



Workshop on High Performance Computing for Materials Characterization, Design and Discovery - SMR3966

Quantum ESPRESSO Hands-on session

(Basic SCF + post-processing + convergence test calculations)

Rafael González, Carlos Pinilla, Camilo Espejo (Based on MaX School tutorial - 2021 by Pietro Delugas, et. al.)





Super Computación y Cálculo Científico UIS









- 1. How to run basic PWscf (pw.x) calculations
- 2. How to run post-processing calculations to plot molecular-orbitals and chargedensity (pp.x), DOS (dos.x), and band-structure (bands.x)
- 3. How to calculate low-dimensional systems (example1.benzene/ and example2.graphene/)
- 4. How to make basic convergence tests (example3.Si/)
- 5. How to deal with metals (example4.Al/)

About Quantum ESPRESSO



More info about Quantum ESPRESSO can be found in:

- https://www.quantum-espresso.org/
- Quantum ESPRESSO (QE) documentation:
 - on-line manuals at www.quantum-espresso.org/resources/users-manual
 - Doc/ sub-directories in the QUANTUM ESPRESSO distribution
 - input data description: most programs contained in QE have their own input file description in the form of hyperlinked INPUT_***.html files (where *** stands for the name of the program)

Hands-on material



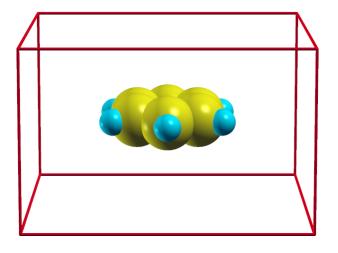
QE Hands-on examples material is contained within its own directory:

- example1.benzene/ hands-on exercise 1
- example2.graphene/ ... example5.Fe/
- All directories contain a **README.md** file with instructions how to run exercise(s)
- Naming of files is described in README-filenames.md
- To help recognizing for which program a given input file is intended, the filename starts with the name of the program, i.e.:
 - pw.*.in input file for pw.x program
 - pp.*.in input file for pp.x program
 - etc.

Disclaimer: *many examples use lousy convergence thresholds to speed-up calculations*

1. How to describe a molecule with Quantum ESPRESSO

With Quantum ESPRESSO we can describe a molecule by putting it in a big box.



- move to example1.benzene/ directory
- look at the input file pw.benzene.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 terminal execute more pw.benzene.scf.in
- instructions for how to run the example are in README.md

Disclaimer: the box used in this example is very small as to speed-up calculations

SMR3966: Workshop on High Performance Computing for Materials Characterization, Design and Discovery Example: example1.benzene/

1. How to calculate and plot molecular orbitals

Here are the two needed input files for calculation of signed molecular-orbital densities of benzene (i.e., $sign(\psi_i(\boldsymbol{r}))|\psi_i(\boldsymbol{r})|^2$), opened with emacs using QE-emacs-modes:

