

# Hands on Session I:

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## Overview

- discussion of the required files
- lot's of examples
  - O atom
  - O<sub>2</sub> dimer
  - CO
  - H<sub>2</sub>O
- tasks
  - relaxation
  - vibrational frequencies
  - MD

## *The very first step: a single atom*

- files required to do all calculations presented in this session can be found in

~vw/1\_1\_description\_of\_job1

~vw/1\_2\_description\_of\_job2

first digit corresponds to the number of the hands on session, second one to the example

- required files INCAR, KPOINTS, POSCAR, POTCAR
  - POTCAR pseudopotential file  
generated by concatenation of individual POTCAR files from the data-base
  - KPOINTS Brillouin zone sampling  
describes which k-points are used
  - POSCAR structural data  
basis vectors and positions
  - INCAR steering the calculations

## *The POTCAR file*

- in this course, you can copy the POTCAR files from

```
~vw/potpaw_PBE/element_name/POTCAR
```

```
~vw/potpaw_PBE/0/POTCAR
```

or simply type

```
makepaw_PBE 0
```

or copy all input files from `~vw/1_1_0atom`

```
mkdir 0atom ; cd 0atom ; cp ~vw/1_1_0atom/* .
```

- what information can be found in the POTCAR file:
  - pseudopotential description
  - **data that is required to regenerate the potential**
  - number of valence electrons
  - atomic mass
  - required energy cutoff

## *The KPOINTS file*

- determines how many k-points are used to sample the Brillouin zone
- for molecules or atoms **only a single k-point is required**

KPOINTS file:

Gamma-point only

```
1          ! one k-point
```

```
rec        ! in units of the reciprocal lattice vector
```

```
0 0 0 1    ! 3 coordinates and weight
```

- for atoms and molecules the Bloch theorem does not apply, hence there is no need to use more than one single k-point  
when more k-points are used, only the interaction between the atoms (which should be zero) is described more accurately

## *The POSCAR and INCAR files*

- determines the lattice vectors (Bravais lattice) and the coordinates (position of the atoms)
- a single atom POSCAR file:

```
0 atom in a box
```

```
1.0          ! universal scaling parameters
```

```
8.0 0.0 0.0  ! lattice vector  a(1)
```

```
0.0 8.0 0.0  ! lattice vector  a(2)
```

```
0.0 0.0 8.0  ! lattice vector  a(3)
```

```
1           ! number of atoms
```

```
cart       ! positions in cartesian coordinates
```

```
0 0 0
```

- INCAR steers the calculations:

```
SYSTEM = 0 atom in a box
```

```
ISMEAR = 0
```

## Running vasp

type:

vasp

vasp.4.6.2 07Jul02

POSCAR found : 1 types and 1 ions

LDA part: xc-table for Pade appr. of Perdew

POSCAR, INCAR and KPOINTS ok, starting setup

WARNING: wrap around errors must be expected

entering main loop

	N	E	dE	d eps	ncg	rms	rms(c)
DAV:	1	0.39156E+02	0.39157E+02	-0.95953E+02	14	0.335E+0	
DAV:	2	0.39499E+01	-0.35207E+02	-0.34385E+02	28	0.480E+0	
DAV:	3	-0.15830E+00	-0.41082E+01	-0.39042E+01	14	0.376E+0	
DAV:	4	-0.31026E+00	-0.15195E+00	-0.13836E+00	14	0.660E+0	
DAV:	5	-0.31321E+00	-0.29544E-02	-0.29502E-02	28	0.907E-0	0.286E-01
DAV:	6	-0.31407E+00	-0.86398E-03	-0.18767E-03	14	0.397E-0	0.142E-01
DAV:	7	-0.31422E+00	-0.14198E-03	-0.21862E-04	14	0.149E-0	0.480E-02
DAV:	8	-0.31427E+00	-0.55085E-04	-0.26453E-05	14	0.469E-0	

1 F= -.31427624E+00 E0= -.16001392E+00 d E =-.308525E+00

writing wavefunctions

## OSZICAR *and* stdout file

initial charge corresponds to the charge of isolated overlapping atoms (POTCAR)  
for 4 steps the charge remains fixed, then the charge is updated (rms(c) column)

