

# Smarter Computing: NCI GADI GPU Research Tips

## Research Seminar: Optimizing the Use of Computing Resources

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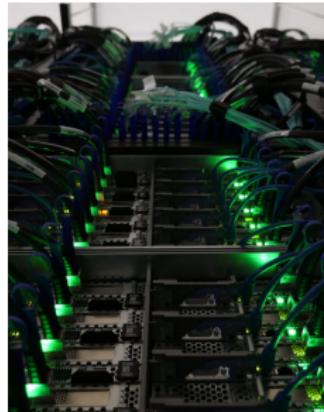
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# Introduction

# HPC system - Gadi

Gadi contains > 250,000 CPU cores, 930 Terabytes of memory & 640 GPUs.

- 160 nodes: 4 Nvidia V100 GPUs, 2 Intel Xeon processors each.
- 90 Petabytes of data storage capacity, reaching a total aggregate IO performance of around 450 GB/second.
- More than 200 supported software packages.
- PBS (Portable Batch System) is responsible for optimising the job scheduling and managing the workload of the cluster.



# HPC system - Gadi (cont.)

Make sure the efficient use of Gadi!



# Access to GADI - Two Approaches

- **Prerequisites**

Ensure OpenSSH is installed on your local machine.

## 1 Standard Terminal Access

```
$ ssh <yourAcc>@gadi.nci.org.au
```

## 2 SSH Config for Quick Access

1 Generate an SSH key on your local machine.

2 Append the following lines to the SSH config file in `~/.ssh/config`

```
Host gadi
```

```
    HostName gadi.nci.org.au
```

```
    User <yourAcc>
```

```
    IdentityFile ~/.ssh/id_ed25519
```

3 Set up public key authentication on Gadi:

```
$ ssh-copy-id -i ~/.ssh/id_ed25519.pub gadi
```

4 Afterwards, access Gadi without entering a password:

```
$ ssh gadi
```

# Access to GADI - Login Nodes

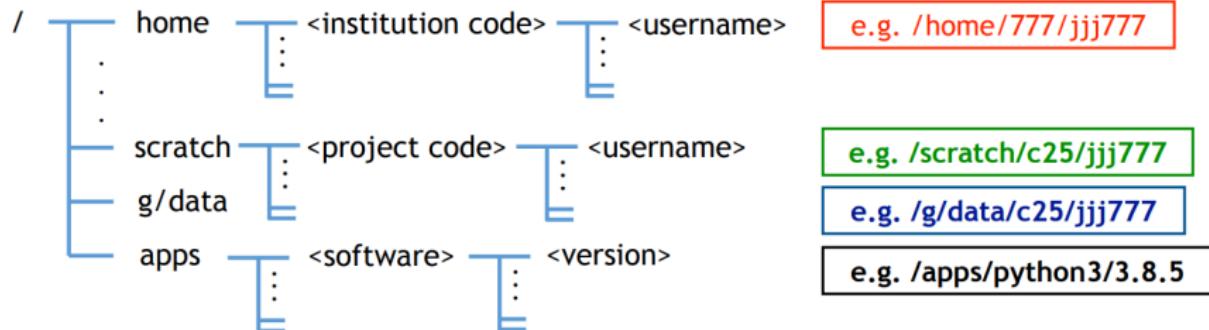
```
<welcome message: Mesage of the Day>
=====
Last login: Wed Nov 22 17:00:04 2023 from 150.203.68.254
[qc2666@gadi-login-05 ~]$ pwd
/home/135/qc2666
[qc2666@gadi-login-05 ~]$ cat $HOME/.config/gadi-login.conf
PROJECT xj17
SHELL /bin/bash
```

- Processes exceeding **30 minutes** of CPU usage or **4 GiB** of memory will be terminated.
- Every user have **10 GiB** in Home directory.
- With backups in **\$HOME/.snapshot**