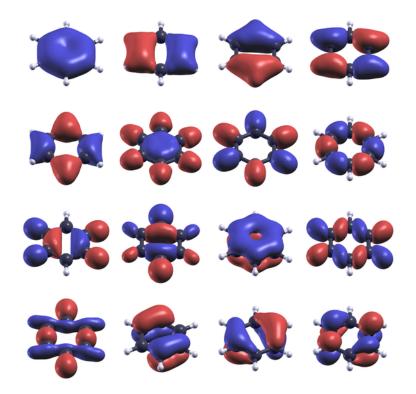
File Edit Options Buffers Tools Help &CONTROL prefix='benzene', / &SYSTEM ibrav = 6 A = 11.0 C = 7.0 nat = 12, ntyp = 2, nbnd = 16 ecutwfc = 20.0, ecutrho = 200.0, assume_isolated = 'martyna-tuckerman', / &ELECTRONS / ATOMIC_SPECIES C 1.0 C.pbe-rrkjus.UPF H 1.0 H.pbe-rrkjus.UPF H 1.0 H.pbe-rrkjus.UPF ATOMIC_POSITIONS angstrom H 5.5000000 7.98563953 3.5 C 5.5000000 7.98563953 3.5 C 6.7089386 6.19812524 3.5 H 7.6529918 6.74309454 3.5 C 6.7089386 4.80187470 3.5 H 7.6529918 4.25690561 3.5 C 5.5000000 4.10479662 3.5 H 3.3470081 4.25690556 3.5 C 4.2910612 4.80187468 3.5 H 3.3470082 6.74309458 3.5 K_POINTS gamma :: pw.benzene.scf.in All (18,2) (QE-pw.x) Beginning of buffer		emacs@cl	\odot \otimes \otimes
<pre>prefix='benzene', SSYSTEM ibrav = 6 A = 11.0 C = 7.0 nat = 12, ntyp = 2, nbnd = 16 ecutwfc = 20.0, ecutrho = 200.0, assume_isolated = 'martyna-tuckerman', SELECTRONS ATOMIC_SPECIES C 1.0 C.pbe-rrkjus.UPF H 1.0 H.pbe-rrkjus.UPF ATOMIC_POSITIONS angstrom H 5.5000000 7.98563953 3.5 C 5.5000000 6.89520922 3.5 C 6.7089386 6.19812524 3.5 H 7.6529918 4.25690561 3.5 C 5.5000000 4.10479062 3.5 H 7.6529918 4.25690561 3.5 C 5.5000000 3.01436043 3.5 C 4.2910612 4.80187468 3.5 H 3.3470081 4.25690556 3.5 C 4.2910613 6.19812528 3.5 H 3.3470082 6.74309458 3.5 K_POINTS gamma -: pw.benzene.scf.in All (18,2) (QE-pw.x)</pre>	File Edi	t Options Buffers Tools Help	
<pre>SSYSTEM ibrav = 6 A = 11.0 C = 7.0 nat = 12, ntyp = 2, nbnd = 16 ecutwfc = 20.0, ecutrho = 200.0, assume_isolated = 'martyna-tuckerman', // ATOMIC_SPECIES C 1.0 C.pbe-rrkjus.UPF H 1.0 H.pbe-rrkjus.UPF ATOMIC_POSITIONS angstrom H 5.5000000 7.98563953 3.5 C 5.5000000 6.89520922 3.5 C 5.5000000 6.89520922 3.5 C 6.7089386 6.19812524 3.5 H 7.652918 6.74309454 3.5 C 6.7089386 4.80187470 3.5 H 7.652918 4.25690561 3.5 C 5.5000000 4.10479062 3.5 H 7.652918 4.25690561 3.5 C 4.2910612 4.80187468 3.5 H 3.3470081 4.25690556 3.5 C 4.2910613 6.19812528 3.5 H 3.3470081 4.25690556 3.5 C 4.2910613 6.19812528 3.5 H 3.3470082 6.74309458 3.5 K_POINTS gamma -: pw.benzene.scf.in All (18,2) (QE-pw.x)</pre>	pr		Ô
<pre>ntyp = 2, nbnd = 16 ecutwfc = 20.0, ecutrho = 200.0, assume_isolated = 'martyna-tuckerman', / KELECTRONS / ATOMIC_SPECIES C 1.0 C.pbe-rrkjus.UPF H 1.0 H.pbe-rrkjus.UPF ATOMIC_POSITIONS angstrom H 5.5000000 7.98563953 3.5 C 5.5000000 6.89520922 3.5 C 6.7089386 6.19812524 3.5 H 7.6529918 6.74309454 3.5 C 6.7089386 4.80187470 3.5 H 7.6529918 4.25690561 3.5 C 5.5000000 4.10479062 3.5 H 5.5000000 3.01436043 3.5 C 4.2910612 4.80187468 3.5 H 3.3470081 4.25690556 3.5 C 4.2910613 6.19812528 3.5 H 3.3470082 6.74309458 3.5</pre>	&SYST ib A	prav = 6 = 11.0	
