

Introduction to Andes

OLCF Data Analysis and Visualization Cluster

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What is Andes for?

Andes prioritizes small jobs and schedules in such a way that there should always be nodes open for the work of smaller tasks

"Andes is a 704-compute node commodity-type linux cluster. The primary purpose of Andes is to provide a conduit for large-scale scientific discovery via pre/post processing and analysis of simulation data generated on Summit."





What type of tasks can I do on Andes?

Any type of ancillary analysis or processing work that does not require the large-scale leadership class mandate resources on Summit are ideal for running on Andes!

- **Preprocessing**: cleaning, concatenating, or reshaping large data, or otherwise preparing input for a large-scale run, etc.
- **Postprocessing**: running scripts to analyze, reshape, or restructure model output.
- Data Visualization: Running Vislt, Paraview, TurboVNC. Creating plots and videos using scripts written in R, Python, Julia, etc.
- Debugging, troubleshooting, and lots more!



System Overview: Connecting to Login Nodes

To connect to Andes, ssh to andes.olcf.ornl.gov using your OLCF username and Passcode (PIN+Tokencode)

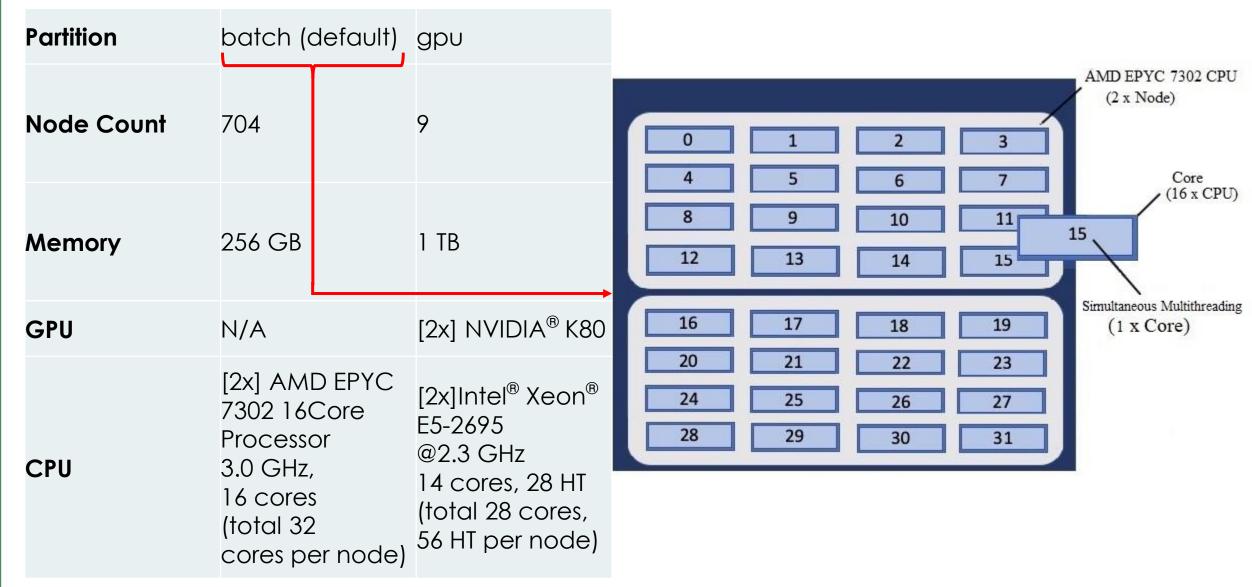
\$ ssh username@andes.olcf.ornl.gov

- There are 8 login nodes (identical to the batch partition compute nodes) which provide an environment for editing, compiling, and launching codes onto the compute nodes.
- All Andes users access the system through these login nodes, and as such, any CPU- or memory-intensive tasks on these nodes could interrupt service to other users.

PLEASE REFRAIN FROM DOING ANY ANALYSIS OR VISUALIZATION TASKS ON THE LOGIN NODES.



System Overview: Compute Nodes





System Overview: File System

- Users can access the moderate security enclave NFS, Alpine, and HPSS filesystems from Andes.
- These are the same filesystems accessible on Summit, making paired analysis from work on Summit more convenient.

	Area	Path	Permissions	Quota	Backups	Purged	Retention	On Compute Nodes
NFS -	User Home	/ccs/home/[userid]	User set	50 GB	Yes	No	90 days	Read-only
	Project Home	/ccs/proj/[projid]	770 (rwxrwx)	50 GB	Yes	No	90 days	Read-only
ſ	Member Work	/gpfs/alpine/[projid]/scratch/[userid]	<u>700 (rwx)</u>	50 TB	No	90 days	<u>N/A</u>	Read/Write
Alpine -	Project Work	/gpfs/alpine/[projid]/proj-shared	770 (rwxrwx)	50 TB	No	90 days	<u>N/A</u>	Read/Write
L	World Work	/gpfs/alpine/[projid]/world-shared	775 (rwxrwxr-x)	50 TB	No	90 days	<u>N/A</u>	Read/Write
ſ	Member Archive	/hpss/prod/[projid]/users/[userid]	700 (rwx)	100 TB	No	No	90 days	No
HPSS -	Project Archive	/hpss/prod/[projid]/proj-shared	770 (rwxrwx)	100 TB	No	No	90 days	No
	World Archive	/hpss/prod/[projid]/world-shared	775 (rwxrwxr-x)	100 TB	No	No	90 days	No



