



ADOPTING SLURM

TRANSITIONING FROM PBS SCHEDULER

Ondřej Meca

Senior Innovation Technology Developer



```
git clone git@code.it4i.cz:infralab/presentations/slurm23.git
```

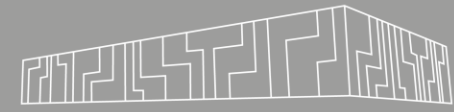
```
/scratch/project/dd-23-116/slurm
```



simple applications:

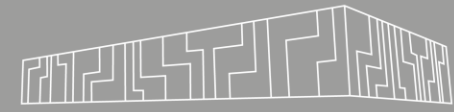
- `src/work.cpp`
 - sequential application (duration 100sec)
- `src/parallel.cpp`
 - hybrid application (duration 40sec)
- `src/mapping.cpp`
 - hybrid application (just print mapping of procs/threads to cores)

build: `sh make.sh`



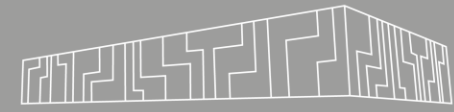
Two modes:

- Interactive
 - `salloc -A DD-23-116 -p skoleni-Adopting-Slurm-CPU`
- Batch
 - `sbatch script.slurm`



Basic parameters

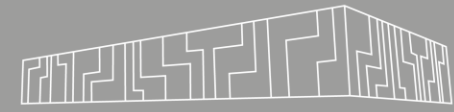
- **-N, --nodes**
- **-n, --ntasks**
 - total number of processes in the job
- **-c, --cpus-per-task**
 - number of threads per process
- **-t, --time**
 - Acceptable time formats include "minutes", "minutes:seconds", "hours:minutes:seconds", "days-hours", "days-hours:minutes" and "days-hours:minutes:seconds"
- **-D, --chdir**
- **-A, --account**
- **-J, --job-name**



hostname.slurm

- set account, partition, name
- print hostname
- output to slurm-JOBID.out

- try to run it with different parameters (N, n)



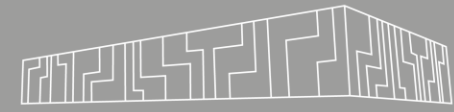
parameters.slurm

- print environment parameters of the job
- output file with many parameters available in Slurm



work.sequential.slurm, work.sequential.long.slurm

- simple sequential work (2x)
- check job info
- how many steps in the job?
- how to kill the job/step?



job.pbs, job.slurm

- check how PBS can be set in Slurm
- where are the jobs executed?
- how to set OMP_NUM_THREADS?



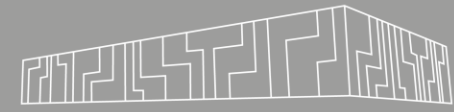
work.parallel.slurm

- simple parallel computation
- how to set number of MPI procs?
- how to set number of threads?
- check the job by htop



work.multiple.slurm

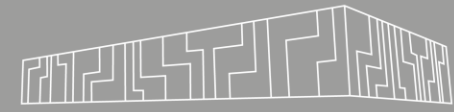
- several sequential computation on the same node
- check the syntax
- check the htop
- why parameter `-c` is used?
- check how steps are executed by `sacct`



work.array.slurm

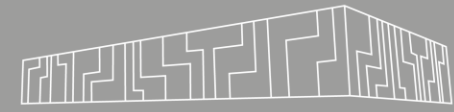
- array job
- several independent jobs spawned by Slurm
- the behavior is the same as creating several separated jobs
 - each job gets its own resources (compute node)

- check JOB_ID, ARRAY_JOB_ID, ARRAY_TASK_ID



work.dependent.slurm

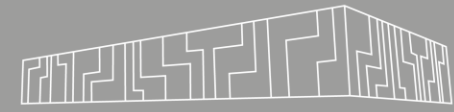
- job dependency
- run the script and check execution by sacct and squeue



work.partial.slurm

- ask for a particular number of GPUs (only on Karolina)

- docs.it4i.cz/general/karolina-slurm



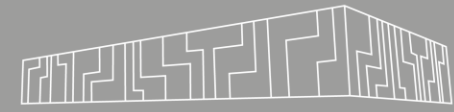
mapping.percore.slurm

- default mapping (with `-N` and `-n` parameters)
- each process is pinned to single core
- can be used for compute bound applications



mapping.pernode.slurm

- without `-n` parameter
- cores of whole node is available
- be aware of NUMA
- set the number of threads (default is the number of cores)



mapping.persocket.slurm

- use `-c`, `--cpus-per-task` parameter
- check the number of cores per socket

mapping.persocket.cyclic.slurm

- different mapping of MPI processes



Ondřej Meca
ondrej.meca@vsb.cz

IT4Innovations National Supercomputing Center
VSB – Technical University of Ostrava
Studentská 6231/1B
708 00 Ostrava-Poruba, Czech Republic
www.it4i.cz

VSB TECHNICAL
UNIVERSITY
OF OSTRAVA

IT4INNOVATIONS
NATIONAL SUPERCOMPUTING
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