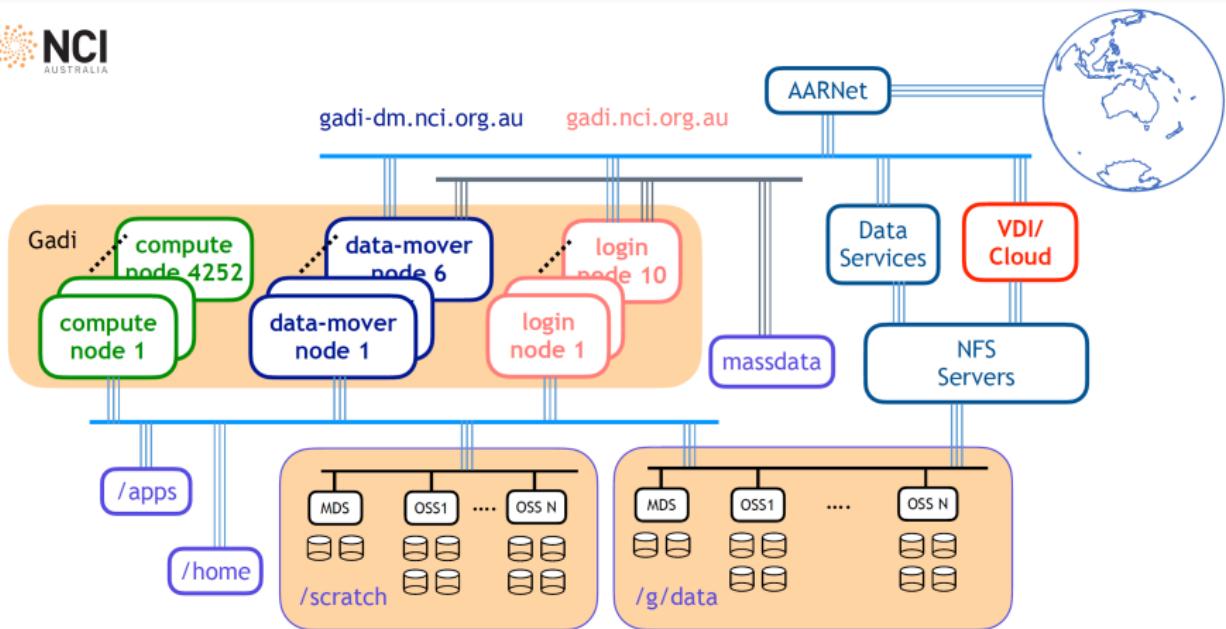
# Navigating through Gadi



## Navigating through directories on login nodes



# HPC Hardware Layout



# Environmental Settings

# Environment Modules

Environment Modules<sup>2</sup> provide packages/licenses in multiple versions, allowing users to easily switch between them.

- **To look for specific Applications/Licenses:**

```
$ module avail <app_name>
```

- **To load/unload a module:**

```
$ module load <app_name>/<version>
```

```
$ module unload <app_name>/<version>
```

- **To list all modules:**

```
$ module list
```

- **To remove all modulefiles from the current shell env:**

```
$ module purge
```

- \* **NCI-data-analysis module**

- Multiple GPU nodes: DeepSpeed
- Running Data Analysis Python Notebooks

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<sup>2</sup>Check [here](#) for more commands.

# Conda/Virtual Environment

Conda/Virtual environments facilitate the efficient management of specific dependencies, accommodating those with low data storage demands or those not included in the modules.

For Conda:

**1 Install a specific version of [Anaconda/Miniconda](#):**

```
$ wget <URL_Here>  
$ bash Anaconda3-xx.sh
```

**2 Create the conda environment and install necessary packages<sup>3</sup>.**

- \* Export all packages from a conda environment.

```
$ conda list --explicit > pkgs.txt
```

- \* Install packages in a new conda environment.

```
$ conda create --name <NEWENV> -- file pkgs.txt
```

---

<sup>3</sup>check [here](#) for more information.