N	iteration count
E	total energy
dE	change of total energy
d eps	change of the eigenvalues (fixed potential)
ncg	number of optimisation steps $\mathbf{H}\psi$
rms	total residual vector $\sum_{nk} w_k f_{nk} (\mathbf{H} - \epsilon_{nk}) \psi_{nk}$
rms(c)	charge density residual vector



## OUTCAR *file*

individual parts are separated by lines

---

- first part: reading INCAR, POTCAR, POSCAR
- nearest neighbor distances and analysis of symmetry
- information on what was parsed from INCAR
- verbose job information
- information on lattice, k-points and positions
- information on the basis set (number of plane waves)
- non local pseudopotential information
- information for each electronic step (one line in OSZICAR)

```
POTLOK:  VPU time    0.93: CPU time    0.93
SETDIJ:  VPU time    0.01: CPU time    0.01
EDDAV :  VPU time    0.82: CPU time    0.83
DOS   :  VPU time    0.00: CPU time    0.00
```

```
-----
LOOP:   VPU time    1.76: CPU time    1.76
```

```
eigenvalue-minimisations :    14
total energy-change (2. order) : 0.3915659E+02 (-0.9595269E+02)
number of electron      6.0000000 magnetization
augmentation part      6.0000000 magnetization
```

Free energy of the ion-electron system (eV)

```
-----
alpha Z          PSCENC =          0.27135287
Ewald energy     TEWEN  =          -91.92708002
-1/2 Hartree    DENC   =         -281.84385690
-V(xc)+E(xc)    XCENC  =           26.11949869
PAW double counting =       245.99840262      -247.84808825
entropy T*S     EENTRO =          -0.18330906
eigenvalues     EBANDS =          -43.69352752
atomic energy   EATOM  =          432.26319604
```

```
-----
free energy     TOTEN  =          39.15658846 eV
```

```
energy without entropy =          39.33989752  energy(sigma->0) =          39.24824299
```

- eigenvalues

```
k-point 1 :      0.0000      0.0000      0.0000
band No.  band energies      occupation
   1      -23.8345      2.00000
   2       -8.8952      1.33333
   3       -8.8952      1.33333
   4       -8.8952      1.33333
   5       -0.4860      0.00000
   6        1.8485      0.00000
   7        1.8486      0.00000
```

- information on charge + some more timing informations

```
soft charge-density along one line, spin component      1
      0      1      2      3      4      5      6      7      8
x      5.4925      5.1765      4.3721      3.3741      2.4214      1.6318      1.0341      0.6112      0.3290
```

- information on the energy and stress tensor

FREE ENERGIE OF THE ION-ELECTRON SYSTEM (eV)

-----  
 free energy TOTEN = -0.314276 eV

energy without entropy= -0.005752 energy(sigma->0) = -0.160014

FORCE on cell =-STRESS in cart. coord. units (eV/reduce length):

Direction X Y Z XY YZ ZX

-----  
 Alpha Z 0.27 0.27 0.27  
 Ewald -30.64 -30.64 -30.64 0.00 0.00 0.00  
 Hartree 93.89 93.89 93.89 0.00 0.00 0.00  
 E(xc) -27.94 -27.94 -27.94 0.00 0.00 0.00  
 Local -147.85 -147.85 -147.85 0.00 0.00 0.00  
 n-local -20.54 -20.54 -20.54 0.00 0.00 0.00  
 augment 5.55 5.55 5.55 0.00 0.00 0.00  
 Kinetic 126.50 126.50 126.50 0.00 0.00 0.00

-----  
 Total -0.77 -0.77 -0.77 0.00 0.00 0.00

in kB -2.41 -2.41 -2.41 0.00 0.00 0.00

external pressure = -2.41 kB Pullay stress = 0.00 kB

- final timing information

## *Some comments on this particular run*

- the relevant energy for molecules and atoms is `energy without entropy`

```
energy without entropy= -0.005752 energy(sigma->0) = -0.160014
```

three degenerate p orbitals are occupied by 2/3 electrons causing a unphysical electronic entropy

```
entropy T*S EENTRO = -0.30852464
```

