Tutorial 5

Let them roll!

Hands on (ab initio) Molecular Dynamics

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Hands-on Tutorial Workshop 2011 on *Ab Initio* Molecular Simulations Berlin, 19.07.2011

Molecular Dynamics: the idea

- "Computer experiment"
 - Prepare the sample: select model system and numerically solve **Newton's equations** of motion.
 - Perform measurement
- Here: Born-Oppenheimer Molecular Dynamics
 - Converge electronic density for each step
 - Calculate ab initio forces on atoms [remember J. Wieferink' s lecture on July 13th]

It should be very simple

Newton's equation:

$$F(x,t) = m\ddot{x}$$

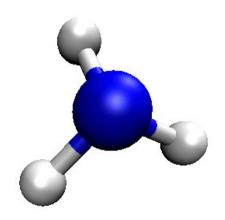
- Naive approach:
- Use just simple integrator (e.g. the Euler integrator)

$$x(t_1) = x(t_0) + v(t_0)\Delta t_1 + \frac{F}{2m}\Delta t_1^2$$

$$x(t_2) = x(t_1) + v(t_1)\Delta t_2 + \frac{F}{2m}\Delta t_2^2$$
:

- Have fast / not so accurate force evaluation ?
- Use big time step fast evolution over time

Would it work?



 $\Delta t = 3fs$

- Not really ...
- It is simple, but details and accuracy are important

Which conditions should we simulate?

- Natural ensemble: microcanonical
 - time-independent Hamiltonian:
 energy is conserved
- Also possible to simulate other ensembles: canonical (discussed further on), NPT, NPH, etc.

A model MD program

- 1.Read essential parameters (temperature, # of atoms, time step, etc.)
- 2.Initialize system positions and velocities

Central

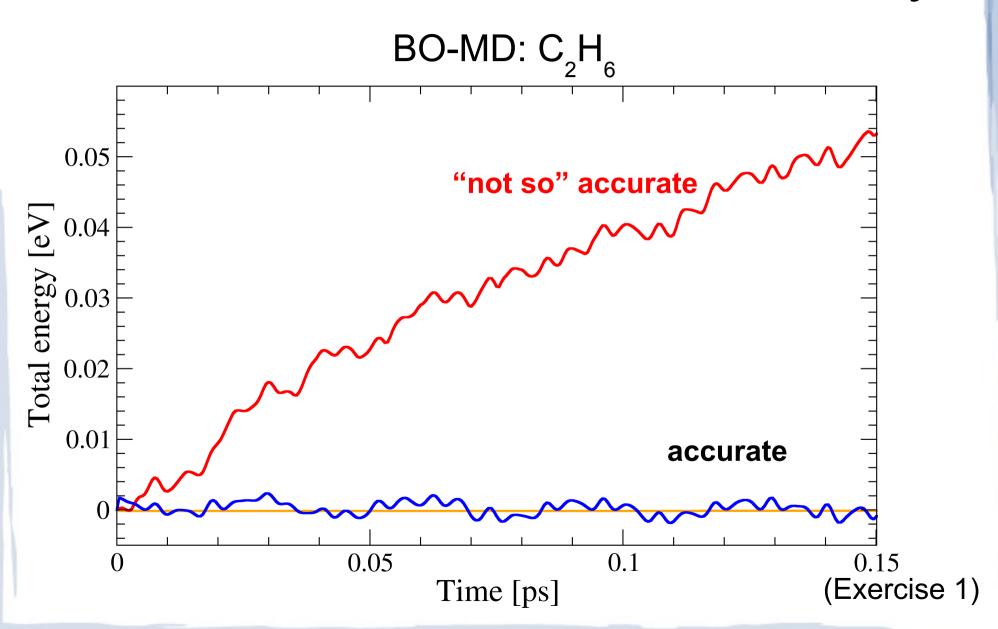
loop

- 3. Evaluate forces
- 4.Integrate equations of motion
- 5.Stop after a given time enough statistics for your measurement

Details of the central loop (Exercise 1 and 2)

- The force evaluation
 - In Born-Oppenheimer Molecular Dynamics (BOMD), forces are evaluated at electronic self consistency.
 - Stability of MD simulations depends critically on the accuracy of the forces.
 - When forces are not accurate enough, there is an undesired systematic deviation from the Born-Oppenheimer surface.

