

Getting started with the scientific cluster

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Introduction : What is a high-performance computing cluster ?

Typical processor (CPU) in a personal laptop :

109 Products [COMPARE ALL](#)

Product Name	Launch Date	# of Cores	Max Turbo Frequency	Cache	Processor Graphics ‡
<input type="checkbox"/> Intel® Core™ i5 processor 14401E (24M Cache, up to 4.70 GHz)	Q3'24	6	4.7 GHz	24 MB Intel® Smart Cache	Intel® UHD Graphics 730
<input type="checkbox"/> Intel® Core™ i5 processor 14401TE (24M Cache, up to 4.50 GHz)	Q3'24	6	4.5 GHz	24 MB Intel® Smart Cache	Intel® UHD Graphics 730
<input type="checkbox"/> Intel® Core™ i5 processor 14501TE (24M Cache, up to 5.10 GHz)	Q3'24	6	5.1 GHz	24 MB Intel® Smart Cache	Intel® UHD Graphics 770

Introduction : What is a high-performance computing cluster ?

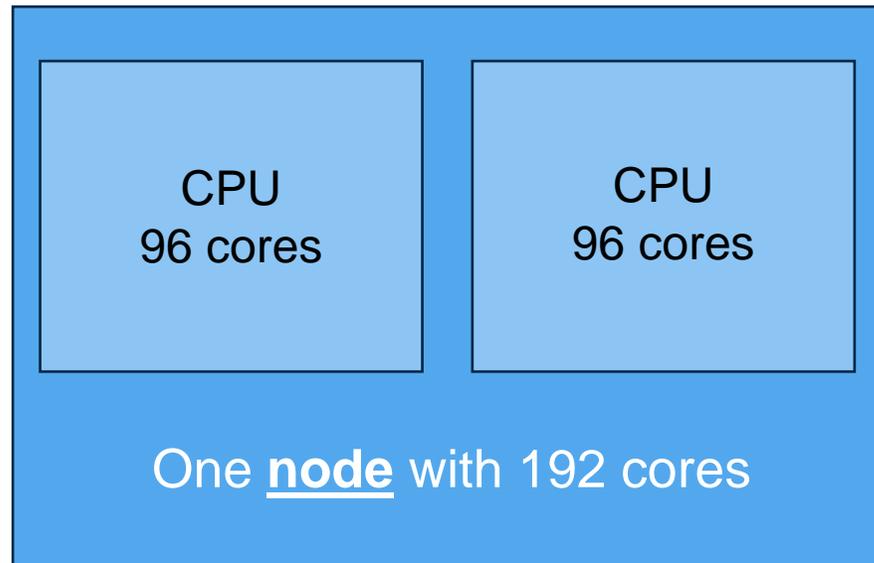
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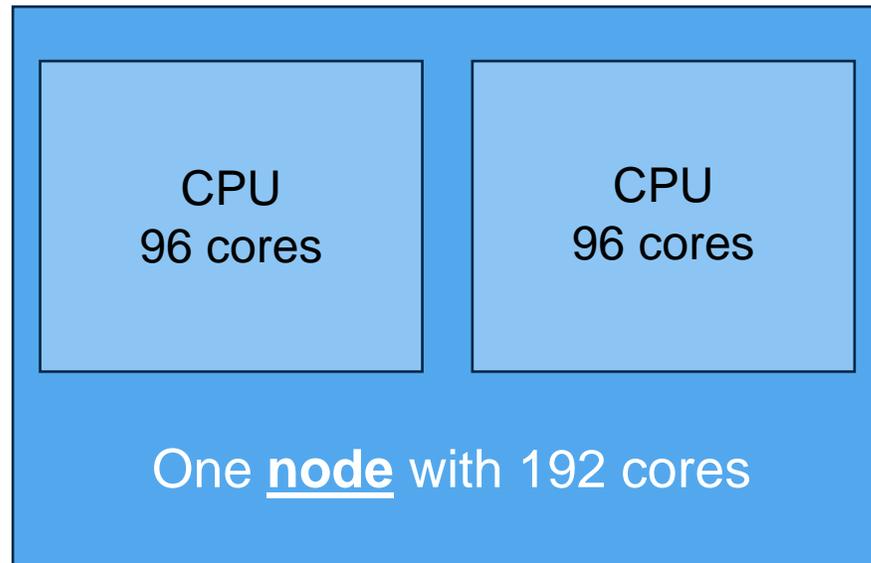
Euler is a shared system among ~4000 users that has, at this point, has ~160k cores. Here is a schematic of a typical architecture :



Typical clockspeed: ~2.3-2.4 GHz
(much slower than recent personal workstations)

Introduction : What is a high-performance computing cluster ?

Euler is a shared system among ~4000 users that has, at this point, has ~160k cores. Here is a schematic of a typical architecture :



Typical use cases of an HPC cluster :

- Parallelisation of applications
- Usage of special hardware (GPU)
- Marginally, running 1-core codes for long times

Outlook

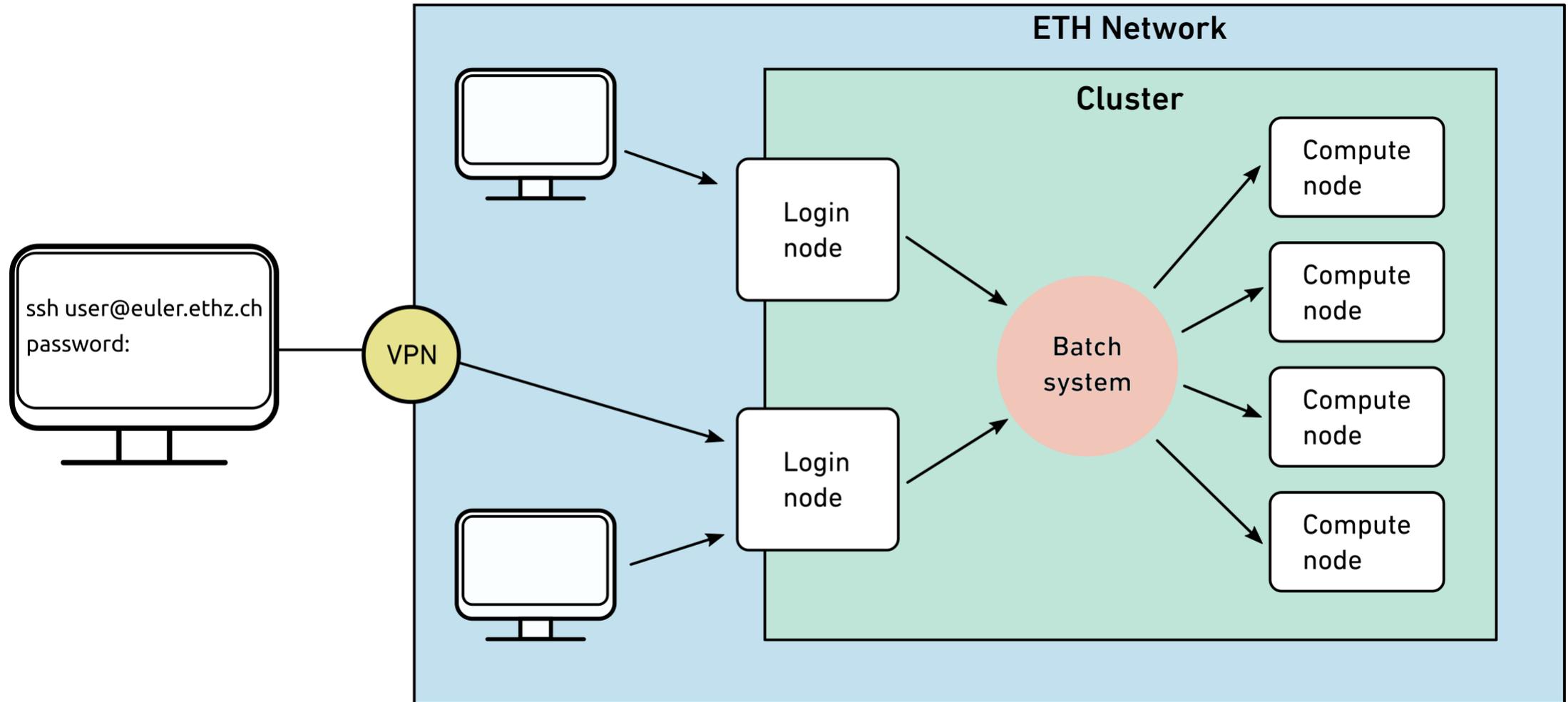
- Accessing the cluster
- Storage and data transfer
- Modules and applications
- Using the batch system