- a tiny value for SIGMA=0.01 would reduce the entropy but might slow convergence (default is SIGMA=0.2)

SIGMA controls the electronic temperature, which is not a very meaningful quantity for molecules and atoms

- the total energy is found to be essentially zero

VASP subtracts from any calculated energy the energy of the atom in the configuration for which the pseudopotential was generated

all pseudopotentials were generated using non spin polarized reference atoms

## *Restart vasp in same directory*

vasp.4.6.2 07Jul02

POSCAR found : 1 types and 1 ions

LDA part: xc-table for Pade appr. of Perdew

found WAVECAR, reading the header

POSCAR, INCAR and KPOINTS ok, starting setup

WARNING: wrap around errors must be expected

the WAVECAR file was read successfully

initial charge from wavefunction

entering main loop

	N	E	dE	d eps	ncg	rms	rms(c)
DAV:	1	-0.314277E+00	-0.31428E+00	-0.14320E-06	14	0.899E-03	0.157E-03
DAV:	2	-0.314277E+00	0.64237E-07	-0.19000E-07	7	0.353E-03	
1 F=		-.31427731E+00	E0=	-.16001499E+00	d E	=-.308525E+00	

writing wavefunctions

when vasp is restarted the **WAVECAR** file is read and the run is continued from the previous wavefunctions (converging rapidly)

## Spin polarized calculation

- the O atom is an open shell system with 2 unpaired electrons
- add ISPIN=2 to the INCAR file remove WAVECAR and restart vasp  
(alternatively copy all input files from `~vw/1_2_Oatomspin`)

```
vasp.4.6.2 07Jul02
```

```
POSCAR found : 1 types and 1 ions
```

```
...
```

```
entering main loop
```

	N	E	dE	d eps	ncg	rms	rms(c)
DAV:	1	0.38975372E+02	0.38975E+02	-0.10098E+03	32	0.259E+02	
DAV:	2	0.31791299E+01	-0.35796E+02	-0.35789E+02	64	0.438E+01	
DAV:	3	-0.11905610E+01	-0.43697E+01	-0.36660E+01	32	0.327E+01	
DAV:	4	-0.12616637E+01	-0.71103E-01	-0.69167E-01	32	0.508E+00	
DAV:	5	-0.12625234E+01	-0.85968E-03	-0.85961E-03	48	0.504E-01	0.653E+00

```
...
```

```
DAV: 11 -0.16719490E+01 0.16543E-04 -0.47746E-04 32 0.131E-01  
1 F= -.16719490E+01 E0= -.15948179E+01 d E =-.154262E+00 mag= 1.9986
```

```
writing wavefunctions
```

## Spin polarized calculation

- eigenstates for spin up and spin down are calculated “separately”  
in LSDA they interact only via the effective local potential  
spin-up and spin-down potential
- in the OUTCAR file, one can see two spin components
- the spin component 1 has 2 more electrons corresponding the a magnetization of  $2 \mu_B$

```
k-point 1 :      0.0000      0.0000      0.0000
band No.  band energies      occupation
   1      -25.0761      1.00000
   2      -10.0715      1.00000
   3      -10.0715      1.00000
   4      -10.0715      1.00000
   5       -0.3997      0.00000
   6       1.6965      0.00000
   7       1.9499      0.00000
   8       1.9499      0.00000
```

```
spin component 2
k-point 1 :      0.0000      0.0000      0.0000
band No.  band energies      occupation
   1      -21.8260      1.00000
   2       -7.0425      0.33333
   3       -7.0425      0.33333
   4       -7.0425      0.33333
   5       -0.4479      0.00000
   6       1.9043      0.00000
   7       1.9043      0.00000
   8       1.9043      0.00000
```



## Symmetry broken O atom

- in the GGA, most atoms are characterized by a symmetry broken solution  
VASP however symmetrizes the charge-density according to the determined symmetry of the cell  
check the OUTCAR file, which symmetry is VASP using
- to lower the symmetry simply change the lattice parameters to 7.0 8.0 and 9.0 in the POSCAR file (alternatively copy all input files from `~vw/1_3_Oatomspinlow`):

```
7.0 0.0 0.0 ! lattice vector a(1)
0.0 7.5 0.0 ! lattice vector a(2)
0.0 0.0 8.0 ! lattice vector a(3)
```

and reduce SIGMA to SIGMA=0.01 (INCAR file)

