

# RUNNING CP2K CALCULATIONS

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# Overview

- How to run CP2K
- CP2K Input file
  - The Basics
  - The How – FORCE\_EVAL
  - The What – MOTION
- Basis Sets and Pseudopotential libraries
- CP2K Output
  - Controlling what gets written
  - Overview of an output file
- Restarting a calculation



# How to run CP2K

- CP2K binaries:
  - cp2k.version where version is usually one of:
    - sopt – Serial, optimised
    - ssmp – Single process + symmetric multiprocessor (OpenMP)
    - popt – Parallel (MPI), optimised
    - psmp – Parallel (MPI) + symmetric multiprocessor (OpenMP)
- Available from <http://www.cp2k.org/download>
  - Linux binaries (released versions)
  - Also in Linux package managers
  - Source code (released versions and latest trunk), GPL
  - May be pre-installed, e.g. NSCCS, ARCHER ...



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# How to run CP2K

- Basic command line options:

- `cp2k.sopt -i input_file -o output_file`
  - By default, output goes to the standard output
  - Output to file appends (beware!)
  - Input file is the last argument if not otherwise specified

- Other useful options:

- `cp2k.sopt --version`
- `cp2k.sopt --check input_file`
- `cp2k.sopt --html-manual`
- `cp2k.sopt --help`



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# How to run CP2K

- Typical files associated with a CP2K run:
  - Input (required):
    - e.g. H2O-32.inp (main input file, name and extension are arbitrary)
  - Optional inputs:
    - POTENTIAL (psuedopotential library)
    - BASIS\_SET (basis set library)
    - Structure file (e.g. psf, xyz, crd ...)
    - ...
  - Outputs:
    - PROJECT-1.restart (input file to restart calculation)
    - PROJECT-pos-1.xyz (trajectory for MD or GEO\_OPT)
    - PROJECT-1.ener (MD energies, temperature, cons. Q ...)
    - PROJECT-1.cell (cell parameters for NPT MD or CELL\_OPT)
    - PROJECT-RESTART.wfn (orbitals for restart)



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# CP2K Input file: The Basics

- Full documentation available online:
  - <http://manual.cp2k.org>
  - Or generate with --html-manual
- Sections – 13 (optional) top level sections

```
&BEGIN section_name [params]  
...  
&END [section_name]
```

- Keywords
  - KEYWORD value
  - KEYWORD [ON|OFF] [YES|NO] [TRUE|FALSE] ...
  - KEYWORD
- Nesting
  - Sections may others sections and keywords



# CP2K Input file: The Basics

- Basic pre-processing syntax

@INCLUDE 'filename'	– copy in text from file
@SET VAR value	– define a variable
\$VAR	– replaced with variable value
@IF / @ENDIF	– simple logic
! or #	– comments

- Units

- Numerical entries have a default unit (see manual)
- Specify other units by hand e.g.

ABC [nm] 100 100 100 (or bohr, default is angstrom)

EMAX\_SPLINE [eV] 50 (or Ry, joule, default is hartree)

- Also combinations e.g. [hartree\*bohr^-2]



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# CP2K Input file: The Basics

- GLOBAL section (required)

```
&GLOBAL
```

```
    PROJECT H2O-32
```

```
    RUN_TYPE MD
```

```
    PRINT_LEVEL HIGH
```

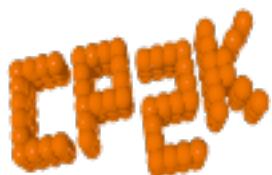
```
&TIMINGS
```

```
    THRESHOLD 0.000001
```

```
&END
```

```
WALLTIME 3600
```

```
&END GLOBAL
```



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# CP2K Input file: The How

- FORCE\_EVAL section (required)

```
&FORCE_EVAL
    METHOD QS    (or FIST, QMMM ...)
    &DFT
    ...
    &END DFT
    &SUBSYS
    ...
    &END SUBSYS
&END FORCE_EVAL
```



# CP2K Input file: The How

```
&DFT
  BASIS_SET_FILE_NAME GTH_BASIS_SETS
  POTENTIAL_FILE_NAME POTENTIAL
  &MGRID
    CUTOFF 280
    REL_CUTOFF 30
  &END MGRID
  &QS
    EPS_DEFAULT 1.0E-12
    WF_INTERPOLATION PS
    EXTRAPOLATION_ORDER 3
  &END QS
  &SCF
    SCF_GUESS ATOMIC
    &OT ON
      MINIMIZER DIIS
    &END OT
    &PRINT
      &RESTART OFF
    &END
  &END SCF
  &XC
    &XC_FUNCTIONAL Pade
  &END XC_FUNCTIONAL
  &END XC
&END DFT
```



