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Job Manager for local execution of QuantumATK scripts

Job Manager for local execution of QuantumATK scripts

Version: 2017.0

Downloads & Links

- [PDF version](#)
- [PDF version for QuantumATK 2016](#)
- [silicon.py](#)
- [cnt.py](#)
- [mpi_check.py](#)

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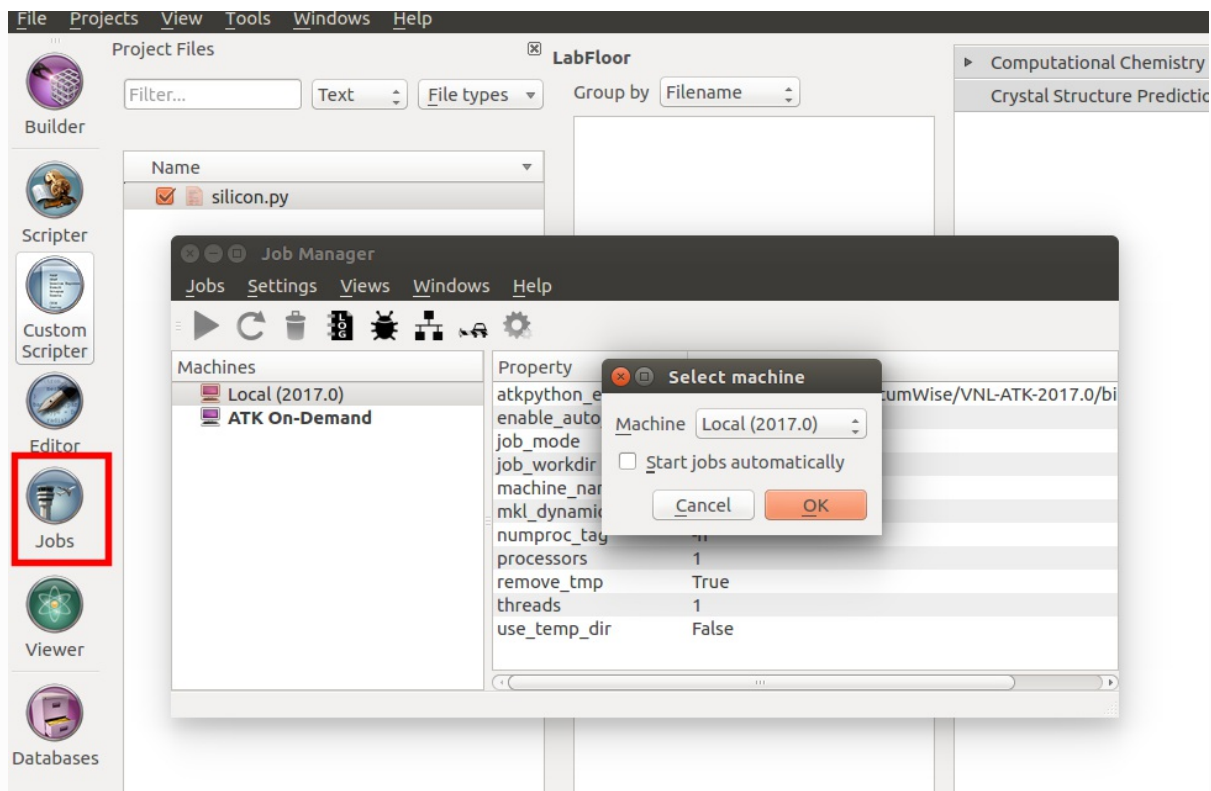



Execute QuantumATK simulations via the Job Manager

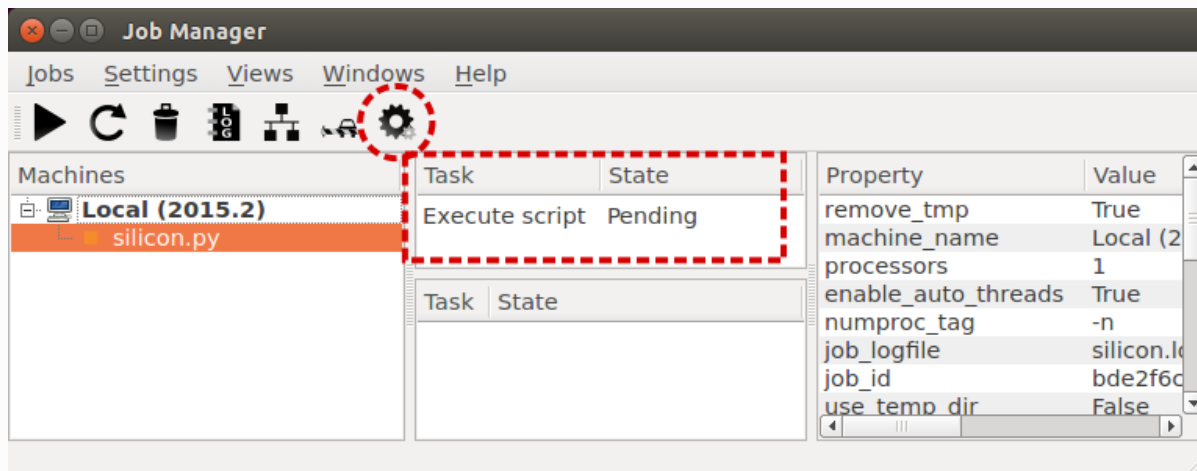
In this section you will learn how to use the **Job Manager** for local execution of QuantumATK scripts. Specifically, you will learn about queuing, running and managing QuantumATK jobs.

Create a new empty project and download the example script [silicon.py](#), which runs an ATK-DFT calculation with very many k-points (31x31x31).

Drop it on the  **Job Manager** and select a local machine for the job execution.



