



Betriebseinheit für technisch-wissenschaftliche Infrastruktur

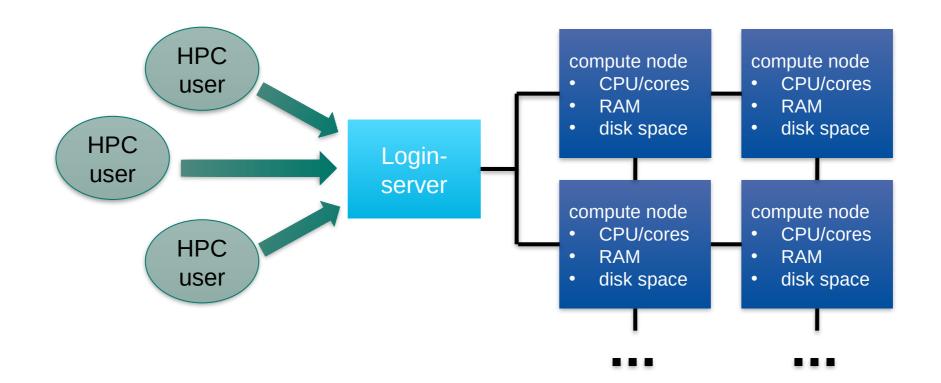
Introduction to High-Performance Computing

Session 02 Basic Cluster Usage and Job Scheduler



Basic Usage HPC Cluster

• HPC cluster from users perspective





Basic Usage HPC Cluster

- many users share a single HPC cluster (resource)
- requires management of the resources
 - for fair sharing
 - for efficient usage
- possible strategies
 - users find free resource and use it
 - part of the resource is reserved for a (group of) user(s)
 - Resource Manager and Job Scheduler

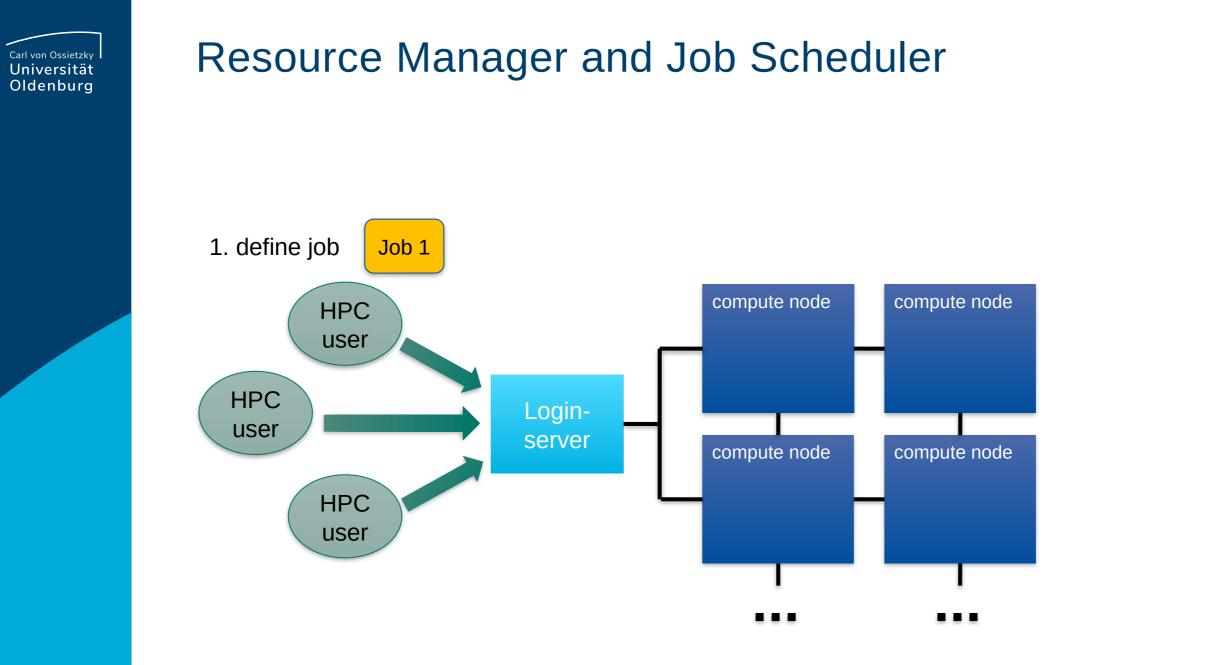


- RM provides low-level functionality for managing jobs
 - start, hold, cancel, and monitor jobs
 - functionality needed by the job scheduler