Basis and PP library files

Parameters for the realspace multi-grids

Quickstep options

Control of SCF procedure, including minimisation scheme

Exchange-Correlation Functional (LDA)

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# CP2K Input file: The How

```
&SUBSYS
  &CELL
    ABC 9.8528 9.8528 9.8528
  &END CELL
  # 32 H2O (TIP5P,1bar,300K) a = 9.8528
  &COORD
    O      2.280398      9.146539      5.088696
    O      1.251703      2.406261      7.769908
    O      1.596302      6.920128      0.656695
    ...
    H      0.837635      8.186808      8.987268
    H      8.314696     10.115534      2.212519
    H      8.687134      8.667252      2.448452
  &END COORD
  &KIND H
    BASIS_SET TZV2P-GTH
    POTENTIAL GTH-PADE-q1
  &END KIND
  &KIND O
    BASIS_SET TZV2P-GTH
    POTENTIAL GTH-PADE-q6
  &END KIND
&END SUBSYS
```



Cell definition

Particle coordinates

Could also @include an external file or parse other formats via

```
&TOPOLOGY
  COORD_FILE_NAME
&END TOPOLOGY
```

Definitions of atomic kinds

Could specify charge, mass ...

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# CP2K Input file: The What

- MOTION section

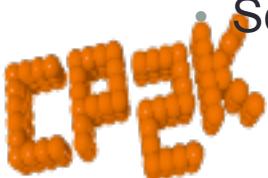
```
&MOTION  
  &MD  
    ENSEMBLE NVE  
    STEPS 10  
    TIMESTEP 0.5  
    TEMPERATURE 300.0  
  &END MD  
&END MOTION
```

- Also used to control Geometry Optimisation, NEB, Monte Carlo, ...



# Basis Sets and PP libraries

- CP2K uses Goedecker-Teter-Hutter, seperable Pseudopotentials
  - Several sets of PPs and corresponding optimised basis sets are available
  - See `cp2k/data` or online:  
<http://sourceforge.net/p/cp2k/code/HEAD/tree/trunk/cp2k/data>
- POTENTIAL, GTH\_POTENTIALS
  - Wide range of PPs for at many elements - LDA (PADE), PBE, BLYP ...
- BASIS\_SET, GTH\_BASIS\_SET, BASIS\_MOLOPT
  - Various qualities / size of basis
  - Make sure Basis and PP match (functional and number of electrons)
  - Some documentation and references at head of each file



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# CP2K Output: Controlling what gets written

- The PRINT\_LEVEL keyword in &GLOBAL
  - SILENT, LOW, MEDIUM (default), HIGH, DEBUG
  - HIGH can give more information if you are interested
    - Also gives some per-process logging in parallel jobs
  - For long MD runs (e.g. classical), recommend using LOW
- Fine grained control is available via print-keys
  - Most input sections contain a &PRINT sub-section
  - Each &PRINT sub-section has further subsections for each quantity that may be printed

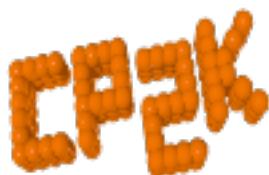


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# CP2K Output: Controlling what gets written

- For example, the &PRINT section in &MOTION contains
  - &CELL
  - &FORCES
  - &TRAJECTORY
  - &VELOCITIES
  - ...
- Each section has parameters (and defaults) for which print level it is output
  - &TRAJECTORY defaults to LOW
  - &VELOCITIES defaults to HIGH



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# CP2K Output: Controlling what gets written

- Can also specify frequency of printing via &EACH subsection e.g.

```
&PRINT  
  &CELL  
    &EACH  
      MD 100  
    &END EACH  
  &END CELL  
&END PRINT
```

- Control over filenames, file formats etc. at each &PRINT section

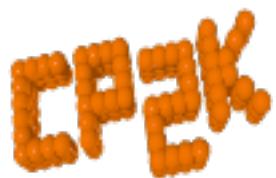


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# CP2K Output: Overview of an output file

...



