

# DFT hands-on using quantum espresso



**First edition of the African School on Computational Materials for  
Sustainable Energy Technologies (CMSET 2025)**

**November 26, 2025**

- Download cif file using material project database
- Convert Cif file to input file for quantum espresso
- Read and visual input and output file
- Convergence test for Ecut, Ecutrho and kpoint mesh
- Cell relaxation
- Total and Partial Density of state (PDOS)
- Band structure

# Download the NaCl Crystal Structure (CIF Format) from Public Databases

Material project

Crystallography Open Database (COD)

Material cloud

AFLOW

Material project

Start exploring materials

Type your chemical formula: NaCl

The Materials Project

Apps About Community ML API

Home / Apps / Materials Explorer / Cl-Na / NaCl / mp-22851

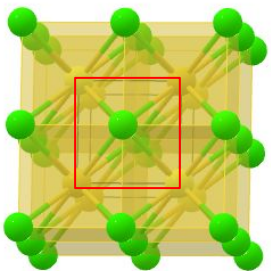
## Materials Explorer

References Documentation

**NaCl**  
mp-22851

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- Crystal Structure
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Energy Above Hull 0.154 eV/atom

Space Group Pm $\bar{3}$ m

Band Gap 3.97 eV

Predicted Formation Energy -1.884 eV/atom

Magnetic Ordering Non-magnetic

Total Magnetization 0.00  $\mu$ B/f.u.

Experimentally Observed Yes

Download cif file

# Understanding CIF Format: Key Structural Parameters

# generated using pymatgen

data\_NaCl

\_symmetry\_space\_group\_name\_H-M Pm-3m

\_cell\_length\_a 3.50219000

\_cell\_length\_b 3.50219000

\_cell\_length\_c 3.50219000

\_cell\_angle\_alpha 90.00000000

\_cell\_angle\_beta 90.00000000

\_cell\_angle\_gamma 90.00000000

\_symmetry\_Int\_Tables\_number 221

\_chemical\_formula\_structural NaCl

\_chemical\_formula\_sum 'Na1 Cl1'

\_cell\_volume 42.95553287

\_cell\_formula\_units\_Z 1

loop\_

\_symmetry\_equiv\_pos\_site\_id

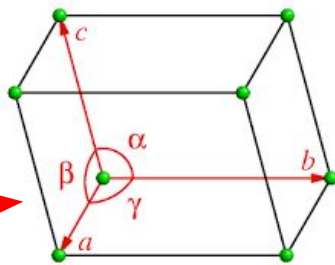
\_symmetry\_equiv\_pos\_as\_xyz

1 'x, y, z'

2 '-x, -y, -z'

3 '-y, x, z'

4 .....



loop\_

\_atom\_type\_symbol

\_atom\_type\_oxidation\_number

Na+ 1.0

Cl- -1.0

loop\_

\_atom\_site\_type\_symbol

\_atom\_site\_label

\_atom\_site\_symmetry\_multiplicity

\_atom\_site\_fract\_x

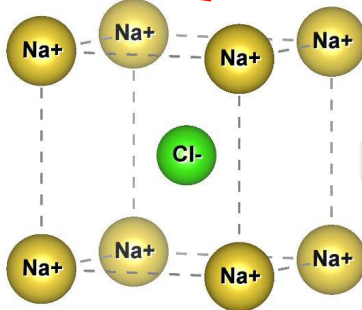
\_atom\_site\_fract\_y

\_atom\_site\_fract\_z

atom\_site occupancy

Na+ Na0 1 0.00000000 0.00000000 0.00000000 1

Cl- Cl1 1 0.50000000 0.50000000 0.50000000 1



You can use vesta software for cif file visualisation

# Converting CIF file to input file for quantum espresso (.in format)

cif2cell tool

Open terminal and type

```
cif2cell file_name.cif -p quantum-espresso -o file_name.in
```

Material cloud website

Work

Quantum ESPRESSO input generator and structure visualizer



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EXPLORE

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Work

Tools

Quantum ESPRESSO input generator and structure visualizer

## Quantum ESPRESSO input generator and structure visualizer

► About the Quantum ESPRESSO input generator and structure visualizer

► Instructions

► Acknowledgements

Upload a crystal structure

Pick an example structure

Upload a file:

Select the file format:

Select the protocol:<sup>[?]</sup>

Select the XC functional:

Select the magnetism/smearing:<sup>[?]</sup>

Refine cell (using spglib):

► Advanced settings<sup>[?]</sup>

Browse... NaCl.cif

CIF File (.cif) [parser: ase]

balanced

PBE

non-magnetic metal (fractional occupations)

No

Insert cif file

Exchange-correlation  
Functional

Magnetism of material and its  
electronic nature

By continuing, you agree with the terms of use of this service.

Generate the PWscf input file

# Input for quantum espresso

## &CONTROL

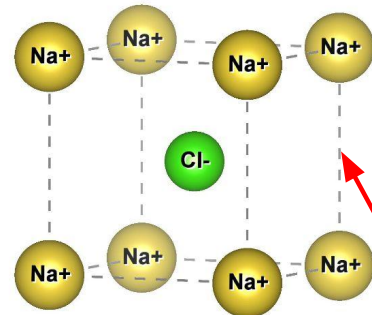
```
calculation='scf',  
outdir='.',  
prefix='NaCl',  
pseudo_dir='/home/max/pseudo',  
verbosity='low',  
tprnfor=.true., tstress=.true.,  
/
```

Type of the calculation:  
scf, relax, vc-relax

Calculate forces and  
stress

## &SYSTEM

```
ibrav = 0  
A = 3.4  
nat = 2  
ntyp = 2  
ecutwfc=40,  
ecutrho=200,  
input dft='pbe',  
occupations='smearing',  
smearing='mv',  
degauss=0.005d0,  
/
```



Number and type of  
atoms in unit cell

## &ELECTRONS

```
conv_thr=1d-08,  
mixing_beta=0.7d0,  
/
```

Convergence threshold  
for the energy

Occupations “fixed” for  
semiconductor/insulator and  
“smearing” for metals

## &IONS

```
ion_dynamics='bfgs',  
/
```

Ionic relaxation

## &CELL

```
cell_dynamics='bfgs',  
press=0.d0,  
press_conv_thr=0.5d0,  
/
```

Cell relaxation

## ATOMIC\_SPECIES

```
Cl 35.45150 cl_pbe_v1.4.uspp.F.UPF  
Na 22.98900 na_pbe_v1.5.uspp.F.UPF
```

## CELL\_PARAMETERS {alat}

```
1.0000000000000000 0.0000000000000000 0.0000000000000000  
0.0000000000000000 1.0000000000000000 0.0000000000000000  
0.0000000000000000 0.0000000000000000 1.0000000000000000
```