- JS provides functionality to define and submit jobs
 - interface to RM functionality for the user
 - jobs are scheduled for optimal usage of resource, taking into account fair sharing and other requirements (priority)

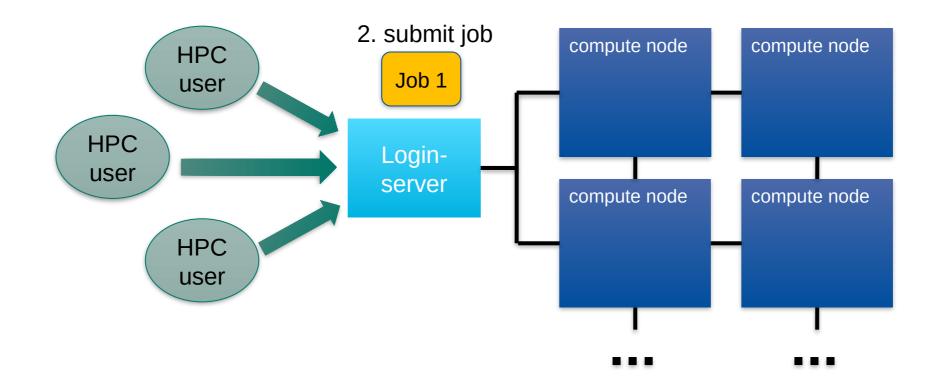
• typically RM and JS are in one application







