pi$ is the angular coordinate and $\xi$ is the stereographic projection. The set of points in the $x$-$y$ plane that are in the ring centered at the origin is mapped with the following equations:

$$x = 4R^2 \frac{\xi}{(\xi^2)^2 + 4R^2}, \quad y = R \frac{(\xi^2) - 4R^2}{(\xi^2)^2 + 4R^2}.$$

The variable $\xi$ is obtained from the point $p \in S^1$ by projecting with a straight line from the point (0,R) through $p$ onto line $y=-R$. The map in Eq. (40) follows from simple geometric arguments. Two PIMC simulations aimed at testing Eqs. (23), (28), and (29) are carried out; one simulation uses $k_m = k_m$ corresponding to the primitive algorithm and the other uses $k'_m = 4k_m$ for the RRS. Both simulations use $k_m = 8$ and are comprised of $1 \times 10^6$ “warmup” moves followed by $1 \times 10^6$ moves over which data points are collected. Both simulations are repeated independently ten times to estimate the statistical error. The statistical error is estimated from the standard deviation to obtain a 95% confidence interval. In Fig. 5 the heat capacity of the particle in a ring computed with the primitive algorithm is compared with the same
quantity computed with the RRS method and with the exact result obtained by diagonalization. The particle in a ring is a simple and convenient system that has proved essential in testing new PIMC methods in non-Euclidean spaces.

Additional numerical tests are performed in the three-dimensional inertia ellipsoid. \( I_\lambda (\lambda=1,2,3) \) represents the eigenvalues of the moment of inertia tensor; the appropriate metric tensor for a rigid top is

\[
g_{\mu \nu} = I_\lambda J_{\mu \lambda} J_{\nu \lambda} J_{\nu \mu},
\]

where the symbol \( J_{\mu \lambda} \) represents a set of 18 independent functions obtained by writing the kinetic energy of a rigid asymmetric top in the laboratory frame in terms of the Euler angles \( \theta, \phi, \psi \). The symbol \( f_{\mu \lambda}^\mu \) represents the Jacobian matrix element for the transformation between the Euler angle \( q^\mu \) and the stereographic projection \( q^\mu \). The \( \xi^1, \xi^2, \xi^3 \rightarrow \theta, \phi, \psi \) map is

\[
\xi^3 = \sin \frac{\theta}{2} \sin \frac{\phi - \psi}{2} \left( \frac{1 + \cos(\theta/2)\sin((\phi + \psi)/2)}{1 - \cos(\theta/2)\sin((\phi + \psi)/2)} + 1 \right).
\]

(47)

We use geometric arguments similar to those for \( S^1 \) to derive the map and its inverse from Eulerian angles to the stereographic projections by using the four-dimensional quaternion space and the spherical constraint. To simplify the notation further we introduce seven auxiliary quantities,

\[
d^1 = \sqrt{16(\xi^1)^2 + [(\xi^1)^2 + (\xi^2)^2 + (\xi^3)^2 - 4]^2},
\]

(48)

\[
d^2 = \sqrt{(\xi^2)^2 + (\xi^3)^2},
\]

(49)

\[
d^3 = (d^2)^2, \quad d^4 = (d^1)^2,
\]

(50)

\[
d^5 = 8(\xi^1)^2 - 4[(\xi^1)^2 + (\xi^2)^2 + (\xi^3)^2 - 4],
\]

(51)

\[
d^6 = (\xi^1)^2 + (\xi^2)^2 + (\xi^3)^2 + 4,
\]

(52)

\[
d^7 = (\xi^1)^2 + (\xi^2)^2 + (\xi^3)^2 - 4.
\]

(53)

