



Vlaanderen
is supercomputing

VSC HPC introduction course

<https://hpcleuven.github.io/HPC-intro>

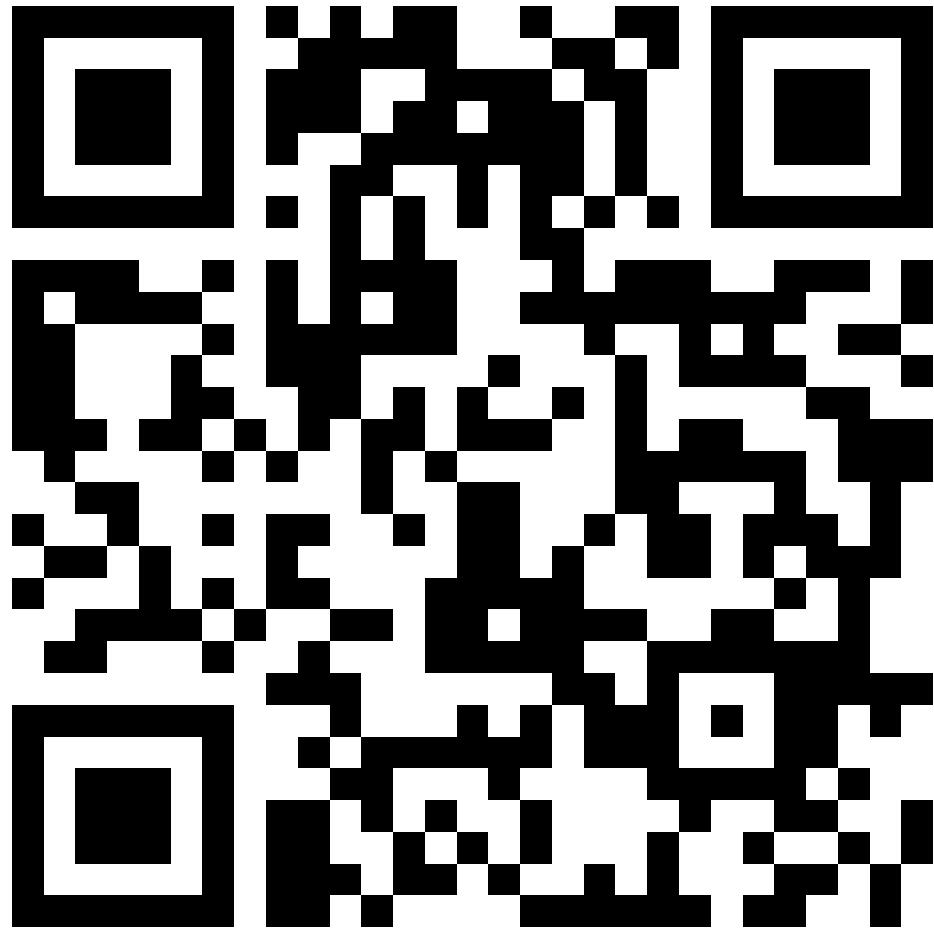


Training material

- ❑ Everything is available on Github:
<https://hpcleuven.github.io/HPC-intro/>

- ❑ Video recordings
 - Scan the QR code
 - Recommended videos: ~2 hrs
 - Optional videos: ~1 hr

By registering and participating in VSC trainings, you accept to abide by the
[VSC training Code of Conduct](#)



What is High Performance Computing?

- ❑ Using supercomputers to solve advanced computational problems
- ❑ Reduce the computation time from days, years, decades or centuries to minutes, hours, days or weeks
- ❑ The key is parallelism
- ❑ Access to specialized hardware
(GPU, large memory, high-speed interconnect)



What is High Performance Computing?



The concept is simple: **parallelism** = employing multiple processors for a single problem

Training outline

Part 1: Basics

- VSC
- Tier-2 clusters
- Storage
- Accounting
- Login

Open OnDemand

SSH client & terminal

NX

- Data transfer

FileZilla

rsync

Globus

Part 2: Hands-on

- Using Open OnDemand
- Software
- Miniconda
- Batch jobs
- Demo: try things yourself



VSC
(Vlaams Supercomputer Centrum)

VSC PARTNERSHIP



Supported by



Visit <https://vscentrum.be>

Extensive set of courses
every academic year

Links to the VSC **account page** and the VSC **documentation**
(answers >80% of your questions!)

The screenshot shows the homepage of the VSC website. At the top left is the VSC logo with the text "VLAAMS SUPERCOMPUTER CENTRUM" and "Vlaanderen is supercomputing". A navigation bar below the logo includes links for Home, About VSC, Systems & Services, Showcase, News & Events, VSC Training, User Portal, and Access. To the right of the navigation is a search bar with a magnifying glass icon. The main content area features a large, bold, black text block that reads: "Welcome to VSC Flanders' most highly integrated high-performance research computing environment". Below this text is a "Read More" button. The background of the main content area has a dark, abstract design with glowing red and white vertical streaks.

VLAAMS
SUPERCOMPUTER
CENTRUM

Vlaanderen
is supercomputing

Home About VSC Systems & Services Showcase News & Events VSC Training User Portal Access

ch...

Welcome to VSC

Flanders' most highly integrated

high-performance research

computing environment

[Read More](#)



Welcome to the User Portal

Here you can find the gateway to the User documentation of the Vlaams Supercomputer Centrum

User documentation page

Manage your VSC-account
partner institute account required

VSC account page

<https://docs.vscenrtum.be>

Welcome to the VSC documentation

The VSC documentation offers extensive *how-to* guides and technical information about the services provided by the [Vlaams Supercomputer Centrum](#).

Accounts and access
How to get your VSC account and access the different VSC services and platforms.

Research Data
Data transfer and storage in the VSC infrastructure.

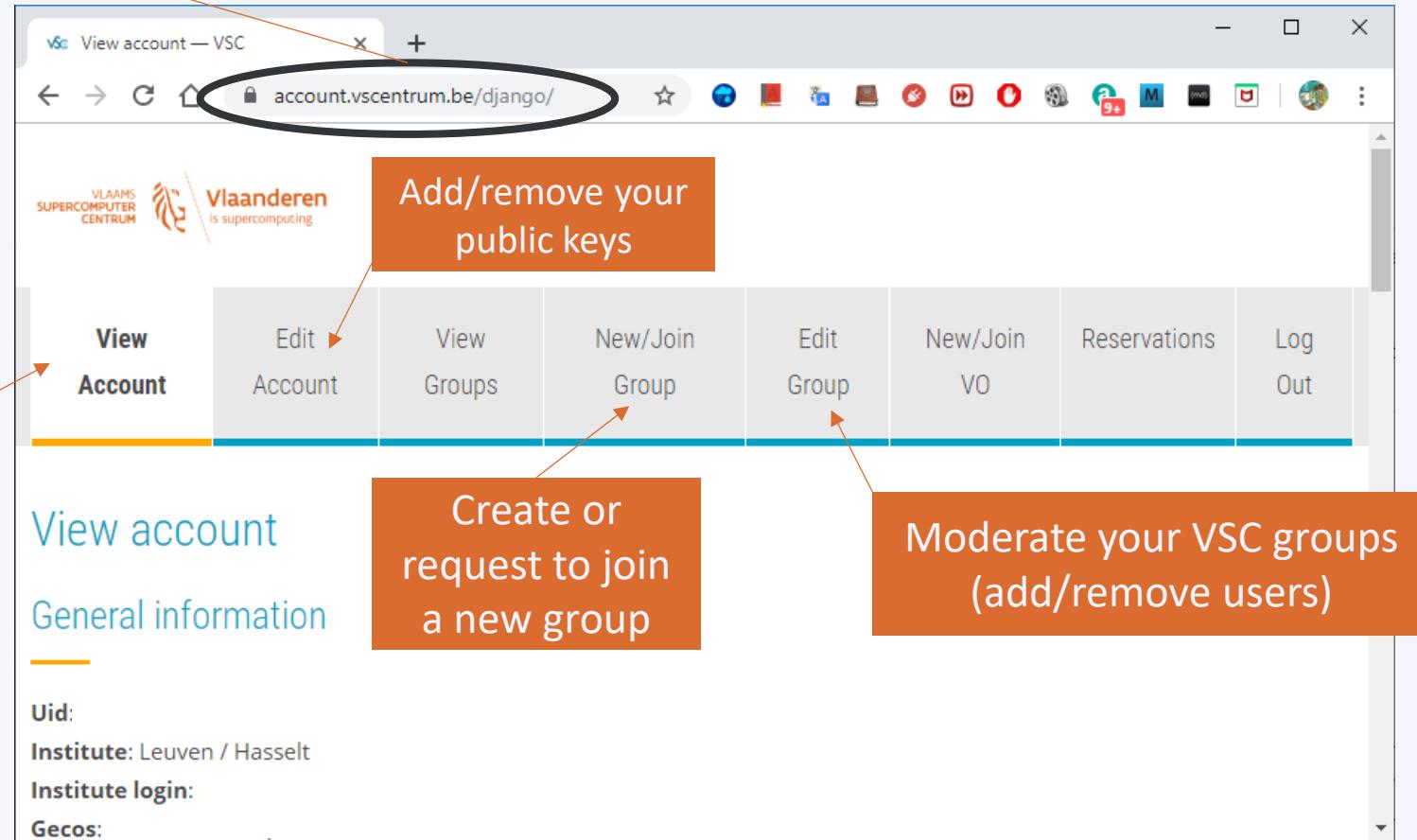
Compute
The high-performance computing (HPC) platform provides multiple tiers of parallel processing

Tier-1 Cloud
The VSC Cloud component provides *on-demand* resources in a more flexible and cloud-like

Tier-1 Data
The VSC Data component enables research data to remain close to the computing infrastructure

Enter your keywords here; e.g. SSH key

Manage your VSC account at
<https://account.vscentrum.be>



Support and services

Basic support

- Helpdesk (hpcinfo@kuleuven.be)
- Monitoring and reporting

Application support

- Installation and porting
- Optimisation and debugging
- Benchmarking
- Workflows and best practices

Training

- Documentation and tutorials
- Scheduled trainings and workshops
- Workshops on request
- One-to-one sessions

Become a VSC user

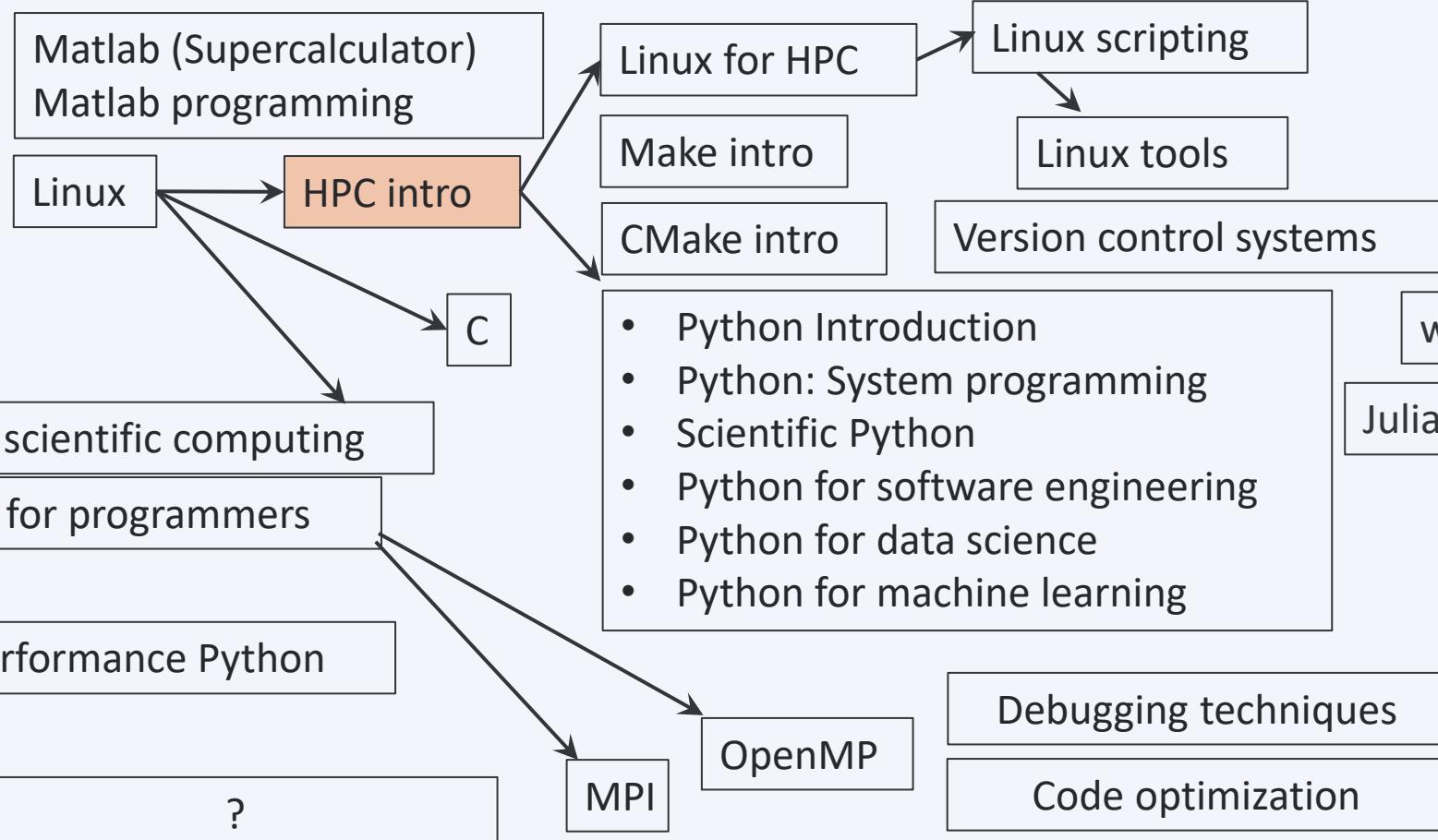
- (Optional) Create a secure (4096 bit) [SSH key pairs](#)
Upload it on the account page: <https://account.vscentrum.be>
- You need to [request a VSC account](#)
Normally gets processed swiftly
- Request [introductory credits](#) (2M free credits for 6 months)
- Request [project credits](#) (for supervisors and project leaders)
You need to create a VSC group
Add users to the group to give them access to use credits
Fill out the request form
- Extra storage requests
Scratch extension: free of charge
Staging fileset: 20 € per TB per year
- All service costs (compute and storage) are described in the ICTS service catalog: <https://icts.kuleuven.be/sc>
Click on [High Performance Computing](#) (NL/EN)

The screenshot shows a web browser window titled 'View account — VSC'. The URL is 'account.vscentrum.be/django...'. The page header includes the Vlaams Supercomputer Centrum logo and the text 'Vlaanderen is supercomputing'. A navigation menu at the top has 'View Account' selected. Below the menu, there are sections for 'View account' and 'General information'.

The screenshot shows a web browser window titled 'ICTS Servicecatalogus'. The URL is 'icts.kuleuven.be/sc'. The page header includes the KU Leuven logo and the text 'ICTS SERVICECATALOGUS'. A navigation menu is visible. The main content area features a section titled 'WAT IS DE ICTS SERVICECATALOGUS?' and another titled 'OPGELET' with a note about netto prijzen. On the right side, there is a graphic of a computer mouse and a search bar labeled 'ZOEKEN'.