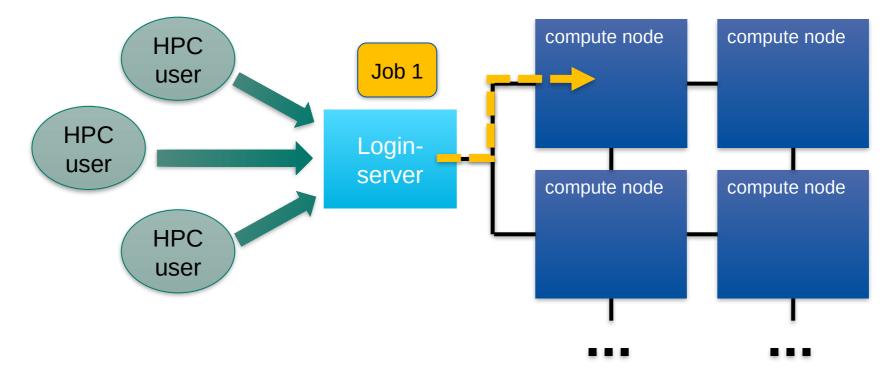




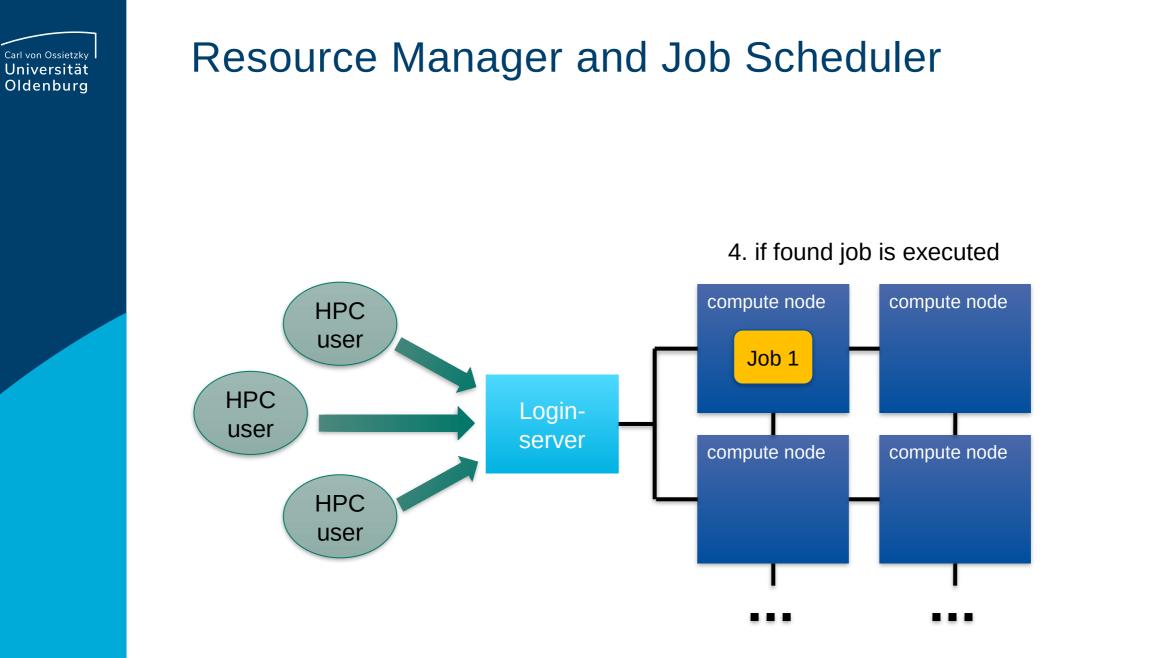




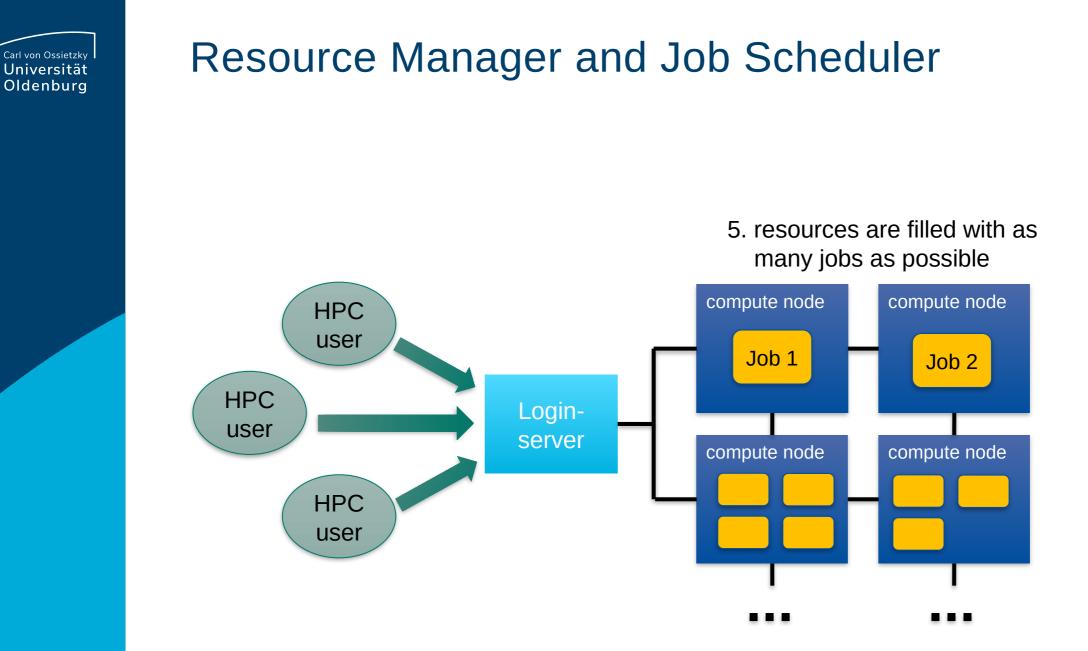
3. JS checks available resources





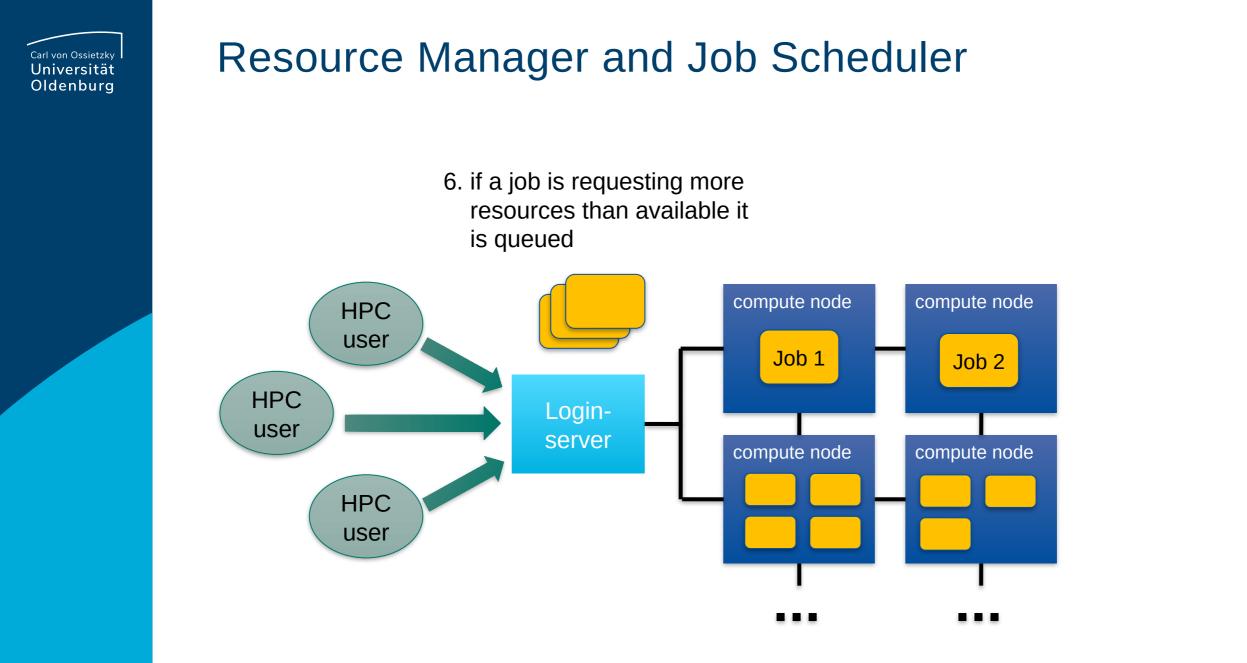








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Tasks of a Job Scheduler

- handling job requests by users (submission, deletion,...)
- prioritize jobs based on the set rules and policies
- place jobs in queue until resources become available
- organize workload on the HPC system for optimal load
- send jobs to the execution host (compute node)
- monitor running jobs
- log files
 - stdout and stderr of jobs
 - accounting information of finished jobs
- terminate job if it use more resources than requested



- there many Resource Manager and Job Scheduler applications available
 - PBS/Torque
 - SLURM (used on the current HPC clusters)
 - LSF

. . .

- SGE (was used on the old HPC clusters)
- LoadLeveler

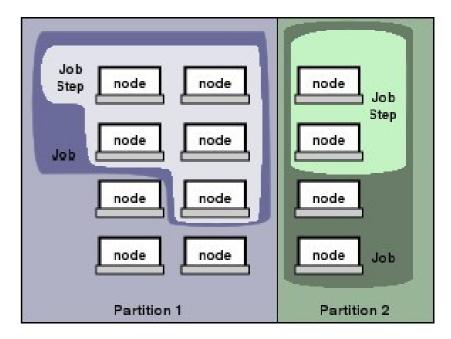
the examples in this course will use SLURM but the principles are the same for all Job Schedulers

(see e.g. http://slurm.schedmd.com/rosetta.pdf)



SLURMs System View

- compute nodes are the basic resource
- compute nodes are organized in partitions
 - logical sets
