

Workshop Agenda – Feb 25th 2015

Time	Presenter	Title
09:30	T. König	Talk – bwHPC Concept & bwHPC-C5 - Federated User Support Activities
09:45	R. Walter	Talk – bwHPC architecture (bwUniCluster, bwForCluster JUSTUS, ForHLR Phase I)
10:00	A. Fuchs	Talk – Cluster: Access, Data Transfer and Storage, GUI
10:30		<i>Break</i>
10:45	R. Barthel	Talk – File System, Software System (modulefiles), Batch System
11:10	A. Fuchs	Tutorial – bwUniCluster: Access, Data Transfer, Compiling, Modulefiles, Batch Job Scripting
11:50		<i>Lunch Break</i>
13:00	R. Barthel	Talk – Advanced Bash Scripting
13:30	R. Barthel	Tutorial – Advanced (Batch) Job Scripting
14:15		<i>Break</i>
14:30	A. Fuchs	Tutorial – Compiling, Makefile, Parallelising
15:15		User Forum – Solving User Cases
16:00		<i>End</i>





bw|HPC – C5

bwUniCluster Tutorial

Access, Data Transfer, Compiling, Modulefiles, Batch Job Scripting

Annika Fuchs



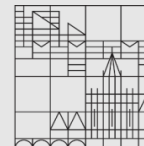
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www.bwhpc-c5.de

Login

- Username <username>
 - Same username as your user account at university.
 - Users from other universities than KIT have to prefix their username by the organization's token, e.g. ho_anfuchs
 - Host <host>
 - bwUniCluster: `uc1.scc.kit.edu`
-
- | | |
|--|--|
| <ul style="list-style-type: none">■ Linux / OS X<ul style="list-style-type: none">■ open terminal:
> <code>ssh <username>@<host></code> | <ul style="list-style-type: none">■ Windows<ul style="list-style-type: none">■ use SSH-Client, e.g. PuTTY■ connect to <host>:
> Login as: <code><username></code> |
|--|--|

Basic commands

\$ <code>pwd</code>	show path of working directory
\$ <code>mkdir <dirname></code>	make directory
\$ <code>cp <sourcefile> <targetfile></code>	copy file
\$ <code>mv <sourcefile> <targetfile></code>	move file
\$ <code>rm <filename></code>	remove file
\$ <code>man <command></code>	show command's manual

Data Transfer

- From localhost to cluster:
 - use `scp` (secure copy) or `sftp` (secure file transfer program)
 - Read manual for options/syntax questions (`man scp`, `man sftp`)
- Linux / OS X
 - Open terminal at your computer:
 - `$ scp <sourcefile> <username>@<host>:<targetfile>`
 - or
 - `$ sftp <username>@<host>:<targetdir>`
 - `$ put <sourcefile>`
- Windows
 - use SCP/SFTP-Client, e.g. WinSCP
 - connect to `<username>@<host>`
 - copy data by drag&drop mechanism

Module Environment

- Users require different software in different versions.
- Software is installed and can be used by loading corresponding modules.

> module avail	show all installed software packages
> module avail compiler	show all available compilers
> module load <modulepath>	load a module in list
> module unload <modulepath>	remove a module from list
> module list	show all loaded modules
> module show <modulepath>	show environment variables of module
> module help <modulepath>	show usage information of module

From \$HOME to \$WORK

- Compute nodes read&write in \$WORK very much faster than in \$HOME directory.
- **DO NOT COMPUTE IN \$HOME !!**
- \$HOME:
 - Source code
- \$WORK:
 - Program input (e.g. initial and boundary conditions)
 - Program output

If lifetime of \$WORK is too short, create a workspace.
But **never** compute in \$HOME!

Exercise

- Download source code from indigo
- Copy source code to bwUniCluster
- Log on bwUniCluster
- Load module file corresponding to the compiler of choice
- Compile the source code, e.g. sequential version with Intel-Compiler:

```
$ icc -o hello hello.c
```
- Move your binary in \$WORK

Source code is written in C and Fortran90 and provided in a sequential version or with OpenMP, MPI or hybrid parallelization.

