



#### MAX School on Advanced Materials and Molecular Modelling with QUANTUM ESPRESSO

# QE-2021: Hands-on session – Day-2

(Convergence-tests calculations)

















# QE-2021: Hands-on session – Day-2



Topics of Day-2 hands-on session:

- 1. How to make basic convergence tests (example1.Si/)
- 2. How to deal with metals (example2.Al/)
- 3. How to deal with ultrasoft pseudopotentials and with spin polarization (example3.Fe/)

To get the latest version of the exercises, move to Day-2/ directory and execute:

• git pull

# 1. Bulk system: Silicon



Self-consistent calculation (and a series of tests) for Silicon in the diamond structure:

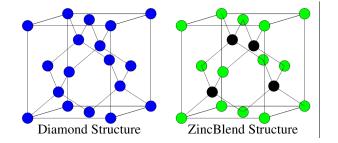
- move to Day-2/example1.Si/ directory
- look at the input file pw.si.scf.in. It is composed of three "namelists" &CONTROL (note that calculation = 'scf' is the default value), &SYSTEM, &ELECTRONS, followed by three "cards" ATOMIC\_SPECIES, ATOMIC\_POSITIONS, K\_POINTS
- in the &CONTROL namelist notice the following two variables (they are commented):
  - outdir: temporary directory for large files. Must be writable, will be created if not existent. You may set environment variable ESPRESSO\_TMPDIR instead.
  - pseudo\_dir: directory where pseudopotential (PP) files are kept. It must exist, be readable, and contain the required PP file (in this example, Si.pz-vbc.UPF for Silicon). You may set environment variable ESPRESSO\_PSEUDO instead.

(note that for the hands-on exercises we rely on ESPRESSO\_TMPDIR and ESPRESSO\_PSEUDO environmental variables, hence we don't need to set outdir and pseudo\_dir variables)

# Providing atomic structure in input



How is the crystal structure defined? This is a very simple case: the diamond lattice is an fcc (face-centered cubic) lattice with two atoms per unit cell. You need to specify:



- What is the Bravais lattice?ibrav=2, meaning fcc lattice
- How many and which parameters are needed to completely define Bravais lattice? just one: celldm(1)=10.2, lattice parameter a in a.u.
- How many atoms there are in the unit cell? nat=2: two atoms
- How many different atomic species are present?ntyp=1: one species
- Which ones, described by which pseudopotential?
   See card ATOMIC\_SPECIES
- Where the atoms are located in the unit cell? See card ATOMIC\_POSITIONS: here, in Cartesian axes, in units of a ("alat")

Notice that there are several alternative methods to specify an atomic structure!





**k-points** are described in the **K\_POINTS** card. One has to choose

Whether to provide a list of k-points or a uniform grid

• If a list is chosen: provide a list of k-points in the Irreducible BZ and corresponding symmetry weights; the latter do not need to add up to 1, they are normalized by the code

Frequently Asked Question: where do I find special k-points and their weights? Answer: 1) in papers, 2) use an auxiliary code kpoints.x, 3) use uniform grids

• If a uniform grid is chosen: specify Monkhorst-Pack parameters (*Phys. Rev. B* **13**, 5188 (1976)) and offsets along the three directions (uniform k-point grids are covered in more detail in section 1.2 below).

# Running the pw.x code



For serial (single processor) execution you can use

• pw.x -in pw.si.scf.in > pw.si.scf.out

(note: input redirection pw.x < pw.si.scf.in works but it is not recommended on parallel machines) Look at the directory specified by outdir (in our case \$ESPRESSO\_TMPDIR) and its content:

• 1s \$ESPRESSO\_TMPDIR silicon.save silicon.xml

(to see only these files, you may need to use ls \$ESPRESSO\_TMPDIR/silicon.\*)

The directory contains a data directory (silicon.save/) with binary data files for further processing and an XML file (silicon.xml) with general information on the run. The name of the various files is determined by the value of the prefix variable and by their content.

Do not run two instances of pw.x that access the same outdir with the same prefix! Unpredictable behavior may follow (the directory is used for temporary files as well). In case of trouble, clean outdir.