https://scicomp.ethz.ch/wiki/Main_Page

Outlook

- Accessing the cluster
- Storage and data transfer
- Modules and applications
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Accessing the cluster

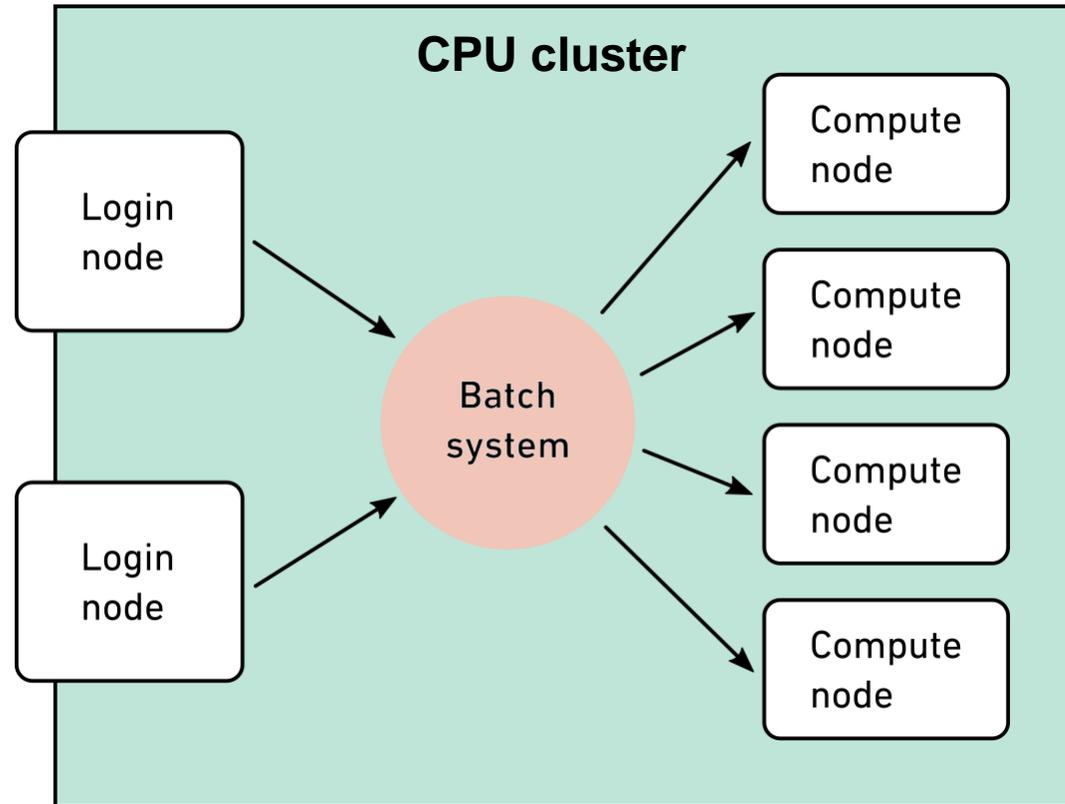


Access > Who can use the cluster > CPU cluster

All ETH members (they can use the cluster as guest users with limited resources)

Shareholders that invested into a share of the cluster resources

External collaboration partners of shareholders



Access > Prerequisites

- A valid ETH account
- Local computer with an SSH client
 - Linux and macOS contain SSH client as part of the operating system
 - Windows users need to install a third party SSH client
 - MobaXterm (<https://mobaxterm.mobatek.net/>) is a free open source SSH client that we recommend
- An X11 server for graphical user interface (optional)
 - Linux (<https://www.xorg.com>)
 - macOS (<https://xquartz.org>)
 - Windows (included in MobaXterm)

Access > Legal compliance

- The HPC clusters are subject to ETH's acceptable use policy for IT resources (Benutzungsordnung für Telematik, BOT, <https://rechtssammlung.sp.ethz.ch/Dokumente/203.21en.pdf>), in particular:
 - Cluster accounts are **strictly personal**
 - DO NOT share your account (password, ssh keys) with anyone
 - DO NOT use someone else's account, even if they say it's OK
 - If you suspect that someone used your account:
 - change your password at <https://password.ethz.ch>
 - contact cluster-support@id.ethz.ch
- Consequences
 - In case of abuse, the offender's account may be blocked temporarily or closed
 - System administrators are obliged by law to investigate abusive or illegal activities and report them to the relevant authorities

Access > How to access the clusters > ETH members

1. Start your SSH client
2. Use `ssh` command to connect to the login node of Euler

```
ssh username@euler.ethz.ch
```

3. Use your ETH credentials to login
4. First login
 - On first login a verification code is sent to your email address (username@ethz.ch)
 - By entering the verification code, your account is created automatically
 - New users must accept the cluster's usage rules

https://scicomp.ethz.ch/wiki/New_account_request_process_for_HPC_clusters

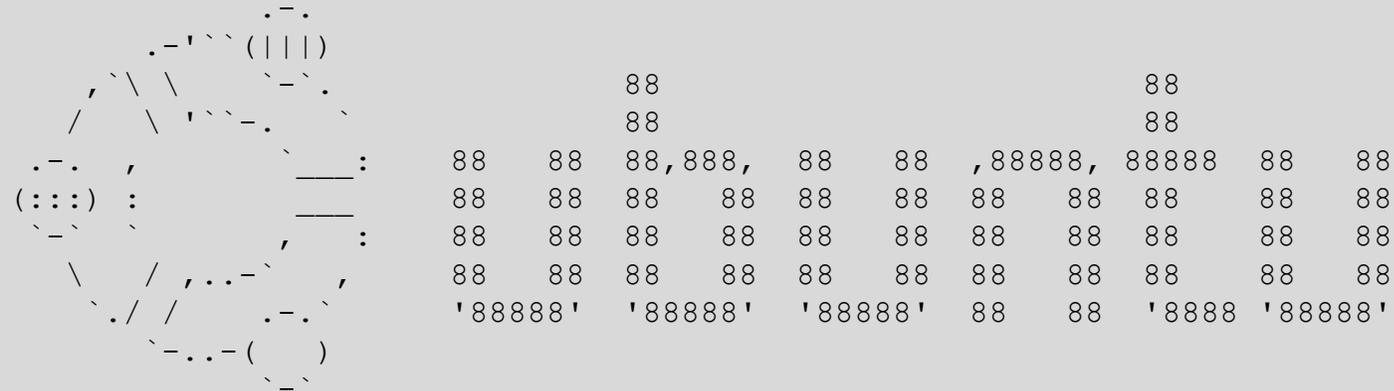
Access > SSH connection > Linux, macOS

```
nmarounina 13:13:24 >>ssh nmarounina@euler.ethz.ch
```