Then, the Jacobian is

\[
J_{\mu \nu} = \begin{pmatrix}
-\frac{16d^2\xi^1}{d^1d^6} & \frac{2d^2\xi^2}{d^1d^6} & \frac{2d^2\xi^3}{d^1d^6} \\
\frac{d^5}{d^3} & \frac{8\xi^1\xi^2}{d^3} + \frac{\xi^3}{d^3} & \frac{8\xi^1\xi^3}{d^3} - \frac{\xi^2}{d^3} \\
\frac{d^5}{d^3} & \frac{8\xi^2\xi^2}{d^3} - \frac{\xi^3}{d^3} & \frac{8\xi^2\xi^3}{d^3} + \frac{\xi^2}{d^3}
\end{pmatrix},
\]

(54)

while the nonvanishing \( f_{\mu \lambda}^\mu \) functions are

\[
f_{11}^1 = \left( \frac{4\xi^1\xi^2 - \xi^3 d^1}{d^1d^2} \right)^2,
\]

(55)

\[
f_{11}^1 = \left( \frac{4\xi^1\xi^2 + \xi^3 d^1}{d^1d^2} \right)^2,
\]

(56)

\[
f_{12}^1 = \frac{f_{21}^1 = - f_{12}^2 = - f_{22}^1 = \frac{8(4\xi^1\xi^2 - \xi^3 d^1)(4\xi^1\xi^3 + \xi^2 d^1)}{(d^1d^2)^2d^3d^2}},
\]

(57)

\[
f_{22}^1 = \frac{64(4\xi^1\xi^3 + \xi^2 d^1)}{(d^1d^2)^4},
\]

(58)

\[
f_{22}^2 = \frac{64(4\xi^1\xi^2 - \xi^3 d^1)}{(d^1d^2)^4},
\]

(59)

\[
f_{22}^3 = \frac{2d^4}{(d^1d^2)^2 - 1},
\]

(60)

\[
f_{22}^3 = \frac{2d^4}{(d^1d^2)^2 - 1},
\]

(61)

\[
f_{22}^3 = 1.
\]

(62)

Clearly, \( g_{\mu \nu} \) is analytical, as are its gradient and its Hessian; however, the analytical expressions for these as functions of
\(\xi^a\) are formidable. Therefore, it is simpler to compute \(J_{\mu',\nu'}^a\) and \(\mathcal{J}_{\mu',\nu'}\) and translate the sum in Eq. (41) directly into code.

The eigenvalues of the inertia tensor \(I_k\) are chosen to be identical to those of rigid water, namely, \(I_1, I_2, I_3 = 12.614, 8588, 4026\) a.u., respectively. The rotations are hindered using an external scalar field represented by the function

\[
V(\xi^1, \xi^2, \xi^3) = -V_0 \frac{\xi^1 + \xi^2 + \xi^3}{(\xi^1)^2 + (\xi^2)^2 + (\xi^3)^2 + 4}.
\]