## ATOMIC\_POSITIONS {crystal}

```
Cl 0.5000000000000000 0.5000000000000000 0.5000000000000000  
Na 0.0000000000000000 0.0000000000000000 0.0000000000000000
```

## K\_POINTS {automatic}

```
5 5 5  
0 0 0
```

Complete description of input file in the website:

[https://www.quantum-espresso.org/Doc/INPUT\\_PW.html](https://www.quantum-espresso.org/Doc/INPUT_PW.html)

# Using XCrySDen to Visualize Quantum ESPRESSO Input and Output Files

Open terminal and type:

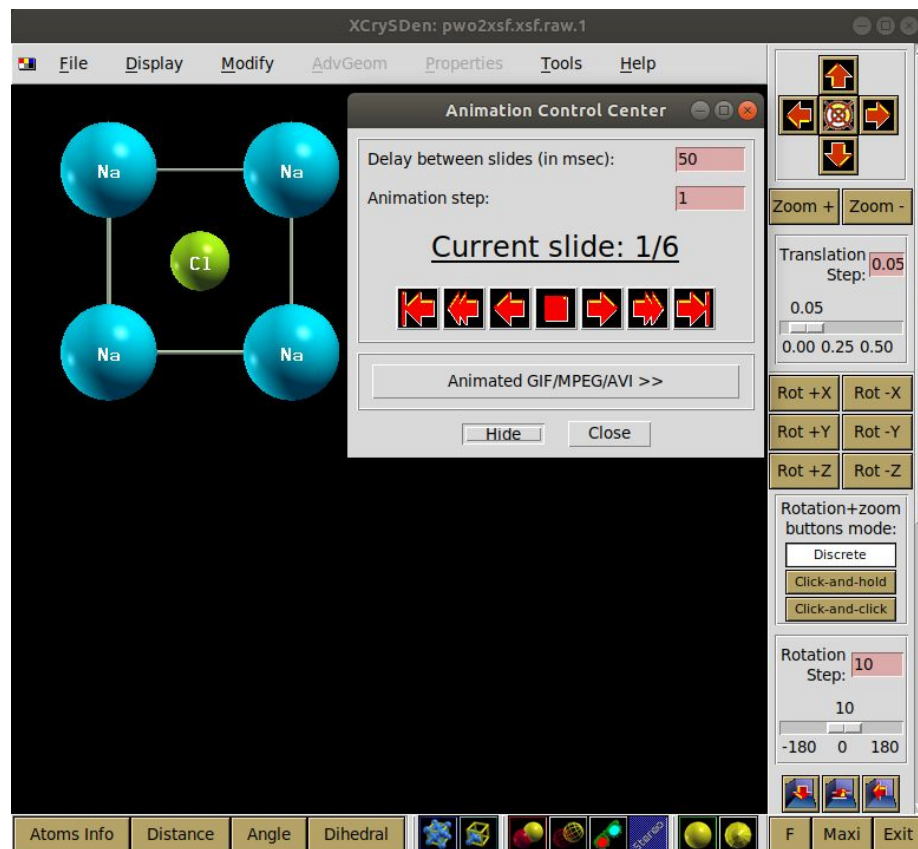
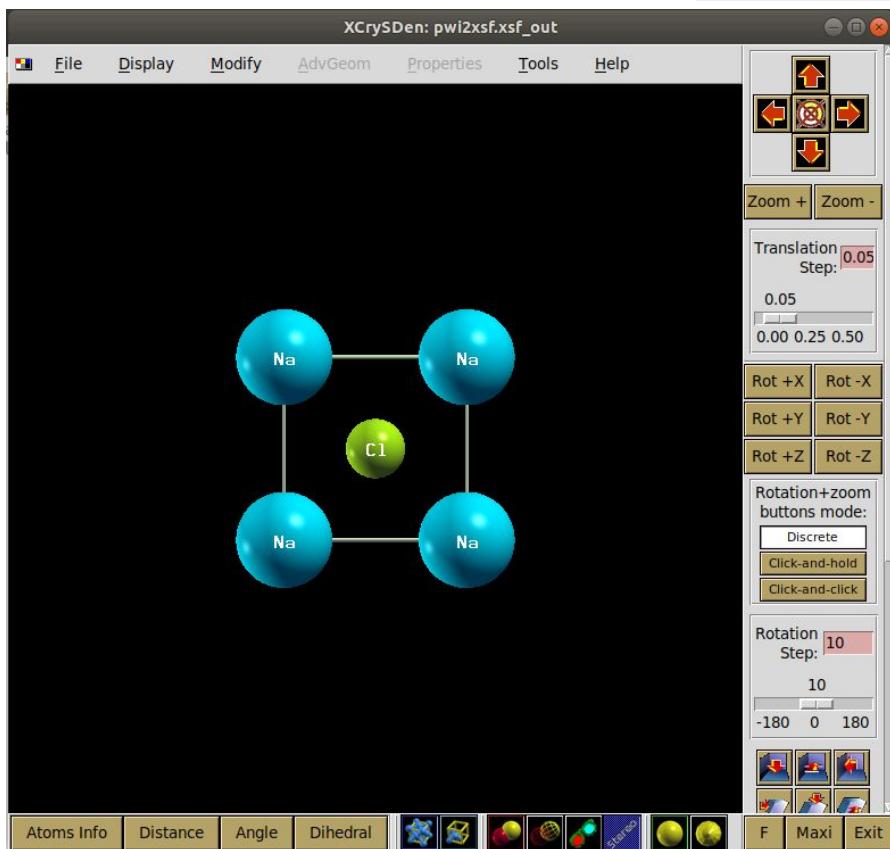
**conda activate visualise**

Open terminal and type **xcrysdn**

**file**

**Open PWscf**

**Open PWscf Input file or Output file**



# Input file and scf loop in quantum espresso

Calculated from ions positions

Initial Guess  
 $\rho(\mathbf{r})$

Will guessed from pseudo potential file or outdir file

ATOMIC\_POSITIONS {crystal}  
Cl 0.50 0.50 0.50  
Na 0.00 0.00 0.00

startingwfc='atomic' —from pseudopotential  
startingwfc='file' — from outdir file