Need of accurate self-consistency



Details of the central loop

- Integrating the equations of motion
 - Verlet algorithm:

$$r(t + \Delta t) = r(t) + v(t)\Delta t + \frac{F(t)}{2m}\Delta t^2 + \ddot{r}\frac{\Delta t^3}{3!} + O(\Delta t^4)$$

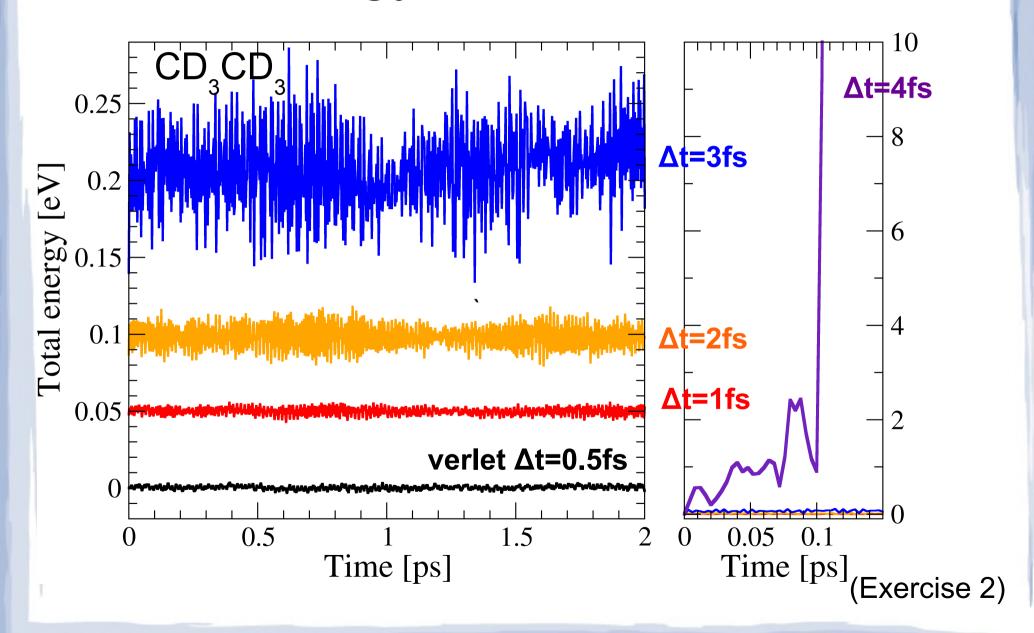
$$r(t - \Delta t) = r(t) - v(t)\Delta t + \frac{F(t)}{2m}\Delta t^2 - \ddot{r}\frac{\Delta t^3}{3!} + O(\Delta t^4) + \frac{F(t)}{m}\Delta t^2 + O(\Delta t^4) + \frac{F(t)}{m}\Delta t^2 + O(\Delta t^4)$$

$$\Rightarrow r(t + \Delta t) \approx 2r(t) - r(t - \Delta t) + \frac{F(t)}{m}\Delta t^2$$

Implemented: Velocity-Verlet

$$v(t + \Delta t) = v(t) + \frac{F(t + \Delta t) + F(t)}{2m} \Delta t$$

How do energy fluctuations look like?



- The idea: couple the system to a thermostat
- Why is it interesting:
 - Experiments are usually done at constant temperature
 - If system suffers a conformational change, energy difference becomes kinetic energy, and temperature in the simulation would change

- Simulating the real coupling:
 - Stochastic thermostat, e.g. Andersen
 - Particle randomly selected, with probability υΔt, has its velocity replaced by one drawn from a Maxwell-Boltzmann distribution at the target temperature (the heat bath).
 - Drawback: since it is stochastic, it destroys the trajectory of the system – not to be used for time dependent properties.

- Simulate the effect of coupling:
 - Velocity rescaling, e.g. Berendsen
 - It does not sample the canonical ensemble
 - Scales velocity with factor λ

$$\lambda = \left[1 + \frac{\Delta t}{\tau} \left(\frac{T_0}{T} - 1\right)\right]^{1/2}$$

- Extended Lagrangian, e.g. Nosé-Hoover
 - Inclusion of the heat bath in the Lagrangian as continuous degrees of freedom

$$\mathcal{L}_{\text{nose}} = \sum_{i} \frac{\mathbf{m}_{i}}{2} s^{2} \dot{\mathbf{r}}_{i} - U\left(\mathbf{r}^{N}\right) + \frac{Q}{2} \dot{s}^{2} + (3N+1) \frac{\ln s}{\beta}$$

- Energy of the system is not conserved anymore, but the energy of the extended system (subsystem + heat bath) is.
- Equations of motion:

$$\mathcal{H}_{\text{nose-hoover}}(\mathbf{p_i}, \mathbf{r_i}, \pi, \eta) = \sum_{i} \frac{\mathbf{p}_i^2}{2m_i} + U(\mathbf{r}^N) + \frac{1}{2Q}\pi^2 + 3N\frac{\eta}{\beta}$$

$$\dot{\mathbf{r}}_i = \frac{\mathbf{p}_i}{m_i}; \ \dot{\mathbf{p}}_i = -\frac{\partial U}{\partial \mathbf{r}_i} - \frac{\mathbf{p}_i \pi}{Q}; \ \dot{\eta} = \frac{\pi}{Q}; \ \dot{\pi} = \left(\sum_i \frac{\mathbf{p}_i^2}{m_i} - \frac{3N}{\beta}\right)$$