VSC training

- Introductory



- Intermediate

- Advanced

- Specialized track

Infosessions:

- Containers
- Notebooks

How to acknowledge VSC in publications

Why?

- a contractual obligation for the VSC
- helps VSC to secure funding
- you will benefit from it in the long run

For KU Leuven users

- add the relevant papers to the “KUL-HPC” virtual collection in Lirias

In het Nederlands

De infrastructuur en dienstverlening gebruikt in dit werk werd voorzien door het VSC (Vlaams Supercomputer Centrum), gefinancierd door het FWO en de Vlaamse overheid.

In English

The resources and services used in this work were provided by the VSC (Flemish Supercomputer Center), funded by the Research Foundation - Flanders (FWO) and the Flemish Government.



Tier-2 clusters

VSC HPC infrastructure



Tier-2 clusters @ KU Leuven

Genius (since 2018)

250 nodes | 8936 cores



Most of Genius will be decommissioned in early 2026

wICE (since September 2022)

186 nodes | 13392 cores



Tier-2 cluster – Genius

Node type	CPU type	# cores	Total mem	Mem per core (MB)	Partition	Planned end of life
Cascadelake	Xeon Gold 6240	36	192 GB	5000	batch	Q1 2026
Skylake large mem	Xeon Gold 6240	36	768 GB	21000	bigmem	Q1 2026
Skylake P100 GPU	Xeon Gold 6140 4xP100 SXM2 16GB	36	192 GB	5000	gpu_p100	2027 (?)
Cascadelake V100 GPU	Xeon Gold 6240 8xV100 SXM2 32GB	36	768 GB	21000	gpu_v100	2027 (?)
Superdome	Xeon Gold 6132 8 sockets	112	6 TB	53500	superdome	Q1 2026
Skylake interactive	Xeon Gold 6140	36	192 GB	5000	interactive	Q1 2026

- **Access:** login.hpc.kuleuven.be
- **Interconnect:** InfiniBand EDR (25 Gb/s)
- **SSD Disks:** 200 GB

- **Interactive partition:**
max resources: 8 cores, 16hr

Remarks

Tier-2 cluster – wlCE

Node type	CPU type	# cores	Total mem	Mem per core (MB)	Partition
Icelake	Xeon 8360Y	72	256 GB	3455	batch_icelake
Sapphire Rapids	Xeon 8468	96	256 GB	2500	batch_sapphirerapids
Icelake large mem	Xeon 8360Y	72	2 TB	28000	bigmem
Icelake large mem	Xeon 8360Y	72	8 TB	111900	hugemem
Icelake A100 GPU	Xeon 8360Y 4xA100 SXM2 80GB	72	512 GB	7000	gpu_a100
Genoa H100 GPU	Epyc 9334 4xH100 SXM5 80GB	64	768 GB	11700	gpu_h100
Icelake interactive	Xeon 8358 1xA100 SXM2 80GB	64	512 GB	7500	interactive

- **Phase 1:** since September 2022
- **Access:** login.hpc.kuleuven.be
- **Interconnect:** InfiniBand HDR (100 Gb/s)

- **SSD Disks:** 960 GB
- **Interactive partition:**
max resources: 8 cores, 1 GPU, 16hr

Remarks

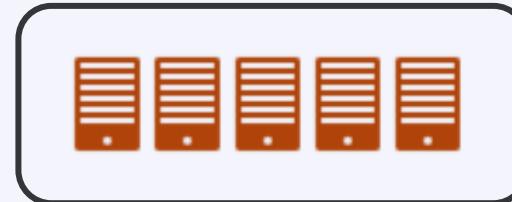
Tier-2 cluster – wlCE

Thin nodes



172x IceLake 72c 256 GB

Large memory nodes



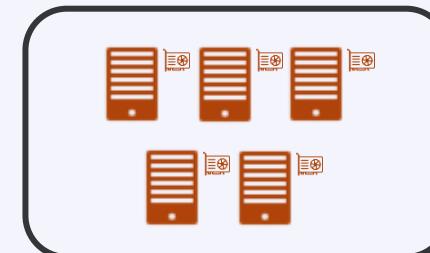
5x IceLake 72c 2048 GB

GPU nodes



4x IceLake 72c 512 GB
4 A100 SXM4 80GB

Interactive nodes

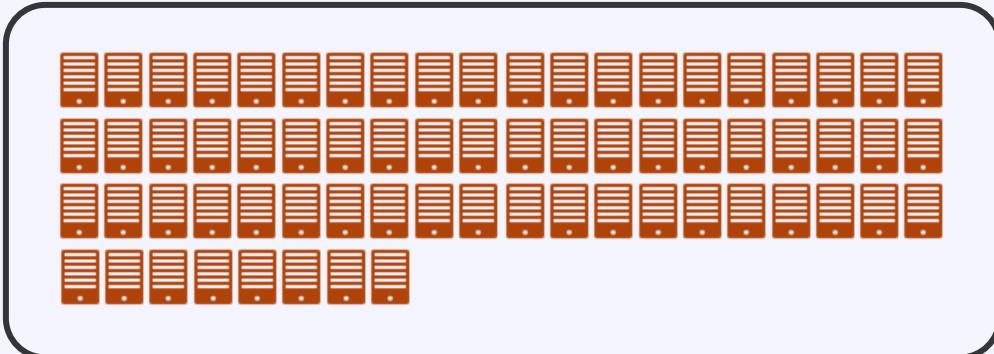


4x IceLake 64c 512 GB
1 A100 80GB

No Dedicated
Login Nodes

Tier-2 cluster – wlCE extension

Thin nodes



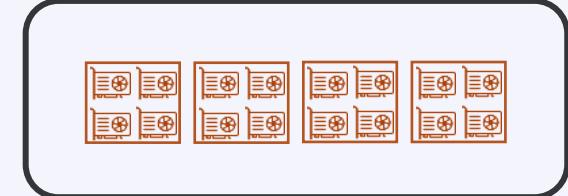
68x Sapphire Rapids 96c 256 GB

Large memory nodes



1x IceLake 72c 8 TB

GPU nodes



4x AMD Genoa 64c 768 GB
4x H100 SXM5 80GB

Storage

Overview of the Tier-2 storage locations

- ✓ Only you own your files (POSIX)
Users can share folders via [VSC groups](#)
- ✓ A VSC account has 3 default storages (free of charge)
 - \$VSC_HOME
 - \$VSC_DATA
 - \$VSC_SCRATCH
- ✓ You can additionally request staging storage
- ✓ Different storage volumes have different:
 - mount point
 - size and performance
 - use case
 - backup and maintenance policy
- ✓ For more info, see the [ICTS Service Catalog](#) (EN/NL)

Storage locations

Example

- ❑ **Avoid using /tmp (\$TMPDIR) on the login nodes**

It is only 10 GB and is needed by the operating system
Your application can crash if using /tmp

- ❑ You are automatically logged into your home folder upon login. Best to immediately switch to some other storage, e.g.

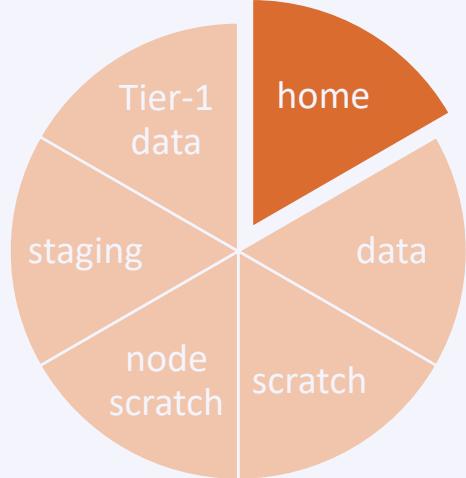
```
$ cd $VSC_DATA
```

- ❑ Always check your storage balance using myquota command

```
$ myquota
file system $VSC_HOME
    Blocks: 1479M of 3072M
    Files: 12934 of 100k
file system $VSC_DATA
    Blocks: 12G of 75G
    Files: 1043k of 10000k
file system $VSC_SCRATCH
    Blocks: 15M of 1.5T
```

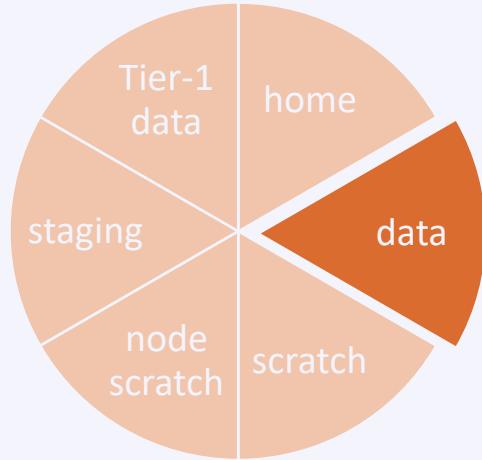
- ✓ [Request form for extra storage](#)
- ✓ [More info in the service catalog](#)

Storage locations



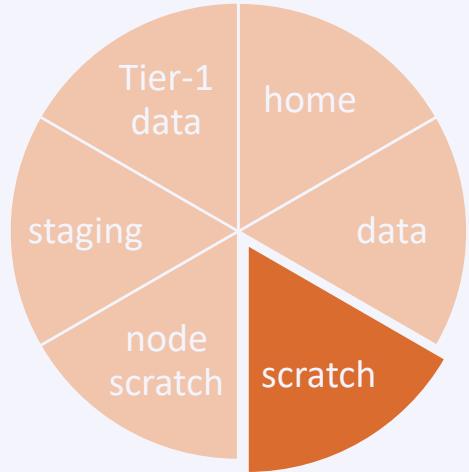
Storage	Home folder
Env. variable	\$VSC_HOME
Filesystem type	NFS
Access	Global
Backup	Hourly, daily, weekly (until last month) inside .snapshot folders
Default quotum	3 GB
Quotum increase	Not possible
Usage	Only storing SSH keys, config files, ...
Remarks	<ul style="list-style-type: none">- Stay away from using it- Can easily overflow:<ul style="list-style-type: none">+ Your jobs may crash+ Login issues

Storage locations



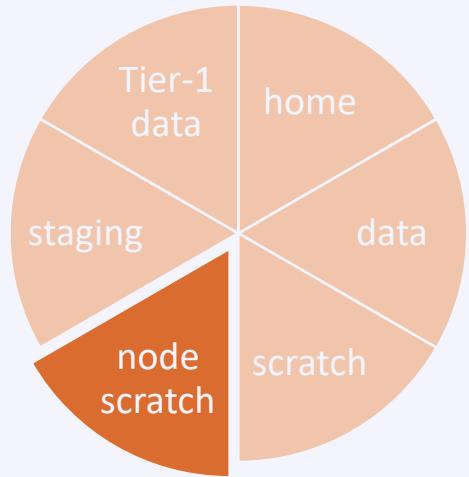
Storage	Data folder
Env. variable	\$VSC_DATA
Filesystem type	NFS
Access	Global
Backup	Hourly, daily, weekly (until last month) inside .snapshot folders
Default quotum	75 GB
Quotum increase	On purchase
Usage	Your data, code, software, results
Remarks	<ul style="list-style-type: none">- Permanent storage for initial/final results- Not optimal for intensive or parallel I/O

Storage locations



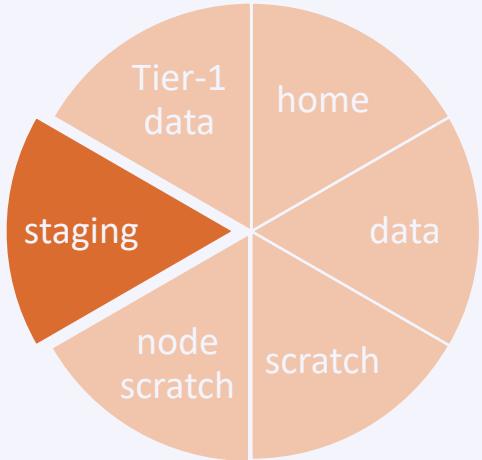
Storage	Scratch folder
Env. variable	\$VSC_SCRATCH
Filesystem type	Lustre
Access	Local
Backup	Files get deleted 30 days after last access
Default quotum	500 GB
Quotum increase	Free
Usage	Intensive, parallel I/O, temporary files
Remarks	<ul style="list-style-type: none">- Recommended storage for all jobs- Copy scratch files to VSC_DATA or local storage after jobs are done- Deleted files cannot be recovered- Avoid folders with >10,000 files

Storage locations



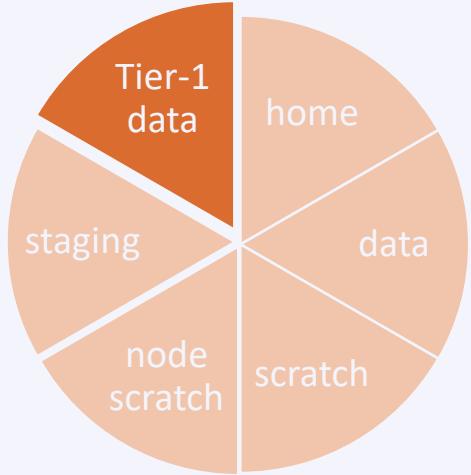
Storage	Node scratch folder
Env. variable	\$VSC_SCRATCH_NODE
Filesystem type	Ext4
Access	On compute node, only at runtime
Backup	None
Default quotum	591 GB
Extension	Read about beeOND
Usage	Temporary storage at runtime
Remarks	<ul style="list-style-type: none">- Fastest I/O, attached to the node- Is cleaned after job terminates- Copy the data to your home, scratch, or staging before job ends

Storage locations



Storage	Staging folder
Path	/staging/leuven/stg_000xx
Filesystem type	Lustre
Access	On demand, only Tier-2@KUL
Backup	None
Default quotum	None
Extension	On purchase, from 1 TB
Usage	Permanent; share with a group
Remarks	<ul style="list-style-type: none">- Accessible from login/compute nodes- Fast, parallel I/O

Storage locations



- VSC offers [Tier-1 Data](#) as a large storage service
- Only for active data (no archiving)
- Supports metadata (search & discover)
- Share within VSC (permission management)
- Share outside VSC (via [Globus](#))
- Stage-in/stage-out data workflow
- Free of charge for academic use
- Starting volume is 1 PB
- Submit a proposal to get access
You can apply at any time

Extra (KU Leuven) storage locations



Dataset



Archived package

- In production
- For **active** data
- Data should stage in/out
- Has backup
- See the [ICTS service catalog](#)

- In development
- For **inactive** data
- Storage until 10 years
- Has backup
- See the [ICTS service catalog](#)

Also see the [KU Leuven storage guide](#) for even more storage possibilities

ManGO





Compute credits



Tier-2 credit system

What?

- Credits are a measure for compute resources
- Mandatory for accessing these resources
- Traceability and accountability

Why?

- Incentive to optimize your code and workflow

How?

