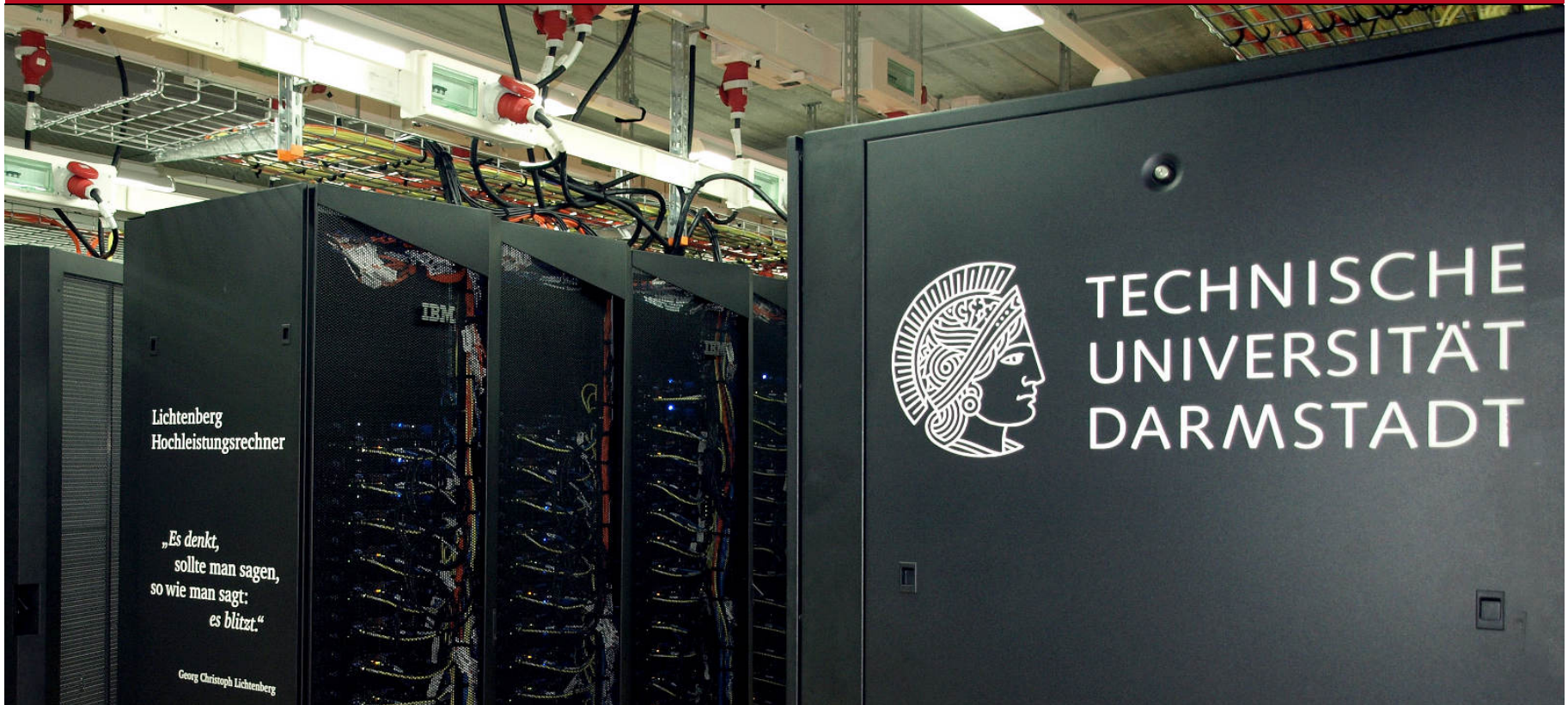


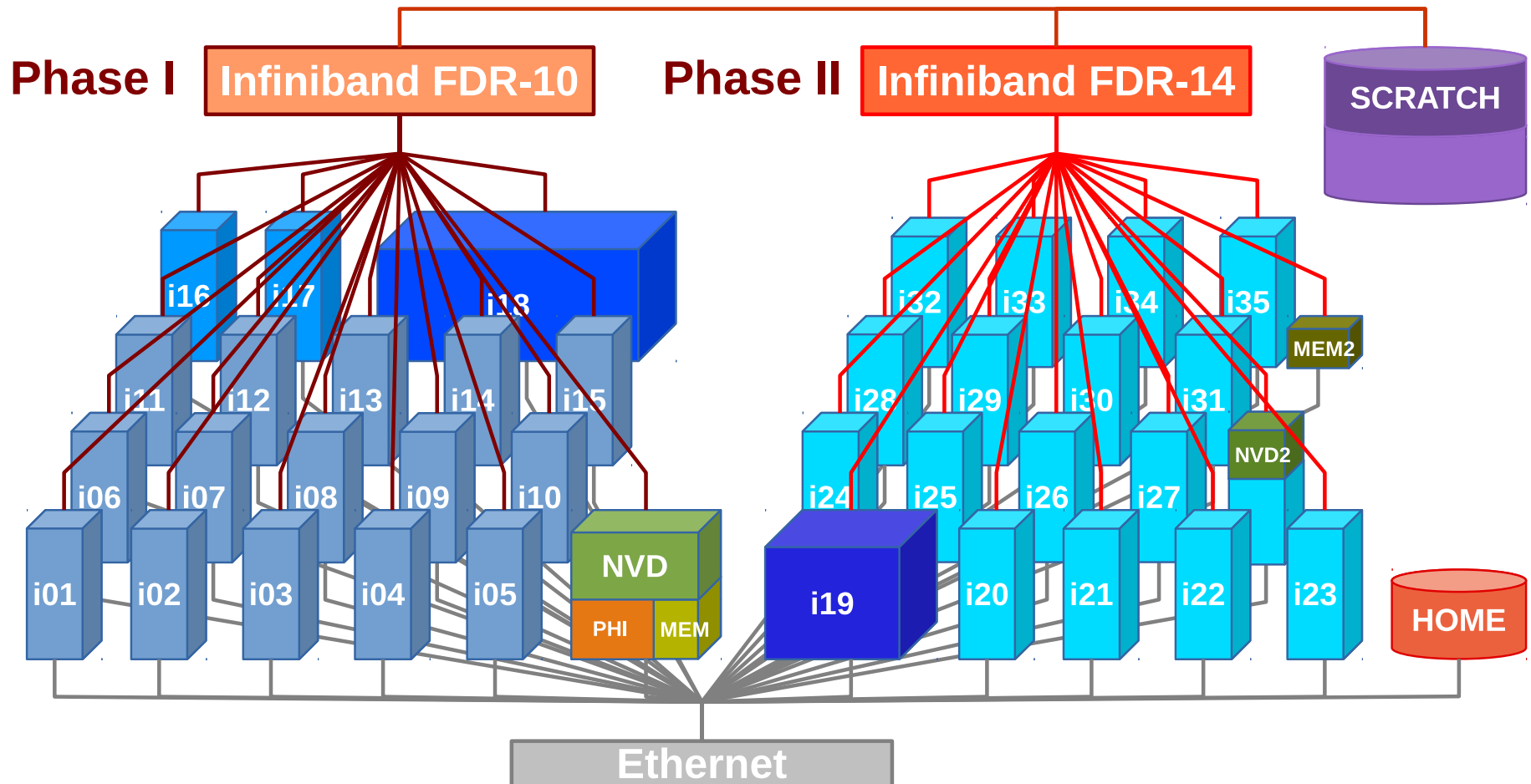
Introduction to the Lichtenberg Cluster



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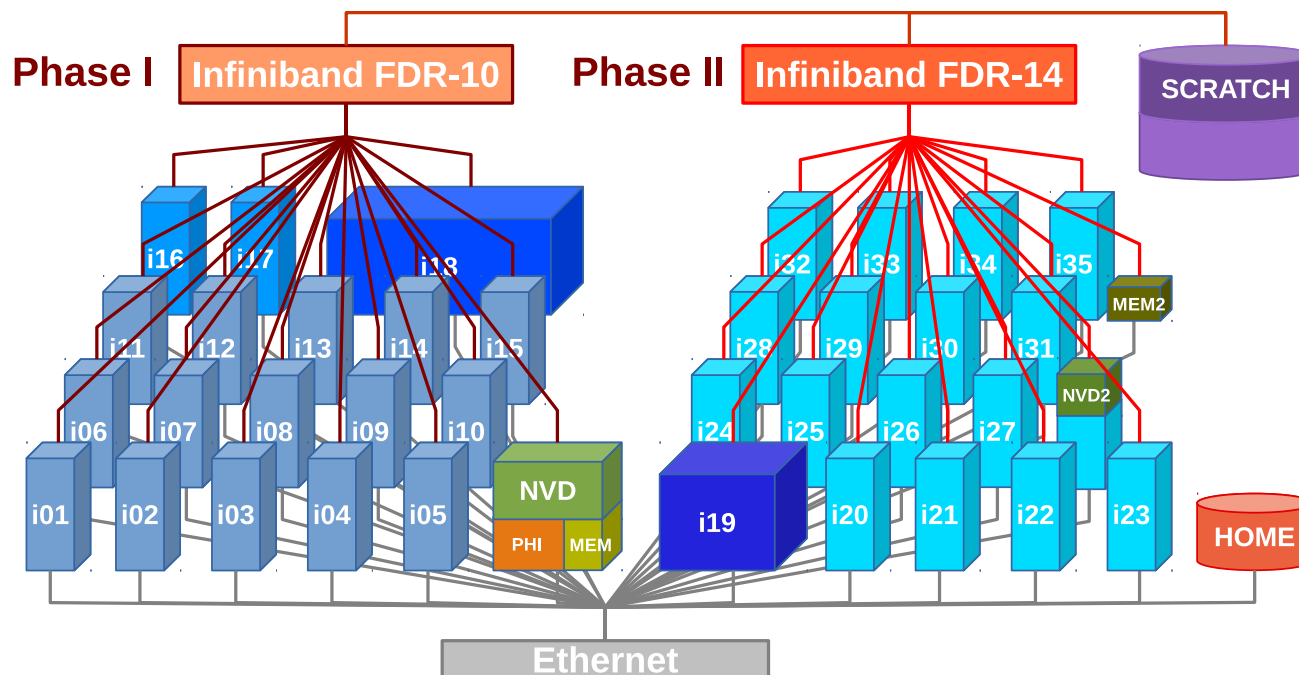
One Cluster – Multiple Islands



One Cluster – Multiple Islands (2)



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- Cluster is divided into 2 phases
- Each phase is divided into several islands
- Rule of thumb:
