

Practical Example: High Performance Computing

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The Operating System UNIX

- UNIX Development
 - 60s: MULTICS: MULTiplexed Informaion and Computing Service
 - Developed by MIT, Bell Labs and General Electric
 - Written in Programming Language 1 (PL1) and Assembly
- Thompson and Ritchie later create UNIX, written in C

Linux – Basic Commands (Revisit)

- `command parameter1 parameter2 ...`
- *man* – **man**ual. Displays manuals for linux packages (try *man man*). *Displays useful information about how to use the package.*
- *pwd* – **p**rint **w**orking **d**irectory. Displays the current working directory

Linux – Basic Commands (Revisit)

- `ls` – **list**. Displays the content of a directory
 - `ls -l ~/Pictures`

```
-rw-r--r--    1 martin martin 101685 Apr 29 13:23 Selection_001.png
-rw-r--r--    1 martin martin  39835 Apr 29 14:37 Selection_002.png
-rw-r--r--    1 martin martin  76985 Apr 29 14:48 Selection_003.png
-rw-r--r--    1 martin martin  27096 May  2 14:04 Selection_004.png
-rw-r--r--    1 martin martin  62783 May  3 11:58 Selection_005.png
-rw-r--r--    1 martin martin  65201 May  6 16:11 Selection_006.png
drwxrwxr-x    2 martin martin  4096 Apr 24 11:41 Temp
```



Access
rights

Links

Owners/
Groups

Size

Last
Modified

Filename

Linux – Basic Commands (Revisit)

- `ls` – **list**. Displays the content of a directory

```
-rw-r--r--    1 martin martin 101685 Apr 29 13:23 Selection_001.png
```

- Access rights:
 - Type: (**d**)irectory, (**l**)ink, (**-**) a file
 - Rights: (**r**)ead, (**w**)rite, e(**x**)ecute
 - Sequence: *user group anybody*
- Owner/Group:
 - Owner – first column: *martin*
 - Group – second column: *martin*
- Either can be changed with *chmod* and *chgrp* respectively
- Run executable with full path or from directory with “*./file*”

Linux – Basic Commands (Revisit)

- *cd* – **c**hange **d**irectory. Switches to a new directory, supplied as a parameter
- *mkdir* – **m**ake **d**irectory. Creates a new directory
- *rm* – **r**emove. Removes a specified file or directory (“-r”)
- *cp* – **c**opy. Copy a file or directory (“-r”)
- *scp* – **s**ecure **c**opy. Copy a file to or from a remote source
- *rsync* – a fast copying tool. Also works for remote copy

Linux – Basic Commands (Revisit)

- *ln* – **link**. Link a file into a new directory
- *echo* – Prints a string to standard output
- *cat* – **con**catenate. The content of a file is printed to standard output
- *wc* – **w**ord **c**ount. Counts the number of words, rows (“-l”) pr characters (“-c”) in a file.

Linux – Basic Commands (Revisit)

- pipe (“|”) - connect commands
- output (“>”) and input (“<”) for a specific program.
Output can also concatenate to existing content, without deleting (“>>”)

Linux – Environment Variables

Variable	Description
PATH	Colon separated list of directories, which will be searched through when entering a name of executable
HOME	The pathname of the home directory.
SHELL	The currently used shell program
USER	The current username

```
$ echo $PATH  
/usr/local/sbin
```

```
$ export PATH=~/.local/bin:$PATH  
$ echo $PATH  
/home/martin/.local/bin:/usr/local/sbin
```

Linux – Vim

- A further development of Vi
- Open-Source editor for use inside the Terminal
- Open file with “\$ vim filename”
 - Type “i” for insert (edit)
 - ESC to go back to entry menu
 - :q – exit without saving
 - :wq – save and exit
 - :q! - force exit without saving

Linux – Bash Scripts – Exercise

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

```
echo Download some sample data
```

```
wget https://www.costalab.org/wp-content/uploads/2019/05/data2.zip
```

```
echo Unzipping the data
```

```
unzip data2.zip
```

```
echo Removing the archive
```

```
rm data2.zip
```

```
echo Data ready to work with
```

- Write in a file with `.sh` – extension
- Change the file mods bits
 - `chmod 770 somefile.sh`
- Execute the file with `./somefile.sh`

RWTH Compute Cluster

- High Performance Computing Infrastructure
- Linux-Based
- Highly Parallelizable
- All RWTH - affiliates granted access, but with limited resource allocation
- More information on:
<https://doc.itc.rwth-aachen.de/display/CC/General>