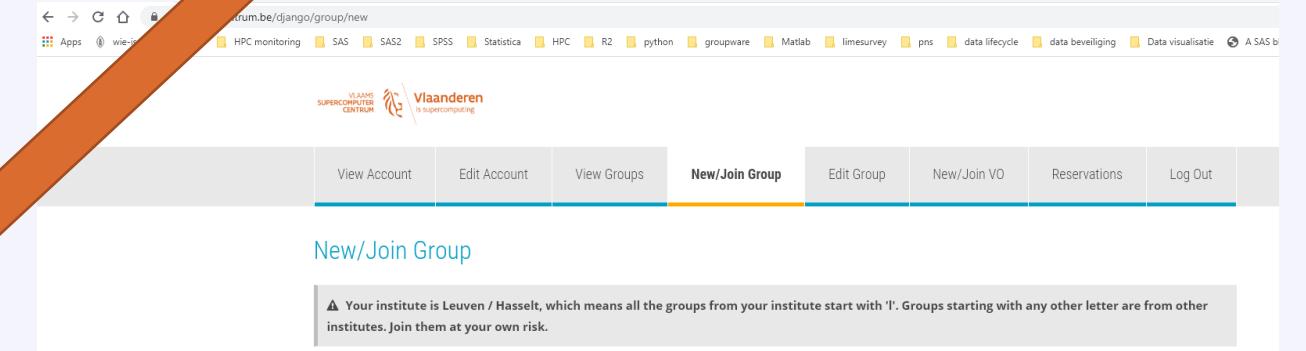
- 1 million credits = 3.5 EUR
- Pay-as-you-go model
- Credit rates on the right = for fully allocated nodes
(if you request less cores, you also get billed less)
- See the [ICTS service catalog](#) for more info

Node types	Credits per node-hour
Genius Skylake / Cascadelake	10 000
Genius Skylake Bigmem	12 000
Genius Skylake 4 GPUs	20 000
Genius Skylake interactive node	Free
Genius Cascadelake 8 GPUs	39 900
Genius Superdome	126 000
wICE IceLake thin node	11 000
wICE IceLake Large Memory	19 000
wICE IceLake GPU node	45 000
wICE IceLake interactive node	Free
wICE IceLake hugemem node	80 000
wICE Sapphire Rapids thin node	20 000
wICE AMD Zen-4 4 GPUs	150 000

Tier-2 credit system – how to request credits

- Request your introduction credits
 - Every new user can request them once
 - Free of charge
 - 6 months
- Request permission to join a group
- Request project credits
 - Create group on the VSC account page
 - Start the name preferably with **p_**
 - Your group will start with **lp_**
 - Creator of the group can add and remove users
- Tier-2 credits can be used on all (Leuven) Tier-2 clusters

Check the [ICTS service catalog](#)



A screenshot of a web browser showing the 'Create new group' form. The URL is 'account.vscentrum.be/django/group/new'. The page has a header with the Vlaams Supercomputer Centrum logo and the text 'Vlaanderen is supercomputing'. Below the header is a navigation bar with tabs: View Account, Edit Account, View Groups, New/Join Group (which is highlighted in blue), Edit Group, and New/Join VO. A red circle highlights the 'New/Join Group' tab. The main content area has a heading 'Create new group' and a note: '⚠ We will automatically prepend the letter 'l' to your groupname.' Below this is a form field labeled 'Groupname *' with a red asterisk. An orange arrow points from the 'Request project credits' section towards this 'Create new group' form.

Tier-2 credit system – how

- Submit your jobs with a suitable credit account

```
#SBATCH -A lp_myproject
```

```
sbatch -A lp_myproject ... myjob.slurm
```

- Before submitting the job, Slurm checks if the account has enough available credits for the job (in addition to any other already-running jobs)
- Once a job start running, it starts consuming credits from the account

Tier-2 credit system – manage your credits

- We provide various *sam*-commands to facilitate credit related queries

Command	Purpose
sam-balance [-A account]	List active projects and available credits


```
tier2-p-login-3$ sam-balance -A lp_hpcinfo
ID          Name           Balance      Reserved      Available
===== ====== ====== ====== ======
96141      lp_hpcinfo    9782636        0            9782636
tier2-p-login-3$ 
```

Tier-2 credit system – manage your credits

- We provide various *sam*-commands to facilitate credit related queries

Command	Purpose
sam-list-allocations -A <account-name>	List the credit deposits and refunds (if any)


```
tier2-p-login-3$ sam-list-allocations -A lp_hpcinfo
AllocID    AccountID Account          Timestamp           Credits
=====  ======  ======  ======  ======
80745      96141    lp_hpcinfo        2023-02-14T10:40:50  2000000
203565     96141    lp_hpcinfo        2023-04-17T16:42:26  2000000
227730     96141    lp_hpcinfo        2023-05-03T09:10:29  2000000
313474     96141    lp_hpcinfo        2023-06-19T17:33:59  8410000
tier2-p-login-3$ 
```

Tier-2 credit system – manage your credits

- We provide various *sam*-commands to facilitate credit related queries

Command	Purpose
sam-quote sbatch [sbatch args] [myjob.slurm]	Estimates the (max) amount of credits a job will use

```
tier2-p-login-3$ sam-quote sbatch --cluster=wice hello.slurm  
10980  
tier2-p-login-3$ █
```

Tier-2 credit system – manage your credits

- We provide various *sam*-commands to facilitate credit related queries

Command	Purpose
sam-statement -A <account-name> -s <start-date> -e <end-date>	Overview of credits used by each job in some time window

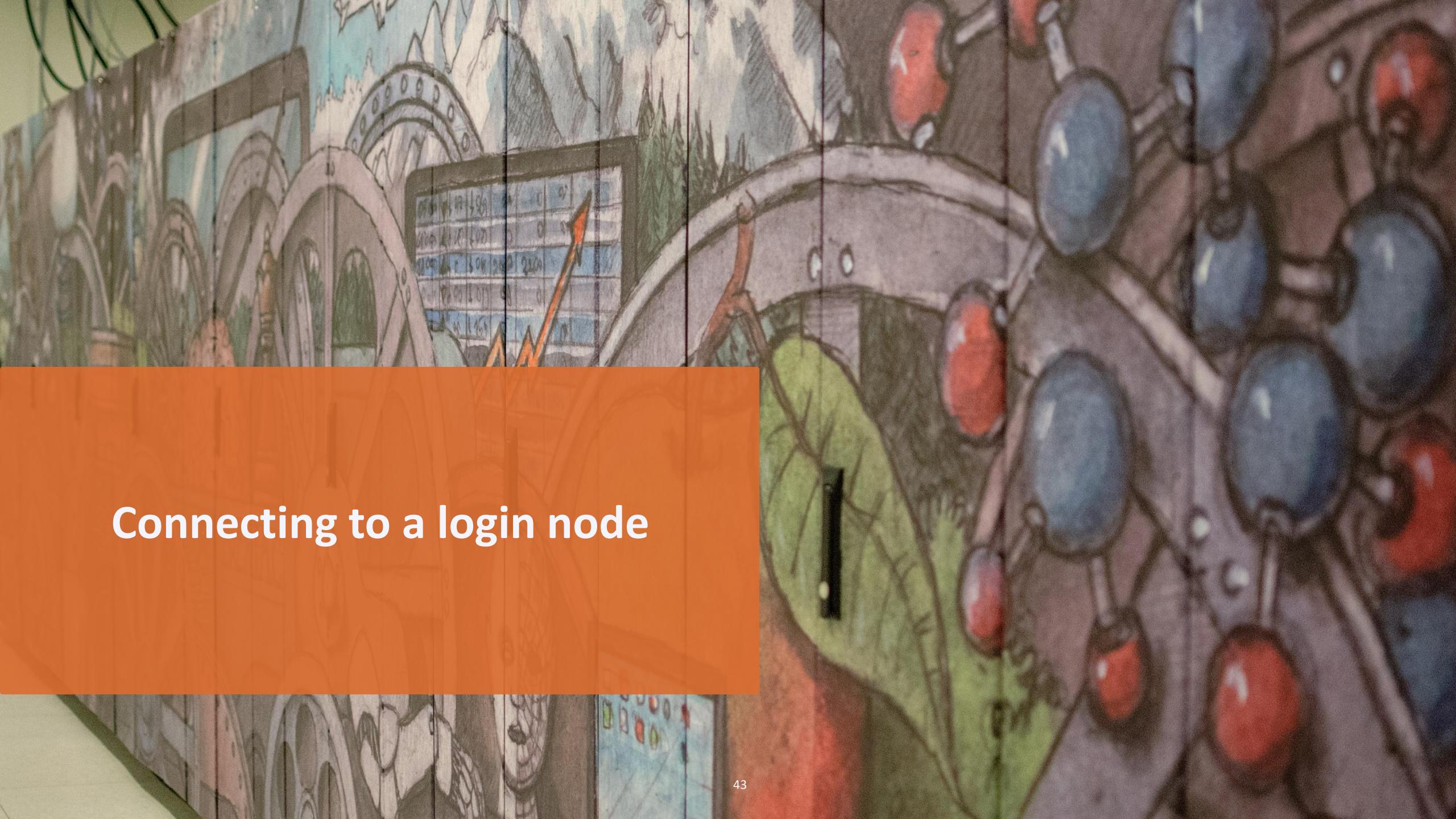
```
tier2-p-login-3$ sam-statement -A lp_hpcinfo -s 2023-09-01 -e 2023-10-25
#####
# Includes Account=lp_hpcinfo
# Generated on Mon Oct 30 11:01:29 2023.
# Reporting fund activity from 2023-09-01 to 2023-10-25
#
#####

Credits deposited in the given period: 0
Credits refunded in the given period: 0
Credits consumed in the given period: 0
-----
JobID      Cluster  Account          User      Partition  Credits
=====  ======  ======  ======  ======  ======
60739746   wice    lp_hpcinfo      vsc30446 batch    0
60740123   wice    lp_hpcinfo      vsc30446 batch    0
60740125   wice    lp_hpcinfo      vsc30446 batch    0
60746730   wice    lp_hpcinfo      vsc30446 interactive 0
60773547   wice    lp_hpcinfo      vsc30446 interactive 0
60774141   wice    lp_hpcinfo      vsc30446 batch    0
60774160   wice    lp_hpcinfo      vsc30446 batch    0
60783857   wice    lp_hpcinfo      vsc30446 interactive 0
```

Tier-2 credit system – manage your credits

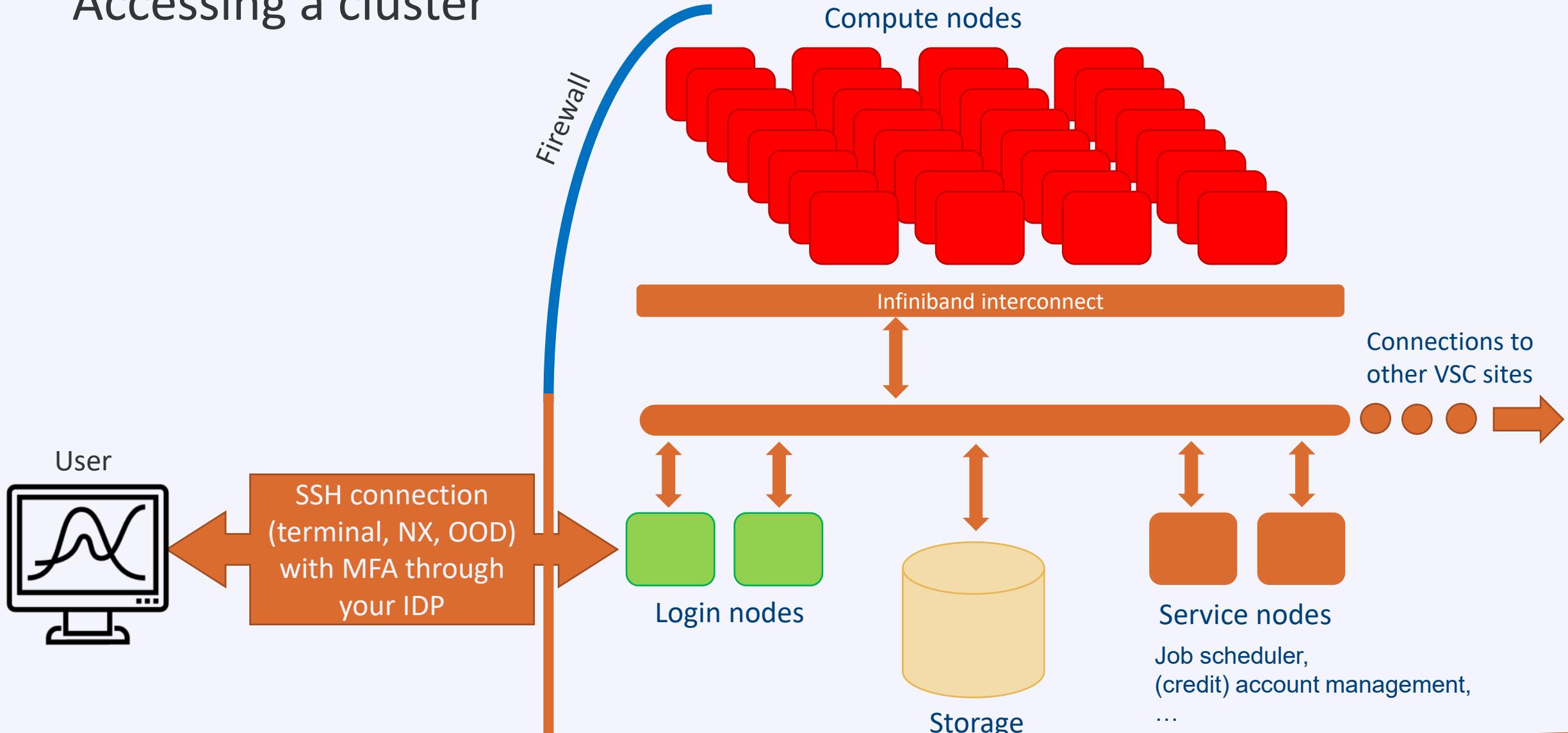
- We provide various *sam*-commands to facilitate credit related queries

Command	Purpose
sam-balance	Lists active projects and available credits
sam-list-allocations -A <account-name>	Lists the credit deposits and refunds (if any)
sam-quote sbatch [sbatch args] [myjob.slurm]	Estimates the (max) number of credits a job will use
sam-statement -A <account-name> -s <start-date> -e <end-date>	Overview of credits used by each job in some time window



Connecting to a login node

Accessing a cluster



How to login?

