



ilifu Online Training

Jeremy Smith
Ilifu User Training Workshop – Basic Training
30 March 2021





- Support channels
- Introduction to the ilifu research facility services
- Directory structure
- Software Environment
 - Singularity containers
 - Modules
- Using JupyterHub
- Introduction to SLURM
 - Submitting a job on SLURM
 - Interactive sessions on SLURM
- Best Practices



Getting help



- Support contact support@ilifu.ac.za
- User documentation <u>http://docs.ilifu.ac.za/#/</u>
- Ilifu System Status https://status.ilifu.ac.za/
- Training videos http://www.ilifu.ac.za/il/accessing-facilities/training







Cloud-based infrastructure for data-intensive research

- Support variety of different scientific projects and requirements
- Data management: storage, transfer
- Flexible compute environment







Cluster & Job Scheduler



• JupyterHub service – development environ.



Containerised software environment



• Other services: data transfer, CARTA





Computing environment - interface

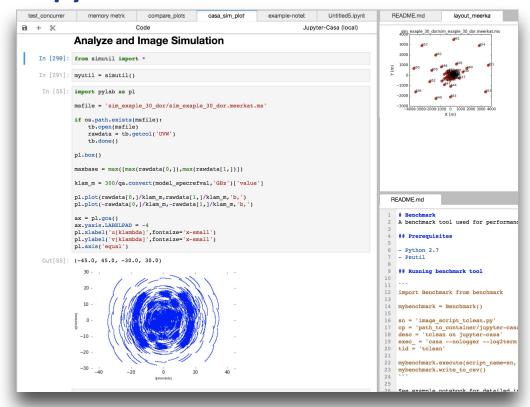


ssh - shell terminal

```
https://ubuntu.com/advantage
 System information as of Fri Aug 23 11:36:57 SAST 2019
 System load: 0.49
                                  Users logged in: 8
 Usage of /: 35.9% of 21.15GB IP address for ens3: 192.168.100.39
                                  IP address for ens4: 10.102.26.97
                                 IP address for ens5: 10.102.28.133
* 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
* Canonical Livepatch is available for installation.
  - Reduce system reboots and improve kernel security. Activate at:
    https://ubuntu.com/livepatch
170 packages can be updated.
75 updates are security updates.
ast login: Fri Aug 23 09:08:21 2019 from 196.11.235.232
jeremy@slurm-login:~$ sinfo
PARTITION
                 AVAIL TIMELIMIT NODES STATE NODELIST
                                           mix slwrk-[106-113]
                    up 14-00:00:0
                                     14 alloc slwrk-[101,104-105,114-124]
                                          idle slwrk-[102-103,125-160]
JupyerSpawnerONLY
                        infinite
                                           mix slwrk-[201-202,205,209]
JupverSpawnerONLY
                                      4 alloc slwrk-[206-208.210]
                        infinite
JupyerSpawnerONLY up infinite
                                      2 idle slwrk-[203-204]
eremy@slurm-login:~$ sbatch compute job.sh
```

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

JupyterHub



https://jupyter.ilifu.ac.za







- Common areas:
 - /USETS

 only 40 TB shared among all users, for scripts and small files don't place data here, capping /users storage capacity can prevent access to the cluster for all users.
 - Scratch storage:
 - /scratch/users or /scratch3/users
 - storage space for processing, temporary storage only, i.e. use this space during processing, and then clear all files immediately after processing. Remove unnecessary data and move data that you want to keep to project folder.
 - /software
- Remaining storage is separated by group:
 - IDIA, CBIO, Ilifu (DIRISA projects)







IDIA structure:

• /idia/users

- user's private workspace, may store data products that are not ready to move to shared project space (limited to 10TiB per user)

/idia/projects

- project specific directories. These directories are for sharing data and resources within project groups. Raw data associated with a project will also be available from the project folder. Raw data should always be read-only.