Calculate Effective Potential  
 $v_{\text{eff}}(\mathbf{r}) = V_{\text{en}}(\mathbf{r}) + \int \frac{\rho(\mathbf{r}')}{|\mathbf{r}-\mathbf{r}'|} d\mathbf{r}' + V_{\text{xc}}[\rho(\mathbf{r})]$

exchange-correlation specified in input otherwise read from pseudopotential

Solved using iterative diagonalization

Solve Kohn-Sham Equations  
 $[-\frac{\hbar^2}{2m_e} \nabla_i^2 + v_{\text{eff}}] \psi_i = \epsilon_i \psi_i$

diagonalization='david'

Evaluate the Electron Density & Total Energy  
 $\rho(\mathbf{r}) = \sum_i |\psi_i(\mathbf{r})|^2 \rightarrow E_{\text{tot}}[\rho(\mathbf{r})] = \dots$

input\_dft='pbe',

Converged?

mixing old and new density for better convergence stability

Calculate converged energy, forces and stress

conv\_thr=1d-08,

mixing\_beta=0.7d0,

tpnfor=.true.,  
tstress=.true.,

Output Quantities

$\rho_0(\mathbf{r}), E_i[\rho_0(\mathbf{r})] \rightarrow \text{Forces, Eigenvalues, ...}$



# Run input file and Read output file

Open input file directory

activate quantum espresso: conda activate qespresso

run input using `pw.x -input input.in > output.out`

```
bravais-lattice index      =          0
lattice parameter (alat)  =      6.4251  a.u.
unit-cell volume          =    265.2365 (a.u.)^3
number of atoms/cell      =          2
number of atomic types    =          2
number of electrons       =     16.00
number of Kohn-Sham states=      12
kinetic-energy cutoff      =     40.0000  Ry
charge density cutoff     =     200.0000  Ry
convergence threshold     =     1.0E-08
mixing beta               =     0.7000
number of iterations used =          8  plain    mixing
Exchange-correlation      = PBE ( 1  4  3  4  0  0)
```

```
celldm(1)=  6.425069  celldm(2)=  0.000000  celldm(3)=  0.000000
celldm(4)=  0.000000  celldm(5)=  0.000000  celldm(6)=  0.000000
```

```
crystal axes: (cart. coord. in units of alat)
a(1) = (  1.000000  0.000000  0.000000 )
a(2) = (  0.000000  1.000000  0.000000 )
a(3) = (  0.000000  0.000000  1.000000 )
```

```
reciprocal axes: (cart. coord. in units 2 pi/alat)
b(1) = (  1.000000  0.000000  0.000000 )
b(2) = (  0.000000  1.000000  0.000000 )
b(3) = (  0.000000  0.000000  1.000000 )
```

- For an insulator is number of valence electrons /2
- For a metal, 20% more (minimum 4 more)

celldm(1) = a  
celldm(2) = b/a,  
celldm(3) = c/a,  
celldm(4) = cos(bc),  
celldm(5) = cos(ac),  
celldm(6) = cos(ab)

# Reading output file

```
number of k points= 10| Marzari-Vanderbilt smearing, width (Ry)= 0.0050
cart. coord. in units 2pi/alat
k( 1) = ( 0.0000000 0.0000000 0.0000000), wk = 0.0160000
k( 2) = ( 0.0000000 0.0000000 0.2000000), wk = 0.0960000
k( 3) = ( 0.0000000 0.0000000 0.4000000), wk = 0.0960000
k( 4) = ( 0.0000000 0.2000000 0.2000000), wk = 0.1920000
k( 5) = ( 0.0000000 0.2000000 0.4000000), wk = 0.3840000
k( 6) = ( 0.0000000 0.4000000 0.4000000), wk = 0.1920000
k( 7) = ( 0.2000000 0.2000000 0.2000000), wk = 0.1280000
k( 8) = ( 0.2000000 0.2000000 0.4000000), wk = 0.3840000
k( 9) = ( 0.2000000 0.4000000 0.4000000), wk = 0.3840000
k(10) = ( 0.4000000 0.4000000 0.4000000), wk = 0.1280000
```

As the k-point weight increase, the symmetry in this kpoint increase

```
iteration # 1      ecut= 40.00 Ry      beta=0.70
Davidson diagonalization with overlap
ethr = 1.00E-02, avg # of iterations = 5.0
```

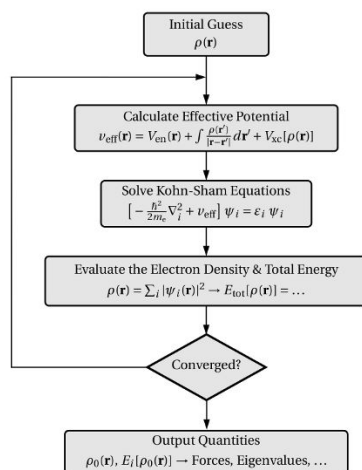
Threshold (ethr) on eigenvalues was too large:  
Diagonalizing with lowered threshold

```
Davidson diagonalization with overlap
ethr = 3.31E-04, avg # of iterations = 2.3
```

negative rho (up, down): 1.598E-02 0.000E+00

total cpu time spent up to now is 1.7 secs

```
total energy      = -128.86376078 Ry
Harris-Foulkes estimate = -128.89603018 Ry
estimated scf accuracy < 0.05107874 Ry
```



Kohn-Sham Hamiltonian matrix solved for each k-point

Scf loop has been solved for the first time

# Reading output file

The total energy is the sum of the following terms:

```
one-electron contribution = -74.18279152 Ry
hartree contribution      =  42.36507042 Ry
xc contribution           = -23.92644874 Ry
ewald contribution        = -73.13425203 Ry
smearing contrib. (-TS)   =  0.00000000 Ry
```

**Eigenvalues of Kohn-Sham  
Hamiltonian for each kpoint**

```
      k = 0.0000 0.0000 0.0000 ( 1141 PWs)   bands (ev):
-47.0643 -18.9626 -18.9626 -18.9626 -10.6534  2.4857  2.4857  2.4857
  7.3475  11.4677  11.4677  19.6121
```

```
! total energy      = -128.87842187 Ry
  Harris-Foulkes estimate = -128.87842187 Ry
  estimated scf accuracy < 7.6E-11 Ry
```

**Converged total energy**

```
atom   1 type   1 force =  0.00000000  0.00000000  0.00000000
atom   2 type   2 force =  0.00000000  0.00000000  0.00000000

Total force =  0.000000 Total SCF correction =  0.000000
```