Other famous extended Lagrangian

- Car-Parrinello MD
- Wave function extrapolation: Kühne-Parrinello, Niklasson.
- NPT ensemble: Andersen, Parrinello-Rahman-Nosé

- Stochastic velocity rescaling (Bussi-Donadio-Parrinello)
 - Conserved (pseudo-)hamiltonian

$$\tilde{H}(t) = H(t) - \int_0^t \left(\overline{K} - K(t') \right) \frac{dt'}{\tau} - 2 \int_0^t \sqrt{\frac{K(t')\overline{K}}{N_f}} \frac{dW(t')}{\sqrt{\tau}}$$

 In practice, after every velocity verlet step, new velocities are calculated. These velocities are rescaled by a factor α, such that:

$$\alpha^{2} = e^{-\Delta t/\tau} + \frac{\overline{K}}{N_{f}K} \left(1 - e^{-\Delta t/\tau}\right) \left(R_{1}^{2} + \sum_{i=2}^{N_{f}} R_{i}^{2}\right) + 2e^{-\Delta t/2\tau} \sqrt{\frac{\overline{K}}{N_{f}K}} \left(1 - e^{-\Delta t/\tau}\right) R_{1}$$

Thermostats comparison (Ex. 3)

Andersen

- Static averages in the canonical ensemble, but dynamical quantities are meaningless
- Good for equilibration: excites "equipartition-wise" all the modes.

Berendsen

- Does not sample any known ensemble.
- If τ small (strong coupling), yields completely unreliable trajectories, at *seemingly* the right temperature.

Nosé-Hoover

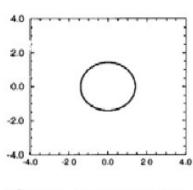
- Rigorously samples the canonical ensemble
- Dynamical quantities (e.g. autocorrelations) are typically seen to be reliable
- But: has ergodicity issues (fixable via Nose-Hoover chains)

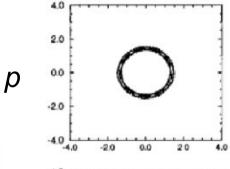
Bussi-Donadio-Parrinello

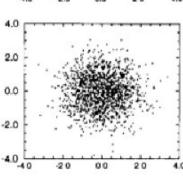
- Rigorously samples the canonical ensemble
- Dynamical quantities are reliable
- No ergodicity issues

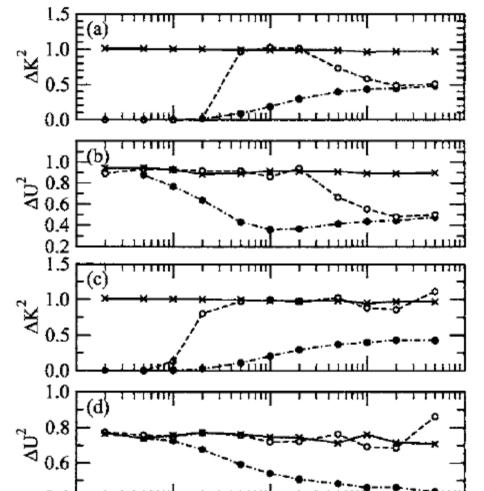
On Ergodicity

Harmonic oscillator









0.1

τ(ps)

0.01

 \times BDP

o NH

Berendsen

Lennard-Jones solid @20K

Lennard-Jones liquid @120K

What can we use MD for, anyway?

- Measures of:
 - Diffusion coefficients in liquids
 - Rotational-vibrational spectra with anharmonic contributions
 - Evaluating the excess free energy via thermodynamic integration
 - (Fast) rates and phenomenology of reactions
 - Out-of-equilibrium quantities, e.g. heat diffusion
- Furthermore:
 - Together with enhanced sampling techniques,
 - e.g. replica exchange, constrained dynamics (Blue-Moon ensemble), transition path sampling, transition interface sampling, forward flux sampling...
 - measures of free energy differences
 - -> phase diagrams, reaction rates`

Calculating vibrations via MD (Ex. 4)