Examine output file pw.si.scf.out, look how self-consistency proceeds:

```
$ grep -e "total energy" -e estimated pw.si.scf.out
                                   -15.79103344 Ry
    total energy
    estimated scf accuracy
                                     0.06376674 Rv
    total energy
                              = -15.79409289 Ry
    estimated scf accuracy
                                     0.00230109 Rv
    total energy
                              = -15.79447822 Ry
                                     0.00006291 Ry
    estimated scf accuracy
                              = -15.79449510 Ry
    total energy
                                     0.00000448 Ry
    estimated scf accuracy
                              = -15.79449593 Ry
    total energy
    estimated scf accuracy <
                                     0.0000005 Ry
    The total energy is the sum of the following terms:
```

Notice that there are 8 electrons in the cell: 2 (pseudo-)atoms/cell with 4 electrons. The system is a non-magnetic insulator, so just the lowest 4 (= 8/2) valence bands (Kohn-Sham states) are computed.

# 1. Convergence tests for Si bulk



Convergence tests for Si bulk consist of the following steps:

- 1. convergence with respect to basis-set, i.e., kinetic energy cutoff (variable ecutwfc)
- 2. convergence with respect to k-points (card K\_POINTS)
- 3. with converged ecutwfc and k-points, determine the lattice parameter of Si bulk
- 4. Bonus: with converged parameters (ecutwfc, k-points, and lattice parameter), calculate a band structure of Si-bulk

# 1.1 Convergence w.r.t. the kinetic energy cutoff

The kinetic energy cutoff ecutwfc (in Ry) determines the size of the Plane-Wave (PW) basis set used to expand wave-functions (i.e. Kohn-Sham orbitals) (the default for the charge density is ecutrho=4\*ecutwfc, which is OK for norm-conserving PPs)

- A manual test of convergence w.r.t. kinetic energy cutoff entails the following tasks (BEWARE: we will not do it manually!)
  - 1. change ecutwfc in the pw.si.scf.in input to, e.g., 16, 20, 24, 28, 32 Ry
  - 2. for each value of ecutwfc, run pw.x and collect the final total energy
  - 3. collect the data in a file, say si.etot\_vs\_ecut (i.e. each line should contain two values: ecutwfc and "total-energy")
  - 4. plot the energies collected in si.etot\_vs\_ecut using your preferred plotting program, for instance:
    - gnuplot
      gnuplot> plot 'si.etot\_vs\_ecut' with lines
- because such a manual procedure is very cumbersome we use scripts instead

# 1.1 Convergence w.r.t. kinetic energy cutoff (II

To make convergence tests easier and faster, scripts are commonly used. To this end, Unix shell-scripts have been traditionally used, yet there are other more "fancy" alternatives, e.g., PWTK scripts.

- A Unix shell-script is located in ex1.ecutwfc.classic/sub-directory (file: ecutwfc.sh)
- A PWTK script is located in ex1.ecutwfc/ sub-directory (file: ecutwfc.pwtk)

```
Unix shell-script
#!/bin/sh
rm -f si.etot_vs_ecut.dat
for ecut in 12 16 20 24 28 32
   cat > pw.si.scf.$ecut.in << EOF
&CONTROL
   prefix='silicon',
 &SYSTEM
   ibrav = 2,
   celldm(1) = 10.2,
   nat = 2,
   ntyp = 1,
   ecutwfc = $ecut,
&ELECTRONS
ATOMIC_SPECIES
  Si 28.086 Si.pz-vbc.UPF
ATOMIC POSITIONS
  Si 0.00 0.00 0.00
  Si 0.25 0.25 0.25
K_POINTS automatic
 4 4 4 1 1 1
EOF
   pw.x -in pw.si.scf.$ecut.in > pw.si.scf.$ecut.out
   grep -e 'kinetic-energy cutoff' -e ! pw.si.scf.\ecut.out \
       awk '/kinetic-energy/ {ecut=$(NF-1)}
                            {print ecut, $(NF-1)}' >> si.etot_vs_ecut.dat
done
```

```
load_fromPWI ../pw.si.scf.in

set fid [open si.etot_vs_ecut.dat w]

foreach ecut {12 16 20 24 28 32} {
    SYSTEM "ecutwfc = $ecut"
    runPW pw.Si.scf.$ecut.in

    puts $fid "$ecut [::pwtk::pwo::totene pw.Si.scf.$ecut.out]"
} close $fid
```

