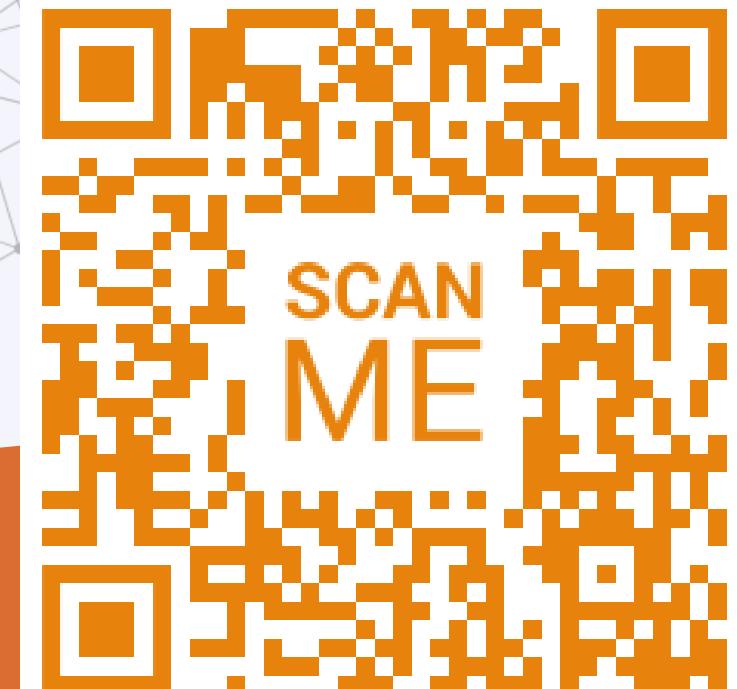




Vlaanderen
is supercomputing

Start with vsc

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



Basics

✓ Request membership to lp_hpcinfo group (account.vscentrum.be)

✓ Login via OpenOnDemand: www.ondemand.hpc.kuleuven.be

Features: File transfer, File editor, interactive apps, Jobs' overview

✓ Check disk quota:

```
$ myquota
```

✓ Check the credits

```
$ sam-balance
```

and

```
$ sam-list-allocations
```

✓ Software modules:

```
$ module {avail|list|load|unload|purge}
```

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

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 --cluster=wice <JobID> \$ slurm_jobinfo <JobID>	Detailed job info (very useful to diagnose issues)
\$ squeue --cluster=wice --long	Status of all recent jobs
\$ squeue --cluster=wice --start	Give a rough estimate of start time
\$ sinfo --cluster=wice	Info about the state of available partitions and nodes
\$ sacct --cluster=wice --batch --job <JobID>	Show minimal info about a queue or partition (-p)
\$ slurmtop \$ scontrol --cluster=wice show node <hostname>	Overview of the cluster 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

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 resource usage (`htop`)

- ✓ 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

- ✓ Start an interactive job on Genius or wICE

```
srun -M genius -A <account> -n 1 --pty /bin/bash -l
```

- ✓ Install miniconda in your \$VSC_DATA directory

- \$ wget https://repo.continuum.io/miniconda/Miniconda3-latest-Linux-x86_64.sh
- \$ DIR_INSTALL=\$VSC_DATA/miniconda3_\${VSC_INSTITUTE_CLUSTER}
- \$ bash Miniconda3-latest-Linux-x86_64.sh -b -p \${DIR_INSTALL}

- ✓ Add a PATH to conda:

- \$ export PATH="\${VSC_DATA}/miniconda3_\${VSC_INSTITUTE_CLUSTER}/bin:\${PATH}"

- ✓ Check if conda is 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

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/>

VSC agenda: training sessions, events (VSC and KU Leuven user days)

Systems status page:

<https://status.vscentrum.be>

*Stay Connected
to vsc*