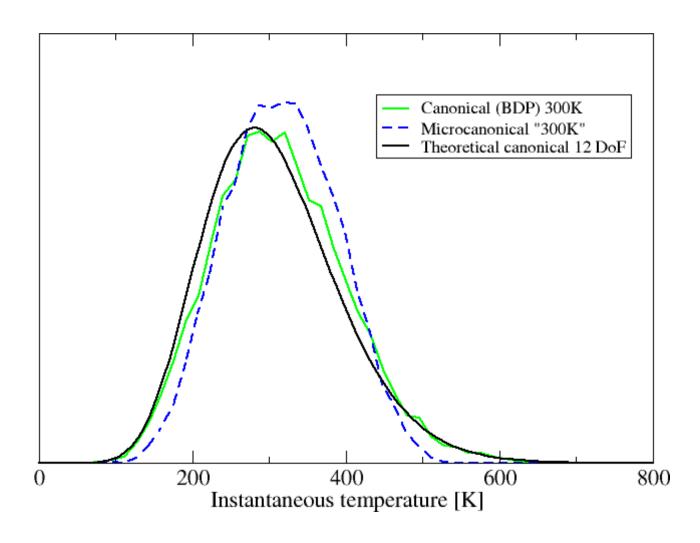
- Time autocorrelation functions can give information about vibrations
 - From Fermi's golden rule, the dipole time auto correlation function gives the intensities of IR active frequencies

$$I(\omega) = \int_{-\infty}^{\infty} dt \, e^{i\omega t} \langle \vec{M}(t) \cdot \vec{M}(0) \rangle_t$$

- Velocity time autocorrelation gives all frequencies of vibration. $\mathrm{VDOS}(\omega) = \sum_{i=1}^{N} \int_{-\infty}^{\infty} dt \, \mathrm{e}^{i\omega t} \langle \vec{v}_i(t) \cdot \vec{v}_i(0) \rangle_t$
- Possible to assign to individual atoms displacements and project on eigenmodes.

Calculating vibrations via MD (Ex. 4)

Canonical versus microcanonical sampling



Calculating vibrations via MD (Ex. 4)

Advantages:

- It doesn't rely on the harmonic approximation
- Take into account anharmonic effects present in experiments
- Simulate at different temperatures

Challenges:

- Needs a lot of statistics difficult for big systems, especially if ab initio
- Nuclei are classical, but in reality quantum effects for light nuclei are needed – ongoing work in the community

Excess free energy (Ex. 5)

Thermodynamic-path integration.

The absolute free energy cannot be calculated in a finite time MD.

But, the difference in free energy from a reference system, or condition of the system (e.g. very low *T*) can be often evaluated.

For example, by integrating: $\frac{\partial (\beta \mathcal{F})}{\partial \beta} = \langle \mathcal{U} \rangle_{NVT}$

Pay attention to possible hysteresis along the integration path!

Exercises overview

- Exercise 1- NVE ensemble
 - → Investigate 2 different self consistency accuracy settings



14:30 - 14:50 (20 min)

- Exercise 2 NVE ensemble
 - Velocity verlet with 1fs and 3fs time step
 - Ethane vs. Heavy Ethane



14:50 - 15:10 (20 min)

- Exercise 3 NVT ensemble
 - → Investigate 4 thermostats: Berendsen, Andersen and Nosé-Hoover, Bussi-Donadio-Parrinello



15:10 - 15:40 (30 min)

- Exercise 4 Application: CD₃CD₃
 - Harmonic vibrations
 - Anharmonic vibrations: dipole and velocity autocorrelations
 - → Temperatures: 600K first 2 rows, 450K next 2 rows, 300K last 2 rows.



15:40 - 17:00 (80 min)

Exercises overview

- Exercise 5 Evaluating (excess) free energy.
 - → Parallelizing over groups, each group calculates the average energy at a given temperature
 - Postprocessing done by one of us (possibly on the real data calculated by the participants ... and on-air)