The value of \(V_0\) is 0.04 hartree. \(V_0\) is in the same order of magnitude of the cohesive energy of a water molecule in the bulk. \(V_0\) is about six times larger than the binding energy of the water dimer and about two times larger than the typical cohesive energy of a water molecule in the octamer. The heat capacity of the system at various temperatures is presented in Fig. 6 for a number of values of core path variables \(k_m\). The simulations are carried out between 20 and 1000 K, in regular increments of 20 K using Metropolis sampling. The heat capacity is estimated using the RRS with \(k'_m = k_m\) and the numerical derivative estimator in Eq. (29). A total of 42 independent samples are collected at every temperature using the Metropolis algorithm. Each sample consists of \(1 \times 10^{10}\) moves used to reach the asymptotic distribution, followed by \(1 \times 10^{10}\) moves over which the potential energy, the total energy, and the heat capacity are collected. Each move consists of a configuration change \(\xi^1, \xi^2, \xi^3 \rightarrow (\xi^1, \xi^2, \xi^3)\) and the translation of a randomly selected path variable for each coordinate \(\alpha^1_k, \alpha^2_k, \alpha^3_k \rightarrow (\alpha^1_k, \alpha^2_k, \alpha^3_k)'\). \(k\) is an integer selected using a uniform discrete distribution in \((1 \leq k \leq k_m')\). The data graphed in Fig. 6 have a number of reassuring features. The \(k_m = 0\) curve is the classical estimate of the heat capacity. The \(k_m = 0\) value of \(C_v/k_B\) clearly approaches the proper equipartition value. The classical value of \(C_v/k_B\) decreases systematically as the temperature increases, indicating that the hindered rotations gradually develop a free character at high temperatures. The heat capacity estimators for \(k_m > 0\) behave very well statistically and show a clear convergence pattern that one anticipates for such a system. All the \(k_m > 0\) data merge into the classical data at high temperatures. The system behaves classically only above 500 K. This temperature is large compared to the freezing and boiling temperature of bulk water. The convergence information gathered with this simpler model is indicative of the convergence pattern that a bulk water simulation may display. The \(k_m = 4\) simulation is converged for \(T > 260\) K. The \(k_m = 16\) simulation converges for \(T > 80\) K. The \(k_m = 32\) simulation is probably converged for \(T > 60\) K; a cluster of rigid water molecules may require relatively larger values of \(k_m\) to converge at these temperatures.

### C. Simulations of water clusters

#### 1. The manifold for \(n\) asymmetric rigid tops

Let \(q^a\) represent the \(6n\)-dimensional vector associated with a configuration point in the space \((R^3 \otimes I^3) \otimes (R^3 \otimes I^3) \otimes \cdots (R^3 \otimes I^3)\), where \(\otimes\) is the Cartesian product, \(R^3\) is the three-dimensional Euclidean space for the center of mass of a top, and I symbolizes the inertia ellipsoid, the configuration space for the orientations. In deriving the algorithms and for the implementations of our code, we use the following ordering for the coordinates: \(q^{6(i-1)+1}, q^{6(i-1)+2}, \) and \(q^{6(i-1)+3}\) \((1 \leq i \leq n)\) represent, respectively, the \(x, y,\) and \(z\) coordinates of the center of mass for the \(i\)th top; \(q^{6(i-1)+4}, q^{6(i-1)+5},\) and \(q^{6(i-1)+6}\) \((1 \leq i \leq n)\) represent the three stereographic projection coordinates for the \(i\)th top. With these definitions the metric tensor takes the following block-diagonal form:

\[
g_{\mu\nu} = \begin{pmatrix} g^{(1)}_{11} & 0 & 0 \\ 0 & \cdots & \cdots \\ 0 & 0 & g^{(n)}_{nn} \end{pmatrix}.
\]

The molecular metric tensor \(g^{(i)}\) is represented by a \(6 \times 6\) block-diagonal matrix,

\[
g^{(i)} = \begin{pmatrix} m & 0 & 0 \\ 0 & m & 0 \\ 0 & 0 & m \\ 0 & 0 & 0 \end{pmatrix} G^{(i)}.
\]

with \(m\) representing the mass of the molecular top and \(G^{(i)}\) representing the metric tensor associated with the orientations. The six independent elements of \(G^{(i)}\) are calculated using Eq. (41). Given the sparse nature and block-diagonal structure of \(g_{\mu\nu}\) it is not difficult to design algorithms that only require \(O(n)\) operations for the computation of \(g_{\mu\nu} \dot{q}^\mu \dot{q}^\nu\),

\[
g_{\mu\nu} \dot{q}^\mu \dot{q}^\nu = \sum_{i=1}^{n} g^{(i)}_{\mu\nu} \dot{q}^{6(i-1)+\mu} \dot{q}^{6(i-1)+\nu}.
\]

The evaluation of \(g_{\mu\nu} \dot{q}^\mu \dot{q}^\nu\) requires CPU times comparable to those needed for the computation of an empirical potential surface in the equidimensional space. Analytical expressions for the Hessian elements of \(g_{\mu\nu}\) can be obtained. However, the computation of the Hessian of \(g_{\mu\nu}\) is just as expensive as the Hessian of a typical empirical potential surface of equal dimension. Therefore, the present formulation of the RRS method in curved spaces is considerably cheaper than the previous one.