# **PWTK** scripting: basics



The basic philosophy is to keep the syntax close to original input syntax!

```
pw.x input file
```

```
&CONTROL
   calculation = 'scf'
&SYSTEM
   ecutwfc = 25.0
   ecutrho = 200.0
ATOMIC POSITIONS alat
   Si
         0.00 0.00
                     0.00
   Si
         0.25 0.25 0.25
K POINTS automatic
           1 1 1
```

pw.Si.in

Run from terminal as:

```
pw.x -in pw.Si.in > pw.Si.out
```

# script

```
CONTROL {
   calculation = 'scf'
SYSTEM {
   ecutwfc = 25.0
   ecutrho = 8*25.0
ATOMIC POSITIONS alat {
   Si
         0.0
              0.0
                   0.0
         1/4 1/4
                   1/4
   Si
K POINTS automatic {
runPW pw.Si.in
```

Si bulk.pwtk

Run from terminal as:

```
pwtk Si bulk.pwtk
```

## **PWTK** scripting: basics



- PWTK scripts are basically Tcl-scripts, hence they use Tcl-syntax
- namelists and cards have the same names as in QE (with a few exceptions), but they are all written in **upper-case**. Their content is encapsulated in curly braces:

```
CONTROL { calculation = 'scf', outdir = '/tmp/pwscf/'}
ATOMIC_POSITIONS { ... }
```

instead of curly braces, one can also use double-quotes ("..."), e.g.:
 SYSTEM " celldm(1) = \$a "

- the difference between curly braces {...} and double-quotes "..." is that inside double-quotes the variable \$a is substituted by its value, whereas inside curly braces the \$a is treated literally (i.e. no substitution)
- real numbers can be specified as mathematical expressions (e.g. ecutrho = 8\*25.0)
- indices of <a href="mailto:ntyp-type">ntyp-type</a> array variables, such as <a href="mailto:starting\_magnetization">starting\_magnetization</a>(i), can be specified with atomic labels, e.g.:

```
SYSTEM { starting_magnetization(Fe) = -0.8 } where Fe is one among atomic species defined in the ATOMIC_SPECIES card.
```





- namelist variables can be set on the fly (we will use this heavily)
- the order of namelists and cards is not important; PWTK knows how to construct proper input files
- namelists and cards can be called many times, but there is a big difference how multiple calls are handled for namelists and cards: cards are handled in overwrite mode, whereas namelists are handled in a kind of append mode. For example, the following is OK:

```
CONTROL { calculation = 'scf' }
CONTROL { outdir = '/tmp/qe' }
and is equivalent to:
CONTROL { calculation = 'scf' , outdir = '/tmp/qe' }
```

• to unset a namelist variable, set it to an empty-value; for example to unset the outdir variable, use:

```
CONTROL { outdir = }
```





- load\_fromPWI loads input data from an existing pw.x input file
- pwo\_totene returns the converged total energy from pw.x output
- seq like the Unix seq command (returns a sequence of numbers)
- runPW constructs pw.x input file and runs a calculation
- runPP similar as runPW but for the pp.x program
- runDOS ... for the dos.x program
- runPROJWFC ... for the projwfc.x program
- PWTK web-site:

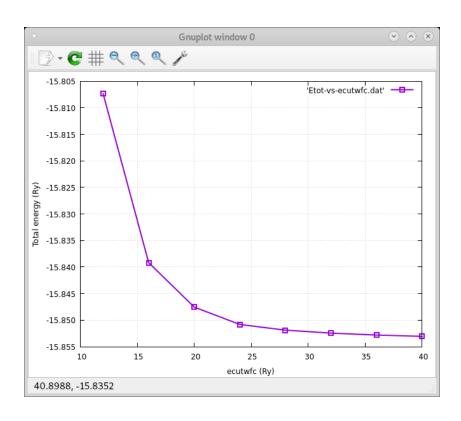
```
http://pwtk.ijs.si/ or http://pwtk.quantum-espresso.org/
```

PWTK documentation is available at:

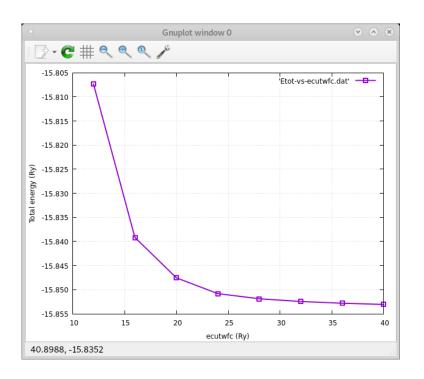
http://pwtk.ijs.si/documentation.html

# 1.1 Convergence w.r.t. the kinetic energy cutoff (III)

- To run the convergence test via the Unix shell-script move to ex1.ecutwfc.classic/sub-directory (read the README.md file) and execute
  - ./ecutwfc.sh
- To run the convergence test via the PWTK script move to ex1.ecutwfc/ sub-directory (read the README.md file) and execute
  - pwtk ecutwfc.pwtk



# 1.1 Convergence w.r.t. the kinetic energy cutoff (IV)



#### Notes:

- Convergence w.r.t. cutoff is a property of the *pseudopotential(s)* used.
- Convergence of the *absolute energy* is typically slower than convergence of *interesting physical properties*, e.g. structure.
- Absolute values of total energy do not have any physical meaning (and depend upon the specific pseudopotential): only energy differences have meaning

# 1.2 Convergence w.r.t. k-points



A sufficiently dense grid of k-points is needed in order to account for *periodicity*.

To test the convergence w.r.t. k-points, you need to edit the K\_POINTS card and request *automatic* Monkhorst-Pack grids:

```
K_POINTS automatic
nk1 nk2 nk3     k1 k2 k3
```

then step-wise increase nk1=nk2=nk3 to, e.g., 2, 4, 6, 8 (keep k1=k2=k3=1) and run pw.x calculation for each value of nk1=nk2=nk3.

For example, with PWTK this can be achieved with the following snippet:

```
load_fromPWI pw.si.scf.in

foreach k {2 4 6 8} {
    K_POINTS automatic "$k $k $k 1 1 1"
    runPW pw.si.scf.$k.in
}
```





Description of the K\_POINTS card for *automatic* mode:

```
K_POINTS automatic
nk1 nk2 nk3     k1 k2 k3
```

The first three nk1 nk2 nk3 numbers mean "there are nk1,nk2,nk3 grid points along crystal axis 1,2,3"; the second three k1 k2 k3 numbers, either 0 or 1, mean "grid starts from 0" or "displaced by half a step" along crystal axis 1,2,3

#### Also note that:

- Convergence is not necessarily monotonic: there is no variational principle w.r.t.
   number of k-points
- The 2 2 2 1 1 1 Monkhorst-Pack grid is the same as the "two Chadi-Cohen points" (see: D.J. Chadi and M.L. Cohen, Phys. Rev. B 8, 5747 (1973))



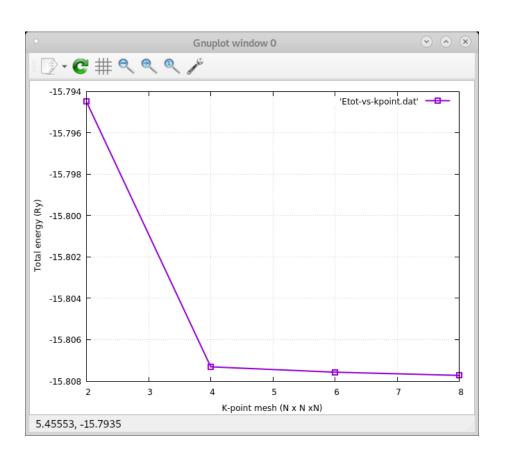


The PWTK script for testing the convergence with respect to k-points is located in Day-2/example1.Si/ex2.kpoints/ directory (see README.md for detailed instructions).

Within this directory execute:

• pwtk kpoints.pwtk

You should get a plot like this one:







Equilibrium in Si is determined by the minimum-energy lattice parameter alone: there are no forces on atoms due to symmetry (you can verify this by setting tprnfor=.true. in namelist &CONTROL and looking for forces reprinted at the end).

To find the lattice parameter:

- Choose suitable values for ecutwfc and k-point grid (e.g. 30 Ry and 4 4 4 1 1 1)
- Run pw.x for values of celldm(1) ranging from 9.7 to 10.7 in steps of 0.1 a.u.

