

HP-SEE PARADOX Cluster job management

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HP-SEE

High-Performance Computing Infrastructure for South East Europe's Research Communities

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HP-SEE High-Performance Computing Infrastructure for South East Europe's Research Communities

- Access to login node
- Preparing job submitting scripts
- Submitting jobs:
 - Serial jobs
 - OpenMP jobs
 - MPI jobs
 - Hybrid jobs
 - Application specific jobs (NAMD)
- Batch system job control and monitoring

Access to login node



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- PARADOX login node : ui.ipb.ac.rs
- Access via ssh (remote login program)
 - \$ ssh username@ui.ipb.ac.rs
 - If you need a graphical environment you have to use the -X option: Sh username@ui.ipb.ac.rs -X

Access to login node



File transfer

- To transfer files between **ui.ipb.ac.rs** and your local machine, you can use the **scp** command. Create an archive with the directories you want to copy (it will be faster to transfer):
- \$ tar -cvzf archivename.tgz directoryname1 directoryname2
 or in the case of a file:
- \$ tar -cvzf archivename.tgz filename
- Transfer the archive to your home directory at ui.ipb.ac.rs:
- \$ scp archivename.tgz username@ui.ipb.ac.rs:

Login on **ui.ipb.ac.rs**:

- \$ ssh username@ipb.ac.rs
- Uncompress the archive in your target directory:
- \$ tar -xvzf archivename.tgz destinationdirectory

Access to login node



There are two file systems available to users at ui.ipb.ac.rs:

- /home
- /nfs
- File systems have directories like /home/<USERNAME> or /nfs/<USERNAME>
- /nfs is shared between all nodes on the cluster
- It is required to put all executables and data used by jobs in this directory.
- There is another local file system available on each worker node: /scratch which is used for temporary storage of running jobs

Preparing job submitting scripts



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Portable batch system

- Job submissions
- Resources allocations
- Jobs launching
- PBS server ce64.ipb.ac.rs
- command qsub (ui.ipb.ac.rs)
- Queue hpsee is available for user's job submission
- To submit a job, you first have to write a shell script which contains:
 - A set of directives (beginning with #PBS) which describe needed resources for your job
 - Lines necessary to execute your code

Preparing job submitting scripts



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PBS script

#!/bin/bash

#PBS -q hpsee

#PBS -l nodes=1:ppn=1

#PBS -1 walltime=00:10:00

#PBS -e \${PBS_JOBID}.err

#PBS -o \${PBS_JOBID}.out

cd \$PBS_0_WORKDIR

chmod +x job.sh

./job.sh

Job script

#!/bin/bash
date
hostname
pwd
sleep 10

Preparing job submitting scripts



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- #!/bin/bash Specifies the shell to be used when executing the command portion of the script.
- #PBS -q <queue> Directs the job to the specified queue. Queue hpsee should be used.
- #PBS -o <name> Writes standard output to <name>. \$PBS_JOBID is an environment variable created by PBS that contains the PBS job identifier.
- □ **#PBS** -e <name> Writes standard error to <name>.

Preparing job submitting scripts



- #PBS -1 walltime=<time> Maximum wall-clock time.
 <time> is in the format HH:MM:SS.
- #PBS -1 nodes=1:ppn=1 Number of nodes to be reserved for exclusive use by the job and number of virtual processors per node (ppn) requested for this job.
- cd \$PBS_0_WORKDIR Change to the initial working directory
- chmod +x job.sh setting execute permission on job.sh file
- ./job.sh execute job

Preparing job submitting scripts



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Job submiting is done by executing qsub in /nfs/username/somefolder

- □ \$ qsub job.pbs
- □ The qsub command will return:
 - <JOB_ID>.ce64.ipb.ac.rs



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Serial jobs

http://wiki.ipb.ac.rs/index.php/Serial_job_example

#!/bin/bash

#PBS -q hpsee

- #PBS -1 nodes=1:ppn=1
- #PBS -1 walltime=00:10:00
- #PBS -e \${PBS_JOBID}.err
- #PBS -o \${PBS_JOBID}.out

cd \$PBS_0_WORKDIR

chmod +x job.sh

./job.sh





OpenMP

http://wiki.ipb.ac.rs/index.php/OpenMP_job_example

#!/bin/bash

#PBS -q hpsee

- #PBS -1 nodes=1:ppn=6
- **#PBS** -1 walltime=10:00:00

#PBS -e \${PBS_JOBID}.err

#PBS -o \${PBS_JOBID}.out

```
cd $PBS_O_WORKDIR
chmod +x job
export OMP_NUM_THREADS=6
./job
```



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MPI jobs

http://wiki.ipb.ac.rs/index.php/MPI_job_examples

```
#!/bin/bash
#PBS -q hpsee
#PBS -1 nodes=3:ppn=2
#PBS -1 walltime=10:00:00
#PBS -e ${PBS JOBID}.err
#PBS -o ${PBS JOBID}.out
cd $PBS O WORKDIR
chmod +x job
cat $PBS NODEFILE
${MPI_MPICH_MPIEXEC} ./job
#${MPI_MPICH2_MPIEXEC} --comm=pmi ./job # If mpich2-1.1.1p1 is used
#${MPI OPENMPI MPIEXEC} ./job
```

If mpich-1.2.7p1 is used # If openmpi-1.2.5 is used





Hybrid jobs:

http://wiki.ipb.ac.rs/index.php/Hybrid_job_example

#!/bin/bash

#PBS -q hpsee

#PBS -1 nodes=4:ppn=8

#PBS -1 walltime=10:00:00

#PBS -e \${PBS_JOBID}.err

#PBS -o \${PBS_JOBID}.out

export OMP_NUM_THREADS=8

```
cd $PBS_0_WORKDIR
chmod +x job
```

\${MPI_OPENMPI_MPIEXEC} -np 4 -npernode 1 ./job



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Application specific jobs (NAMD):

http://wiki.ipb.ac.rs/index.php/Application_specific_job_example_(NAMD)

#!/bin/bash

#PBS -q hpsee

- #PBS -1 nodes=4:ppn=8
- **#PBS** -1 walltime=10:00:00
- #PBS -e \${PBS_JOBID}.err
- #PBS -o \${PBS_JOBID}.out

```
cd $PBS_O_WORKDIR
chmod +x script.sh
./script.sh
```

Batch system job control and monitoring



To check the status of job use the following command:

\$ qstat <JOB_ID>

Alternatively check the status of all your jobs using the following syntax of the qstat command:

\$ qstat -u <user_name>

To get detailed information about your job use the following command:

\$ qstat -f <JOB_ID>

If, for some reason, you want to cancel a job following command should be executed:

\$ qdel <JOB_ID>

Batch system job control and monitoring



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qstat	list information about queues and jobs
qstat -q	list all queues on system
qstat -Q	list queue limits for all queues
qstat -a	list all jobs on system
qstat –au <i>userID</i>	list all jobs owned by user userID
qstat -s	list all jobs with status comments
qstat -r	list all running jobs

Batch system job control and monitoring



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qstat- f jobID	list all information known about specified job
qstat -n	in addition to the basic information, nodes allocated to a job are listed
qstat –Qf <queue></queue>	list all information about specified queue
qstat -B	list summary information about the PBS server
qdel jobID	delete the batch job with <i>jobID</i>
qalter	alter a batch job
qsub	submit a job





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