Introduction to ZECM (a.k.a. ZECM) HPC Cluster

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Cluster infrastructure

Hardware equipment - Cluster partitions

- 3 partitions: *standard*, *smp* and *gpu*
 - **smp partition**: 3 big(!) nodes, each with 3 TB memory and 64 cores (2.10GHz, Intel Xeon E7-4850 v4, AVX2)
 - standard partition: 132 nodes, each with 250 GiB memory and 20 cores (2.20GHz, Intel Xeon E5-2630 v4, AVX2)
 - gpu partition: 21 nodes, each with 500 GiB memory, 2x 16GB Nvidia Tesla P100 and 20 cores (2.20GHz, Intel Xeon E5-2630 v4, AVX2)
- fast node interconnect: Intel's proprietary Omni-Path architecture
- Distributed Parallel File System for high-bandwidth network disk storage: Fraunhofer's BeeGFS

Software equipment

- Workload/scheduler facility SLURM (Simple Linux Utility for Resource Management): managing workload on cluster of computing nodes grouped by partitions; accessed via queues by jobs demanding for resource allocations
- STUBL (SLURM Tools and UBiLities): Collection of user-friendly wrappers for bare SLURM commands

Software equipment

Available Quantum Chemistry packages:
 TURBOMOLE 7.3 (Kaupp's group)
 GAUSSIAN 16 (OpenMP, but no LINDA)
 MRCC (in planning)

- Available Developer modules: Compiler suites: GNU 7.2.x and Nvidia CUDA 9.0.x/9.2.x compilers; MPI: Open MPI 3.0.x/3.1.x
- Cluster logon via: ssh username@gateway.hpc.tu-berlin.de

Concepts of SLURM workload manager

SLURM: Design aspects

- Purpose Fair sharing of compute resources by quoting wrt. to jobs and users on clusters (and federations)
- Jobs and Steps consuming computing resources via allocation: cores, runtime, main memory, accelerator cards, further general resources . . .
- Queueing job's priority: Weighted by aspects of runtime;
 available and used resources accounted per user/group/project

SLURM: Functioning and lexical terms

- General terms: job (general resource allocation), job step (separate tasks/processes within job), task (an OS-process), core (no hyper-threading per default)
- A job creates a context for one or several tasks, each running on a single core. A job can be divided into several steps each launched by srun.
- Compared to Sun's Gridendgine: SLURM without concept of parallel environments, direct implementation of distributed jobs (tasks per node, see below)

Using SLURM with interactive and batch jobs

SLURM commands - Essentials

- General command line interfaces (see **SLURM** manpages):
 - 1. Resource allocation: salloc
 - Running jobs: sbatch (batch jobs via scripts) srun (interactive jobs and job tasks)
 - 3. Cluster/Queue status: sinfo, squeue sqstat and snode (STUBL)
 - 4. Job status/cancelling: sstat, scancel
 - 5. Job and Cluster configuration: scontrol

Launching interactive jobs - An Example

```
srun -p[standard|smp|gpu] -t20:00 --mem=2[T|G|M|K] -n8 -N4
--ntasks-per-node=2 --gres=gpu:tesla:1 --pty /bin/bash -il
```

- 1. --mem: resident memory per node in TiB, GiB, ...
- 2. -t: wall time of your job
- 3. -p: selection of partition
- 4. --mem: resident memory per node
- 5. -n: overall number of tasks (processes) (digit, no range allowed)
- 6. -N: number of nodes (digit or range allowed; e.g.: -N2-4)
- 7. --ntasks-per-node: tasks (processes) per node (digit, no range allowed)
- 8. --gres: general resources (here GPU)
- 9. --pty: interactive shell on pseudo terminal

General batch job script

Writing scripts - Directives with sbatch:

```
#!/bin/bash --login

#SBATCH -J JOB_NAME
# IMPORTANT: memory limit; otherwise all node memory allocated
#SBATCH --mem=1GB # 1GiB resident memory per NODE
# #SBATCH --mem-per-cpu=1GB # 1GiB resident memory per CORE
#SBATCH -N 4 # number of nodes (digit or range allowed)
#SBATCH -N 8 # overall number of tasks (digit; no range allowed)
# #SBATCH --ntasks-per-node=2 # tasks per node (digit; no range allowed)
# #SBATCH -t 00:30:00 # 30 minutes wall clock time
# #SBATCH --gres=gpu:tesla:1 # no of GPU accelerators
#SBATCH -p smp # runnable on hosts of partition smp
```

General batch job script

Continuation - Directives with sbatch:

```
# setting CWD if
# different from directory where sbatch having been launched
# #SBATCH -D /path/to/target/dir
# general batch job output
# formatting:: %N: hostname; %j: $SLURM_JOBID; %x: jobname
#SBATCH -o stdout-%x-%j-%N.out
#SBATCH -e stderr-%x-%j-%N.out
# mail options
# #SBATCH --mail-type=<BEGIN|END|FAIL|ALL|NONE>
# #SBATCH --mail-user=name@postbox.de
# only if needed
# export MODULEPATH=/path/to/module/directory:$MODULEPATH
module load [your_module]
[your command] < [input] > [output]
```

Advanced SLURM features

- Array jobs (available with sbatch): -a <id_range[:step]> (e.g. -a 0-12:4)
- Job dependencies: -d <[after|afterany|afterok|...]:job_id[:job_id]>
 (see man srun)
- Exclusive Jobs and Job Steps: --exclusive=<user|group> (read man srun carefully before using)
- Heterogeneous job types: see SLURM manpages and also https://slurm.schedmd.com/heterogeneous_jobs.html

Running batch jobs - An example with TURBOMOLE

Batch job script - An example with TURBOMOLE

TURBOMOLE with MPI:

```
#!/bin/bash --login
#SBATCH -J dscf MPI
# IMPORTANT: memory limit; otherwise all node memory allocated
#SBATCH --mem=2GB
# #SBATCH --mem-per-cpu=1GB
#SBATCH -N 4
#SBATCH -n 8
# #SBATCH --ntasks-per-node=2
#SBATCH -t. 00:30:00
#SBATCH -p smp
export MODULEPATH=/home/units/Fak_II/quantenchemie/\
turbomole/modules: $MODULEPATH
export PARA_ARCH=MPI
module load turbomole 7.3.0
dscf control > dscf.out
```

Working on cluster - Best practice

Best practice on ZECM HPC

- write temporary data to directory /scratch/[your_user_name] (actually on BeeGFS, no node-local scratch)
- Always define the number of tasks per job via -N even when running intra-node jobs (cf. man srun)
- Always define your job's memory requirements via switches:
 --mem= or --mem-per-cpu= (standard for --mem-per-cpu predefined on this cluster)
- new job types: plan to run short test jobs in order to acquire necessary amount of resident main memory and probable runtime

Literature and references

Finding further information

Further references ZECM HPC and SLURM

```
Official documentation ZECM HPC (login with TUB account): 
https://hpc.tu-berlin.de
```

Documentation ZECM HPC in team IT's Wiki:

 $http://it.chem.tu-berlin.de/wiki/doku.php?id=hpc:hpc_ZECM:start$

 ${\sf ISIS\ course/forum\ (login\ with\ TUB\ account):}$

 $https://isis.tu-berlin.de/course/view.php?id{=}13680$

Official **SLURM** documentation with online manual pages:

https://slurm.schedmd.com

Additional **SLURM** documentation:

http://www.lrz.de/services/compute/linux-cluster/batch_parallel