RWTH Selfservice

- Use Selfservice (<https://idm.rwth-aachen.de/selfservice/>) to manage university accounts and create HPC-account to user the Cluster

<input type="checkbox"/>	Service/Account	Username
<input type="checkbox"/>	RWTH Single Sign-On (former CRIFUS-Office/Student)	mm844865
<input type="checkbox"/>	Hochleistungsrechnen RWTH Aachen	mm844865
<input type="checkbox"/>	Hochleistungsrechnen RWTH Aachen	mm844865
<input type="checkbox"/>	Hochleistungsrechnen RWTH Aachen	mm844865
<input type="checkbox"/>	Hochleistungsrechnen RWTH Aachen	mm844865
<input type="checkbox"/>	Life- und Lemporal LP	mm844865
<input type="checkbox"/>	PC-Pool	mm844865
<input type="checkbox"/>	RWTH Service	mm844865@uni.rwth-aachen.de
<input type="checkbox"/>	RWTH Service	mm844865@uni.rwth-aachen.de
<input type="checkbox"/>	RZ-Ticketbot	mm844865
<input type="checkbox"/>	TIM Service	mm844865
<input type="checkbox"/>	WL/AN/PH	mm844865

SSH and Cluster login

- Use established *secure shell protocol (ssh)* connection to log in to the front-end cluster nodes.

```
$ ssh <Username>@<Servername(or IP)>
```

SSH and Cluster login

Server-name	OS	Purpose
login18-1.hpc.itc.rwth-aachen.de	Linux (CentOS 7)	Front-End Dialogue System for CLAIX2018
login18-2.hpc.itc.rwth-aachen.de	Linux (CentOS 7)	Front-End Dialogue System for CLAIX2018
login18-3.hpc.itc.rwth-aachen.de	Linux (CentOS 7)	Front-End Dialogue System for CLAIX2018
login18-4.hpc.itc.rwth-aachen.de	Linux (CentOS 7)	Front-End Dialogue System for CLAIX2018
login.hpc.itc.rwth-aachen.de	Linux (CentOS 7)	Front-End Dialogue System for CLAIX2016
login2.hpc.itc.rwth-aachen.de	Linux (CentOS 7)	Front-End Dialogue System for CLAIX2016
login18-g-1.hpc.itc.rwth-aachen.de	Linux (CentOS 7)	GPU-System (2018)
login18-g-2.hpc.itc.rwth-aachen.de	Linux (CentOS 7)	GPU-System (2018)
login-g.hpc.itc.rwth-aachen.de	Linux (CentOS 7)	GPU-System (2016)

SSH and Cluster login

- Use established *secure shell protocol (ssh)* connection to log in to the front-end cluster nodes.

```
$ ssh <Username>@<Servername(or IP)>
```

- In the case of the RWTH Cluster:

```
$ ssh <TIM>@<login18-1.hpc.itc.rwth-aachen.de>
```


SSH and Cluster login

- Special nodes for intensive IO operations

copy18-1.hpc.itc.rwth-aachen.de	CentOS
copy18-2.hpc.itc.rwth-aachen.de	CentOS
copy.hpc.itc.rwth-aachen.de	CentOS

- Use this nodes if you ever need to transfer big files to the cluster

Cluster – available file systems

Name	Path	Backup	Quota (file)	Quota (#files)
\$HOME	<i>/home/</i> <TIM-Kennung>	yes	1000 GB	1000
\$WORK	<i>/work/</i> <TIM-Kennung>	no	250 GB	1000
\$HPCWORK	<i>/hpcwork/</i> <TIM-Kennung>	no	1000 GB	50000

Cluster – mount drive

- Use *sshfs* to mount remote directory to local machine:
 - `$ sshfs <TIM>@copy18-1.hpc.itc.rwth-aachen.de:<Path> <local Path>`
- Unmount with:
 - `$ sudo umount -l <local Path>`
- Only works if you are inside eduroam main-network or through RWTH VPN
 - <https://doc.itc.rwth-aachen.de/display/VPN/VPN+%28Ubuntu%29+Cisco+OpenConnect>

Cluster – Batch Jobs

- Since 2018 new front-end nodes, new jobs-handling system – SLURM
- SBATCH [options] command [arguments]
- Slots

Parameter	Function
-c, --cpus-per-task <numcpus>	Number of threads/processes for an OpenMP/Hybrid script
-n, --ntasks <numtasks> --ntasks-per-node <numtasks>	Number of threads/processes for an MPI job
-N, --nodes <numnodes>	Number of nodes/hosts for the job