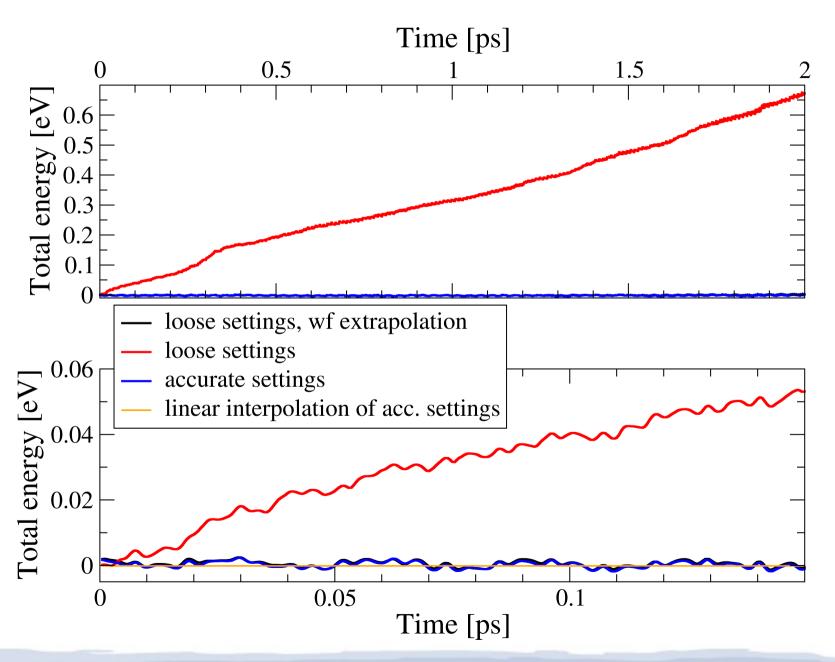


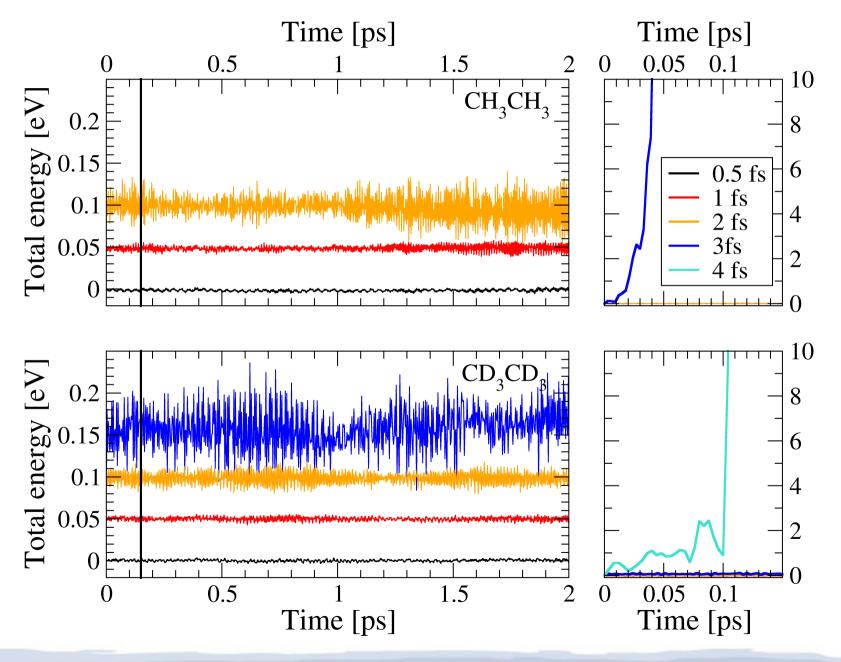
17:00 - 17:30 (30 min)

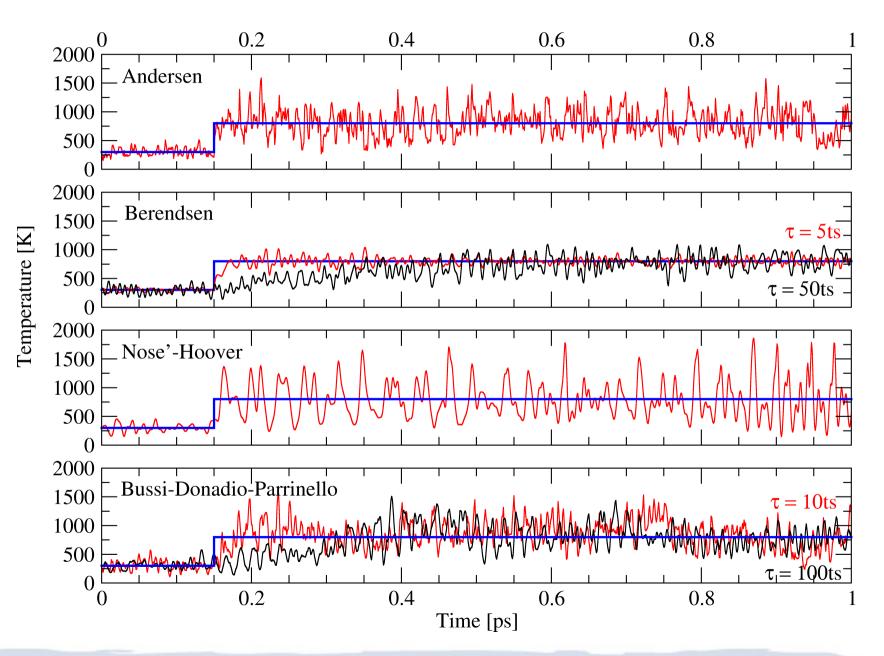
- Small project –Estimation of Heat flux
 - Mimicking the "Laser-Flash Measurements": ab initio heat conduction of a 1D (infinite) -(CD₂)-chain
 - Introduction on phonons (provided)
 - Generation of non equilibrium conditions



Setting
17:30 – 18:00
Calculations
over night
Discussion
tomorrow at ...