- rerunning VASP you will find a much lower energy

```
vasp.4.6.2 07Jul02
...
DAV: 17 -0.190131780202E+01 -0.53122E-04 -0.35327E-06 32 0.128E-02
1 F= -.19013178E+01 E0= -.19013178E+01 d E =-.938468E-52 mag= 1.9997
```

## *Let's add another atom: the O<sub>2</sub> dimer*

- copy the required files and start VASP (see footnote)

- POSCAR:

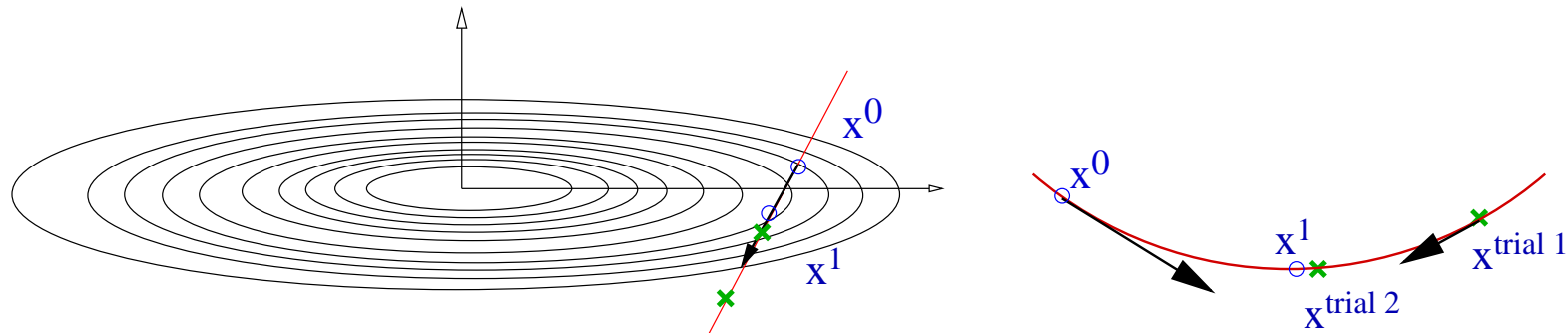
```
0 atom in a box
1.0          ! universal scaling parameters
8.0 0.0 0.0  ! lattice vector a(1)
0.0 8.0 0.0  ! lattice vector a(2)
0.0 0.0 8.0  ! lattice vector a(3)
2           ! number of atoms
cart       ! positions in cartesian coordinates
0 0 0      ! first atom
0 0 1.22   ! second atom
```

- INCAR:

```
SYSTEM = O2 dimer in a box
ISMEAR = 0 ! Gaussian smearing
ISPIN  = 2 ! spin polarized calculation
NSW    = 5  ! 5 ionic steps
IBRION = 2 ! use the conjugate gradient algorithm
```

## Relaxing the $O_2$ dimer

- we have inserted that geometry relaxation should be performed:  
in this case 5 ionic steps (NSW = 5) should be done at most  
for the relaxation a conjugate gradient algorithm is used IBRION = 2
- CG requires a line minimizations along the search direction



this is done using a variant of Brent's algorithm

- trial step along search direction (gradient scaled by POTIM)
- quadratic or cubic interpolation using energies and forces at  $\vec{x}_0$  and  $\vec{x}_1$  allows to determine the approximate minimum
- continue minimization, if app. minimum is not accurate enough

