Guided Diffuison Quantum Monte Carlo for Calculating Zero Point Energies

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Introduction

- ▶ DQMC solves the Schrödinger equation and gives the quantum-mechanical ground state energy to then calculate the zero point energy (ZPE).
- ► A Gaussian guiding wave function is used to improve the performance.
- ▶ DFTB+ is used for calculation the energy.[1]

The Unguided DQMC Algorithm [2]

- The stationary solution of the diffusion equation satisfies the Schrödinger equation and has the propagator $G(x,y;\Delta t) = \frac{1}{\sqrt{2\pi\Delta t}} e^{-(x-y)^2/(2\Delta t)} e^{-\Delta t(V(y)-E_T)}.$
- ▶ We simulate a ensemble of walkers to reach this stationary solution and the Schrödinger equation is solved.
- The propagator consists of a diffusion term $\frac{1}{\sqrt{2\pi\Delta t}}e^{-(x-y)^2/(2\Delta t)}$ and a branching term $e^{-\Delta t(V(y)-E_T)}$.
- ▶ The diffusion term is simulated by randomly displacing the walker from its previous position according to a Gaussian distribution.
- ► The branching term updates the weight of a walkers according to which the walker survives, reproduces or dies.
- ▶ The trial energy E_T is then adjusted to keep the population of walkers stable.
- ▶ With time, the trial energy E_T converges.



The Guiding Wave Function [3]

- ► Harmonic approximation of the potential energy surface with the Hessian matrix.
- ► Gaussian guiding wave functions along the normal modes $\psi_{\rm T}({\rm x}) = \mathcal{N} {\rm e}^{-{\rm x}^2/(2\sigma^2)}$.
- ► The width of the Gaussian is given by $\sigma^2 = \frac{1}{\omega m}$, where ω^2 is the corresponding eigenvalue of the Hessian matrix.
- This is incorporated to the algorithm with a drift velocity $v(x) = \nabla_x \psi_T(x)$ which is added to the pure diffusion and a kinetic energy term in the local energy $E_L(x) = (H\psi_T(x))/\psi_T(x)$ which replaces the potential energy in the branching step.

The Guiding Wave Function

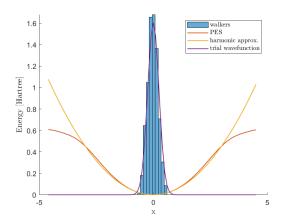


Fig. 1. Typical dimension in the coordinate system of the normal modes.

Input and Output

Input:

- ► Equilibrium geometry of the atoms (must not be perfectly accurate as geometry will be optimized by DFTB+ in the beginning)
- ► Masses of the atoms

Output:

- Zero point energy
- ► Walker positions give a sample of the nucleonic wave function

Results C₂H₆ (first test case)

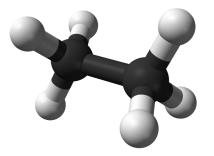


Fig. 2. Input structure of Ethane

First test case to develop the algorithm and check the result with a literature value.

Results C₂H₆ (first test case)

Simulation with 1000 walkers for 2000 time steps

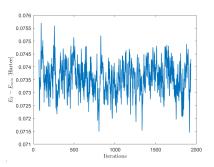


Fig. 3. Fluctuation of the ZPE after equilibration.

ZPE = 0.07356 hartee with a standard deviation of 0.00060 hartee. The literature value is 0.073927 hartree [4].

Results C₂H₆ (first test case)

Simulation with 1000 walkers for 2000 time steps

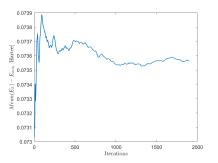


Fig. 4. Convergence of the ZPE after equilibration.

Simulation time: 14 hours. ≈ 0.02 seconds per energy calculation.

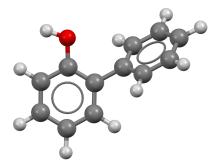


Fig. 5. Input structure of 2-Phenylphenol

Slightly larger molecule to test the performance of the algorithm.

$\underset{C_{12}H_{10}O}{Results}$

Simulation with 1000 walkers for 2000 time steps

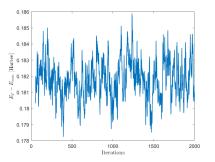


Fig. 6. Fluctuation of the ZPE after equilibration.

ZPE = 0.18184 hartee with a standard deviation of 0.00117 hartee.

$\underset{C_{12}H_{10}O}{Results}$

Simulation with 1000 walkers for 2000 time steps

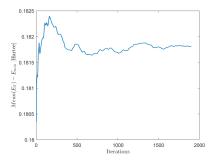


Fig. 7. Convergence of the ZPE after equilibration.

Simulation time: 57 hours. ≈ 0.11 seconds per energy calculation.

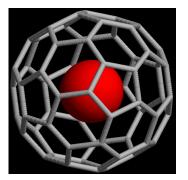


Fig. 8. Input structure of the dihydrogen endofullerene

The dihydrogen endofullerene is expected to show strong anharmonic effects on the zero point energy.

Results $_{\rm H_2 @ C_{60}}$

Simulation with 1000 walkers for 2000 time steps

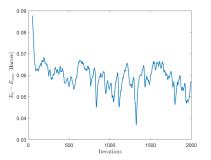


Fig. 9. Fluctuation of the ZPE after equilibration.

 $\mathrm{ZPE} = 0.05914$ hartee with a standard deviation of 0.00569 hartee.

$\underset{H_2 @ C_{60}}{Results}$

Simulation with 1000 walkers for 2000 time steps

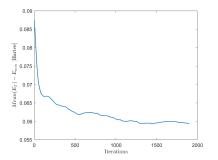


Fig. 10. Convergence of the ZPE after equilibration.

Simulation time: 51 hours. ≈ 0.10 seconds per energy calculation.

Conclusions and Improvements

Conclusions:

- ▶ DQMC delivers moderately accurate results for the ZPE in reasonable time.
- ▶ DQMC is very versatile and can calculate the ZPE of most molecules.

Improvements:

- ► Adjustments for periodic structures
- ▶ Parallelization
- Making it more user-friendly

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