Instructions for compiling and running ONETEP on Michael

1. Setting up the environment

You will need to load the correct set of modules. The provided config file is guaranteed to work with the Intel compiler v19.1.0.166 (curiously, known as compilers/intel/2020) and Intel MPI 2019 Update 6. To do this, add the following lines verbatim to the end of your ~/.bashrc file:

Then log out and log back in again for the changes to take effect. The first line takes care to unload all the modules so that they do not interfere with the ones we're using. Subsequent lines load the required modules – some prerequisites, git, the Intel compiler and Intel MPI.

2. Compilation

Use the config file conf.michael. Issue the command below while in the directory of your ONETEP installation:

```
make onetep ARCH=michael
```

This should compile and link a working binary, which will be placed in the bin subdirectory. If you tend to do this more often you might like the following command more:

```
make onetep ARCH=michael -j | utils/colorise
```

It compiles many files in parallel (so it's faster) and colorises output to make it more readable.

3. Testing

If you have just compiled a new binary for Michael, it would be prudent to test it by running a suite of quality-checks ("QCs") bundled with ONETEP. These test a number of important functionalities by running short jobs and comparing results against known good values. To run the QC-test suite, copy the QC-test submission script qcsubmit.michael.2021 (provided in the directory hpc_resources/Michael of your ONETEP installation) to the tests directory of your ONETEP installation. Then submit it to the batch system by changing to the tests directory and issuing

```
qsub jobsubmit.michael.2021
```

There is no need to edit this file, unless you are using a budget (allocation) that is not Faraday_MSM_sky. If so, adjust the line marked with [*]. The tests should complete within two hours of starting. Once the testing job starts, you should see, in the same directory, a file called

QC_tests.onnnnn.out, where nnnnn is the ID (number) of your job. This contains the log of how your job is running – diagnostic messages from the batch system and the submission script, and any output from testcode – the python script that actually runs the tests. If everything goes well, you should see subsequent test numbers (there are about 85 of them), followed by "Passed" or, occasionally "WARNING". The warnings can usually be ignored. If there are any errors, the test number will be followed by "FAILED". If this happens, go to the directory of this particular test and examine the test.out.*, test.err.* and (potentially) *.error_message files. They should give you an idea about what went wrong. Once the tests complete, you will see a file called *DONE in your tests directory. This does not necessarily mean that all tests completed successfully. You should examine the QC tests-onnnn.out file for a line like this:

```
All done. 105 out of 105 tests passed (12 warnings).
```

If all tests passed – you're good to go.

4. Running

Use the provided submission script: jobsubmit.michael.2021. Place it in a directory where you have input files. Make sure you only run one calculation in a directory. Do not run multiple calculations (multiple . dat files) in a single directory – this will lead to a mess and the provided script disallows it.

Adjust the script to your liking by editing its first lines (see next page). Submit it with: qsub jobsubmit.michael.2021. Details on how to monitor and control submitted and running jobs can be found on the Michael support web page at https://www.rc.ucl.ac.uk/docs/Clusters/Michael/

Things to adjust:

- 1. The name of your job. It has no effect on how it's run, but it will help you distinguish it from any other jobs you might have.
- 2. Priority of the job.
 - a. Use 'Gold' to submit a paid job that will take Gold from a particular project budget.
 - b. Use 'Free' to submit a free job that will not use up any Gold.
 - c. Use 'Test' to submit testing jobs that will not use up any Gold, and will have higher priority than normal free jobs, but are limited to 2 nodes (48 cores) and 1 hour of walltime.
- 3. Name of the allocated project. All jobs need to specify what project they belong to, whether they are paid or free. To see the name of your project(s) and how much allocation that budget has, run the command budgets.
- 4. Type of the node. We strongly suggest you always specify which type of node you intend your job to run on, to avoid unintentionally wasting cores if your total number does not cleanly fit on that node size. The old nodes are K-type (24 cores) while the new nodes with hyperthreading are A-type (40 cores). Jobs never run across a mix of node types it will be all K nodes or all A nodes.
- 5. MPI/OMP architecture. Specify the *total* number of **CPU cores** (MPI processes × OMP threads) you want to use, summed over all nodes. Also specify the number of OpenMP *threads* spawned from each MPI process. I suggest using 4. The parallel runner script on Michael (gerun) will figure out the correct number of MPI tasks (processes), both per node and in total.
- 6. RAM in integer number of Gigabytes (G) per core. Max RAM per node is 128G. So If you ask for 128G RAM and 24 cores, that will actually run on 24 nodes using only one core per node. This