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# Restarting a calculation

- If you need to restart your job...
  - Hardware failure
  - Batch system time limit
  - Need more MD sampling
  - ...
- CP2K dumps a restart input file which can be directly re-run
  - cp2k.sopt -i PROJECT-1.restart
  - Continuous numbering of MD steps
  - Stores all state variables (incl. extended system)
  - Use SCF\_GUESS RESTART

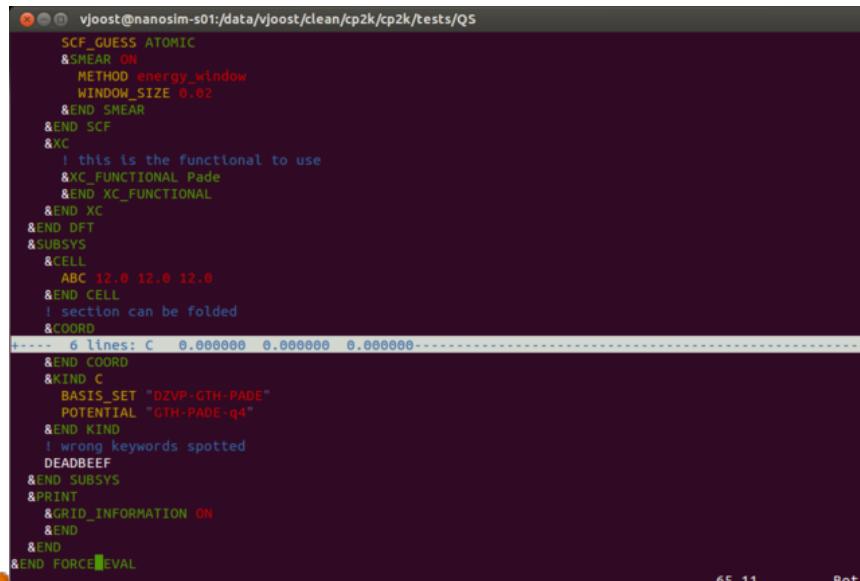


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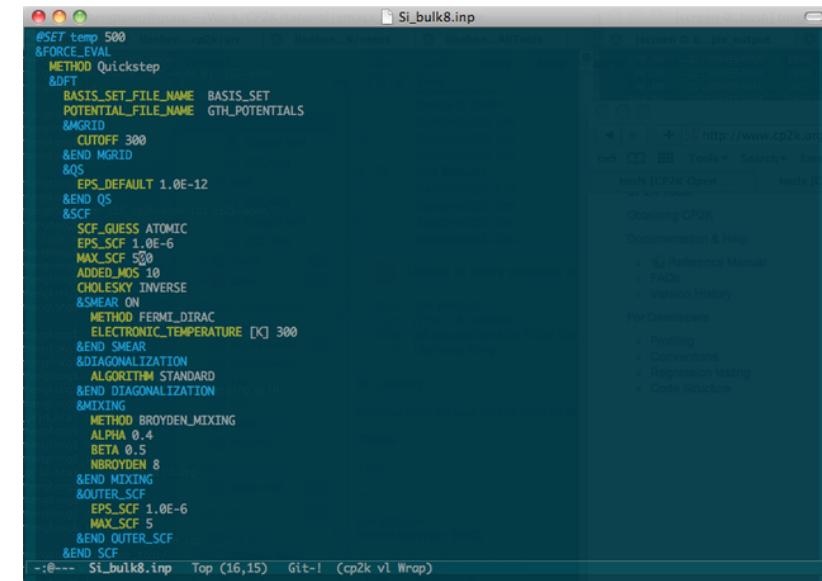