It is important to clarify that our use of the space \((R^3 \otimes I^3) \otimes (R^3 \otimes I^3) \otimes \cdots (R^3 \otimes I^3)\) does not introduce bias in the distribution function of the constrained system. In
move blocks in the non-Euclidean manifold \((R^3 \otimes I^3) \otimes (R^3 \otimes I^3) \otimes \cdots (R^3 \otimes I^3)\). Each block consists of \(1 \times 10^6\) warmup moves followed by \(1 \times 10^6\) simulation moves. The standard error in the mean is used to estimate the 95% confidence interval. The confidence interval is represented graphically by error bars in Fig. 8. Each move consists of a translation or a rotation of a randomly selected molecule, and a translation of one path variable for each coordinate, e.g., \(a_k^6(r-1)+1\), \(a_k^6(r-1)+2\), \(a_k^6(r-1)+3\) \(\rightarrow\) \((a_k^6(r-1)+1\), \(a_k^6(r-1)+2\), \(a_k^6(r-1)+3\)). The integer \(k\) is in \((1 \leq k \leq k_m)\) and is selected randomly from a uniform discrete distribution in the set. The simulations employ parallel tempering with 40 walkers at evenly spaced temperatures between 50 and 500 K. To define the cluster, a constant volume continuous Lee-Barker-Abraham spherical cavity is employed,

\[
V_{LBA} = \sum_{i=1}^{8} \frac{(r_{O_i} - R_{CM})^{20}}{2R_C},
\]

where \(r_{O_i}\) is the location of the oxygen atom for the \(i\)th water molecule, \(R_{CM}\) is the center of mass of the cluster, and \(R_C\) is 9.0 bohr.

The \(k_m=24\) graph is in the uniform convergence asymptote above 100 K, while the heat capacity is still converging nonuniformly below 100 K with \(k_m=24\) core coefficients. The data sets in Fig. 7 are graphed without error bars to make the graphs less cluttered and to make the convergence pattern stand out. The quantum heat capacities in Fig. 8 are computed with \(k_m=24\), which are converged within the statistical fluctuations for \(T \geq 100\) K for the octamer. The difference between the classical and the quantum simulations remains substantial at 500 K for all sizes; nevertheless, it is clear in both Figs. 7 and 8 that the two estimates converge at some higher \(T\).

D. Characterization of the random walks for \((H_2O)_8\)

Given the large discrepancies between the classical and the quantum simulations, we explore the simple point charge (SPC) potential energy surface for the dimer through the octamer to find all the important minima; in turn these are used

![Figure 7](image_url)  
**FIG. 7.** The heat capacity of the water octamer for various values of \(k_m\) [Eq. (29)] showing the convergence profile of the estimator.

![Figure 8](image_url)  
**FIG. 8.** The classical (white circles) and quantum (\(k_m=24\), black squares) heat capacity of water clusters obtained with the RRS method (Sec. II) and the finite difference estimator in Eq. (29).
to characterize the random walks using a Structural Comparison Algorithm (SCA).\textsuperscript{52} The quantum effects are most conspicuous for the octamer, for which the pronounced melting feature in the classical simulation is “washed away” by quantum fluctuations. Therefore, as a check, we minimize all the sizes from the dimer through the octamer and compare our minima with those found by other groups with similar potentials.\textsuperscript{50,53–58} We use the minima of the octamer to characterize the configurations in several random walks. To minimize the structures, we use the full $R^n$ space mapped by Cartesian coordinates, with the potential along the covalent stretching and bending degrees of freedom obtained with the F2 extension.\textsuperscript{59} The candidate structures are generated with the genetic algorithm, and these are quenched with a $T=0$ Brownian dynamics algorithm, which in Euclidean spaces with Cartesian coordinates takes a simple form,

$$\dot{x}^\mu + \gamma x^\mu V + \gamma x^\mu = 0. \quad (68)$$

Here, $\gamma$ is the coefficient of friction. The genetic algorithm is implemented to create 100 candidate structures using simple slicing-rejoining operators. The candidates are quenched, sorted by energy in ascending order, and compared with SCA if the energy difference between two minima is less than $1 \times 10^{-4}$ hartree.