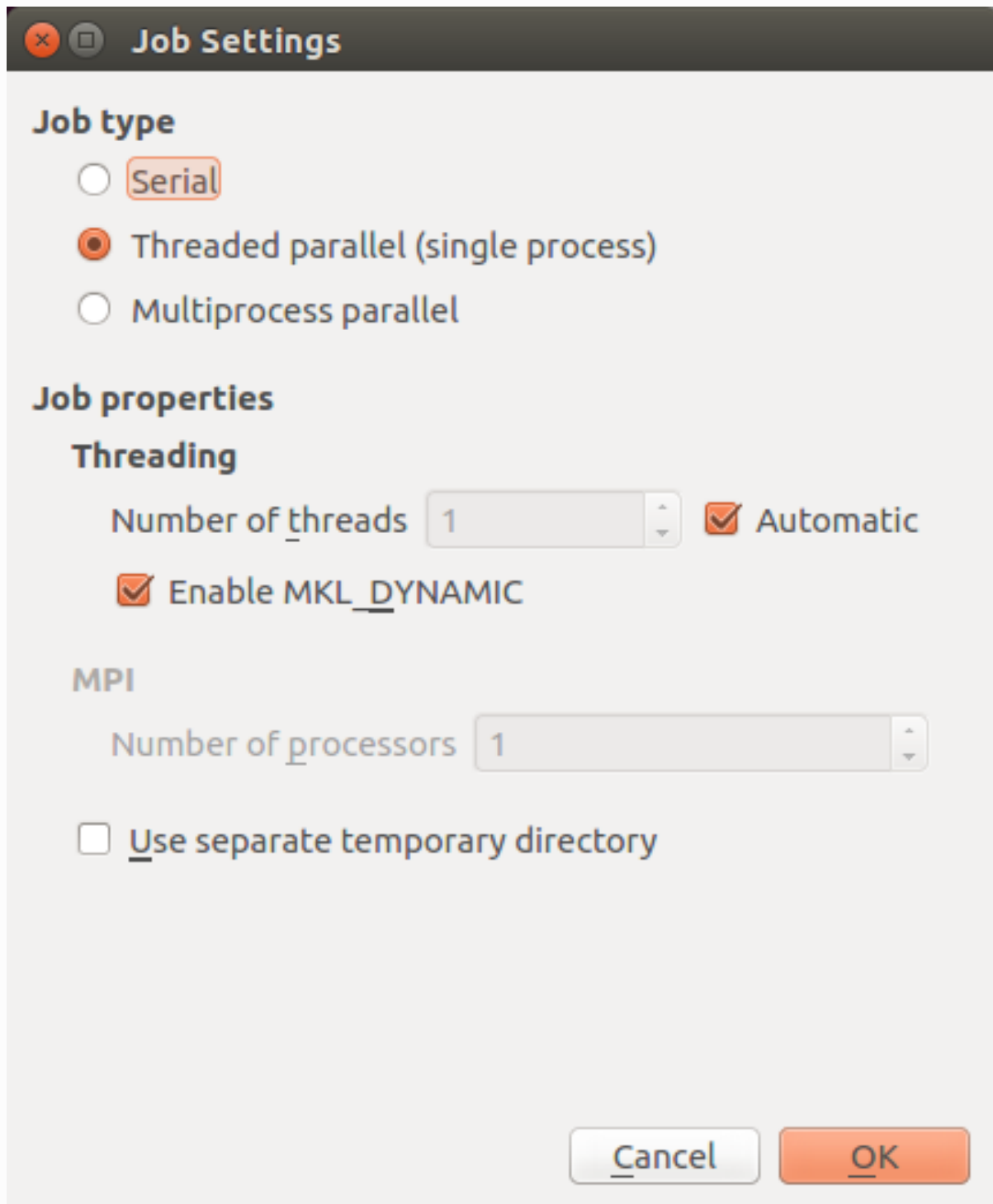
The job is now in the task state “pending” with “Threaded parallel (Single process)” default settings. Click the Job Settings  icon to edit the job settings.



The Job Settings widget has three basic panels:

- **Job type;**
 - Serial
 - Threaded parallel (single process)
 - Multiprocess parallel
- **Job properties;**
 - Threading
 - MPI
- **Use separate temporary directory** (set this to allow using a non-default working directory for running the job. Note, results will not be appended to existing hdf5 or nc files, but instead a new file

will be created.)



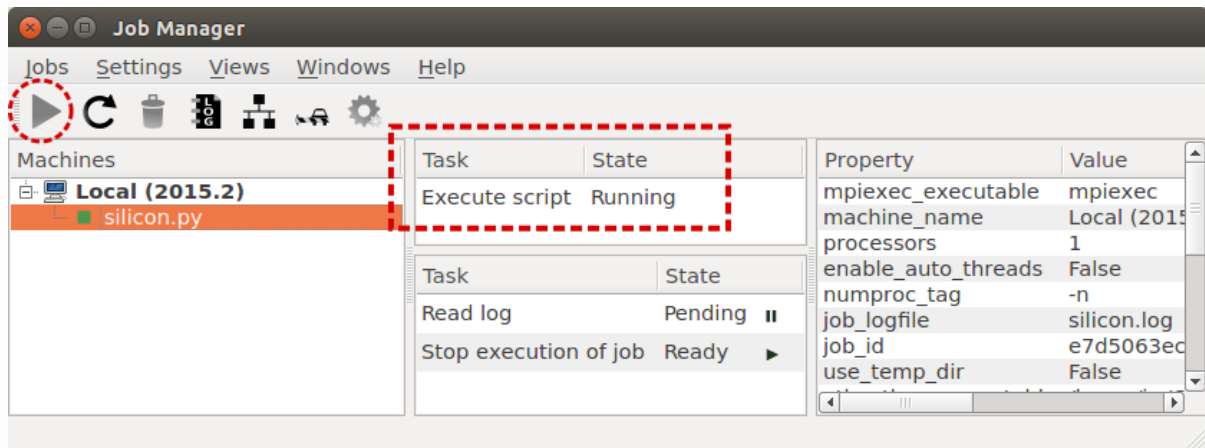
The image shows a 'Job Settings' dialog box with a dark header bar containing a close button (x), a maximize button (square), and the title 'Job Settings'. The main area is light gray and contains the following sections:


- Job type**
 - Serial
 - Threaded parallel (single process)
 - Multiprocess parallel
- Job properties**
 - Threading**
 - Number of threads: 1 (spin box) Automatic
 - Enable MKL_DYNAMIC
 - MPI**
 - Number of processors: 1 (spin box)
 - Use separate temporary directory

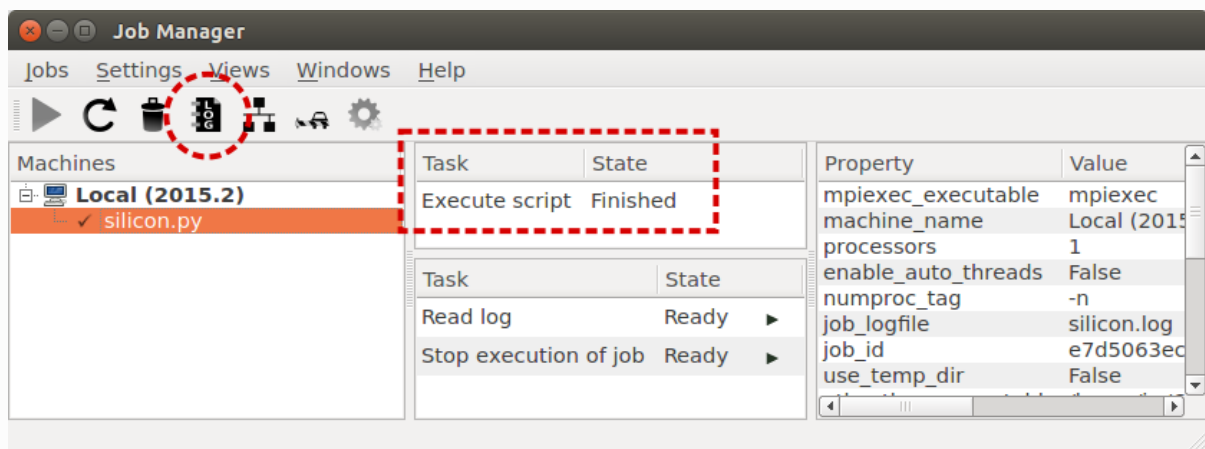
At the bottom right, there are two buttons: 'Cancel' (light gray) and 'OK' (orange).