Open OnDemand
<ul style="list-style-type: none"><input type="checkbox"/> Web-based login via your browser<input type="checkbox"/> https://ondemand.hpc.kuleuven.be<input type="checkbox"/> Multi-Factor Authentication (MFA): Login through your institute's IDP<input type="checkbox"/> No SSH key needed
NX
<ul style="list-style-type: none"><input type="checkbox"/> Graphical desktop<input type="checkbox"/> Requires a NoMachine client (available for Linux/MacOS/Windows)<input type="checkbox"/> Suitable for applications with a GUI (MATLAB, VisIt, ParaView, ...)<input type="checkbox"/> Nvidia Quadro GPU for visualization

SSH client
<ul style="list-style-type: none"><input type="checkbox"/> Windows: PuTTY, MobaXterm, ... MacOS/Linux: any terminal<input type="checkbox"/> Optional: for VSC clusters <i>not</i> hosted at KU Leuven you need an (open)SSH key pair (upload the public key to the VSC account page and wait \geq 30mins)<input type="checkbox"/> Hosts: login.hpc.kuleuven.be:22 nx.hpc.kuleuven.be (NX)<input type="checkbox"/> Recommended to use an SSH agent<input type="checkbox"/> Can open GUIs if X11 forwarding is enabled

Using the login nodes

- To develop code
- To check your storage and credit balance
- To manage jobs (submit, check status, debug, resubmit, ...)
- To move data around
 - within VSC: use data, scratch, staging
 - outside VSC: copy/sync from/to your local storage (e.g. Globus)
- To pre- or postprocess your data
- To visualize your data
- To share files and folders

Tips

Do NOT execute CPU- or memory-intensive tasks here

Warning

Login nodes are shared resources

Submit jobs instead (you can e.g. compile software on compute nodes via an interactive job)

Use `slurmtop` to see how busy the cluster is

Connecting via a terminal

(Linux and Mac)

- ✓ Use ssh to connect:

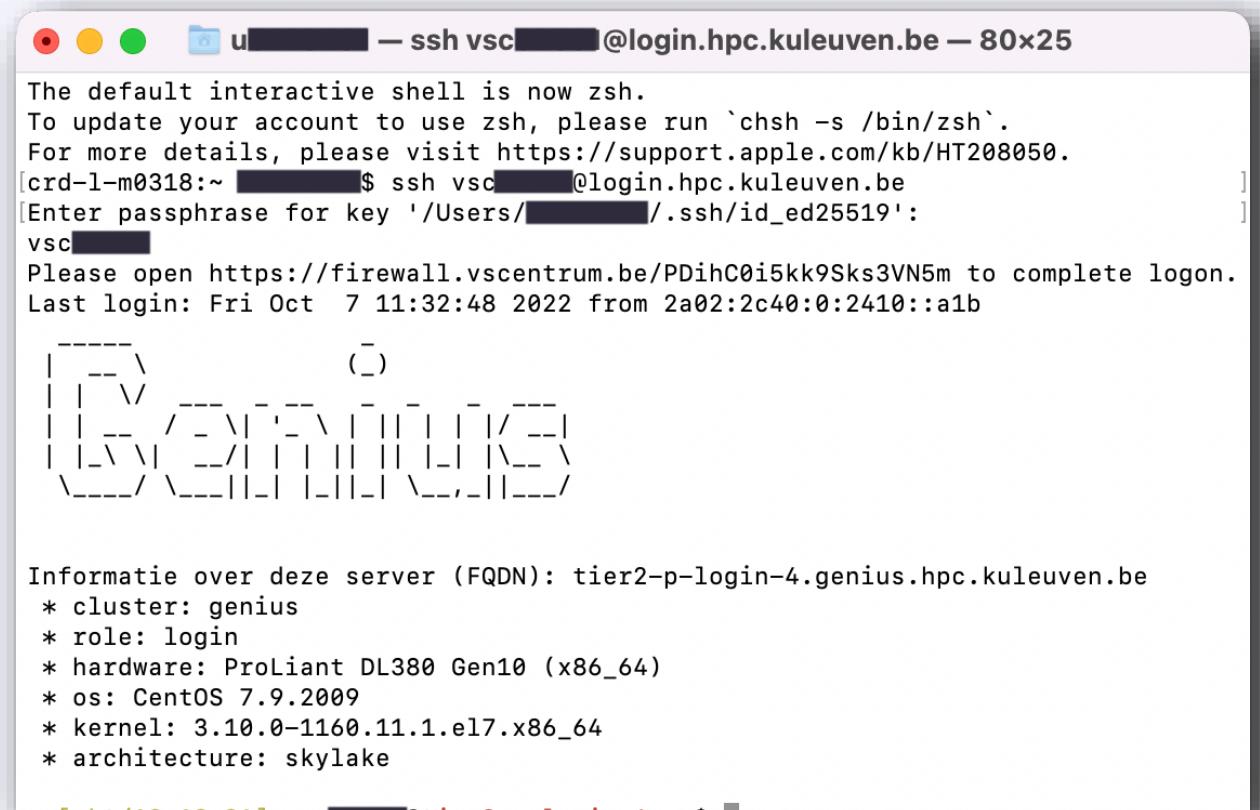
```
$ ssh -A vscXXXXX@<host_name>
```

- ✓ If key not found:

```
$ ssh -i </path/to/keyfile> ...
```

If asked for **password**, please stop connecting and contact support, otherwise after a few attempts you will be blocked for 24h.

Hostname:
login.hpc.kuleuven.be



The default interactive shell is now zsh.
To update your account to use zsh, please run `chsh -s /bin/zsh`.
For more details, please visit <https://support.apple.com/kb/HT208050>.
[crd-l-m0318:~ vscXXXXX\$ ssh vscXXXXX@login.hpc.kuleuven.be
[Enter passphrase for key '/Users/vscXXXXX/.ssh/id_ed25519':
vscXXXXX
Please open <https://firewall.vscentrum.be/PDihC0i5kk9Sks3VN5m> to complete logon.
Last login: Fri Oct 7 11:32:48 2022 from 2a02:2c40:0:2410::a1b
[REDACTED]
Informatie over deze server (FQDN): tier2-p-login-4.genius.hpc.kuleuven.be
* cluster: genius
* role: login
* hardware: ProLiant DL380 Gen10 (x86_64)
* os: CentOS 7.9.2009
* kernel: 3.10.0-1160.11.1.el7.x86_64
* architecture: skylake
✓ [okt/12 13:31] vscXXXXX@tier2-p-login-4 ~ \$

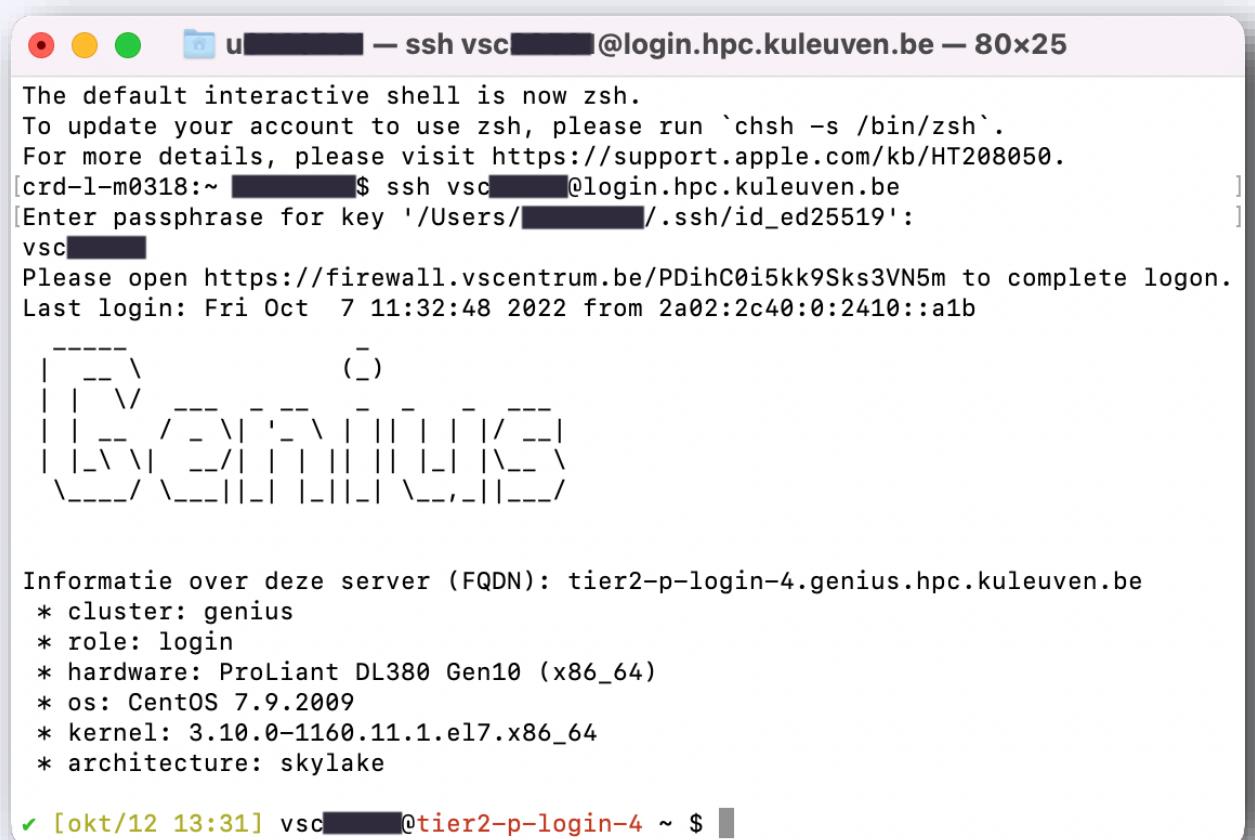
Connecting via a terminal

(Linux and Mac)

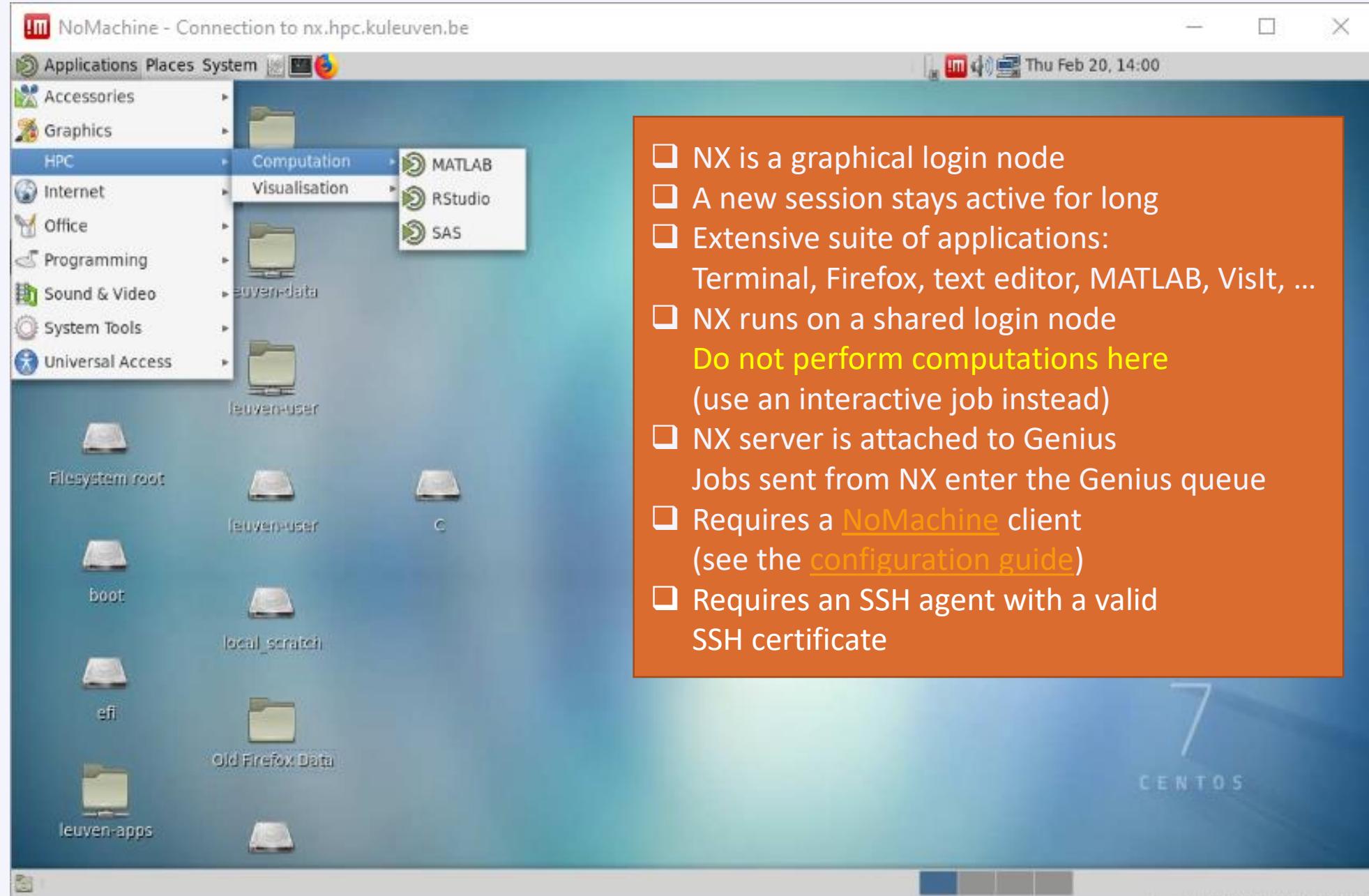
With an SSH agent

- ✓ Check your SSH agent
Is your SSH key found?
\$ ssh-add -l
 - ✓ If your SSH agent is not running:
\$ eval \$(ssh-agent)
 - ✓ If your key is not found,
add it to the agent:
\$ ssh-add </path/to/keyfile>
 - ✓ Use ssh to connect:
\$ ssh -A vscXXXXXX@<hostname>