**Forces after converging scf loop**

# Reading output file

```
negative rho (up, down): 1.517E-02 0.000E+00
total stress (Ry/bohr**3)
0.00021242 0.00000000 0.00000000
0.00000000 0.00021242 0.00000000
0.00000000 0.00000000 0.00021242
```

	(kbar)	P=
31.25	0.00	0.00
0.00	31.25	0.00
0.00	0.00	31.25

Hydrostatic pressure  
on the cell

`pw.x -input input.in > output.out`

```
Parallel routines
fft_scatter :      0.10s CPU      0.10s WALL ( 6305 calls)

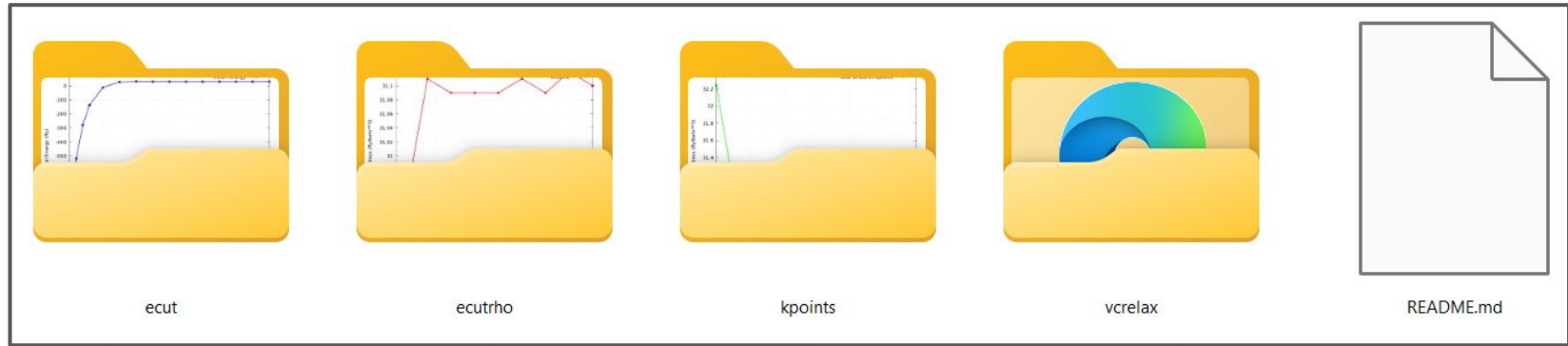
PWSCF :      4.76s CPU      5.08s WALL
```

`mpirun --bind-to core --map-by core -np 2 (or 4 or 6) pw.x -input input.in > output.out`

```
Parallel routines
fft_scatter :      0.13s CPU      0.14s WALL ( 6301 calls)

PWSCF :      3.00s CPU      3.12s WALL
```

# Optimisation steps and folders



**ECut optimisation**

**Ecutrho optimisation**

**K-point meshes optimisation**

**Cell relaxation**

**Readme file**

# Cutoff-Energy (Ecut) Convergence Study

ecut.\$ecut.in

```
&CONTROL
  calculation='scf',
  outdir='.',
  prefix='NaCl',
  pseudo_dir='.././../pseudo',
  verbosity='low',
  tprnfor=.true.,
  tstress=.true.,
/
&SYSTEM
 ibrav = 0
  A = 3.4
  nat = 2
  ntyp = 2
  ecutwfc = ${ecut},
  ecutrho = 200,
  input_dft='pbe',
  occupations='smearing',
  smearing='mv',
  degauss=0.005d0,
/
&ELECTRONS
  conv_thr=1d-8,
  mixing_beta=0.7d0,
/
&IONS
  ion_dynamics='bfgs',
/
&CELL
  cell_dynamics='bfgs',
  press=0.d0,
  press_conv_thr=0.5d0,
/
```

CELL\_PARAMETERS {alat}

```
1.0 0.0 0.0
0.0 1.0 0.0
0.0 0.0 1.0
```

ATOMIC\_SPECIES

```
Cl 35.45150 cl_pbe_v1.4.uspp.F.UPF
Na 22.98900 na_pbe_v1.5.uspp.F.UPF
```

ATOMIC\_POSITIONS {crystal}

```
Cl 0.5 0.5 0.5
Na 0.0 0.0 0.0
```

K\_POINTS {automatic}

```
5 5 5 0 0 0
```

## Set cutoffs

for ecut in 20 22 24 26 30 35 40 45 50 55  
60 65 70 75 80; do

## Target parameter for accurate optimization



## Hydrostatic pressure

	(kbar)	P= -924.42
-924.42	0.00	0.00
0.00	-924.42	0.00
0.00	0.00	-924.42

## Total energy

total energy = -128.49580518 Ry  
Harris-Foulkes estimate = -128.49580519 Ry  
estimated scf accuracy < 9.1E-09 Ry

The total energy is the sum of the following terms:

one-electron contribution = -73.12275949 Ry  
hartree contribution = 41.52353047 Ry  
xc contribution = -23.76232413 Ry  
ewald contribution = -73.13425203 Ry  
smearing contrib. (-TS) = -0.00000000 Ry

convergence has been achieved in 6 iterations

# Cutoff-Energy (Ecut) Convergence Study

We will use Gnuplot to make the plot. Here is the script code for plotting:

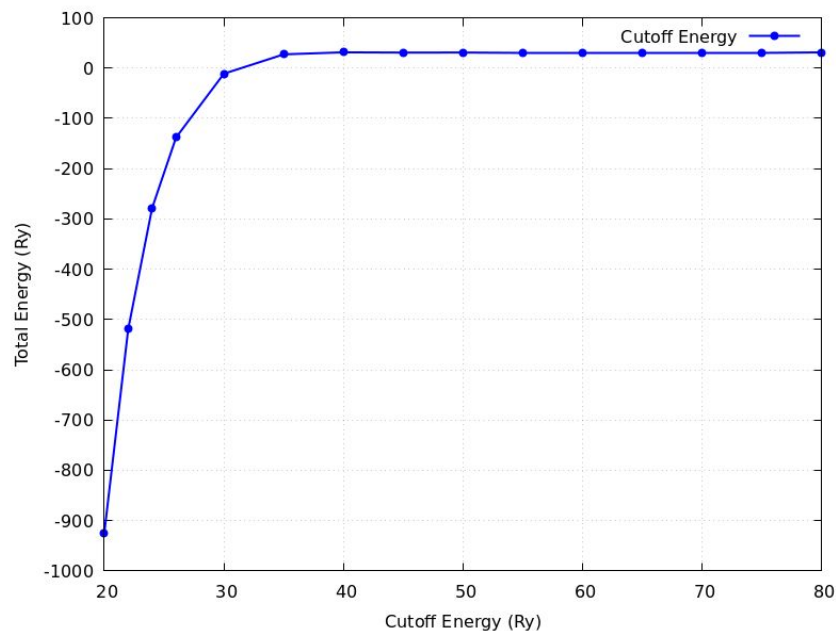
calc-pres.dat

20	-924.42
22	-518.80
24	-278.47
26	-138.08
30	-11.19
35	27.21
40	31.25
45	30.79
50	30.88
55	31.13
60	31.13
65	30.99
70	30.94
75	31.01
80	31.14

