

Getting Started with the Arkansas Research Platform

SAU Campus
February 20th

Day 1

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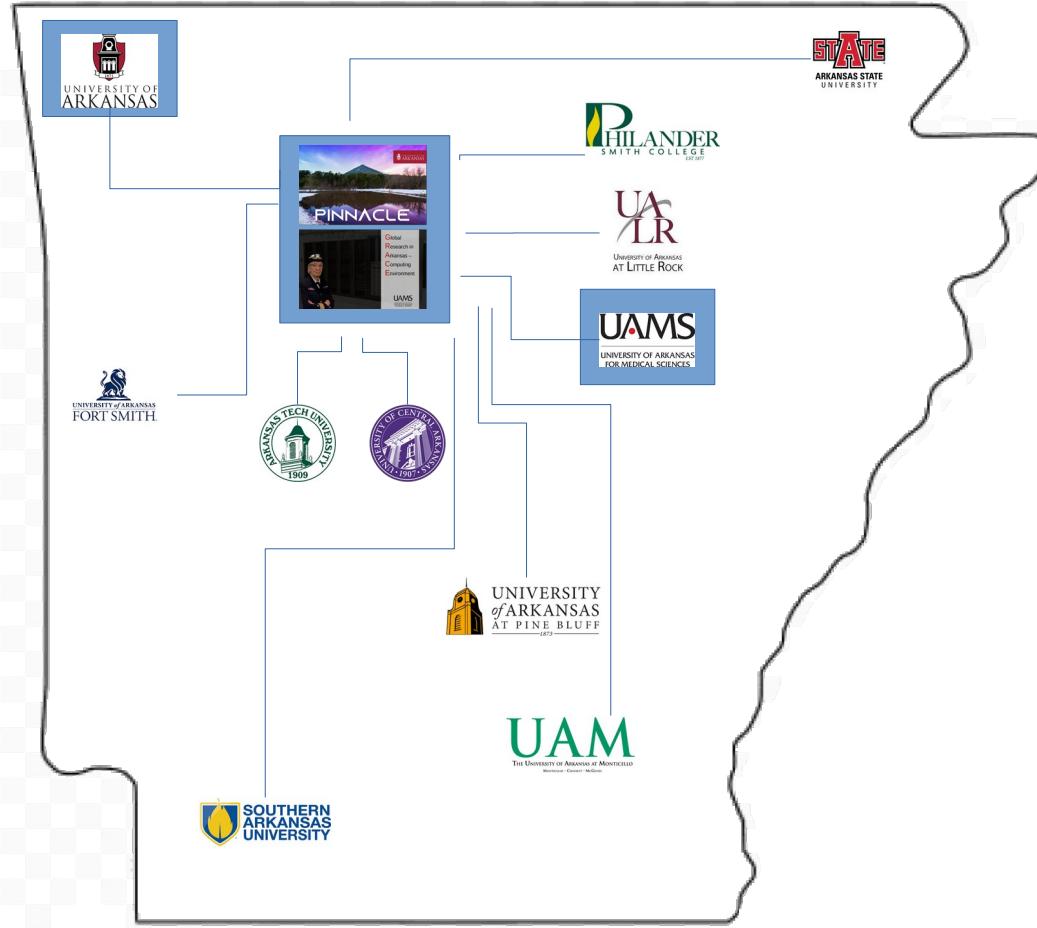
Day 1 Schedule

8:30 am	Registration & Set-Up Check with each participant to ensure ability to connect to internet Make sure all required programs have been downloaded and installed
9:00 am	Welcome & Introductions Brief introductions: Name, institution, 1-sentence research interests
9:15 am	HPC Overview High Performance Computing - explained in 30 min
9:45 am	What is Arkansas Research Platform (ARP) ? Brief description of ARP hardware resources When to use HPC resources Programs that can leverage HPC resources Getting familiar with the vocabulary Cluster Functional diagrams
10:15 am	The Essentials Getting an account on Pinnacle / Grace Logging into Open OnDemand portal OOD Overview
10:45 am	Running Jobs - Open Ondemand Batch job (run, modify, rerun) Interactive jobs (Jupyter notebook, VMD)
11:45 am	LUNCH
12:45 pm	Moving Data Open Ondemand File Upload/Download Terminal access in Open OnDemand wget, scp, rsync, FileZilla (GUI)
2:00 pm	Globus Log in/create Globus ID account Globus endpoints Transfer data between Pinnacle and Public endpoint Install Globus Connect Personal Transfer data between Pinnacle and personal device
4:00 pm	Reflection and Closing Remarks for Day 1 Where are you stuck?