<pre>ecutrho = 200.0, assume_isolated = 'martyna-tuckerman', / &ELECTRONS /] ATOMIC_SPECIES C 1.0 C.pbe-rrkjus.UPF H 1.0 H.pbe-rrkjus.UPF ATOMIC_POSITIONS angstrom H 5.5000000 7.98563953 3.5 C 5.5000000 6.89520922 3.5 C 6.7089386 6.19812524 3.5 H 7.6529918 6.74309454 3.5 C 6.7089386 4.80187470 3.5 H 7.6529918 4.256905561 3.5 C 5.5000000 4.10479062 3.5 H 5.5000000 3.01436043 3.5 C 4.2910612 4.80187468 3.5 H 3.3470081 4.25690556 3.5 C 4.2910613 6.19812528 3.5 H 3.3470082 6.74309458 3.5 K_POINTS gamma -: pw.benzene.scf.in All (18,2) (QE-pw.x)</pre>	nt	typ = 2,	
SELECTRONS ATOMIC_SPECIES C 1.0 C 1.0 H 1.0 H.pbe-rrkjus.UPF ATOMIC_POSITIONS angstrom H 5.5000000 H 5.5000000 C 6.7089386 C 6.7089386 C 6.7089386 A.80187470 3.5 C 6.7089386 C 5.5000000 J 1.0479062 C 5.5000000 J 1.0479062 A 2.500000 J 1.0479062 C 4.2910612 J 4.80187468 J 3.470081 A 2.560556 H 3.3470082 A 7.4309458 J A H 3.3470082 A <td>ec as</td> <td>cutrho = 200.0,</td> <td>erman',</td>	ec as	cutrho = 200.0,	erman',
H 1.0 H.pbe-rrkjus.UPF ATOMIC_POSITIONS angstrom H 5.5000000 7.98563953 3.5 C 5.5000000 6.89520922 3.5 C 6.7089386 6.19812524 3.5 H 7.6529918 6.74309454 3.5 C 6.7089386 4.80187470 3.5 H 7.6529918 4.25690561 3.5 C 5.5000000 4.10479062 3.5 H 5.5000000 3.01436043 3.5 C 4.2910612 4.80187468 3.5 H 3.3470081 4.25690556 3.5 C 4.2910613 6.19812528 3.5 H 3.3470082 6.74309458 3.5 K_POINTS gamma -: pw.benzene.scf.in All (18,2) (QE-pw.x)	&ELEC	SPECIES	
H 5.5000000 7.98563953 3.5 C 5.5000000 6.89520922 3.5 C 6.7089386 6.19812524 3.5 H 7.6529918 6.74309454 3.5 C 6.7089386 4.80187470 3.5 H 7.6529918 4.25690561 3.5 C 5.5000000 4.10479062 3.5 H 5.5000000 3.01436043 3.5 C 4.2910612 4.80187468 3.5 H 3.3470081 4.25690556 3.5 C 4.2910613 6.19812528 3.5 H 3.3470082 6.74309458 3.5 K POINTS gamma	Н	1.0 H.pbe-rrkjus.UPF	
C 6.7089386 4.80187470 3.5 H 7.6529918 4.25690561 3.5 C 5.5000000 4.10479062 3.5 H 5.5000000 3.01436043 3.5 C 4.2910612 4.80187468 3.5 H 3.3470081 4.25690556 3.5 C 4.2910613 6.19812528 3.5 H 3.3470082 6.74309458 3.5 K_POINTS gamma -: pw.benzene.scf.in All (18,2) (QE-pw.x)	H C	5.5000000 7.98563953 3.5 5.5000000 6.89520922 3.5	
C 4.2910612 4.80187468 3.5 H 3.3470081 4.25690556 3.5 C 4.2910613 6.19812528 3.5 H 3.3470082 6.74309458 3.5 K_POINTS gamma -: pw.benzene.scf.in All (18,2) (QE-pw.x)	C H	6.70893864.801874703.57.65299184.256905613.5	
C 4.2910613 6.19812528 3.5 H 3.3470082 6.74309458 3.5 K_POINTS gamma -: pw.benzene.scf.in All (18,2) (QE-pw.x)	C	4.2910612 4.80187468 3.5	
K_POINTS gamma -: pw.benzene.scf.in All (18,2) (QE-pw.x)	С	4.2910613 6.19812528 3.5	
	- ;	pw.benzene.scf.in All (18.2	• (OE-pw.x)
			· · · · · · · · · · · · · · · · · · ·

