

VSC HPC Introduction

ICTS KU Leuven

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Material

- ☐ Everything is on Github:
 - https://hpcleuven.github.io/HPC-intro/
- ☐ Video Recordings
 - Scan the QR code
 - Recommended videos: ~ 2 hrs
 - Optional videos: ~1 hr



What is High Performance Computing?

- using supercomputers to solve advanced computation problems
- ☐ Reduce the computation time from days, years, decades, or centuries to minutes, hours, days, or weeks
- The key is parallelism



In practice, it is more like ...



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

Outline

- What is the VSC?
- What is a cluster?
- Genius Cluster
- Storage
- Login nodes & MFA
- Connection Setup
- Software environment
- How to submit jobs?
- Dedicated hardware
- How to choose resources?
- Optional material
 - Linux in brief
 - Conda for Python and R
 - Worker Framework







VSC HPC Environments







Tier-1



Hortense











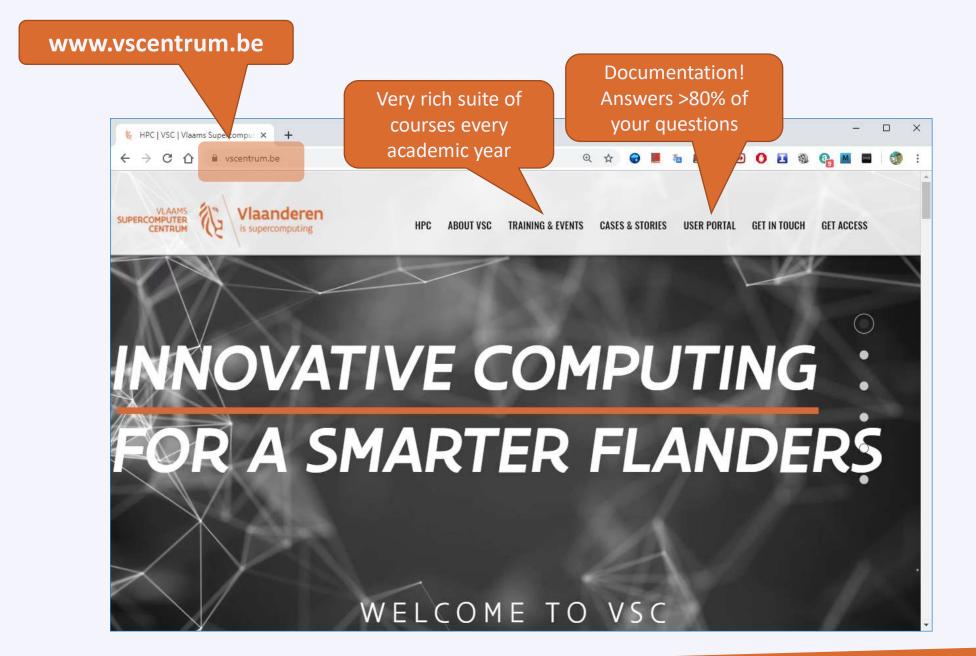


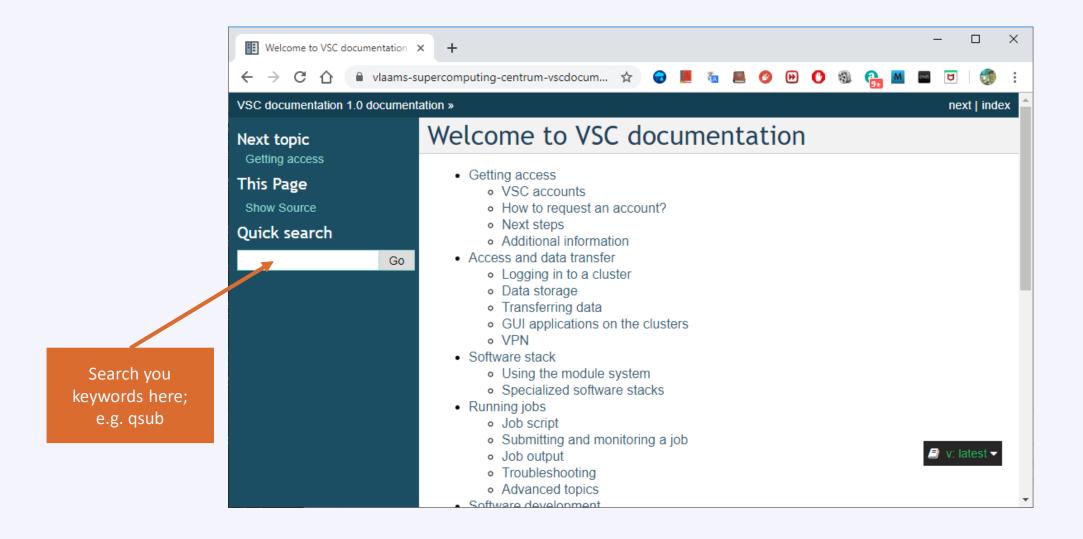
HOPPER/LEIBNIZ





GENIUS/WICE





Support and Services

Basic support

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

Application support

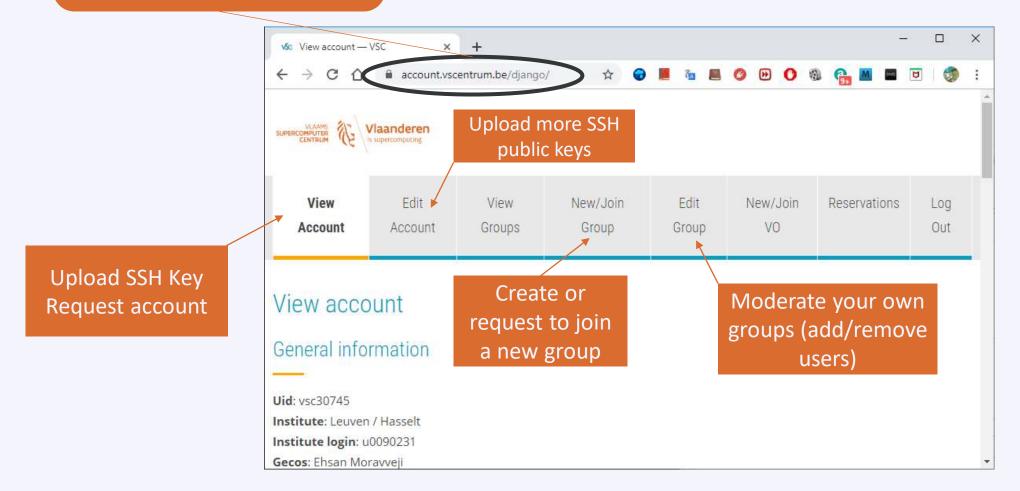
- Installation and porting
- Optimisation and debugging
- Benchmarking
- Workflows and best practices

Training

- Documentation and tutorials
- Scheduled trainings / workshops
- On request workshops
- One-to-one sessions

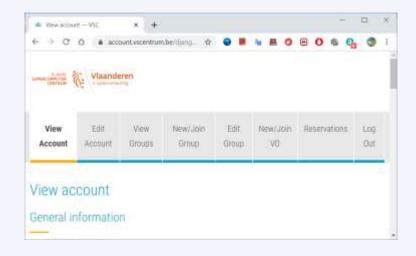
To manage your VSC account:

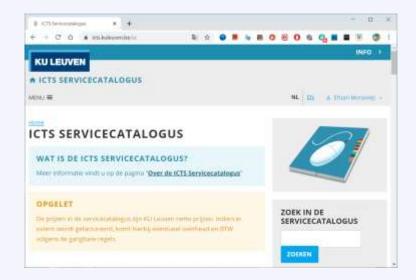
account.vscentrum.be

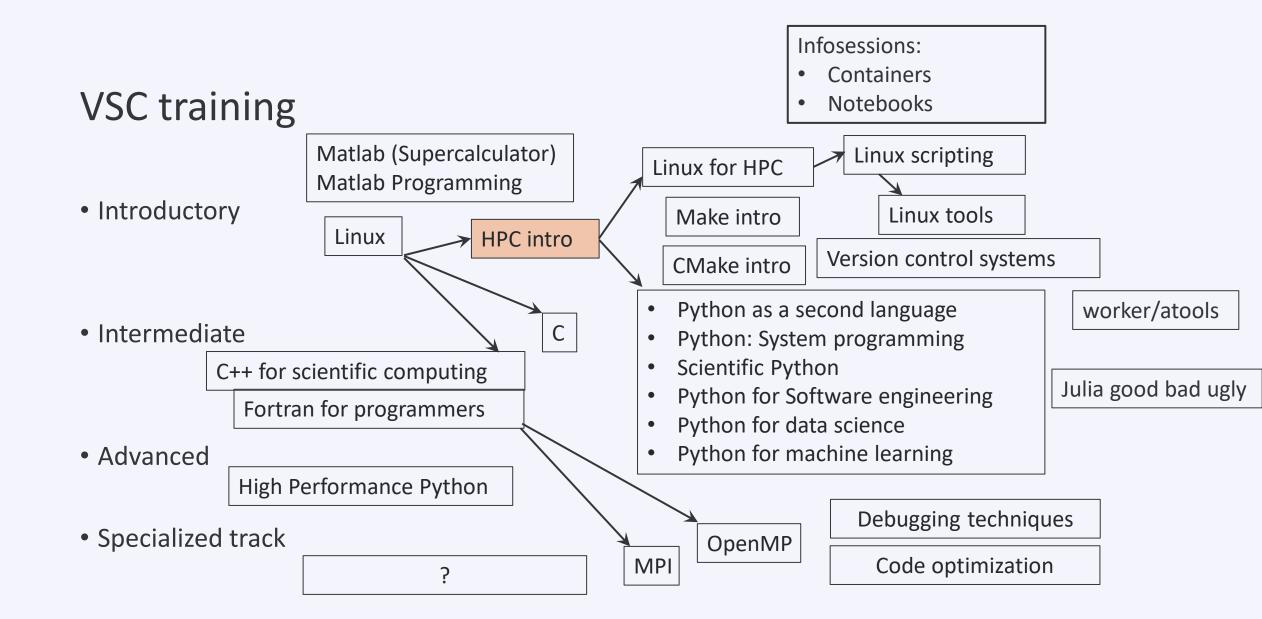


Become a VSC user

☐ Create a secure (4096 bit) SSH key pairs Upload it on the account page: www.account.vscentrum.be ☐ You need to <u>request a VSC account</u> Normally processed swiftly ☐ Request introductory credits (2M free credits for 6 months) ☐ Request <u>project credits</u> (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 all explained Go to ICTS service catalogus: https://icts.kuleuven.be/sc Click on High Performance Computing (NL/EN)







To Acknowledge VSC in publications

Why?