## Relaxing the O<sub>2</sub> dimer

```
DAV:   1      0.511277926223E+02    0.51128E+02   -0.31305E+03    60    0.528E+02
...
DAV:  11     -0.985454093746E+01   -0.88826E-04   -0.59757E-05    44    0.746E-02
  1 F= -.98545409E+01 E0= -.98545409E+01 d E =-.985454E+01 mag=      2.0000
curvature:   0.00 expect dE= 0.000E+00 dE for cont linesearch 0.000E+00
trial: gam= 0.00000 g(F)= 0.111E+00 g(S)= 0.000E+00 ort = 0.000E+00 (trialstep = 0.100E+01)
search vector abs. value= 0.111E+00
bond charge predicted
...
  2 F= -.96306943E+01 E0= -.96306943E+01 d E =0.223847E+00 mag=      2.0000
trial-energy change:   0.223847  1 .order   0.186756   -0.110518   0.484030
step:  0.1406(harm= 0.1859) dis= 0.00726 next Energy=   -9.862210 (dE=-0.767E-02)
bond charge predicted
...
  3 F= -.98624278E+01 E0= -.98624278E+01 d E =-.788682E-02 mag=      2.0000
curvature:  -0.09 expect dE=-0.448E-05 dE for cont linesearch -0.448E-05
trial: gam= 0.00000 g(F)= 0.484E-04 g(S)= 0.000E+00 ort =-0.231E-02 (trialstep = 0.828E+00)
search vector abs. value= 0.484E-04
reached required accuracy - stopping structural energy minimisation
```

## *CG: What does all this mean?*

- the quantity `trial-energy` change is the change of the energy in the trial step
- the first value after `1.order` is the expected energy change calculated from the forces  $((\mathbf{F}(\text{start}) + \mathbf{F}(\text{trial}))/2 \times \text{change of positions})$

**central difference**

second and third value corresponds to  $\mathbf{F}(\text{start}) \times \text{change of positions}$  and  $\mathbf{F}(\text{trial}) \times \text{change of positions}$

- the value `step`: is the estimated size of the step leading to a line minimization along the current search direction

**harm** is the optimal step using a second order (or harmonic) interpolation

- the trial step size can be controlled by the parameter `POTIM`  
the value `step`: times the present `POTIM` is usually optimal
- the final positions after the optimisation are stored in `CONTCAR`  
you can copy `CONTCAR` to `POSCAR` and continue the relaxation

## *Let's add another species: the CO molecules*

- copy required files and start VASP

- POSCAR:

```
...  
1 1          ! number of atoms for each species  
cart        ! positions in cartesian coordinates  
0 0 0       ! first atom  
0 0 1.12    ! second atom
```

- POTCAR is created by the concatenation of two individual POTCAR files corresponding to O and C; e.g.:

```
cat ~vw/potpaw_PBE/O/POTCAR ~vw/potpaw_PBE/C/POTCAR >POTCAR
```

- a similar relaxation as in the previous case is performed

but in this case more steps are required, since the first estimate for the minimum is not very accurate

the trial steps are much too long (POTIM parameter)

## Relaxing the CO dimer

```
1 F= -.14764188E+02 E0= -.14764188E+02 d E =-.147642E+02
curvature: 0.00 expect dE= 0.000E+00 dE for cont linesearch 0.000E+00
trial: gam= 0.00000 g(F)= 0.822E+00 g(S)= 0.000E+00 ort = 0.000E+00 (trialstep = 0.100E+01)
search vector abs. value= 0.822E+00
...
2 F= -.12657048E+02 E0= -.12657048E+02 d E =0.210714E+01
trial-energy change: 2.107140 1 .order 1.312507 -0.821770 3.446784
step: 0.1925(harm= 0.1925) dis= 0.02710 next Energy= -14.843291 (dE=-0.791E-01)
...
3 F= -.14747873E+02 E0= -.14747873E+02 d E =0.163154E-01
curvature: -0.10 expect dE=-0.909E-01 dE for cont linesearch -0.909E-01
ZBRENT: interpolating
opt : 0.0929 next Energy= -14.802370 (dE=-0.382E-01)
...
4 F= -.14797047E+02 E0= -.14797047E+02 d E =-.328587E-01
curvature: -0.04 expect dE=-0.341E-03 dE for cont linesearch -0.341E-03
trial: gam= 0.00000 g(F)= 0.844E-02 g(S)= 0.000E+00 ort =-0.833E-01 (trialstep = 0.819E+00)
search vector abs. value= 0.844E-02
reached required accuracy - stopping structural energy minimisation
```