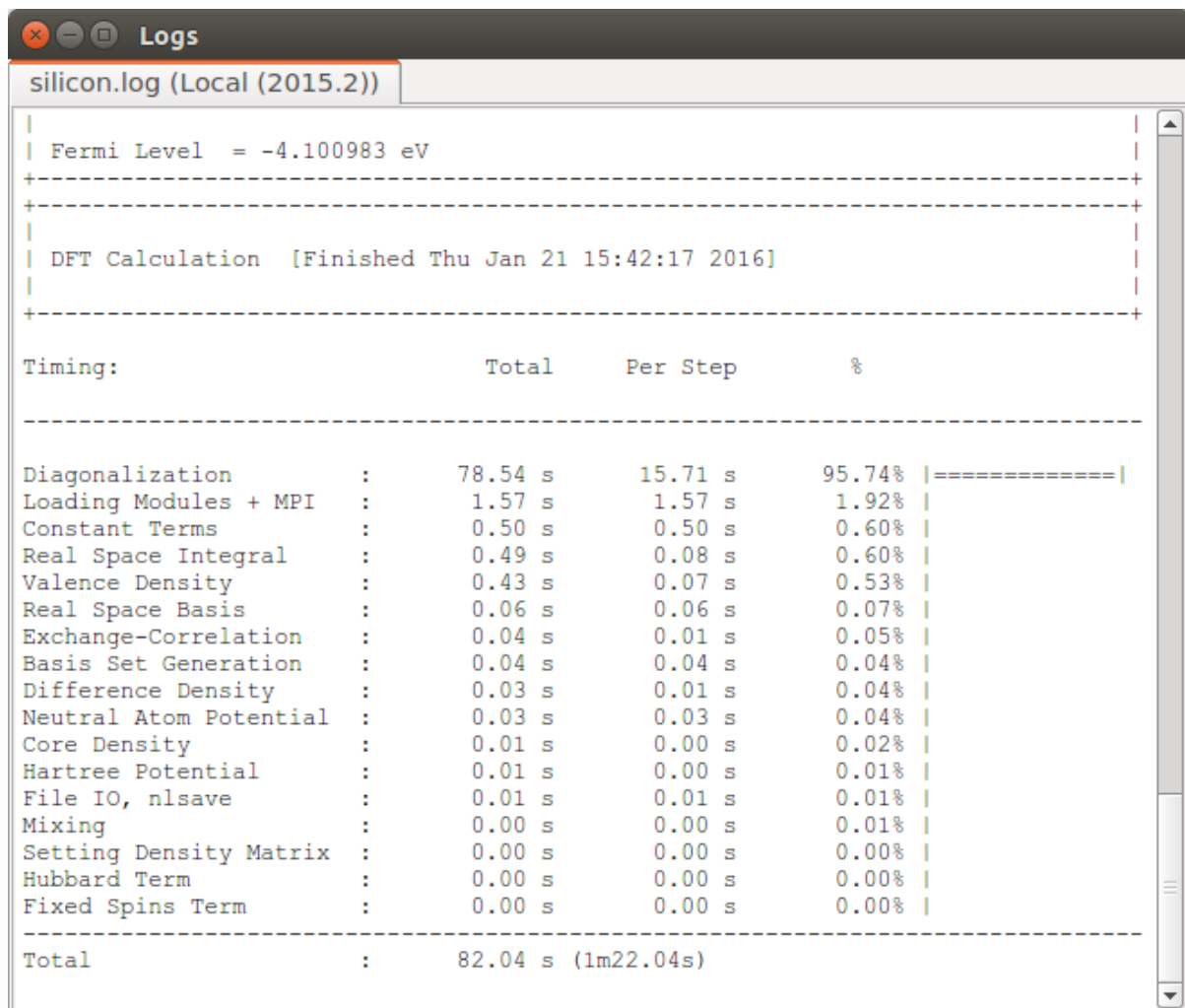
Set these settings according to your needs, and click OK.

Back in the **Job Manager**, click the **Run** ► icon to start the job. The task state changes from “Pending” to “Running”.



The job finishes after ca. 1 minute (2.5 GHz CPU). Note that the task state changes to "Finished". You can inspect the job log file by clicking the LOG icon: 





