HIGH PERFORMANCE RESEARCH COMPUTING

ACES: Introduction to CryoSPARC for Cryo-EM Data Processing in Collaboration with the Laboratory for Biomolecular Structure and Dynamics



High Performance Research Computing

Fall 2024



CryoSPARC on ACES

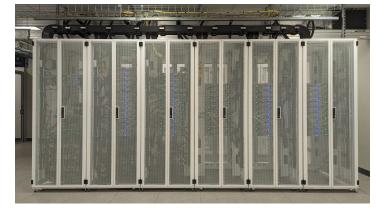
- CryoSPARC Academic License ID
- Resources and Limitations
- CryoSPARC on the ACES Portal
- Group data directories
- CryoSPARC Interactive Tutorial

CryoSPARC Academic License ID

- CryoSPARC provides a personal Academic License ID free of charge for TAMU staff and students
- A CryoSPARC Academic License ID is required to launch the CryoSPARC ACES portal app
- Use your academic email address for the Academic License
 - https://cryosparc.com/download

Resources for Running CryoSPARC

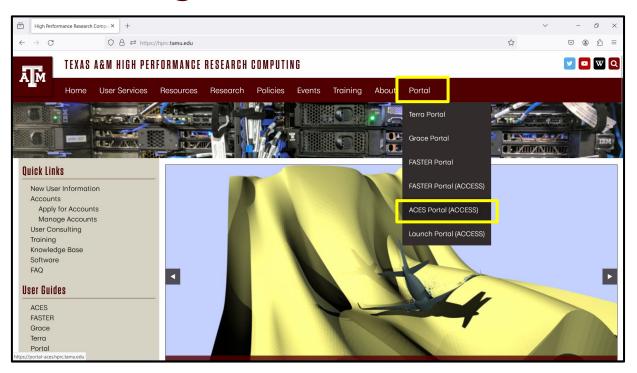
- CryoSPARC is available as an Interactive App on the ACES Portal
 - o https://portal-aces.hprc.tamu.edu
- ACES hardware overview (can change since ACES is a composable resource cluster)
 - 30 x H100 GPUs (2 x H100 GPUs per node)
 - 6 x A30 GPUs (6 x A30 GPUs per node)
 - o 70 x CPU only nodes
 - 10 x other specialized nodes not for CryoSPARC
- Specify enough time to allow your processing to complete
- If you launch a job for 24 hours and you finish your work in 12 hours, delete your job to make GPUs resources available for other jobs



Resource Limitations

- Can only launch one CryoSPARC portal session at a time
 - o can run multiple CryoSPARC jobs within a portal session
- CPUs
 - Some CryoSPARC tasks, such as importing images, do not require a GPU
- GPUs
 - The GPU queue can get busy at times and it may take over an hour for your job to launch
 - Cancel your pending job if you will be away from your computer for a long time and it hasn't started yet
- Submitted jobs can have delayed start times due to unavailability of GPUs or scheduled system maintenance

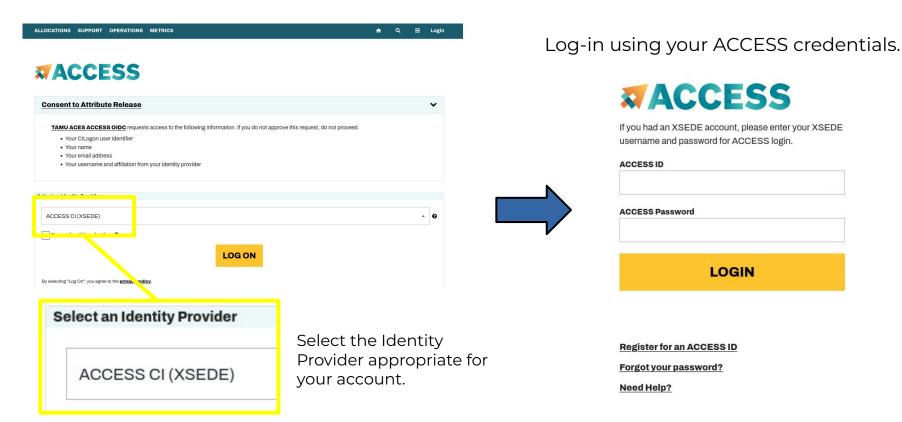
Accessing the HPRC ACES Portal



HPRC webpage: <u>hprc.tamu.edu</u>

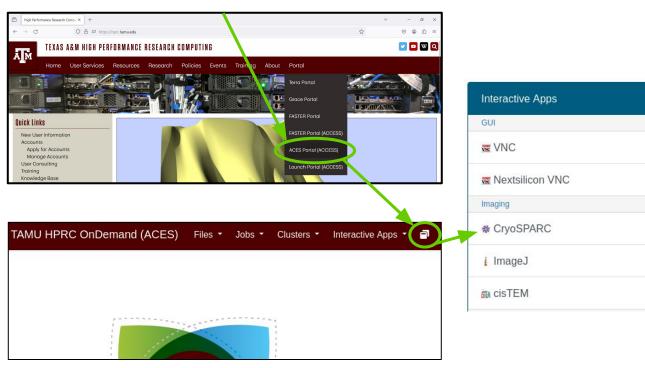


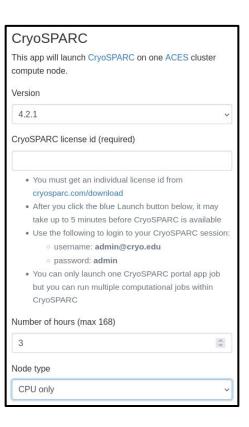
Accessing ACES via the Portal (ACCESS)



