

Structure of the QE [Input File](#)

Quantum Espresso's input file consists of 3 major sections (&control, &system, &electrons) and 2 optional sections (&ions, &cell) depending on the type of calculation to be performed. Sections must appear in the order given below,

```
&CONTROL
&SYSTEM
&ELECTRONS
&IONS
&CELL
```

A “/” indicates the end of a section.

After listing all required sections, additional information about your molecule or the crystal system has to be specified such as

```
ATOMIC_SPECIES
ATOMIC_POSITIONS
K_POINTS
CELL_PARAMETERS
```

For a full list of sections and other options refer to the [QE website](#).

Find below an example input file for a self-consistent calculation of single layer MoS₂.

Example QE Input File for Monolayer MoS₂

```
1 &control
2  calculation = 'scf',
3  restart mode = 'from_scratch',
4  pseudo\_dir = './',
5  prefix = 'MoS2',
6  outdir = './',
7  nstep = 100,
8 /
9
10 &system
11 ibrav = 4,
12 a = 3.16000e+00,
13 c = 2.00000e+01,
14 nat = 3,
15 ntyp = 2,
16 ecutwfc = 60,
17 occupations = 'fixed',
18 smearing = 'gauss',
19 vdw corr = 'DFT-D',
20 /
21
22 &electrons
23 conv thr = 1.0d-9,
24 /
25
26
27 ATOMIC SPECIES
28 Mo 95.94000 Mo_ONCV_PBE-1.0.upf
29 S 32.06600 S_ONCV_PBE-1.0.upf
30
31 ATOMIC POSITIONS (angstrom)
32 S 0.002804304 0.001613133 8.429890591
33 S 0.002804304 0.001613133 11.570109409
34 Mo 1.582007393 0.913365735 10.000000000
35
36 K POINTS automatic
37 51 51 1 0 0 0
```

For further details please click on the highlighted input parameter.

In line 2 of the input file, we list the type of calculation we ask to perform. Available types of calculations include,

- *SCF* - Single point energy calculation without changing atomic coordinates or the cell parameters, i.e., no structure relaxation.
- *NSCF* - Non-self-consistency calculation to extract the desired properties such as the DOS in case you need a denser electronic momentum mesh (k-mesh), based on the previous *SCF* calculation.
- *BANDS* - Calculation of electronic bands along a path of high-symmetry k-points based on the previous *SCF* calculation.
- *RELAX* - Optimization of the atomic coordinates to minimize the forces while keeping the cell fixed. This calculation requires section &ions. All options for a single *SCF* calculation apply, plus a few others.
- *VC-RELAX* - Optimization of both atomic coordinates and the cell parameters. Structure relaxation is important to minimize forces acting on the atoms as well as to reduce the stress in the cell. This calculation requires section &cell.

Important Convergence Parameters

Accurate DFT calculations require specific parameters for the material of interest and their convergence. These parameters for any type of DFT calculation include the planewave kinetic energy cutoff (*ecutwfc*), the Brillouin zone sampling [k-points](#), and the self-consistency energy convergence threshold (*conv_thr*). To obtain accurate results, one should optimize these parameters with energy minimization.

Please see the following links below for a hands-on exercise to better understand these parameters and other related calculations.