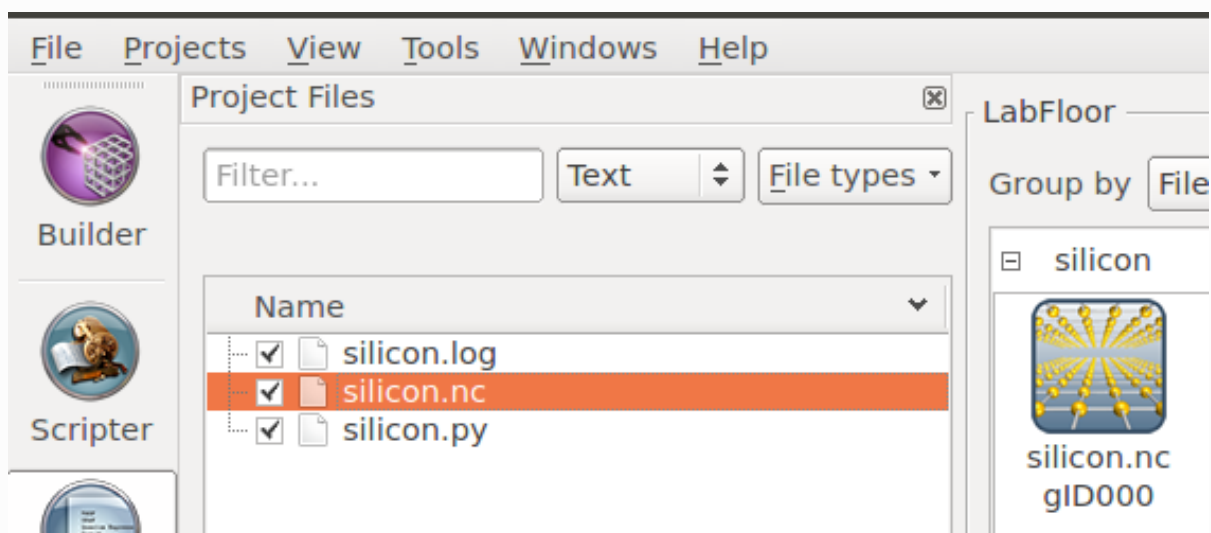
```

silicon.log (Local (2015.2))
|
| Fermi Level = -4.100983 eV
+-----+
+-----+
| DFT Calculation [Finished Thu Jan 21 15:42:17 2016]
+-----+
+-----+

Timing:                Total      Per Step      %
-----
Diagonalization       :    78.54 s    15.71 s    95.74% |=====|
Loading Modules + MPI :     1.57 s     1.57 s     1.92% |
Constant Terms        :     0.50 s     0.50 s     0.60% |
Real Space Integral   :     0.49 s     0.08 s     0.60% |
Valence Density       :     0.43 s     0.07 s     0.53% |
Real Space Basis      :     0.06 s     0.06 s     0.07% |
Exchange-Correlation  :     0.04 s     0.01 s     0.05% |
Basis Set Generation  :     0.04 s     0.04 s     0.04% |
Difference Density    :     0.03 s     0.01 s     0.04% |
Neutral Atom Potential:     0.03 s     0.03 s     0.04% |
Core Density          :     0.01 s     0.00 s     0.02% |
Hartree Potential     :     0.01 s     0.00 s     0.01% |
File IO, nlsave      :     0.01 s     0.01 s     0.01% |
Mixing                :     0.00 s     0.00 s     0.01% |
Setting Density Matrix:     0.00 s     0.00 s     0.00% |
Hubbard Term         :     0.00 s     0.00 s     0.00% |
Fixed Spins Term     :     0.00 s     0.00 s     0.00% |
-----
Total                  :     82.04 s (1m22.04s)

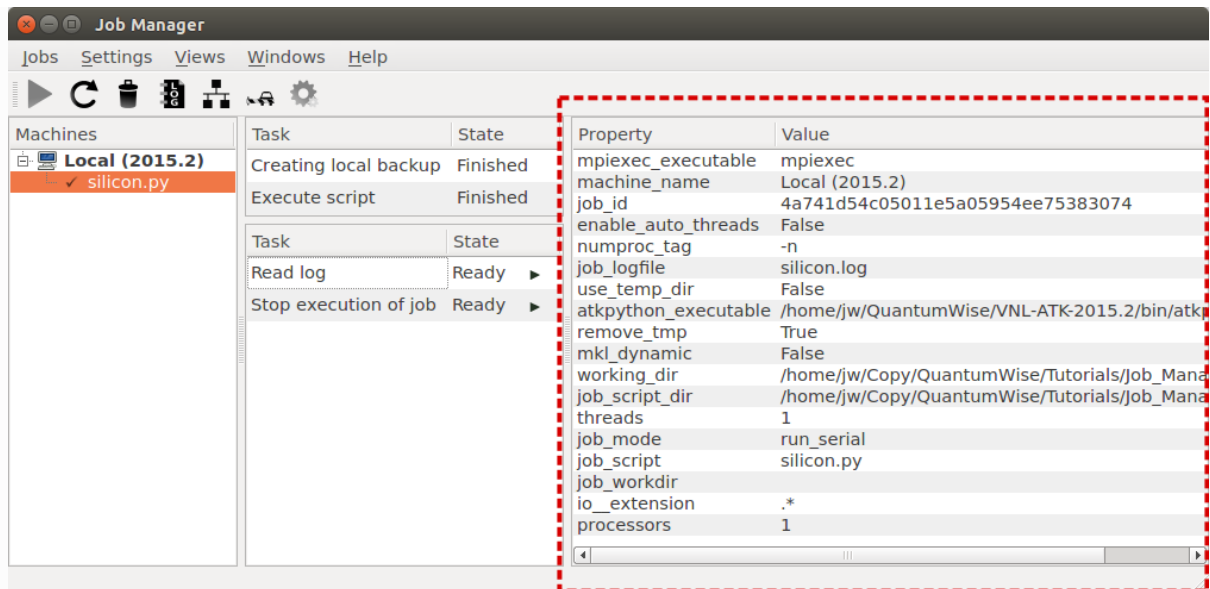
```


The job output of course appears on the QuantumATK LabFloor after job execution.

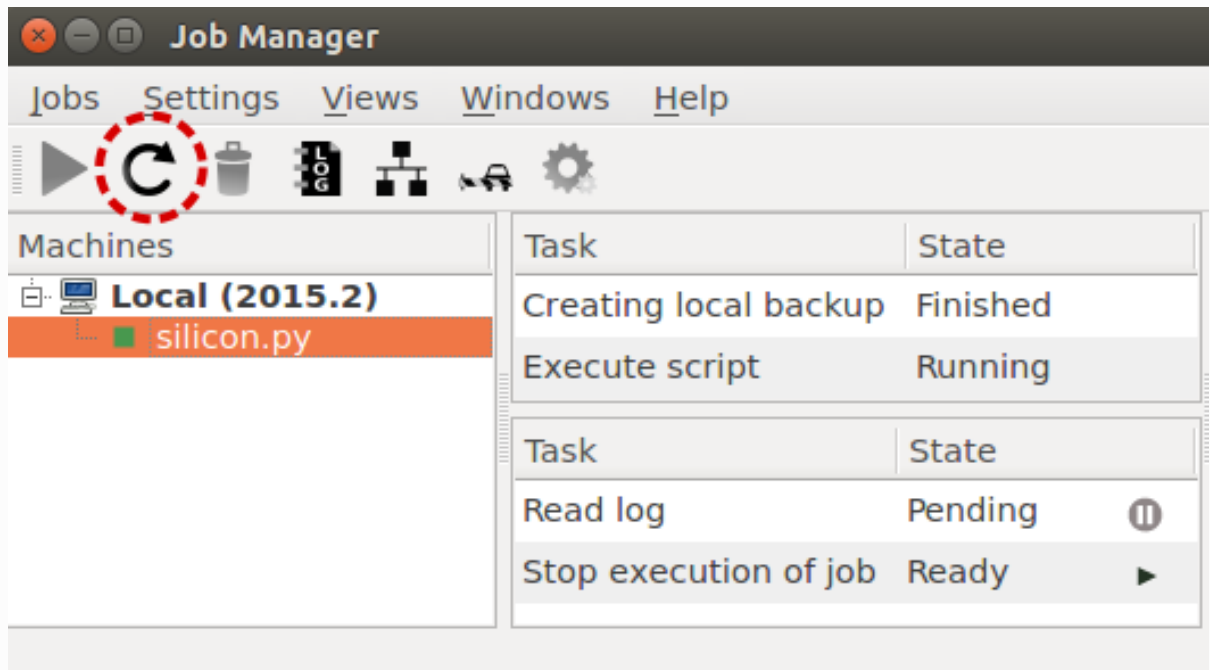


Back in the **Job Manager**, the Property–Value list shows all details of the settings used for job execution, including

- path to the QuantumATK executable;
- name of the Python script and the log file.
- threading and other prallelization options.