emacs@cl	 × ×
File Edit Options Buffers Tools Help	
<pre>SINPUTPP prefix = 'benzene', filplot = 'psi2.benzene',</pre>	^
<pre>plot_num = 7, kpoint = 1, kband(1) = 1, kband(2) = 16, lsign = .true.,</pre>	
/ &PLOT	
<pre>fileout = '.xsf', iflag = 3, nfile = 1, output_format = 5, weight(1) = 1.0,</pre>	
/	
-: pp.benzene.psi2.in All (1,0) (QE-	-pp.x)
	*

1. How to calculate and plot molecular orbitals

To plot signed molecular-orbital densities $(sign(\psi_i(\boldsymbol{r}))|\psi_i(\boldsymbol{r})|^2)$, we need to:

- calculate Kohn-Sham states with pw.x (i.e. make an SCF calculation)
- instruct pp.x to write them in a suitable format to specified files
- plot orbitals with xcrysden
- See README.md for detailed instructions.



1. How to plot molecular orbitals with xcrysden

- Execute in the terminal:
 - pw.x < pw.benzene.scf.in > pw.benzene.scf.out
 - pp.x < pp.benzene.scf.in > pp.benzene.scf.out

The resulting molecular orbitals (i.e., $sign(\psi(\boldsymbol{r}))|\psi(\boldsymbol{r})|^2$) are written to psi2.benzene_*.xsf

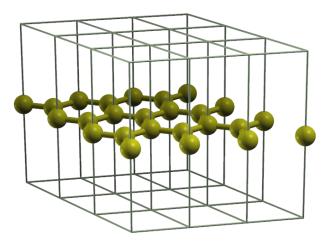
- Plot one of the generated XSF files with xcrysden, e.g.:
 - xcrysden --xsf psi2.benzene_K001_B006.xsf
 - and follow these instructions:
 - use the menu Tools-->Data Grid; a new window opens, press [OK]
 - an isosurface-control window appear; specify the Isovalue, say 0.005 and press [Submit]
 - click the Render +/- isovalue radiobutton and again press [Submit]
 - rotate and zoom the structure according to your preference
 - save the displayed state via the menu File-->Save Current State (e.g., save to my-display.xcrysden)
 - try this with other orbitals, e.g.:
 - xcrysden --xsf psi2.benzene_K001_B005.xsf --script my-display.xcrysden
- To plot all orbitals, execute: ./plot-psi2.sh

2. How to calculate a 2D-periodic system: graphene

A 2D-periodic system (e.g., a graphene sheet) is modelled by adding a vacuum layer in the 3rd direction.