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Arkansas Research Platform



Arkansas Research Platform is a collaboration of higher education institutions within the state of Arkansas which provides computing resources and data storage to all its members (free of charge to students, faculty and staff). The core components are the **Pinnacle cluster** managed by the Arkansas High Performance Computing Center (UAF, Fayetteville) and the **Grace cluster** managed by UAMS High Performance Computing Center (UAMS, Little Rock).

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Pinnacle Cluster - Fayetteville



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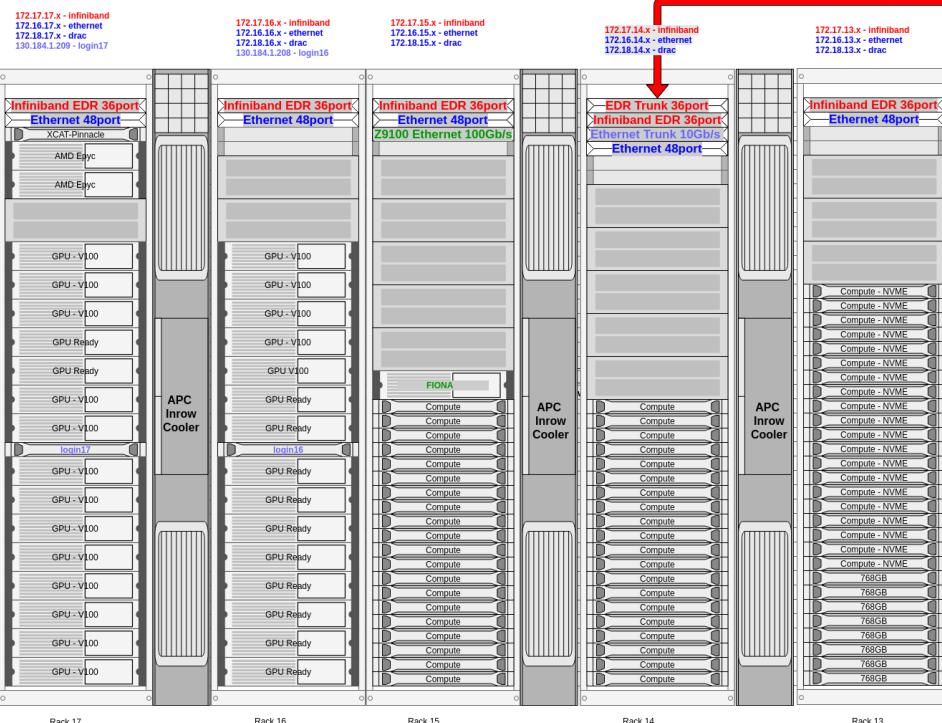
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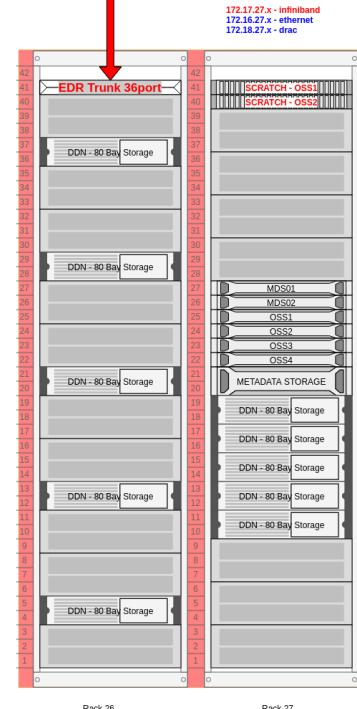
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Pinnacle Cluster - Fayetteville