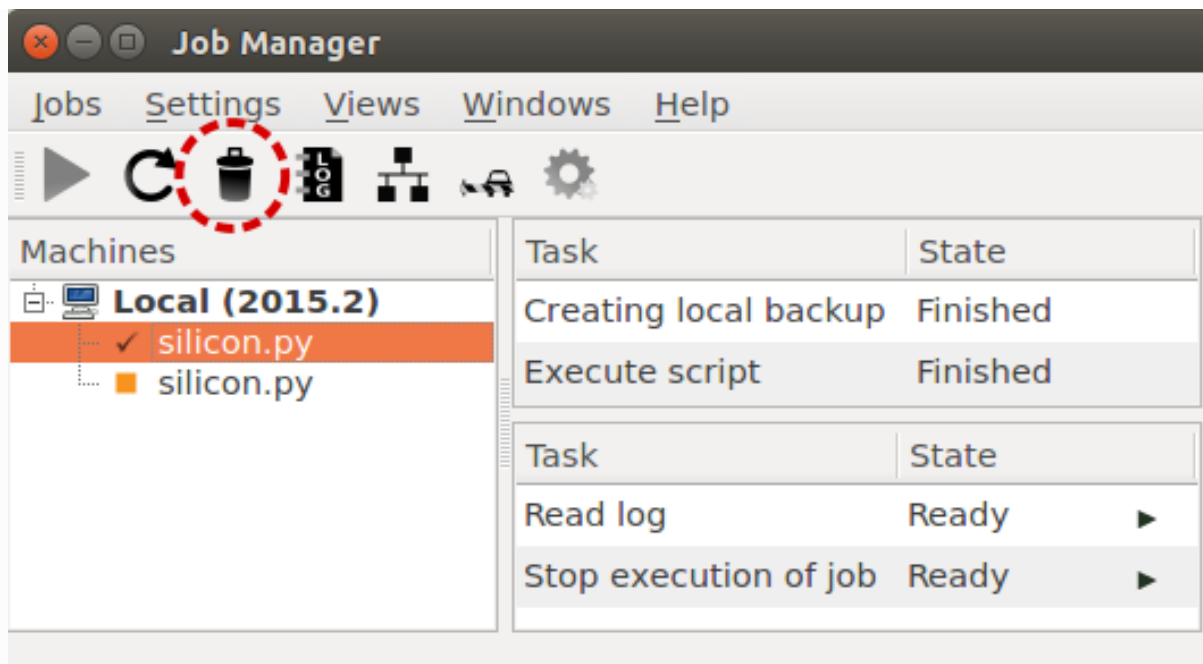
You can use the Resubmit  icon to resubmit a script. Note that any changes that have been made to the script will be picked up by the new job.



Note

Remember that the default job type is “Threaded parallel (Single process)”. You can change this to “Serial” or “Multiprocess parallel” before starting the job.

Use the Trash  icon to remove jobs from the job queue.



Serial execution

In the **Job Settings** window select a **Serial** job type as shown in the below figure to run on a single process with no threading. In fact, note that threading is turned off (number of threads is 1), and MPI parallelization is not available.

✕
□
Job Settings

Job type

Serial
 Threaded parallel (single process)
 Multiprocess parallel

Job properties

Threading

Number of threads Automatic
 Enable MKL_DYNAMIC

MPI

Number of processors

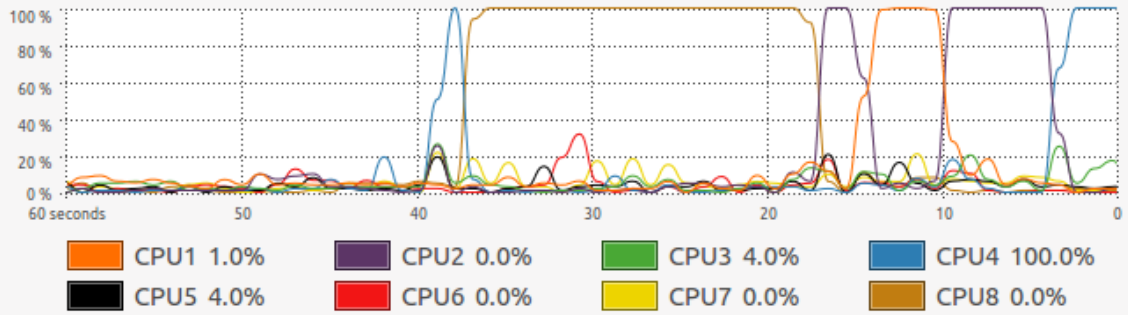
Use separate temporary directory

If you check the system load during local execution in serial, you should see that the serial job launches only a single computing task on a single CPU core.

Process Name	User	% CPU	▲	ID	Memory	Priority
atkpython_exec	jw	12		5067	483.3 MiB	Normal
compiz	jw	0		2134	585.5 MiB	Normal