Shell and Programming Environment

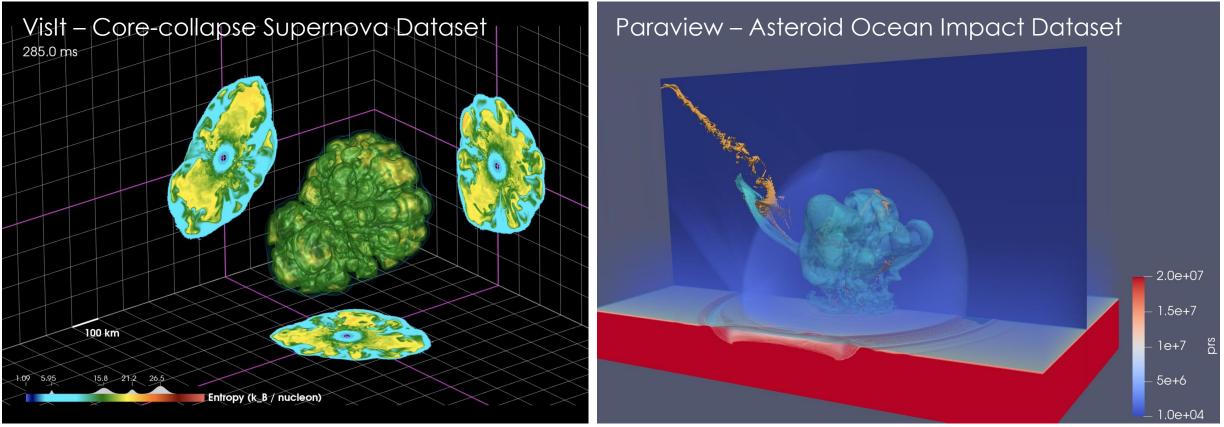
- Shells: bash (default), tsch, csh, ksh. You can request a change in shell by submitting a ticket to <u>help@olcf.ornl.gov</u>.
- Lmod: Use the module command to manage your modules

Command	Description
module -t list	Shows a terse list of the currently loaded modules.
module avail	Shows a table of the currently available modules
module spider <string></string>	Searches all possible modules according to <string></string>
module load <modulename> []</modulename>	Loads the given <modulename>(s) into the current environment</modulename>
module help <modulename></modulename>	Shows help information about <modulename></modulename>
module show <modulename></modulename>	Shows the environment changes made by the <modulename> modulefile</modulename>
module use <path></path>	Adds <path> to the modulefile search cache and MODULESPATH</path>
module unuse <path></path>	Removes <path> from the modulefile search cache and MODULESPATH</path>
module purge	Unloads all modules

Software: Type module avail to see all available software. A definitive guide to Python on Andes can be found in our docs.

Visualization with Paraview and Visit

VisIt and Paraview are interactive, parallel analysis and visualization tools for exploring large scientific datasets. Both are available on Andes



See the Software -> Visualization Tools section of the docs for in-depth guides on installation and use



Python on Andes

Andes is the best place to run Python

- It is the easiest place to install if you need customization
- It plays nicely with the Slurm scheduler

```
$ module avail python
------ /sw/andes/modulefiles/core -----
python/3.7-anaconda3
$ module load python
$ conda list
```

If a package you require isn't available, you can create and load a custom environment and install yourself.

Step by step instructions on custom environments and running python on Andes are in the docs under Software -> Python COAK RIDGE Wational Laboratory

Compilers

The **intel** (default), **pgi**, and **gcc** compilers are available on Andes, and can be loaded or changed using the module command

<pre>\$ module avail intel</pre>
/sw/andes/spack-envs/base/modules/site/Core/sw/andes/spack-envs/base/modules/site/Core
intel/19.0.3 (D) intel/2021.4.0
<pre>\$ module avail pgi</pre>
/sw/andes/spack-envs/base/modules/site/Core/sw/andes/spack-envs/base/modules/site/Core
pgi/19.10
<pre>\$ module avail gcc</pre>
/sw/andes/spack-envs/base/modules/site/Core/sw/andes/spack-envs/base/modules/site/Core
gcc/6.5.0 gcc/9.3.0 (D) gcc/10.1.0 gcc/10.3.0

The following wrapper programs are cognizant of your currently loaded modules:

- mpicc to invoke the c compiler
- **mpicc**, **mpicxx**, or **mpic++** to invoke the c++ compiler
- mpif77 or mpif90 to invoke appropriate versions of the fortran compiler



Compilers, cont.