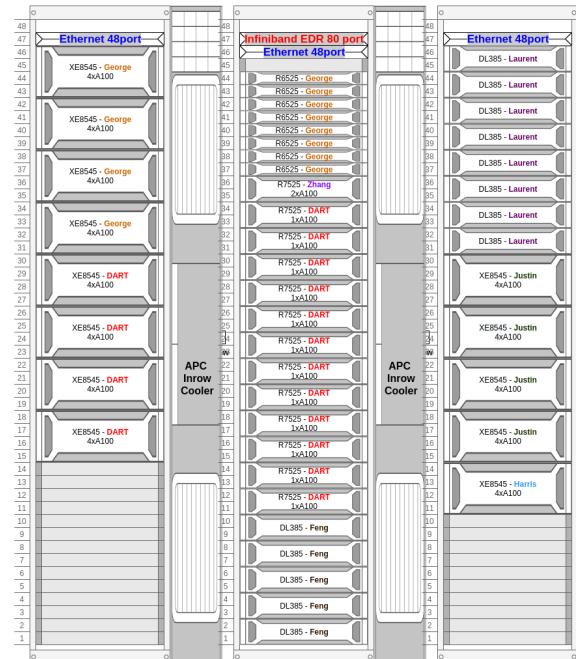
Pinnacle Cluster



Lustre Storage



Pinnacle 2 - DART Expansion



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Pinnacle Cluster – Fayetteville

Compute Nodes

100 2x Intel Xeon 6130, 2.1GHz, 32cores, 192GB RAM
50 2x AMD Epyc 7543, 2.8GHz, 64cores, 1024GB RAM
248 other/misc configuration

GPU equipped Nodes: 56

13 4x Nvidia A100
1 2x Nvidia A100
18 1x Nvidia A100
1 2x Nvidia V100
20 1x Nvidia V100
1 4x Nvidia TitanV
2 4x Nvidia K80

Large Memory Nodes (>1TB): 4

3 2TB RAM
1 3TB RAM

Storage

2.3 PB main storage (DDN SFA14k exascaler)
0.02PB scratch storage (HPE nvme/lustre)
1 PB archive storage (Dell/lustre)
2 PB object store (Dell/EMC ECS)

Network

Internal 100Gbps Infiniband interconnect
External 100Gbps Ethernet connection

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Pinnacle Cluster – Fayetteville

Node Count		CPUs		Memory	GPUs
124		AMD a6136	32cores	64GB	None
70		Intel i6130	32cores	192GB	None
55		Intel i2650v2	16cores	64GB	None
19		Intel i6130	32cores	192GB	1xNVidia V100
18		AMD a7543	64cores	1024GB	1xNVidia A100
14		Intel i2650v4	24cores	128GB	None
13		AMD a7543	64cores	1024GB	4xNVidia A100
12		Intel i2620	8cores	32GB	None
10		Intel i6230	40cores	384GB	None
8		AMD a7543	32cores	512GB	None
8		AMD a7543	64cores	256GB	None
7		Intel i6128	24cores	768GB	None
5		AMD a7543	64cores	512GB	None
3		AMD o6378	64cores	512GB	None
3		Intel i2667v4	16cores	256GB	None
3		AMD a7543	32cores	1024GB	None
2		Intel i2650v2	16cores	128GB	4xNVidia K80
2		AMD a7351	32cores	256GB	None
2		Intel a7110	64cores	112GB	None
1		Intel i4166	24cores	192GB	4xNVidia TitanV
1		Intel i6130	32cores	768GB	2xNVidia V100
1		AMD a7543	64cores	512GB	2xNVidia A100
1		Intel i6230	40cores	192GB	1xNVidia V100
1		Intel i2620	8cores	92GB	None
1		Intel i2620	8cores	8GB	None
1		Intel i2620	8cores	82GB	None
1		Intel i4627v2	32cores	768GB	None
1		Intel i2620	8cores	72GB	None
1		Intel i2620	8cores	62GB	None
1		Intel i2620	8cores	52GB	None
1		Intel i2650v4	24cores	512GB	None
1		Intel i2620	8cores	42GB	None
1		Intel i4860v2	96cores	3072GB	None
1		AMD a7402	48cores	256GB	None
1		Intel i4627v2	128cores	2048GB	None
1		AMD a7543	64cores	2048GB	None
1		AMD a7402	48cores	2048GB	None
1		Intel i6128	24cores	192GB	None
1		AMD a7452	64cores	1024GB	None