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Swiss Federal Institute of Technology Zurich

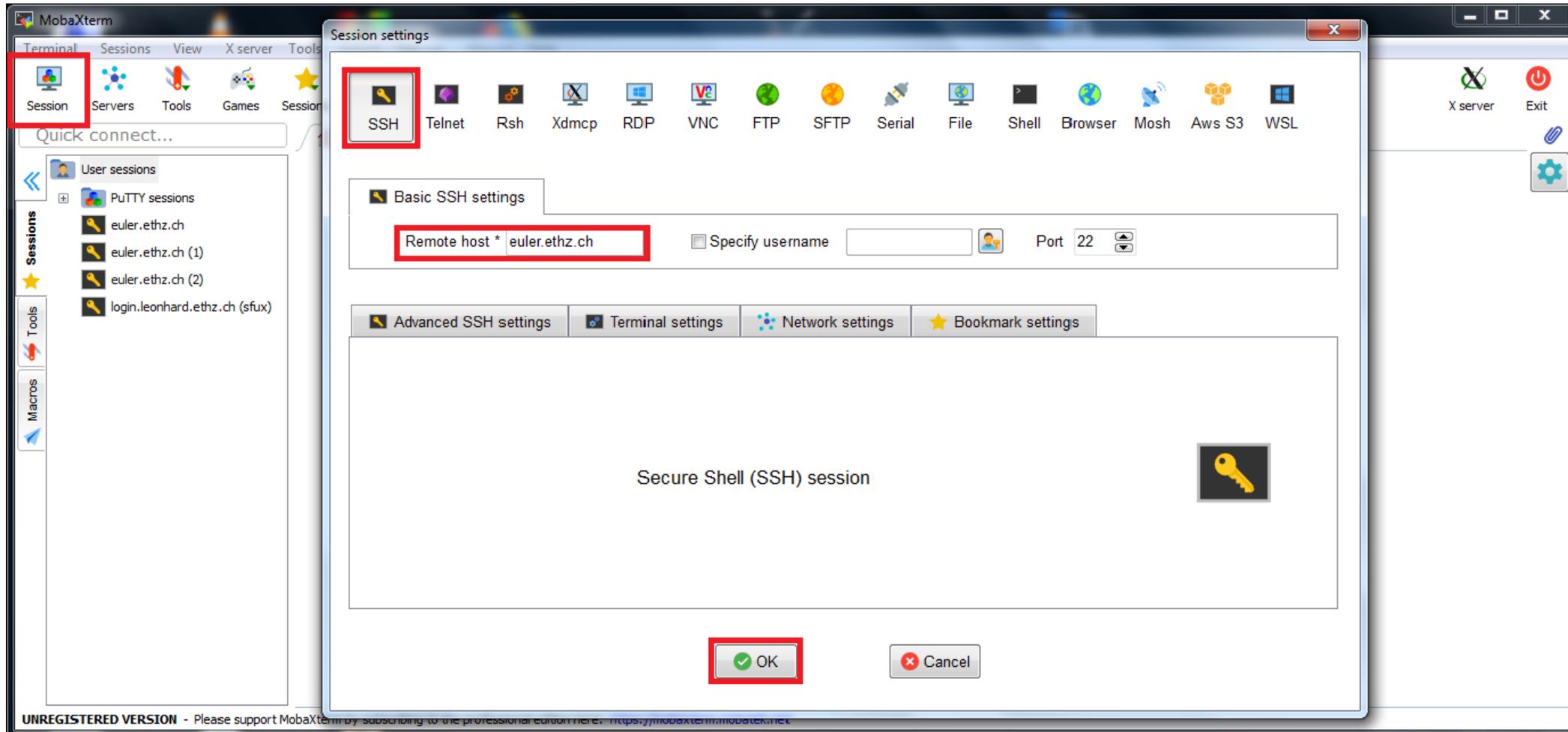
E U L E R C L U S T E R



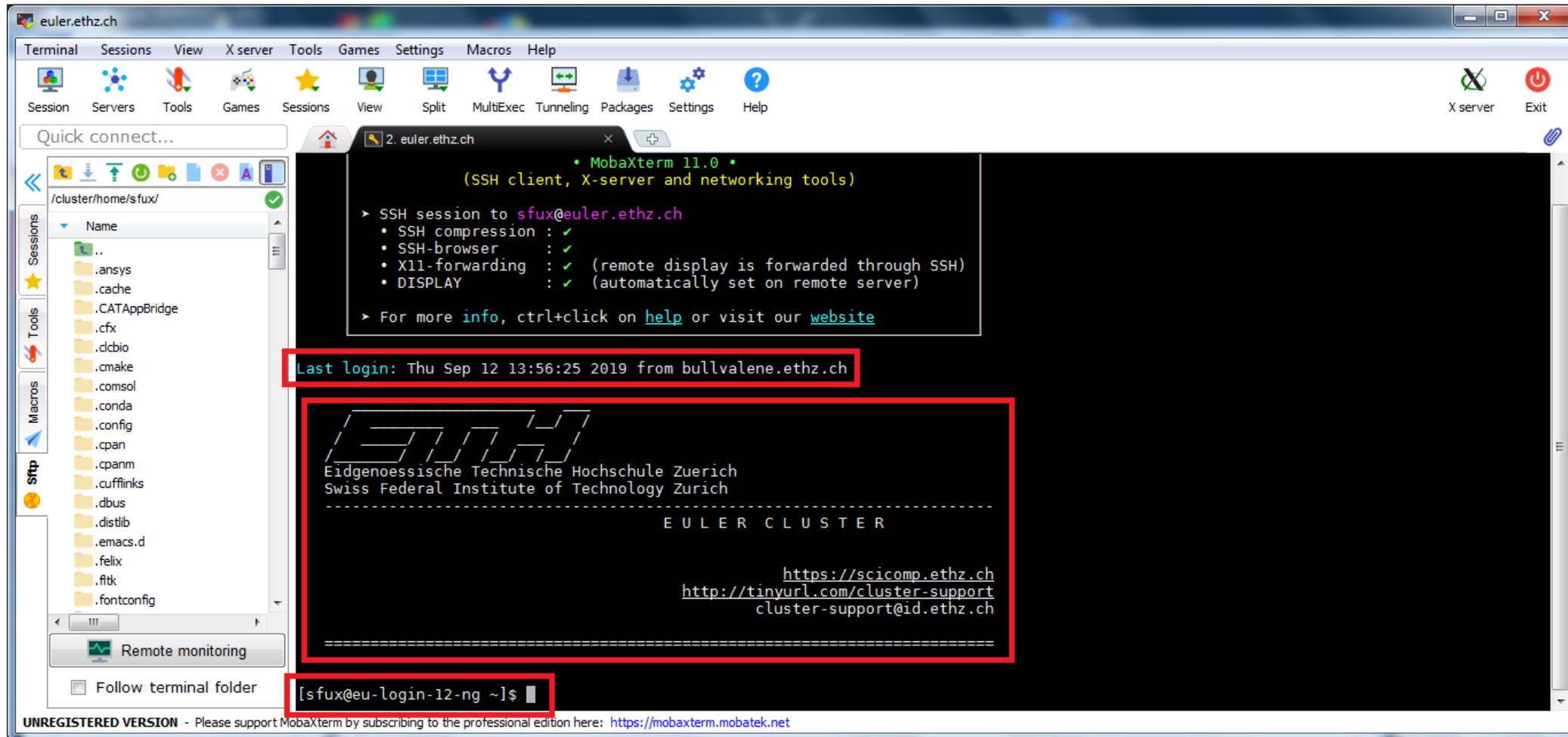
Documentation: <https://scicomp.ethz.ch>
User support: <https://smartdesk.ethz.ch> / cluster-support@id.ethz.ch
Helpdesk: https://scicomp.ethz.ch/wiki/Euler_Helpdesk
Mailing list: <https://sympa.ethz.ch/sympa/info/cluster-news>