Hostname:
login.hpc.kuleuven.be



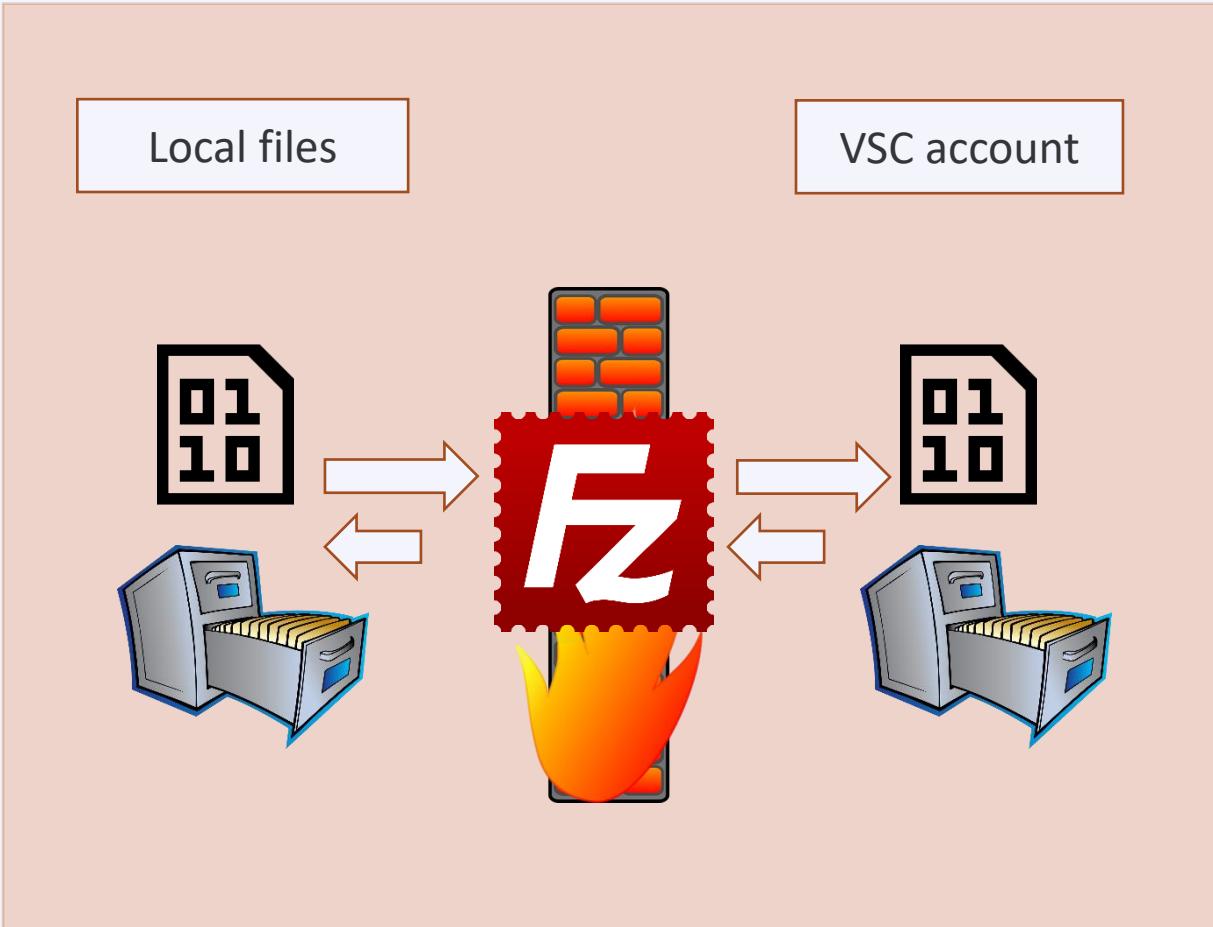
NX





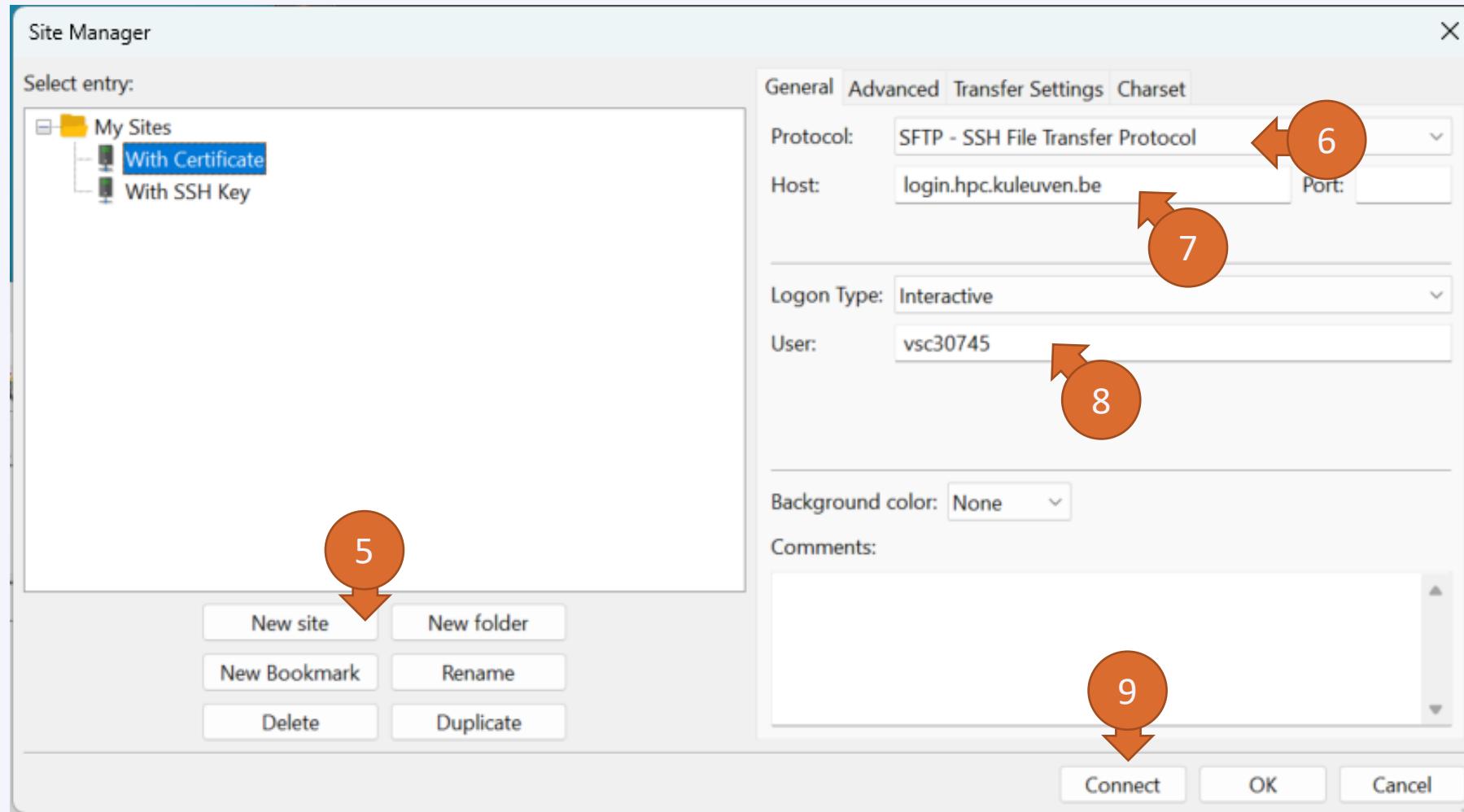
Data transfer

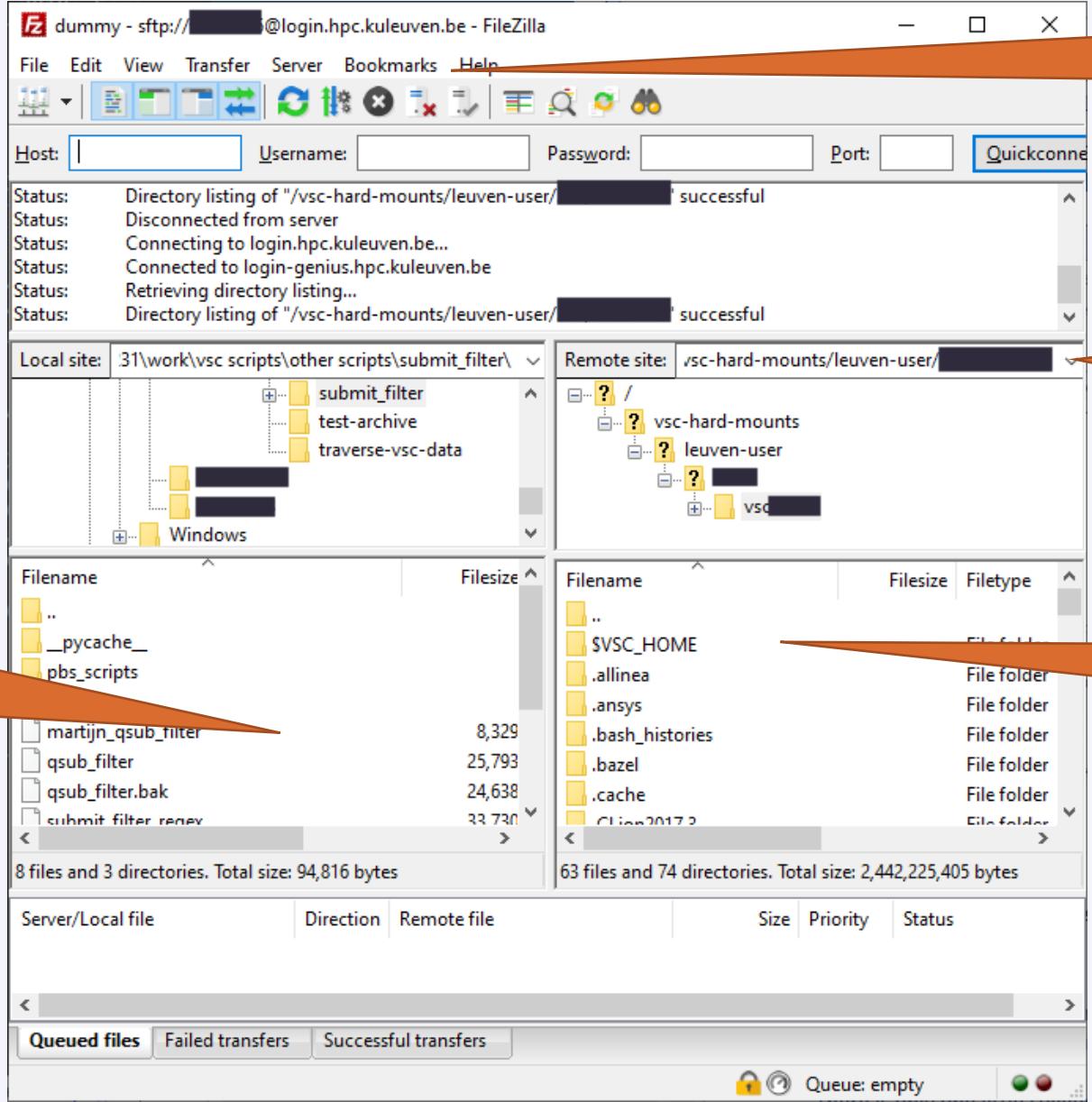
File transfer tools



Application	OS
FileZilla	Windows, Linux, Mac
WinSCP	Windows
rsync, scp	Linux, Mac
Globus	Window, Linux, Mac

File transfer with FileZilla





Content on your local machine (e.g. laptop)

It can be convenient to bookmark your data and scratch folders

Navigate here to to your \$VSC_DATA folder, e.g. /data/leuven/399/vsc39934

Content of your chosen folder

WARNING: by default you will land in your \$VSC_HOME which is typically **NOT** where you want to store data

File transfer with rsync

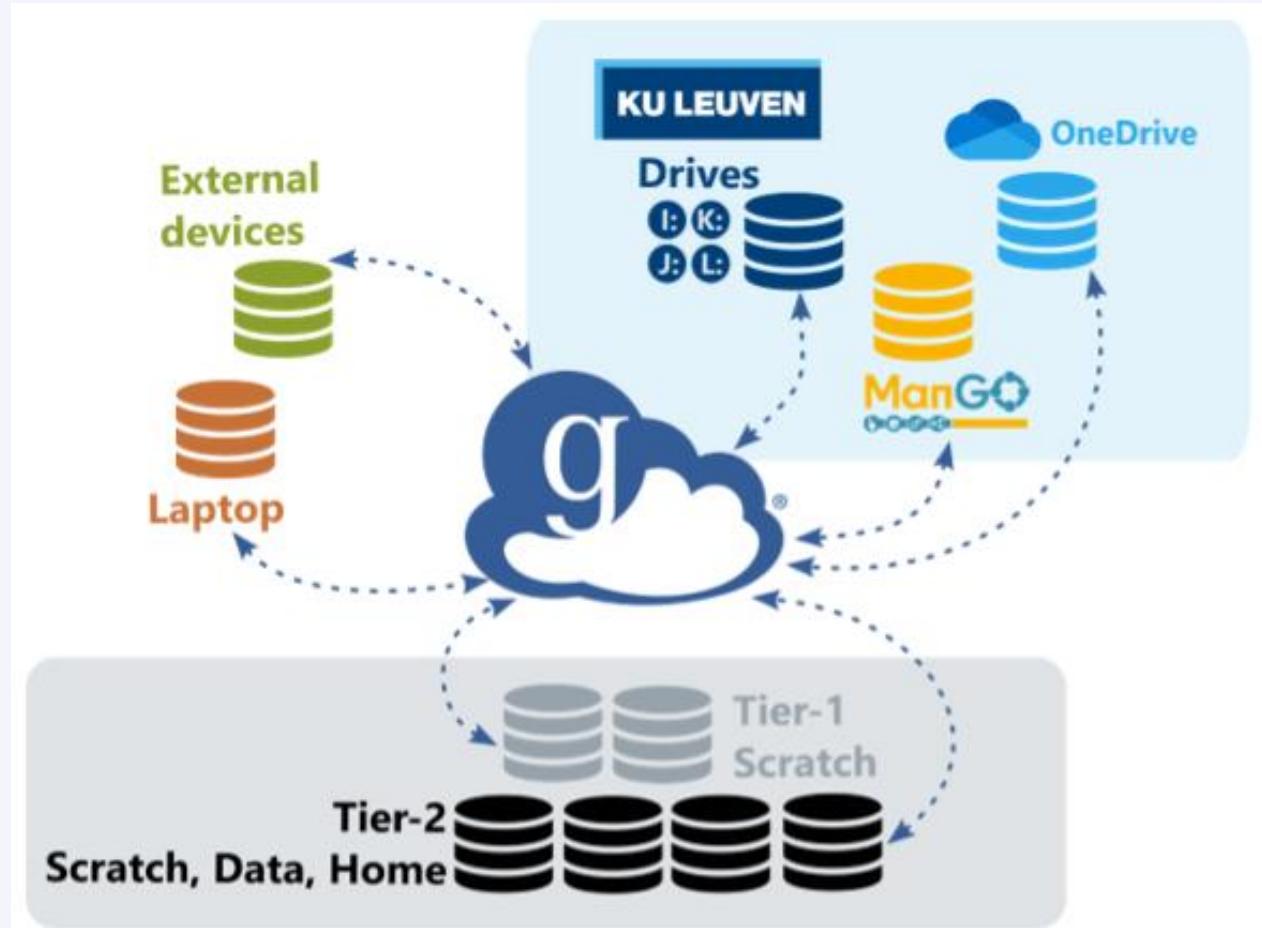
- ❑ For Linux, Mac and Windows WSL users
- ❑ Synchronize source and destination
- ❑ Listens to your SSH agent
- ❑ Example usage:

```
$ rsync -av /path/to/source vsc3XXXX@login.hpc.kuleuven.be:/path/to/destination
```

File transfer with Globus

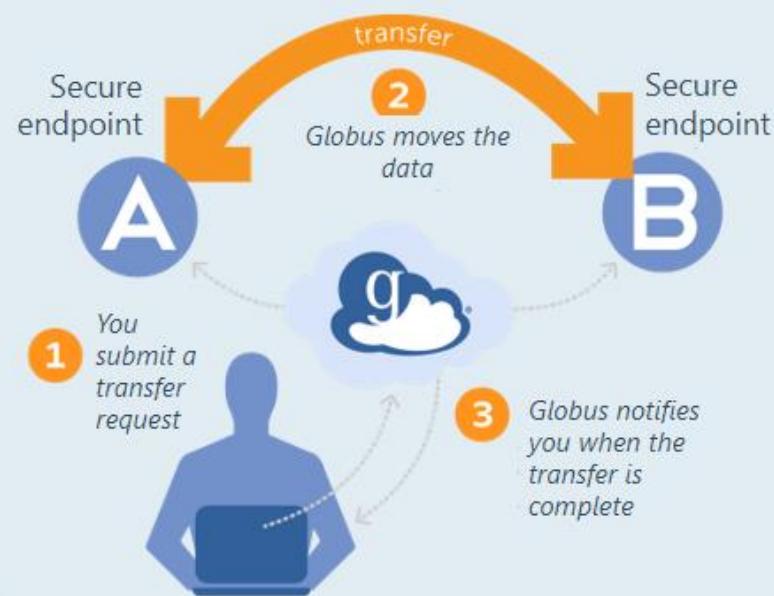
- Web interface for transferring, sharing and searching within data
- Schedule and resume (large) transfers
- Workflow:
 - Define “endpoint” on the source
 - Define “endpoint” on the destination
 - Transfer between endpoints
- Endpoints on all VSC sites:
 - all Tier-2 clusters: home, data, scratch*
 - all Tier-1 clusters: scratch, projects