When building threaded codes with Openmp, compiler-specific flags must be included to ensure a proper build.

For example, if compiling C code with the various compiler modules, add the following compiler specific flags:

- For pgi, add "-mp" to the build line.
 \$ mpicc -mp test.c -o test.x
 \$ export OMP_NUM_THREADS=2
- For gnu, add "-fopenmp" to the build line.
 \$ mpicc -fopenmp test.c -o test.x
 \$ export OMP NUM THREADS=2
- For intel, add "-qopenmp" to the build line.

\$ mpicc -qopenmp test.c -o test.x
\$ export OMP_NUM_THREADS=2



Running Jobs with Slurm

- Only a limited number of simultaneous basic tasks (editing files, compiling codes) should be done on the login nodes. For compute- or memory-intensive tasks, users must submit as jobs to the compute nodes.
- Unlike Summit, Andes uses the Slurm batch scheduler.

Task	LSF (Summit)	Slurm
View batch queue	jobstat	squeue
Submit batch script	bsub	sbatch
Submit interactive batch job	bsub -Is \$SHELL	salloc
Run parallel code within batch job	jsrun	srun

- There are two ways to submit to the computes:
 - 1. Request nodes for an interactive batch job with salloc
 - 2. Write and submit a batch script with **sbatch**

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Interactive Jobs

• To submit an interactive job, use the **salloc** command:

\$ salloc -A abc123 -p gpu -N 4 -t 1:00:00	0 3
	-
• The job will wait for nodes to be	-
available. An interactive promp	,† -
will appear once the job starts.	-

salloc	Start an interactive session
-A	Charge to the abc123 project
-p gpu	Run in the gpu partition
-N 4	request (4) nodes
-† 1:00:00	for (1) hour

• Use the **sbatch** --**test-only** command to see when a job of a specific size could be scheduled:

\$ sbatch --test-only -N2 -t1:00:00 batch-script.slurm

sbatch: Job 1375 to start at 2022-08-29T10:54:01 using 64 processors on nodes andes[499-500] in partition batch



Writing a Batch Script

Consider the following batch script

<pre>#!/bin/bash #SBATCH -A ABC123 #SBATCH -J RunSim123 #SBATCH -o %x-%j.out #SBATCH -e %x-%j.err #SBATCH -t 1:00:00 #SBATCH -t 1:00:00 #SBATCH -p batch #SBATCH -N 128 #SBATCH -N 128 #SBATCHmail-type=ALL #SBATCHmail-type=ALL</pre>	<pre>Interpreter line -A - Project to charge -J - Job Name -o - stdout, %x=job_name, %j=job_number -e - stderr -t - walltime in HH:MM:SS -p - partition (queue) -N - number of nodes Email notifications (ALL, BEGIN, END, etc.) Email address to notify</pre>
cd \$MEMBERWORK/abc123/Run.456	Change into the run directory
srun -N # -n # -c # -cpu-bind=cores ./a.out	Run the job

Where the srun command takes the following common options: -N [#nodes] -n [#ntasks] -c [#cores_per_task] -cpu-bind=[cores,threads,no]

Submit your batch script on the command line with sbatch myscript.slurm



Monitoring and Modifying Jobs

Command	Action/Task	LSF Equivalent
squeue	Show the current queue	bjobs
sinfo	Show node/partition info	bqueues or bhosts
sacct	Show queued and historical job info. and steps	bacct
scancel	Cancel a job or job step	bkill
scontrol	View or modify job configuration.	bstop, bresume, bmod

To see all jobs currently in the queue:

\$ squeue −1

To see all of your queued jobs: \$ squeue -1 -u \$USER

To see all your queued jobs (plus steps) in the queue **\$ sacct -u \$USER**

To see all steps submitted to job 123456: **\$ sacct -j 123456** To cancel a job in the queue: \$ scancel -j <job_id_#>

To place a queued job on hold: \$ scontrol hold <job_id_#>

To release a held job: \$ scontrol release <job_id_#>

To see additional details for a job: \$ scontrol show job <job_id_#>



Queue Policies

There are two compute node partitions: "batch" and "gpu". Each has different queue policies for the number of jobs that can be run and wall times per number of nodes requested.

Batch Partition:

Bin	Node Count	Duration	Policy
А	1 - 16 Nodes	0 - 48 hr	
В	17 - 64 Nodes	0 - 36 hr	max 4 jobs running and 4 jobs eligible per user in bins A, B, and C
С	65 - 384 Nodes	0 - 3 hr	

GPU Partition:

Node Count	Duration	Policy
1-2 Nodes	0 - 48 hrs	max 1 job running per user

Note: to access the gpu partition, specify -p gpu in your script or interactive node request.



Questions?

• Full documentation for using Andes can be found here: <u>https://docs.olcf.ornl.gov/systems/andes_user_guide.html</u> <u>https://docs.olcf.ornl.gov/software/python/index.html</u>

• Trouble? Submit a help-desk ticket to <u>help@olcf.ornl.gov</u>



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