
Workshop "OMNI - New Features and Changes"

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September 30th 2021

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Motivation

- OMNI cluster in service for half a year
 - First hardware installed a year ago
- Review of what has been accomplished, what is yet to do
- Common problems we noticed with users

Agenda

- Filesystem changes
- Common pitfalls we have noticed
- Module system, software licenses, terms of use
- Using the GPUs
- Ongoing work (e.g. Jupyter)
- Questions and Discussion

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File systems

- **Home:** as before, but changed permissions
 - Group spaces under `/group`, explained later
- **Work:** as before (`ws_allocate...`)
- **Fast (Burst Buffer, SSDs):** new, use like workspaces
- **tmp:** used in background, modified by us
 - Make sure you do not use `/tmp` directly, better `$TMP`
- **Storage services (NAS/XNAS):** new, only on login nodes, can and should be used for transfer only
 - NAS for persons, XNAS for groups
 - Mounted under `/nas` and `/xnas`

<https://cluster.uni-siegen.de/omni/usage/file-systems/?lang=en>

Change in home permissions

- New permissions for home directories since August 2021
- Other users cannot see inside your home anymore: `rwX --- ---`
 - Changed from HoRUS to OMNI: previously `rwX r-x r-x`
- Advantage: personal data is now protected
 - Example: sensitive data used for data science
- Disadvantage: files in your home cannot be reached by colleagues
 - If you need to give access, you can change it:
`chmod 755 <your home directory>`

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SLURM Pitfalls

Common problem we regularly see with new users:

- Job that does not use full node (e.g. one task, one core)
- Job then fails with “not enough memory” error

Reason: OMNI nodes configured for shared use

- If you use 1/64th of cores, you only get 1/64th of memory (4 GB)

<https://cluster.uni-siegen.de/omni/usage/queuing-a-job/?lang=en>

SLURM Pitfalls: memory

- Solution: request memory in job script

```
#SBATCH --mem=X
```

- Where X is value in Megabytes
- Maximum is 240000 (240 GB)
- Other units can be used: `mem=120G` will request 120 Gigabytes
- `mem=0` will request all memory on the node

<https://cluster.uni-siegen.de/omni/usage/slurm/?lang=en>

SLURM pitfalls: parallel execution

- Several users thought they run parallel computations, but didn't
- Common example: multiple Python processes
- Reminder: `srun` inside of job script will launch parallel tasks
 - `SLURM_NTASKS` processes by default
 - Use options to reduce number/modify using SLURM parameters
- **Bad idea:** *“It's slow? I need to use more processes!”* blindly

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Module environment

- Our modules come from multiple different sources:
 - Bright cluster manager (GPU stack)
 - OpenHPC (C and Fortran compilers, MPI)
 - Software vendors (Intel, MATLAB)
- Complex structure
 - Interdependencies
 - Mutual incompatibilities
- Brief overview here to reduce confusion

<https://cluster.uni-siegen.de/omni/usage/modules/?lang=en>

Module environment

- Incompatible modules cannot be loaded together
 - **Hidden from view** in `module avail`
 - Can still be searched with `module spider`
- GPU stack is incompatible with almost everything else
- Regular stack: four main “branches”
 - Almost everything depends on either GCC or Intel compiler
 - A lot of things depend on either OpenMPI or Intel MPI (and those depend on GCC or Intel compiler)
 - Default: GCC (`gnu9` module) and OpenMPI (`openmpi4` module)

Regular stack

```

js056352@hpc-login02 ~]$ module avail
----- /opt/ohpc/pub/moduledeps/gnu9-openmpi4 -----
adios/1.13.1      fftw/3.3.8      mfem/4.2        netcdf/4.7.3    phdf5/1.10.6    py3-s
boost/1.75.0     geopm/1.1.0     mumps/5.2.1     omb/5.6.2       pnetcdf/1.12.1  scala
dimemas/5.4.2   hypre/2.18.1    netcdf-cxx/4.3.1  opencoarrays/2.9.2  ptscotch/6.0.6  scala
extrae/3.7.0     imb/2019.6      netcdf-fortran/4.5.2  petsc/3.14.4    py3-mpi4py/3.0.3  score
----- /opt/ohpc/pub/moduledeps/gnu9 -----
R/3.6.3          hdf5/1.10.6     impi/2020.4 (D)  metis/5.1.0     mvapich2/2.3.4  openmpi4/4.0.5 (L)
gsl/2.6          impi/2019.5     likwid/5.0.1    mpich/3.3.2-ofi  openblas/0.3.7  pdtoolkit/3.25.1
----- /cm/shared/ohpc/modulefiles -----
autotools        gnu9/9.3.0      (L)             intel/19.0.5.100010_cm9.0_6a80743563  os
charliecloud/0.15  hwloc/2.1.0     intel/19.1.3.100008_cm9.0_f654bdadee (D)  papi/5.7.0
cmake/3.19.4      intel/19.0.5_2019  libfabric/1.12.1  (L)             paraver/4.8.2
----- /cm/shared/omni/modulefiles -----
DefaultModules (L)  ansys/2021R1    ls-dyna/FLB/r10.1.0  ml/stack        paraview/5.9.0
GpuModules         gromacs/2018.6  ls-dyna/FLB/r12.0.0 (D)  nfft/3.5.2      poweracoustics
abaqus/2021        gromacs/2021.2 (D)  ls-dyna/UTS/r10.1.0  nlopt/2.6.2     powerflow/6-20
abinit/9.4.1       group_test      ls-dyna/UTS/r12.0.0 (D)  nwchem/7.0.0    quantum-espres
ansys/2019R3       hyperworks/14.0  matlab/2020b        octave/6.3.0    quantum-espres
ansys/2020R2 (D)  hyperworks/2019.2 (D)  miniconda3/4.9.2    openfoam/8      quantum-espres