SCA provides us with a measure of distance between two configurations that are being compared. This comparison is obtained by attempting to find the best superposition of two structures through a set of translations, rotations, and a sorting process. Typically a running configuration is compared with a minimum. In the case of the genetic algorithm, we compare the newly quenched candidates with all the minima already known. This eliminates multiple copies of a minimum included in the genetic pool that one uses for the next generation of “children structures.” The key difference with out implementation is that the comparison process allows both enantiomers of optically active minima to contribute to the “gene pool.” Many minima lack an improper rotation axis and consequently display optical isomerism. If all the degenerate replicas of an optically active minimum are eliminated from the pool, only one of the two enantiomers is used in the generation of new candidates. The impact of eliminating enantiomers from the gene pool could have on the efficiency or even the success of a global optimization is not known, but one can anticipate that when a large number of atoms are involved this detail may become important. We run 40 generations for the hexamer, heptamer, and octamer. However, the global minimum for all the sizes from the dimer through the octamer is found very quickly by the genetic algorithm.

The structures we find for the global minima agree qualitatively with those reported in a number of global minimization studies carried out with related potential functions.\textsuperscript{50,53–58} In particular, we find all the cubic isomers reported recently by Nigra \textit{et al.}\textsuperscript{50} In agreement with Nigra \textit{et al.},\textsuperscript{50} we find that the $D_{2d}$ cube is lower in energy than the $S_4$ cube; we find the $C_2$, $C_i$, and five $C_1$ cubes with slightly different energetic orders. We focus our comparison on Nigra \textit{et al.}\textsuperscript{50} since the authors report a characterization of the classical parallel tempering walk similar to the one we carry out. The $C_2$, $C_i$, and $C_1$ cubes have energies comparable to other open minima of the octamer potential energy surface and contribute only slightly in the melting range. Our melting peak in the classical calculation is at $T_{\text{max}}=142.3$ K instead of the $T_{\text{max}}=178.5$ K reported by Nigra \textit{et al.} The potential used by the authors of Ref. 50 is a sophisticated model,\textsuperscript{53} known as the Matsuoka-Clementi-Yoshimine (MCY) potential, constructed by fitting CI computations on the water dimer, and is quite different from the SPC model we use here; discrepancies are to be expected. Therefore, it is rewarding to find that the characterization of the walks generated on the SPC surface that we report in the following is in excellent qualitative agreement with the results obtained by the MCY surface.

We use SCA to characterize the walk in the following manner. Let $x^i_t$ represent the Cartesian coordinates of atom $i$ in a configuration under analysis after a systematic set of translation, rotations, and permutations is carried out to find the best possible overlap with the global minimum configuration. $(x_0^i)_{i=1}^{3n}$ represents the Cartesian coordinates of the atoms in the global minimum. Then,

