QXMD Hands-on Session

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Outline for QXMD Hands-on

- 1. Build QXMD
- 2. Optimize Water Structure
- 3. Adiabatic Quantum Molecular Dynamics (QMD)
- 4. Non-adiabatic Quantum Molecular Dynamics (NAQMD)

Open Your Google Colab: colab.research.google.com





1. Build QXMD

```
qxmdmpi
data/
control/
CONFIG
```

IN.PARAM: *parallel

filename

*start

*approximation for Exc

*SCF iterations

*molecular dynamics

*supercell

*electronic bands

*dump wavefunctions

*atoms

.....

IN.VELOC

The manual for QXMD can be found at: https://usccacs.github.io/QXMD DEV/

2. Optimize Water Structure

*molecular dynamics

ifmd: 1

dtmd: 0.1d0 nstop: 100

ioptmze: -1: do not optimize atomic coordinates

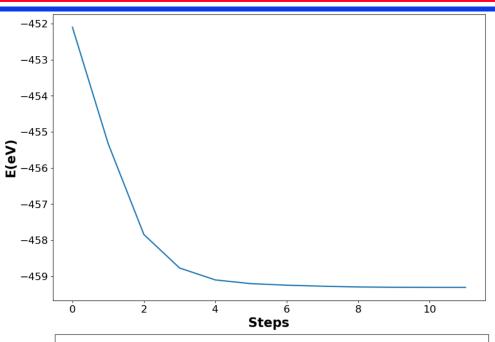
0: Conjugate gradient

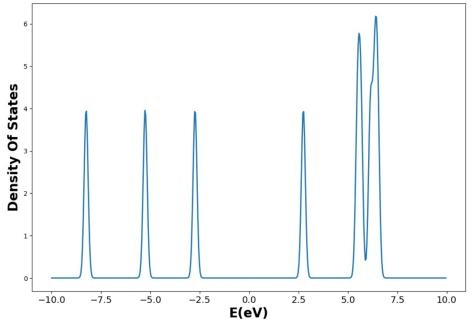
1: Projected velocity Verlet

2: Quasi-Newton method with BFGS formula

tol_energy: 1.d-07

tol_force: 5.d-04





3. Adiabatic Quantum Molecular Dynamics (QMD)

Time(fs)

*molecular dynamics

ifmd: 2:NVE

3:NVT

4:NPT

dtmd: 10.0d0

nstop: 300

treq: 300

liscale: .false.