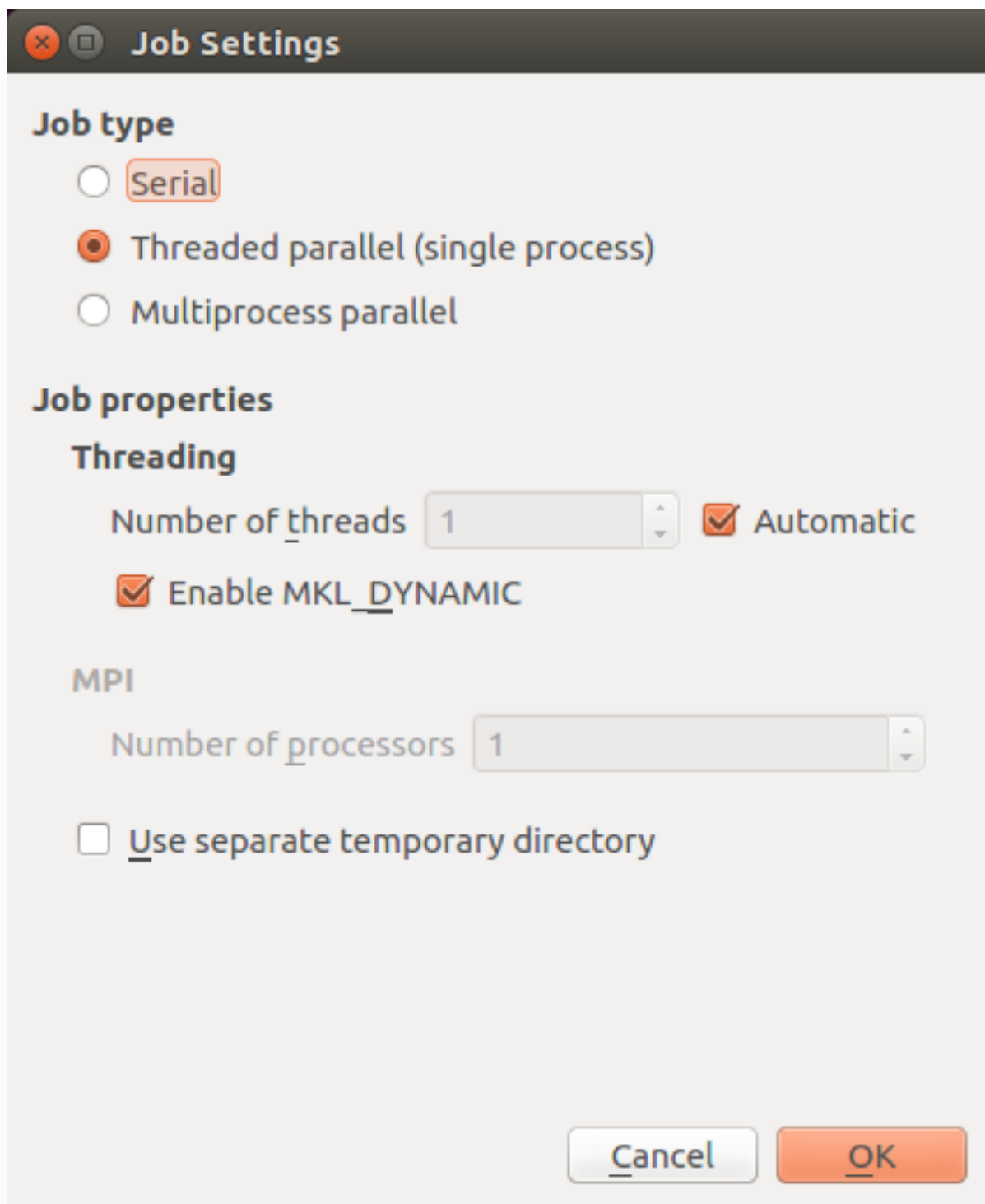
Only one core is used at a time, but the hardware process manager may move the task between cores from time to time.

CPU History



Threading

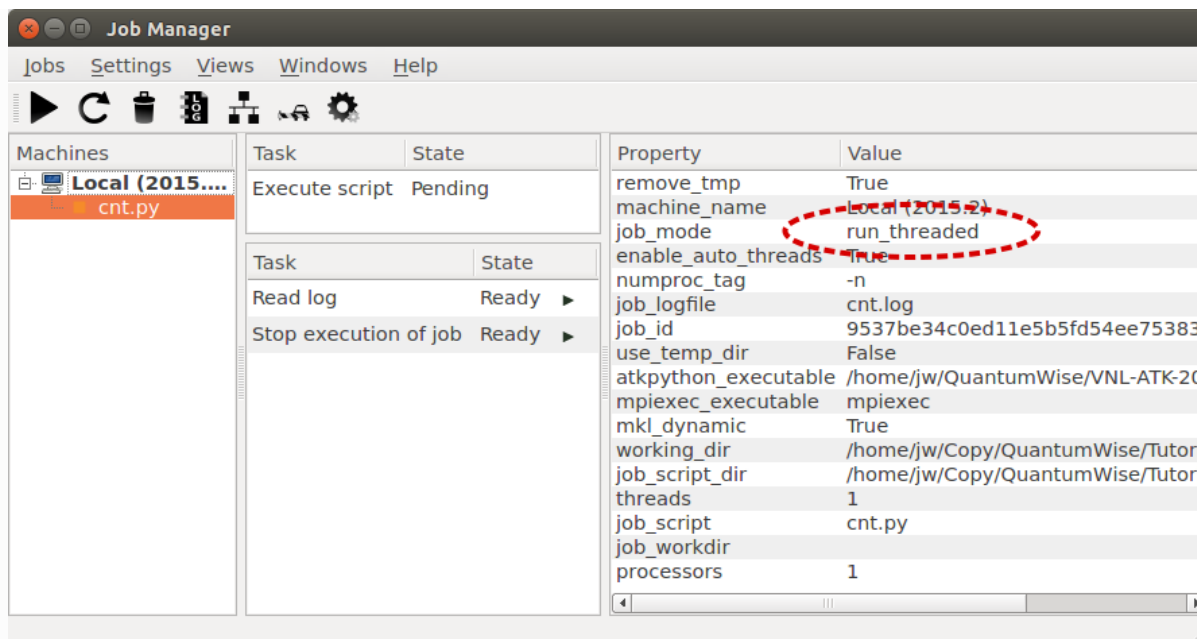
In the **Job Settings** window select a **Threaded parallel (single process)** job type as shown in the below figure to run on a single process with threading.