ACES CryoSPARC Portal App

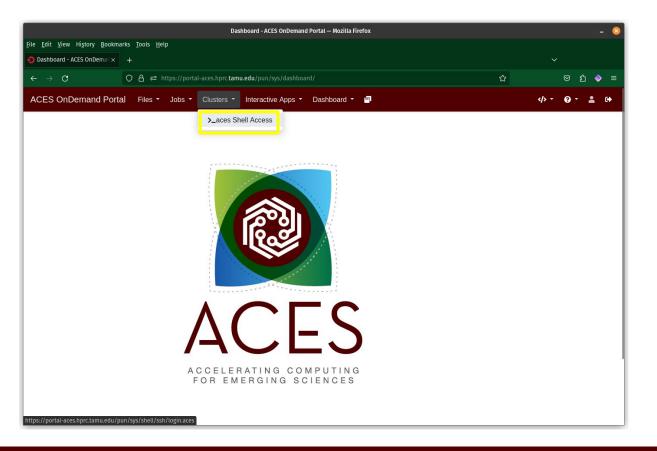
portal-aces.hprc.tamu.edu





Launch CryoSPARC the first time using **CPU only** in order to initialize the database.

Shell Access via the ACES Portal





Display GPU Availability on ACES

See GPU configuration and current availability using the ACES portal shell access

- https://portal-aces.hprc.tamu.edu
- select "Clusters" then ">_aces Shell Access"
- at the command line prompt, enter gpuavil
- the CONFIGURATION will change frequently since ACES is a composable resource cluster

Based on this output, there are 9 compute nodes on ACES that have exactly 2 x H100 GPUs attached to each gpuavail CONFIGURATION NODE NODE TYPE COUNT qpu:h100:2 9 gpu:pvc:4 qpu:h100:4 qpu:a30:2 AVAILABILITY NODE GPU GPU GPU CPU GB MEM NAME TYPE COUNT AVAIL AVAIL AVAIL h100 248 ac027 72 h100 96 488 ac048 h100 408 ac049 ac050 h100 488 ac064 a30 96 488

https://hprc.tamu.edu/kb/Software/useful-tools/gpuavail



ACES Maintenance

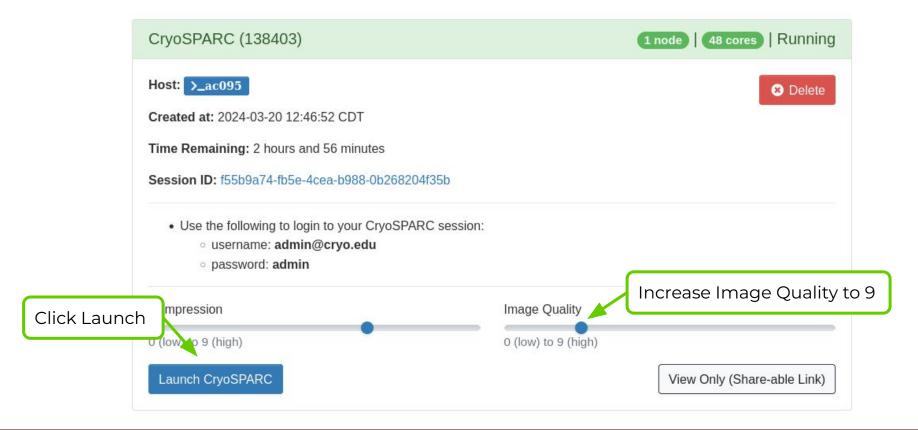
- ACES will be offline during routinely scheduled maintenances which may occur a few times each year
- Submitted jobs will remain PENDING and not run if the job script wall time overlaps with the maintenance start time
- When maintenance is complete, scheduled jobs will automatically begin running based on priority
- Use the maintenance command to see if a maintenance is scheduled and to display the time until the start of the maintenance

```
maintenance
The scheduled 35 hour ACES maintenance will start in:
   3 days 21 hours 32 minutes
```

Scheduled jobs will not start if they overlap with this maintenance window.