1 island \triangleq 32 compute nodes \triangleq 512 (phase I) / 768 (phase II) CPU cores

HARDWARE

Phase I (706+70 nodes):

•Processors:

- 2 Intel Xeon E5-2670
(**Sandy Bridge**) processors
 $\triangleq 2 \cdot 8 = \mathbf{16 \text{ CPU cores}}$
- **2.6 GHz**
(up to 3.3 GHz in turbo mode)

•Main Memory:

- **32 GB RAM** (some have 64 GB)

•Network:

- Gigabit Ethernet
- FDR-10 InfiniBand

Phase II (596+32 nodes):

•Processors:

- 2 Intel Xeon E5-2680 v3
(**Haswell**) processors
 $\triangleq 2 \cdot 12 = \mathbf{24 \text{ CPU cores}}$
- **2.5 GHz**
(up to 3.3 GHz in turbo mode)

•Main Memory:

- **64 GB RAM**

•Network:

- Gigabit Ethernet
- FDR-14 InfiniBand

Login Nodes

4 nodes (hardware similar to phase I):

- Processors:

- 4 Intel Xeon E5-4650
(**Sandy Bridge**) processors
 $\triangleq 4 \cdot 8 = \mathbf{32 \text{ CPU cores}}$
- **2.7 GHz**
(up to 3.3 GHz in turbo mode)

- Main Memory:

- **128 GB RAM**

- Network:

- 2 · 10 Gigabit Ethernet
- 2 · FDR-10 InfiniBand

8 nodes (hardware similar to phase II):

- Processors:

- 2 Intel Xeon E5-2680 v3
(**Haswell**) processors
 $\triangleq 2 \cdot 12 = \mathbf{24 \text{ CPU cores}}$
- **2.5 GHz**
(up to 3.3 GHz in turbo mode)

- Main Memory:

- **128 GB RAM**

- Network:

- 2 · 10 Gigabit Ethernet
- FDR-14 InfiniBand

Purpose of the Login Nodes

- Transferring files to the cluster
 - Editing files on the cluster
 - Compiling software
 - Testing software (time limit: 30 minutes)
 - Debugging small test cases
-
- Submitting computations to the batch system (explained later)
 - Checking status of compute jobs



ACCESS TO THE CLUSTER

Set Password for Cluster Access

1. Go to <https://www.idm.tu-darmstadt.de/ando>
2. Log in with TU-ID
3. Click on tab “HLR”
4. Set password

Connecting to the Cluster

Open an SSH connection to one of the login nodes:

- `lcluster1.hrz.tu-darmstadt.de` – `lcluster4.hrz.tu-darmstadt.de`
 - Sandy Bridge, 32 cores, 128 GB RAM
- `lcluster5.hrz.tu-darmstadt.de` – `lcluster11.hrz.tu-darmstadt.de`
 - Haswell, 24 cores, 128 GB RAM

Linux

- `ssh <tu-id>@lcluster9.hrz.tu-darmstadt.de`
- `ssh -X -C -c blowfish <tu-id>@lcluster9.hrz.tu-darmstadt.de`

X11 forwarding

Cipher (speedup)

Compression (speedup)

Windows

- Cygwin (<https://www.cygwin.com>): `ssh` as with Linux above
- PuTTY (<http://www.chiark.greenend.org.uk/~sgtatham/putty/download.html>)

File Transfer to the Cluster

Copy files between own computer and cluster

- Linux
 - `scp <local_src> <tu-id>@lcluster9.hrz.tu-darmstadt.de:<remote_dst>`
 - `scp <tu-id>@lcluster9.hrz.tu-darmstadt.de:<remote_src> <local_dst>`
- Windows
 - Cygwin (<https://www.cygwin.com>): `scp` as with Linux above
 - PuTTY (<http://www.chiark.greenend.org.uk/~sgtatham/putty/download.html>)

File Transfer to the Cluster (2)

Alternative to copying files between cluster and own computer is to mount remote cluster file system via SSH on own computer

- Linux
 - SSHFS
- Windows
 - Win-sshfs
 - NetDrive



SOFTWARE

- Linux distribution CentOS 7.5
- Some programs are included in the operating system
 - You can just use these programs, e.g., text editors such as **vim**, **nano**, **mc**, **gedit** for directly editing files on the login nodes
 - Only a limited number of packages is provided by the operating system

- A lot of software is provided by the module system
 - Compilers (GCC, Intel, ...)
 - Libraries (MPI, CUDA, ...)
 - Tools (SVN, Git, GDB, ...)
 - Application software
- Use **module avail** or **module whatis** to obtain a complete list of all available modules
- Use **module help <modulename>** to get further details about specific module

- To load a module, use the command `module load <modulename>`
 - Read the hints that might appear
 - Sometimes you have to load additional modules
 - Typically, the module name has to contain the version number, e.g. `ansys/17.1`
 - For some modules, a default version is set
E.g., it may be sufficient to write `module load gcc`, but not `module load openmpi` – choose `module openmpi/gcc` or `openmpi/intel`
- To show all currently loaded modules, use `module list`
- To remove a module, use `module unload <modulename>`
- To remove all modules, use `module purge`



QUESTIONS & DEMO LOGIN



USING THE CLUSTER

- Do not confuse the login nodes with the cluster
 - There are only few login nodes but >1000 compute nodes
 - Login nodes are the point of access to the cluster
 - Connecting to compute nodes not possible
- **Use the batch system to submit jobs from the login nodes**
 - Write **job script** file with all job parameters and commands and submit the script
- The cluster runs 24/7 (including weekends)