With PWTK this can be achieved with the following snippet:

```
load_fromPWI pw.si.scf.in

foreach alat [seq 9.7 0.1 10.7] {
    SYSTEM "celldm(1) = $alat"
    runPW pw.si.scf.$alat.in
}
```





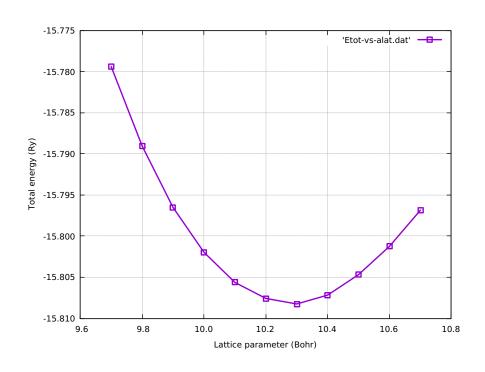
The corresponding PWTK script is located in Day-2/example1.Si/ex3.alat/directory (see README.md for detailed instructions).

Within this directory execute:

• pwtk alat.pwtk

The experimental lattice parameter for Si is 5.47 Å or 10.26 a.u.. This is a case where plain simple LDA yields remarkable results. You may experiment changing cutoff, k-points, pseudopotential, ...

You should find that:



- The energy vs lattice parameter E(a) curves are shifted down rather uniformly with increasing cutoff and are not strongly dependent on k-points.
- Structural properties and energy differences converge faster than total energies.





Use the code ev.x to fit your results to a phenomenological equation-of-state (EOS, e.g. Murnaghan) and to get accurate values for the lattice parameter and for the bulk modulus.

The ev.x code prompts for some data and reads a data-file like the one produced by the alat.pwtk script (the data-file is Etot-vs-alat.dat). For cubic systems a data-file should contain the following rows:

$$a_1$$
  $E(a_1)$ 
 $a_2$   $E(a_2)$ 
 $a_3$   $E(a_3)$ 

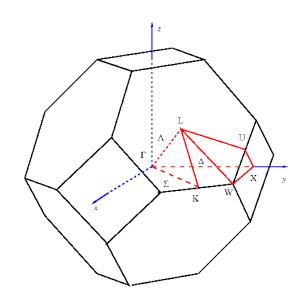
. . .

### 1.4 Band Structure of Silicon



The scheme to calculate the bands (spaghetti plot) is the following:

- 1. SCF pw.x calculation (calculation = 'scf')
- 2. "bands"-type non-SCF pw.x calculation (fixed-potential) with:
  - calculation = 'bands'
  - the number of Kohn-Sham states explicitly set (variable nbnd)
  - a suitable path of k-points specified in K\_POINTS (in this example we use the  $L-\Gamma-X-W-K-L$  path)



3. bands.x calculation, which, among others, produces data-files for the spaghetti plot

Important: outdir and prefix must be the same for "bands" and "scf" pw.x calculations and for the bands.x calculation
Important: the k-point path must be continuous in k-space





The input for the bands.x program is the following:

```
&BANDS

prefix='...', outdir='...', filband = 'Si.bands.dat', lsym=.true.
/
```

Two files are produced: Si.bands.dat.gnu, directly plottable with gnuplot, and Si.bands.dat, for further processing by the auxiliary command plotband.x.

If option lsym=.true., bands.x performs a symmetry analysis. An additional file Si.bands.dat.rep is generated, containing information on symmetry labels of the various bands.

The snippet for running all three calculations manually is:

- pw.x -in pw.Si.scf.in > pw.Si.scf.out
- pw.x -in pw.Si.bands.in > pw.Si.bands.out
- bands.x -in bands.Si.in > bands.Si.scf.out

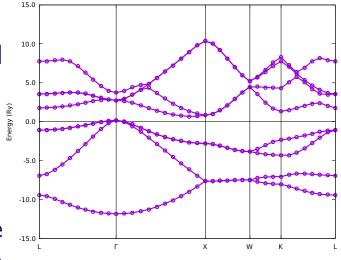
but we will use a PWTK script instead.

# 1.4 Band Structure of Silicon (III)



To execute the PWTK script that will perform all the needed calculations for plotting the bands, proceed as follows:

- move to directory
   Day-2/example1.Si/ex4.bands/ and read the README.md file
- set suitable values for celldm(1), ecutwfc, and K\_POINTS
- and execute: pwtk bands.pwtk
- you may set the Efermi value to the top of the occupied bands in the gnuplot file plot.gp (see the instructions in README.md); then re-plot spaghetti with: gnuplot plot.gp



Remark: in PWTK, once outdir and prefix are set, they are automatically inherited for subsequent calculations.

