



# MaX School on Advanced Materials and Molecular Modelling with Quantum ESPRESSO

# QE-2021: Hands-on session – Day-1

(Basic SCF + post-processing calculations)















# QE-2021: Hands-on session - Day-1



Topics of Day-1 hands-on session:

- 1. Installation/compilation of Quantum ESPRESSO (example0.QE-compilation)
- 2. How to run basic PWscf (pw.x) calculations
- 3. How to run post-processing calculations to plot molecular-orbitals and charge-density (pp.x), DOS (dos.x), and band-structure (bands.x)
- 4. How to calculate low-dimensional systems (example1.benzene and example2.graphene)

# **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)

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#### Hands-on material



Hands-on material for each day is contained within its own directory:

- Day-1/ hands-on exercises for Day-1
- Day-2/, Day-3/ ... Day-10/ hands-on exercises for Day-2 to Day-10

Please go to the Day-1/ directory and execute: git pull this will update the hands-on exercises to the latest version from the GitLab.

- All directories contain a README.md file with instructions how to run exercise(s)
- Naming of files is described in README-filenames.md (in Day-1/).
- 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

# 0. Compilation of Quantum ESPRESSO



Please go to the Day-1/ directory, then execute:

```
$ cd example0.QE-compilation/
$ tar zxvf qe-6.7-ReleasePack.tgz
$ cd qe-6.7/
```

Now read the INSTALL.md file therein. It contains installation (compilation) instructions. In essence, compilation consists of:

```
$ ./configure [options]
$ make target
```

(remark: make alone prints a list of acceptable targets)

Today we will only compile pw.x program (for the sake of time constraint), hence:

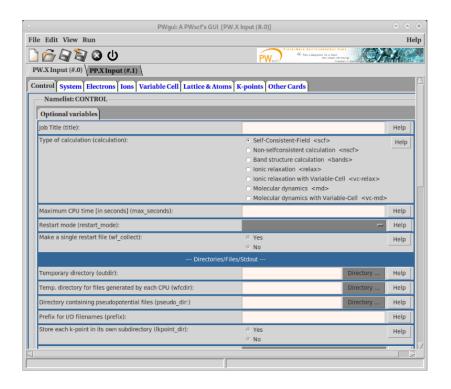
```
$ ./configure
$ make pw
```

... and wait for a while as compilation takes some time. If all went OK, the compiled pw.x program (along with some other executables) is now located in bin/ directory.

# Preparation of Quantum ESPRESSO input files

A few tools are available that aid at editing Quantum ESPRESSO input files:

- PWgui a QE input file builder GUI (pwgui)
- QE-emacs-modes makes editing of input files easier with emacs editor. It provides syntax highlighting, basic auto-indentation, and several utility commands; its manual is available in the QE sub-directory GUI/QE-modes/Doc/user\_guide.pdf



```
~ A X
                emacs@catalyst.ijs.si
File Edit Options Buffers Tools Help
   calculation = 'scf'
              = 'silicon'
   outdir
              = '/temp/tone/pw/Si'
   tstress
   tprnfor
               = .true.
   ibrv = 2 ! mistyped variable is not highlighted
   A = 5.43
   nat = 2
   ntyp = 1
   ecutwfc = 18
&ELECTRONS
  conv_thr = 1.0d-8
ATOMIC SPECIES
  Si 28.086 Si.pz-vbc.UPF
ATOMIC POSITIONS alat
          0.00 0.00
0.25 0.25
K POINTS automatic
  444 111
-:--- scf.Si.in
```

#### **About QE-emacs-modes**

QE-emacs-modes package provides syntax highlighting, auto-indentation, and several utility commands, in particular:

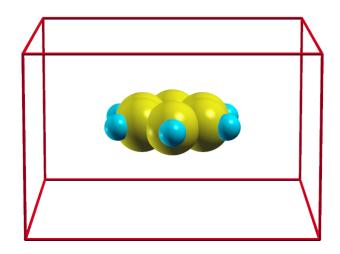
- Alt-x prog-insert-template inserts a respective input file template
- Alt-x prog-NAMELIST inserts a blank namelist named NAMELIST
- Alt-x prog-CARD inserts a blank namelist named CARD
- Alt-x prog-variable inserts a namelist variable named variable
- Alt-x prog-mode toggles the respective mode
- Alt-x indent-region indents region

#### where

- prog is one of qe, pw, neb, cp, ph, ld1, or pp (these stands for pw.x, neb.x, ... Quantum ESPRESSO (QE) programs)
- NAMELIST is the uppercase name for a given QE namelist
- CARD is the uppercase name for a given QE card
- variable is the lowercase name for a given namelist variable

#### 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 Day-1/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
- 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

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### 1. How to calculate and plot molecular orbitals