- allows you to have sparse process placement when you do actually need that much RAM per process.
- 7. TMPDIR space per node (default is 10G).
- 8. The walltime you are asking for in hh:mm:ss. The maximum on Michael is 48:00:00. Don't exceed this value or your job will never start. Normally using lower values allows faster job turnaround the scheduler is able to stick short jobs into the allocation more easily.
- 9. Path to the onetep_launcher script. Change it to the location of onetep_launcher. It is located in the utils subdirectory of the ONETEP installation. The script should be able to find the correct ONETEP executable automatically (e.g. by examining . . /bin, relative to the utils directory it itself is in).

```
# -----
# Edit this file according to your job requirements.
# Copy it to the job directory, change to that directory and submit it by issuing:
# qsub jobsubmit.michael.2021
# -----
#1 Give a name to the job
#$ -N jobname
#2 Select priority
#$ -P Gold #select a paid 'Gold' job or 'Free' job or 'Test' job (1hr on 48 cores)
#3 Mention allocated project name
#$ -A Faraday MSM sky #Adjust this according to name of the allocated project
#4 Select the type of node to run the job (K: 24-core node or A: 40-core node)
#$ -ac allow=K
#5 Specify parallel mpi/omp architecture
# E.g. requesting 2 K-type nodes (48 cores) with 4 OMP threads per MPI process
#$ -pe mpi 48
                  #Specify total number of CPU cores for the job
threads_per_process=4 #Specify number of OMP threads spawned from each MPI process
#6 Request RAM per core (must be an integer)
#$ -1 mem=1G
#7 Request TMPDIR space per node (default is 10 GB)
#$ -1 tmpfs=10G
#8 Request walltime
#$ -1 h_rt=24:00:00
#9 Path to ONETEP launcher
onetep_launcher="$HOME/Scratch/onetep_ab/utils/onetep_launcher"
#-----You need not change anything below this line------
```

Specifications from the file ~/hpc resources/Michael/jobsubmit.michael.2021

Once you adjusted the above, you are ready to submit your job. The submission script, together with onetep_launcher, will take of everything. There is no need for you to set the stack size,

OMP_STACKSIZE, or OMP_NUM_THREADS. Also, do not add any threads_ keywords to your input file. It's all handled by the script, really.

The submission script will try to communicate any error or success conditions by creating files with filenames beginning with % (so they are easy to spot in the file listing) in the directory of your run. These include:

%NO DAT FILE – there are no . dat files in your job directory. Job did not start.

%MORE_THAN_ONE_DAT_FILE — there are multiple . dat files in your job directory. Job did not start. This script only works correctly in a one-dat-file-per-run set-up.

***ONETEP_LAUNCHER_MISSING** — the path you provided does not point to a valid onetep launcher. It either does not exist or is not executable. Job did not start.

%GERUN_ERROR – gerun, the MPI wrapper script on Michael reported an error. You job did not start or failed. Examine the standard error file for any error messages.

***ONETEP_ERROR_DETECTED** – ONETEP failed gracefully, producing an error message. Job ran, but failed. Examine the .error message file to see what happened.

***ONETEP_DID_NOT_COMPLETE** – Job likely started, but it doesn't look like it's completed, even though there's no error message. It's likely your job deadlocked (got stuck), crashed non-gracefully or ran out of walltime. Examine the standard error file for any error messages and the ONETEP output for any hints on what might have happened.

%DONE – your calculation ran to completion with no apparent errors.

5. Crucial files.

Assuming your ONETEP input is called myinput. dat, these are some important files in your run.

- myinput.out: your ONETEP output.
- myinput.err: the standard error output of your job. On Michael, once the job starts, it is immediately polluted by diagnostic information from gerun, such as the number of MPI processes (tasks) and OMP threads used, the names of the machines (nodes) on which your job is running and the mpirun command that gerun came up with to start your job. This will be all, if everything goes well. Otherwise, this file will also contain error messages from the OS, the batch system, gerun, MPI or the Fortran RTL. You might wish to examine this file in case of any issues.
- myinput.error_message: if ONETEP fails gracefully, this will contain an error message.
- jobname.onnnn and jobname.ponnnn, where jobname is the name of your job and nnnn is the ID (number) of your job. This contains the log of how your job ran diagnostic messages from the batch system and the submission script.
- jobname.ennnn and jobname.pennnn, where jobname is the name of your job and nnnn is the ID (number) of your job. This contains any errors or messages of successful completion from the batch system and the submisstion script.
- \$modules_loaded: The list of modules actually loaded when the job is run is echoed here. You might want to examine it if you have any doubts if they match your expectations.