Last login: Mon Aug 19 16:04:29 2024 from id-sis-vpn-1-a-159.ethz.ch

Access > SSH connection > Windows



Access > SSH connection > Windows



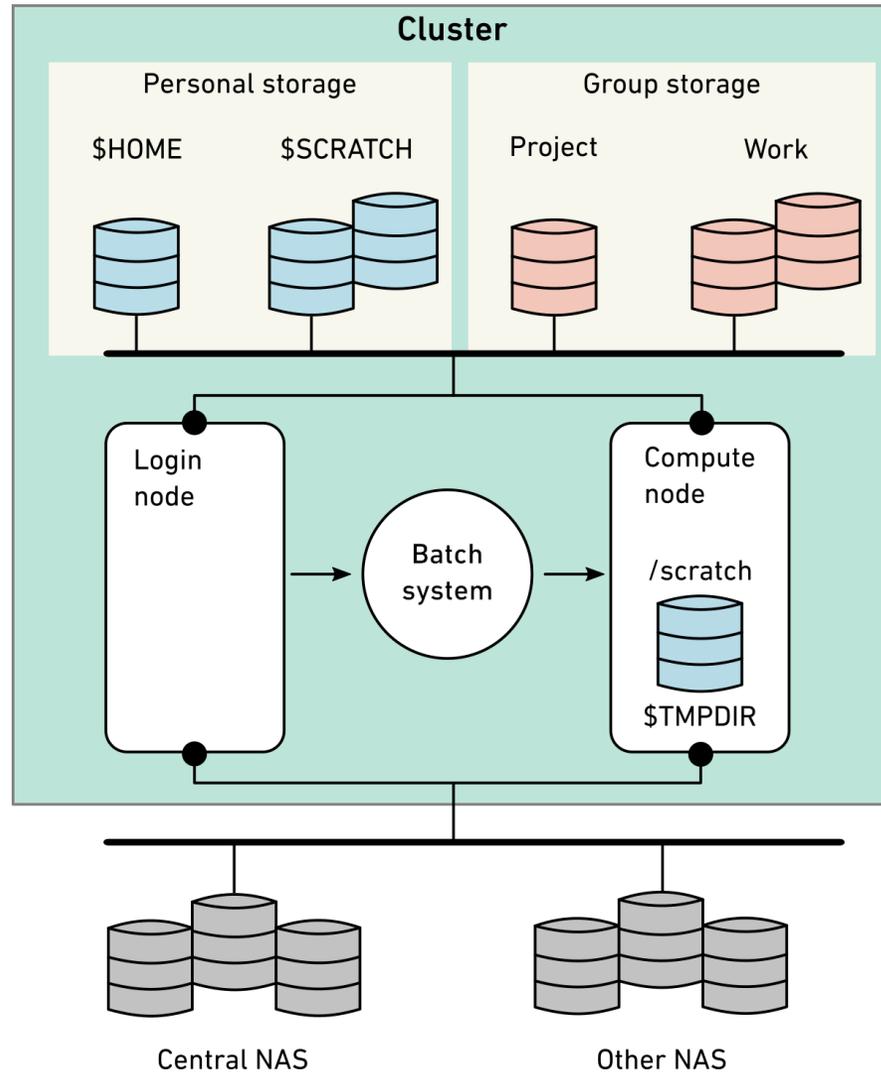
Access > SSH keys

- SSH keys allows passwordless login
 - Useful for file transfers and automated tasks
 - When used properly, SSH keys are much safer than passwords
- The procedure to create and use the SSH keys is detailed here :
https://scicomp.ethz.ch/wiki/Accessing_the_clusters#SSH_keys

Outlook

- Accessing the cluster
- Storage and data transfer
- Modules and applications
- Using the batch system

Data > Available storage systems



- Cluster wide storage systems
 - Home (personal)
 - Global scratch (personal)
 - Work (group)
 - Project (group)
- Local storage inside the compute node
 - Local scratch
- External storage
 - NAS
 - CDS
 - LTS

https://scicomp.ethz.ch/wiki/Storage_systems#External_storage

Data > Personal storage (every user) > Home

```
$ cd $HOME  
$ pwd  
/cluster/home/username
```

- Safe, long-term storage for critical data (program source, scripts, etc.)
- Accessible only by the user (owner); other people cannot read its contents
- Disk quota of 45/50 GB and a maximum of 450'000/500'000 files (soft/hard quota). Quota can be checked with the command `lquota`
- Contents saved every hour/day/week using snapshot. Users can access these snapshots in the hidden `.snapshot` directory

https://scicomp.ethz.ch/wiki/Storage_systems#Home

Data > lquota

```
nmarounina@eu-login-43:~$ lquota
```

Storage location:	Quota type:	Used:	Soft quota:	Hard quota:
/cluster/home/nmarounina	space	12.80 GB	45.00 GB	50.00 GB
/cluster/home/nmarounina	files	121385	450000	500000
/cluster/shadow	space	4.10 kB	2.15 GB	2.15 GB
/cluster/shadow	files	4	50000	50000
/cluster/scratch/nmarounina	space	71.50 GB	2.50 TB	2.70 TB
/cluster/scratch/nmarounina	files	378585	1000000	1500000

Data > Personal storage (every user) > Global scratch vs. local scratch

Global scratch

```
$ cd $SCRATCH
$ pwd
/cluster/scratch/username
```

- Fast, short-term storage for computations running on the cluster
- Created automatically upon first access and visible (mounted) only when accessed
- Disk quota of 2.5/2.7 TB and a maximum of 1m/1.5m files (soft/hard quota). Quota can be checked with the command `lquota`
- Strict usage rules; see `$SCRATCH/___USAGE_RULES___` for details
- No backup

Data > Group storage (only shareholders) > Project vs. Work

Project

`/cluster/project/groupname`

- Similar to home, but for groups
- Safe, long-term storage for critical data

Work

`/cluster/work/groupname`

- Similar to global scratch, but without purge
- Fast, short- or medium-term storage for large computations
- Visible (mounted) only when accessed

- Shareholders can buy as much space as they need
- The access rights are managed by the owner
- Quota can be checked with `lquota`
- Backed up multiple times per week

Data > File system comparison

File system	Life span	Snapshot	Backup	Max size	Small files	Large files
/cluster/home	Permanent	Yes	Yes	50 GB	+	0
/cluster/scratch	2 weeks	-	-	2.7 TB	0	++
/cluster/project	4 years	Optional	Yes	Flexible	+	+
/cluster/work	4 years	-	Yes	Flexible	0	++
local /scratch	Job	-	-	800 GB	++	0
central NAS	Flexible	Yes	Optional	Flexible	+	+