$$\Delta_0 = \sum_{i=1}^{3n} |x^i_0 - x^i_t| \quad (69)$$

represents the “structural distance” from the global minimum for $(x^i_t)_{i=1}^{3n}$. Histograms of $\Delta_0$ can be used to detect features in the density matrix arising from as solid-solid transformations, melting, and similar thermally induced phenomena. Figure 9 contains two sets of such histograms for $\Delta_0$ sampled in the $T$ interval below the melting peak of the classical heat capacity. The graphs in Fig. 9(a) are obtained by measuring $10^3$ configurations during the classical walk after $1 \times 10^5$ warmup moves. The graphs in Fig. 9(b) are obtained with $k_m=24$. The broad feature between 10 and 40 bohrs in both figures is comprised of cubic structures; these are either the $D_{2d}$ or $S_4$ isomers. The $S_4$ isomer produces a shoulder of this feature above 30 bohrs. In the classical simulation [Fig. 9(a)] this shoulder is detectable at 50 K and gradually gains intensity. The first detected configurational jump for a cube to one of the open structures (typically intersecting rings) in the classical simulation is at 96.2 K. This peak in the histograms of $\Delta_0$ is well differentiated from the broad features associated with cubic structures.

The isomeric distributions we measure in the classical simulation agree qualitatively with those reported by Nigra \textit{et al.}\textsuperscript{50} in a number of key features. We see a significant $D_{2d} \rightarrow S_4$ isomerization present at 50 K. Nigra \textit{et al.} reported the onset of $D_{2d} \rightarrow S_4$ isomerizations at 40 K. Similarly, we find that the other cubic structures (the $C_2$, $C_i$, and $C_1$ cubes) play a small role in the liquid region. This finding agrees with the reported probabilities for this group of structures. In Fig. 8 of Ref. 50, Nigra \textit{et al.} reported the onset of liquid structures above 120 K. We find the onset of liquid structures around 96 K, explaining the difference in the peak temperature for the melting.

Probably, the quantum distributions of $\Delta_0$ [Fig. 9(b)] are not converged at 73.1 K and below. The abrupt appearance of the feature at 75 bohrs at 73.1 K is suggestive of the lack
of convergence for the two distributions of $\Delta_0$ at 61.6 and 50.0 K. Nevertheless, we can see a pattern developing, even at the colder temperatures. The $D_2h \rightarrow S_4$ equilibrium is shifted to the right considerably compared to the classical simulation. Similarly, the cube $\rightarrow$ liquid phase change is shifted. At 84.6 K we can say with confidence that the octamer is melting. However, the quantum distributions of $\Delta_0$ continue to display distinguishable albeit broad features at higher temperatures. The feature centered at 40 bohrs that grows systematically at 40 bohrs between 96.2 and 119 K contains a mixture of $S_4$ and $C_2$, $C_4$, and $C_1$ cubes, whereas the feature that grows around 60 bohrs from 84.6 K contains cubic structures with a significant amount of obliqueness and cubic structures with one missing vertex. The remarkable differences in the distributions of $\Delta_0$ between classical and quantum simulations explain the large differences visible in the thermodynamic properties. One would anticipate that quantum fluctuations would simply broaden the classical distributions of $\Delta_0$ and shift isomerization features to lower temperatures. That shifting and broadening can be gleaned in Fig. 9. However, one would not anticipate the appearance of new features that remain well resolved at elevated temperatures inside the liquid coexistence region. In fact, this has been observed recently in other hydrogen bonded systems.42

IV. CONCLUSIONS

The two main objectives for the work reported in this article are to illustrate the efficiency gained by using holonomic constraints in PIMC simulations of molecular condensed matter and the exploration of the quantum thermodynamic properties of small water clusters.

It is well known that rigid body simulations improve the efficiency of thermodynamic estimators in Metropolis simulations; however, little is known about how holonomic constraints affect the convergence of PIMC simulations. We have investigated a simple one-dimensional harmonic chain using analytical solutions of the thermodynamic path integral for finite Trotter number. For this system the analytical solutions of the path integral expression for the heat capacity as a function of $T$ and the Trotter number (or a number of path variables) clearly support our hypothesis: The choice to use holonomic constraints in PIMC simulations is dictated by the efficiency of a simulation as it relates to convergence properties rather than statistical fluctuations. The present investigation shows that if the smallest frequency in the intramolecular set is at least a factor of 10 greater than the largest frequency in the intermolecular set, then the adiabatic approximation is not only convenient; it is necessary to produce numerical convergence with reasonable means at low temperatures. The numerical tests with the harmonic chain reveal that the primitive algorithm simulation in the adiabatic limit converges better than the cubically convergent estimator for the flexible case. The fully flexible Cartesian simulation becomes unfeasible at low temperatures, where the thermodynamics are dominated by intermolecular rearrangements and the “molecules” are predominantly in the ground vibrational state. Since the results obtained from the harmonic chain and the conclusions that can be drawn are for 1000 point masses, there is evidence that these results may apply to bulk molecular condensed matter as well as clusters.

