

CONTENT

- Introduction
- Specification & Node architecture
- Step by step on how to connect to the raplab-hackathon Cluster
 - STEP 1: Activate your Axis Account
 - STEP 2: Connecting to the Cluster
- Modules & environment
- Storage policies
- Common scheduler commands
 - Interactive mode
 - Batch mode (submission script)
- Containers
 - Singularity
- Transfer files to/from the Curiosity cluster

INTRODUCTION

About the Cluster:

The hackathon cluster consists of a single CPU-only login node and 10 DGX-A100 compute nodes.

NVIDIA and external users access the cluster via SSH to the login node. Cluster accounts are controlled via a web-based service called Axis.

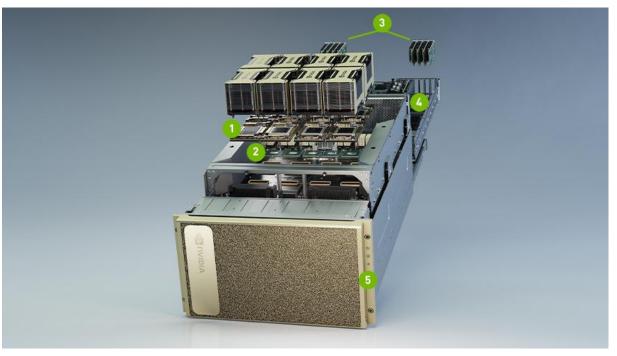
Join the OpenACC User Group SLACK workspace:

http://www.openacc.org/community#slack

CURIOSITY CLUSTER

Specs per node (total of 10 nodes)

- 1 8x NVIDIA A100 GPUs with 640 GB
 Total GPU Memory
 12 NVLinks/GPU, 600 GB/s GPU-to-GPU
 Bi-directonal Bandwidth
- 2 6x NVIDIA NVSwitches
 4.8 TB/s Bi-directional Bandwidth, 2X
 More than Previous Generation NVSwitch
- 3 10X NVIDIA Connectx-6 200 Gb/s Network Interface 500GB/s Peak Bidirectional Bandwidth
- Dual 64-Core AMD CPUs and 2 TB System Memory
 3.2X More Cores to Power the Most Intensive AI Jobs
- 5 30 TB Gen4 NVME SSD 50 GB/s Peak Bandwidth, 2X Faster than Gen3 NVME SSDs



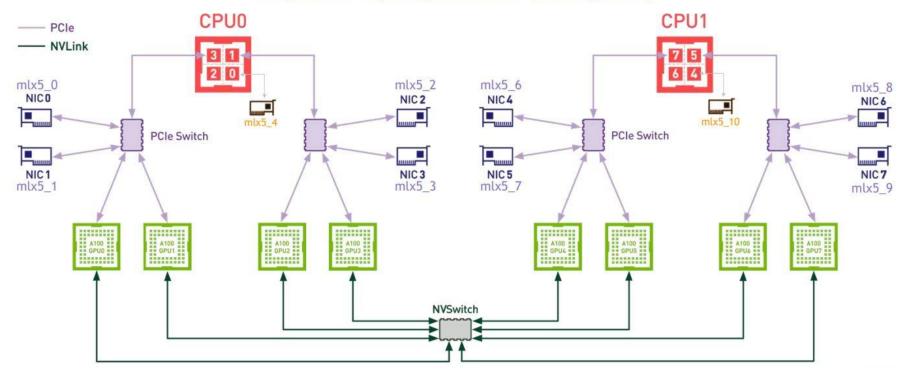
Node architecture

10 DGX-100A

DGX A100

High-level Topology Overview (with options)

Data plane (can be used as eth or IB)
Compute plane (IB)