Retention time

Snapshots: up to 1 week

Backup: up to 90 days

Data > Copying data from/to the cluster (command line)

Secure copy (`scp`) is most commonly used to transfer files

```
scp [options] source destination
```

Examples: All the following examples need to be run on your local computer

- Upload a file from your workstation to Euler

```
scp local_file username@euler.ethz.ch:/path/to/remotedir
```

- Download a file from Euler to your workstation

```
scp username@euler.ethz.ch:/path/to/remote_file /path/to/localdir
```

- Copy a whole directory

```
scp -r localdir username@euler.ethz.ch:remotedir
```

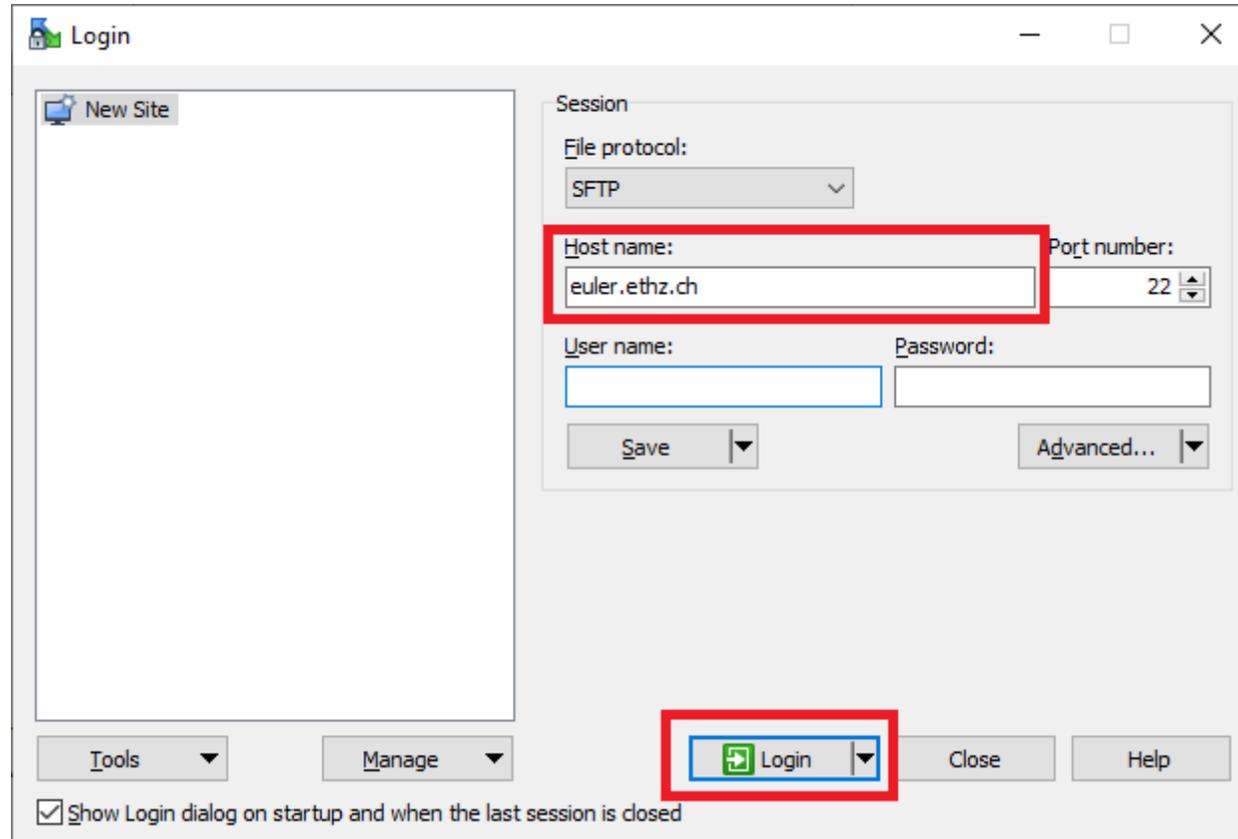
Alternatives to `scp`: `sftp`, `rsync`, `svn`, `git`, `wget`

Data > Copying data from/to the cluster (graphical user interface)

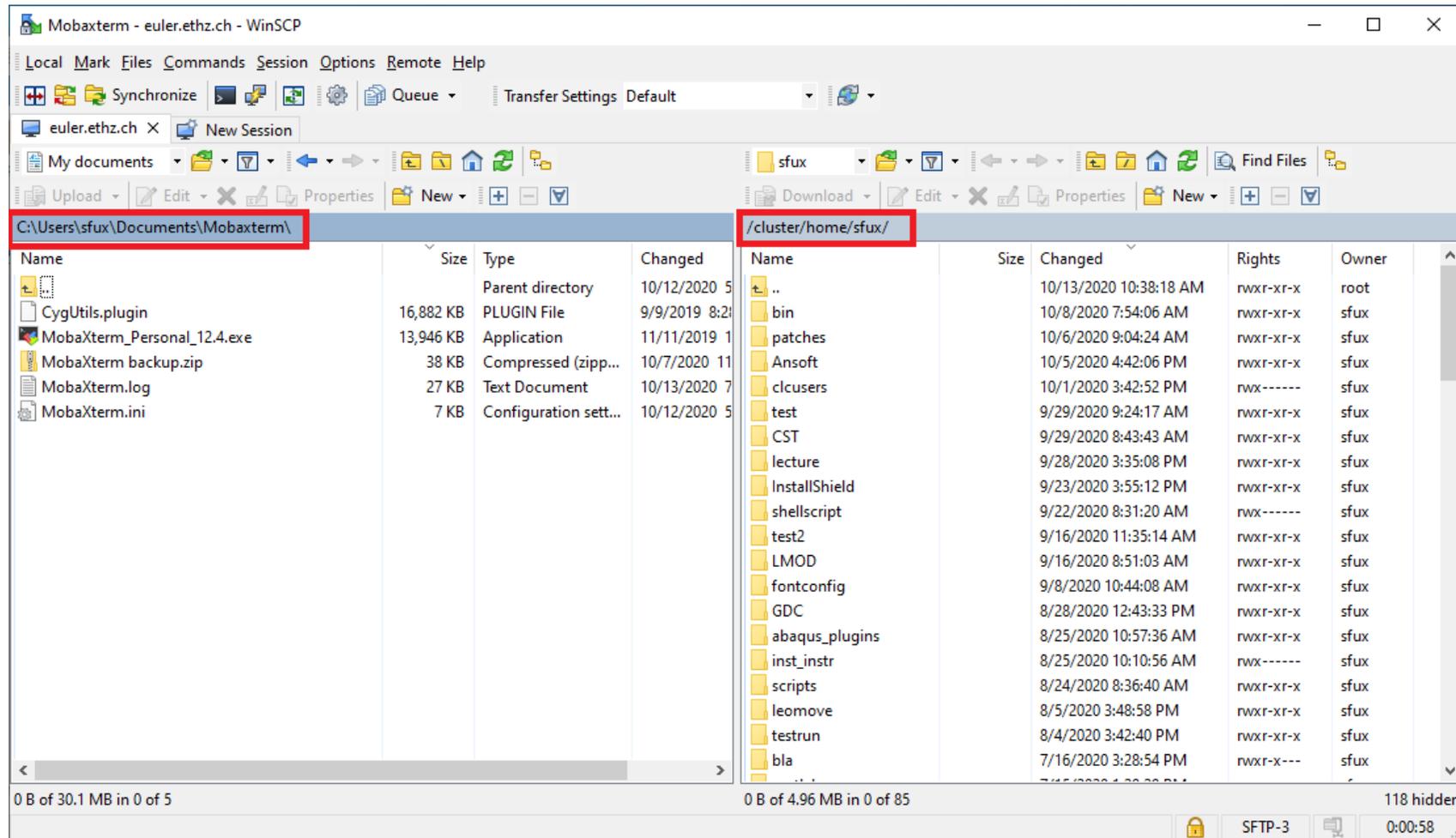
Graphical file transfer programs

Linux	macOS	Windows
FileZilla	FileZilla Cyberduck	WinSCP PSCP FileZilla Cyberduck

