SETTING UP A CP2K CALCULATION

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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

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How to run CP2K

- CP2K binaries:
  - `cp2k.version` where version =
    - sopt – Serial, optimised
    - 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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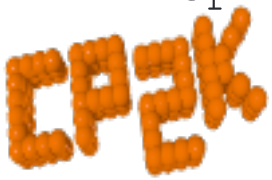
How to run CP2K

- 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)
    - `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

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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
```

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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
  &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, …

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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` sub-section 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

…

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)
  - May want to use `SCF_GUESS RESTART`

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Any questions?


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