Introduction to Digital Research Alliance of Canada Resources

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High Performance Clusters





Figure: Narval and Niagara Clusters

Interconnect

The InfiniBand Mellanox HDR network with a blocking factor of 33:7

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Resources

Website: Alliance Canada

Wiki: Technical Documentation

Account Roles

Advanced Research Computing: ARC Presentation

Research Data Management: Lunaris Platform

Research Software: Canadian Research Software Platforms

Training: Calender

Help: Technical Support

Login

MacOS/Linux

SSH: SSH Documentation

```
[name@server ~]$ ssh -Y sudhipv@narval.computecanada.ca
```

Windows:

PuTTY: Connecting with PuTTY

MobaXterm: Connecting with MobaXterm

Other Packages:

Cyberduck
VS Code Documentation
Carleton VS Code Tutorial

Environment

Login Nodes and Compute Nodes Project Folder and Scratch Folder

[sudhipv@beluga2 ~] \$ diskusage_report		
Description	Space	# of files
/home (user sudhipv)	15G/50G	295k/500k
/scratch (user sudhipv)	1335M/20T	3904/1000k
<pre>/project (group sudhipv)</pre>	0/2048k	0/1025
<pre>/project (group def-asarkar)</pre>	16G/1000G	190k/505k
<pre>/project (group def-maclab)</pre>	458G/1000G	36k/505k
<pre>/project (group rrg-asarkar)</pre>	17G/25T	3384/505k

Running Jobs

Wiki: Running Jobs

Slurm Scheduler: What is a Scheduler?

Slurm Commands: Slurm Quickstart Guide

Job Status

Slurm commands sq, sacct **Detailed usage status Portal** Screen command blog, cc

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Interactive Job

```
$ salloc --time=1:0:0 --mem-per-cpu=3G --ntasks=1 --account=def-asarkar
salloc: Granted job allocation 1234567
$ python helloworld.py # do some work
$ exit # terminate the allocation
salloc: Relinquishing job allocation 1234567
```

File Transfer

Personal to Cluster: SSH, Cyberduck, PuTTY, VS Code

Between Clusters: Globus File Transfer

GPU

NVIDIA Tesla GPUs: P100, V100, A100, H100

Multi Instance GPUs: MiG



Links to Linux Commands

cheatsheet1 cheatsheet2 bash

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