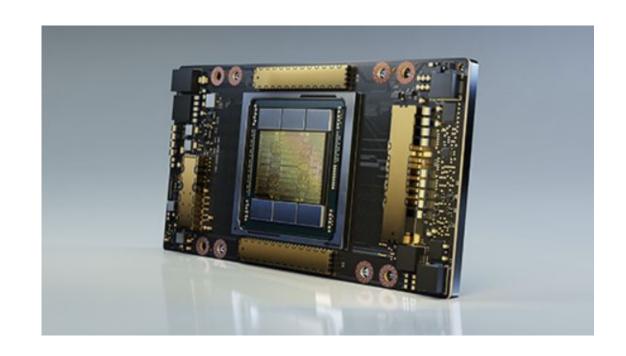
Node architecture

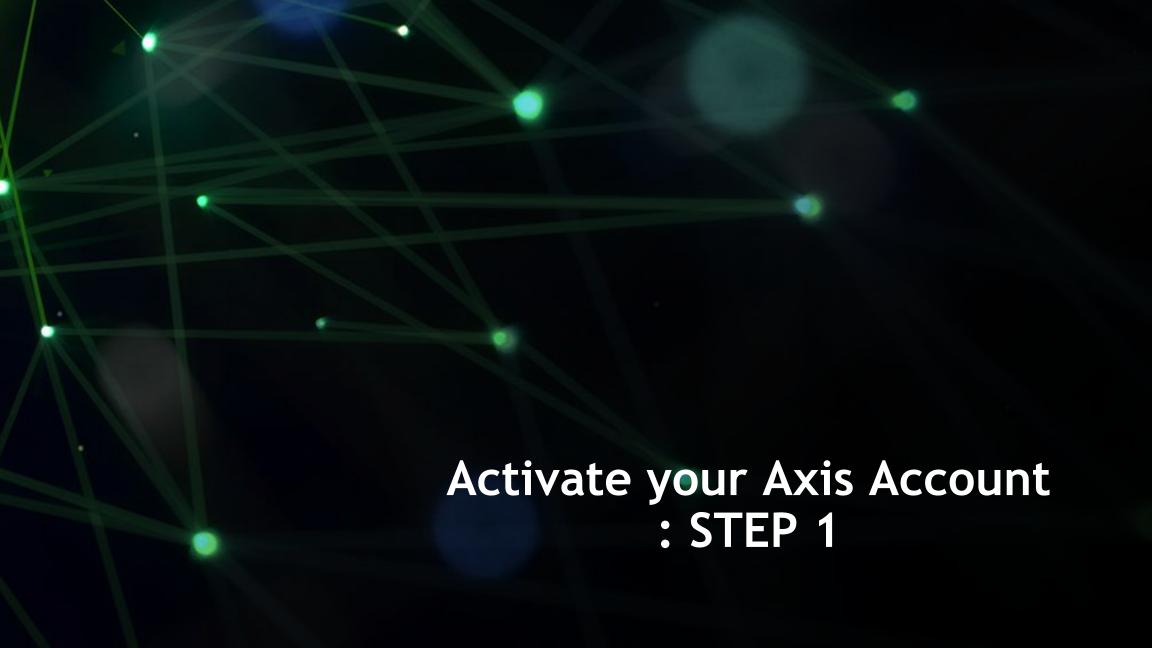
NVIDIA AMPERE A100

- 4 Tensor cores;
- 64 Single-precision (FP32) cores;
- 64 integer (INT32) cores;
- 32 double precision (FP64) cores;
- 256 KB of register file (RF);
- Up to 164 KB shared memory (configurable).

Tesla A100 peak computational throughput is:

- 312/624 Peak Tensor TOPS/TFLOPS
- 9.7 Peak FP64 TFLOPS
- 19.5 Peak FP32 TFLOPS



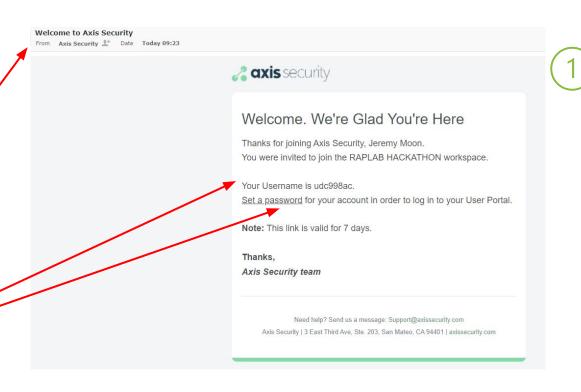


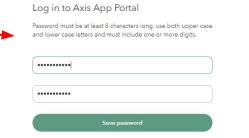
2: The welcome email will list your username.