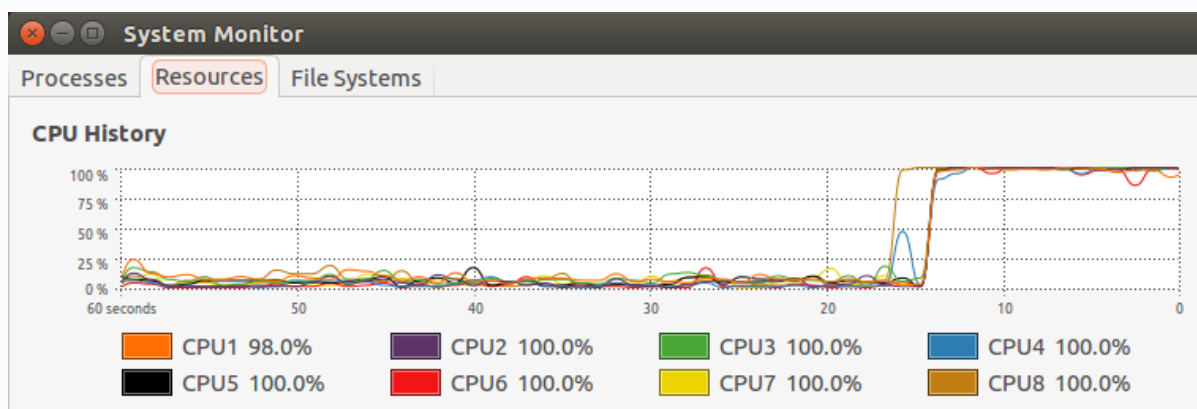
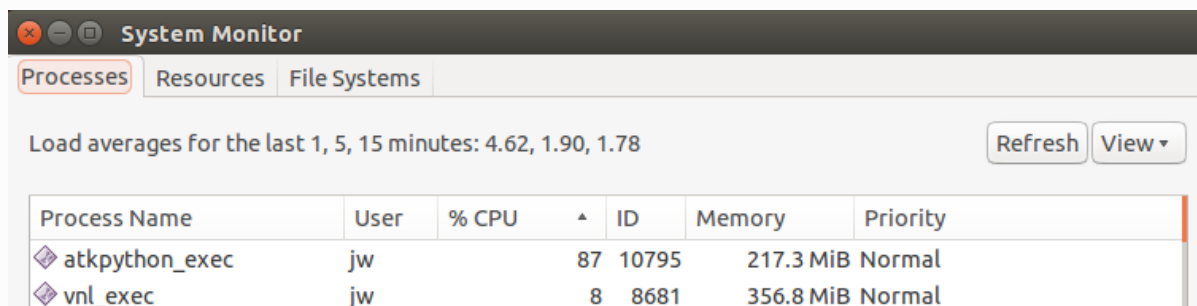
Threading is one way to parallelize a computational job. QuantumATK uses Intel [MKL](#) for [openMP](#) threading. Note that we do in general recommend [MPI parallelization](#) over threading for parallelizing DFT calculations. However, threading is often more efficient for parallelizing ATK-ForceField calculations.

Download the script [cnt.py](#), which uses ATK-ForceField to calculate the dynamical matrix of a multiwall carbon nanotube.

Execute it using the **Job Manager**, and choose job type “Threaded parallel (single process)”. It should be pretty fast.



If you check the system load during execution of the calculation, you should see that only a single **atkpython** process is started, even though several cores appear to be busy. This is because the work load of the one process is split into a number of threads that may be distributed on more cores.



Download the script [cnt.py](#) to test the performance of a ATK-ForceField simulations using threading. This specific example will calculate the dynamical matrix of a multiwall carbon nanotube. If you also run the calculation in serial, you will see that the wall-clock time used for evaluating ATK-ForceField forces may decrease significantly when threading is switched on. In the example shown below, the time spent on force calculations is roughly halved.

```

1 ==> cnt_threading.log <==
2 Timing:
3 -----
4 Forces          :      14.22 s      0.00 s      60.63% |=====|
5 Loading Modules + MPI :      1.57 s      1.57 s      6.69% ||
6 File IO, nlsave  :      0.10 s      0.05 s      0.43% |
7 -----
8
9 ==> cnt_serial.log <==
10 Timing:
11 -----
12 Forces          :      22.37 s      0.01 s      79.53% |=====|
13 Loading Modules + MPI :      1.55 s      1.55 s      5.52% ||
14 File IO, nlsave  :      0.06 s      0.03 s      0.21% |
15 -----

```

MPI parallelization

📌 Important

If you are running ATK 2016 or earlier you need to have MPI installed on your local machine.

Both the **Linux** and **Windows** versions are compiled against Intel MPI library. Since ATK 2017 Intel's `mpiexec.hydra` is provided on both Windows and Linux versions - **this is the recommended way to run QuantumATK in parallel.**

In **Job Settings** choose *Multiprocess parallel* and e.g. 4 MPI processes.

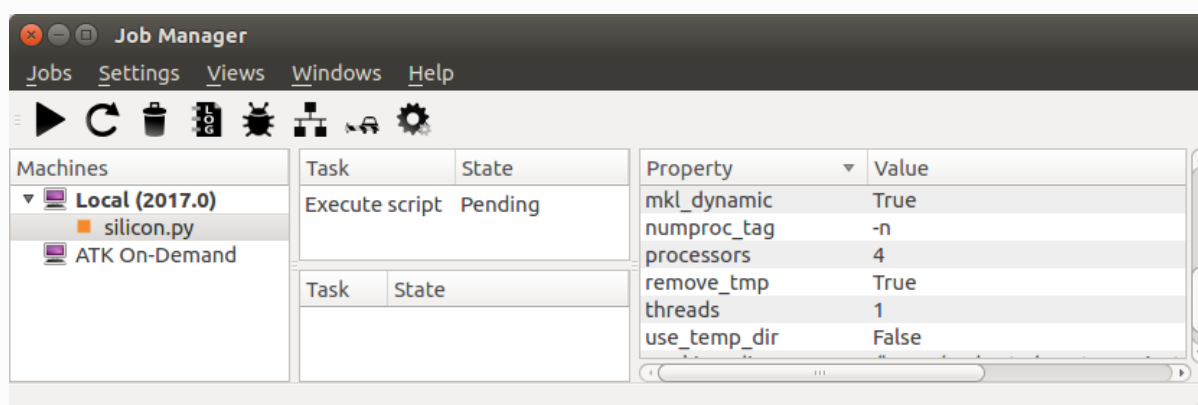
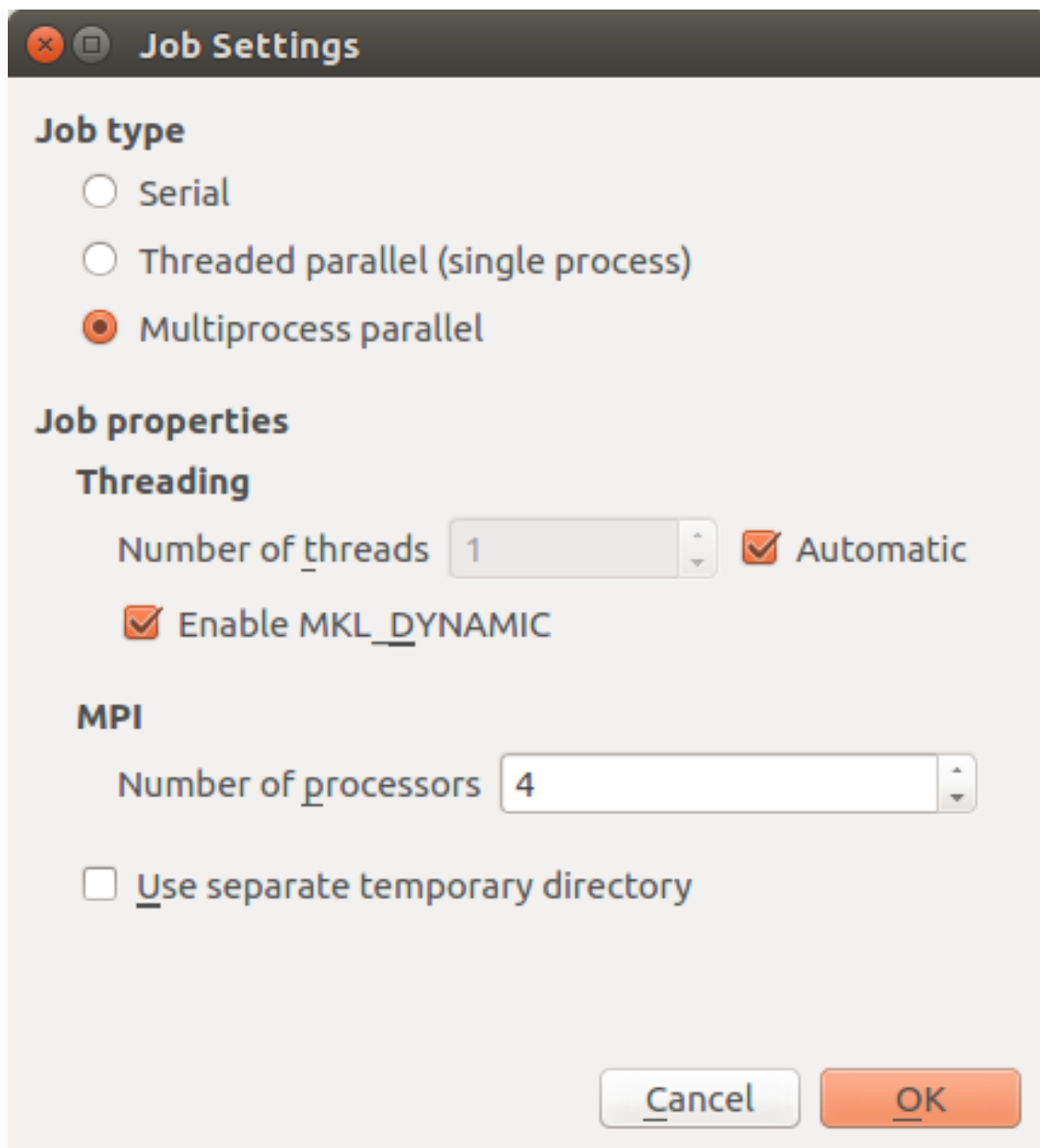