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

At KU Leuven

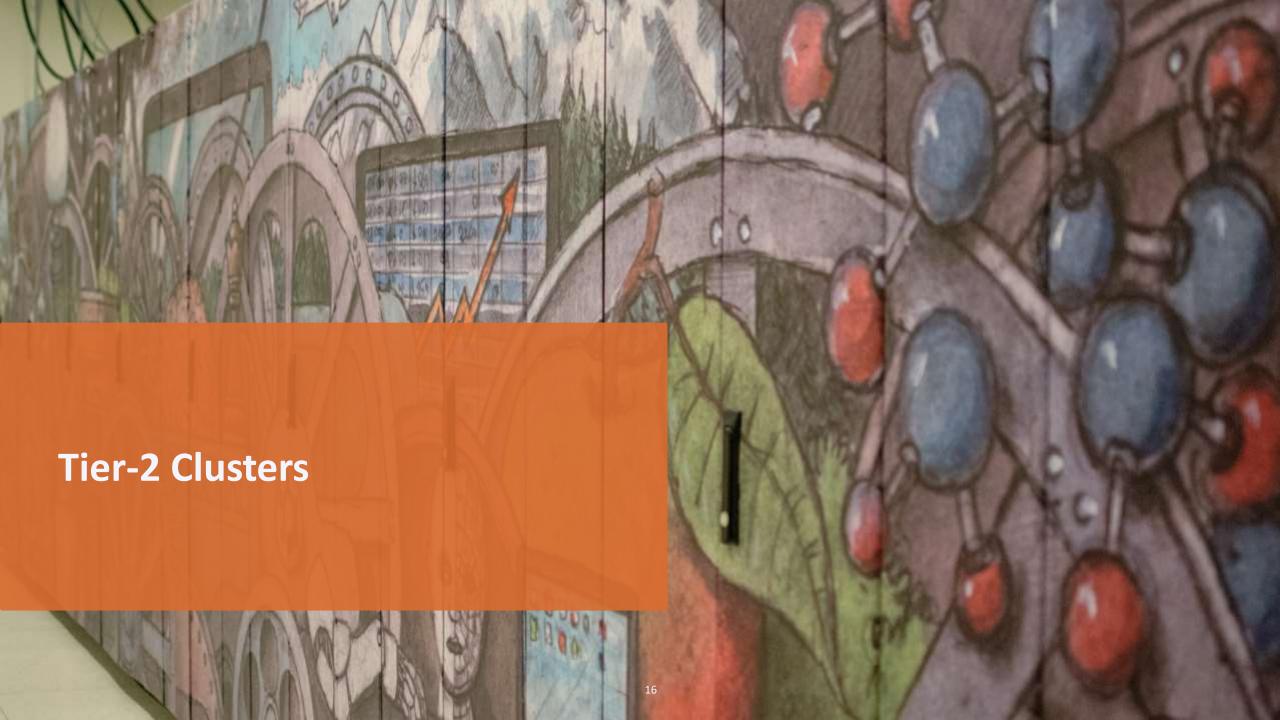
☐ add the relevant papers to the virtual collection "High Performance Computing" in Lirias

In het nederlands

De rekeninfrastructuur en dienstverlening gebruikt in dit werk, werd voorzien door het VSC (Vlaams Supercomputer Centrum), gefinancierd door het FWO en de Vlaamse regering – departement EWI.

In English

The computational 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 — department EWI.

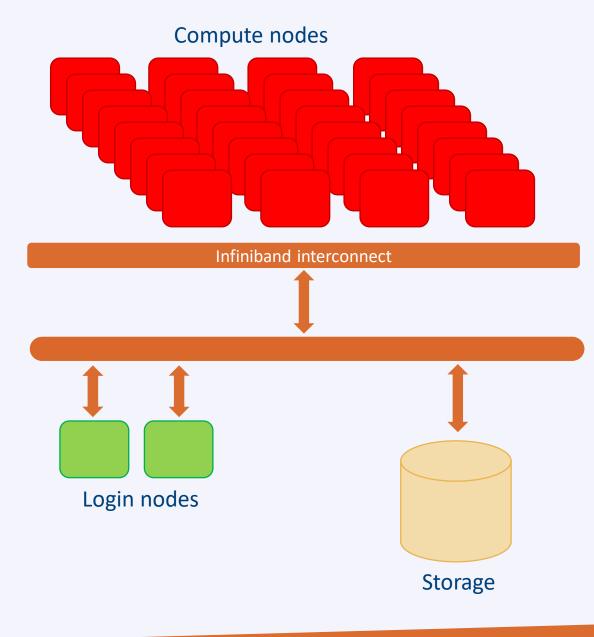


What is a Cluster?

A cluster is comprised of many powerful interconnected compute nodes

Some Terminology:

	What	Components
Cluster	The primary part of a cluster are its many powerful interconnected compute nodes.	Compute nodes Login nodes Storage space Interconnect Network
Node	A single computer connected with other nodes via an interconnect network	CPU, Storage, Memory,
CPU	Processing Component of a computer, also called the processor .	Controllers, Cache Memory, Processing Cores
Core	Processing element of the CPU.	Control Unit, Arithimic- Logic Unit, Memory.



Tier-2 Clusters @ KU Leuven

Genius (since 2018) 250 nodes: 8,936 cores



wICE (since 9/2022) 186 nodes; 13,392 cores



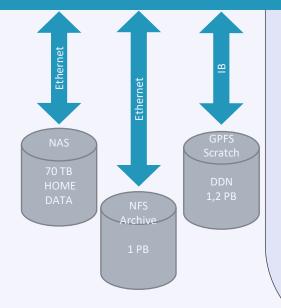
wICE (since 9/2022) 186 nodes; 13,392 cores



IB EDR

SkyLake Cascade Lake Superdome 20 nodes 86+10 nodes 2 nodes 120 nodes 720 cores 3456 cores 72 cores 4320 cores 2x Intel 2x Intel 2x Intel Skylake 2x Intel Skylake Cascadelake Cascadelake 18 cores 18 cores 18 cores 18 cores **AMD** 192 GB 768 GB 4 x NVIDIA P100 192 GB

Belnet



2 nodes
72 cores

2x Intel Skylake
18 cores

2x Intel Skylake
18 cores

2x Intel Skylake
18 cores

1x NVIDIA
P600

Watch out with login scripts!

login-genius.hpc.kuleuven.be

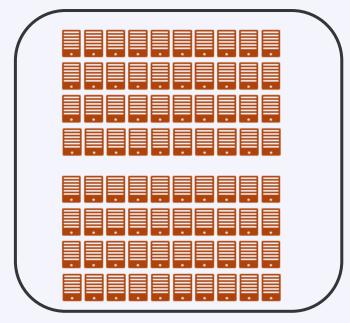
Genius

KU Leuven

wICE Overview



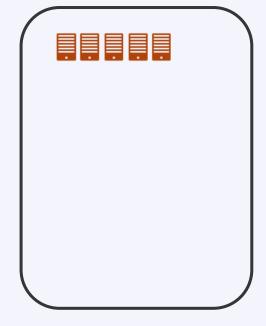
Compute nodes



Thin nodes

+ 172x Icelake 72c 256 GB

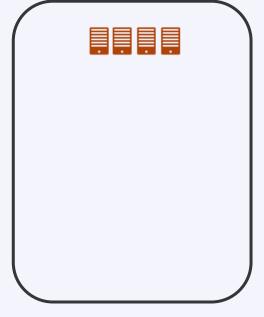
Large memory nodes



BigMem

+ 5x Icelake 72c 2 TB

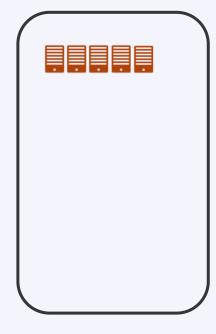
GPU nodes



GPUs

+ 4x Icelake 72c 512 GB 4x A100 80 GB

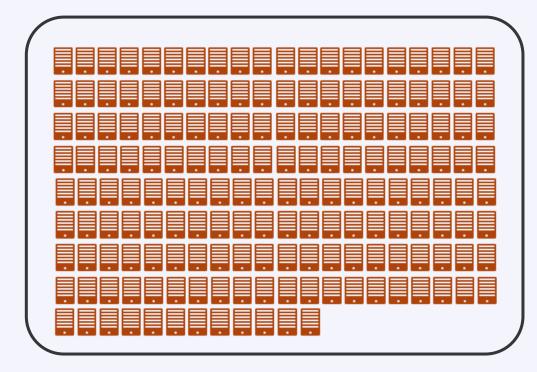
Interactive nodes



+ 5x Icelake 64c 512 GB 1x A100 80 GB

Tier-2 Overview wICE

Compute nodes



172x IceLake 72c 256 GB

Large memory nodes



5x IceLake 72c 2048 GB

Interactive nodes



5x IceLake 64c 512 GB 1 A100 80GB

GPU nodes



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



Tier-2 Cluster: wICE

Type of node	CPU type	Inter- connect	# cores	installed mem	local discs	# nodes
Icelake	Xeon 8358	IB HDR-100	72	256 GB	960 GB	172
Icelake large mem	Xeon 8358	IB HDR-100	72	2048 GB	960 GB	5
Icelake GPU	Xeon 8358 4xA100 SXM2 80GB	IB HDR-100	72	512 GB	960 GB	4
Icelake Interactive	Xeon 8358 1xA100 SXM2 80GB	IB HDR-100	64	512 GB	960 GB	5



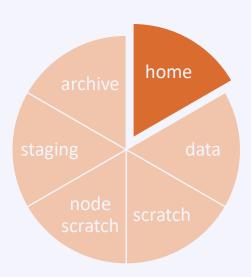
Overview of the storage infrastructure

- Your files are owned only by you.
 Other VSC users have no permission to read/write/execute your files (POSIX)
- 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
- More info on <u>ICTS Service Catalog</u> (EN/NL)

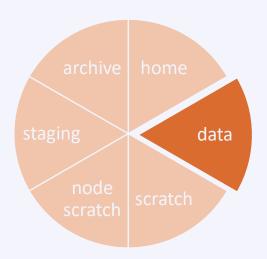
Request form for extra storage More information Do not use / tmp It is only 10 GB and is reserved for the OS and root processes Your application can crash if using /tmp You are automatically logged into your home folder upon login. Make sure you immediately go to your other storages, e.g. \$ cd \$VSC DATA Always check your storage balance using myquota command

Example

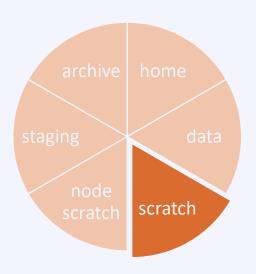
```
$ myquota
file system $VSC_HOME
    Blocks: 1479M of 3072M
    Files: 12934 of 100k
file system $VSC_DATA
    Blocks: 12G of 75G
    Files: 1043k of 10000k
file system $VSC_SCRATCH
    Blocks: 15M of 1.5T
```



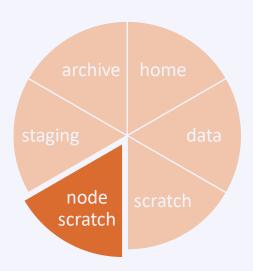
Storage	home folder
Env. Variable	\$VSC_HOME
Filesystem Type	NFS
Access	Global
Backup	Hourly, daily, weekly (until last month) inside the .snapshot folder.
Default Quota	3 GB
Extension	Not possible
Usage	Only storing SSH keys, config files
Remarks	Stay away from using itCan easily overflow:+ Your jobs may crash+ Login issues