 - may overlap
- resources are allocated to jobs
 - jobs may contain multiple job steps



(see https://slurm.schedmd.com/overview.html)



Partitions

- in SLURM job limits are defined for each partition
 - partitions know about and manage available resource of the compute nodes
 - other limits (e.g. maximum run time) can be imposed
 - jobs are placed in a partition only if the requested resources fit
 - jobs can be placed in more than one partition (different partitions may have access to different resources)
 - you need to specify at least one partition
 - if you do not specify resources defaults will be used
 - information about partitions with **scontrol**



Partitions

- partitions can be considered job queues
 - each node type has its own partition
 - partitions define the available resources and set defaults

Partition	NodeType	Node Count	CPUs	Default RunTime	Default Memory	Misc
mpcs.p	MPC-STD	158	24		10 375M	
mpcl.p	MPC-LOM	128	24	2h	5 000M	
mpcb.p	MPC-BIG	30	16		30G	2x GTX 1080 in mpcb[001-4]
mpcp.p	MPC-PP	2	40		50G	
mpcg.p cfdg.p	MPC-GPU CFD-GPU	9 3	24		10 375M	1-2x Tesla P100 GPU
cfdl.p	CFD-LOM	160	24		2 333M	
cfdh.p	CFD-HIM	81	24		5 000M	
carl.p	combines mpcl.p and mpcs.p					
eddy.p	combines cfdl.p and cfdh.p					