• /idia/software

- software containers and the IDIA Pipelines software is stored here

/idia/software/containers







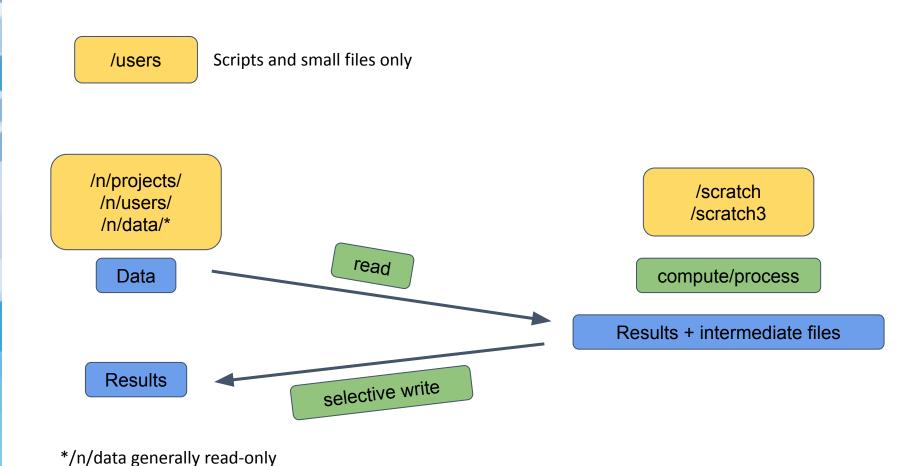
Similar structure for /cbio and /ilifu groups,

- /cbio/users
- /cbio/projects
- /cbio/soft
- /ilifu/users
- /ilifu/software
- Exception for ilifu projects:
 - /ilifu/astro/projects
 - /ilifu/bio/projects



Directory Structure - workflow







Singularity containers





- Encapsulated software environments
- A software stack that contains everything required to run an application/workflow, including files, environmental variables, libraries and dependencies
- Containers accessible across platforms and services, allowing sharing of application environments







Supported Containers:

- CASA
- KERN suite
- Astronomy container (ASTRO-PY3.simg)
- Python 2.7, Python 3.6, R
- Project containers:
 - MeerLICHT,
 - Simba
 - HI intensity map
- /idia/software/containers; /cbio/soft; /ilifu/software/containers



Singularity containers (S)





Open container as an interactive shell:

singularity shell /path/to/container

Example:

\$ singularity shell /idia/software/containers/ASTRO-PY3.simg

Run a script/workflow using a container environment:

singularity exec /path/to/container <software> <script/input params>

\$ singularity exec /idia/software/containers/casa-stable.img casa -c myscript.py





Software environment - modules

module avail

| LAPACK/3.9.0 | | | /modules/common 3/login.old | | drmaa/1.1.1 | c | penBLAS/0.3. | 9 | perlbrew/ | perlbrew |
|--------------------------|---------------|---------------------------|----------------------------------|---------|---------------------------|---------|---|---------|---------------|----------|
| uby/2.6.6 | | | ,, , | | | | ,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,, | | P, | F |
| R/RStudio1.2.5042-R4.0.0 | | anaconda3/login | | | homebrew/2.4.13 | | openmpi/2.1.6 | | python/2.7.18 | |
| R/RStudio1.2.5042-R4.0.4 | | anaconda3/2020.07 (D) | | | java/jre-1.8.0_261 | c | openmpi/3.1.6 | | python/3.7.7 | |
| R/3.6.3 | | cuda/10.0 | 0.130_410.48 | | java/openjdk-14.0.1 | (D) c | penmpi/4.0.3 | 3 | python/3. | 8.2 |
| R/4.0.0 | | cuda/10. | 1.243_418.87.00 | | julia/1.5.3 | c | penmpi/4.0.5 | 5 | python/3. | 8.3 |
| R/4.0.2 | | cuda/10.2 | 2.89_440.33.01 | | maven/3.6.3 | c | penmpi/4.1.0 | (D) | python/3. | 8.6 |
| R/4.0.3 (D) | | cuda/11.0.2_450.51.05 (D) | | | mono/6.8.0.123 | | per1/5.33.0 | | python/3.9.0 | |
| | | /software | e/modules/bio | | | | | | | |
| | | | genomestrip/2.00.1958 | | | prsice- | prsice-2/2.3.1d treePL | | homebrew | |
| bebio/1.2.3 | canvas/1.40.0 | 1.1613 | genomestrip/2. | 00.1958 | plink/2.00a2.3 | F | 2/2.3.14 | CTCCTT/ | | |
| | |).1613 | genomestrip/2. | 00.1958 | plink/2.00a2.3 | Passass | 2,2.3.14 | CICCII) | | |
| | | | penomestrip/2.0 htslib/1.10.2 | 00.1958 | plink/2.00a2.3 popgen/0.1 | samtool | | | s/0.1.16 | vep/101 |
| p/singularity | canvas/1.40.0 | | | 00.1958 | | _ | | | s/0.1.16 | vep/101 |

Where:

D: Default Module







module avail

module help <module>

```
~$ module help python
----- Module Specific Help for "python/3.9.0" -----
This module configures Python 3.9.0 for use
```

- module load <module>
- module purge
- module --help



JupyterHub

IDİA

https://jupyter.ilifu.ac.za