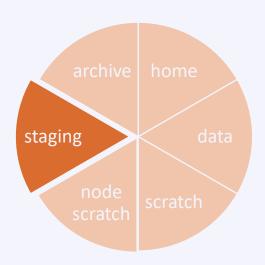
Storage	data folder
Env. Variable	\$VSC_DATA
Filesystem Type	NFS
Access	Global
Backup	Hourly, daily, weekly (until last month) inside the .snapshot folder.
Default Quota	75 GB
Extension	On purchase
Usage	Your data, code, software, results
Remarks	 Permanent storage for initial/final results Not optimal for intensive or parallel I/O



Storage	scratch folder	
Env. Variable	\$VSC_SCRATCH	
Filesystem Type	Lustre	
Access	Global	
Backup	delete after 30 days from last access	
Default Quota	500 GB	
Extension	Free	
Usage	Intensive, parallel I/O, temporary files	
Remarks	 Recommended storage for all jobs Copy scratch files to VSC_DATA or local storage after jobs are done Deleted files cannot be recovered 	

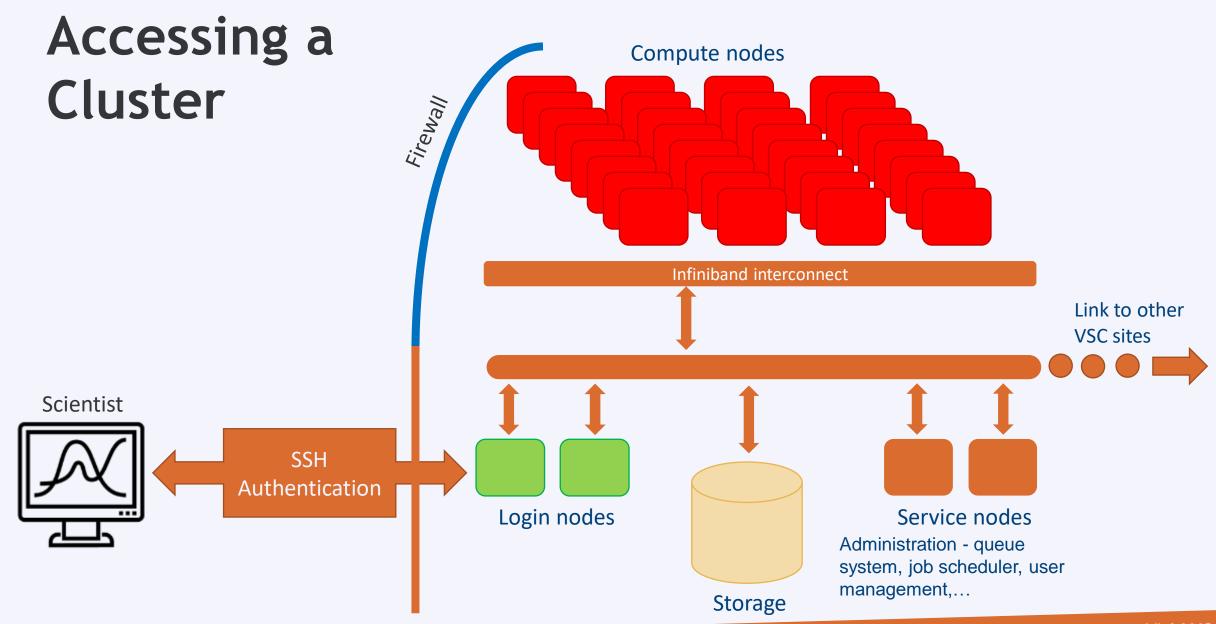


Storage	Node scratch folder
Env. Variable	\$VSC_SCRATCH_NODE
Filesystem Type	Lustre
Access	On compute node, only at runtime
Backup	None
Default Quota	591 GB
Extension	Read about beeOND
Usage	Temporary storage at runtime
Remarks	 Fastest I/O, attached to the node Is cleaned after job terminates Copy the data to your home, scratch, or staging before job ends

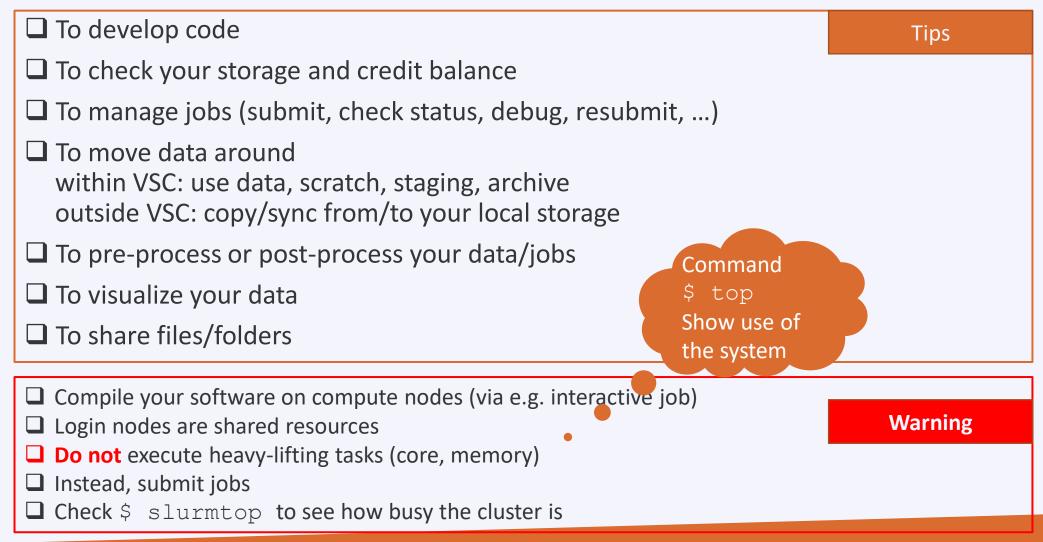


Storage	Staging folder
Path	/staging/leuven/stg_000XX
Filesystem Type	Lustre
Access	On demand, only Tier-2@KUL
Backup	None
Default Quota	None
Extension	On purchase, from 1 TB
Usage	Permanent; share with a group
Remarks	 Accessible from login/compute nodes Fast, parallel I/O





Using Login Nodes



Login Hosts on Different Machines/partitions

Windows: <u>PuTTY</u> or <u>MobaXterm</u> or NX

Linux/Mac: terminal or NX

• To login, you need an active VSC number and a hostname

\$ ssh -X vscXXXXX@<hostname>

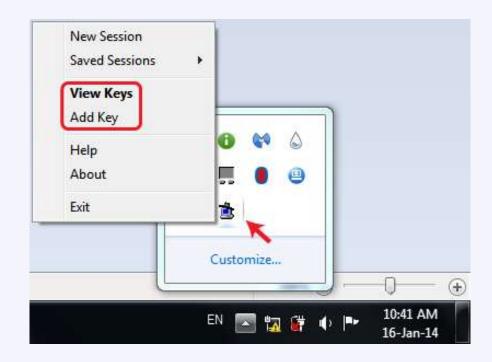
Cluster / Partition	<hostname></hostname>	Remark(s)
Genius	login.hpc.kuleuven.be	Recommended
	<pre>login{1,2}-tier2.hpc.kuleuven.be</pre>	No GPU
	<pre>login{3,4}-tier2.hpc.kuleuven.be</pre>	Nvidia Quadro P6000
wICE		Accessible only via Genius



Activate Your SSH Agent

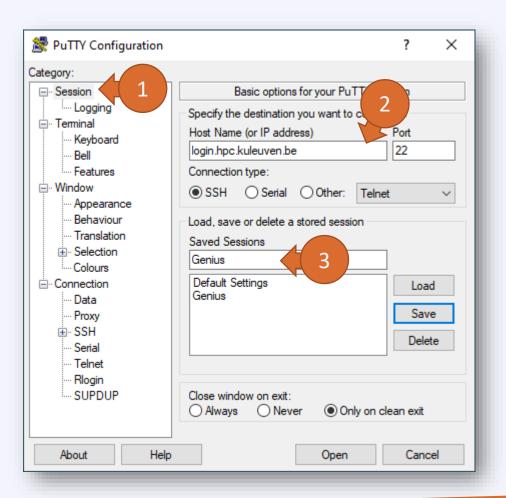
(Windows only)

- When you install PuTTY, the agent called Pageant is automatically installed
- Open Pageant; it hides inside your bottom-right tray
- Click on "Add Key" and brows to your private key(s) folder
- After choosing a key, you are asked for a passphrase
- If successful, agent always remembers your SSH keys and Certificates (Multi-Factor Authentication)

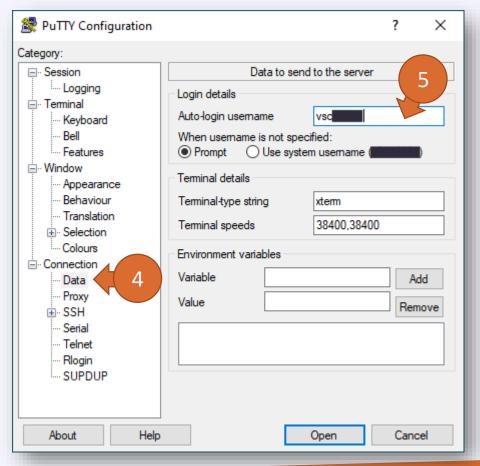


Setup PuTTY in 12 Clicks

(Windows only)

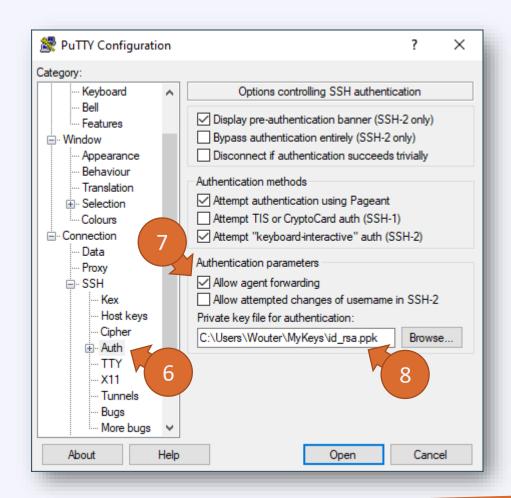


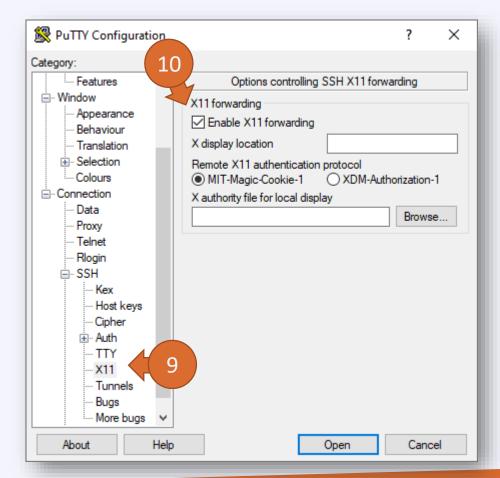
Host Name: login.hpc.kuleuven.be



Setup PuTTY in 12 Clicks

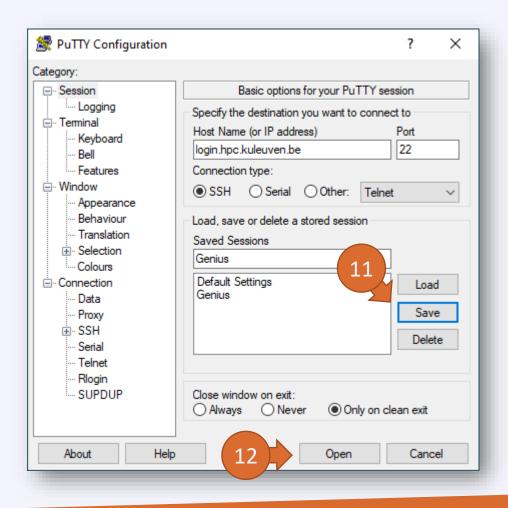
(Windows only)