Job Script Example



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```
#!/bin/bash
```

```
#SBATCH -J my_job
#SBATCH --mail-user=my@mail.com
#SBATCH --mail-type=ALL
#SBATCH -e /home/<tu-id>/somewhere/Job_Name.err.%j
#SBATCH -o /home/<tu-id>/somewhere/Job_Name.out.%j
#SBATCH --mem-per-cpu=250
#SBATCH -t 00:05:00
#SBATCH -n 4
#SBATCH --account=kurs00025
#SBATCH --partition=kurs00025
#SBATCH --reservation=kurs00025
```

Job parameters

```
echo "This is Job $SLURM_JOB_ID"
```

```
module load gcc
module load openmpi/gcc
mpirun hostname
```

Commands to run your program

Job Script Example (2)



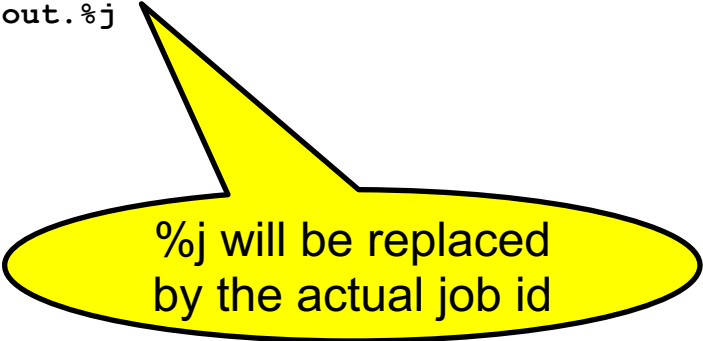
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```
#!/bin/bash

#SBATCH -J my_job
#SBATCH --mail-user=my@mail.com
#SBATCH --mail-type=ALL
#SBATCH -e /home/<tu-id>/somewhere/Job_Name.err.%j
#SBATCH -o /home/<tu-id>/somewhere/Job_Name.out.%j
#SBATCH --mem-per-cpu=250
#SBATCH -t 00:05:00
#SBATCH -n 4
#SBATCH --account=kurs00025
#SBATCH --partition=kurs00025
#SBATCH --reservation=kurs00025
```

```
echo "This is Job $SLURM_JOB_ID"
```

```
module load gcc
module load openmpi/gcc
mpirun hostname
```



%j will be replaced
by the actual job id

You should always set the following parameters:

- **-n <tasksnum>**: Number of tasks (processes)
Default: one CPU core per task
- **--mem-per-cpu=<memory>**: Maximum memory per core in MB
- **-t <time>**: Time limit of the job (in minutes)
Maximum time limit is 5 min for SPP
- **--partition=kurs00025**
- **--account=kurs00025**
- **--reservation=kurs00025**

Additional Job Parameters

- **-J <jobname>**: Name of the job (does not have to be unique)
- **--mail-user=<email>**: Email address
- **--mail-type=<BEGIN | END | ALL | [...]>**: Send an email
- **-o <filename>**: Write standard output ('stdout') of the job to a file
- **-e <filename>**: Write standard error ('stderr') of the job to a file
- **-c <number>**: Number of CPU cores per task
Important for OpenMP with several cores per process

Batch System Commands

- **sbatch** **<JobScriptFile>**: Submit the job
- **squeue**: List all your jobs
- **sjobs** **<JobID>**: Print detailed information about a job
- **scancel** **<JobID>**: Kill a particular job
- **scancel -u** **<TU-ID>**: Kill all your own jobs

Job was submitted but does not start

- Job does not start when it exceeds limits
- Time limit is 5 min
- Memory per core should not exceed 1600 MB
- Memory per task must not exceed 26 GB
- We have three exclusive nodes with 16 cores and 26 GB each
- When job limits are correct, job only starts when resources on the three nodes are available
- Job start might be delayed due to jobs of other students

Further Information



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<http://www.hhlr.tu-darmstadt.de>



QUESTIONS & DEMO JOB