Data > Copying data from/to the cluster > WinSCP



Data > Copying data from/to the cluster > WinSCP

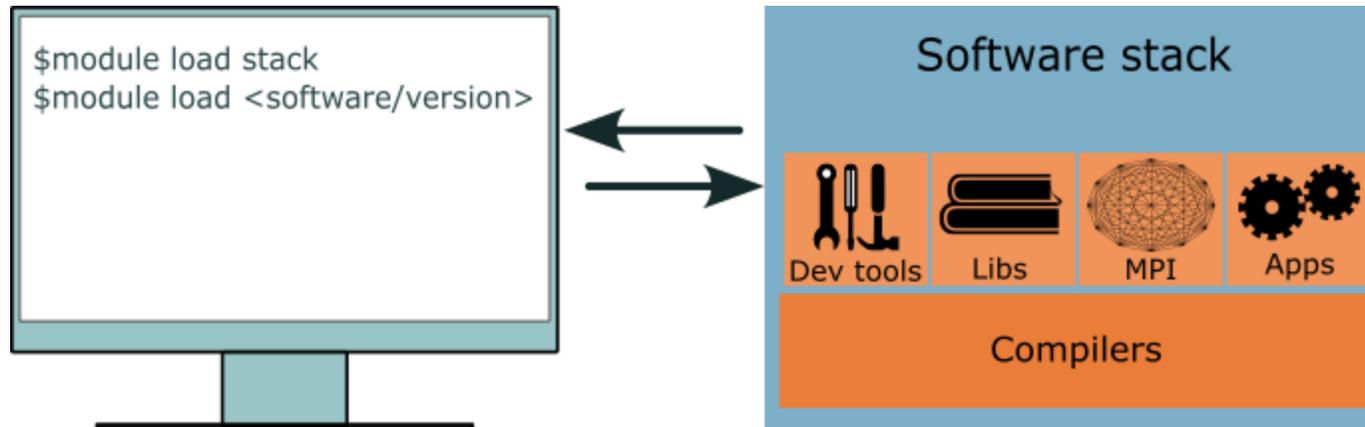


Outlook

- Accessing the cluster
- Storage and data transfer
- **Modules and applications**
- Using the batch system

Modules

Euler is shared among thousands of users. It needs to enable everyone to use the version of the software that they need, at the moment that they need it. The solution is to use the so-called **modules**.

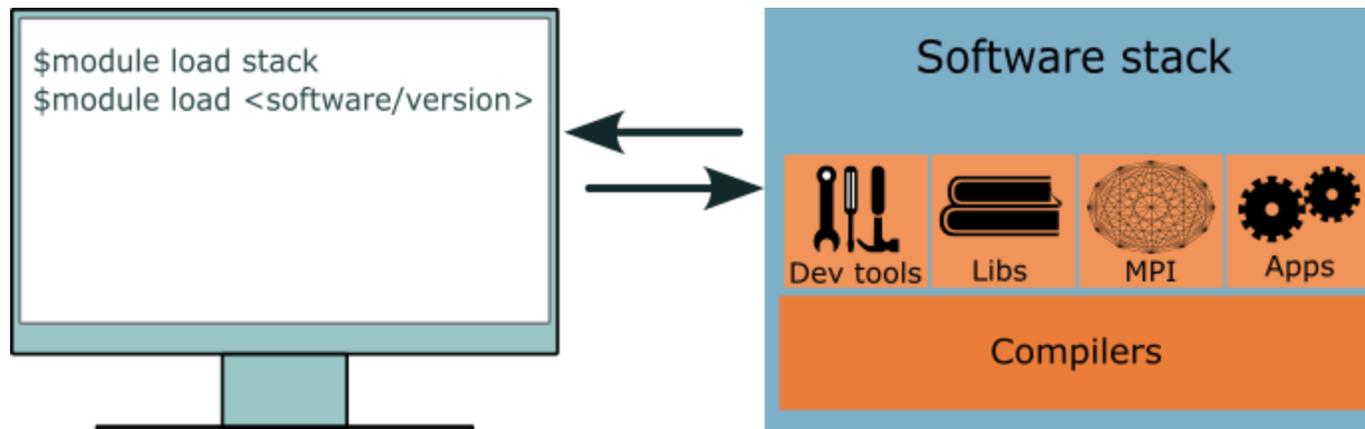


A module on Euler can include :

- Development tools
- Scientific libraries
- Communication libraries (MPI)
- Third-party applications

Modules

Euler is shared among thousands of users. It needs to enable everyone to use the version of the software that they need, at the moment that they need it. The solution is to use the so-called **modules**.



Advantages:

- Automatic configuration
- Different versions of the same software can co-exist and can be selected explicitly
- You can easily try out different tools, switch between versions, to find out which one works best for you

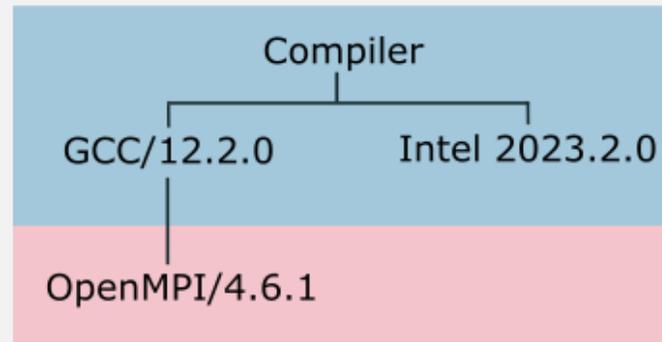
Modules: Hierarchy and Software Stacks

Core

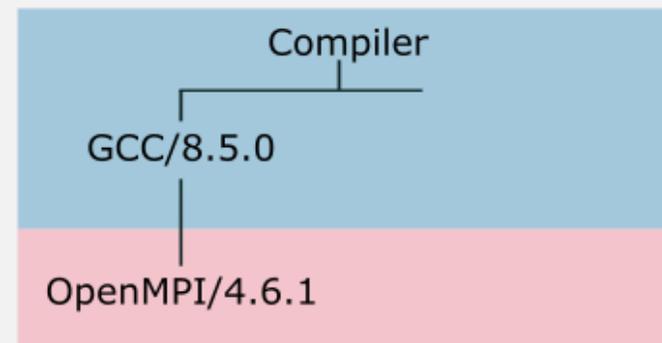
Commercial Software

Precompiled binaries

stack/2024-06



stack/2024-04



Module hierarchy with 3 layers:

- Core layer

```
$ module load comsol/6.2
```

- Compiler layer

```
$ module load stack/2024-06 gcc/12.2.0 python/3.11.6
```

- MPI layer

```
$ module load stack/2024-06 gcc/12.2.0 openmpi/4.6.1 openblas
```

Modules: General recommendations

- Use module spider to find all software of a given name on Euler:

```
nmarounina@eu-login-39:~$ module spider python
```

```
-----  
python:  
-----
```

```
  Versions:
```

```
    python/3.8.18-c3ikxoi  
    python/3.8.18-mcsq152  
    [...]  
    python/3.11.6_cuda-oe7bpyk  
    python/3.11.6_cuda  
    python/3.11.6_rocm-oe7bpyk  
    python/3.11.6_rocm  
    python/3.11.6-m4n2ny4  
    python/3.11.6-oe7bpyk  
    python/3.11.6
```

- From that list, avoid modules with hashes in their names (e.g. `python/3.11.6-oe7bpyk`)
- If given a choice, and if unsure about which version to pick, choose always the most recent one

Modules > Commands (demonstration)

Load Python module in GCC/6.3.0 toolchain

```
nmarounina@eu-login-39:~$ module load stack gcc/12.2.0 python/3.11.6
```

List available python module

```
nmarounina@eu-login-39:~$ module avail python
```

```
----- gcc/12.2.0 -----  
python/3.9.18      python/3.10.13      python/3.11.6 (L,D)      python_cuda/3.9.18  
python_cuda/3.11.6 (D)
```

List all currently loaded modules

```
nmarounina@eu-login-39:~$ module list
```

Currently Loaded Modules:

```
1) stack/2024-06  2) gcc/12.2.0  3) python/3.11.6
```

Modules > Commands

module

get info about module sub-commands

module spider

list all modules available on the cluster

module key *keyword*

list all modules whose description contains *keyword*

module help *name*

get information about module *name*

module show *name*

show what module *name* does (without loading it)

module unload *name*

unload module *name*

module purge

unload all modules at once

Modules > How to install applications locally?

Users can install additional applications in their home directory, but only if the quotas (space: 50 GB, files/directories: 500'000) are not exceeded

- Avoid anaconda installations as they often conflict with the files/directories quota. Alternatively, you can create a Python virtual environment.
- For Python and R, packages can easily be installed locally

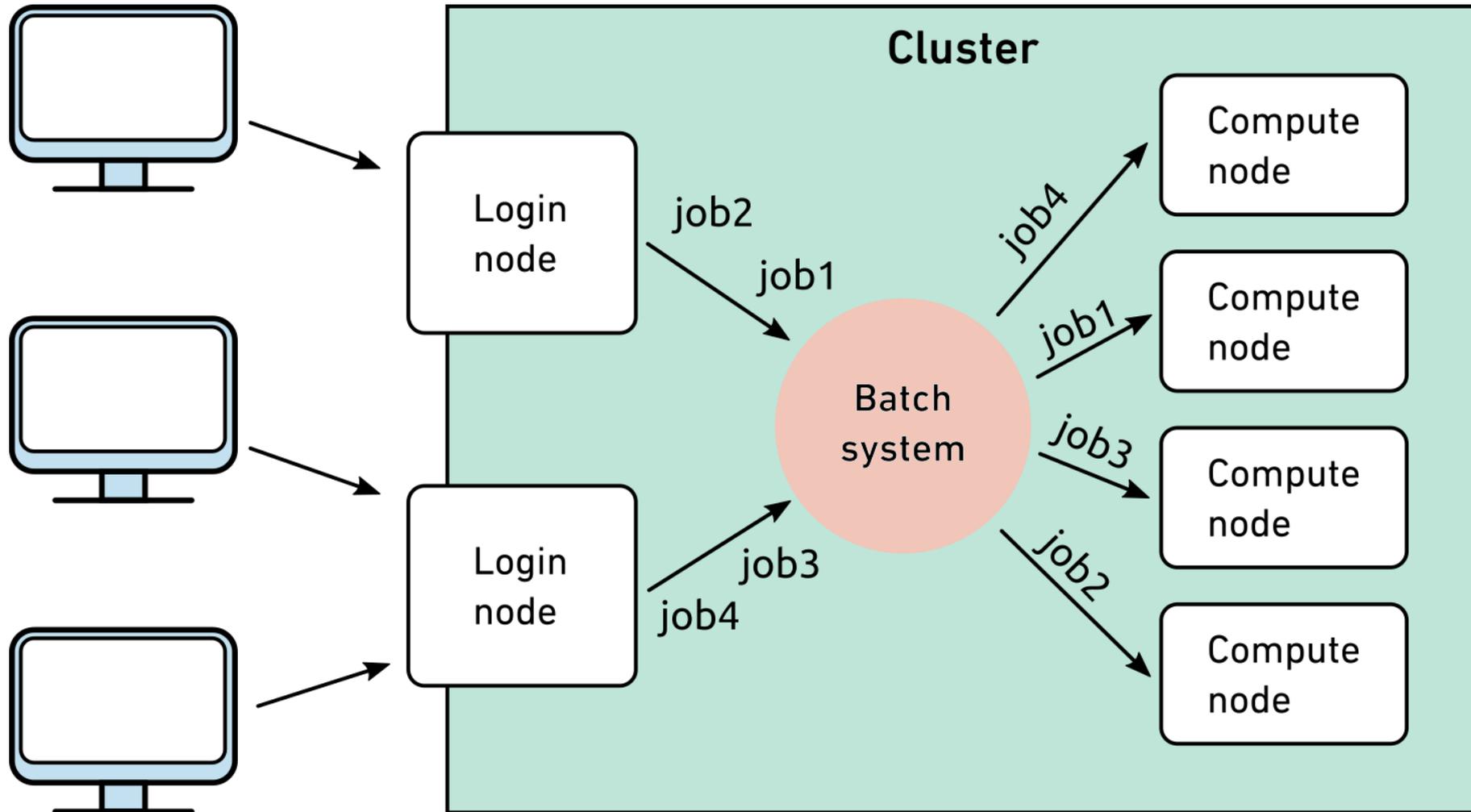
```
$ pip3 install --user packagename
```

- For python, consider creating virtual environments using venv, poema, ...

Outlook

- Accessing the cluster
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- Modules and applications
- Using the batch system

Batch > Overview



Batch > Overview

- The batch system of Euler is called **SLURM** (Simple Linux Utility for Resource Management)
- SLURM manages all resources available on the cluster and allocates them to user jobs
 - Ensures that resources are used as efficiently as possible
 - Calculates user/job priorities based on a fair share principle
- All computations must be submitted to the batch system
 - There is no other way to access the cluster's compute nodes
- Please do not run computations on the login nodes
 - Login nodes may only be used for file transfer, compilation, code testing and debugging, and quick pre- and post-processing

Batch > Basic job submission

- When you submit a job via `sbatch`, you specify the desired resources for it (in terms of how long the job should run, how many cores, GPUs, etc..)
- The batch system then analyzes the requested resources and dispatches the job to a batch queue. The job will run once the desired resources are available
- If all goes well, `sbatch` :
 - Tells the job's unique identifier ("job ID") – e.g. "1010171"
 - Will automatically create an output stderr/stdout file `slurm-jobID.out`
- The jobID is important to check, monitor or terminate a job
- If you report a problem with a job (pending, running or done) to cluster support, then **always** provide the corresponding jobID and the `slurm-jobID.out` file