Fig. 22 The Property-Value list shows the name of the MPI executable and that 4 processors are used for MPI.

Running from the command line

If you wish to run QuantumATK in parallel from the command line you can use the `mpiexec.hydra` binary

shipped with QuantumATK and located in the folder `libexec/mpiexec.hydra` present in your installation folder.

In this case you can run parallel jobs with:

```
$ QW_INSTALLATION_PATH/libexec/mpiexec.hydra -n 4 atkpython atk_script.py
```

Hint

Prepend `QW_INSTALLATION_PATH/libexec/` in your `PATH`.


```
$ export PATH=QW_INSTALLATION_PATH/libexec:$PATH
```

This way you will automatically pick up the `mpiexec.hydra` binary shipped with QuantumATK:

```
$ mpiexec.hydra -n 4 atkpython atk_script.py
```

Machine Manager

It may sometimes be convenient to have a predefined local machine that is set up with MPI parallelization as default mode. You can easily add such a machine yourself.

In the Job Manager main window, click  to open the **Machine Manager**, and click **New** ▶ **Local**.

Then edit the default job settings of the new machine in the window that pops up:

- Name the machine, e.g. "Local (2017.0) - 4 MPI".
- Select *Multiprocess Parallel* as job type.
- Make sure threading is turned off (*Number of threads* = 1)
- Choose the default number of processors, e.g. 4.
- Click *OK* to add the new machine to the Machine Manager.

Job Settings

Machine name Local (2017.0) - 4 MPI

Job type

- Serial
- Threaded parallel (single process)
- Multiprocess parallel

Job properties

Threading

Number of threads 1 Automatic
 Enable MKL_DYNAMIC

MPI

Number of processors 4

Use separate temporary directory

Cancel

OK

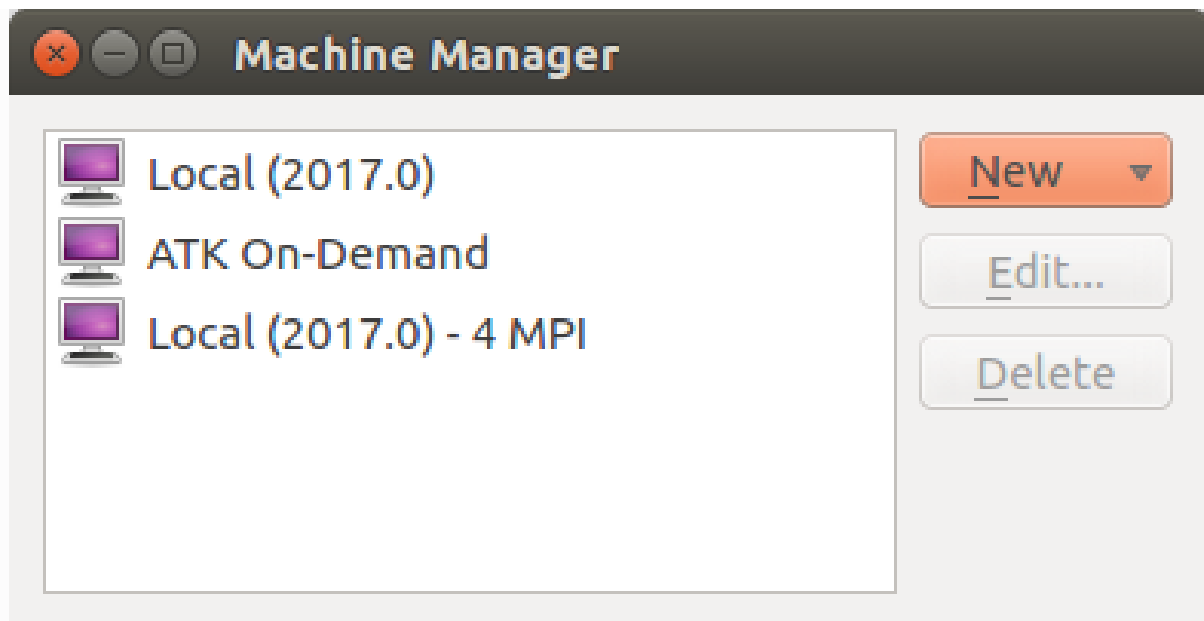


Fig. 23 You can add as many custom machines to the Machine Manager as you like.

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