# Conda/Virtual Environment (cont.)

For Virtualenv:

- 1 **Install<sup>4</sup> the package and make directory.**

```
$ pip3 install virtualenvwrapper  
$ mkdir ~/.virtualenvs
```

- 2 **Edit the .bashrc file by appending:**

```
export WORKON_HOME=$HOME/.virtualenvs  
source $HOME/.local/bin/virtualenvwrapper.sh
```

- 3 **Run \$ source ~/.bashrc**

- 4 **Load the desired python version and create the environment.**

```
$ module load python3/3.8.5  
$ mkvirtualenv --system-site-packages <NEWENV>
```

- 5 **Run \$ pip install <package> to install the necessary packages.**

- 6 **Run \$ workon <NEWENV> to activate.**

---

<sup>4</sup>check [here](#) for more information.

# Job Submission & Monitoring

# Submission Script

## Sample Script<sup>5</sup>:

```
#!/bin/bash
#PBS -P xj17
#PBS -q gputolta
#PBS -l ngpus=1
#PBS -l ncpus=12
#PBS -l mem=16GB
#PBS -l jobfs=10GB
#PBS -l walltime=00:05:00
#PBS -l wd
#PBS -l storage=gdata/xj17+scratch/xj17
cd /g/data/xj17/qc2666/demo
module load pytorch/1.10.0

# Activate Conda
# NOTE: Replace <ENV> with your actual conda environment name
#export CONDA_ENV='/home/135/qc2666/miniconda3/bin/activate'
#source $CONDA_ENV <ENV>

# Activate Virtualenv
# NOTE: Replace <ENV> with your actual virtualenv environment name
#export VIRTUAL_ENV='/home/135/qc2666/.virtualenvs/<ENV>/bin/activate'
#source $VIRTUAL_ENV

python3 main.py
```

<sup>5</sup>check [here](#) for more PBS Directives Explained.

# Queue Types

Queue		Max queueing jobs per project	Charge rate per resource*hour *	PBS_NCPUS	Max PBS_MEM/node †	Max PBS_JOBFS/node †	Default walltime limit
<b>gpuvolta</b>	<b>gpuvolta(route)</b>	1000	3 SU	multiple of 12	382 GB	400 GB	48 hours for 1-96 CPU cores
	<b>gpuvolta-exec</b>	50					24 hours for 144-192 CPU cores
<b>dgxa100</b>	<b>dgxa100(route)</b>	50	4.5 SU	multiple of 16	2000 GB	28 TB	48 hours for 16-128 cores
	<b>dgxa100-exec</b>	50					5 hours for 144-256 cores

# Submit jobs

- **Run \$ nci\_account -P <project code> to check the available resources and storage allocations.**
- **Run \$ qsub <submit\_job.sh> to submit the job.**
- **Run \$ qdel <job\_id> to delete the job.**
- **Run \$ for file in submit\_job\*.sh; do qsub \$file; done to submit multiple job scripts start with submit\_job.**
- \* **Running Array jobs.**

# Compute Grant and Job Debiting

$$\begin{aligned}\text{Job Cost (SU)} &= \text{Queue Charge Rate}^6 \\ &\times \text{Max [NCPUs, Memory Proportion]} \\ &\times \text{Walltime Used (Hours)}\end{aligned}$$

Memory Proportion = Mem requested/Mem per core  
 Mem per core = Mem Per Node/NCPUs per node for queue

Queue	CPUs request	GPUs request	Mem request	Walltime Usage	Cost
gpuvolta	12	1*	90GB	5 hours	$3 \times 12 \times 5 = 180 \text{ SU}$
gpuvolta	12	1*	380GB	5 hours	$3 \times 12 \times \max[1, (380/12) \times (48/382)] \times 5 = 3 \times 12 \times \max[1, 3.97905...] \times 5 = 716.23 \text{ SU (rounded)}$

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<sup>6</sup>Charge rate of **gpuvolta** is 3 SU per hour, check [here](#) for more information.

# Monitoring Commands

- **Displaying Job Status:**

For a simple check: \$ qstat

For detailed information:

```
$ qstat -swx <job_ID>
```

```
$ qstat -Esw
```

- **Job state**

- Q - job is queued, eligible to run or routed.
- R - job is running.
- E - Job is exiting after having run.
- H - Job is held.
- F - job is finished.