3: Click on the Set a password link.

4: This will bring up a windows for you to set your password for AXIS.

5: Upon successfully setting your password you will be sent to the login page. Your AXIS account has now been activated.





2

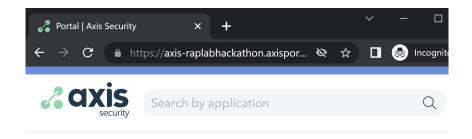


- → Login to Axis with your credentials
 - Link: https://axis-raplabhackathon.axisportal.io/apps
 - Use Chrome browser or make sure your browser does not block pop ups

1. Login to Axis with your credentials: https://axis-raplabhackathon.axisportal.io/apps

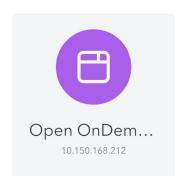
*Use Chrome or make sure your browser does not block pop ups!

2. Click on the "Hackathon Login" app



Sort by: Last Opened



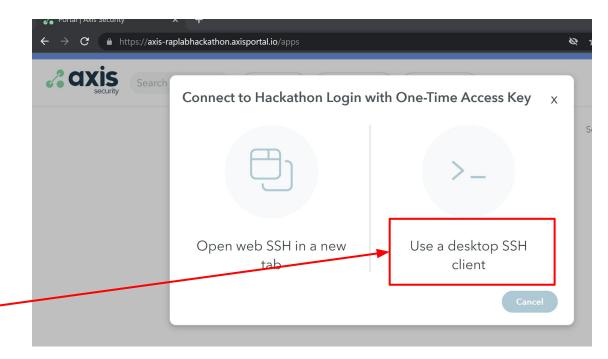


There are 2 methods to connect to the cluster.

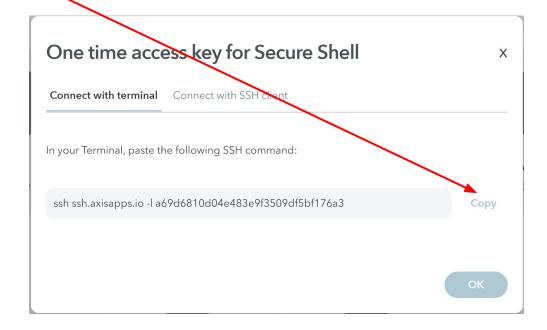
- Using a desktop client or terminal.
- 2) Web based SSH.

For Example for using a desktop client:

Click on the "Use a desktop SSH client"



Click "Copy" and open your preferred terminal (Powershell etc)



 Enter the command you copied in the last step and paste it into your terminal and click your enter key.

 You are now logged into the Cluster.

```
PS C:\Windows\System32\WindowsPowerShell\v1.0> ssh ssh.axisapps.io
                                                                   -l a69d6810d04e483e9f3509df5bf176a3
Welcome to Ubuntu 20.04.4 LTS (GNU/Linux 5.4.0-เซซ-generic x86 64)
  Documentation: https://help.ubuntu.com
  Management:
                  https://landscape.canonical.com
                  https://ubuntu.com/advantage
  Support:
 System information as of Tue 19 Apr 2022 08:08:58 PM UTC
 System load: 0.09
                                 Users logged in:
 Usage of /: 51.5% of 3.44TB IPv4 address for docker0: 172.17.0.1
                                 IPv4 address for ens10f0: 10.150.168.212
 Memory usage: 5%
                                 IPv4 address for ens2f1: 10.0.224.247
 Swap usage:
 Processes:
               1012
  Super-optimized for small spaces - read how we shrank the memory
  footprint of MicroK8s to make it the smallest full K8s around.
  https://ubuntu.com/blog/microk8s-memory-optimisation
 updates can be applied immediately.
To see these additional updates run: apt list --upgradable

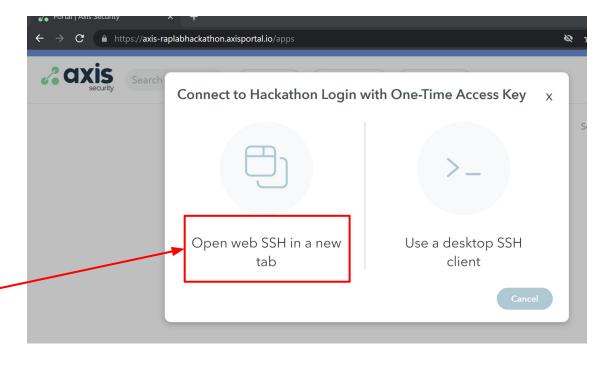
'** System restart required ***
Last login: Tue Apr 19 20:01:51 2022 from 10.150.169.22
Open OnDemand
Jsername:
assword:
 eremym-nvidia-com@rl-cpu-r82-u02:~$
```