The graphs in Fig. 4, however, point to the next challenge for the present developments in the statistical thermodynamic theory of condensed molecular matter. Clearly, between $1 < T^* < 2$ the system does not behave classically, and the adiabatic approximation is not satisfactory. Figure 4 suggests that perhaps one only needs to perform a flexible PIMC simulation for $T^* > 1$ and a rigid simulation below this value. However, in more complex models of condensed matter there may be more than one high frequency mode. Furthermore, even the rigid simulation may require parallel tempering runners at temperatures above $T^* = 1$ to overcome potential energy barriers. These complications arise in the quantum simulations of water clusters. With the parameters of the SPC/F2 in Lobaugh and Voth,50 one obtains the following frequencies for water. The fundamental for the $\text{H}–\text{O}–\text{H}$ bend is at 1504 cm$^{-1}$, the $\text{O}–\text{H}$ symmetric stretch is at 3322 cm$^{-1}$, and the asymmetric $\text{O}–\text{H}$ stretch is at 3601 cm$^{-1}$. These frequencies are substantially redshifted from the gas phase data since the SPC/F2 has been designed for molecular dynamics simulations of bulk water. These fre-
APPENDIX: TRANSFORMING FROM EULER ANGLES TO CARTESIAN COORDINATES AND BACK

Most empirical intermolecular potential energy surface models are relatively simple expressions of the Cartesian coordinates for each atom. Consequently, it is important to find the Cartesian coordinates of all the atoms in the molecular top $i$ when the center of mass and the three stereographic projections are known. There are two ways to convert the Cartesian coordinates of all atoms to Euler angles, and then to stereographic projections, each yielding equivalent metrics in the inertia ellipsoid. The active map of the body fixed frame is considered in this Appendix. The transformation of the passive map can be handled in a similar way. The Euler angles can be obtained from the stereographic projections by using Eqs. (42)–(44). The Cartesian coordinates of three points in a rigid body in the center of mass coordinate system can be obtained from the three Euler angles once a reference configuration in the body frame common to all tops is specified. For water clusters we take the following reference configuration:

$$
\begin{bmatrix}
 x \\
 y \\
 z
\end{bmatrix}
= \begin{bmatrix}
 H_1 & 0 & \alpha_c & \gamma_c \\
 O & 0 & \beta_c & 0 \\
 H_2 & 0 & \alpha_c & -\gamma_c
\end{bmatrix},
$$

where for the rigid model of water $\alpha_c=0.9865$, $\beta_c = -0.1243$, and $\gamma_c = 1.529$ bohrs. The reference configuration defines the space axes; this is a configuration point in which the inertia tensor is diagonal. These space axes can be obtained by simply transforming back from the reference body frame to the running configuration, and the inverse mapping can be obtained as well. The problem can be stated as follows. Let the element $\mathbf{R} \in O(3)$ be the rotation matrix that takes the reference configuration to an arbitrary running configuration (namely, the body coordinates in the center of mass Cartesian space). When expressed in terms of Euler angles the representation of the element $\mathbf{R} \in O(3)$ is as follows:

$$
\mathbf{R} = \begin{bmatrix}
 \cos \phi \cos \psi - \cos \theta \sin \phi \sin \psi & \sin \phi \cos \psi + \cos \theta \cos \phi \sin \psi & \sin \psi \sin \theta \\
 -\cos \phi \sin \theta & -\sin \phi \sin \theta & \cos \theta \\
 \sin \phi \sin \psi & \cos \phi \sin \psi & \cos \phi \cos \psi - \cos \theta \sin \phi \sin \psi
\end{bmatrix}.
$$