• move to example2.graphene/ directory

look at the input file pw.graphene.scf.in; graphene has a 2-atom hexagonal unit cell in the xy plane: ibrav=4, celldm(1)=4.654, celldm(3)=some suitably large value, e.g. 3.0;



(remember: celldm(1) in Bohr radii, celldm(3)=c/a; alternatively: A=2.463, C=7.388 in Å)

atomic positions:
ATOMIC_POSITIONS (alat)
C 0.000000 0.000000 0.000000
C 0.000000 0.5773503 0.000000

or, equivalently: ATOMIC_POSITIONS (crystal) C 0.000000 0.000000 0.000000 C 0.333333 0.6666667 0.000000

```
k-points: use a dense grid in the xy plane only, e.g.
K_POINTS (automatic)
9 9 1 0 0 0
(a uniform 9×9×1 grid, centered on k = (0,0,0))
```

2. Graphene: DOS and bands (spaghetti)

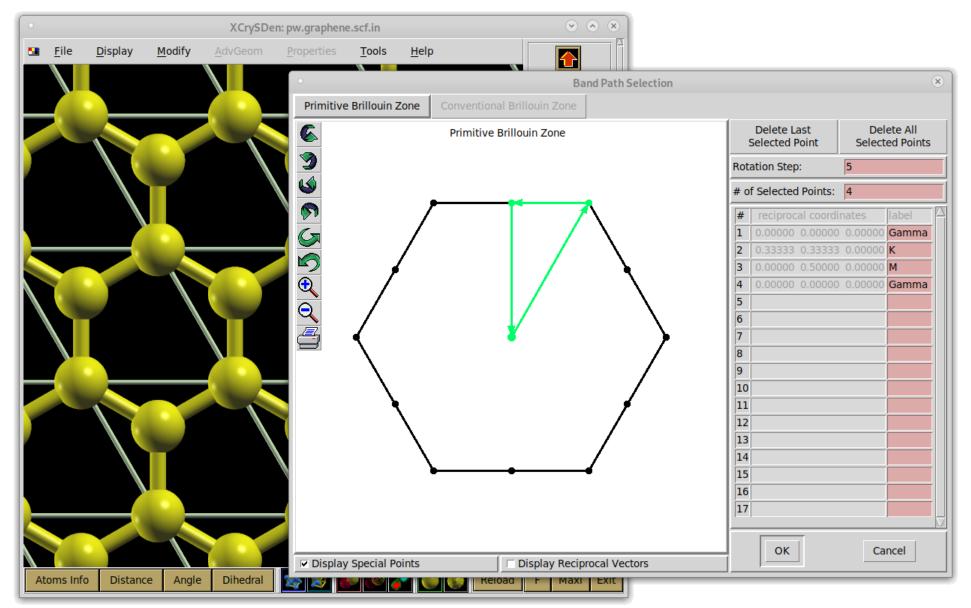
- DOS is typically calculated by a pw.x SCF calculation followed by a pw.x non-SCF calculation (calculation = 'nscf') with a denser k-point grid, and finally using dos.x post-processing code.
- to calculate the bands (spaghetti plot), the pw.x SCF calculation is followed by a pw.x "bands"-type non-SCF calculation (calculation = 'bands'), for which we need a suitable path of k-points. The most difficult (?) part is to figure out a suitable path of k-points.

You may either use the "k-path selection" tool of xcrysden or the SeeK-path web site at http://materialscloud.org/tools/seekpath.