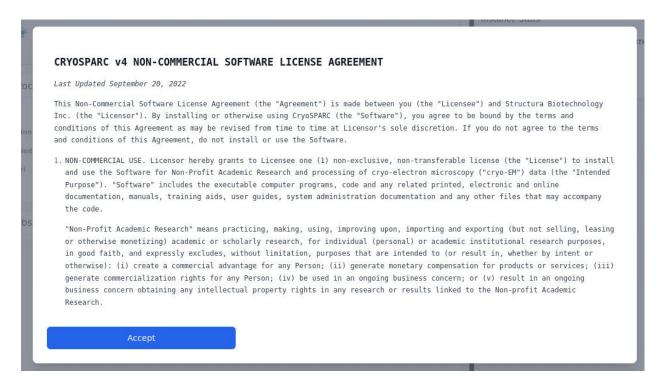


ACES CryoSPARC Running Job





Read and Accept the License

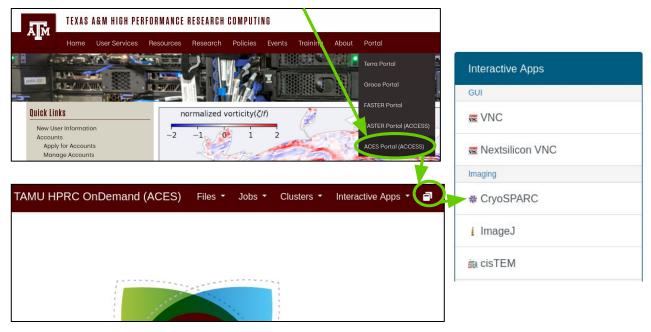


then exit and start a new CryoSPARC job this time with a GPU

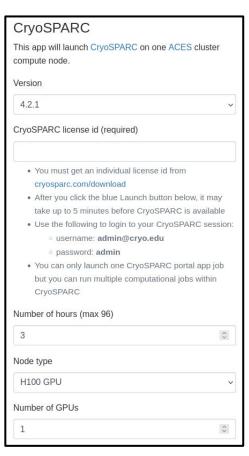


ACES CryoSPARC Portal App

portal-aces.hprc.tamu.edu



Delete your CryoSPARC "CPU only" job and launch a new CryoSPARC job using 1 x H100 GPU for the interactive tutorial

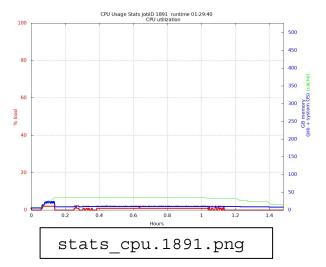


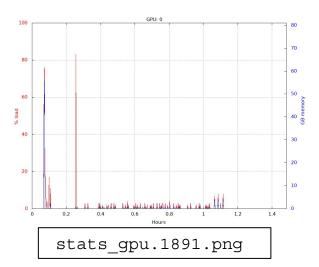
CryoSPARC Visualization of Resource Utilization

Review CPU and GPU usage for a Job

The **jobstats** utility automatically runs for each CryoSPARC portal job and monitors CPU and GPU resource usage and creates a graph for each.

- CPU stats monitors all cores regardless of how many were configured for the job
 - CryoSPARC jobs use a fraction of total cores and memory if selecting one of many GPUs on a compute node to allow users to request the other GPUs on the compute node
- GPU stats will create a graph for each GPU that was configured for the job



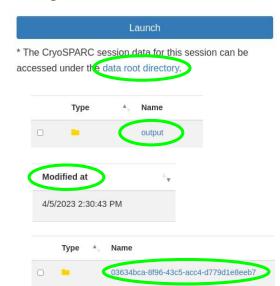


https://hprc.tamu.edu/kb/Software/useful-tools/jobstats



Review CPU and GPU usage for a CryoSPARC Job

- At the bottom of the CryoSPARC portal app, access the CryoSPARC data root directory
- 2. Click 'output'
- Click 'Modified at' to sort newest session on top
- 4. Click the Name of the session you want to view based on the date
- 5. Click the .png image files to view the CPU and GPU usage stats
 - a. CPU usage stats are for the entire GPU node so it will include any other jobs running on that node
 - b. GPU stats are exclusive to your job





3.

5.

Group Data Directories

- Group data directories can be used to share input files or to have a group work space
- Send a request to the HPRC helpdesk to create a group
 - o help@hprc.tamu.edu
 - provide a group name and usernames of members
 - group directories have their own disk quotas separate from individual users
- You will need to mount your group directory when launching the CryoSPARC portal app

Optional group directory to mount	
	You must already be a member of the group
•	Example values:
	∘ /junjiez
	 /scratch/group/davislab



https://hprc.tamu.edu

HPRC Helpdesk:

help@hprc.tamu.edu Phone: 979-845-0219 Take our short course survey!



https://u.tamu.edu/hprc_shortcourse_survey

Help us help you. Please include details in your request for support, such as, Cluster (ACES, FASTER, Grace, Launch), NetID (UserID), Job information (JobID(s), Location of your jobfile, input/output files, Application, Module(s) loaded, Error messages, etc), and Steps you have taken, so we can reproduce the problem.

CryoSPARC Tutorial presented by:

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Laboratory for Biomolecular
Structure and Dynamics

