



# Introduction to Gadi — Part I

Get Full Access to All Resources on Gadi

NCI Training

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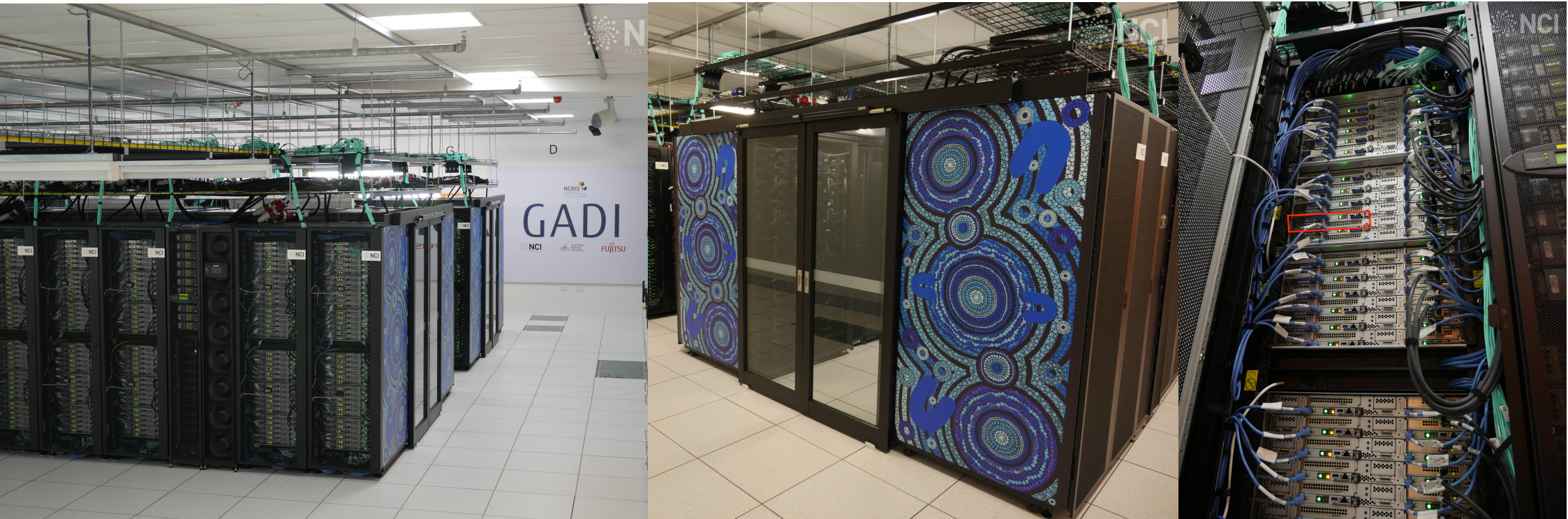
# Agenda

- First half of [Welcome to Gadi](#) opus page
- Three sections
  - Login
  - Storage and data transfer
  - Jobs
- Exercise + Key points + Q&A