Basic Usage of SLURM



Examples and Exercises

- · download file to your local computer
- copy the .tgz- or .tar.gz-file to the cluster
 - for Linux use **scp**-command

local\$ scp <localfile> abcd1234@carl.hpc.uni-oldenburg.de:[<remote/path>]

- for Windows use WinSCP or mobaXterm
- unpack the files from the archive
 - on the cluster, e.g.

carl\$ tar -zxvf exercise.tgz

- output is a list of unpacked files



Important SLURM Commands

Command	Used for			
sinfo	information about SLURM nodes and partitions			
squeue	overview of jobs in the scheduler queue			
sacct	accounting information about jobs			
sbatch	submit jobs to the scheduler			
srun	allocate resources if needed and launch a job step within an job allocation			
scancel	delete queued or running jobs			
scontrol	manage jobs (limited) and more			

to get information about commands visit <u>https://slurm.schedmd.com/documentation.html</u> or use

\$ man <command>



sinfo

• information about nodes and partitions

```
$ sinfo -p mpcs.p
PARTITION AVAIL TIMELIMIT
                         NODES STATE NODELIST
mpcs.p up 21-00:00:0 1 drain mpcs025
mpcs.p up 21-00:00:0 61
                                   mix mpcs[002,004,007,009,015,018-
019,022,024,026-027,029-031,034,036-038,041,044,046-050,052-
053,069,072,075,078-082,084-087,089-092,099-102,104-107,110-112,114-
116, 119, 123, 149, 152]
            up 21-00:00:0 76 alloc mpcs[001,005-006,011-014,016-
mpcs.p
017,020-021,023,032-033,039-040,042-043,045,051,054-068,071,073-
074,083,088,093-098,108-109,113,117,120-122,133-148,151,153-158]
                             20
                                idle
            up 21-00:00:0
mpcs.p
mpcs[003,008,010,028,035,070,076-077,103,118,124-132,150]
```

- give idea about used and free resources on cluster
- the state of nodes can be **idle** (no jobs running, free to use), **alloc** (no free resources), **mix** (jobs running but some free resources), or **drain** (node not available)



squeue

• get information about jobs in the scheduler queue

\$ squeue						
JOBID	PARTITION	NAME	USER	ST	TIME	NODES NODELIST(REASON)
2580499_	all_nodes	ofparamt	hoga9120	PD	0:00	16 (ArrayTaskLimit)
1196528					18-21:40:13	
1229276	carl.p	crystal_	wexo7212	R	16-03:57:31	1 mpcs023
1229277	carl.p	crystal_	wexo7212	R	16-03:56:11	1 mpcs093
1229278	carl.p	crystal_	wexo7212	R	16-03:54:47	1 mpcs016

- use the option **-u \$USER** to only show your own jobs
- the option -l gives additional information, output can also be adjusted as needed
- jobs can be shown depending on partition, state, ...



sacct

• accounting information about jobs

<pre>\$ sacct -j 23 JobID ExitCode</pre>	303252 JobName	Partition	Account	AllocCPUS	State
2303252 0:0	HelloClus+	mpcs.p	hrz	8	COMPLETED
2303252.bat+ 0:0	batch		hrz	2	COMPLETED
2303252.0 0:0	orted		hrz	3	COMPLETED

- per default **sacct** shows all jobs of user on the current day



sbatch

 allows to submit a job with sbatch [options] <job-script>

- one mandatory option is -p to set the partition

\$ cat HelloWorld_v1.sh
#/bin/bash

```
# execute these commands
sleep 10
echo "Hello World from $(hostname)"
$ ./HelloWorld_v1.sh
Hello World from hpcl001
$ sbatch -p carl.p HelloWorld_v1.sh
Submitted batch job 2582937
$ squeue -u $USER
$ ls
... slurm-2582937.out ...
```



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sbatch

- options allow to specify requested resources and other settings
 - options have long formation and sometimes short format as well

```
$ sbatch -p carl.p --time=0:10:00 -o HelloWorld.o%j
HelloWorld_v1.sh
Submitted batch job 2582942
$ squeue -u $USER
    JOBID PARTITION NAME USER ST TIME NODES NODELIST
    2582942    carl.p HelloWor lees4820 R 0:03 1 mpcs019
$ ls
. . . HelloWorld.o2582942
$ cat HelloWorld.o2582942
Hello World from mpcs019
$
```