# **Auxiliary program plotband.x**



#### plotband.x prompts for terminal input:

```
• plotband.x
Input file > Si.bands.dat
Reading 8 bands at 39 k-points
Range: -5.6940 16.4680eV Emin, Emax > -5.6940 16.4680
high-symmetry point: -0.5000 0.5000 0.5000 x coordinate
                                                         0.0000
high-symmetry point: 0.0000 0.0000 0.0000 x coordinate
                                                         0.8660
high-symmetry point: 0.0000 0.0000 1.0000 x coordinate
                                                         1.8660
high-symmetry point: 0.0000 0.5000 1.0000 x coordinate
                                                         2.3660
high-symmetry point: 0.0000 0.7500 0.7500 x coordinate
                                                         2.7196
high-symmetry point: -0.5000 0.5000 0.5000
                                                         3.3320
                                          x coordinate
output file (gnuplot/xmgr) > Si.bands.plot
bands in gnuplot/xmgr format written to file Si.bands.plot
output file (ps) > (press Return)
```

If symmetry analysis was performed in the previous step, the output is written to several plottable files Si.bands.plot.N.M, where N labels the high-symmetry lines, M labels irreducible representations.

# 2. A metallic example: Aluminum



Aluminum is even simpler than Silicon: one atom per unit cell in an fcc lattice. **BUT:** it is a metal, only valence bands and a few k-points will not suffice.

- move to the Day-2/example2.Al/ directory
- read the pw.x input file pw.al.scf.in
- notice the presence of new variables: occupations, smearing, degauss;
- run pw.x as:
  - pw.x -in pw.al.scf.in > pw.al.scf.out
- in the output file notice that
  - the number of bands (Kohn-Sham states) is automatically set to a value larger than the number of electrons divided by 2
  - the Fermi energy is computed.

# 2.1 Convergence with respect to k-points, degauss, and smearing

This is a "three-dimensional" convergence test, where the number of k-points and values of degauss and smearing variables are varied. In particular, we will vary:

- smearing variable, possible values: 'gauss' (or 'g'), 'marzari-vanderbilt'
   (or 'm-v'), 'methfessel-paxton' (or 'm-p')
- degauss variable, in range from 0.003 to 0.1
- k-points using the *automatic* grids of 4 4 4, 8 8 8, 12 12 12, and 16 16 16

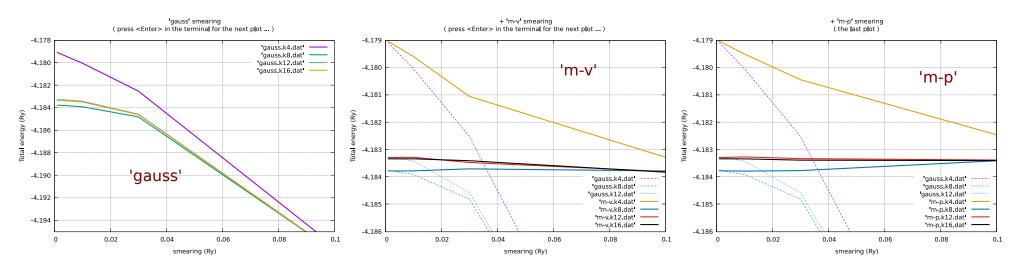
With PWTK this can be achieved with the following snippet:

# 2.1 Convergence with respect to k-points, degauss, and smearing

- move to Day-2/example2.Al/ex1.degauss/ directory
- execute: pwtk degauss.pwtk

Notice how much slower the convergence is for metals than for insulators!

Both m-v and m-p depend much less upon degauss and allow for faster and safer convergence than simple Gaussian broadening. For Al and m-v or m-p smearing, good convergence is achieved for a 12 12 k-point grid and degauss  $\sim 0.01$  to 0.05 Ry.



Beware that you cannot reduce the broadening too much: the energy levels must have some overlap, or else the advantage of broadening is lost!