* You can also access staging storage through this endpoint (Leuven Tier-2 only)

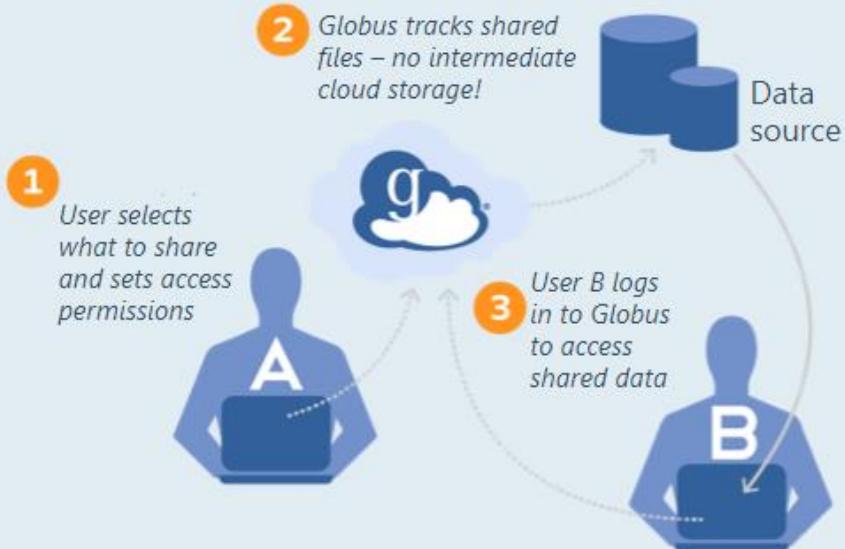




Transfer data



Share data





An orange rectangular overlay covers the bottom left portion of the image, containing the text "Open OnDemand".

Open OnDemand



Open OnDemand

- ✓ Access clusters via your web browser
- ✓ <https://ondemand.hpc.kuleuven.be>
- ✓ Login via your institute's IDP
- ✓ File browser
- ✓ Interactive apps + integrated shell
- ✓ Create, start and monitor jobs
- ✓ Develop, compile and test your code
- ✓ VSCode and PyCharm editors
- ✓ Fluent, Jupyter Lab, Rstudio, MATLAB, ParaView, Tensorboard, Gnuplot

Please close your interactive sessions
when you no longer actively use them!



KU Leuven OnDemand Apps ▾ Files ▾ Jobs ▾ Clusters ▾ Interactive Apps ▾ My Interactive Sessions

KU LEUVEN

Open OnDemand provides an integrated, single access point for all your HPC resources.

Pinned Apps A featured subset of all available apps

Active Jobs System Installed App	Home Directory System Installed App	Job Composer System Installed App	Login Server Shell Access System Installed App
MATLAB System Installed App	ANSYS FLUENT System Installed App	Visual Studio Code System Installed App	Gnuplot System Installed App
Interactive Shell System Installed App	JupyterLab System Installed App	ParaView System Installed App	PyCharm System Installed App
RAPIDS System Installed App	RStudio Server System Installed App	TensorBoard System Installed App	



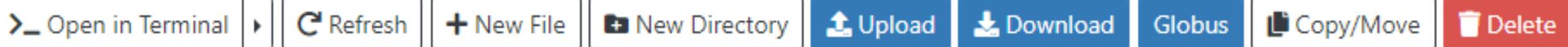
Open OnDemand

Access your VSC storage

- ✓ Use the “Files” tab to access your VSC_HOME and VSC_DATA
- ✓ VSC_SCRATCH is not available from this menu

The screenshot shows the KU Leuven OnDemand web interface. At the top, there is a navigation bar with tabs for "KU Leuven OnDemand", "Apps", "Files" (which is currently selected), "Jobs", "Clusters", "Interactive Apps", and a file icon. Below the navigation bar, there is a banner with the text "KU LEUVEN". A dropdown menu is open under the "Files" tab, showing two options: "Home Directory" and "Data Directory /data/leuven/307/vsc30745".

File/Folder management



- ✓ OK for transferring small files/folders
- ✓ Open any (sub)folder in terminal (will open a new tab)
- ✓ Deleted files/folders can be retrieved from (hourly, daily, weekly) snapshots



Open OnDemand

Monitor your active jobs

KU Leuven OnDemand Apps ▾ Files ▾ **Jobs ▾** Clusters ▾ Interactive Apps ▾ ? ▾

① Active Jobs
Job Composer

Your Jobs ▾ All Clusters ▾

Active Jobs

Show 50 entries Filter:

ID	Name	User	Account	Time Used	Queue	Status	Cluster	Actions
55337242	PostProcessing	vsc30745	lp_hpcinfo	00:02:03	batch	Completed	genius	



Open OnDemand: Login Server Shell Access

- ✓ Starts a shell in a new browser tab
- ✓ You land on a Genius login node
(from which you can also access wlCE)
- ✓ You land on your VSC_HOME
cd to data/scratch/staging immediately
- ✓ Do NOT compute on the login nodes
- ✓ Try:
\$ sam-balance
\$ myquota

Host: login.hpc.kuleuven.be Themes: Default

Login shell

```
[vsc30745@login ~]$
```

Informatie over deze server (FQDN): tier2-p-login-1.genius.hpc.kuleuven.be

- * cluster: genius
- * role: login
- * hardware: ProLiant DL380 Gen10 (x86_64)
- * os: Rocky 8.8
- * kernel: 4.18.0-477.10.1.el8_8.x86_64
- * architecture:
- * architecture-suffix:

On 25-09-2023 the operating system of the Genius cluster was upgraded to Rocky8.
The following documentation page provides an overview of important changes:
https://docs.vsczentrum.be/leuven/genius_2_rocky.html
Last login: Tue Nov 7 23:42:32 2023 from 2a02:2c40:0:2410::a1c

```
vsc30745@tier2-p-login-1 ~
$ [red]
```



Open OnDemand: Interactive Shell

- ✓ Start a shell as a job (on a compute node)
- ✓ Recommended to:
 - Install your own software (in VSC_DATA)
 - Test/run your program interactively
 - Debug a program
- ✓ Choose the relevant cluster, partition and resources (cores, memory, GPU)

Interactive Shell

Interactive Apps

Servers
Interactive Shell
Jupyter Lab
Nvidia Rapids
RStudio Server
Tensorboard
code-server

Cluster: wice

Account: lpt2_sysadmin

Partition: interactive

Number of hours: 1

Number of cores: 1

Required memory per core in megabytes: 3400

Number of nodes: 1

Number of gpu's: 0

[type:] Specify the total number of GPUs slices for the job. An optional GPU type specification can be supplied. For example "A100:3" or "3".

Reservation (optional):

Name of an existing reservation in which the job should run

I would like to receive an email when the session starts

Launch

* The Interactive Shell session data for this session can be accessed under the [data root directory](#).



Open OnDemand: Jupyter Lab

Jupyter Lab

- ✓ [Create a notebook](#) as a job (on a compute node)
- ✓ Suitable for:
 - pre/post-processing your data
 - prototyping
 - adding figures and text for pedagogical purposes
- ✓ Choose the appropriate cluster, partition and resources (cores, memory, GPU)
- ✓ User your own Python/R kernels (next slide)

Jupyter Lab

This app will launch a Jupyter Lab server on one or more nodes.

Servers

Interactive Shell

Jupyter Lab

Nvidia Rapids

RStudio Server

Tensorboard

code-server

Cluster

genius

Account

lpt2_sysadmin

Partition

gpu_p100

batch(_long) or bigmem or interactive or gpu(_p100_v100) or dedicated...

Enable nvidia rapids
Only available on w/CE interactive or gpu nodes with a GPU requested

Number of hours

1

Number of cores

9

Required memory per core in megabytes

3400

Number of nodes

1

Number of gpu's

1

[type:] Specify the total number of GPUs slices for the job. An optional GPU type specification can be supplied. For example "A100:3" or "3".

Reservation (optional)

Name of an existing reservation in which the job should run

I would like to receive an email when the session starts

Pre-run scriptlet

Bash commands to be executed before starting application, \$node,_arch variable containing for example 'skylake' is available

Launch

* The Jupyter Lab session data for this session can be accessed under the [data root directory](#).



Open OnDemand: Custom R/Python kernels

Custom kernels

- ✓ The default kernels may not offer the R/Python/... interpreter version or packages you want
- ✓ You can create a miniconda env and add it to your OnDemand kernels

Step 1: Miniconda

Start an Interactive Shell

Go to your data folder

```
$ cd ${VSC_DATA}
```

Download Miniconda

```
$ wget https://repo.continuum.io/miniconda/Miniconda3-latest-Linux-x86\_64.sh
```

Install Miniconda in your VSC_DATA

```
$ bash Miniconda3-latest-Linux-x86_64.sh -b -p ${VSC_DATA}/miniconda3
```

Permanently add the path to Miniconda to your ~/.bashrc

```
$ echo 'export PATH="${VSC_DATA}/miniconda3/bin:${PATH}"' >> ~/.bashrc
```



Open OnDemand: Custom R/Python Kernels

Step 2: Environment setup

Create a new environment <env_name> with all packages you need, e.g.

```
$ conda create --name=<env_name> numpy scipy ...
```

Enable your new environment

```
$ source activate <env_name>
```

Add the ipykernel package

```
$ conda install -c conda-forge ipykernel
```

Step 3: Register your kernel

```
$ python -m ipykernel install --user --env PYTHONPATH "" \  
--name <env_name> --display-name <env_name>
```

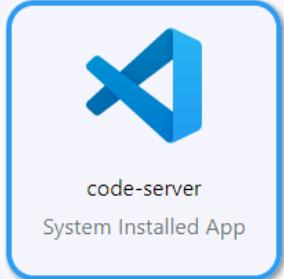
Step 4: Launching Jupyter Lab

Now start a new Jupyter Lab session

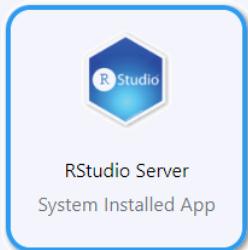
Your new kernel should be available there



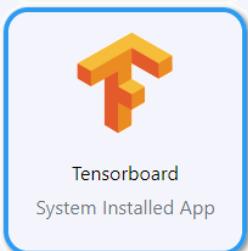
Open OnDemand: Other apps



Start Visual Studio Code server as a job
Develop, deploy and debug your workflow directly on HPC
Use of the interactive partitions is free of charge and is a good choice here
Supports many languages + Github integration



Start RStudio IDE as a job
Install your own packages in VSC_DATA



Track machine learning metrics and visualize data during the workflow
You need to provide a log folder



Open OnDemand: Other apps



Start PyCharm IDE as a job

Develop, deploy and debug your Python workflow directly on HPC

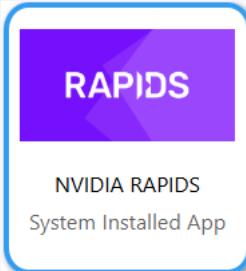
Use Python environment based on Conda or virtual environments



Use Fluent for CFD simulations (single-node)

You need to be in a license group before being able to see the app logo

Contact your supervisor to help you joining an existing license group



Use RAPIDS for accelerated data types and functionalities from NumPy, Pandas (and more) on GPUs



Open OnDemand: Resources

E.g. to start a Jupyter notebook

- ✓ Pick a valid Slurm credit account
- ✓ Default partition for most apps: *interactive*
(*interactive*: max 8 cores, 1 GPU instance, 16 hr)
- ✓ Default resources:
1 core, 1 hour, 3400 MB RAM, no GPU

Interactive Apps

Servers
● Interactive Shell
Jupyter Lab
● RStudio Server
● Tensorboard
● code-server
Work in progress
● ParaViewWeb - Work in progress
● cryosparc

Jupyter Lab

This app will launch a Jupyter Lab server on one or more nodes.

Account

lp_myproject

Partition

interactive

batch(_long) or bigmem or interactive or gpu or dedicated...

Number of hours

3

Number of cores

1

Required memory per core in megabytes

3400

Number of nodes

1

Number of gpu's

0

[type:] Specify the total number of GPUs slices for the job. An optional GPU type specification can be supplied. For example "A100:3" or "3".

Reservation (optional)

Name of an existing reservation in which the job should run

I would like to receive an email when the session starts

Launch

Software

Software: Centrally installed modules

- ✓ OS: Rocky Linux 8.10
- ✓ Toolchains:
 - *intel* (icc, icpc, ifort; Intel MPI; Intel MKL)
 - *foss* (gcc, g++, gfortran; OpenMPI; ScaLAPACK, OpenBLAS, FFTW)
 - Toolchain year on Genius from 2018a onwards; on wICE from 2021a onwards
- ✓ Note: not all toolchains are compatible with each other (conflicting dependencies)
For example, **never mix ...-foss-... and ...-intel-...** modules

Command	Remark
module av	List all available modules
module av Python	List all Python-related modules
module spider Python	Get more info
module load Python/3.12.3-GCCcore-13.3.0	Load a specific module
module list	List all loaded modules and their dependencies
module unload Python/3.12.3-GCCcore-13.3.0	Unload a module (but dependencies still stay)
module purge	Remove all modules from your work session

Software: Your specific needs

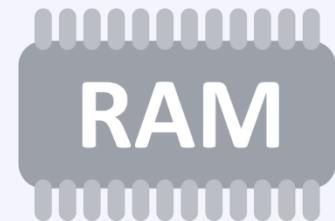
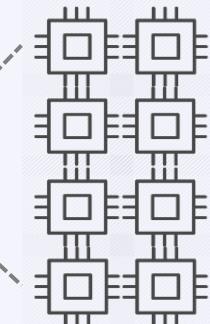
- ✓ If you need a software package that is not yet available as a module, you can ask us to provide it
 - ✓ For Python and R packages, however, we prefer that you install it yourself
 - ✓ Read more about [Python Package Management](#)
 - ✓ Read more about [R Package Management](#)
- ✓ If you want to compile software yourself:
 - ✓ Best install it in your VSC_DATA
 - ✓ Use *intel* or *foss* toolchains
 - ✓ Compile your code on a compute node (with interactive job)
 - ✓ If you run into problems, don't hesitate to contact us



Starting to compute

Resource glossary

- ✓ **Cluster:** which machine to use? Genius or wICE?
- ✓ **Nodes:** how many compute servers to allocate?
- ✓ **Cores:** how many cores per node to allocate?
- ✓ **Memory:** how much CPU memory to allocate?
- ✓ **Partition:** batch, gpu, bigmem, ...?
- ✓ **Walltime:** how long to allocate the resources?
- ✓ **Storage:** how much storage (data, scratch, etc) will the job require?
- ✓ **Credits:** how many compute credits will the job consume?