- List of all available nodes:

<https://doc.itc.rwth-aachen.de/display/CC/Hardware+of+the+RWTH+Compute+Cluster>

Cluster – Batch Jobs

- Further job parameters

Parameter	Function
-J --job-name=<jobname>	A name for the current job
--mem-per-cpu=<size>	Required RAM per allocated CPU
-o, --output=<filename>	Name for a report file, containing the standard output of the job
-t, --time=d-hh:mm:ss	Time for job execution, after this time the job is killed
-A, --account=<project>	Submit a job for a specific project
--gres=gpu:<type>:2	Requesting two GPUs per node

Cluster – Batch Jobs

- **sbatch** jobscript.sh – To run a job, specified in jobscript.sh
- At the end of the jobscript.sh file add a line for execution a script of programm
- Might need to load some modules prior, e.g.:
 - *module load python/3.6.0*
 - *module avail* – to list all available modules
- Set environment variable inside a cluster job script
 - *export PATH=/usr/local_rwth/sw/cuda/9.0.176/bin:\$PATH*

Cluster – Example Scripts

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

```
### Job name
```

```
#SBATCH --job-name=MYJOB
```

```
### File for the output
```

```
#SBATCH --output=MYJOB_OUTPUT
```

```
### Time your job needs to execute, e. g. 15 min 30 sec
```

```
#SBATCH --time=00:15:30
```

```
### Memory your job needs per node, e. g. 1 GB
```

```
#SBATCH --mem=1G
```

```
### The last part consists of regular shell commands:
```

```
### Change to working directory
```

```
cd /home/usr/workingdirectory
```

```
### Execute your application
```

```
myapp.exe
```

Cluster – Monitor Jobs and Resources

- Use ***squeue*** to monitor current jobs in progress:
 - `squeue -u <TIM>` for only jobs by specific users

- Use ***scancel*** to stop current jobs from executing
 - `scancel -u <TIM>` for killing jobs by specific user
 - `scancel -n <Jobname>` for killing a specific job

- Use ***quota*** to see occupied space on the hard-drives and used number of files.

- Use ***sinfo*** to see updated accounting information. Currently no monthly CPU-time quotas in place.

Cluster – GPU Nodes

- Login to a GPU-Node by:
ssh <TIM>@login18-g-1.hpc.itc.rwth-aachen.de
- See GPU usage with nvidia-smi

Tue May 7 17:24:20 2019

```
+-----+
| NVIDIA-SMI 418.43          Driver Version: 418.43          CUDA Version: 10.1          |
+-----+-----+-----+-----+-----+
| GPU  Name                Persistence-M| Bus-Id        Disp.A | Volatile Uncorr. ECC |
| Fan  Temp   Perf          Pwr:Usage/Cap|      Memory-Usage | GPU-Util  Compute M. |
+-----+-----+-----+-----+-----+
|   0   Tesla V100-SXM2...    Off          | 00000000:62:00:0 Off |             0         |
| N/A   42C    P0           54W / 300W |  0MiB / 16130MiB |    0%    E. Process  |
+-----+-----+-----+-----+-----+
|   1   Tesla V100-SXM2...    Off          | 00000000:89:00:0 Off |             0         |
| N/A   42C    P0           54W / 300W |  0MiB / 16130MiB |    0%    E. Process  |
+-----+-----+-----+-----+-----+

+-----+-----+-----+-----+-----+
| Processes:                                                       GPU Memory |
|  GPU       PID    Type   Process name                               Usage    |
+-----+-----+-----+-----+-----+
| No running processes found                                     |
+-----+-----+-----+-----+-----+
```

