

ilifu Support Astronomer, IDIA, Department of Astronomy, University of Cape Town Adjunct Fellow, Western Sydney University



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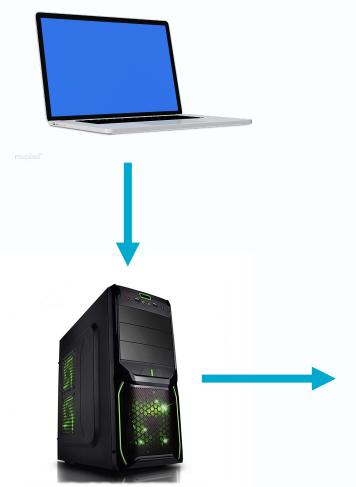




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ilifu: from laptop to VM to cluster









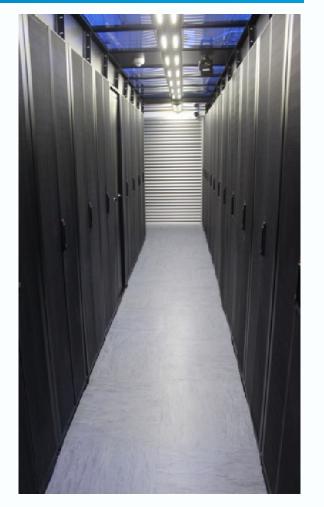




ilifu

- ilifu (<u>http://ilifu.ac.za</u>)
 - Tier 2 Data Intensive Research Facility
 - Joint Cloud Platform for Astronomy and Bioinformatics
 - Cluster w/ ~85 nodes (32 CPUs, 256/512 GB)
 - ~6 PB usable storage (BeeGFS & CephFS)
 - 10 Gb/s network to South African National Research Network (SANReN)







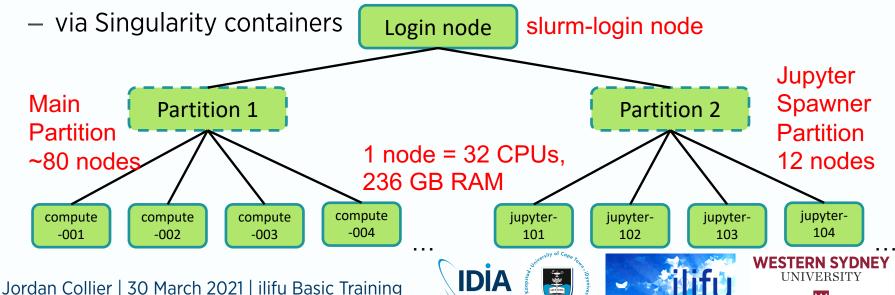






SLURM

- http://docs.ilifu.ac.za/#/getting_started/submit_job_slurm
- Login node (job submission & management)
 - where you land when you log in (also known as "head node")
 - run SLURM commands/submit jobs, but not software/heavy processes
- Compute nodes
 - Where your processes run (also known as "worker nodes")



SLURM

- http://docs.ilifu.ac.za/#/getting started/submit job slurm
- ssh <username>@slurm.ilifu.ac.za
- https://jupyter.ilifu.ac.za
- Partitions: Main (~80 nodes), HighMem (2), GPU (4), Jupyter (12)

ssh - shell terminal

JupyterHub System information as of Fri Aug 23 11:36:57 SAST 2019 Login node slurm Users logged in: 8
IP address for ens3: 192.168.100.39 Analyze and Image Simulation 35.9% of 21.15GB In [298]: from simutil import IP address for ens4: 10.102.26.97 Memory usage: 5% IP address for ens5: 10.102.28.133 Swap usage: nafile - 'sim exaple 30 dor/sim exaple 30 dor.meerkat.m Keen to learn Istio? It's included in the single-package MicroK8s. https://snapcraft.io/microk8s Get cloud support with Ubuntu Advantage Cloud Guest: http://www.ubuntu.com/business/services/cloud klam n = 300/ga.convert(model specrefyal.'GHz')('value') Canonical Livepatch is available for installation. Reduce system reboots and improve kernel security. Activate at: https://ubuntu.com/livepatch 1 node = 32 CPUs,pl.xlabel('u[klambda]',fontsize='x-snall')
pl.ylabel('v[klambda]',fontsize='x-snall')
pl.axis('equal') 70 packages can be updated. updates are security updates. Out [55]: 4-45.0. 45.0. -30.0. 30.00 **236 GB RAM** eremy@slurm-login:~\$ sinfo AVAIL TIMELIMIT NODES STATE NODELIST up 14-00:00:0 mix slwrk-[106-113] alloc slwrk-[101,104-105,114-124] slwrkslwrkup 14-00:00:0 38 idle slwrk-[102-103,125-160] mix slwrk-[201-202,205,209] 104 201 infinite alloc slwrk-[206-208,210] infinite IDIA UNIVERSITY