Default resources

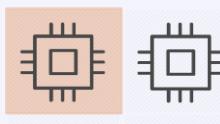
- ✓ **Cluster:** the cluster associated with the node you're on (Genius or wICE)
- ✓ **Nodes:** 1
- ✓ **Cores:** 1
- ✓ **Memory:** depends on which node(s) you use
- ✓ **Partition:** batch
- ✓ **Walltime:** typically 1 hour
- ✓ **Storage:** no default
- ✓ **Credits:** no default



Serial application

1 process on 1 core

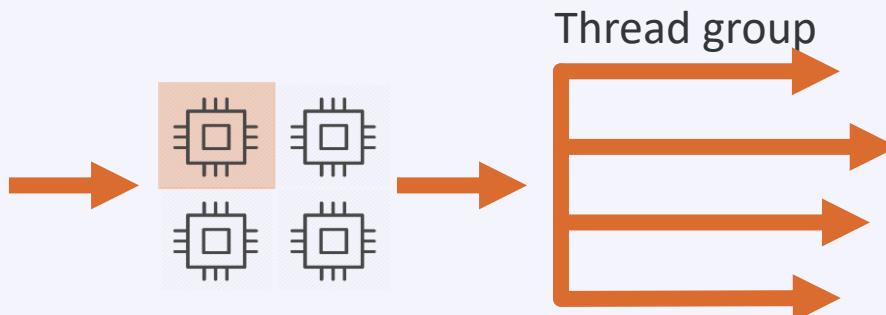
Single process



One node

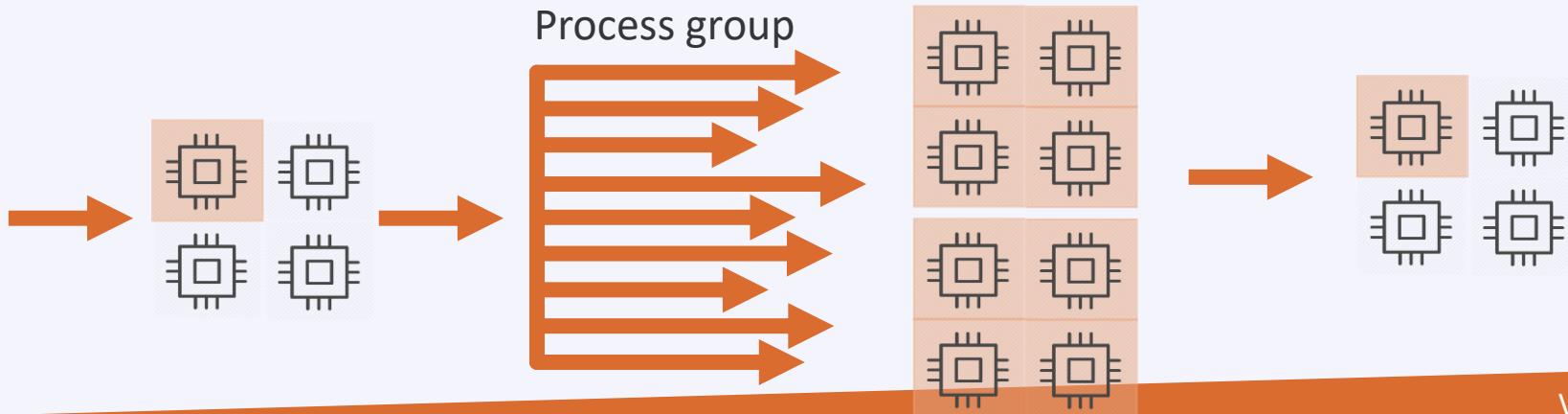
Multithreaded application

N threads on N cores
on 1 node

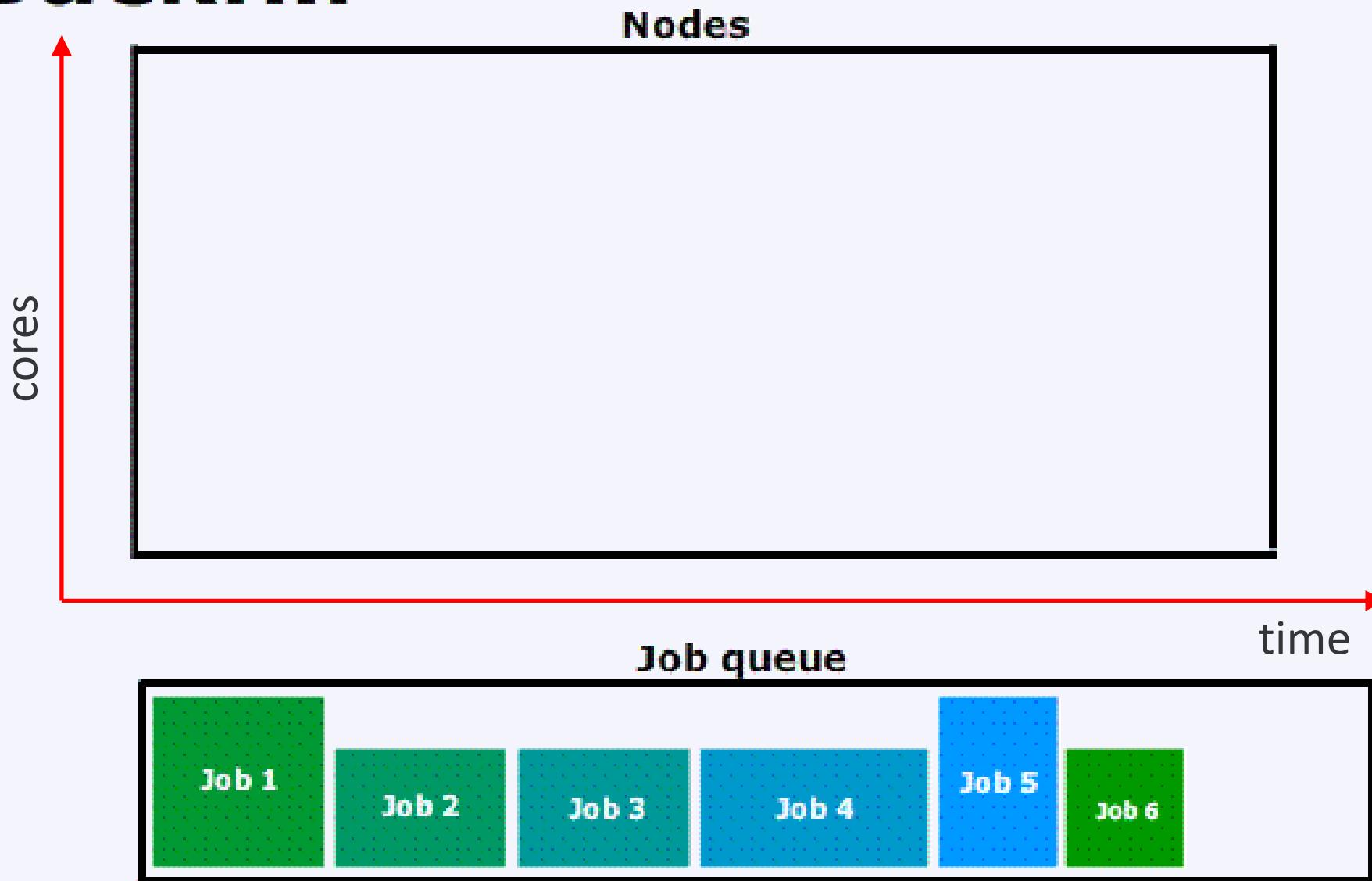


Distributed application

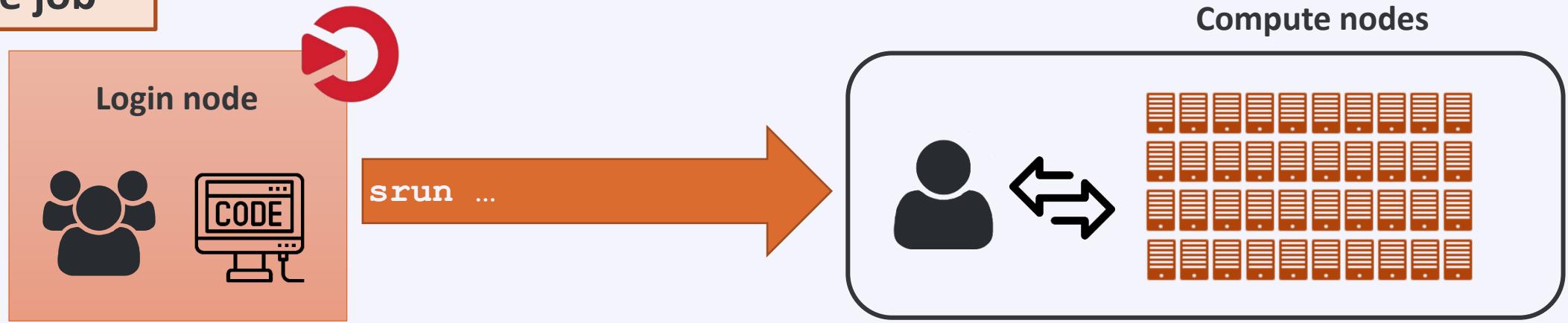
Many processes on
multiple cores,
possibly spread over
multiple nodes



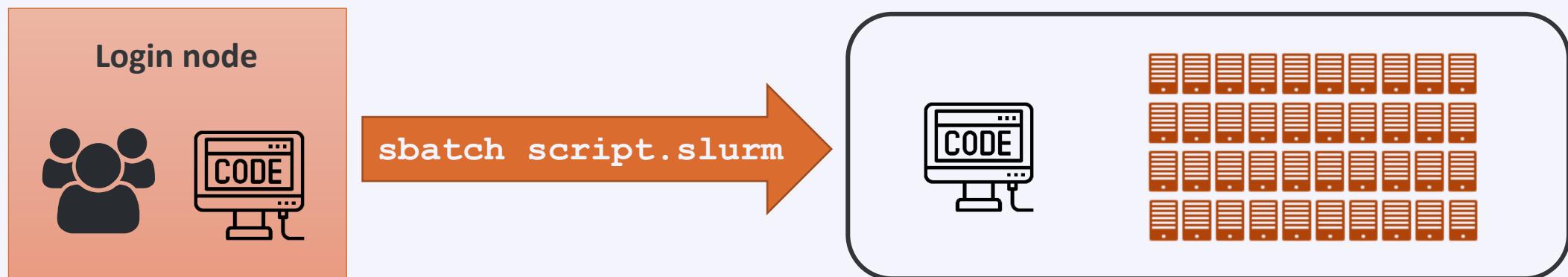
Backfill



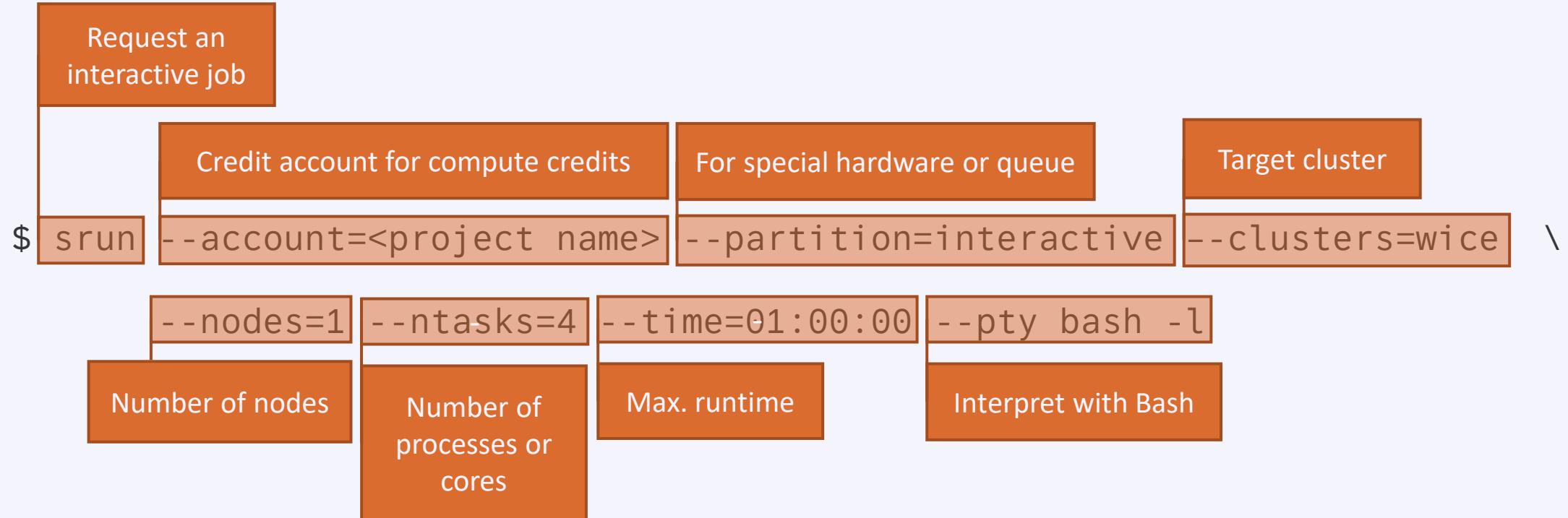
Interactive job



Batch job



Example interactive job on wICE



Remark

- Specifying <project name> for credits is mandatory, e.g. **-A lp_hpcinfo**
- Implicit defaults for an interactive job on wICE are
`-n 1 -t 01:00:00 -p batch --mem-per-cpu=3400M`

Slurm command-line tools

Command	Purpose
\$ sbatch ...	Submit a batch job
\$ srun ...	Submit an interactive job
\$ scancel --cluster=wice <JobID>	Cancel a specific pending or running job
\$ scontrol show job --clusters=wice <JobID> \$ slurm_jobinfo <JobID>	Detailed job info (very useful to diagnose issues)
\$ squeue --clusters=wice --long	Status of all recent jobs
\$ squeue --clusters=wice --start	Give a rough estimate of when your jobs will start
\$ sinfo --clusters=wice	Info about the state of available partitions and nodes
\$ sacct --clusters=wice --batch --job <JobID>	Show the jobsript used for a given batch job
\$ slurmtop	Overview of the cluster
\$ scontrol --clusters=wice show node <hostname>	Get detailed information about the status of a node
\$ sam-balance	Overview of all your available credit projects
\$ sam-list-allocations	Detailed overview of your credit allocation history

Slurm job submit options

Slurm option	Remarks
--nodes=X --ntasks-per-node=Y --ntasks=X*Y --ntasks=X --cpus-per-task=Y	or or (e.g. Hybrid MPI + OpenMP)
--partition=<partition_name>	Default: "batch" partition
--mem-per-cpu=<size><M>	Min memory per core, e.g. 5000M (Genius)
--time=<dd-hh:mm:ss>	e.g. 1-12:30:00
--job-name=<job_name>	
--output=<file_template>	STDOUT; default="slurm-%j.out"
--error=<file_template>	Default: redirect to STDOUT
--mail-type=FAIL,BEGIN,END	
--mail-user=<email-address>	
--export=<ALL, key=value>	Additional variables to pass to the job
--account=<account_name>	Credit account (mandatory)

