



ilifu Online Training

Session 1: Introduction to slurm

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IDIA



Job scheduling & cluster management tool

- Framework: Login node; Compute nodes; Scheduler; and Accounting
- **Login node**
 - Accessed via ssh (`$ ssh <username>@slurm.ilifu.ac.za`)
 - Submit jobs and manage work directories
- **Compute nodes**
 - Where your analysis / simulation runs (in a slurm job)
 - Software available via singularity containers or modules
- **Scheduler and Accounting Database**
 - Manage Jobs, Partitions and Queues
 - Accounting

Login node

Run SLURM & bash commands
cd, mkdir, ls, etc

Jupyter/Dev. node

Development space
New code / workflows / routines
Debugging / testing software

Main partition

Stable, computationally
heavy processing

HighMem/GPU

For single-high memory jobs
that can't be split into
multiple jobs for MPI

Main, Jupyter, Devel

32 core, ~232GiB RAM

GPU

32 core, ~232GiB RAM,
Nvidia GPUs

HighMem

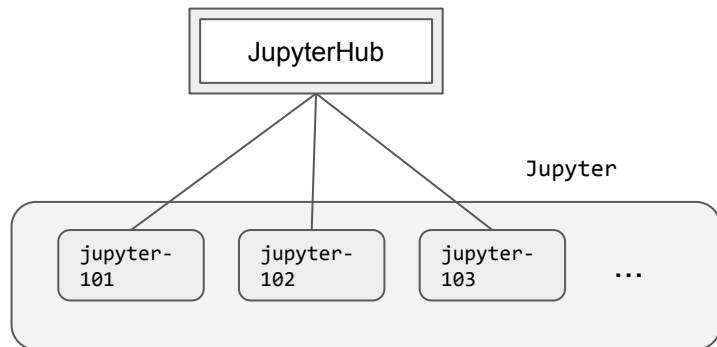
32 core, 503GiB RAM
96 core, 1.5TiB RAM

85 + 12 Nodes

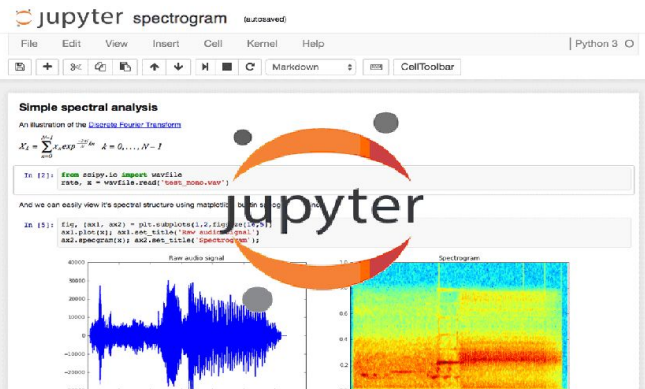
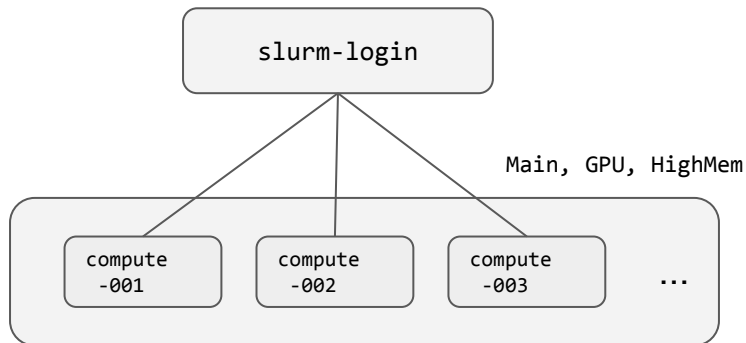
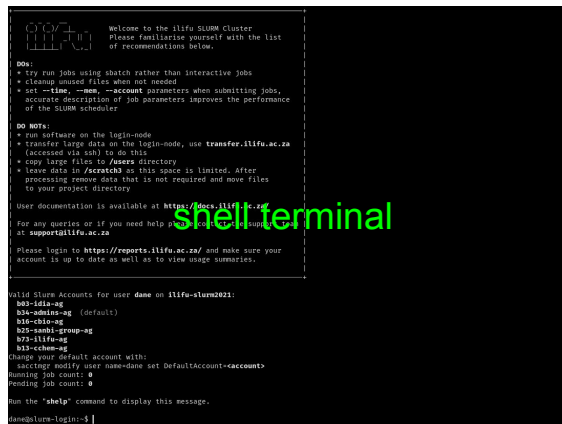
8 nodes