Total #of compute nodes: 398
Total #of cores: 12,912

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Grace Cluster – Little Rock



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Grace Cluster – Little Rock

Compute Nodes

96	2x Intel Xeon E5-2680v4, 2.40GHz, 56cores, 128GB RAM
80	Intel Xeon Phi 7230F, 1.30GHz, 64 cores, 384GB RAM
16	Intel Xeon Phi 7230F, 1.30GHz, 64 cores, 192GB RAM
3	2x Intel Xeon E5-2680v3, 2.50GHz, 48cores, 128GB RAM, 2x Nvidia P100 GPUs
1	2x Intel Xeon E5-2680v3, 2.50GHz, 48cores, 128GB RAM, 2x Nvidia K40 GPUs

Other Nodes

7	management/login/storage interface
---	------------------------------------

Storage

1.9PB	high speed storage (DDN GS14KX GridScaler, IBM Spectrum Scale 5)
1.0PB	NAS intermediate storage
4.2PB	object store (Dell/EMC ECS)

Network

Internal	100Gbps Omnipath interconnect
External	redundant 10Gbps Ethernet connections

Total #of compute nodes: 196

Total #of cores: 11,712

Coming soon:

10 additional nodes, 640 AMD Epyc cores, 9 A100 GPUs, 100Gbs ethernet

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When to use HPC resources

- ? - word processing
- ? - editing spreadsheets
- ? - editing images
- ? - plotting graphs
- ? - storing personal images
- ? - storing personal movies
- ? - bitcoin mining
- ? - playing multi-player online games
- ? - GPU accelerated programs
- ? - machine learning
- ? - running parallel multi-node simulations
- ? - visualizing large data sets
- ? - moving large data sets
- ? - processing large data sets

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When to use HPC resources

- NO - word processing
- NO - editing spreadsheets
- NO - editing images
- NO - plotting graphs
- NO - storing personal images
- NO - storing personal movies
- NO - bitcoin mining
- NO - playing multi-player online games
- YES - GPU accelerated programs
- YES - machine learning/AI
- YES - parallel multi-node simulations
- YES - visualizing large data sets
- YES - moving large data sets
- YES - processing large data sets

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HPC Applications

Software on Pinnacle

Molecular dynamics: namd, lammps

Quantum chemistry: quantum espresso, VASP, Gaussian, PQS, NWChem

Multiphysics: COMSOL, Ansys

Data processing programming languages: Python, R

Simulation frameworks: Mathematica, Matlab

Statistics: SAS

Bioinformatics: abyss, bamtools .. spades, samtools, trinity...
[\(Bioinformatics Software on Pinnacle\)](#)

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HPC Vocabulary

cluster - multiple computers connected by a network, with access to the same file system.

node - a single computer

login node - a gateway node to the cluster. Shared by all users. No computationally intensive tasks are permitted

compute node - node which can run computationally intensive tasks

queueing system - software which organizes and schedules jobs submitted by users to be executed on compute nodes

job - a computational task submitted by a user to the queueing system

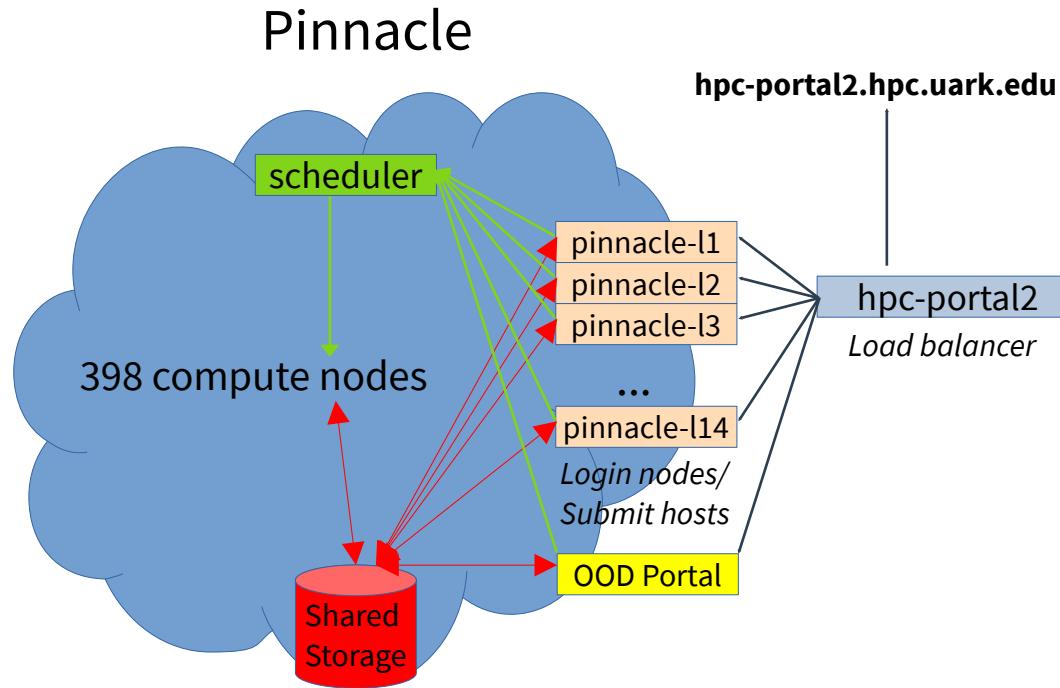
cpu/core/thread/socket - central processing unit (CPU) is a microprocessor on a single integrated circuit with 2 or more processing units (cores). Core can run a single instruction at a time. With Hyperthreading the same core can run 2 instructions at once. Socket is an interface on the motherboard which receives a CPU. HPC motherboards usually have 2 - 4 sockets.