SLURM – user commands

- https://slurm.schedmd.com/
- \$ sinfo #shows partitions and resources
- \$ squeue #shows all jobs in SLURM queue
- \$ squeue -u <username> #shows your jobs
- \$ sbatch slurm_job_script.sh #submit job to SLURM queue
- \$ sbatch --help #describes input parameters
- \$ scancel <jobid> #cancels job
- \$ sacct #shows status of recent jobs that have run, or are running









SLURM – running a job

```
$ cat slurm job script.sh
#!/bin/bash
#SBATCH -- job-name=demo job
#SBATCH --time=00:01:00
#SBATCH --mem=4GB
#SBATCH --output=demo-job-%j.out
#SBATCH --error=demo-job-%j.err
#SBATCH --account=b03-idia-aq
module load python/3.7.7
python myscript.py
$ sbatch slurm job script.sh #submit job to SLURM queue
```









SLURM – running a job

```
$ cat slurm job script.sh
#!/bin/bash
#SBATCH -- job-name=demo casa
#SBATCH --time=01:00:00
#SBATCH --mem=4GB
                                               Describe job
#SBATCH --output=demo-job-%j.out
                                               parameters/resources
#SBATCH --error=demo-job-%j.err
#SBATCH --account=b03-idia-aq
                                                      container software
                                                                               script
echo "Submitting demo SLURM job"
singularity exec /idia/software/containers/casa-stable-5.7.0.simg casa -c myscript.py
                                                               what's being executed with
$ sbatch slurm job script.sh #submit job to SLURM queue
                                                               above parameters
```









SLURM – running a job

```
$ cat slurm job script.sh
#!/bin/bash
#SBATCH --job-name=demo casa
#SBATCH --time=01:00:00
#SBATCH --mem=4GB
#SBATCH --output=demo-job-%j.out
#SBATCH --error=demo-job-%j.err
#SBATCH --account=b03-idia-aq
module load casa/5.7.0
casapy -c myscript.py
$ sbatch slurm job script.sh #submit job to SLURM queue
```









SLURM – running an interactive job

http://docs.ilifu.ac.za/#/getting_started/submit_job_slurm
 ?id=interactive-slurm-session

```
$ srun --pty bash #opens bash shell session on compute
#node with default 3 hours and ~7GB RAM
```

```
$ srun --pty --time=01:00:00 --mem=64GB singularity exec
/idia/software/containers/casa-stable.img casa
```

```
#opens interactive CASA session on compute node,
#with 1 hour walltime, 64GB RAM,
#using Singularity CASA container
```

\$ srun --help

#view srun help docs for input parameters





SLURM – running an interactive job

With X11 support for viewing GUI

```
$ ssh -Y <username>@slurm.ilifu.ac.za
#important to include the -Y parameters when
logging into the SLURM login node, for X-forwarding
$ srun --x11 --pty bash
#opens bash shell session on compute node
#with X11 support; default 3 hours, ~7GB RAM
```

\$ srun --x11 --pty --time=00:10:00 --mem=1GB --qos qos-interactive bash









Demo Time!









Best practices

- Don't run software / heavy processes / scp on the login node
 - Only submit jobs and run SLURM commands (sbatch, srun, squeue, etc)
 - Use transfer.ilifu.ac.za to transfer data (external/internal), not login node
- Before running a large job, identify the available resources
 - Use sinfo. Don't hog the cluster. Reduce your allocation if possible
 - Increase likelihood of jobs running with less memory and less walltime
- Use sbatch (srun / screen / tmux are volatile)
- Cleanup files that aren't needed
 - Old raw data, temporary products, scratch data, etc
- Don't place large files in your home directory (/users)
- Use Singularity (you cannot install software on the nodes)









THANK YOU

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