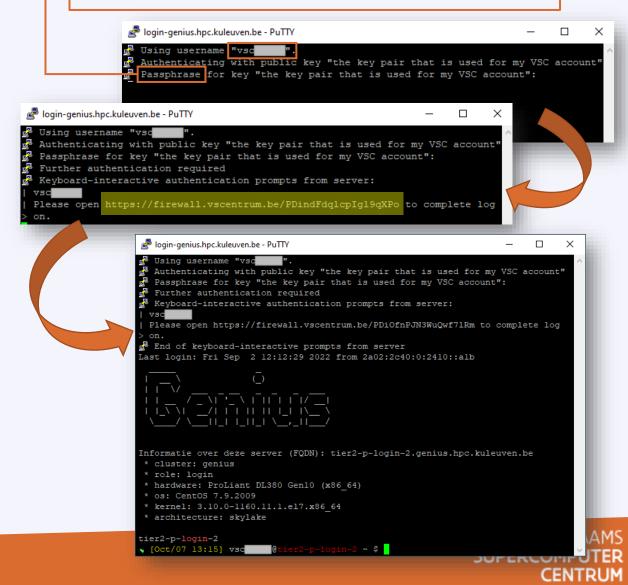


Setup PuTTY in 12 Clicks

(Windows only)



If PuTTY asks for **password**, exit immediately, and check the path to your private key. Else you will be blocked for 24h.



Connecting via Terminal

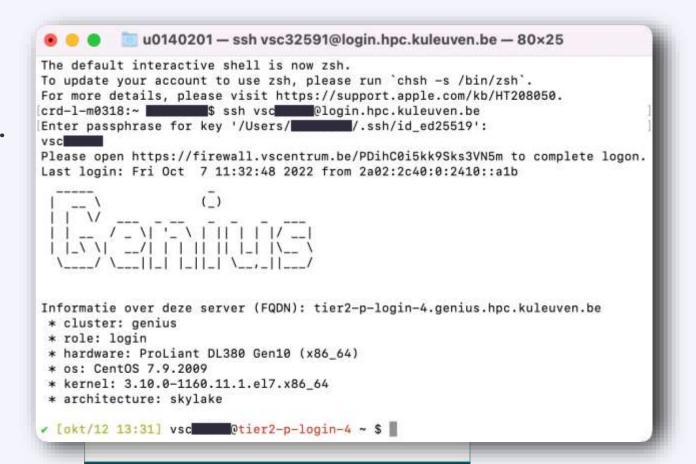
(Linux and Mac)

• Use ssh to connect:

• If key not found:

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

Host Name: login.hpc.kuleuven.be

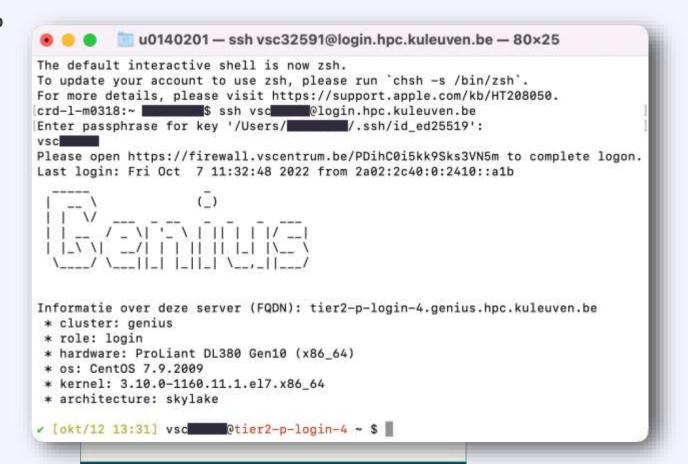


Connecting via Terminal

With SSH Agent (Linux and Mac)

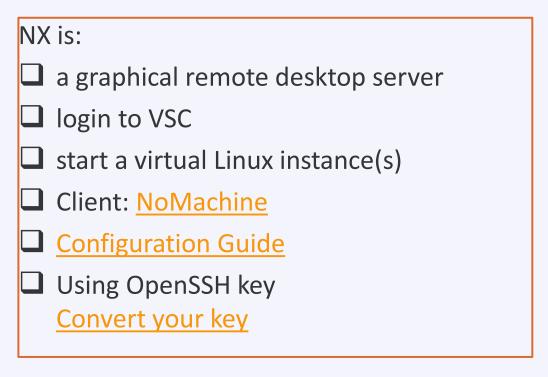
- Check your SSH Agent. Is your SSH key found? \$ ssh-add -1
- If your SSH Agent is not running: eval \$(ssh-agent)
- If your key is not found, add it to the Agent: \$ ssh-add </path/to/keyfile>
- Use ssh to connect: \$ ssh vscXXXXX@<hostname>

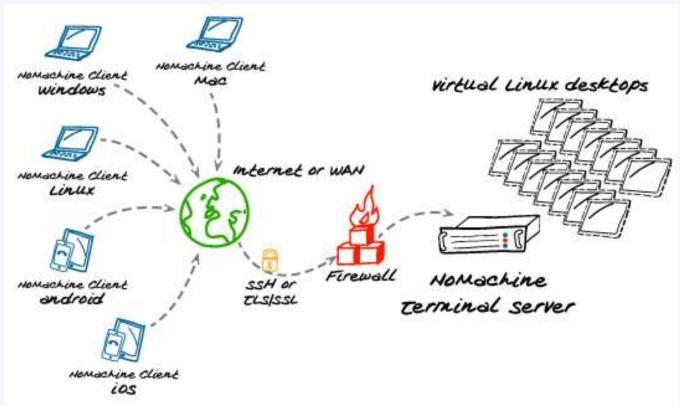
Host Name: login.hpc.kuleuven.be





NX – The Graphical Login

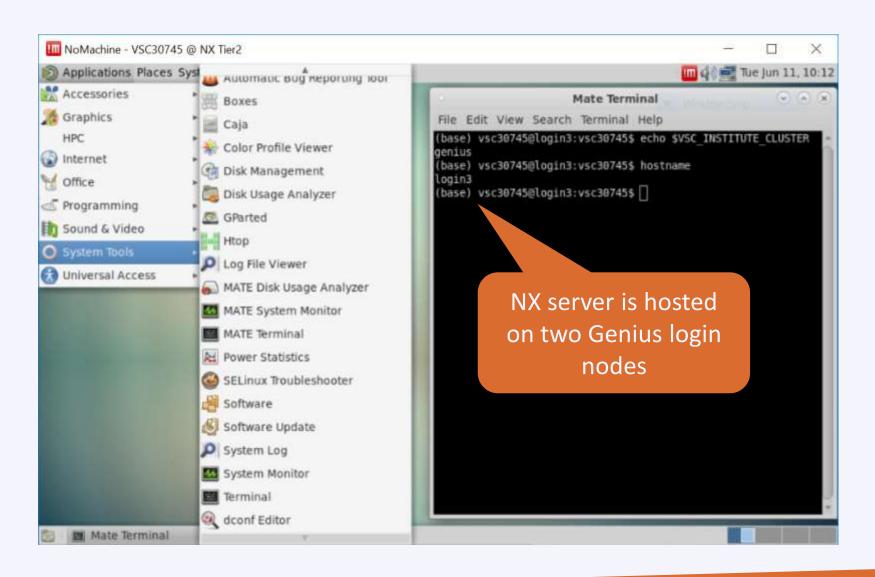


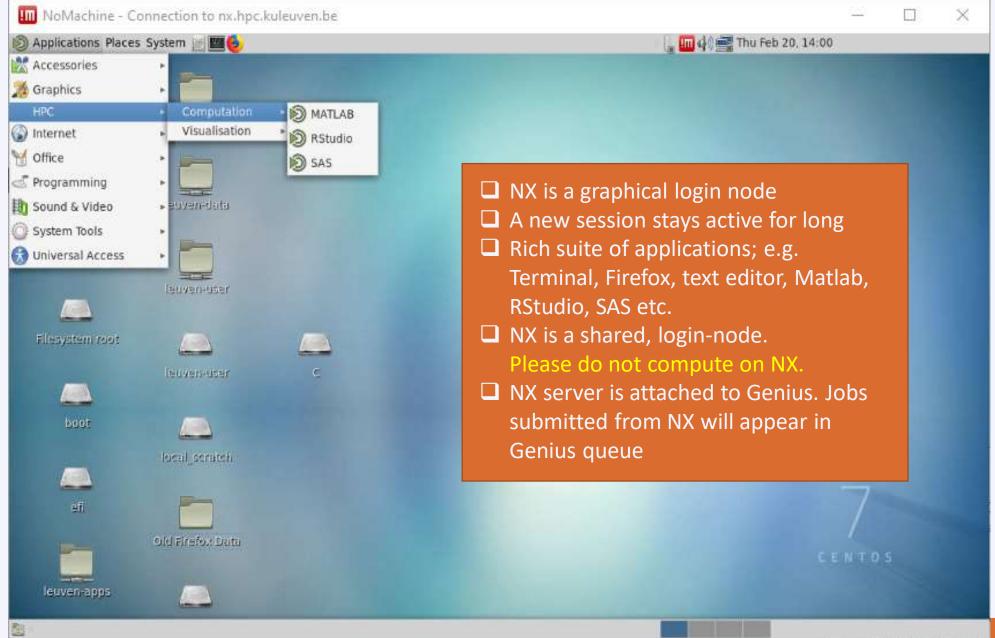


Advantages of NX

☐ Graphical login
A session stays alive/active even if you close your client (e.g. laptop)
Resume a session, and immediately keep working where you left off
☐ More interactive jobs
☐ Available apps:
Internet browser, terminal,
Text editor, PDF reader, Image viewer,
Matlab, RStudio, SAS
☐ Remark:
NX is shared among tens of users
Do not compute/test your code there
Instead, submit jobs from NX to compute nodes