There are 2 methods to connect to the cluster.

- Using a desktop client or terminal.
- Web based SSH.

2)Using the web based SSH:

Click on the "Open web SSH in a new tab"



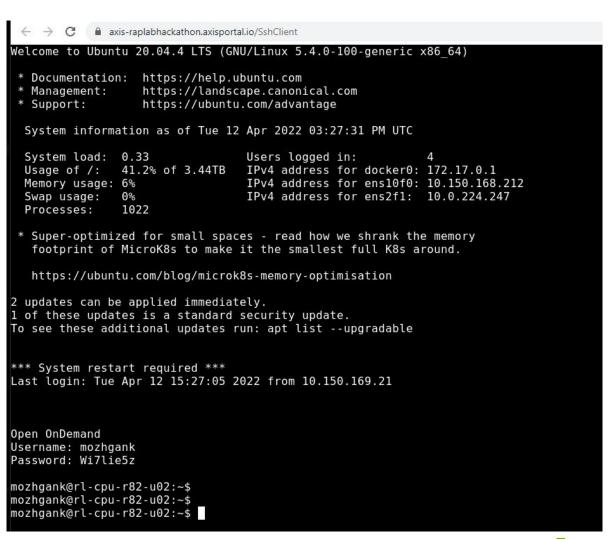
The main advantages of web-based SSH can be summarized as follows: Accessibility: Web-based SSH as described in this article requires no local installation of client software. It is thus possible to access SSH servers through a web browser from anywhere

Be patient ...

Connecting 10.150.168.212:22

Connected to Axis. Waiting for response...

You are now connected to the cluster





Common module commands

List currently loaded modules.

module list

List available packages.

module avail

Description of specified module.

module help [modulefile]

Displays information about specified module, including environment changes, dependencies, software version and path.

```
module show [modulefile]
```

Loads module or specifies which dependencies have not been loaded.

```
module load [modulefile]
```

Unloads specified module from environment.

```
module unload [modulefile]
```

Unloads all loaded modules

```
module purge
```

Overview environment

Current pre-installed modules

```
/sw/modules/nv-tools -----
nsight-compute/2021.3
              nsight-compute/2022.2.1 (D)
                                nsight-systems/2021.5.1 nsight-systems/2022.4.1 (D)
  compilers/nvhpc-21.5-mpi
                compilers/nvhpc-22.5-mpi (D)
                                  cuda/11.4.4
                                           cuda/11.6.1 gcc/9.3.0/python/3.9.10
compilers/nvhpc-21.9-mpi
                cuda/11.0.3
                                  cuda/11.5.1
                                           cuda/11.7.0
                                                        (D) gcc/9.3.0/python/3.9.13 (D)
compilers/nvhpc-22.2-mpi
                cuda/11.3.1
                                  cuda/11.6.0
                                           gcc/9.3.0/cmake/3.23.2
              conda/2022.9
          ------/sw/software/rootless-docker/modulefiles
rootless-docker
Where:
D: Default Module
```



Storage policy

- Central storage is via NFS
 - If you compile on the compute node then please use /tmp or local scratch
 - If you have heavy I/O then please please use /tmp or local scratch
- There is no quota on \$HOME
 - If you create lot of temporary files then please use /tmp on the compute node
 - If you create big profile traces then delete them as soon as you do not need them
- The /tmp location on compute node is not persistent!
 - When you job ends or die, /tmp gets cleaned up by the scheduler/OS
- When the hackathon ends and the account deleted, all \$HOME are purged. The is no backup
 - Save your files.