Options for **SBATCH**

https://slurm.schedmd.com/sbatch.html

Option	Short Form	Description
job-name=JobName	-J JobName	sets a name for job which is display in the queue
 partition= <partition></partition>	-p <partition></partition>	(comma-separated list of) partition(s) where the job should run, no default
output= <filename> error=<filename></filename></filename>	-o <filename> -e <filename></filename></filename>	output files for STDOUT and STDERR , default is to join in slurm-%j.out
ntasks= <nt></nt>	-n <nt></nt>	number of tasks (e.g. for MPI parallel jobs)
cpus-per-task= <nc></nc>	-c <nc></nc>	number of CPU-cores for each task, can be used for thread-parallel jobs (OpenMP)
mem-per-cpu= <m></m>		memory per CPU-core, optional
mem= <m></m>		memory per node, exclusive with above
mail-type= <mt> mail-user=</mt>		mail settings





sbatch

- alternatively, sbatch options are specified in job script
 - SLURM options begin with **#SBATCH** (a special comment)
 - then similar to cmd-line option, e.g **#SBATCH** -p carl.p
 - cmd-line options overwrite specifications in script

\$ sbatch HelloWorld_v2.sh
Submitted batch job 2583091
\$





HelloWorld v2.sh

\$ cat HelloWorld_v2.sh #!/bin/bash

######## SLURM options begin

general settings #SBATCH --partition=carl.p #SBATCH --job-name=HelloWorld **#SBATCH** --output=HelloWorld.o%j

requested resources #SBATCH --time=0:10:00 # max runtime #SBATCH --mem=1G

max memory

######## SLURM options end

```
# execute these commands
sleep 10
echo "Hello World from $(hostname)"
```



sbatch

what happens when a job is submitted?

- during the execution of **sbatch**
 - SLURM makes a copy of your job script (changes after submission have no effect)
 - if SLURM accepts job a job ID is returned
 - SLURM may also reject a job, should return error message

- after execution of **sbatch**
 - SLURM computes job priority (many factors are counted)
 - places the job in the queue
 - executes the job script when resoures become available



scontrol

\$ scontrol show part mpcs.p

PartitionName=mpcs.p AllowGroups=carl,hrz AllowAccounts=ALL AllowQos=ALL AllocNodes=ALL Default=NO QoS=N/A DefaultTime=02:00:00 DisableRootJobs=YES ExclusiveUser=N0 TraceTime=0 Hidden=NO MaxNodes=UNLIMITED MaxTime=21-00:00:00 MinNodes=1 LLN=N0 MaxCPUsPerNode=24 Nodes=mpcs[001-158] PriorityJobFactor=1 PriorityTier=1 RootOnly=N0 ReqResv=N0 OverSubscribe=N0 PreemptMode=OFF State=UP TotalCPUs=3792 TotalNodes=158 SelectTypeParameters=NONE DefMemPerCPU=10375

MaxMemPerNode=249000 Introduction to HPC - Session 02



Job Control

- delete a job
 - use scancel <jobID>

- change job details
 - in principle e.g. with scontrol update jobid=<jobid> TimeLimit=0:05:00
 - limitations on what can be changed, also dependent on state of job
 - examples for possible (and useful) changes:
 - reduce **TimeLimit** (only admins can increase)
 - change **Partition** while job is pending



Information about Finished Jobs

- output from job script is written to SLURM output file
 - per default **STDOUT** and **STDERR** are written to the same file
 - default name of output file is slurm-<jobid>.out
 - behavior can be modified with options --output and --error

- running and finished jobs can also be analyzed with **sacct**
 - get information about runtime, CPU time, memory usage
 - See https://wiki.hpcuser.uni-oldenburg.de/index.php?title=Information_on_used_Resources



Exercises



Exercises

- 1. Try the **HelloWorld** Example
- 2. Use the **sacct** command to analyse job

Use the job with id 26445950

- a. How long was the job running? On which nodes?
- b. How much memory was used?
- c. What else can you say about the job?
- 3. Use the **squeue** and **sinfo** commands to get information about the cluster
 - a. How many jobs are running? How many are pending?
 - b. What is the status of the nodes in the partition **mpcl.p**?