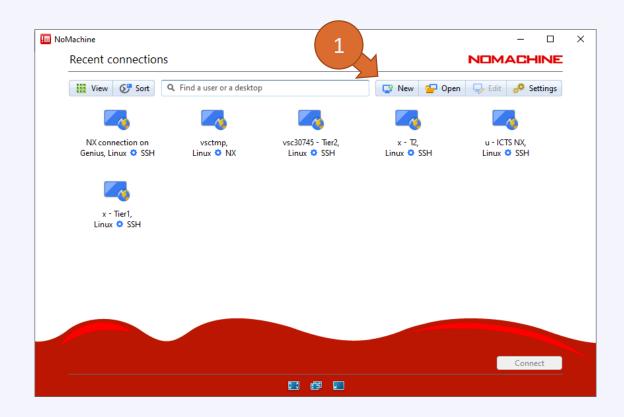
NX virtual desktop

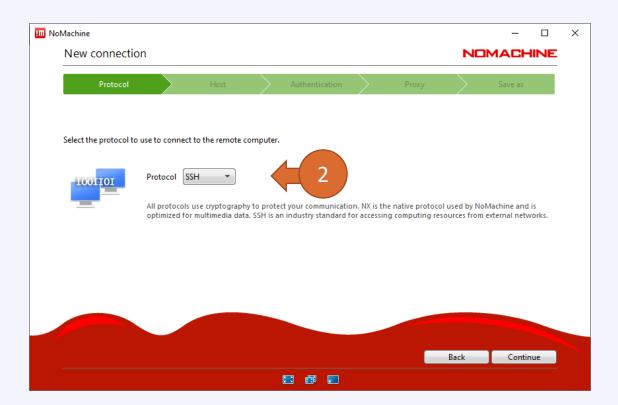


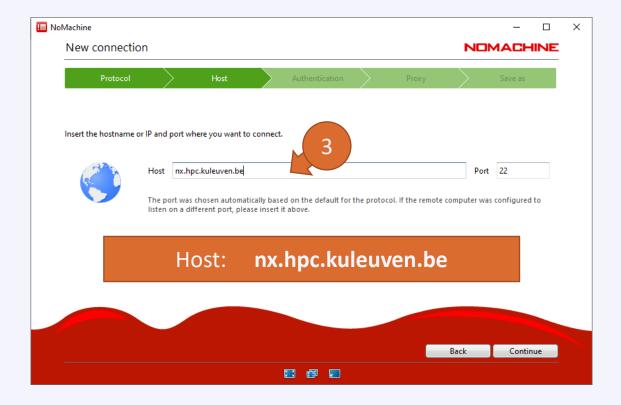


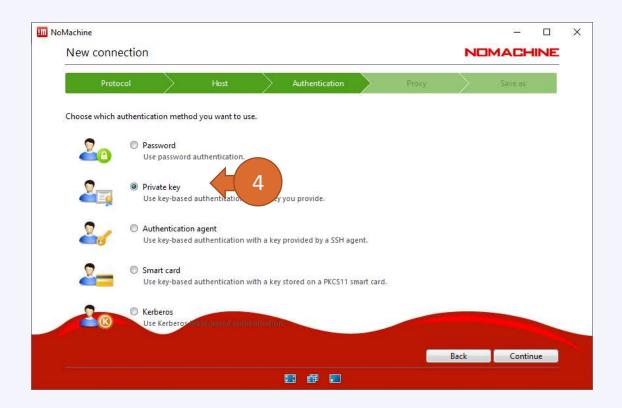
NX: available software

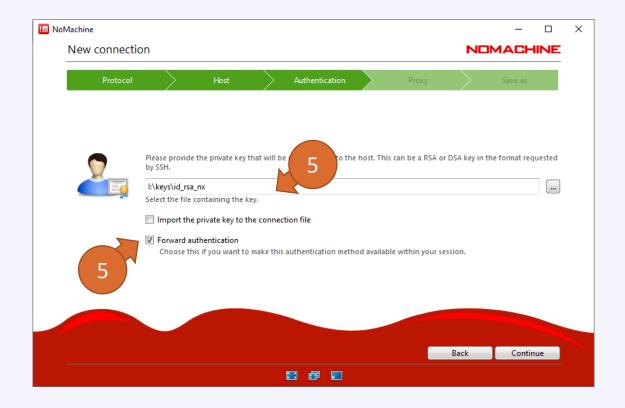
☐ Accesories: Gedit, Vi IMproved, Emacs (dummy version), Calculator
☐ Graphics : gThumb (picture viewer), Xpdf Viewer
☐ Internet: Firefox
☐ HPC: Computation: Matlab (2018a), RStudio, SAS
☐ Visualisation : Paraview, VisIt, VMD
☐ Programming : Meld Diff Viewer (visual diff and merge tool)
☐ System tools: File Browser, Terminal
☐ Additionally: Gnuplot (graphing utility), Filezilla (file transfer tool), Evince (PDF, PostScript, TIFF, XPS, DVI Viewer)
☐ Software launched though modules from Terminal.



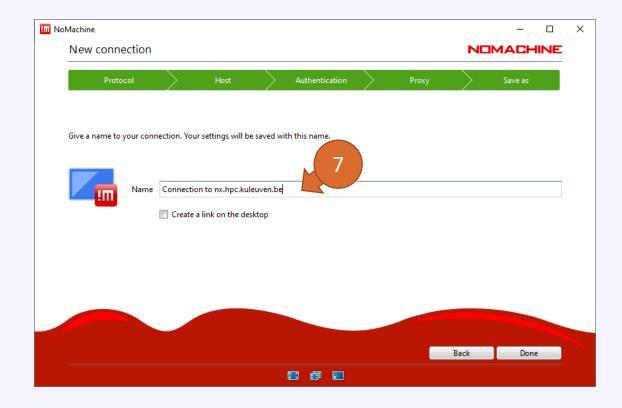




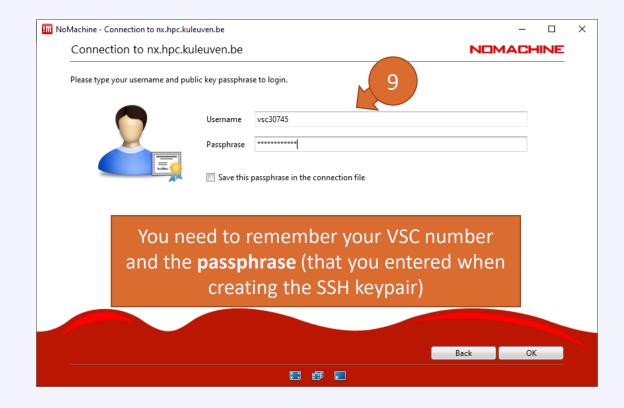


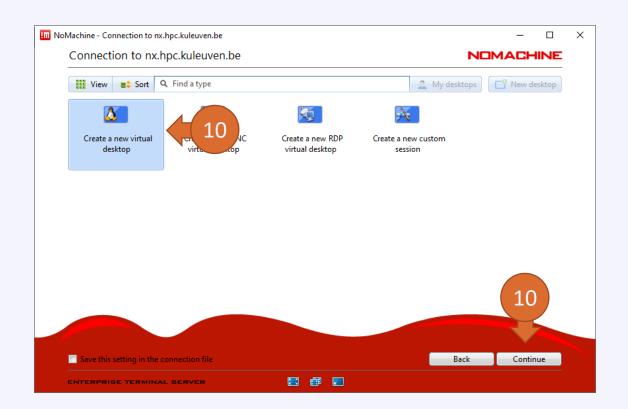






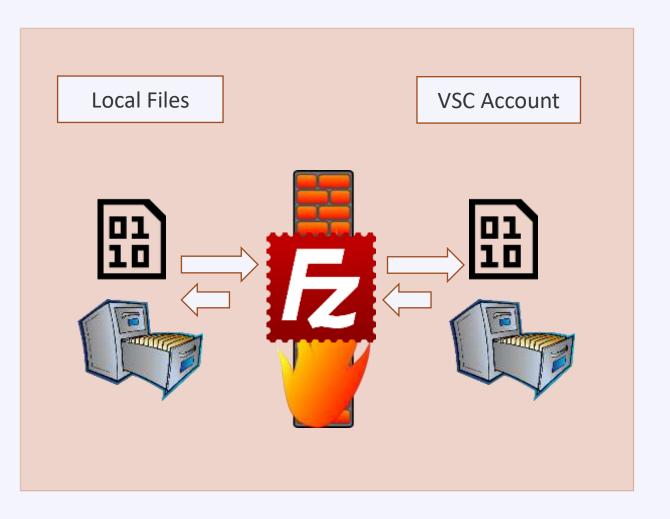






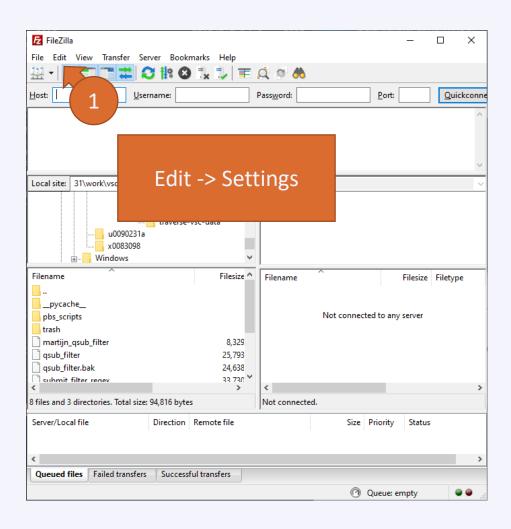


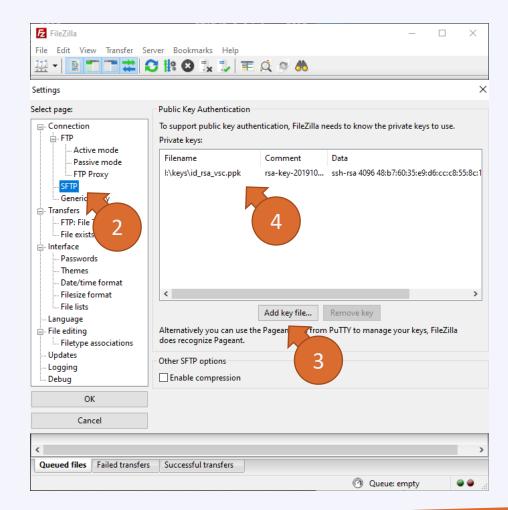
File transfer



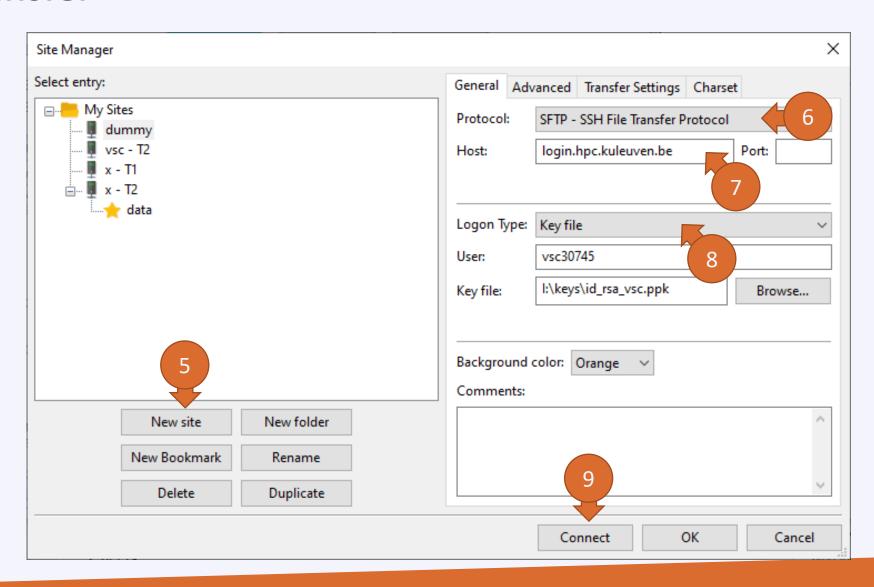
Application	OS
<u>FileZilla</u>	Windows, Linux, Mac
WinSCP	Windows
rsync, scp	Linux, Mac
Globus	Window, Linux, Mac

