



bw|HPC – C5

# bwUniCluster Tutorial

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

Mehmet Soysal



UNIVERSITÄT  
HEIDELBERG  
ZUKUNFT  
SEIT 1386

Hochschule  
für Technik  
Stuttgart



**Hochschule Esslingen**  
University of Applied Sciences

Universität  
Konstanz



UNIVERSITÄT  
MANNHEIM



Universität Stuttgart

EBERHARD KARLS  
UNIVERSITÄT  
TÜBINGEN



**KIT**  
Karlsruher Institut für Technologie



ulm university universität  
**uulm**



Funding:

Baden-Württemberg

MINISTERIUM FÜR WISSENSCHAFT, FORSCHUNG UND KUNST

[www.bwhpc-c5.de](http://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, xy1234
  - Host <host>
    - bwUniCluster: `uc1.scc.kit.edu`
- 
- |  |  |
|--|--|
| <ul style="list-style-type: none"><li>■ Linux / OS X<ul style="list-style-type: none"><li>■ open terminal:<br/>&gt; <code>ssh &lt;username&gt;@&lt;host&gt;</code></li></ul></li></ul> | <ul style="list-style-type: none"><li>■ Windows<ul style="list-style-type: none"><li>■ use SSH-Client, e.g. PuTTY</li><li>■ connect to &lt;host&gt;:<br/>&gt; <b>Login as:</b> <code>&lt;username&gt;</code></li></ul></li></ul> |
|--|--|

# Basic commands

\$ <code>pwd</code>	show path of working directory
\$ <code>mkdir &lt;dirname&gt;</code>	make directory
\$ <code>cp &lt;sourcefile&gt; &lt;targetfile&gt;</code>	copy file
\$ <code>mv &lt;sourcefile&gt; &lt;targetfile&gt;</code>	move file
\$ <code>rm &lt;filename&gt;</code>	remove file
\$ <code>man &lt;command&gt;</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`)
  - From Linux to Linux Systems you can also use `rsync`
- 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.

> <b>module avail</b>	show all installed software packages
> <b>module avail compiler</b>	show all available compilers
> <b>module load &lt;modulepath&gt;</b>	load a module in list
> <b>module unload &lt;modulepath&gt;</b>	remove a module from list
> <b>module list</b>	show all loaded modules
> <b>module show &lt;modulepath&gt;</b>	show environment variables of module
> <b>module help &lt;modulepath&gt;</b>	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

<https://indico.scc.kit.edu/indico/event/132/material/0/>  
/opt/bwhpc/kit/workshop/

- 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:

```
$ module load compiler/intel
$ icc -o hello hello.c
$ ./hello
```

- 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
#sleep 60
```

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

# Parallel Jobs (MPI)

■ `mpicc -o hello_mpi hello_mpi.c`

```
#!/bin/bash
#MSUB -l nodes=2:ppn=4
#MSUB -l walltime=05:00
#MSUB -l pmem=1000mb
#MSUB -q multinode
#MSUB -N hello_mpi
```

```
module load mpi/impi
```

```
mpirun hello_mpi
#sleep 60
```

- 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)

```
icc -openmp hello_omp.c -o hello_omp
```

```
#!/bin/bash
#MSUB -l nodes=1:ppn=8
#MSUB -l walltime=05:00:00
#MSUB -l pmem=1000mb
#MSUB -q singlenode
#MSUB -N hello_omp

EXECUTABLE=./hello_omp
export OMP_NUM_THREADS=${MOAB_PROCCOUNT}

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

./hello_omp
```

■ 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
```