This definition yields nine equivalences between the Euler angles and the Cartesian coordinates of each atom; from these it is straightforward to derive the following relationships between the trigonometric functions and the center of mass coordinates:

$$
\cos \theta = \frac{z_{H_1} - z_{H_2}}{2 \gamma_c},
$$

$$
\sin \theta = \frac{1}{2 \gamma_c} \sqrt{(x_{H_1} - x_{H_2})^2 + (y_{H_1} - y_{H_2})^2},
$$

$$
\sin \psi = \frac{x_{H_1} - x_{H_2}}{2 \gamma_c \sin \theta},
$$

$$
\cos \psi = \frac{y_{H_1} - y_{H_2}}{2 \gamma_c \sin \theta}.
$$
\[
\cos \phi = -\frac{z_O}{\beta_z \sin \theta}, \quad (A7)
\]
\[
\sin \phi = \frac{1}{\cos \psi} \left( \frac{x_{H1} + x_{H2}}{2a} - \cos \theta \cos \phi \sin \psi \right). \quad (A8)
\]

Some of these equations become infinite for those special cases (rare events in random walks), where \( \sin \theta = 0 \). Two possibilities must be considered:

Case 1: \( \theta = 0 \). The rotation matrix becomes

\[
R = \begin{pmatrix}
\cos(\phi + \psi) & \sin(\phi + \psi) & 0 \\
-\sin(\phi + \psi) & \cos(\phi + \psi) & 0 \\
0 & 0 & 1
\end{pmatrix}. \quad (A9)
\]

The body coordinates now are

\[
x_{H1} = x_{H2} = \alpha_x \sin(\phi + \psi),
\]
\[
x_O = \beta_z \sin(\phi + \psi),
\]
\[
y_{H1} = y_{H2} = \alpha_x \cos(\phi + \psi),
\]
\[
y_O = \beta_z \cos(\phi + \psi),
\]
\[
z_{H1} = -z_{H2} = y_z; \quad z_O = 0. \quad (A14)
\]

Case 2: \( \theta = \pi \). Then, the element of O (3) is

\[
R = \begin{pmatrix}
\cos(\phi - \psi) & \sin(\phi - \psi) & 0 \\
-\sin(\phi - \psi) & -\cos(\phi - \psi) & 0 \\
0 & 0 & -1
\end{pmatrix}. \quad (A15)
\]

The center of mass space fixed frame now gives the following coordinates for the atoms:

\[
x_{H1} = x_{H2} = \alpha_x \sin(\phi - \psi),
\]
\[
x_O = \beta_z \sin(\phi - \psi),
\]
\[
y_{H1} = y_{H2} = -\alpha_x \cos(\phi - \psi),
\]
\[
y_O = -\beta_z \cos(\phi - \psi),
\]
\[
z_{H1} = -z_{H2} = -y_z; \quad z_O = 0. \quad (A20)
\]

Both special cases can be considered without loss of generality as a single rotation about the space \( \phi \) axis by \( \phi \) radians. Therefore, one arrives at

\[
\sin \phi = x_{H1}/\alpha_x,
\]
\[
\cos \phi = \pm y_{H1}/\alpha_y
\]

for the \( \theta = 0 \) and \( \pi \), respectively. A translation of the center of mass for each top completes the transformation from the coordinates of the center of mass—projection space to the full Euclidean space \( (R^3 \otimes I^3) \otimes (R^3 \otimes I^3) \cdots (R^3 \otimes I^3) \rightarrow R^n \) which contains the Cartesian coordinates of each atom.