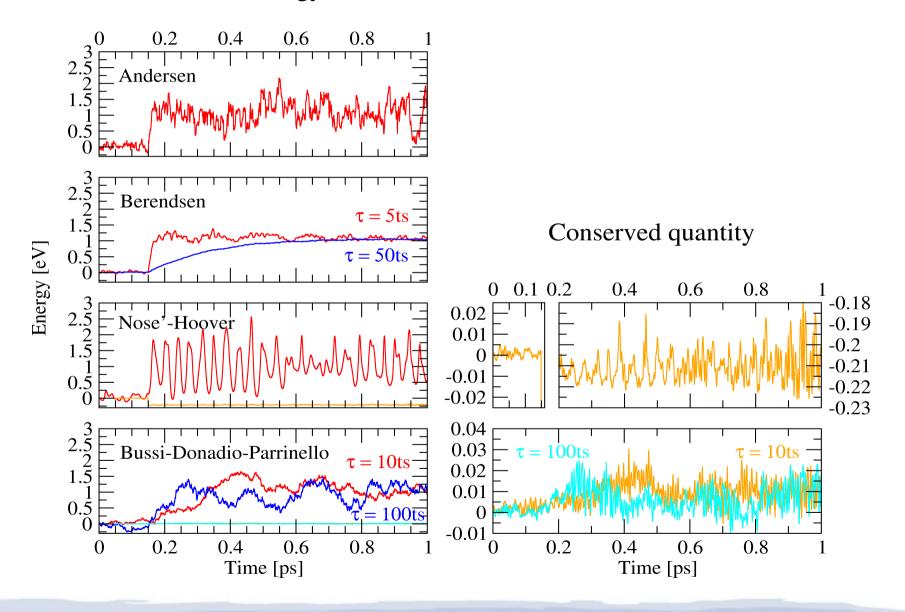


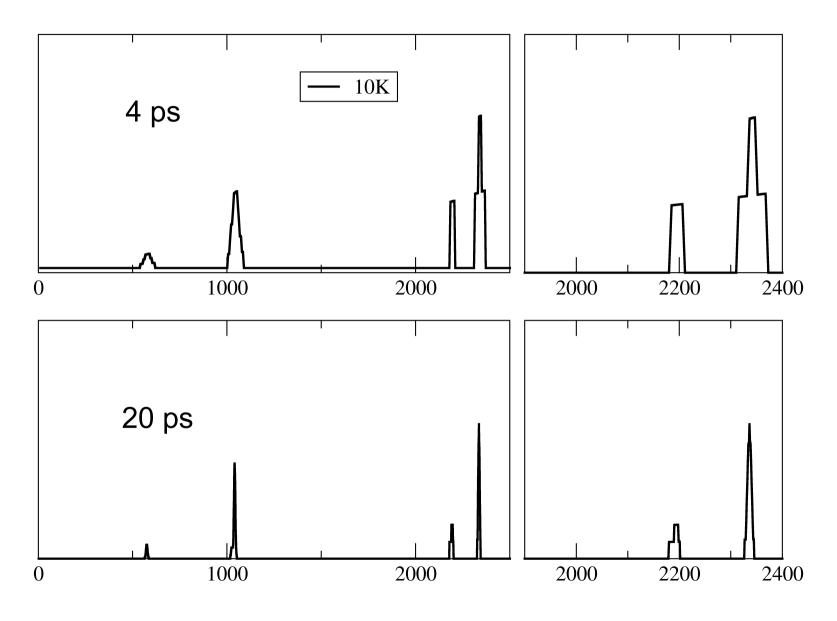


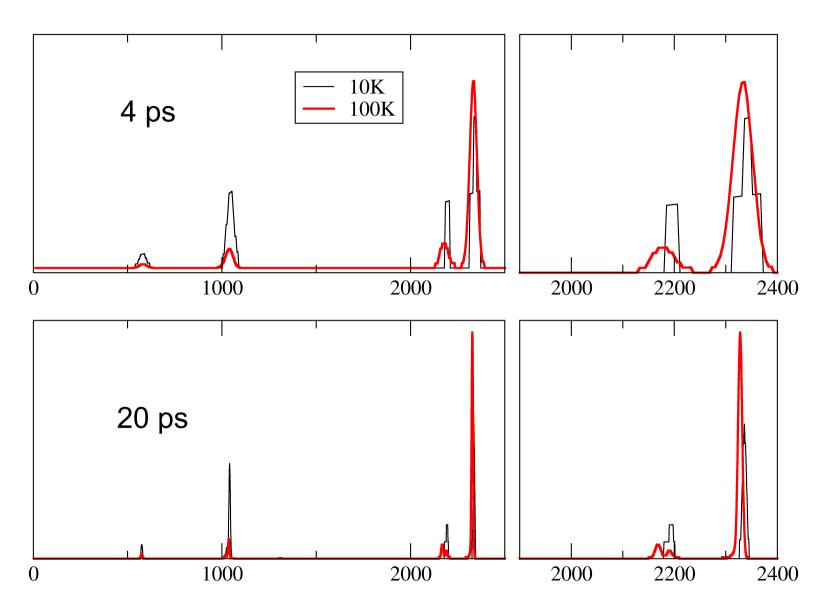


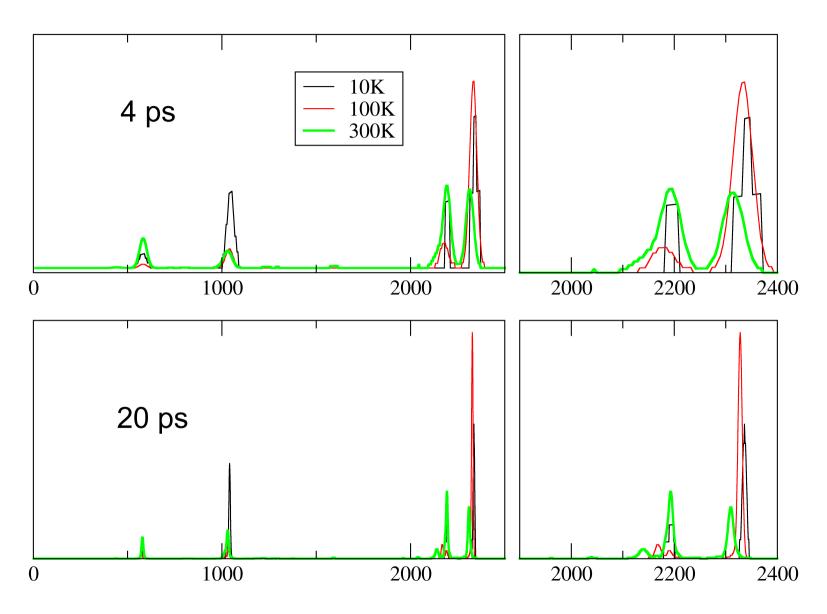
Comments on exercise 3 (ctd.)