Slurm Common Commands

For full list of common Slurm command, follow this <u>link</u>.

Command	Description	Detailed
sinfo	Reports the state of the partitions and nodes managed by Slurm	PARTITION: the name of the partition AVAIL: whether the partition is up or down TIMELIMIT: the maximum length a job will will run in the format Days-Hours:Minutes:Seconds NODES: the number of nodes of that configuration STATE: down* if jobs cannot be ran, idle if it is are available for jobs, alloc if all the CPUs in the partition are allocated to jobs, or mix if some CPUs on the nodes are allocated and others are idle. NODELIST: specific nodes associated with that partition.
sacct	Lists the jobs that are running or have been run.	
squeue	Lists the state of all jobs being run or scheduled to run. Use squeue -u username to view only the jobs from a specific user	JOBID: number id associated with the job PARTITION: name of partition running the job NAME: name of the job ran with sbatch or sinteractive USER: who ordered the job to be ran ST: State of the job, PD for pending, R for running TIME: how long the job has been running in the format Days-Hours: Minutes:Seconds NODES: number of nodes allocated to the job NODELIST(REASON): either the name of the node running the job of the reason the job is not running such as JobHeldAdmin (job is prevented from running by the administrator).
scancel	Signals or cancels a job. One or more jobs separated by spaces may be specified.	

Job Submission

Resource Requests

To run your job, you will need to specify what resources you need. These can be memory, cores, nodes, GPUs, etc. There is a lot of flexibility in the scheduler to get specifically the resources you need.

Options	Description
nodes	The number of nodes for the job (computers)
mem	The amount of memory per node that your job need
-n	The total number of tasks you job requires
gres gpu:#	The number of GPUs per node you need in you job
gres=gpu:type:#	you can also specify the type of gpu. We have mostly p100s, but also 2 v100s for testing
qos	the QOS you want to run in, currently normal or debug
mem-per-cpu=	The amount of memory per cpu your job requires
-N	The minimum (and maximum) number of nodes required
ntasks-per-node=#	tasks per node.
exclusive	this will get you exclusive use of the node
constraint=	constrain to particular nodes. use skylake or broadwell for particular processor types

Scheduler (SLURM) policies

Each user can (by default)...

- Run one job at a time
- Request max 8 hours walltime
- Request 2 GPU and 32 cores (HyperThreading is enabled and MIG Disabled)
- Each team is limited to 2 GPU

If you are ready for benchmarks, scaling 1~8 GPUs, exploring NVLink effects...

- Request to mentors/sysadmins to see a possibility to enable the user access to a full node (8 GPUs and 128 cores)
- Walltime remains 8 hours but, if you have very long tests, an extension can be granted

Common scheduler commands

Interactive mode, 1 GPU

```
srun --ntasks=5 --nodes=1 --cpus-per-task=2 \
    --partition=gpu --time=4:00:00 --gres=gpu:1 \
    --pty /bin/bash
```

*Please note this is a sample run. The number of tasks will depend on the number of GPU core assigned per team.

Scheduler (SLURM) policy

Batch mode, 1 GPU (serial job)

```
#!/bin/bash
#SBATCH --partition=gpu
#SBATCH --nodes=1
                             # Max is 1
#SBATCH --ntasks=16
                             # Max is 16 (1/8 of 2x 64 AMD EPYC CPUs)
                             # Max is 2 (Clustered Multithreading is on)
#SBATCH --cpus-per-task=2
#SBATCH --gres=gpu:1
                             # Max is 1 (1 single A100)
#SBATCH --time=4:00:00
                             # Max is 4 hours
module purge > /dev/null 2>&1
#module load ...
export MYEXE=...
export OMP PROC BIND=true
export OMP NUM THREADS=5
./${MYEXE} 2>&1 | tee out.${SLURM_JOBID}
```