plot\_ecut.gp

```
# plot_ecut.gp
# Gnuplot script to plot total energy vs. cutoff
energy
# Set output to PNG image file
set terminal pngcairo size 800,600
set output "ecut_convergence.png"
set xlabel "Cutoff Energy (Ry)"
set ylabel "Total Stress (Ry/Bohr**3)"
set style line 1 lc rgb "blue" lw 2 pt 7
# Plot the data
plot "ecutpres.dat" using 1:2 with linespoints
linestyle 1 title "Cutoff Energy«
# Close the output file
set output
```

Hydrostatic pressure as a function of Ecut energy



Based on Hydrostatic convergence test, an ecut value of 60 Ry is appropriate for a convergence within ~1-2 kbar.



# Charge Density cutoff (Ecutrho) Convergence Study

ecut.\$ecutrho.in

```
&CONTROL
  calculation='scf',
  outdir='.',
  prefix='NaCl',
  pseudo_dir='.././../pseudo',
  verbosity='low',
  tprnfor=.true.,
  tstress=.true.,
/
&SYSTEM
  ibrav = 0
  A = 3.4
  nat = 2
  ntyp = 2
  ecutwfc = 60
  ecutrho = ${ecutrho}
  input_dft='pbe',
  occupations='smearing',
  smearing='mv',
  degauss=0.005d0,
/
&ELECTRONS
  conv_thr=1d-8,
  mixing_beta=0.7d0,
/
&IONS
  ion_dynamics='bfgs',
/
&CELL
  cell_dynamics='bfgs',
  press=0.d0,
  press_conv_thr=0.5d0,
/
```

```
CELL_PARAMETERS {alat}
  1.0 0.0 0.0
  0.0 1.0 0.0
  0.0 0.0 1.0

ATOMIC_SPECIES
  Cl 35.45150 cl_pbe_v1.4.uspp.F.UPF
  Na 22.98900 na_pbe_v1.5.uspp.F.UPF

ATOMIC_POSITIONS {crystal}
  Cl 0.5 0.5 0.5
  Na 0.0 0.0 0.0

K_POINTS {automatic}
  5 5 5 0 0 0
```

**Set ecutrho**

for ecutrho in 120 180 240 300 360 420  
480 540 600; do



# Charge Density cutoff (Ecutrho) Convergence Study

We will use Gnuplot to make the plot. Here is the script code for plotting:

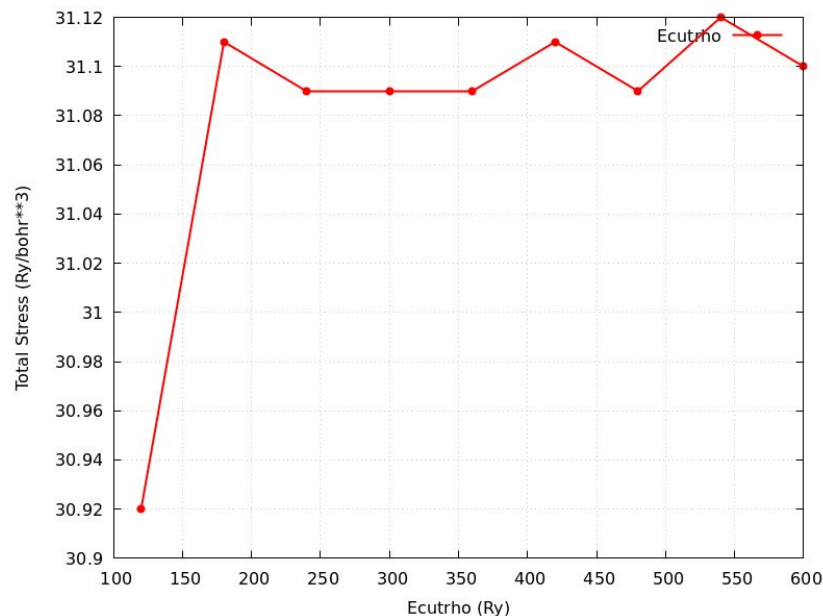
calc-ecutrho.dat

120	30.92
180	31.11
240	31.09
300	31.09
360	31.09
420	31.11
480	31.09
540	31.12
600	31.10

plot\_ecutrho.gp

```
# plot_ecutrho.gp
# Gnuplot script to plot total Stress vs.
# ecutrho energy
# Set output to PNG image file
set terminal pngcairo size 800,600
set output « ecutrho-results.png"
set xlabel « Ecutrho (Ry) "
set ylabel "Total Stress (Ry/bohr**3)"
set style line 1 lc rgb "red" lw 2 pt 7
# Plot the data
plot « file.dat" using 1:2 with linespoints
linestyle 1 title " Ecutrho »
# Close the output file
set output
```

Hydrostatic pressure as a function of Ecutrho energy



Based on Hydrostatic convergence test, an ecutrho value of 300 Ry is appropriate for a convergence within ~1-2 kbar.

# K-point meshes Convergence test

ecut.\$ecutrho.in

```
&CONTROL
  calculation='scf',
  outdir='.',
  prefix='NaCl',
  pseudo_dir='.././../pseudo',
  verbosity='low',
  tprnfor=.true.,
  tstress=.true.,
/
&SYSTEM
  ibrav = 0
  A = 3.4
  nat = 2
  ntyp = 2
  ecutwfc =60,
  ecutrho = 300,
  input_dft='pbe',
  occupations='smearing',
  smearing='mv',
  degauss=0.005d0,
/
&ELECTRONS
  conv_thr=1d-8,
  mixing_beta=0.7d0,
/
&IONS
  ion_dynamics='bfgs',
/
&CELL
  cell_dynamics='bfgs',
  press=0.d0,
  press_conv_thr=0.5d0,
/
```

```
CELL_PARAMETERS {alat}
  1.0 0.0 0.0
  0.0 1.0 0.0
  0.0 0.0 1.0

ATOMIC_SPECIES
  Cl 35.45150 cl_pbe_v1.4.uspp.F.UPF
  Na 22.98900 na_pbe_v1.5.uspp.F.UPF

ATOMIC_POSITIONS {crystal}
  Cl 0.5 0.5 0.5
  Na 0.0 0.0 0.0

K_POINTS {automatic}
  ${mesh} ${mesh} ${mesh} 0 0 0
```