## Vibrational frequencies of the CO dimer

- SYSTEM = CO dimer in a box  
ISMEAR = 0 ! Gaussian smearing  
IBRION = 5 ! vibrational spectrum  
NFREE = 2 ! use central differences  
POTIM = 0.02 ! 0.02 stepwidth  
NSW = 1 ! ionic steps must be larger 0 (that's all)

- POSCAR:

```
sel          ! selective degrees of freedom are changed
cart         ! positions in cartesian coordinates
 0 0 0       F F T ! first atom
 0 0 1.143   F F T ! second atom
```

the selected degrees of freedom are displaced once in the direction  $\hat{x}$  and once  $-\hat{x}$  by  $0.02 \text{ \AA}$

in the present case this makes 4 displacements plus the equilibrium positions (i.e. a total of five ionic configurations)



SECOND DERIVATIVES (NOT SYMMETRIZED)

```

-----
                1Z          2Z
1Z  -114.847733  114.847733
2Z   114.305971 -114.305971

```

Eigenvectors and eigenvalues of the dynamical matrix

```

-----
1 f =   63.876494 THz   401.347846 2PiTHz 2130.690412 cm-1   264.172038 meV
      X           Y           Z           dx           dy           dz
      0.000000  0.000000  0.000000           0           0   -0.655709
      0.000000  0.000000  1.143000           0           0    0.755014
2 f/i=   0.074763 THz    0.469753 2PiTHz   2.493841 cm-1    0.309197 meV
      X           Y           Z           dx           dy           dz
      0.000000  0.000000  0.000000           0           0   -0.755014
      0.000000  0.000000  1.143000           0           0   -0.655709

```

Eigenvectors after division by SQRT(mass)

Eigenvectors and eigenvalues of the dynamical matrix

```

-----
1 f =   63.876494 THz   401.347846 2PiTHz 2130.690412 cm-1   264.172038 meV
      X           Y           Z           dx           dy           dz
      0.000000  0.000000  0.000000           0           0   -0.163927
      0.000000  0.000000  1.143000           0           0    0.217854
2 f/i=   0.074763 THz    0.469753 2PiTHz   2.493841 cm-1    0.309197 meV
      X           Y           Z           dx           dy           dz

```

...

## *H<sub>2</sub>O molecules*

- POSCAR

```
H2O _2
0.52918 ! scaling parameter
 15 0 0
 0 15 0
 0 0 15
1 2
select
cart
      0.00      0.00      0.00 F F F
      1.10     -1.43      0.00 T T F
      1.10      1.43      0.00 T T F
```

all coordinates are scaled by the factor 0.529

- INCAR:

```
PREC = Normal      ! standard precision
ENMAX = 400        ! cutoff should be set manually
ISMEAR = 0 ; SIGMA = 0.1
IBRION = 1         ! use DIIS algorithm to converge
NFREE = 2          ! 2 independent degrees of freedom
NSW = 10           ! 10 ionic steps
EDIFFG = -0.02    ! forces smaller 0.02 A/eV
```

## *H<sub>2</sub>O molecules: comments*

- PREC = Normal should be used in vasp.4.6  
slightly more balanced setup than the default PREC = Medium
- I strongly urge to set the energy cutoffs manually in the INCAR file, as it gives you more control over the calculations
- for the ionic optimisation the DIIS algorithm is used  
this algorithm builds an approximation of the Hessian matrix and converges usually faster than the conjugate gradient algorithm  
it is however recommended to set the independent degrees of freedom manually  
EDIFFG determines when to terminate relaxation  
positive values: energy change between steps must be less than EDIFFG  
negative values:  $|\vec{F}_i| < |\text{EDIFFG}| \quad \forall i = 1, N_{ions}$

## *Interpreting the eigenstates of CO*

- the PROCAR file gives valuable information of the character of the one electron states
  - LORBIT 10 DOSCAR and l decomposed PROCAR file
  - LORBIT 11 DOSCAR and lm decomposed PROCAR file
- we use LORBIT=11 to distinguish  $p_x$  and  $p_z$  states
- copy the required input files, and check them using an editor  
execute vasp again