```

Regular stack

Depends on compiler and MPI

Depends on compiler

OpenHPC, no dependencies

Unrelated to OpenHPC

```

- /opt/ohpc/pub/moduledeps/gnu9-openmpi4 -
/4.7.3          phdf5/1.10.6          py3-s
.6.2           pnetcdf/1.12.1        scala
parrays/2.9.2  ptscotch/6.0.6        scala
/3.14.4        py3-mpi4py/3.0.3      score

----- /opt/ohpc/pub/moduledeps/gnu9 -----
mvapich2/2.3.4  openmpi4/4.0.5 (L)
openblas/0.3.7  pdtoolkit/3.25.1

----- /cm/shared/ohpc/modulefiles -----
010_cm9.0_6a80743563          os
008_cm9.0_f654bdadee (D)    papi/5.7.0
(L)                            paraver/4.8.2

----- /cm/shared/omni/modulefiles -----
.1.0          int/stack          paraview/5.9.0
.0.0 (D)      nfft/3.5.2         poweracoustics
.1.0          nlopt/2.6.2        powerflow/6-20
.0.0 (D)      nwchem/7.0.0       quantum-espres
                octave/6.3.0       quantum-espres
2              openfoam/8          quantum-espres
    
```


After loading GpuModules

```
[js056352@hpc-login02 ~]$ module load GpuModules
Bright Maschine Learning software stack loaded.

Inactive Modules:
  1) GpuModules

Activating Modules:
  1) GpuModules

[js056352@hpc-login02 ~]$ module avail

----- /cm/local/modulefiles -----
boost/1.71.0      cmd      dot      gcc/9.2.0      lua/5.3.5      module-git      null      python3      python3
cluster-tools/9.0  cmjob    freeipmi/1.6.4  ipmitool/1.8.18  luajit        module-info     openldap   python36 (L)  shared

----- /cm/shared/modulefiles -----
DefaultModules
GpuModules (L)
blas/openmpi/gcc/64/1.1patch03
blas/gcc/64/3.8.0
bonnie++/1.98
chainer-py36-cuda10.1-gcc/7.4.0
cm-eigen3/3.3.7
cm-pmix3/3.1.4
cuda10.1/blas/10.1.243
cuda10.1/fft/10.1.243
cuda10.1/nsight/10.1.243
gcc8/8.4.0
gdb/8.3.1
globalarrays/openmpi/gcc/64/5.7
gpytorch-py36-cuda10.1-gcc/1.1.1
hdf5/1.10.1
hdf5_18/1.8.21
horovod-mxnet-py36-cuda10.1-gcc/0.19.4
horovod-pytorch-py36-cuda10.1-gcc/0.19.4
horovod-tensorflow-py36-cuda10.1-gcc/0.19.4
horovod-tensorflow2-py37-cuda10.2-gcc/0.20.3
hpcx/2.4.0
keras-py37-cuda10.2-gcc/2.3.1
keras-py37-mkl-gcc8/2.3.1
lapack/gcc/64/3.8.0
ml-pythondeps-py36-cuda10.1-gcc/3.3.0 (L)
ml-pythondeps-py36-mkl-gcc8/3.3.0
ml-pythondeps-py37-cuda10.1-gcc/4.1.2 (L)
ml-pythondeps-py37-cuda10.2-gcc/4.3.9
ml-pythondeps-py37-mkl-gcc8/4.7.0
mpich/ge/gcc/64/3.3.2
mxnet-py36-cuda10.1-gcc/1.6.0
nccl2-cuda10.1-gcc/2.7.8 (L)
```


Software Licences

Software situation more complex on OMNI than HoRUS

- Cluster is three times the size
- More diverse userbase (data science)
- More software that not everyone is allowed to use