# Tools for building CP2K input

- Plugins are available for your favourite(!) text editors:
  - <https://www.cp2k.org/tools:vim>
  - <https://www.cp2k.org/tools:emacs>
  - Syntax highlighting, indentation, show/hide sections, keywords



```
SCF_GUESS ATOMIC
&SMEAR ON
    METHOD energy_window
    WINDOW_SIZE 0.02
&END SMEAR
&END SCF
&XC
    ! this is the functional to use
    &XC_FUNCTIONAL Pade
    &END XC_FUNCTIONAL
&END XC
&END DFT
&SUBSYS
    &CELL
        ABC 12.0 12.0 12.0
    &END CELL
    ! section can be folded
    &COORD
        6 lines: C  0.000000  0.000000  0.000000-
    &END COORD
    &KIND C
        BASIS_SET "DZVP-GTH-PADE"
        POTENTIAL "GTH-PADE-q4"
    &END KIND
    ! wrong keywords spotted
    DEADBEEF
&END SUBSYS
&PRINT
    &GRID_INFORMATION ON
    &END
&END
&END FORCE_EVAL
```



```
@SET temp 500
&FORCE_EVAL
    METHOD Quickstep
    &DFT
        BASIS_SET_FILE_NAME BASIS_SET
        POTENTIAL_FILE_NAME GTH_POTENTIALS
    &MGRID
        CUTOFF 300
    &END MGRID
    &QS
        EPS_DEFAULT 1.0E-12
    &END QS
    &SCF
        SCF_GUESS ATOMIC
        EPS_SCF 1.0E-6
        MAX_SCF 500
        ADDED_MOS 10
        CHOLESKY_INVERSE
    &SMEAR ON
        METHOD FERMI_DIRAC
        ELECTRONIC_TEMPERATURE [K] 300
    &END SMEAR
    &DIAGONALIZATION
        ALGORITHM STANDARD
    &END DIAGONALIZATION
    &MIXING
        METHOD BROYDEN_MIXING
        ALPHA 0.4
        BETA 0.5
        NBROYDEN 8
    &END MIXING
    &OUTER_SCF
        EPS_SCF 1.0E-6
        MAX_SCF 5
    &END OUTER_SCF
    &END SCF
```

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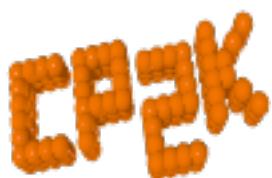
# Tools for building CP2K input

- Python interfaces
  - Atomic Simulation Environment (ASE) – see tomorrow
  - PyCP2K (<https://github.com/SINGROUP/pycp2k>)
    - Object-oriented wrapper following the CP2K input format
    - Auto-completion (for e.g. Spyder IDE)
    - E.g. GLOBAL%RUN\_TYPE is GLOBAL.Run\_type
    - May use ASE for execution

```
23 ##### Write the simulation input #####
24 GLOBAL.Run_type = "ENERGY_FORCE"
25 GLOBAL.Print_level = "LOW"
26 FORCE_EVAL.Method = "Quickstep"
27 FORCE_EVAL.PRINT.FORCEES.Section_parameters = "ON"
28 DFT.Basis_set_file_name = "BASIS_SET"
29 DFT.Potential_file_name = "POTENTIAL"
30 DFT.QS.Eps_default = 1.0E-10
31 SCF.Scf_guess = "ATOMIC"
32 SCF.Eps_scf = 1.0E-7
33 SCF.Max_scf = 300
34 SCF.DIAGONALIZATION.Section_parameters = "ON"
35 SCF.DIAGONALIZATION.Algorithm = "STANDARD"
36 SCF.MIXING.Section_parameters = "T"
37 SCF.MIXING.Method = "BROYDEN_MIXING"
38 SCF.MIXING.Alpha = 0.4
39 SCF.MIXING.Nbroyden = 8
40 SCF.MIXING.Nbroyden = 8
41 KIND = SUBSYS.KIND.add("st") # Section_parameters can be provided as argument.
42 KIND.Basis_set = "DZVP-GTH-PADE"
43 KIND.Potential = "GTH-PADE-q4"
44 calc.create_cell(SUBSYS, lattice)
45 calc.create_coord(SUBSYS, lattice)
46
47 ##### Run the simulation #####
48 calc.run()
~ 3 fewer lines
```

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# Tools for building CP2K input

- CP2K Input Editor
  - Browser-based:
    - <http://cp2k-www.epcc.ed.ac.uk/cp2k-input-editor>
  - Show/hide sections
  - Dropdowns for option selection
  - Tooltip help on every keyword
  - Basic input validation
  - Library of example input template files
  - Supports CP2K releases 2.5-2.7, 3.0, 4.0



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CP2K Input Editor Home Edit Help About

Input Templates:

- Energy\_Forces\_Si\_bulk8
- Geometry\_Optimisation\_H2O

Load CP2K input file

Edit input:

Expand Collapse Show Inactive Hide Inactive

CP2K

- GLOBAL
- MOTION
- FORCE\_EVAL
  - METHOD QUICKSTEP
  - DFT
    - BASIS\_SET\_FILE\_NAME BASIS\_SET
    - POTENTIAL\_FILE\_NAME GTH\_POTENTIALS
  - SCF
    - MAX\_SCF 300
    - EPS\_SCF 1.0E-7
    - SCF\_GUESS ATOMIC
    - DIAGONALIZATION
      - SECTION\_PARAMETERS T
      - ALGORITHM STANDARD
    - MIXING

CP2K

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# Running a CP2K Calculation

Questions?



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