# 2.2 How to plot charge-density



Example Day-2/example2.Al/ex2.chdens/ shows how to calculate the valence and the all electron charge density (the latter requires a PAW potential and a very large cutoff energy)

- move to Day-2/example2.Al/ex2.chdens/ directory (chdens is an acronym for charge-density)
- execute: pwtk 1-chdens.pwtk this script calculates and "plots" the valence charge density; notice that the electron charge is located mainly in interstitial regions (due to the use of a pseudo-potential, there is almost no charge in close vicinity of nuclei; see the next page)
- the scheme to calculate and plot the charge-density is:
  - 1. make an SCF pw.x calculation
  - 2. make a post-processing pp.x calculation (plot\_num=0 for charge density) and instruct the program to write charge density in a suitable format
  - 3. plot the charge density by xcrysden (let's plot density in contour/colorplane style; follow the instructions of tutor and select density range from 0.0 to 0.05)

# 2.2 How to plot charge-density (II)

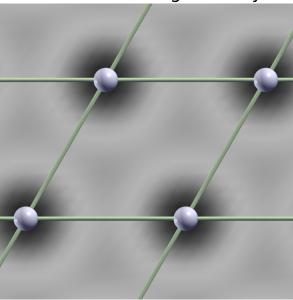


- to calculate all-electron **valence** and **total** charge densities, execute:
  - pwtk 2-chdens-paw.pwtk

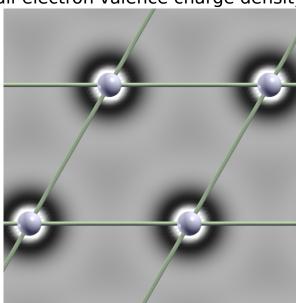
(note that plot\_num=17 for all-electron valence density and plot\_num=21 for all-electron total density)

• comparison between pseudopotential (PP) valence-density vs. all-electron densities (valence and total):

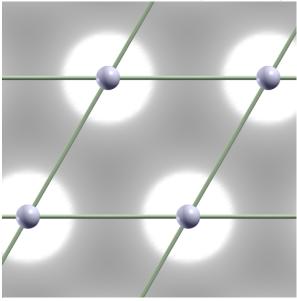
PP valence charge density



all-electron valence charge density



all-electron total charge density



# 3. A magnetic example: Iron bulk



Fe has two remarkable features: it is magnetic and it "calls for" an Ultrasoft PP (USPP) since its localized 3d atomic states are very hard.

- move to the Day-2/example3.Fe/ directory and read the pw.x input file pw.fe\_fm.scf.in
- the structure is bcc (ibrav=3) with one atom per unit cell
- notice (i) variables nspin and of starting\_magnetization, indicating spin-polarized calculation (nspin=2) with unconstrained total magnetization with only the initial guess for the magnetization given and (ii) variables for metallic system (occupations, smearing, degauss)
- notice that this calculation uses GGA (PBE): it is specified inside the PP file (can be guessed from the PP file name), reprinted on output as "Exchange-correlation"
- also notice that with USPP, it is typically needed to set ecutrho > 4\*ecutwfc (it should be at least 8 to 12 times larger)





Run pw.x in the usual way  $(pw.x - in pw.fe_fm.scf.in > pw.fe_fm.scf.out)$ . In the output, notice:

- the number of k-points is doubled w.r.t the non-magnetic case: the first set of k-points contains spin-up states, the second set spin-down states (use verbosity='high' in namelist &CONTROL if there are more than 100 k-points)
- in the output notice such lines:

```
total magnetization = 2.41 Bohr mag/cell absolute magnetization = 2.60 Bohr mag/cell
```

Since there is a single (magnetic) atom per unit cell, the only possible magnetic structure is ferromagnetic.

# 3. Magnetic structures: going antiferromagnetic

- To reach antiferromagnetic states, you need to:
  - introduce a supercell with two sublattices of different species of atoms (even if they are the same, it is important that they are labeled as different)
  - start with opposite initial magnetization for the two sublattices
- Can you write input data for an AFM structure? hint: split bcc into two simple cubic sublattices, ibrav=1, with two atoms at (0,0,0) and  $(\frac{1}{2},\frac{1}{2},\frac{1}{2})$ .