→ Considerably more maintenance overhead

→ We cannot install and maintain everything centrally for everyone

Software Policy

Three-tiered software installation policy:

1. More ways of installing software yourself
 - Singularity (container system like Docker)
 - Package manager `conda` (not just Python)
2. Group spaces (work in progress)
 - Directory where group can install software
 - No guarantees, but we advise and guide
3. Central installation as before
 - If multiple groups use software
 - If we believe maintenance overhead is manageable

Software Policy

- Centrally installed software will need a responsible person among the userbase
- “PAPA” - PriAry ApproAch Partner for the Application
- One for each software application
- Should be experienced user who knows userbase
 - Common: one person responsible for license
- We will start approaching people soon

PAPA responsibilities

Not much extra work (!)

- We will consult you whether we should upgrade version
- Should have an overview over how licensing works
- Helps if you know users personally
- If we notice software is not used for extended period, we might ask whether we can uninstall

PAPA responsibilities

Things being a PAPA does NOT mean:

- You do not need to be an expert user
- You do not need to be a sysadmin for the software
- You do not need to keep an overview over who uses software
- You are not liable for mistakes

Terms of Use

- ZIMT leadership demands stricter enforcement of rules than before
- Each user now has to agree to “Terms of Use” when applying for cluster use in Nutzerkontenverwaltung (unisim.zimt.uni-siegen.de)
- Key terms/restrictions
 - Enforcement of trade embargoes (e.g. Iran citizens need special permissions)
 - Cite OMNI use in publications (just mention in Acknowledgements)
 - Regular ZIMT Terms of Use apply too
- Terms (in German): <http://cluster.uni-siegen.de/Nutzungsbedingungen/>

MATLAB licenses

- We were informed of misunderstanding/changed conditions
- Fak. 4 (Bernd Klose) manages MATLAB licenses
 - Collects money for MATLAB use annually
 - Contacts professors
 - ZIMT not involved in process
- Rule: money collected per parallel usable instance of MATLAB
 - Your PC and the cluster count as two instances
 - Pool workers do not count extra

Unfortunately, that means MATLAB usage on cluster is not free

<https://cluster.uni-siegen.de/omni/application-software/matlab/?lang=en>

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GPU tips and tricks

- 10 GPU-nodes with 24 GPUs in total

```
#SBATCH --partition=gpu or
```

```
#SBATCH --gres=gpu:2
```

- You can exclude nodes:

```
#SBATCH -x gpu-node010
```

–Some software cannot handle running on GPUs other than the first

- Remember different module stack: `module load GpuModules`

Node	# GPUs
gpu-node[001-004]	4
gpu-node[005-008]	1
gpu-node[009-010]	2

<https://cluster.uni-siegen.de/omni/usage/gpus/?lang=en>

GPUs: use efficiently

- GPUs are heavily used → long wait times
- Occasionally: GPU not even used (misconfiguration)
- What can you do?
 - Make sure GPUs are actually faster **including wait time**
 - Split work: only run in GPU queue what really needs GPU
 - Make sure software actually uses GPU
- Many applications do not strictly need GPUs (e.g. Tensorflow)

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Ongoing Work

- Two more cluster features being worked on:
 - Jupyter: web platform for interactive programming, visualization
 - Group Spaces: install software for your group
- Both in “beta state”
- We will officially announce them when ready

Jupyter status

- Jupyter working on OMNI
- Some kernels run as SLURM jobs
 - Same problems as regular jobs: **do not run on front end**
- Not yet stable (regular technical problems)
- Documentation not yet ready (easy to break e.g. Python installs)

Group spaces status

- 3 groups as test cases in temporary setup
- Coordination with other ZIMT departments complete (storage, group management etc.)
- Processes basically fully defined
- Not yet migrated
- Documentation not complete

Group spaces status

Preview of eventual procedure (not final):

- Professor will create ticket asking for group space (not yet possible)
 - Professor needs HPC access
- Group will either be created or existing group used
- Professor will be able to add/remove users via <https://selfservice.zimt.uni-siegen.de>
 - Users still need to apply for HPC access separately
- Each group member can install software in group space

Thank you for your attention!

Questions?

Part 2: Open Discussion

Appendix

How not to run in parallel

```
# Wrong, don't do this. Will not run in parallel
#SBATCH --ntasks 64

python myscript.py
```

```
# Still not ideal (circumvents SLURM)
#SBATCH --ntasks 64

for i in {1..64}
do
    python myscript.py &
done
wait
```

How to run in parallel

```
# Better
#SBATCH --ntasks 64

srun wrapper.sh
```

```
# Corresponding wrapper.sh

# Run a different case in each task
case_id=$SLURM_PROCID

# Now srun task 0 will run case 0 etc.
python myscript.py $case_id
```