Cluster – GPU Nodes

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```
Tue May 7 17:24:20 2019
+-----+
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| GPU   Name                 Persistence-M| Bus-Id        Disp.A | Volatile Uncorr. ECC |
| Fan  Temp  Perf    Pwr:Usage/Cap|      Memory-Usage | GPU-Util  Compute M. |
+-----+-----+-----+-----+-----+
|  0   Tesla V100-SXM2...    Off          | 00000000:62:00:0 Off  |      0%      E. Process |
| N/A   42C    P0     54W / 300W |  0MiB / 16130MiB |              |
+-----+-----+-----+-----+-----+
|  1   Tesla V100-SXM2...    Off          | 00000000:89:00:0 Off  |      0%      E. Process |
| N/A   42C    P0     54W / 300W |  0MiB / 16130MiB |              |
+-----+-----+-----+-----+-----+
+-----+-----+-----+-----+
| Processes:                                                       GPU Memory |
|  GPU       PID    Type   Process name                               Usage      |
+-----+-----+-----+-----+-----+
| No running processes found                                     |
+-----+-----+-----+-----+-----+

```

GPU-id
and type

Cluster – GPU Nodes

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Tue May 7 17:24:20 2019
+-----+
| NVIDIA-SMI 418.43          Driver Version: 418.43          CUDA Version: 10.1          |
+-----+-----+-----+-----+-----+
| GPU   Name                 Persistence-M| Bus-Id        Disp.A | Volatile Uncorr. ECC |
| Fan  Temp  Perf    Pwr:Usage/Cap|      Memory-Usage | GPU-Util  Compute M. |
+-----+-----+-----+-----+-----+
|  0   Tesla V100-SXM2...    Off          | 00000000:62:00:0 Off |             0          |
| N/A   42C    P0      54W / 300W |  0MiB / 16130MiB |    0%    E. Process |
+-----+-----+-----+-----+-----+
|  1   Tesla V100-SXM2...    Off          | 00000000:89:00:0 Off |             0          |
| N/A   42C    P0      54W / 300W |  0MiB / 16130MiB |    0%    E. Process |
+-----+-----+-----+-----+-----+
| Processes:                                                       GPU Memory |
|  GPU       PID    Type   Process name                               Usage      |
+-----+-----+-----+-----+-----+
| No running processes found                                         |
+-----+-----+-----+-----+-----+
```

GPU-id
and type

GPU memory

Cluster – GPU Nodes

- Login to a GPU-Node by:
ssh <TIM>@login18-g-1.hpc.itc.rwth-aachen.de
- See GPU usage with nvidia-smi

Compute model: 1 person

```
Tue May 7 17:24:20 2019
```

NVIDIA-SMI 418.43			Driver Version: 418.43			CUDA Version: 10.1		
GPU	Name	Persistence-M	Bus-Id	Disp.A	Volatile Uncorr.	ECC		
Fan	Temp	Perf	Pwr:Usage/Cap	Memory-Usage	GPU-Util	Compute M.		
0	Tesla V100-SXM2...	Off	00000000:62:00:0	Off	0			
N/A	42C	P0	54W / 300W	0MiB / 16130MiB	0%	E. Process		
1	Tesla V100-SXM2...	Off	00000000:89:00:0	Off	0			
N/A	42C	P0	54W / 300W	0MiB / 16130MiB	0%	E. Process		
Processes:							GPU Memory Usage	
GPU	PID	Type	Process name					
No running processes found								

GPU-id and type

GPU memory

Cluster – Exercise – GPU with python

- Sign in to the cluster
\$ ssh <TIM>@login18-g-2.hpc.itc.rwth-aachen.de
- Check if you're successfully in the group:
\$ groups # nova0028 should be there
- Load python3.6 module and install some libraries locally
\$ module load python/3.6.0
\$ pip3.6 install --user matplotlib tensorflow_gpu keras
- Create a new directory and copy the files needed
\$ mkdir BioInfo
\$ rsync -r /hpcwork/nova0028/BioInfoLab2019/examples BioInfo
\$ cd BioInfo/examples/GPU
- Make start_job.sh executable (\$ chmod 770 <filename>)
- **Edit** (with **vim**) your email address and submit job (\$ sbatch start_job.sh)
- Check status regularly (\$ squeue -u <TIM>)

Cluster – Exercise – CPU with R

- Open R and install some packages

```
$ R
> install.packages("dplyr")
> install.packages("Seurat')
```
- Navigate to the folder

```
$ cd ~/BioInfo/examples/CPU
```
- Make start_job.sh executable (`$ chmod 770 <filename>`)
- **Edit** (with vim) your email address and submit job (`$ sbatch start_job.sh`)
- Check status regularly (`$ squeue -u <TIM>`)

Further Readings

- Linux tutorials:
 - <https://www.tutorialspoint.com/unix/>
 - <https://ryanstutorials.net/linuxtutorial/>
- SLURM – batch system
 - <https://slurm.schedmd.com/documentation.html>
 - <https://doc.itc.rwth-aachen.de/display/CC/Using+the+SLURM+Batch+System>
 - https://hpc-wiki.info/hpc/SLURM#Jobscript_Examples