Scheduler (SLURM) policy

Batch mode, 1 GPU (parallel job)

```
#!/bin/bash
#SBATCH --partition=gpu
#SBATCH --nodes=1
#SBATCH --ntasks=4
#SBATCH --cpus-per-task=1
#SBATCH --gres=gpu:1
#SBATCH --time=4:00:00
module purge > /dev/null 2>&1
module load compilers/nvhpc-21.9-mpi
export MYEXE=hello world
mpicc -o hello world hello world.c
mpirun -np 2 --report-bindings ${MYEXE} 2>&1 | tee out.${SLURM JOBID}
```

Visual profiling tools

```
Nsight Compute (ncu-ui)
```

```
module load nvhpc/22.1 nsight-compute/2021.3
```

Nsight Systems (nsys-ui)

```
module load nvhpc/22.1 nsight-systems/2021.5.1
```

NVIDIA Visual Profiler (nvvp)

```
module load nvhpc/22.1 gcc/9.3.0/python/3.9.10
```



Containers

The Curiosity cluster supports <u>singularity</u> containers.

If you need to create your own containers without sudo. Singularity provides with a --fakeroot option, so when creating custom singularity containers.

NGC Containers

If you are looking for pre-configured containers, please look at GPU-optimized AI, Machine Learning, & HPC Software | NVIDIA NGC, you can also ask your Mentor for assistance.



Running singularity containers

- To transfer the images to the cluster, please visit the "Transfer files to/from hackathon cluster" section.
- You must first request and interactive session using srun command, then get an
 interactive shell in to the image, using the following command: singularity shell
 path/to/imagefile.simg
- To manually mount the existing directories in the image to your own path use -B flag: singularity shell -B path/to/file:/file path/to/imagefile.simg

Create your own Singularity image

```
Bootstrap: docker
From: nvcr.io/hpc/pgi-compilers:ce

%runscript

"$@"

%post

apt-get -y update
apt-get -y install --no-install-recommends python3-pip python3-setuptools zip

rm -rf /var/lib/apt/lists/*
pip3 install --no-cache-dir jupyter

%files

labs /labs

%environment
XDG RUNTIME DIR=
```

To build the container from the example def file (above) on CURIOSITY, use below command:

```
singularity build --fakeroot myimage.simg Singularity
```

To run interactively and using NVIDIA GPUs in the image: singularity shell --nv myimage.simg

To convert Docker containers to singularity, checkout: https://github.com/NVIDIA/hpc-container-maker

Checkout the singularity examples: https://sylabs.io/docs/

Singularity Commands

Build

Build takes a target as input and outputs a Container. Build is a multipurpose tool:

```
$ singularity [global options...] build [local options...] <IMAGE PATH> <BUILD SPEC>
```

Option1: Pull and Build a singularity container from DockerHub

```
$ singularity build docker://tensorflow/tensorflow:latest-gpu
```

Option 2: Pull and Build a singularity container from Definition Script (Below)

```
Bootstrap: docker
From: nvcr.io/nvidia/tensorflow:20.11-tf2-py3

%runscript

"$@"

%post

apt-get -y update
apt-get -y install --no-install-recommends python3-pip python3-setuptools zip build-essential rm -rf /var/lib/apt/lists/*
pip3 install --no-cache-dir jupyter

%files

English/ /labs

%environment
XDG_RUNTIME_DIR=

%labels
```

```
$ sudo singularity build tensorflow.simg Singularity
OR
$ singularity --fakeroot build tensorflow.simg Singularity
```

Run + Exec + Shell

Run: runs the Singularity runfile

```
$ singularity run <singularity container.simg>
```

 Exec: execute container commands from outside the container (like cat /etc/os-release)

```
$ singularity exec <singularity_container.simg> cat /etc/os-relase
```

Shell: opens a shell inside the container

```
$ singularity shell <singularity container.simg>
```

Nvidia GPUs

--nv flag enables the container to setup the environment to use NVIDIA GPUs and the CUDA libraries. Run, exec and shell commands can take the flag.

```
$ singularity build tensorflow.simg docker://tensorflow/tensorflow:latest-gpu
$ singularity run --nv tensorflow.simg
```

Loading local folders

You can use local files from within your container

```
$ echo "hello world" > file1.txt
$ singularity exec tensorflow.simg cat file1.txt
>hello world
```

Singularity binds /home/\$USER, /tmp, and \$PWD at runtime, to bind other directories use:

\$ singularity exec --bind /<host dir>:/test tensorflow.simg ls /test

This example binds the /test directory to a <host dir> inside the container so you can access it, allowing you to do an 1s on it.