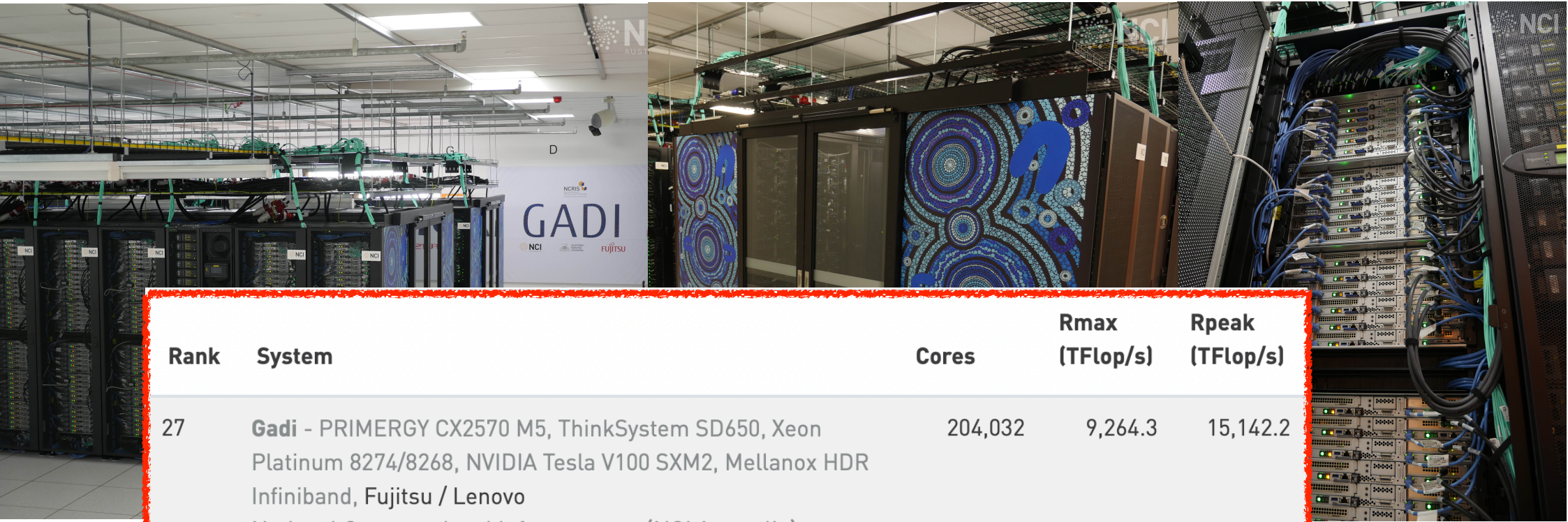
# What is Gadi

- High Performance Computing System operated by NCI
- Supercomputer
  - 10 login nodes
  - 6 data mover nodes
  - 4000+ compute nodes, including 160 GPU ones
  - Mellanox HDR InfiniBand interconnect network
  - 22 PiB storage available on the parallel Lustre filesystems
  - Application software available through environment module
  - PBSPro server handles 3-4M jobs per quarter

