# Submission 01 Moritz Perschke

## Exercise 1

#### Task 1

Study how to submit jobs in SLURM, how to check their state and how to cancel them.

This was mentioned in the proseminar: - to submit a job, use squeue - to check on a jobs status, the alias squ can be used on the cluster, to show the jobs of \$USER - to cancel, use scancel

#### Task 2

Prepare a submission script that starts an arbitrary executable, e.g. /bin/hostname

```
#!/bin/bash
```

```
# Execute job in the partition "lva" unless you have special requirements.

#SBATCH --partition=lva

# Name your job to be able to identify it later

#SBATCH --job-name test

# Redirect output stream to this file

#SBATCH --output=output.log

# Maximum number of tasks (=processes) to start in total

#SBATCH --ntasks=1

# Maximum number of tasks (=processes) to start per node

#SBATCH --ntasks-per-node=1

# Enforce exclusive node allocation, do not share with other jobs

#SBATCH --exclusive
```

### /bin/hostname

The example script does this already.

The output of the script (in output.log in the current dir) is as follows:

```
n001.intern.lcc3.intra.uibk.ac.at
```

### Task 3

In your opionion, what are the 5 most important parameters available when submitting a job and why? What are possible settings of these parameters, and what effect do they have?

I think the example script already includes some of the most important/useful parameters. For example exlusive node allocation, number of tasks and the job name, which I see myself using for almost all jobs in this PS.

I think the --mem-per-cpu or --mem-per-gpu parameter is one of the most important parameters, which is not part of the example script. It might be the case that I know the minimum memory requirement of my job beforehand, and I know that some nodes do not have enough, while others do. In this case this parameter is basically required in order to ensure that my job does not fail.

For a job that reads data from a file, such as measurements, the --input parameter is also essential. This parameter connects the scripts stdin to this file.

On a larger supercomputer, used by multiple projects at the same time, the --clusters parameter might be very important. This parameter specifies the cluster to be used by the job, which could come in very handy in a larger supercomputer. Using this parameter different projects can be assigned different clusters for example, which allows for better sharing of a supercomputer that is meant to be used by different projects at the same time.

For tasks that have specific requirements, the --constraint or -C parameter can ensure the job only runs on nodes with a specific label. These labels are assigned by the administrator. Examples mentioned in the documentation include intel or amd (for vendor specific cpus), graphics (for a node with a gpu) or a specific rack. So if the supercomputer has nodes with an without gpus, and the job requires a gpu, this parameter can ensure the usage of nodes that can adequately handle the job.

There are also a few parameters that deal with 'accounting'. For example --account, which specifies the account to be charged, might be *very* important on an actual scientific cluster. For the lcc3, which by now only exists for teaching purposes (as far as I understand it), I don't see the importance. Noone pays on the lcc3 for resources used, so why woul I want to change the account that gets charged.

### Task 4

How do you run your program in parallel? What environment setup is required? This was also discussed in the proseminar.

```
#!/bin/bash
```

```
# Execute job in the partition "lva" unless you have special requirements.
#SBATCH --partition=lva
# Name your job to be able to identify it later
#SBATCH --job-name test
# Redirect output stream to this file
#SBATCH --output=output.log
# Maximum number of tasks (=processes) to start in total
#SBATCH --ntasks=X
# Maximum number of tasks (=processes) to start per node
```

```
#SBATCH --ntasks-per-node=1
# Enforce exclusive node allocation, do not share with other jobs
#SBATCH --exclusive
```

```
module load openmpi/4.1.4-gcc-12.2.0-6gebvs6 mpiexec -n $SLURM_NTASKS /bin/hostname
```

By loading the mpi module and using it to execute the command in parallel, the "program" is parallelized. Now by increasing the --ntasks=... parameter, the number of /bin/hostname calls can be changed, as every tasks does so.

```
e.g X=3:

n001.intern.lcc3.intra.uibk.ac.at
n002.intern.lcc3.intra.uibk.ac.at
n003.intern.lcc3.intra.uibk.ac.at
X=7:

n001.intern.lcc3.intra.uibk.ac.at
n004.intern.lcc3.intra.uibk.ac.at
n002.intern.lcc3.intra.uibk.ac.at
n003.intern.lcc3.intra.uibk.ac.at
n005.intern.lcc3.intra.uibk.ac.at
n006.intern.lcc3.intra.uibk.ac.at
```

This example also shows that the calls are not happening one after the other quite nicely.

## Exercise 2

When I started this exercise, some people had already entered values into the spreadsheet. At first I used the script above to run the benchmarks, but my times were off by a noticable margin compared to even the highest in the spreadsheet. I then played around with parameters and asked ChatGPT some things that I didn't understand. After a while I realized that instead of using sbatch flags I could use mpiexec flags, wich in combination a more current gcc produced satisfying results.

The used script looks like this:

### #!/bin/bash

```
#SBATCH --partition=lva
#SBATCH --job-name osu_benchmark
#SBATCH --exclusive
#SBATCH --error=errors/cores.log
#SBATCH --output=bandwidth/cores.log
#SBATCH --ntasks=2
```

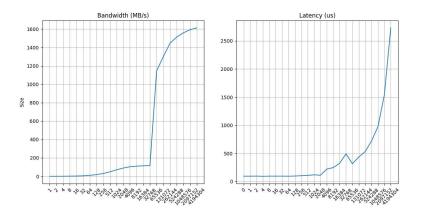


Figure 1: Results of osu\_bw and osu\_latency respectively

```
module load openmpi/3.1.6-gcc-12.2.0-d2gmn55
mpiexec -n $SLURM_NTASKS ./osu_bw
```

In order to run the benchmarks on different cores, sockets or nodes, the --map-by <unit> can be used.

#### #!/bin/bash

```
#SBATCH --partition=lva
#SBATCH --job-name osu_benchmark
#SBATCH --exclusive
#SBATCH --error=errors/nodes.log
#SBATCH --output=bandwidth/nodes.log
#SBATCH --ntasks=2

module load openmpi/3.1.6-gcc-12.2.0-d2gmn55
mpiexec --report-bindings --map-by node -n 2 ../osu_bw
```

To verify the correct placement, the **--report-bindings** option is used, which causes the bindings to be written to stderr.

I am not quite sure why the latency between sockets should be lower for a large packetsize. I assume this is a problem with my benchmark and is not representative of the actual latency.

To measure the consistency of these measurements, I submitted the same job ten times.

Judging by the graph, these measurements appear quite stable

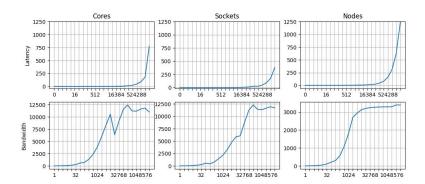


Figure 2: Comparison of Benchmarks across cores, sockets and nodes

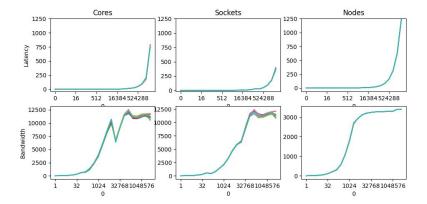


Figure 3: The results of the 10 benchmarks