Making changes to containers

Build as **--sandbox** for writable:

\$ sudo singularity build --sandbox test_tensorflow tensorflow.simg

--fakeroot option: fakeroot lets a user have nearly all the powers of root but only inside the container and does not extend to filesystem. Sudo is not required.

\$ singularity build --fakeroot test_tensorflow tensorflow.simg

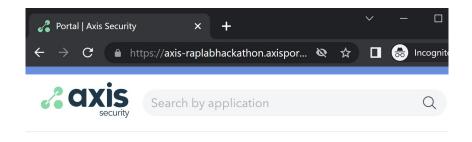


- → Login to Axis with your credentials
 - ◆ Link: https://axis-raplabhackathon.axisportal.io/apps
 - Use Chrome browser or make sure your browser does not block pop ups

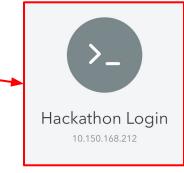
1. Login to Axis with your credentials: https://axis-raplabhackathon.axisportal.io/apps

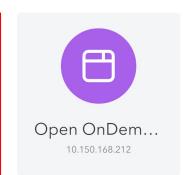
*Use Chrome or make sure your browser does not block pop ups!

2. Click on the "Hackathon Login" app



Sort by: Last Opened

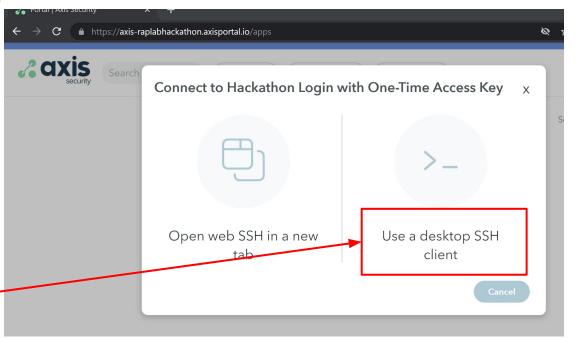




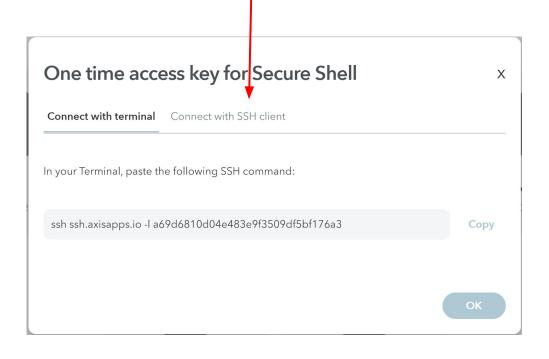
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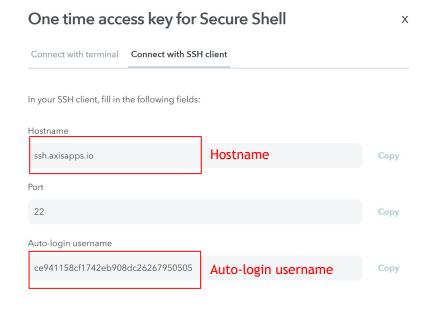
- Using a desktop client or terminal.
- 2) Web based SSH.