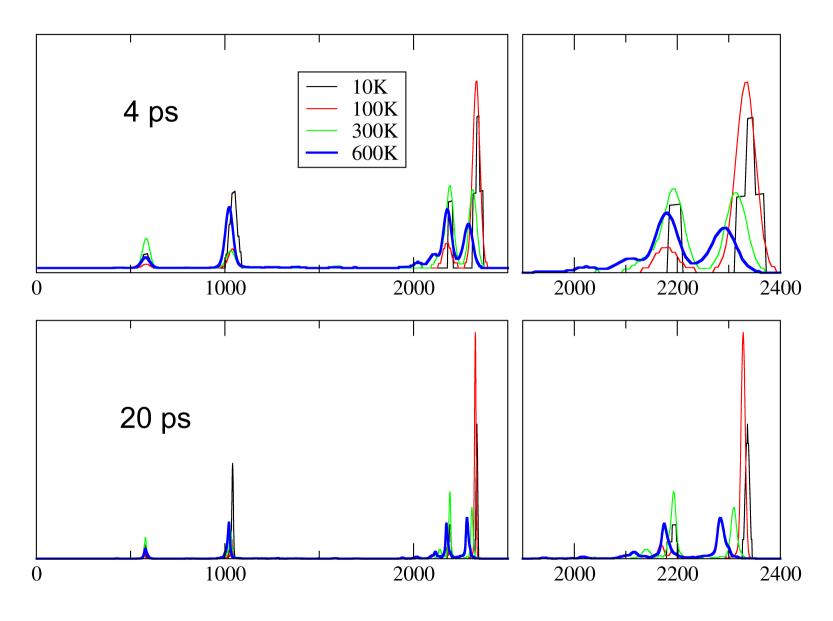
Total Energy

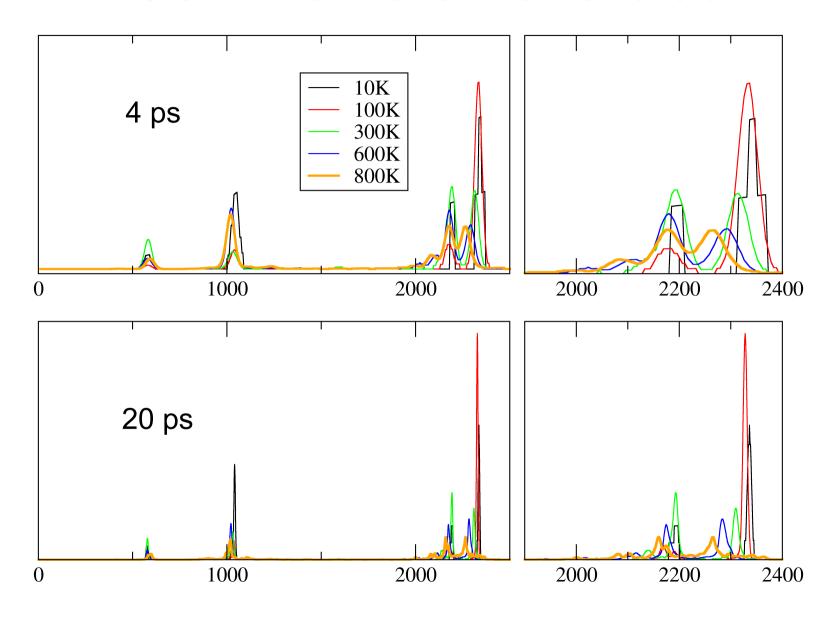


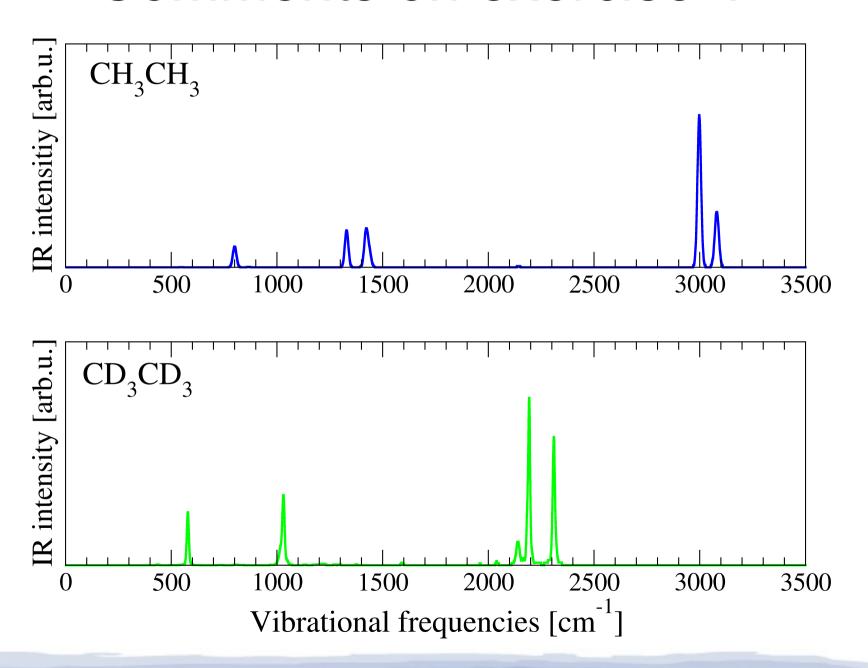