## Set K-points

for mesh in 4 5 6 7 8 9 10 11 12 13 14 15  
16 ; do

# K-point mesh Convergence test

We will use Gnuplot to make the plot. Here is the script code for plotting:

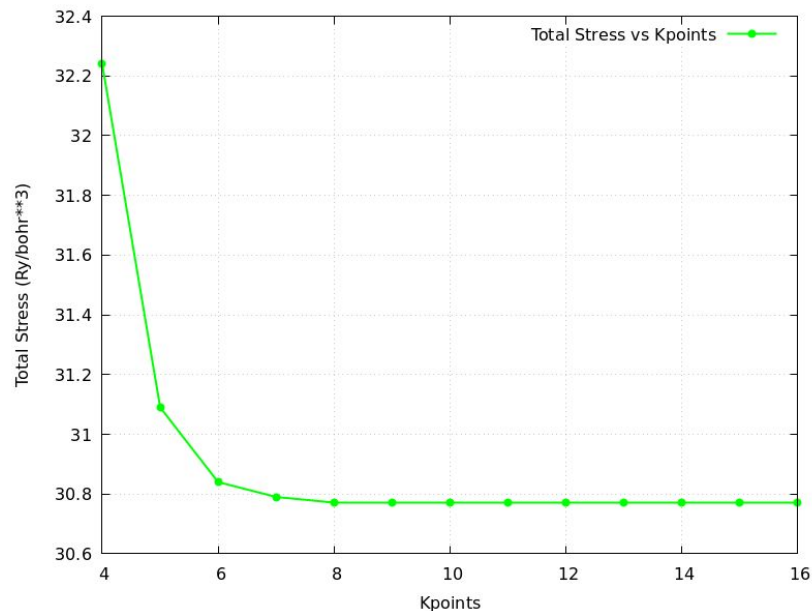
calc-kpoints.dat

4	32.24
5	31.09
6	30.84
7	30.79
8	30.77
9	30.77
10	30.77
11	30.77
12	30.77
13	30.77
14	30.77
15	30.77
16	30.77

plot\_kpoints.gp

```
1 # plot_press.gp
2 # Gnuplot script to plot
3
4 # Set output to PNG image file
5 set terminal pngcairo size 800,600
6 set output "kpoints_convergence.png" # Name of the output
   file
7
8 set xlabel "Kpoints"
9 set ylabel "Total Stress (Ry/bohr**3)"
10
11 set grid
12 set style line 1 lc rgb "green" lw 2 pt 7
13
14 # Plot the data
15 plot "calc-kpoints.dat" using 1:2 with linespoints
   linestyle 1 title "Total Stress vs Kpoints"
16
17 # Close the output file
18 set output
--
```

Hydrostatic pressure as a function of k-point mesh



Based on Hydrostatic convergence test, an kpoint mesh values of 9x9x9 is appropriate for a convergence within ~1-2 kbar.

# Optimization using vc-relax

## vc-relax.in

```
&CONTROL
calculation = 'vc-relax',
outdir      = '.',
prefix      = 'NaCl',
pseudo_dir  = '.././../pseudo',
verbosity   = 'low',
tprnfor     = .true.,
tstress     = .true.,
/
&SYSTEM
ibrav       = 0
A           = 3.4
nat         = 2
ntyp        = 2
ecutwfc     = 60,
ecutrho     = 300,
input_dft   = 'pbe',
occupations = 'smearing',
smearing    = 'mv',
degauss     = 0.005d0,
/
&ELECTRONS
conv_thr     = 1d-8,
mixing_beta  = 0.7d0,
/
&IONS
ion_dynamics = 'bfgs',
/
&CELL
cell_dynamics = 'bfgs',
press        = 0.d0,
press_conv_thr = 0.5d0,
/
```

```
CELL_PARAMETERS {alat}
1.0 0.0 0.0
0.0 1.0 0.0
0.0 0.0 1.0

ATOMIC_SPECIES
Cl 35.45150 cl_pbe_v1.4.uspp.F.UPF
Na 22.98900 na_pbe_v1.5.uspp.F.UPF

ATOMIC_POSITIONS {crystal}
Cl 0.5 0.5 0.5
Na 0.0 0.0 0.0

K_POINTS {automatic}
9 9 9 0 0 0
```

## vc-relax.out

negative rho (up, down): 1.454E-02 0.000E+00

Forces acting on atoms (cartesian axes, Ry/au):

atom	1	type	1	force =	0.00000000	0.00000000	0.00000000
atom	2	type	2	force =	0.00000000	0.00000000	0.00000000

Total force = 0.000000 Total SCF correction = 0.000000

Computing stress (Cartesian axis) and pressure

negative rho (up, down): 1.454E-02 0.000E+00					
	total	stress	(Ry/bohr**3)	(kbar)	P= 0.04
0.00000027	0.00000000	0.00000000	0.04	0.00	0.00
0.00000000	0.00000027	0.00000000	0.00	0.04	0.00
0.00000000	0.00000000	0.00000027	0.00	0.00	0.04

bfgs converged in 6 scf cycles and 5 bfgs steps  
(criteria: energy < 1.0E-04 Ry, force < 1.0E-03Ry/Bohr, cell < 5.0E-01kbar)

End of BFGS Geometry Optimization

Final enthalpy = -128.8814882949 Ry

Begin final coordinates

new unit-cell volume = 291.83131 a.u.^3 ( 43.24494 Ang^3 )  
density = 2.24403 g/cm^3

```
CELL_PARAMETERS (alat= 6.42506885)
1.032363974 0.000000000 0.000000000
0.000000000 1.032363974 0.000000000
0.000000000 0.000000000 1.032363974
```

ATOMIC\_POSITIONS (crystal)

```
Cl 0.500000000 0.500000000 0.500000000
Na 0.000000000 0.000000000 0.000000000
```

End final coordinates

### To run a DOS calculation

1. Perform self-consistent (SCF) calculation using **pw.x**
2. Perform non-self-consistent (NSCF) calculation using **pw.x**
  - use denser k-point mesh than in the SCF step
  - set occupations = 'tetrahedra' in the &SYSTEM card
3. Run **projwfc.x** to obtain the total and projected density of states (PDOS)