• instructions for how to calculate DOS and bands are in README.md



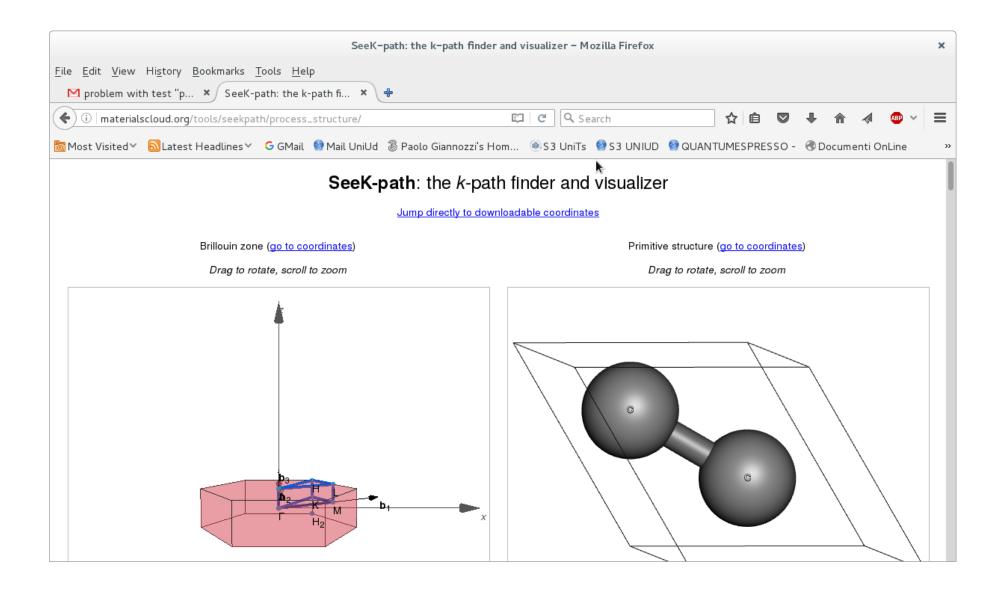
K-path selection tool of xcrysden



(important: to save k-path in Quantum ESPRESSO format, explicitly specify the *.pwscf extension)

SMR3966: Workshop on High Performance Computing for Materials Characterization, Design and Discovery Example: example2.graphene/

SeeK-path @ http://materialscloud.org/tools/seekpath



SMR3966: Workshop on High Performance Computing for Materials Characterization, Design and Discovery Example: example2.graphene/

3. Bulk system: Silicon



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

- move to example3.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 **outdir** and **pseudo_dir** variables which are set in the input files.)

Providing atomic structure in input

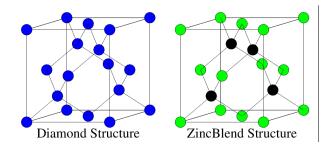
How is the crystal structure defined? This is a very simple case: the diamond lattice is an fcc (face-centered cubic) lattice with two atoms per unit cell. You need to specify:

- 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!





Brillouin zone (BZ) sampling



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 example3.Si) and its content:

•ls ./
silicon.save silicon.xml

(to see only these files, you may need to use ls 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.

Running the pw.x code (II)



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

\$	grep -e "total energy" -e e	estim	nated pw.si.scf.out
	total energy	=	-15.79103344 Ry
	estimated scf accuracy	<	0.06376674 Ry
	total energy	=	-15.79409289 Ry
	estimated scf accuracy	<	0.00230109 Ry
	total energy	=	-15.79447822 Ry
	estimated scf accuracy	<	0.00006291 Ry
	total energy	=	-15.79449510 Ry
	estimated scf accuracy	<	0.00000448 Ry
!	total energy	=	-15.79449593 Ry
	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) \checkmark

To make convergence tests easier and faster, scripts are commonly used. To this end, Unix shell-scripts have been traditionally used.

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

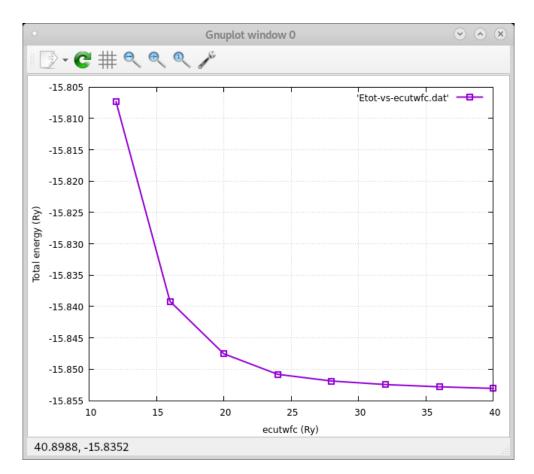
```
Unix shell-script
#!/bin/sh
rm -f si.etot_vs_ecut.dat
for ecut in 12 16 20 24 28 32
do
   cat > pw.si.scf.$ecut.in << EOF
&CONTROL
   prefix='silicon',
&SYSTEM
   ibrav = 2,
   celldm(1) = 10.2,
   nat = 2,
   ntyp = 1,
   ecutwfc = $ecut,
 1
&ELECTRONS
ATOMIC_SPECIES
  Si 28.086 Si.pz-vbc.UPF
ATOMIC_POSITIONS
  Si 0.00 0.00 0.00
  Si 0.25 0.25 0.25
K_POINTS automatic
  4 4 4 1 1 1
EOF
   pw.x -in pw.si.scf.$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
            111
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
```