# Gadi

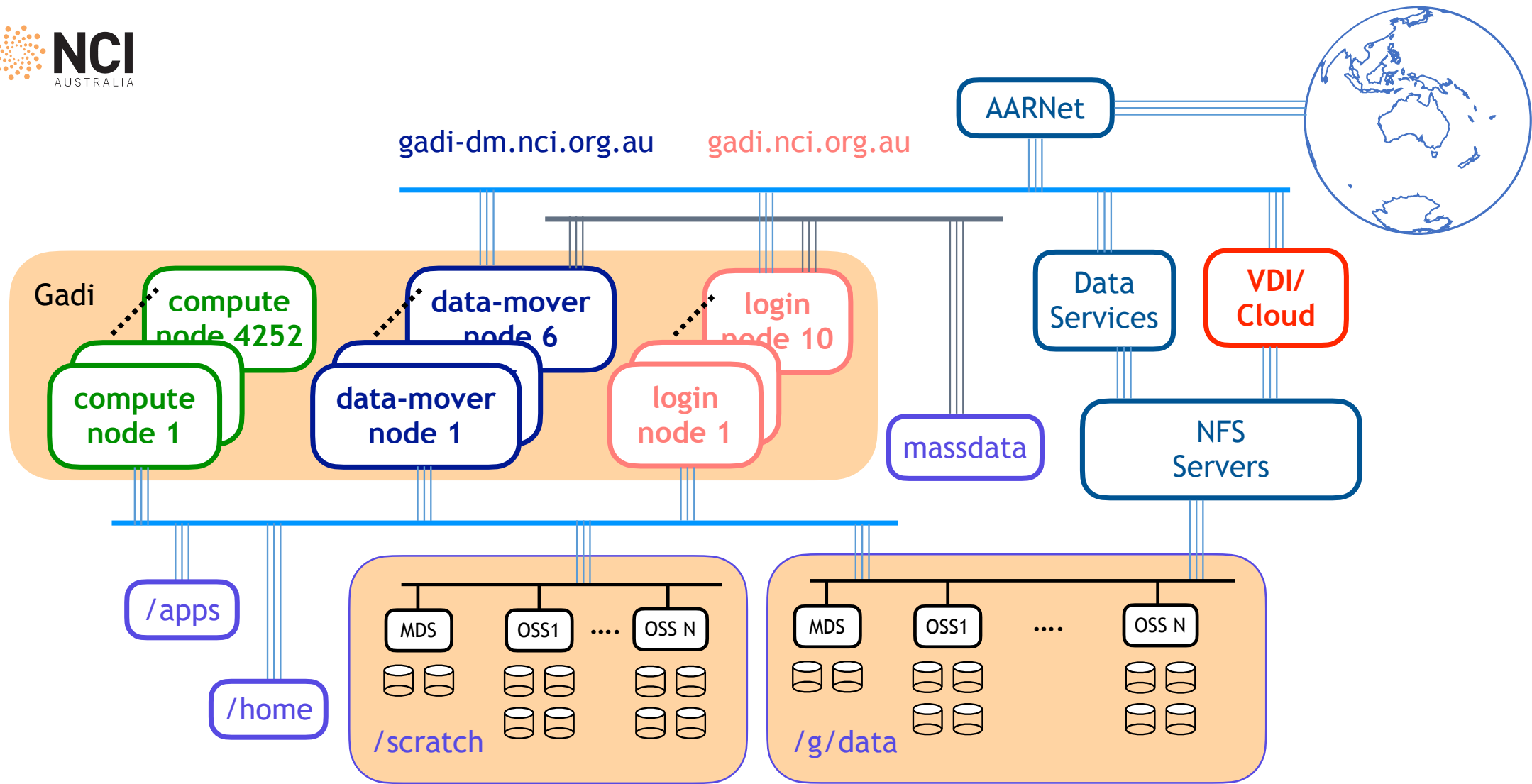


# Gadi



Rank	System	Cores	Rmax (TFlop/s)	Rpeak (TFlop/s)
27	<b>Gadi</b> - PRIMERGY CX2570 M5, ThinkSystem SD650, Xeon Platinum 8274/8268, NVIDIA Tesla V100 SXM2, Mellanox HDR Infiniband, Fujitsu / Lenovo National Computational Infrastructure (NCI Australia) Australia	204,032	9,264.3	15,142.2

<https://www.top500.org/lists/top500/list/2020/11/>





Login and Login Environment

# User, Project, and Shell

## Login to Gadi

- a. Open a terminal and login to one of the ten login nodes by doing

```
ssh <jjj777>@gadi.nci.org.au
```

Your username goes here

- b. X forwarding enabled login and test with xeyes

```
Sue@local:~ $ ssh -Y jjj777@gadi.nci.org.au
jjj777@gadi.nci.org.au's password:
[jjj777@gadi-login-05 ~]$ xeyes
[jjj777@gadi-login-05 ~]$ exit
Sue@local:~ $
```





## Explore Login Environment

- shell: echo \$SHELL
- who you are
  - user: echo \$USER
  - default project: echo \$PROJECT
- where you are
  - gadi-login-xx where xx=[01...10]: hostname
  - directory path: pwd, echo \$HOME

## Edit Login Environment

- To define your own login environment, edit the following two files
  - `~/.config/gadi-login.conf`
  - `~/.bashrc` for `SHELL=/bin/bash`, or `~/.cshrc` for `SHELL=/bin/csh` etc.
- Be careful when making changes because bad edit can lock you out

## Q&A

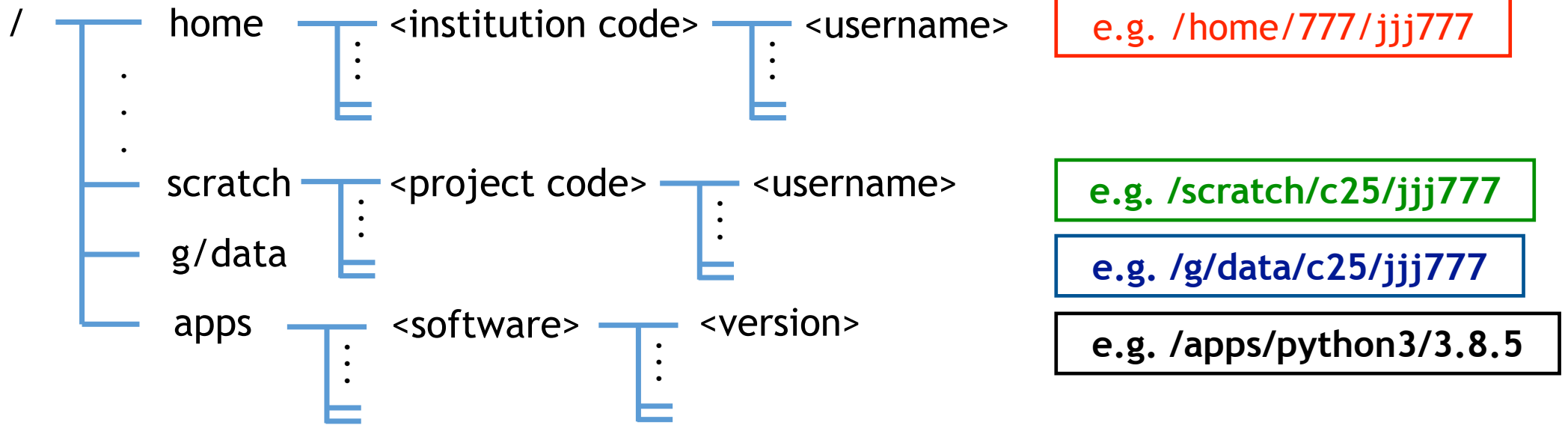
- MOTD
  - login
  - cat /etc/motd
- Usage limit on login node
  - 57 users and 64 login processes on gadi-login-01 at 9:18am this morning
  - Any processes on the login nodes will be terminated immediately if exceeding 30-minute CPU cumulative time limit and/or 4GB memory usage limit.
- Round-robin login
- Other questions?