## PROCAR file:

band 3 # energy -11.46549527 # occ. 2.00000000

ion	s	py	pz	px	dxy	dyz	dz2	dxz	dx2	tot
1	0.000	0.546	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.546
2	0.000	0.157	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.157
tot	0.000	0.703	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.703

band 4 # energy -11.46549510 # occ. 2.00000000

ion	s	py	pz	px	dxy	dyz	dz2	dxz	dx2	tot
1	0.000	0.000	0.000	0.546	0.000	0.000	0.000	0.000	0.000	0.546
2	0.000	0.000	0.000	0.157	0.000	0.000	0.000	0.000	0.000	0.157
tot	0.000	0.000	0.000	0.703	0.000	0.000	0.000	0.000	0.000	0.703

band 5 # energy -8.76451122 # occ. 2.00000000

ion	s	py	pz	px	dxy	dyz	dz2	dxz	dx2	tot
1	0.001	0.000	0.135	0.000	0.000	0.000	0.000	0.000	0.000	0.136
2	0.172	0.000	0.261	0.000	0.000	0.000	0.000	0.000	0.000	0.433
tot	0.173	0.000	0.396	0.000	0.000	0.000	0.000	0.000	0.000	0.569

## *Let's do some a MD for H<sub>2</sub>O*

- **INCAR:**

```
PREC = Normal      ! standard precision
ENMAX = 400        ! cutoff should be set manually
ISMEAR = 0 ; SIGMA = 0.1
```

```
IBRION = 0         ! molecular dynamics
NSW = 100          ! 100 steps
POTIM = 1.0        ! timestep 1 fs
```

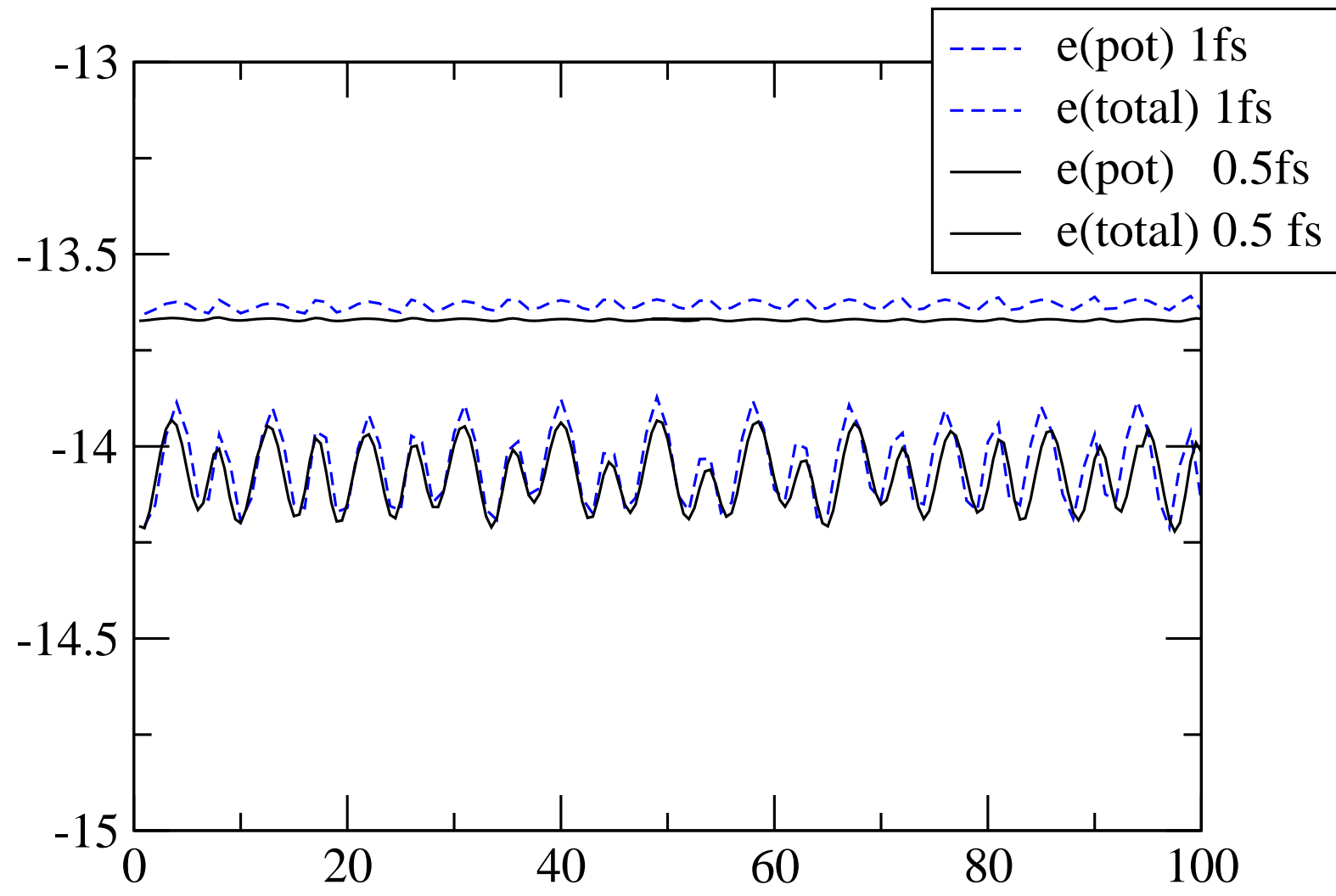
```
SMASS = -3        ! micro-canonical ensemble
TEBEG = 2000 ; TEEND = 2000 ! temperature
```