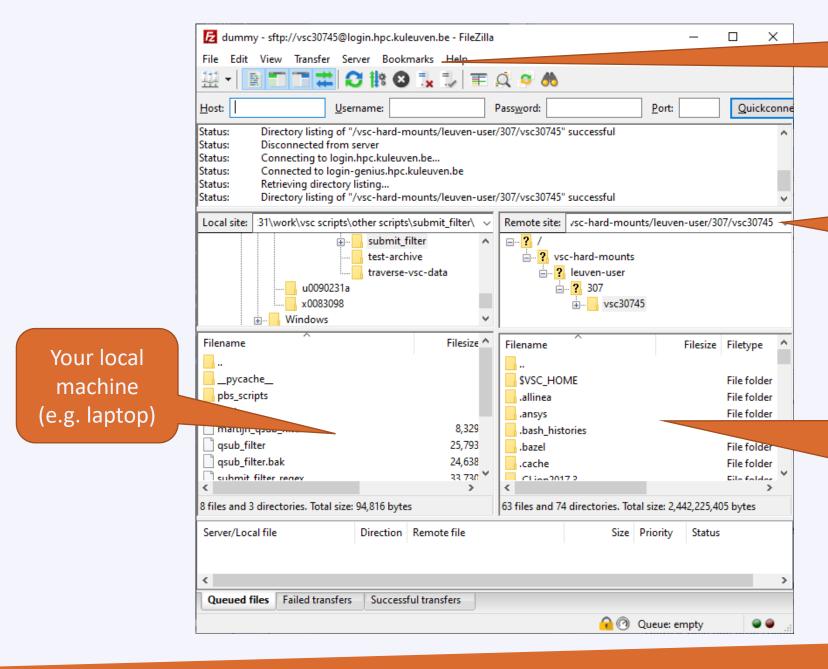
File transfer





File transfer





For convenience, you'd better bookmark your data and scratch folders!

Instead, go to your \$VSC_DATA folder, e.g. /data/leuven/399/vsc39934

Your VSC storage (e.g. \$VSC_DATA, \$VSC_SCRATCH).

Note: by default you arrive at your own \$VSC_HOME. Copy nothing there!



Open OnDemand

- Access wICE via web browser
- o https://ondemand.hpc.kuleuven.be
- Login via KULeuven MFA
- File browser
- Interactive apps + integrated shell
- Create, start and monitor jobs
- Develop, compile and test your code
- VSCode editor
- Jupyter Lab, RStudio and Tensorboard



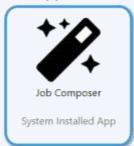
KU LEUVEN

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

Pinned Apps A featured subset of all available apps





















Open OnDemand

- E.g. to start a Jupyter Notebook
- Pick a valid Slurm credit account
- Default partition: batch interactive: Max 8 cores, 1 GPU, 16 hr
- Default resources:1 core, 1 hour, 3400MB RAM, no GPU
- (At the moment) you cannot use scratch and staging

Home / My Interactive Sessions / Jupyter Lab

Interactive Apps	Jupyter Lab	
Servers	This app will launch a Jupyter Lab server on one or more	
● Interactive Shell	nodes.	
	Account	
Jupyter Lab	lpt2_sysadmin	
RStudio Server	Partition	
⋄ Tensorboard	interactive	
≾ code-server	batch(_long) or bigmem or interactive or gpu or dedicated	
Work in progress	Number of hours	
ParaViewWeb - Work in progress	3	
	Number of cores	
₩ cryosparc	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



Software: Available Modules

- OS: Linux CentOS 7.x (Genius) and Rocky Linux 8.x (wICE)
- Toolchains (Genius): Intel 2018a (icc, icpc, ifort; Intel MPI; Intel MKL)
 FOSS 2018a (gcc, g++, gfortran; OpenMPI; ScaLAPACK, OpenBLAS, FFTW)
- wICE: default is 2021a
- Note: Never mix FOSS and Intel compilers (gives dependency conflict)

Command	Remark
module av	List all installed modules
module av Python	List all Python-related modules
<pre>module use /apps/leuven/icelake/2022b/modules/all</pre>	Expose other modules from non-default toolchains, by including them in the search
module spider Python	Get more info
module load Python/3.6.4-intel-2018a	Load a specific module
module list	List all loaded modules and their dependencies
module unload Python/3.6.4-intel-2018a	Unload a module (but dependencies still stay)
module purge	Remove all modules from your work session

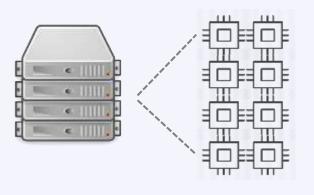
Software: Your Specific Needs

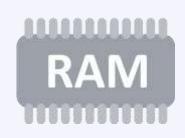
- You can always install your desired software in your \$VSC_DATA
 Use Intel or FOSS toolchains
- Compile your code on a compute node (with interactive job)
- If you cannot, ask us for help
- Python/R packages for AI and ML must be installed by the users themselves
- Read more about Python Package Management
- Read more about <u>R Package Management</u>



Resource Glossary

Nodes: how many compute servers to request?
 Cores: how many cores per node to use?
 Memory requirement: how much memory each core needs?
 Partition: gpu, bigmem, superdome, amd
 Walltime: how long to use resources?
 Storage: how much storage (data, scratch, etc) the job needs?
 Credits: how many compute credits will be consumed?







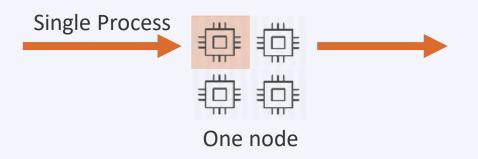






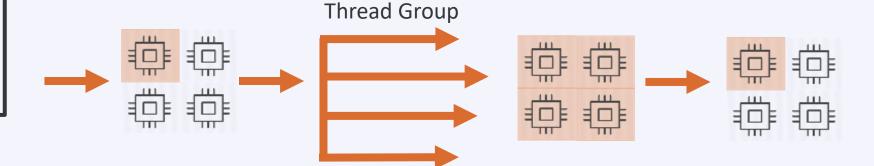
Serial Application

(1 process on 1 core)



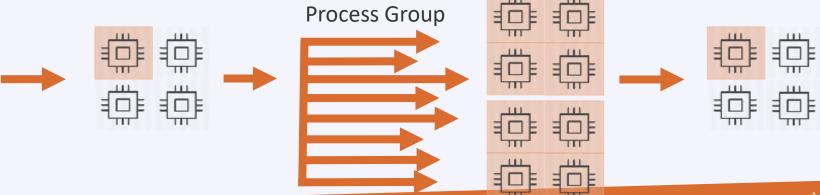
Multi-Core Appl.

(N threads on N cores from **1 node**)



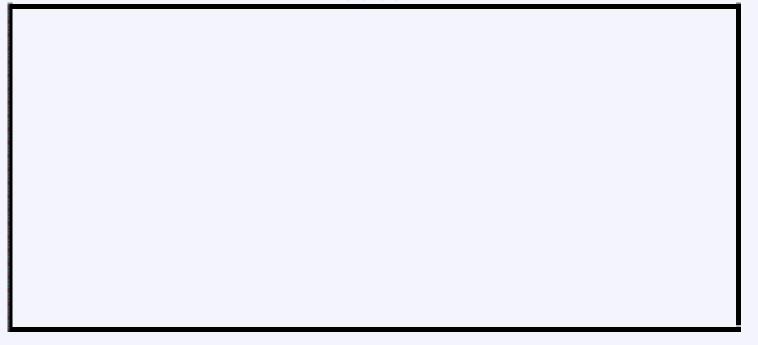
Distributed Appl.

(many processes on multiple cores/nodes)



Backfill



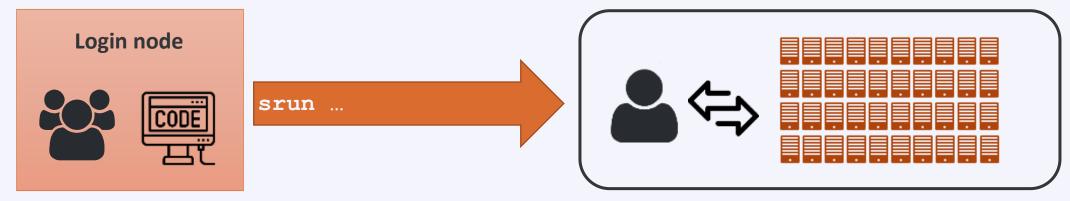


Job queue



Interactive Job

Compute nodes



Batch Job

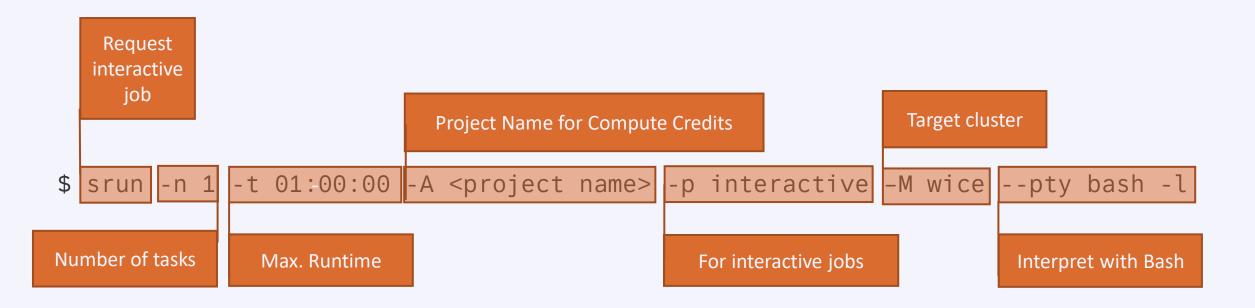


sbatch script.slurm

Compute nodes



Interactive Job on wICE



Remark

- Implicit defaults for an interactive job on wICE are: -n 1, -t 01:00:00, --mem-per-cpu=2000M

Jobs: Slurm submit options

nodes=Xntasks-per-node=Yntasks=X*Yntasks=Xcpus-per-task=Y	or or (for Hybrid MPI/OpenMP only)
partition=gpu	Default: "batch" partition
mem-per-cpu=XG	Default: average memory/core
time=hh:mm:ss	
-J jobname	

Remarks

-o <file template> output -e <file template> Default: same as stdout

--mail-type=FAIL, BEGIN, END

-A account

SLURM

--export=ALL, key=value

--mail-user=<mailaddress>

Additional variables to pass to job Mandatory (credits)

Argument Shorthands

Some of the sbatch, srun and salloc command line arguments have shorthands

Shorthand	Full Argument	Meaning
-A	account	Slurm account name
-a	array	Job array range
- M	cluster	Machine or cluster name
-c	cpus-per-task	Num cores per task (default=1)
-d	dependency	Job dependency (after, afterok, afterany,)
-I	input	STDIN filename
-e	error	STDERR filename
-0	output	STDOUT filename
-G	gpus	Num GPUs per job
-t	time	Maximum walltime
-N	nodes	Minimum num nodes
-n	ntasks	Maximum num tasks
-p	partition	Partition name