QUANTUMESPRESSO

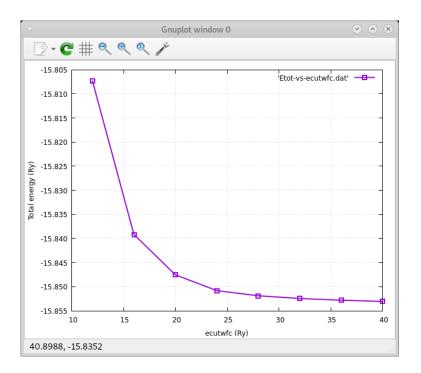
1.1 Convergence w.r.t. the kinetic energy cutoff (III)

- 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



SMR3966: Workshop on High Performance Computing for Materials Characterization, Design and Discovery Example 3.Si/ex1.ecutwfc/

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
}
```

1.2 Convergence w.r.t. k-points (II)



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))

SMR3966: Workshop on High Performance Computing for Materials Characterization, Design and Discovery Example 3.Si/ex2.kpoints/

1.2 Convergence w.r.t. k-points (III)

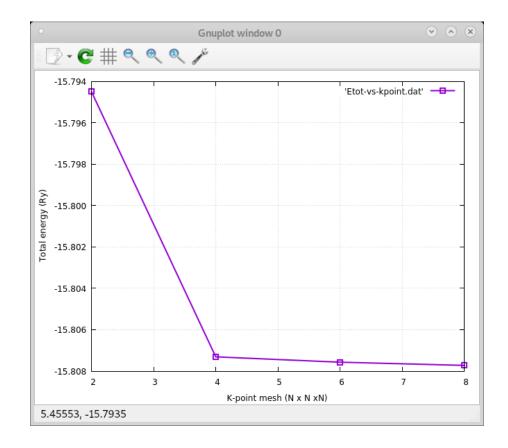


The Unix shell-script for testing the convergence with respect to k-points is located in example3.Si/ex2.kpoints.classic/ directory (see README.md for detailed instructions).

Within this directory execute:

• ./kpoints.sh

You should get a plot like this one:



1.3 Equation of State: silicon



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
}
```

But we are going to use a shell-script.

SMR3966: Workshop on High Performance Computing for Materials Characterization, Design and Discovery Example: example1.Si/ex3.alat/

1.3 Equation of State: silicon (II)

The corresponding Unix shell-script is located in example3.Si/ex3.alat.classic/ directory (see README.md for detailed instructions).

Within this directory execute:

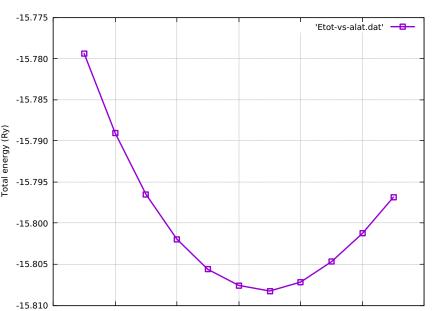
• ./alat.sh

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, kpoints, pseudopotential, ...

You should find that:

- -15.780 -15.785 otal energy (Ry -15.790 -15.795 -15.800 -15.805 -15.810 9.6 9.8 10.0 10.2 10.4 10.6 10.8 Lattice parameter (Bohr)
- 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.