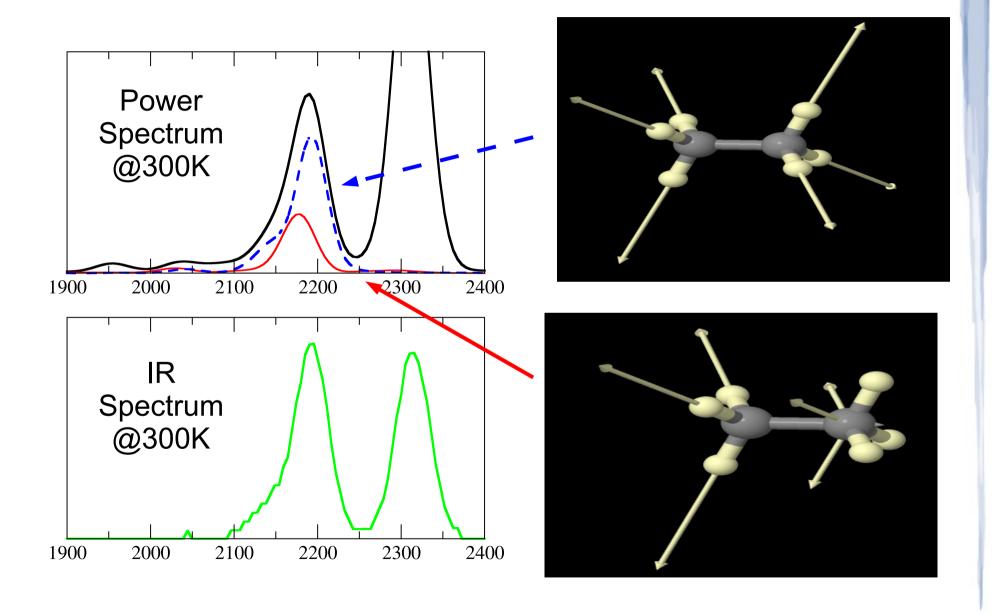












Comparison of accurate-force settings vs loose-force settings with wfe (see Ex.1), in the microcanonical ensemble.