Slurm job submit options

Slurm option	Remarks
--account=<account_name>	Mandatory (credits)
--partition=<partition_name>	Default: "batch" partition
--time=<dd-hh:mm:ss>	e.g. 1-12:30:00
--nodes=X --ntasks-per-node=Y --ntasks=X*Y --ntasks=X --cpus-per-task=Y	or or (e.g. Hybrid MPI + OpenMP)
--mem-per-cpu=<size><M>	Min memory per core, e.g. 5000M (Genius)
--job-name=<job_name>	
--output=<file_template>	STDOUT+STDERR; default="slurm-%j.out"
--mail-type=FAIL,BEGIN,END	
--mail-user=<email-address>	
--export=<ALL, key=value>	Additional variables to pass to the job

Long and short options

Some (not all) of the sbatch, srun and salloc command line options also have a short version

Short version	Long version	Meaning
-A	--account	Slurm account name
-a	--array	Job array range
-M	--clusters	Cluster name
-c	--cpus-per-task	Number of cores per task (default: 1)
-d	--dependency	Job dependency (after, afterok, afterany, ...)
-i	--input	STDIN filename
-e	--error	STDERR filename
-o	--output	STDOUT filename
-t	--time	Maximum walltime
-N	--nodes	Minimum num nodes
-n	--ntasks	Maximum num tasks
-p	--partition	Partition name

Interactive jobs

- ✓ Interactive job with default options (1 core for 1 hour on the *batch* partition)

```
$ srun --clusters=wice --account=lp_hpcinfo --pty bash -l
```

- ✓ Interactive job with X-forwarding

```
$ srun --clusters=wice --account=lp_hpcinfo --x11 --pty bash -l
```

- ✓ Interactive job with multiple cores, on the *interactive* partition

```
$ srun --clusters=wice --account=lp_hpcinfo --nodes=1 --ntasks=4 \
--partition=interactive --pty bash -l
```

- ✓ Interactive job with multiple cores and a GPU device

```
$ srun --clusters=wice --account=lp_hpcinfo --nodes=1 --ntasks=18 \
--partition=gpu --gpus-per-node=1 --pty bash -l
```

- ✓ To join one of your running jobs (connecting to the first compute node in the job's allocation):

```
$ srun -M wice --jobid=<JobID> --overlap -pty bash -l
```

Example Slurm job script

```
#!/bin/bash -l
#SBATCH --clusters=wice
#SBATCH --account=lp_hpcinfo
#SBATCH --nodes=1
#SBATCH --ntasks=4
#SBATCH --mem-per-cpu=3400M
#SBATCH --job-name=hpc_workflow
#SBATCH --mail-type=BEGIN,END,FAIL
#SBATCH --mail-user=my.name@kuleuven.be

module load intel/2024a
module load Python/3.12.3-GCCcore-13.3.0
which python
cd $VSC_SCRATCH/projects/simulations
cp -r $VSC_DATA/input_data .

python modelling.py

cp -r output_data $VSC_DATA
rm -rf ./input_data ./output_data
```

The diagram illustrates the components of the Slurm job script:

- Shebang**: The first line of the script, `#!/bin/bash -l`, is highlighted.
- Resource list**: The resource requirements section, starting with `#SBATCH` directives, is highlighted.
- Loading modules**: The `module load` commands used to load software environments are highlighted.
- Move data**: The `cd` and `cp` commands used to move data between directories are highlighted.
- Execute commands**: The command `python modelling.py` is highlighted.
- Move data**: The final `cp` and `rm` commands used to move data are highlighted.

Batch job workflow

- ✓ Submit the job to the scheduler
- ✓ Receive a unique JobID
- ✓ Error and output files

Job script

```
#!/bin/bash -l  
...  
...
```

Submit command

```
$ sbatch simulation.slurm
```

JobID

**Submitted batch job 60042478 on cluster
wice**

stderr, stdout

```
$ ls *.out  
slurm-60042478.out
```

Output files

- ✓ STDERR and STDOUT are by default redirected to a single file:

slurm-<JobID>.out

- ✓ Contains job info, all errors and warnings, and printouts
- ✓ Worth going through (especially if it's a new kind of job)
- ✓ Address all warnings and errors (if you can)
- ✓ Typical error examples on the right

STDOUT + STDERR

```
$ ls slurm-* .out  
slurm-60042478.out
```

Out of memory

```
slurmstepd: error: Detected 1 oom-kill event(s).  
Some of your processes may have been killed by the  
cgroup out-of-memory handler.
```

Short walltime

```
slurmstepd: error: *** JOB 60042478 ON s28c11n2  
CANCELLED AT 2023-02-08T10:03:43 DUE TO TIME LIMIT  
***
```

Low disk space

```
IOError: [Errno 122] Disk quota exceeded
```

Output files

- ✓ Standard output and error channels can be redirected to other files:

```
#SBATCH --output ...
#SBATCH --error ...
```

stdout

```
$ ls slurm-* .out
slurm-60041238.out
```

Output file

```
SLURM_JOB_ID: 60033947
SLURM_JOB_USER: vscXXXXXX
SLURM_JOB_ACCOUNT: lp_wice_pilot
SLURM_JOB_NAME: testjob
SLURM_CLUSTER_NAME: wice
SLURM_JOB_PARTITION: batch
SLURM_NNODES: 1
SLURM_NODELIST: m28c30n4
SLURM_JOB_CPUS_PER_NODE: 72
Date: Tue Jan 10 17:02:04 CET 2023
Walltime: 00-01:00:00
=====
/apps/leuven/rocky8/icelake/2021a/
softwar
e/intel-compilers/2021.2.0/compile
r/2021.2.0/linux/bin/intel64/icc
cp: cannot stat '/apps/leuven/trai
ning/test': No such file or directo
ry
Hello World
```

Resource summary

stdout

Job monitoring

Example

```
$ scontrol show job --clusters=wice 60049330
  JobId=60049330 JobName=f70.slurm
  UserId=vsc3... (253...) GroupId=vsc3... (253...)
Account=lp_my_project QOS=lp_my_project
JobState=PENDING Reason=QOSGrpBillingMinutes
  SubmitTime=2023-02-16T00:24:58 EligibleTime=2023-02-16T00:24:58
  Partition=batch NodeList= NumNodes=4-4 NumCPUs=288 NumTasks=288 CPUs/Task=1
  TRES(cpu=288,mem=979200M,node=4,billing=733
  MinCPUsNode=72 MinMemoryCPU=3400M
  WorkDir=/vsc-hard-mounts/leuven-data/3.../vsc3.../Odonate_SOM
  StdErr=/vsc-hard-mounts/leuven-data/3.../vsc3.../Odonate_SOM/slurm-60049330.out
  StdIn=/dev/null
  StdOut=/vsc-hard-mounts/leuven-data/3.../vsc3.../Odonate_SOM/slurm-60049330.out
```

Diagnosis

Why does my job remain in a pending state?

Check out the “Reason” on Slurm docs: https://slurm.schedmd.com/resource_limits.html

Other wlCE partitions

GPU

```
#SBATCH --partition=gpu
#SBATCH --nodes=1
#SBATCH --ntasks=18
#SBATCH --gpus-per-node=1
```

Big memory

```
#SBATCH --partition=bigmem
#SBATCH --nodes=1
#SBATCH --tasks-per-node=72
#SBATCH --mem-per-cpu=28000M
```

Interactive partitions

- ✓ Accessible via command line and Open OnDemand
- ✓ To quickly compile, test, debug your (parallel) application
- ✓ To pre- or postprocess your data (incl. visualizations)
- ✓ Short queue time
- ✓ 12 nodes on Genius: 36 cores/node, 192 GB mem
- ✓ 4 nodes on wICE: 64 cores/node, 512 GB mem, 1 physical GPU (split in 7 instances)
- ✓ Max resource per user: 8 cores, 1 GPU instance, 16 hour walltime
- ✓ Free of charge

Interactive job

```
$ srun --account=lp_myproject --clusters=wice \
    --partition=interactive --nodes=1 --time=16:00:00 \
    --gpus-per-node=1 --pty bash -l
```

Debugging partitions

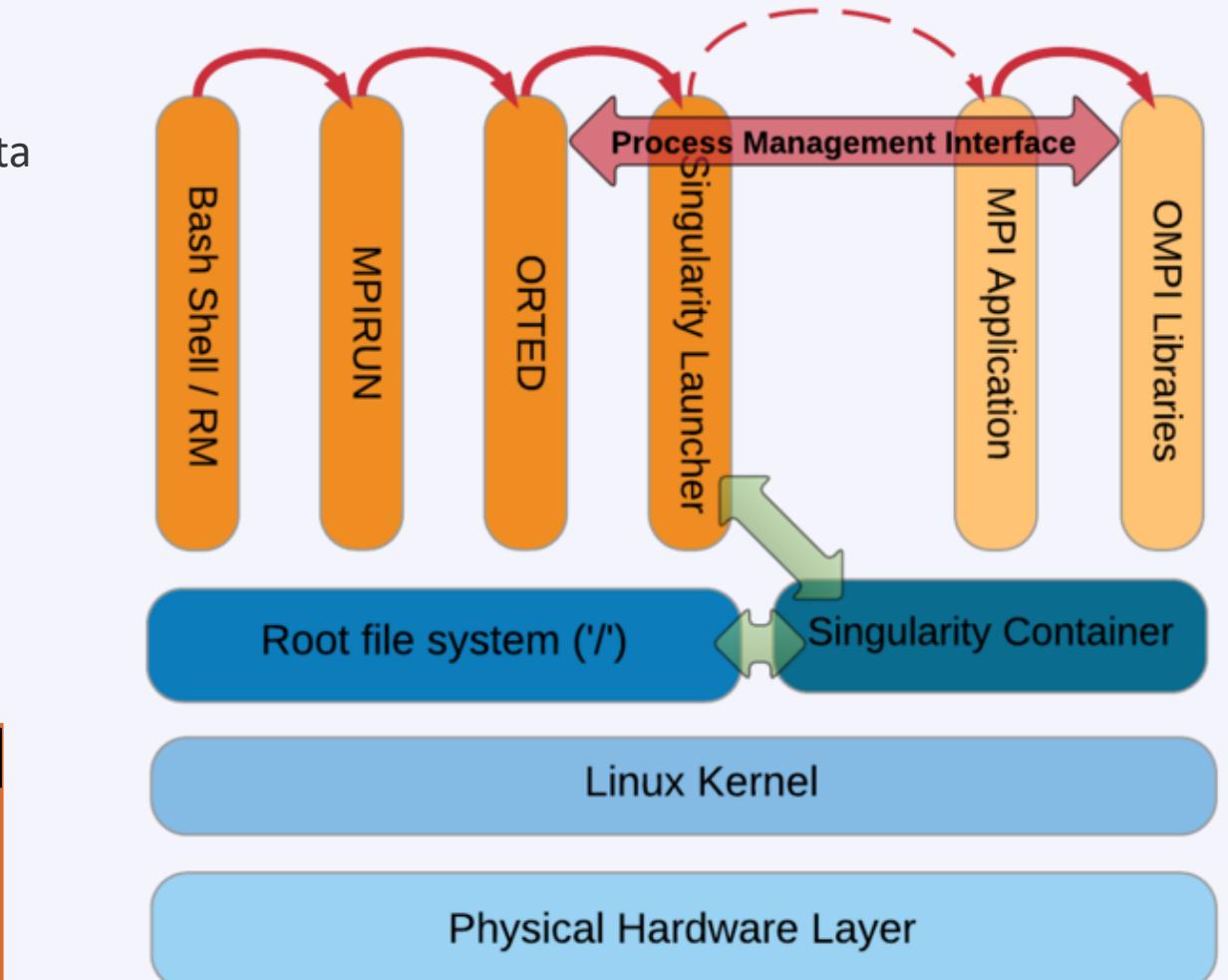
- ✓ Accessible via command line and Open OnDemand
- ✓ Quickly test if your (parallel) application works
- ✓ Short queue time
- ✓ Only one job at a time, max walltime: 1 hr

Cluster	Partition	Resources
Genius	batch_debug	2 Cascadelake nodes
	gpu_p100_debug	1 Skylake node with 4 P100 GPUs
wICE	interactive	4 Icelake nodes, each with 7 MIG instances of 1 A100 GPU
	gpu_a100_debug	1 Icelake node with 1 A100 GPU

Containers: Apptainer (Singularity)

- ✓ What?
 - Self-contained OS image in which you can install software and include data
- ✓ Why?
 - Package managers like apt can be convenient
 - Portability: images can be used on various hardware (but performance may be sub-par)
- ✓ How?
 - Create the image
 - Run it on wICE or Genius
 - Parallelization should work
(multi-node MPI runs may require some care)

```
#!/bin/bash -l
#SBATCH --time=30:00
#SBATCH --nodes=1
#SBATCH --ntasks=72
#SBATCH --clusters=wice
#SBATCH --account=lp_hpcinfo
apptainer run Project.sif ./model.exe
```



High-throughput computing with *atools*

- ✓ You can use atools to handle high-throughput computing workloads
- ✓ module load atools/<version>
- ✓ VSC training: Workflows for Data Science
- ✓ Training material: <https://github.com/gjbx/Workflows-for-HPC>
- ✓ Recording: <https://www.youtube.com/watch?v=GJwaRyi1LVA>
- ✓ User documentation: <https://atools.readthedocs.io/en/latest/>



Demo: try things yourself

Demo: try things yourself

- ✓ Request to become a member of the *lp_hpcinfo* group via account.vscentrum.be
- ✓ Transfer a file with FileZilla
- ✓ Login via NX
- ✓ Check your disk quota
- ✓ Check your available credits
- ✓ Check/load/list/unload/purge module

Demo: try things yourself

- ✓ Copy /apps/leuven/training/HPC_intro/ to your \$VSC_DATA
- ✓ Submit cpujob.slurm to the cluster
- ✓ List all your jobs with squeue -M wice
- ✓ Check the information about the job with slurm_jobinfo -M wice <job_ID>
- ✓ Modify the mpi.slurm script to request 1 node, 72 cores for 30 minutes, with job start/end e-mail notifications
- ✓ Check the status of all the jobs

Demo: job monitoring

✓ Submit an interactive job

Run your program on a compute node

Open a new terminal and ssh to the allocated compute node

Check the resources usage with htop

✓ Submit a GPU batch job

While the job is running, check its details with slurm_jobinfo

and check resource on the compute node using ssh, htop and nvidia-smi

Useful links

Helpdesk (email): hpcinfo@kuleuven.be

Helpdesk (online form): https://admin.kuleuven.be/icts/HPCinfo_form/HPC-info-formulier

VSC website: <https://www.vscentrum.be>

VSC documentation: <https://docs.vscentrum.be>

VSC agenda, training sessions, events (User Day): <https://www.vscentrum.be/vsctraining>

System status page: <https://status.vscentrum.be>

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