A horizontal band with a background of a colorful, abstract microscopic image, possibly showing cellular structures in shades of red, orange, yellow, and blue.

Storage and Data Transfer

# Accessible Directories and their Limits

# Navigating through directories on login nodes



- Does your project have storage allocation on `/g/data/`? : `lquota`
- What's your data footprint on Lustre filesystems? : `nci-files-report -u $USER`



## Remote Transfer through Data-mover Node

Upload data to Gadi from local PC

```
Sue@local:~ $ scp test.hdf5 jjj777@gadi-dm.nci.org.au:/scratch/c25/jjj777/input.hdf5
```

*from/source*

*to/destination*

Download data from Gadi to local PC

```
Sue@local:~ $ scp jjj777@gadi-dm.nci.org.au:/scratch/public/IntroToGadi.pdf ./
```

*from/source*

*to/destination*

- You can also transfer data through `gadi-dm.nci.org.au` using
  - commands such as `sftp`, `rsync`,...
  - applications such as `FileZilla`, `WinSCP`,...
- No interactive sessions on any of the data-mover nodes, try interactive `copyq` jobs

## Q&A

- Shared filesystem vs compute node local disk \$PBS\_JOBFS
  - Shared access at all time vs 'exclusive' access during the running job
  - Size: 22 PiB vs ~400GB
- Home vs project directories vs massdata
  - Access frequency and inodes limit
  - Ownership
- Data located inside the project directory vs data owned by the project
- Apply for more storage space
  - on /scratch from NCI
  - on /g/data and massdata from scheme managers
- Other questions?



Jobs

# Submission and Monitoring

## Submit Jobs

```
cp /scratch/public/yxs900/gutentag.sh ./  
qsub gutentag.sh
```

## Any Errors?

```
$ qsub gutentag.sh  
qsub: Error: You are not a member of project c25.  
You must be a member of a project to submit a job  
under that project.
```

- User has to join a project to use its resources
- Use the -P flag in the command line to overwrite the job submission script

```
qsub -P $PROJECT gutentag.sh
```

## Monitor Jobs

- After submission, query the job status by running
  - `qstat -u $USER`
  - `qstat -sw <jobID>`
  - `qstat -f <jobID>`
- Delete submitted jobs
  - `qdel <jobID1> <jobID2> ...`
- After the job finishes, look up the jobs by running
  - `qstat -fx <jobID>` with the first 24 hours
  - `cat gutentag.sh.o<jobID>`
  - `cat gutentag.sh.e<jobID>`

## Q&A

- How much does the gutentag job cost
  - Queue → charge rate
  - CPU and memory request, walltime usage → resource hour
  - Reserved SU
- qsub, qstat, qdel all launch requests to PBS server
  - run with frequency  $< 0.1\text{Hz}$
- Other questions?

# Any More Questions?

[help@nci.org.au](mailto:help@nci.org.au)

A horizontal banner image showing a colorful, abstract microscopic view of tissue, likely stained with various dyes to highlight different cellular structures and components.

Thank you

Part 2 is coming next week on 18 March 2021