iscnum: 25

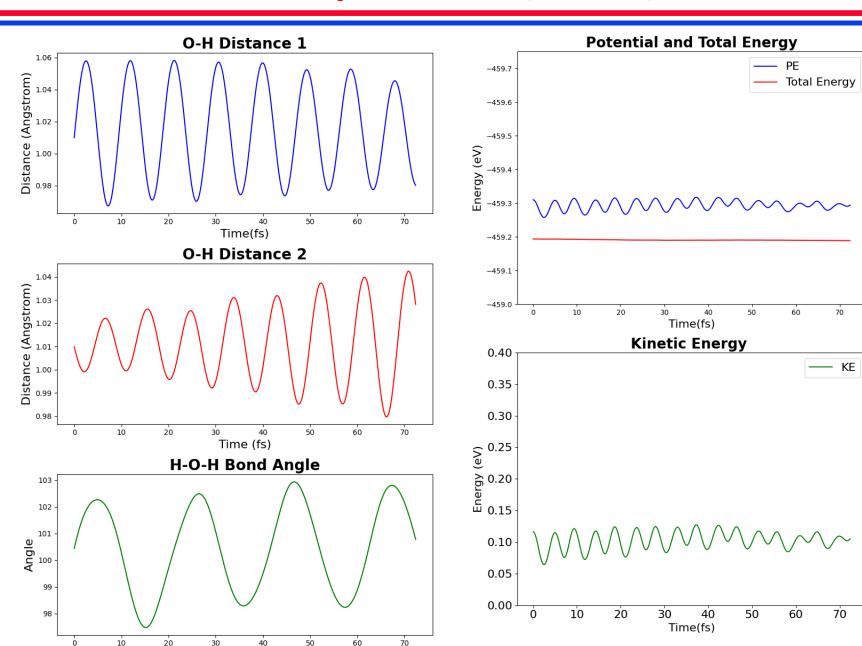
iscstp: 20

ioskip: 1

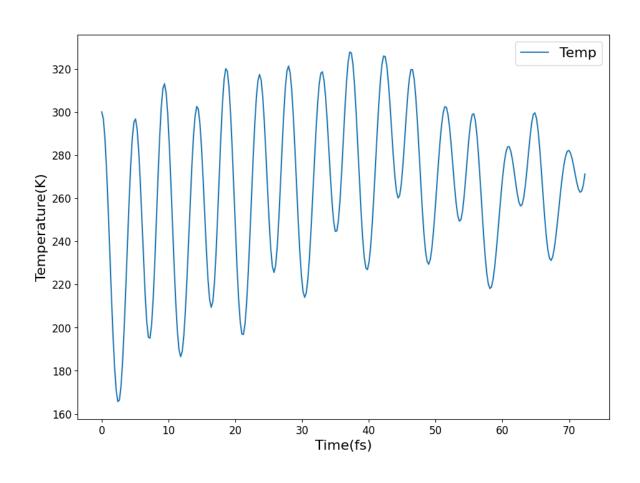
locoor: .true.

lovelo: .true.

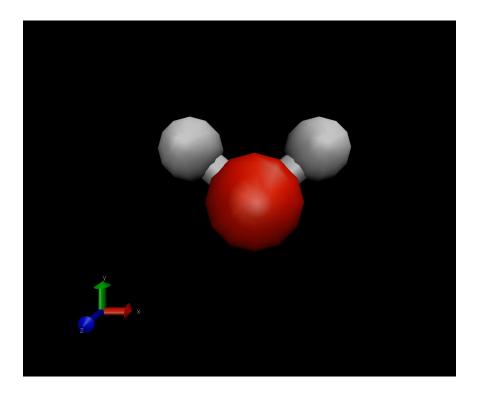
loforc: .true.



3. Adiabatic Quantum Molecular Dynamics (QMD)



Trajectory Visualization



4. Non-adiabatic Quantum Molecular Dynamics (NAQMD)

*molecular dynamics

*TDDFT-MD

ifmd: 2

dtmd: 10.0d0

nstop: 300

treq: 0.1d0

liscale: .false.

iscnum: 25

iscstp: 20

ioskip: 1

locoor: .true.

lovelo: .true.

loforc: .true.

Itddft: .true.

ltddft_fssh : .true.

lfssh_gsscf: .true.

aslh_fssh: 0.8d0

bslh_fssh: 0.13d0

dttddft: 0.04d0

ltddft_start: .false.

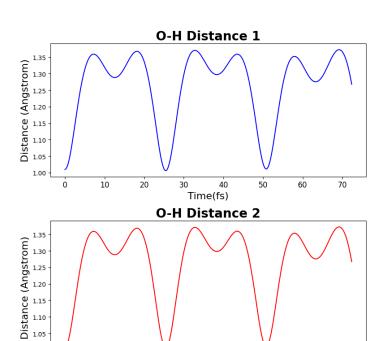
Ifssh_random: .true.

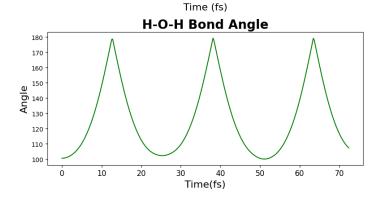
rseed_fssh:1119041003.0000000

nocc_change: 2

numband: 4

occ_new: 1.0 0.0



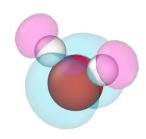


10

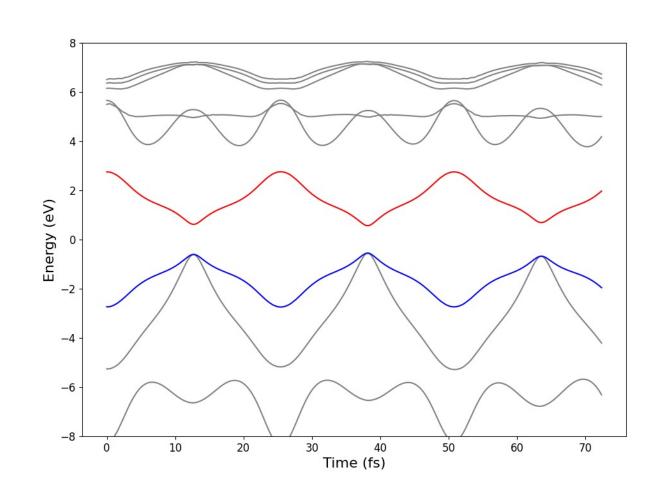
20

4. Non-adiabatic Quantum Molecular Dynamics (NAQMD)

Trajectory Visualization



wavefunction isosurfaces (excited electron (pink), hole (teal)) overlayed on the water molecule.



Recombination Rate (s^{-1}) : 1.644E+09