- 1) Using a desktop client or terminal:
- Click on the "Use a desktop SSH client"



Click on connect with SSH client, and make note of the Hostname and Auto-login username





- Before sending your files to the cluster it is recommended to zip them
- In your terminal (PowerShell or another) enter the following command replacing the hostname and username with the hostname and auto-login username from the previous step.

```
scp event-container.zip <username>@<hostname>:~
```

• And now unzip **to** a folder

```
unzip event-container -d event-container
```

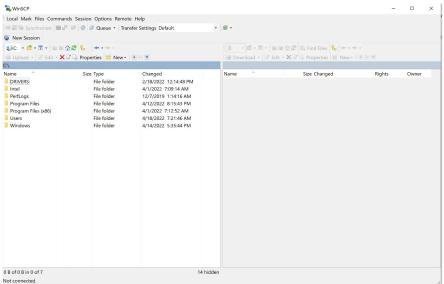
To transfer from Curiosity cluster

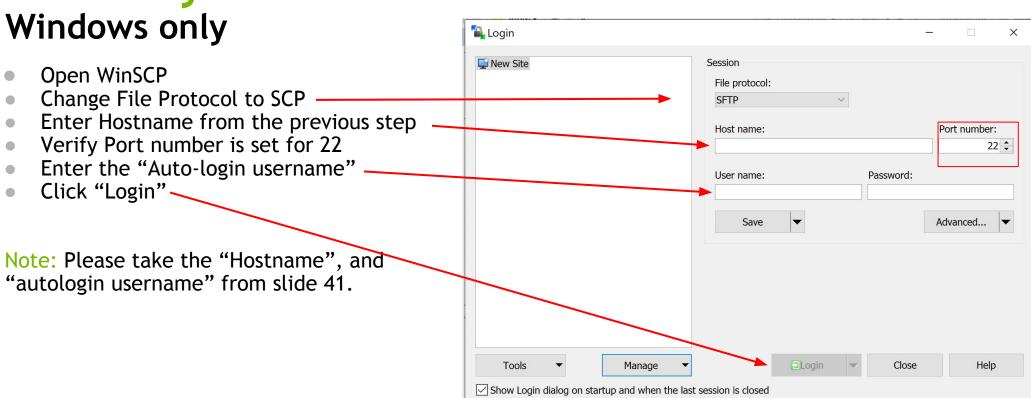
```
scp <username>@<hostname>:~/event-container.zip event-container.zip
```

Windows only

Install WinSCP

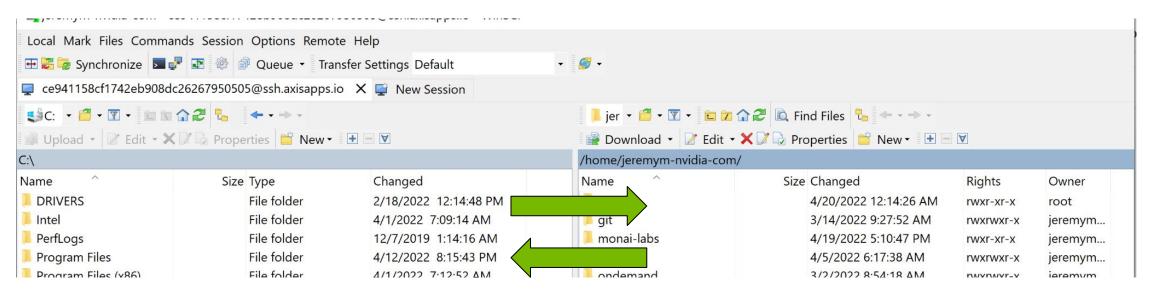
- Download the WinSCP client: https://winscp.net/eng/download.php
- Run the file after it's downloaded, and accept the License Agreement
- C. Click **Next** to select the default installation
- d. Click Install to begin the installation
- Click Finish





Windows only

Now, you can drag and drop files in the panel to transfer files from/to the local machine to the cluster.





Troubleshooting & Workarounds (1/N)

Q: NSight Compute does not operate / start properly issuing a warning that a lock /tmp/nsight-compute-lock is already present or owned by another user

A: NSight Compute uses a global lock (see https://docs.nvidia.com/nsight-compute/ProfilingGuide/index.html#serialization) since serialization across processes is necessary for the collection of HW performance metrics (some GPU and driver objects can only be acquired by a single process at a time).

A workaround to this can be to use a different TMP directory (not **/tmp!**). Before run the toop do the following:

export TMPDIR=/raid/\${SLURM_JOB_ID}/tmp

This change is valid only in the SSH session on the current running interactive job