# Density of States (DOS)

## scf.in

```
&CONTROL
  calculation='scf',
  outdir='.',
  prefix='NaCl',
  pseudo_dir='/home/max/work/pseudo',
  verbosity='low',
  tprnfor=.true.,
  tstress=.true.,
/

&SYSTEM
 ibrav = 0
  celldm(1) = 6.42506885
  nat = 2
  ntyp = 2
  ecutwfc=60,
  ecutrho=300,
  input_dft='pbe',
  occupations='smearing',
  smearing='mv',
  degauss=0.005d0,
/

&ELECTRONS
  conv_thr=1d-08,
  mixing_beta=0.7d0,
/

CELL_PARAMETERS (alat)
  1.032363974  0.000000000  0.000000000
  0.000000000  1.032363974  0.000000000
  0.000000000  0.000000000  1.032363974

ATOMIC_POSITIONS (crystal)
Cl      0.500000000  0.500000000  0.500000000
Na      0.000000000  0.000000000  0.000000000

K_POINTS {automatic}
  9 9 0 0 0
```

pw.x < scf.in > scf.out

## nscf.in

```
&CONTROL
  calculation='nscf',
  outdir='.',
  prefix='NaCl',
  pseudo_dir='/home/max/work/pseudo',
  verbosity='low',
  tprnfor=.true.,
  tstress=.true.,
/

&SYSTEM
 ibrav = 0
  celldm(1) = 6.42506885
  nat = 2
  ntyp = 2
  ecutwfc=60,
  ecutrho=300,
  input_dft='pbe',
  occupations='tetrahedra',
/

&ELECTRONS
  conv_thr=1d-08,
  mixing_beta=0.7d0,
/

CELL_PARAMETERS (alat)
  1.032363974  0.000000000  0.000000000
  0.000000000  1.032363974  0.000000000
  0.000000000  0.000000000  1.032363974

ATOMIC_SPECIES
Cl  35.45150  cl_pbe_v1.4.uspp.F.UPF
Na  22.98900  na_pbe_v1.5.uspp.F.UPF
```

pw.x < nscf.in > nscf.out

```
ATOMIC_POSITIONS (crystal)
Cl      0.500000000  0.500000000  0.500000000
Na      0.000000000  0.000000000  0.000000000

K_POINTS {automatic}
  18 18 18 0 0 0
```

## projwfc.in (PDOS)

```
&PROJWFC
  prefix='NaCl',
  outdir='.',
  Emin=-6,
  Emax=17,
  DeltaE=0.05,
  filpdos='NaCl',
  filproj='NaCl'
/
```

projwfc.x < projwfc.in  
> projwfc.out



# Total and Partial Density of States (PDOS)

## Total Density of States

Output Files: total\_pdos

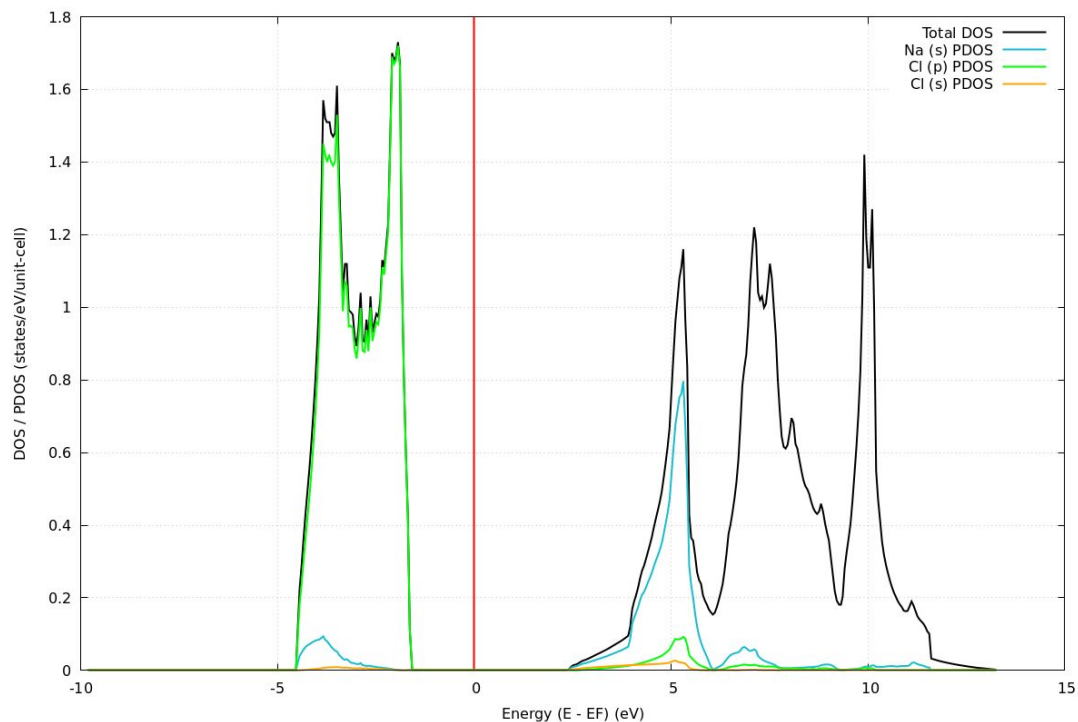
#	E (eV)	dos(E)	pdos(E)
	-6.000	0.000E+00	0.000E+00
	-5.950	0.000E+00	0.000E+00
	-5.900	0.000E+00	0.000E+00
	-5.850	0.000E+00	0.000E+00
	-5.800	0.000E+00	0.000E+00
	-5.750	0.000E+00	0.000E+00

## Partial Density of States

Output Files: NaCl.pdos\_atm#1(Na)\_wfc#2(s)  
NaCl.pdos\_atm#2(Cl)\_wfc#1(s)  
NaCl.pdos\_atm#2(Cl)\_wfc#2(p)

#	E (eV)	ldos(E)	pdos(E)	pdos(E)	pdos(E)
	-6.000	0.000E+00	0.000E+00	0.000E+00	0.000E+00
	-5.950	0.000E+00	0.000E+00	0.000E+00	0.000E+00
	-5.900	0.000E+00	0.000E+00	0.000E+00	0.000E+00
	-5.850	0.000E+00	0.000E+00	0.000E+00	0.000E+00
	-5.800	0.000E+00	0.000E+00	0.000E+00	0.000E+00
	-5.750	0.000E+00	0.000E+00	0.000E+00	0.000E+00
	-5.700	0.000E+00	0.000E+00	0.000E+00	0.000E+00

conda activate visualise  
Gnuplot plot\_bands.gnu



## To run the Band structure

1. Perform self-consistent (SCF) calculation using **pw.x**
2. Perform the non-self-consistent (NSCF) calculation using **pw.x**
  - use k-points along the high-symmetry path in the Brillouin zone
  - for insulators, occupations = 'tetrahedra' in the &SYSTEM card
3. Run **bands.x** to extract the band energies