Submitting jobs via script

- Example: requesting one CPU and 3000 MB of main memory for 5 hours to run the sequential program `hello`

```
#!/bin/bash
#MSUB -l nodes=1:ppn=1
#MSUB -l walltime=5:00:00
#MSUB -l mem=3000mb
#MSUB -q singlenode
#MSUB -N serial-test
#MSUB -m abe
```

Interpreter

Header with msub options

- resource requirements
- queue definition
- notification options,...

```
./hello
```

Execution part

- Submitting the script `jobuc.sh` with MOAB:
> `msub jobuc.sh`

Environment variables in job scripts

Details: http://www.bwhpc-c5.de/wiki/index.php/Batch_Jobs#Environment_Variables_for_Batch_Jobs

■ MOAB variables and own environment variables

	Using MOAB Variables	Defining own variables
Header	<code>#MSUB -o \$(JOBNAME).o\$(JOBID)</code>	<code>#MSUB -v EXEC=./hello</code>
Execution Part	<code>echo „Job \${MOAB_JOBNAME} is running (ID=\${MOAB_JOBID})“</code>	<code>export EXEC=./hello</code>



Parallel Jobs (MPI)

```
#!/bin/bash
#MSUB -l nodes=2:ppn=4
#MSUB -l walltime=05:00
#MSUB -l pmem=1000mb
#MSUB -q multinode
#MSUB -v EXECUTABLE=./hello_mpi
#MSUB -N hello_mpi
#MSUB -o $(JOBNAME).o$(JOBID)

module load mpi/openmpi

echo "Executable ${EXECUTABLE}
running on ${MOAB_PROCCOUNT}
cores"

mpirun ${EXECUTABLE}
```

■ For computations on more than 1 node use queue `multinode`.

■ The corresponding MPI module has to be loaded on the compute nodes.

■ Use `mpirun` to execute the binary.

Parallel Jobs (OpenMP)

```
#!/bin/bash
#MSUB -l nodes=1:ppn=8
#MSUB -l walltime=05:00:00
#MSUB -l pmem=1000mb
#MSUB -q singlenode
#MSUB -v EXECUTABLE=./hello_omp
#MSUB -N hello_omp
#MSUB -o $(JOBNAME).o$(JOBID)

export OMP_NUM_THREADS=${MOAB_PROCCOUNT}

echo "Executable ${EXECUTABLE} running
with ${OMP_NUM_THREADS} threads"

${EXECUTABLE}
```

■ Shared memory restricts to 1 node.

■ Do not define number of threads explicitly. Use MOAB variables.

Keep track of a job

- Submit job script

```
$ msub <jobscript>
```

- If a job (script) is accepted the <jobid> appears at screen.

```
$ checkjob <jobid>    show job details
$ showq               list all my running, idling and blocked jobs by <jobid>
$ showq -n           list all my running, idling and blocked jobs by <jobname>
$ showq -c           list my completed jobs
$ canceljob <jobid>  cancel job
```

GUI via X-Tunnel

Compute at bwUniCluster but display GUI „at home“ (localhost)

■ Modified login

■ Linux / OS X

```
$ ssh -X <username>@<host>
```

■ Windows

- Start X server, e.g. Xming

- PuTTY Configuration: Category SSH > X11

- Check box „Enable X11 forwarding“

■ Submit interactive job (only bwUniCluster)

```
$ msub -I -V -l nodes=1:ppn=1,walltime=02:00:00,mem=4000mb
```

■ Start program, e.g. Matlab:

```
$ module load math/matlab
```

```
$ matlab
```

GUI via VNC (Virtual Network Computing)

- Log on bwUniCluster via terminal/PuTTY

- Submit interactive job

```
$ msub -I -V -l nodes=1:ppn=1,walltime=02:00:00,mem=4000mb
```

- Start VNC server

```
$ module load vis/tigervnc
```

```
$ run_vncserver
```

- Set initial VNC password.

- Follow displayed instructions.

- Start VNC client at localhost

- TightVNC Java Viewer is recommended for Windows users since an SSH client is included.