Jobs: Defaults

If no explicit resources are given your job will start

- with 1 node 1 core,
- with walltime of 1hr
- o in batch partition
- Soon the Slurm accounts will be available also on Genius (with the same account name)

Interactive Jobs

- Interactive job: 1 core for 1 hour (default)

```
$ srun -M wice -A lp hpcinfo --pty /bin/bash -l
```

Interactive job with X-forwarding

```
$ srun -M wice -A lp hpcinfo -x11 --pty /bin/bash -l
```

- Request fraction of a node from interactive partition

```
$ srun -M wice -A lp hpcinfo -N 1 -n 4 -p interactive --pty /bin/bash -l
```

- Request a GPU accelerator

```
$ srun -M wice -A lp hpcinfo -N1 -n 18 -p gpu -G 1 --pty /bin/bash -l
```

```
Example: Slurm Job Script
                                Shebang
#!/bin/bash -1
#SBATCH --cluster wice
#SBATCH --account lp hpcinfo
#SBATCH --nodes 1
#SBATCH --ntasks 4
#SBATCH --mem-per-cpu 4G
                                                         Resource List
#SBATCH --job-name hpc workflow
#SBATCH --mail-type BEGIN, END, FAIL
#SBATCH --mail-user my.name@kuleuven.be
Module --force purge
                                                    Module load(s)
module load intel/2021a
module load Python/3.9.5-GCCcore-10.3.0
which python3
cd $VSC SCRATCH/projects/simulations
                                            Move data
cp -r $VSC DATA/input data .
                                                            Execure commands
python modelling.py
cp -r output data $VSC DATA
                                            Move data
rm -rf ./input data ./output data
```

```
#!/bin/bash -1
```

Command Line

\$ sbatch simulation.slurm

JobID

Submitted batch job 60042478 on cluster wice

stderr, stdout

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

Submit the job to the batch serverReceive a unique JobID

Error and output files

Output File

STDOUT + STDERR

- STDERR and STDOUT are by default redirected to a single file:
- slurm-<JobID>.out
- Contains job info, all errors and warnings, and printouts
- Always study it
- Address all warnings and errors (if you can)
- Typical error examples ...

```
$ 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 File

Always created

- slurm-<job id>.out
- Contains all standard output and error (instead of screen)
- Always study it
- 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: vscXXXXX
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/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 ordirecto
ry
```

Resource Summary

Stdout

Hello World

Inspecting Jobs

Example

```
$ scontrol show job -M wice 60049330
JobId=60049330 JobName=f70.slurm
   UserId=vsc3....(253....) GroupId=vsc3....(253....)
  Account=1p metacommunity models QOS=1p metacommunity models
   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 is my job in pending/hold state?

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

Other Partitions on wICE

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
```

Managing & Monitoring Jobs

Command	Purpose
sbatch	Submit a batch job
srun	Submit an interactive job
scancelcluster=wice <jobid></jobid>	Cancel a specific pending or running job
<pre>scontrol show jobcluster=wice <jobid> slurm_jobinfo <jobid></jobid></jobid></pre>	Detailed job info (very useful to diagnose issues)
squeuecluster=wice -long	Status of all recent jobs
squeuecluster=wice -start	Give a rough estimate of start time
sinfocluster=wice	Info about the state of available partitions and nodes
<pre>sacctcluster=wicebatchjob <jobid></jobid></pre>	Show minimal info about a queue or partition
slurmtopcluster=wice	Overview of the cluster

Interactive Partition

- Accessible via command line and Open OnDemand
- To quickly compile, test, debug your (parallel) application
- To pre-/post-process your data and make visualizations
- Short queue time
- 5 dedicated nodes on wICE
- 64 cores per node, 1 GPU (=7 GPU instances), 512 GB memory
- Max resource per user: 8 cores, 1 GPU instance, 16 hour walltime

Interactive Job

srun -n 1 -p interactive -t 16:00:00 -G 1 --pty bash -l

Singularity Containers

- What?
 - + Self-contained OS & software & data
- Why?
 - + fully resolved dependency chains
 - + portable workflow
- How?
 - + You create the image
 - + Run it on Genius
 - + MPI/OpenMP is supported

Slurm Script

```
#!/bin/bash -l

#SBATCH -t 30:00

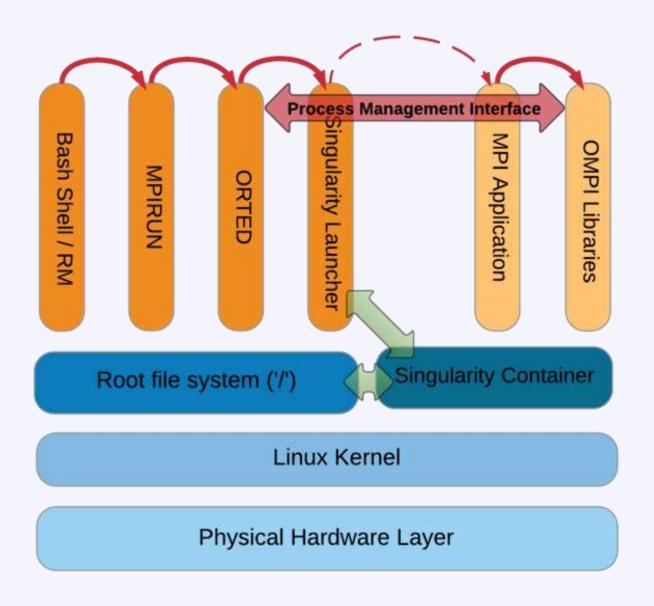
#SBATCH -N 2

#SBATCH -n 72

#SBATCH -M wice

#SBATCH -A lp_hpcinfo

singularity run Project.simg ./model.exe
```



Credits

Credits card concept:

- Pre-authorization: holding the balance as unavailable until the merchant clears the transaction
- Balance to be held as unavailable: based on requested resourced (walltime, nodes)
- Actual charge based on what was really used: used walltime (you pay only what you use, e.g. when job crashes)
- See output file

How to check the rates? Check service catalogue

How to check available credits?

```
$ sam-balance
```

Did the new credits arrive?

\$ sam-list-allocations

Credit Pricing

- o You pay as you go!
- For academic projects:1000k credits = 3.5 EUR
- Credits needed for a job:

#credits (k) = Walltime (hr) × #nodes × Factor

 For shared nodes, you pay a fraction of the costs, based on the ppn specified

Cluster / Partition	Credits/hr
Genius Cascadelake	11.3
Genius Cascadelake 8 GPUs	40
Genius Skylake 4 GPUs	20
Genius Skylake BigMem	12
Genius Skylake	10
Genius / Superdome	10
Genius / AMD nodes	10
wICE / Thinnode	10,99
wICE Bigmem	19
wICE / 1 GPU	11,25
wICE 4GPUs	45
wICE Interactive	-

How to Manage Credits?

Command	Purpose
sam-balance	List active projects and available credits
sam-list-allocations	List the validity dates of different projects/allocations
sam-list-usagerecords	List current charge rates for different nodes
sam-statement -A <account-name></account-name>	Job statements from an account



Conda on wice

https://docs.vscentrum.be/en/latest/software/python_package management.html?highlight=iniconda#install-pythonpackages-using-conda

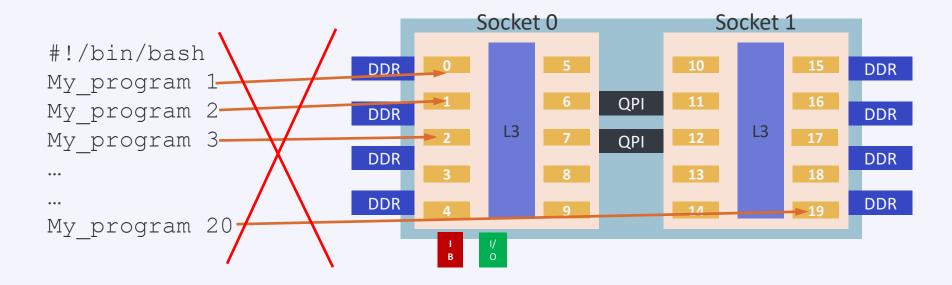
- Good practice would be 2 separate paths of conda installations for each cluster.
- ☑ Beginning of the slurm script matters. Use #!/bin/bash -l not #!/bin/bash
 to load the env properly.
- Universal .bashrc

```
case ${VSC_INSTITUTE_CLUSTER} in
    genius)
    export PATH="${VSC_DATA}/miniconda3/bin:${PATH}"
    ;;
    wice)
    export PATH="${VSC_DATA}/miniconda3-wice/bin:${PATH}"
    ;;
```



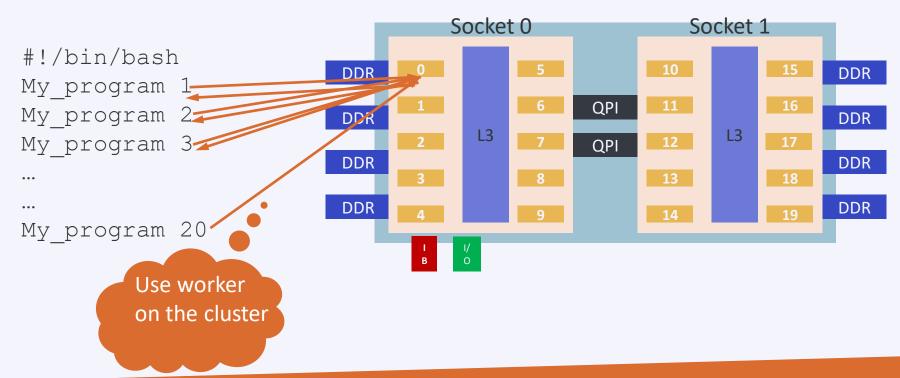
Sequential...

- A shell program consists of a sequential list of commands
- -> bash script on HPC cluster will run sequentially



Sequential...