8 nodes

<https://jupyter.ilifu.ac.za>



`$ ssh <username>@slurm.ilifu.ac.za`

Screenshot of a shell terminal showing the output of the `ssh` command. The terminal displays a welcome message and a list of recommendations for using the cluster. The text "shell terminal" is overlaid in green.

\$ ssh <un>@slurm.ilifu.ac.za → connect to login node (from your machine))

\$ sinfo → shows partitions and resources

\$ squeue → shows all jobs in SLURM queue/partition

\$ squeue -u \$USER → shows your jobs

\$ sbatch slurm_job_script.sh → submit job to SLURM

\$ sbatch --help → information on job submission parameters

\$ scancel <jobid> → cancel running / pending job

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

```
module add python/3.11.2
```

```
python hello_world.py
```

```
-time=0-03:00:00    # 0 days + 3 hours
-mem=3G              # 3 GiB
-ntasks=1            # one task
-nodes=1              # one node
-partition=Main
-account=<your default>
```

Example slurm job script

https://github.com/ilifu/ilifu_user_training/tree/main/introduction/tutorial2

```
#!/bin/bash
#SBATCH --job-name=tutorial2_R_container
#SBATCH --time=00-00:01:00
#SBATCH --mem=4G
#SBATCH --partition=Main
#SBATCH --output=R_container-%j.stdout
#SBATCH --error=R_container-%j.stderr
#SBATCH --mail-user=YOUR_EMAIL_ADDRESS
#SBATCH --mail-type=BEGIN,END,FAIL,TIME_LIMIT_80
#SBATCH --account=ACCOUNTING_GROUP
```

Describe **job** parameters /
resources

```
singularity exec /software/common/containers/RStudio2023.06.1-524-R4.3.1.sif Rscript hello_world.R
```

container

software

Defaults and maximums per partition

https://docs.ilifu.ac.za/#/tech_docs/resource_allocation?id=maximum-allocation

Partition	Node names	Default CPUs	Max CPUs	Default Memory (GiB)	Max Memory (GiB)	Default wall-time	Max wall-time
Main	compute-[002-021]	1	32	3	232	3 hours	14 days
Main	compute-[101-105]	1	48	3	232	3 hours	14 days
Main	compute-[201-260]	1	32	3	251	3 hours	14 days
HighMem	highmem-[001-002]	1	32	15	503	3 hours	14 days
HighMem	highmem-003	1	96	15	1508	3 hours	14 days
GPU	gpu-[001-004]	1	32	7	232	3 hours	14 days
GPU	gpu-005	1	24	7	232	3 hours	14 days
GPU	gpu-006	1	48	7	354	3 hours	14 days
GPU	gpu-007	1	48	7	354	3 hours	14 days
Devel	compute-001	1	32	-	-	3 hours	12 hours

DEMO TIME!

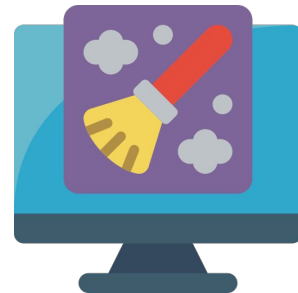
https://github.com/ilifu/ilifu_user_training/tree/main/session1/tutorial2

Do's :

- Run jobs using sbatch rather than interactive jobs
- Identify job resources requirements:
 - No. of nodes and CPUs, amount of RAM and wall-time.
- Remove files that aren't needed
 - /scratch3 folder after data processing is complete
 - Old raw data, temporary products , etc.
- Use Singularity (cannot install software on nodes)
- Use <username>@transfer.ilifu.ac.za for data transfers

Don't:

- Don't run software/heavy processes on login node
- Don't place large files in your home directory (/users)
- Don't transfer using scp/rsync on the login node



Thank you

<https://docs.ilifu.ac.za/>

support@ilifu.ac.za