# Band Structure

scf.in

```
&CONTROL
  calculation='scf',
  outdir='.',
  prefix='NaCl',
  pseudo_dir='/home/max/work/pseudo',
  verbosity='low',
  tprnfor=.true.,
  tstress=.true.,
/

&SYSTEM
  ibrav = 0
  celldm(1) = 6.42506885
  nat = 2
  ntyp = 2
  ecutwfc=60,
  ecutrho=300,
  input_dft='pbe',
  occupations='smearing',
  smearing='mv',
  degauss=0.005d0,
/

&ELECTRONS
  conv_thr=1d-08,
  mixing_beta=0.7d0,
/

CELL_PARAMETERS (alat)
  1.032363974    0.000000000    0.000000000
  0.000000000    1.032363974    0.000000000
  0.000000000    0.000000000    1.032363974

ATOMIC_POSITIONS (crystal)
Cl      0.500000000    0.500000000    0.500000000
Na      0.000000000    0.000000000    0.000000000

K_POINTS {automatic}
  9 9 0 0 0
```

pw.x < scf.in > scf.out

bands.in

```
&CONTROL
  calculation='bands',
  outdir='.',
  prefix='NaCl',
  pseudo_dir='/home/max/work/pseudo',
  verbosity='low',
  tprnfor=.true.,
  tstress=.true.,
/

&SYSTEM
  ibrav = 0
  celldm(1) = 6.42506885
  nat = 2
  ntyp = 2
  ecutwfc=60,
  ecutrho=300,
  input_dft='pbe',
  occupations='smearing',
  smearing='mv',
  degauss=0.005d0,
  nbnd = 12,
/

&ELECTRONS
  conv_thr=1d-08,
  mixing_beta=0.7d0,
/

CELL_PARAMETERS (alat)
  1.032363974    0.000000000    0.000000000
  0.000000000    1.032363974    0.000000000
  0.000000000    0.000000000    1.032363974
```

pw.x < bands.in > bands.out

```
ATOMIC_SPECIES
  Cl  35.45150  cl_pbe_v1.4.uspp.F.UPF
  Na  22.98900  na_pbe_v1.5.uspp.F.UPF

ATOMIC_POSITIONS (crystal)
Cl      0.500000000    0.500000000    0.500000000
Na      0.000000000    0.000000000    0.000000000

K_POINTS {crystal_b}
  8
  0.0000  0.0000  0.0000  30  ! G (Γ)
  0.0000  0.5000  0.0000  30  ! X
  0.5000  0.5000  0.0000  30  ! M
  0.0000  0.0000  0.0000  30  ! G (Γ)
  0.5000  0.5000  0.5000  30  ! R
  0.0000  0.5000  0.0000  30  ! X
  0.5000  0.5000  0.0000  30  ! M
  0.5000  0.5000  0.5000  30  ! R
```

pp.in

```
&BANDS
  outdir = '.'
  prefix = 'NaCl'
  filband = 'NaCl.dat'
/
```

bands.x < pp.in > pp.out

# Special k-points – seek path

Upload your QE input file to <https://www.materialscloud.org/work/tools/seekpath>

## SeeK-path: the k-path finder and visualizer

► What SeeK-path does

► SeeK-path definitions and advantages

Upload your structure

Upload a crystal structure:

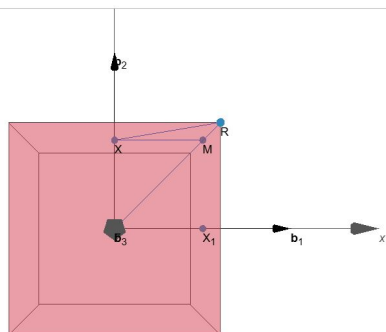
Choisir un fichier

Select here the file format:

Quantum ESPRESSO input [parser: qe-tools] ▼

By continuing, you agree with the terms of use of this service.

Calculate my structure



Supercell:

Camera:

Axes:  ▼

☒ bonds [\[?\]](#) ☐ atom labels [\[?\]](#)

☐ packed cell [\[?\]](#) ☐ space-filling [\[?\]](#)

☐ rotation

## Reciprocal space and Brillouin-zone information

Reciprocal cell vectors (1/Å)

b	x	y	z
b <sub>1</sub>	1.8479956786	0.0000000000	0.0000000000
b <sub>2</sub>	0.0000000000	1.8479956786	0.0000000000
b <sub>3</sub>	0.0000000000	0.0000000000	1.8479956786

Suggested path

Γ—X—M—Γ—R—X|R—M

High-symmetry points (scaled units)

Label	k <sub>1</sub>	k <sub>2</sub>	k <sub>3</sub>
Γ	0.0000000000	0.0000000000	0.0000000000
M	0.5000000000	0.5000000000	0.0000000000
R	0.5000000000	0.5000000000	0.5000000000
X	0.0000000000	0.5000000000	0.0000000000
X <sub>1</sub>	0.5000000000	0.0000000000	0.0000000000

# Band Structure

## SCF file output:

the Fermi energy is 3.7918 eV

## NSCF file output:

```
0.0000 -48.0156
0.0161 -48.0156
0.0323 -48.0156
0.0484 -48.0156
0.0646 -48.0155
0.0807 -48.0155
.....
```

## Postprocessing:

```
Reading collected, re-writing distributed wavefunctions
high-symmetry point: 0.0000 0.0000 0.0000 x coordinate 0.0000
high-symmetry point: 0.0000 0.4843 0.0000 x coordinate 0.4843
high-symmetry point: 0.4843 0.4843 0.0000 x coordinate 0.9687
high-symmetry point: 0.0000 0.0000 0.0000 x coordinate 1.6536
high-symmetry point: 0.4843 0.4843 0.4843 x coordinate 2.4925
high-symmetry point: 0.0000 0.4843 0.0000 x coordinate 3.1774
high-symmetry point: 0.4843 0.4843 0.0000 x coordinate 3.6617
high-symmetry point: 0.4843 0.4843 0.4843 x coordinate 4.1461
```

## Gnuplot plot\_bands.gnu