- **Displaying Queue Status:**

```
$ qstat -Q
```

- **Displaying Server Status:**

```
$ qstat -B
```

# Monitoring Commands (cont.)

- **Check processes inside a job:**

```
$ qps <job_ID>
```

- **Check the status of CPU and memory of a job:**

```
$ nqstat_anu <job_ID>
```

- **Check all jobs in the project:**

```
$ nqstat
```

- **Access the GPU node for expanded monitoring capabilities.**

```
$ qstat -n1 to check the node.
```

```
$ ssh gadi-gpu-v100-0097
```

- **Check here for more detailed instructions.**

# Interactive Jobs

```
[qc2666@gadi-login-03 qc2666]$ qsub -I -qgpuvolta -Pxj17  
-lwalltime=01:00:00,ncpus=12,ngpus=1,mem=8GB,  
storage=gdata/xj17+scratch/xj17  
qsub: waiting for job 103392320.gadi-pbs to start  
qsub: job 103392320.gadi-pbs ready
```

```
[qc2666@gadi-gpu-v100-0006 qc2666]$ module list  
Currently Loaded Modulefiles:  
1) pbs  
[qc2666@gadi-gpu-v100-0006 qc2666]$ exit  
logout
```

# Interactive Login Using a Job Script

```
#!/bin/bash

# This is a simple PBS script that uses the "interactive" mode of PBS.
# See "Interactive-batch Jobs" in the "PBS User Guide".
#
# For interactive use you must submit this job with -I
# qsub -I this_script.sh

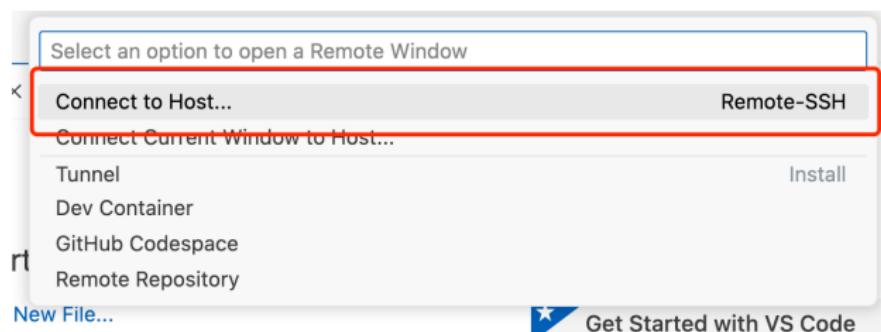
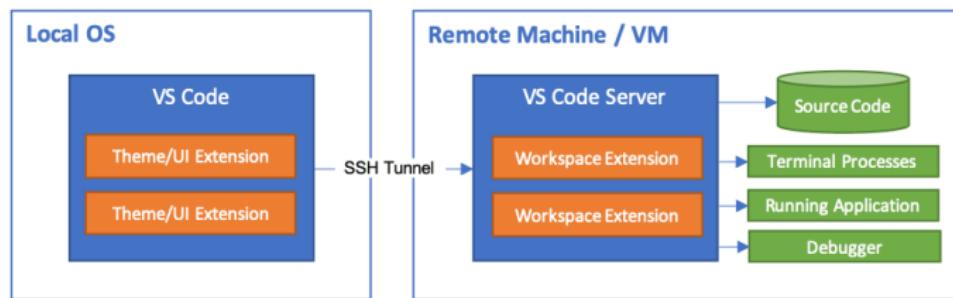
#PBS -P xj17
#PBS -q gpuvolta
#PBS -l ngpus=1
#PBS -l ncpus=12
#PBS -l mem=8GB
#PBS -l jobfs=10GB
#PBS -l walltime=01:00:00
#PBS -l storage=gdata/xj17+scratch/xj17

# Note: don't have any other commands below here!
```

# Operating/Debug Interface

# Remote Development using SSH

VScode or PyCharm



# Remote debug: Jupyter notebook

- 1 Submit a interactive job to access GPU node.

```
$ qsub -I access-gpu.sh
```

- 2 Load or activate the necessary dependencies.