Here are the two needed input files for calculation of molecular orbitals of benzene (actually,  $sign(\psi(\mathbf{r}))|\psi(\mathbf{r})|^2$ ), opened with emacs using QE-emacs-modes:

```
emacs@cl
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
ATOMIC POSITIONS angstrom
  H 5.5000000 7.98563953
      5.5000000
                6.89520922
      6.7089386 6.19812524
     7.6529918
                6.74309454
                4.80187470
     6.7089386
                4.25690561
      7.6529918
     5.5000000
                4.10479062
  H 5.5000000 3.01436043
      4.2910612 4.80187468
  H 3.3470081 4.25690556
    4.2910613 6.19812528
  H 3.3470082
                6.74309458
K_POINTS gamma
-:--- pw.benzene.scf.in All (18,2)
                                        (QE-pw.x)
Beginning of buffer
```

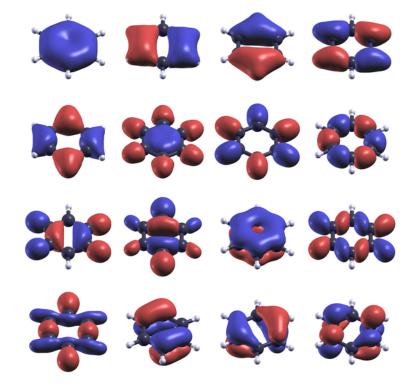
```
(v) (A) (X
                    emacs@cl
File Edit Options Buffers Tools Help
&INPUTPP
    prefix
             = 'benzene',
   filplot = 'psi2.benzene',
   plot num = 7,
    kpoint = 1,
   kband(1) = 1,
   kband(2) = 16,
   lsign = .true.,
&PL0T
   fileout
                 = '.xsf',
   iflag
   nfile
   output format = 5,
   weight(1) = 1.0,
-:--- pp.benzene.psi2.in All (1,0)
                                         (QE-pp.x)
```

# 1. How to calculate and plot molecular orbitals

To plot molecular orbitals (actually the signed square modulus,  $sign(\psi(\mathbf{r}))|\psi(\mathbf{r})|^2$ ), 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.,  $sign(\psi(r))|\psi(r)|^2$ ) 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
```

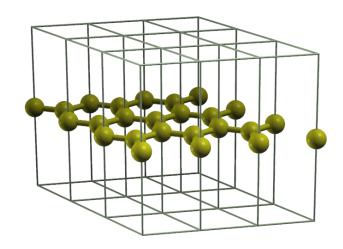
• To plot all orbitals, execute: ./plot-psi2.sh

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# 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 Day-1/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 0.666667 0.000000

 $\bullet$  k-points: use a dense grid in the xy plane only, e.g.

K\_POINTS (automatic)

9 9 1 0 0 0

(a uniform  $9\times9\times1$  grid, centered on  $\mathbf{k}=(0,0,0)$ )

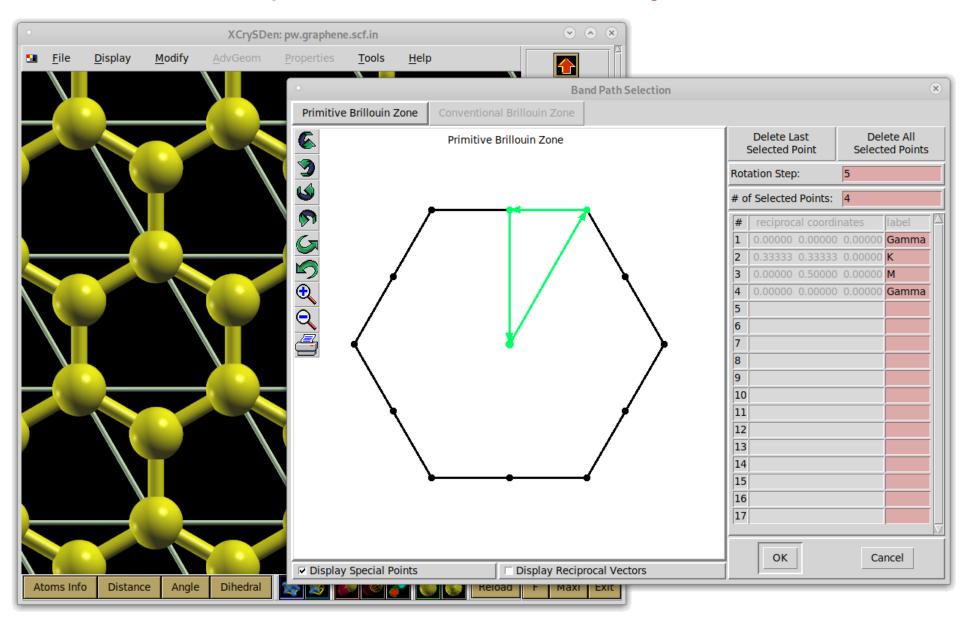
# 2. Graphene: DOS and bands (spaghetti)

- 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)

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