Batch > Basic job submission

- Use `sbatch` to submit a job to the batch system
- `sbatch [Slurm options] --wrap="job"`
- A *job* can be either ...
 - a single Linux command
 - a shell script, passed via “<”
 - a program, with its path
 - a command or program, with its arguments
 - multiple commands, enclosed in quotes
- We'll talk about `sbatch`'s options later

```
pwd  
< script  
/path/to/program  
cmd arg1 arg2  
"cmd1 ; cmd2"
```

Batch > Basic job submission > Examples

```
[sfux@eu-login-03 ~]$ sbatch --wrap="echo hello"  
Submitted batch job 1010112
```

```
[sfux@eu-login-03 ~]$ sbatch < hello.sh  
Submitted batch job 1010113
```

```
[sfux@eu-login-03 ~]$ sbatch --wrap="./bin/hello"  
Submitted batch job 1010114
```

```
[sfux@eu-login-03 ~]$ sbatch --wrap="date; pwd; ls -l"  
Submitted batch job 1010115
```

```
[sfux@eu-login-03 ~]$ sbatch --wrap="du -sk /scratch > du.out"  
Submitted batch job 1010116
```

Batch > Resource requirements

- By default, a job will get 1 core for 1 hour with 1000MB of memory
 - If you need more time and/or processors and/or memory, you must request them
 - Maximum run-time on Euler is 15 days
- These resources are passed to `sbatch` using options :

```
sbatch --time=HH:MM:SS --ntasks=number_of_processors  
--mem-per-cpu=2000 --wrap="command"
```

- If you don't specify any unit for the memory request, then the integer value will be interpreted as MB. If you specify values in GB, then you need to add the suffix "g" (in the example above, you would write 2g instead of 2000)

Batch > sbatch options

<code>--ntasks=N</code>	request N cores (<code>--nodes=1</code> allocates all cores on a single node)
<code>--time=HH:MM:SS</code>	request a runtime of $HH:MM:SS$
<code>--output="filename"</code>	redirect job's standard output to <i>filename</i>
<code>--error="filename"</code>	redirect job's error messages to <i>filename</i>
<code>--mem-per-cpu=YYY</code>	request YYY MB memory per core (or add suffix "g" to specify GBs)
<code>--tmp=YYY</code>	request YYY MB of local scratch space (or add suffix "g" to specify GBs)
<code>--job-name="jobname"</code>	assign a <i>jobname</i> to the job
<code>--account="share"</code>	run job under a particular Euler share " <i>share</i> "
<code>--mail-type=BEGIN</code>	send an email when the job begins
<code>--mail-type=END,FAIL</code>	send an email when the job ends (finishes successfully or fails)

Batch > sbatch GPU options

- `--gpus=N` request N gpus
- `--gpus=MODEL:N` request N gpus of model *MODEL* (for instance `--gpus=rxt_3090:1`)
- `--gres=gpumem:XXg` request a GPU with at least XX GB GPU memory

https://scicomp.ethz.ch/wiki/GPU_job_submission_with_SLURM

Batch > #SBATCH pragmas

- `sbatch` options can be specified either on the command line or inside a job script using the `#SBATCH` pragma, for example

```
#!/bin/bash
#SBATCH --ntasks=24           # 24 cores
#SBATCH --time=8:00:00       # 8-hour run-time
#SBATCH --mem-per-cpu=4000    # 4000 MB per core
cd /path/to/execution/folder
module load stack/2024-06 openmpi/4.1.6
srun myprogram arg1
```

- In this case, the script can be submitted using the “<” operator

```
$ sbatch < script
```

- `sbatch` options specified on the command line override those inside the script

```
$ sbatch --ntasks=48 < script
```

Batch > Job monitoring

squeue

check the state of a job in the queue

myjobs

detailed information about a job

scancel

kill a job

Advanced commands :

scontrol

check resource usage of a job

sstat

check information about a running job

sacct

detailed information about pending, running and finished jobs

Batch > Job monitoring > `squeue`

- After submitting a job, the job will wait in a queue to be run on a compute node and has the pending status (PD). You can check the job status with the `squeue` command

```
[sfux@eu-login-41 ~]$ squeue
```

JOBID	PARTITION	NAME	USER	ST	TIME	NODES	NODELIST (REASON)
1433323	normal.4h	wrap	sfux	PD	0:00	1	eu-g1-026-2
1433322	normal.4h	wrap	sfux	R	0:11	1	eu-a2p-483

Batch > Job monitoring > myjobs

- Detailed information about a job can be provided by myjobs command:

```
$ myjobs -j 6038307
Job information
  Job ID                : 6038307
  Status                : RUNNING
  Running on node       : eu-g3-022
  User                  : nmarounina
  Shareholder group     : es_cdss
  Slurm partition (queue) : gpu.24h
  Command               : script.sbatch
  Working directory     : /cluster/home/nmarounina
Requested resources
  Requested runtime     : 08:00:00
  Requested cores (total) : 12
  Requested nodes       : 1
  Requested memory (total) : 120000 MiB
Job history
  Submitted at          : 2023-01-09T15:56:09
  Started at            : 2023-01-09T15:56:38
  Queue waiting time    : 29 s
Resource usage
  Wall-clock            : 00:00:36
  Total CPU time        : 00:00:00
  CPU utilization       : 0%
  Total resident memory : 2.94 MiB
  Resident memory utilization : 0%
```

Batch > Job monitoring > scancel

```
[sfux@eu-login-15 ~]$ squeue
      JOBID PARTITION      NAME      USER ST      TIME  NODES NODELIST (REASON)
      1525589 normal.24  sbatch    sfux  R      0:11      1 eu-a2p-373
[sfux@eu-login-15 ~]$ scancel 1525589
[sfux@eu-login-15 ~]$ squeue
      JOBID PARTITION      NAME      USER ST      TIME  NODES NODELIST (REASON)
[sfux@eu-login-15 ~]$
```

Options:

job-ID

kill *job-ID*

--name=*jobname*

kill all jobs with name *jobname*

--user=*username*

kill all jobs from user *username*

--state=*state*

kill all jobs in state *state* (states: PENDING, RUNNING or SUSPENDED)

Dos and don'ts

Dos

- Understand what you are doing, i.e. make the effort to inform yourself
- Ask for help if you don't understand what you are doing or you were not able to find the relevant information
- Optimize your workflow to make it as efficient as possible
- Keep in mind that our clusters are shared by many users
- Choose the file system you want to use carefully

Don'ts

- Don't waste CPU time or disk space
- **Don't run applications on the login nodes**
- Don't write large amounts of data to standard output
- Don't create millions of small files
- Don't run hundreds of small jobs if the same work can be done in a single job

Getting help

- Wiki: <https://scicomp.ethz.ch>
- Ticket system
 - <https://smartdesk.ethz.ch> (ETH account authentication)
 - Please describe your problem as accurately as possible
- E-mail
 - cluster-support@id.ethz.ch
 - Please do not send questions to individual members of the team
- Person-to-person
 - In-person Helpdesk every other Wednesday
 - Contact us to set up an appointment at your place
 - Visit us at Binzmühlestrasse 130

Demonstration of our services:

- <https://jupyter.euler.hpc.ethz.ch/> - JupyterHub
- <https://slurm-jobs-webgui.euler.hpc.ethz.ch> – SLURM webGUI
- <https://scicomp.ethz.ch/public/lsla/index2.html> - SLURM command/script generator
- https://scicomp.ethz.ch/wiki/Suggestion_Box - Cluster Suggestion Box

Questions?