SMR3966: Workshop on High Performance Computing for Materials Characterization, Design and Discovery Example: example1.Si/ex3.alat/





1.3 Equation of State: silicon (III)



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.sh 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	$E(a_3)$

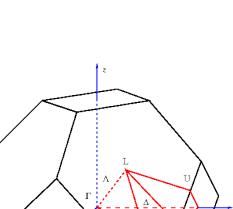
. . .

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





1.4 Band Structure of Silicon (II)



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:

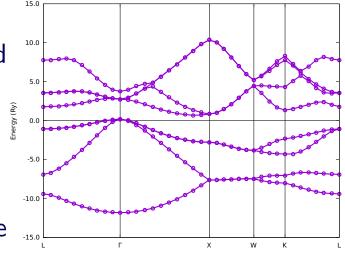
- pw.x -in pw.Si.scf.in > pw.Si.scf.out
- pw.x -in pw.Si.bands.in > pw.Si.bands.out
- bands.x -in pp.Si.bands.in > pp.Si.bands.out

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 example1.Si/ex4.bands/ and read the README.md file
- set suitable values for celldm(1), ecutwfc, and K_POINTS
- and execute: bands.sh
- 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



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 example4.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

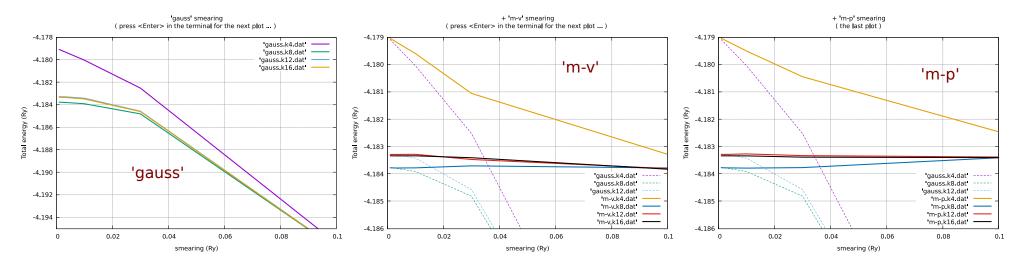
With PWTK this can be achieved with the following snippet: CHECK IF SH SCRIPT

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

- move to example4.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 AI and m-v or m-p smearing, good convergence is achieved for a 12 12 12 k-point grid and degauss ~ 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!

SMR3966: Workshop on High Performance Computing for Materials Characterization, Design and Discovery Example: example4.Al/ex1.degauss/

2.2 How to plot charge-density



Example example4.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 example4.Al/ex2.chdens/ directory (chdens is an acronym for chargedensity)
- 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)

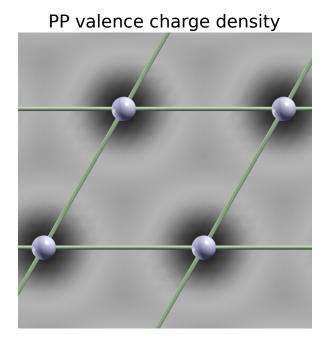
2.2 How to plot charge-density (II)



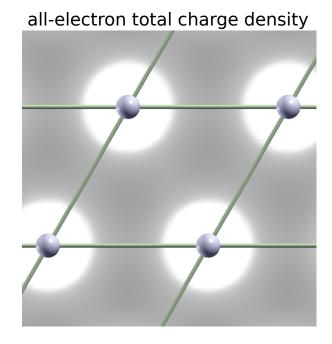
- to calculate all-electron valence and total charge densities, execute:
 - pwtk 2-chdens-paw.pwtk

(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):



all-electron valence charge density



SMR3966: Workshop on High Performance Computing for Materials Characterization, Design and Discovery Example: example4.Al/ex2.chdens/