time step for this system should be around 0.5-0.7 fs

- **POSCAR:** to save time the box size is reduced to 12 a.u.

- **OSZICAR:**

```
1 T= 2134. E= -.13655511E+02 F= -.14207209E+02 E0=.. EK= 0.55170E+00 SP= 0.00E+00 SK= 0.00E+00
2 T= 1971. E= -.13643254E+02 F= -.14152912E+02 E0=.. EK= 0.50966E+00 SP= 0.00E+00 SK= 0.00E+00
3 T= 1336. E= -.13629241E+02 F= -.13974630E+02 E0=.. EK= 0.34539E+00 SP= 0.00E+00 SK= 0.00E+00
4 T= 1011. E= -.13624149E+02 F= -.13885486E+02 E0=.. EK= 0.26134E+00 SP= 0.00E+00 SK= 0.00E+00
5 T= 1307. E= -.13629772E+02 F= -.13967549E+02 E0=.. EK= 0.33778E+00 SP= 0.00E+00 SK= 0.00E+00
```



vi

## Excercises

- How does the energy change when you decrease SIGMA to 0.01 in the INCAR file starting from `~vw/1_1_0atom` ? Why ?
- Try to copy CONTCAR to POSCAR after running the example `~vw/1_4_0dimer`. Why is the calculation so fast ?
- Try to play with the parameter POTIM for the example `~vw/1_4_0dimer`. What is the optimal value ?
- What is the reason for the imaginary frequency in the example `~vw/1_6_C0vib`. Does the behavior improve when the step width (smaller or larger) is changed. Also try to improve the precision to which the groundstate is converged (`EDIFF=1E-5`). What happens if the accuracy of the calculations is improved (`PREC=Accurate`).
- Try to use the conjugate gradient algorithm to the H<sub>2</sub>O molecule (example `~vw/1_7_H2O`).
- Calculate the vibrational frequencies of the H<sub>2</sub>O molecule (example `~vw/1_7_H2O`) after relaxation (example `~vw/1_8_H2Ovib`). Why does one find 3 modes that have



small frequencies. EDIFF=1E-5 gives much improved results than EDIFF=1E-4, can you reproduce this behavior.

The following participants have to share one terminal:

Müller and Sahli (ETH Zürich)

Koza and Poehlmann (Univ. Montpellier)

Mok and Soon (Univ. Singapore)

Calatyyud and Mguig (Univ. P.and M. Curie)

possibly, if we encounter troubles:

Hobbs and Milazzo (Kings College, London)

Garcia-Vergniory and Rodriguez (Univ. Bilbao, Spain)

Cordente and Ricardo Chavez (Univ. Toulouse)