- A shell program consists of a sequential list of commands
- -> bash script on HPC cluster will run sequentially



Parallel Computing

- Serial:
 - one program, on one core
- 'Embarrassingly parallel' problems:
 - lots of runs of one program, with different parameters
- Problems that require 'real' parallel algorithms
 - OpenMP
 - MPI: Message Passing Interface



Use case: parameter exploration

temperature	pressure	humidity
293.0	1.0e05	87
		•••
313.0	1.3e05	75

```
Many single core computations
#!/bin/bash -1
                                                                   _____ob_030.slurm
    #!/bin/bash -
                                                                        job 600.slurm
       #!/bin/bash -1
#SB
        #SBATCH --cluster=wice
#SB
       #SBATCH --nodes=1
    #SB
       #SBATCH --ntasks-per-node=1
cd
        #SBATCH --time=10:00
wea
        #SBATCH -A lp_myproject
    cd
٠.,
    wea
        cd $SLURM SUBMIT DIR
        weather -p 1.0e05 -t 293.0 -h 87
```

Solution: worker with -data

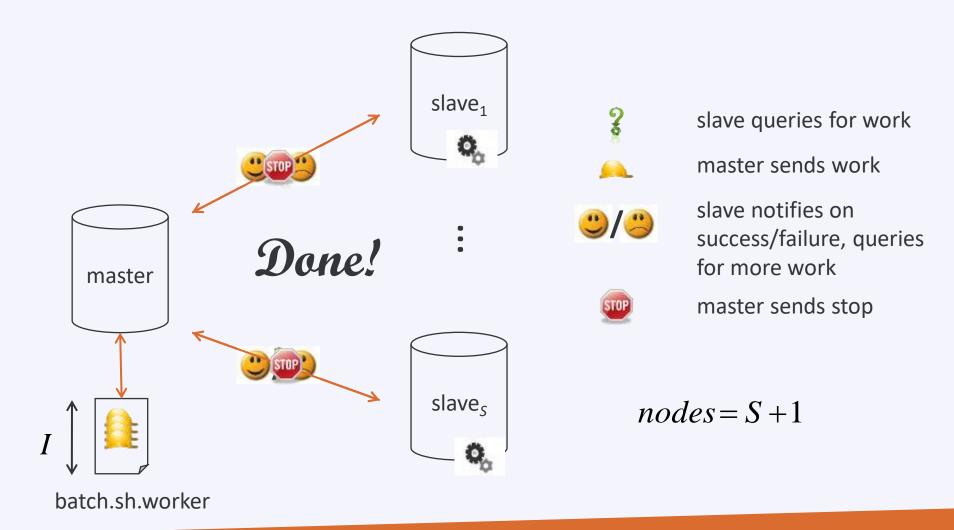
```
temperature
                                                    humidity
                                   pressure
                                                               data.csv
                  293.0
                                   1.0e05
                                                    87
#!/bin/bash -1
                                                            job.slurm
#SBATCH --cluster=wice
#SBATCH --nodes=2
#SBATCH --ntasks-per-node=72
#SBATCH --time=01:00:00
#SBATCH -A lp hpcinfo
cd $SLURM SUBMIT DIR
weather -p $pressure -t $temperature -h $humidity
#From wICE interactive session:
$ module --force purge
$ module use /apps/leuven/icelake/2021a/modules/all
$ module load worker/1.6.12-foss-2021a-wice
$ wsub -data data.csv -batch job.slurm
```

Data exploration: steps

- Write PBS script with parameters
- Create Excel sheet with data
 - Convert to CSV format
- Submit with wsub
 - walltime is time to complete all work items

$$wall time_{job} \ge \frac{N \cdot wall time_{work item}}{nodes \cdot ppn}$$

worker processing: informally



How to use worker well?

- Many work items, i.e., #work items/#proc >> 1
- time(work item) > 1 minute
- Work item is not multithreaded
- Work item is multithreaded
 - will work, but user must be careful to request the right resources
 - Use -threaded <n> flag with wsub
- There be dragons: licensing!

worker & multithreading

- Some software uses multithreading automatically, e.g.,
 - R
 - Matlab
- Will use as many threads as there are cores, regardless of system load
 - 36 cores/node
 - 36 work items/node

Oversubscription: very inefficient!!!

Controlling number of threads

• R, most of the time: OMP_NUM_THREADS=1

- Matlab
 - Use maxNumCompThreads (1) function call
 - Use compiler flag: mcc -singleCompThread ...

```
#!/bin/bash -l
#SBATCH -account=lp_myproject
#SBATCH --cluster=wice
#SBATCH --time=1:00:00
#SBATCH --nodes=2
#SBATCH --ntasks-per-node=72

module load R
cd $SLURM_SUBMIT_DIR

export OMP_NUM_THREADS=1
Rscript program $a $b
```

Help on worker

- See documentation http://worker.readthedocs.io/
- Each command has help, use -h

- All resources have to be specified inside slurm script
- wresume does not work yet. Only wsub is so far supported
- -master flag does not work yet

From wICE (Interactive) Node

```
$ module av worker
----- /apps/leuven/icelake/2021a/modules/all -----
worker/1.6.12-foss-2021a
```

From Genius Login Node

Let us use one of the worker examples: https://github.com/gjbex/worker/tree/development_slurm/examples/bash_example

```
alp.slurm
                                  alpha.csv
#!/bin/bash -1
#SBATCH --nodes=1
                                              letter, counter
#SBATCH --ntasks-per-node 27
                                              a, 1
#SBATCH --time=00:30:00
                                             b, 2
#SBATCH -A lpt2 sysadmin
                                              c, 3
#SBATCH --cluster=wice
                                             d, 4
cd $SLURM SUBMIT DIR
                                              e,5
                                              f,6
alphabet.sh $letter $counter
```

From Genius

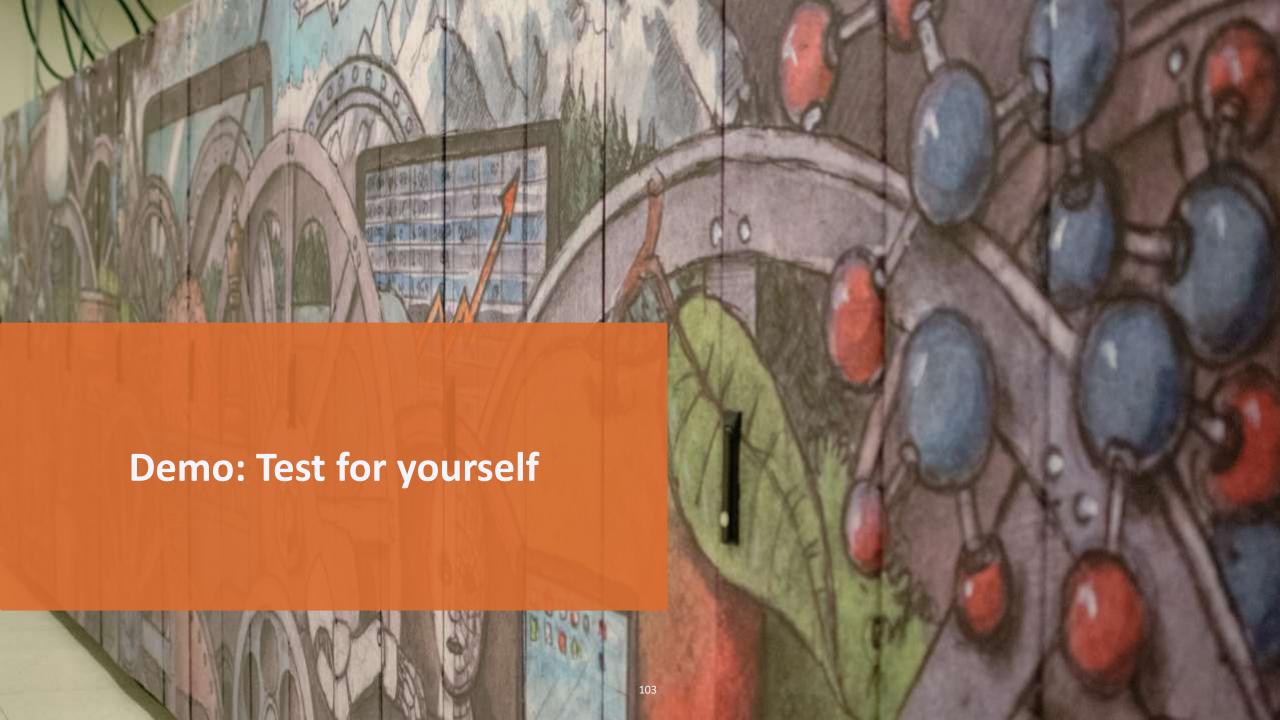
```
#!/bin/bash -l
module purge
module use /apps/leuven/skylake/2021a/modules/all
module load worker/1.6.12-foss-2021a-wice
wsub -batch alp.slurm -data alpha.csv
```

```
total number of work items: 26
Submitted batch job 60008899 on cluster wice
```

From interactive job on wICE

```
#!/bin/bash -l
module --force purge
module use /apps/leuven/icelake/2021a/modules/all
module load worker/1.6.12-foss-2021a
wsub -batch alp.slurm -data alpha.csv
```

```
total number of work items: 26
Submitted batch job 60008889 on cluster wice
```



demo/test yourself

- ♥Request membership to lp_hpcinfo group (account.vscentrum.be)
- **⊘**Login with putty
- ♥Filetransfer with Filezilla
- **⊘**Login with NX
- ♥Check disk quota
- Check the credits
- ♥Check/load/list/unload/purge module

demo/test yourself

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

demo - monitoring

- Submit an interactive job
 Run your program on a compute node
 Open a new terminal and ssh to a compute node
 Check the resources usage (top)
- Submit a batch GPU job

 While the job is running get the information about the node slurm_jobinfo...

 and check usage of resources on the node ssh, top, nvidia-smi

demo – conda installation

- **♥** Install miniconda in your \$VSC_DATA directory
- \$ wget https://repo.continuum.io/miniconda/Miniconda3-latest-Linux-x86_64.sh
- \$ bash Miniconda3-latest-Linux-x86_64.sh -b -p \$VSC_DATA/miniconda3
 - Add a PATH to conda:
- \$ export PATH="\${VSC DATA}/miniconda3/bin:\${PATH}"
 - Check if conda in added to your \$PATH (\$ which conda)
 - Add it to \$PATH in your .bashrc
- \$ echo 'export PATH="\${VSC DATA}/miniconda3/bin:\${PATH}" ' >> .bashrc

demo – conda usage

- Create a conda environment including Jupyter
- \$ conda create -n science jupyter numpy scipy
- Activate this environment
- \$ source activate science
- Add matplotlib package to this environment
- \$ conda install matplotlib
- Return to original environment
- \$ conda deactivate

demo – worker

- ♥ Copy intro training files (/apps/leuven/training/worker/) to your \$VSC DATA
- **⊘** Go to exercise1 directory
- Submit worker job
- Check the output file

Command	Purpose
\$ sbatch	Submit a batch job
\$ srun	Submit an interactive job
<pre>\$ scancelcluster=wice <jobid></jobid></pre>	Cancel a specific pending or running job
<pre>\$ scontrol show jobcluster=wice <jobid> \$ slurm_jobinfo <jobid></jobid></jobid></pre>	Detailed job info (very useful to diagnose issues)
<pre>\$ squeuecluster=wicelong</pre>	Status of all recent jobs
<pre>\$ squeuecluster=wicestart</pre>	Give a rough estimate of start time
<pre>\$ sinfocluster=wice</pre>	Info about the state of available partitions and nodes
<pre>\$ sacctcluster=wicebatchjob <jobid></jobid></pre>	Show minimal info about a queue or partition (-p)
\$ slurmtop	Overview of the cluster
<pre>\$ scontrolcluster=wice show node <hostname></hostname></pre>	Get detailed information about the status of a node
\$ sam-balance	Overview of all your available credit projects
<pre>\$ sam-list-allocations</pre>	Detailed overview of your credit allocation history

VSC

Questions

Helpdesk:

hpcinfo@kuleuven.be or https://admin.kuleuven.be/icts/HPCinfo form/HPC-info-formulier

VSC web site:

http://www.vscentrum.be/

VSC documentation: https://docs.vscentrum.be/en/latest/

VSC agenda: training sessions, events

Systems status page:

http://status.vscentrum.be

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