As a convenience, an antiferromagnetic file is provided (pw.fe\_afm.scf.in)

You can compare the ferromagnetic and antiferromagnetic files by:

• diff pw.fe\_fm.scf.in pw.fe\_afm.scf.in

# 3.1 Convergence tests for ultrasoft pseudopotential(s)

For computational efficiency, it is convenient to keep ecutwfc as low as possible, while ecutrho is less critical (look at the CPU time report at the end of an output: there are very many fftw, depending upon ecutwfc, while a much smaller number of fft depends upon ecutrho)

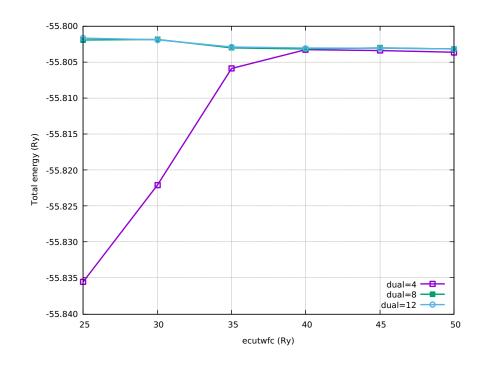
Set the ecutrho/ecutwfc ratio (dual) to 4, 8, 12 and compute the energy vs ecutwfc curve. For dual = 4 it will look funny: energy *increases* with increasing cutoff (see next page), but for a higher dual (i.e. better description of augmentation charge) the normal behavior is observed.

#### The corresponding PWTK snippet is:

# 3.1 Convergence tests for ultrasoft pseudopotential(s)

A PWTK script to make convergence check for USPP is provided, i.e.:

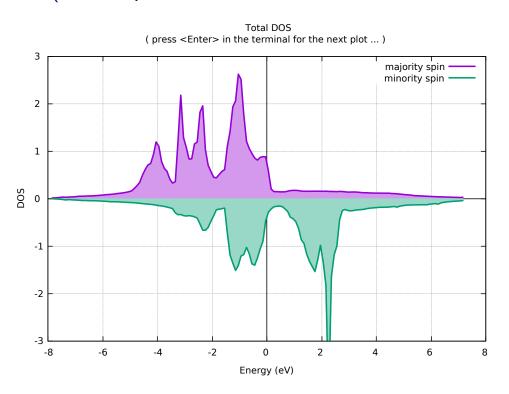
- move to directory:
   Day-2/example3.Fe/ex1.ecut/
- execute: pwtk ecut.pwtk
- you should obtain such a plot (notice that curves for dual=8 and 12 almost coincide)

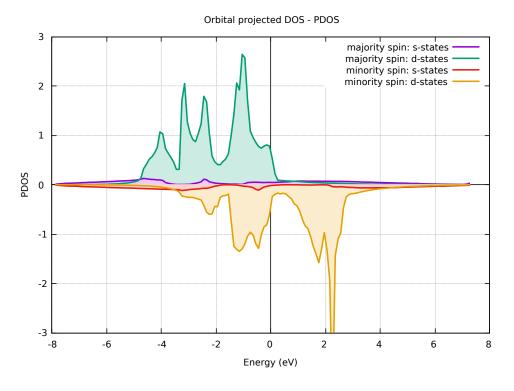


**Homework:** for converged values of both cutoffs and k-points, you may compare the stability of iron in the bcc, fcc, hcp phases (the latter being a slightly more complicated structure)

# 3.2 Density of States (DOS) and Projected DOS (PDOS)

- 1. move to Day-2/example3.Fe/ex2.dos/ directory and
- 2. edit the dos.pwtk script and set ecutwfc and ecutrho to appropriate values
- 3. execute: pwtk dos.pwtk (this script calculates both total DOS and DOS projected (PDOS) to atomic orbitals)





# 3.2 Density of States (DOS) and Projected DOS (PDOS)

The scheme to calculate DOS and PDOS consists of:

- 1. an SCF pw.x calculation (calculation='scf')
- 2. a non-SCF pw.x calculation (calculation='nscf'), where:
  - the same prefix and outdir are used as in the preceding SCF calculation
  - a denser k-point mesh is specified
  - in this example the *linear tetrahedron method* is used (variable occupations='tetrahedra')
- 3. a dos.x calculation to calculate DOS (DOS is written to a file as specified by the fildos variable in the dos.x input; also here the values of prefix and outdir are the same as for SCF pw.x calculation)
- 4. a projwfc.x calculation to calculate PDOS projected to atomic states (this calculation is analogous to dos.x one, inputs are also very similar; you can compare them as:
  - diff dos.Fe.in projwfc.Fe.in

the difference is that dos.x uses the fildos whereas projwfc.x uses the filpdos variable)

Explore the content of fildos and filpdos files!