- 3 cd to the working directory.

- 4 Check the GPU node and Run

```
$ jupyter-notebook --no-browser --ip=gadi-gpu-v100-0097 >  
notebook-output 2>&1 &
```

- 5 Run \$ jupyter notebook list to check the port & token.

- 6 Run

```
$ ssh -N -f -L 127.0.0.1:8888:gadi-gpu-v100-0097:<port>  
yourAcc@gadi.nci.org.au7 on your local machine.
```

- 7 Type 127.0.0.1:8888 in the browser<sup>8</sup> and enter the token.

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<sup>7</sup>You can replace the ssh address with the configured hostname, for example gadi on page 6.

<sup>8</sup>Please verify the GPU node, port, and token on the remote node, and ensure that a corresponding port is available on your local machine

# Code & Data Management

# File Transfer to/from Gadi

## ① Standard SCP Transfer:

- Command: `$ scp <source> <destination>`
- Example: `$ scp input.data gadi-dm.nci.org.au:/home/777/aaaa`

## ② Resumable Transfer with Rsync:

- Command: `$ rsync -avPS <source> <destination>`
- Example:  
`$ rsync -avPS gadi-dm.nci.org.au:/scratch/a00/aaaa/test_dir ./`

③ For instructions on transferring files larger than 500 GB, please refer to the [detailed job submission guide](#).

④ For effective code management, it is recommended to use [GitHub](#).

⑤ Check [here](#) for more detailed instructions.

# Dataset Management

- To view the storage quota details:  
\$ lquota
- For detailed information on the quotas of the group & users:  
\$ nci-files-report -g <Project\_ID>

- (base) [qc2666@gadi-login-08 qc2666]\$ nci\_account -P xj17

Usage Report: Project=xj17 Period=2023.q4

```
=====
```

Grant:	100.00	KSU
Used:	6.67	KSU
Reserved:	3.45	KSU
Avail:	89.88	KSU

Storage Usage Report: Project=xj17

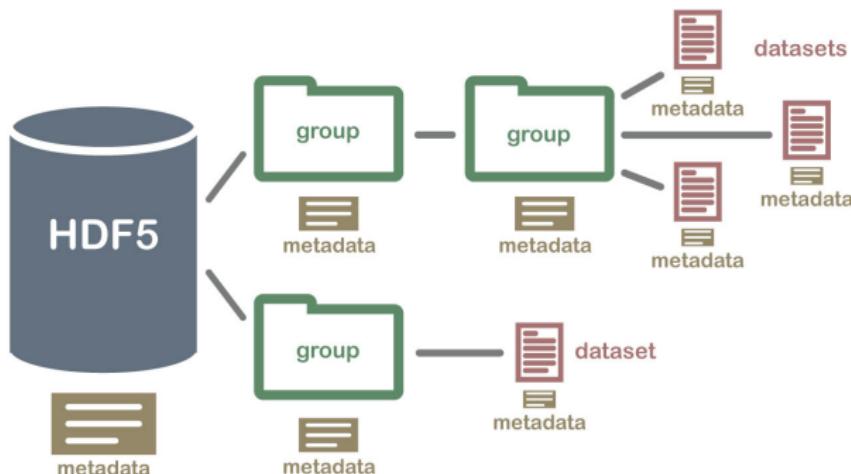
```
=====
```

Filesystem	Used	iUsed	Allocation	iAllocation
scratch2	32.66 MiB	1.91 K	1.00 TiB	202.00 K
gdata6	415.14 GiB	522.15 Kk	4.00 TiB	508.00 K **Over inode quota**

```
=====
```

# Dataset Management (cont.)

HDF5 (Hierarchical Data Format, Version 5), a versatile library and file format designed for storing scientific data, is recommended as a solution to address inode limits issues.



# Thank you!