[HPC Vocabulary - fuller list](#)

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Pinnacle Cluster – functional diagram



scheduler – coordinates all of the nodes in the queue (SLURM queueing system)

submit hosts – nodes from which users submit jobs to the queues

login hosts – nodes accessible from the external network

compute nodes – nodes assigned to queues and run jobs

Getting Accounts on ARP (Pinnacle and Grace)

Pinnacle (UAF, Fayetteville)



Account application: <https://hpc.uark.edu/hpc-support/user-account-requests>
Access: https://hpcwiki.uark.edu/doku.php?id=off-campus_access_dmz
Website: <https://hpc.uark.edu/>

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Web Portal - Open Ondemand

Open OnDemand (OOD) is a web portal interface to HPC resources, which hides the complexities of the HPC scheduling system and Linux/bash command line interface. Also provides a convenient method of displaying desktop/GUI applications running on compute nodes on the users client desktop.

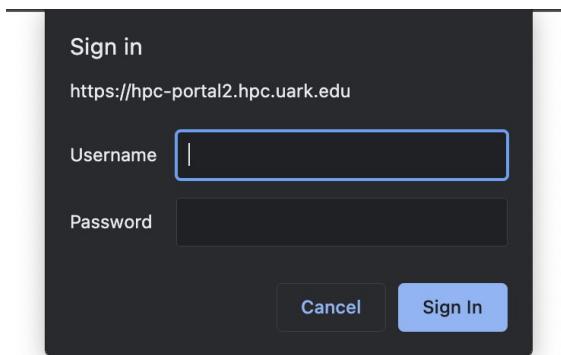
Pinnacle OOD portal

<https://hpc-portal2.hpc.uark.edu/>

Access from outside of UAMS campus network: I

https://hpcwiki.uark.edu/doku.php?id=off_campus_access_dmz

Usernames and passwords supplied in the login instruction emails when Pinnacle accounts were created.



The image shows a dark-themed sign-in form. At the top, it says "Sign in" and displays the URL "https://hpc-portal2.hpc.uark.edu". Below the URL are two input fields: "Username" and "Password". Both fields have a light blue border. At the bottom of the form are two buttons: "Cancel" on the left and "Sign In" on the right, both in white text.

Open Ondemand Portal – Pinnacle

The screenshot shows the OnDemand Pinnacle Portal dashboard. At the top, there's a navigation bar with the AHPCC logo, a search bar containing 'hpc-portal2.hpc.uark.edu/pun/sys/dashboard', and user information 'Logged in as pwolinski' and 'Log Out'. Below the navigation is a banner featuring a landscape image with the word 'PINNACLE' overlaid.

On the left side, there's a portrait of a man in a naval uniform and a banner for 'Global Research in Arkansas – Computing Environment' associated with UAMS (University of Arkansas for Medical Sciences).

On the right side, there's a banner for 'TRESTLES' showing a large wave, and another banner for 'KARPINSKI' showing a server rack.

At the bottom, a footer states: 'OnDemand provides an integrated, single access point for all of your HPC resources.' It also includes the OnDemand logo ('powered by OPEN OnDemand') and the text 'OnDemand version: v1.8.18'.

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Open Ondemand Portal – First HPC job

Submit batch job

Top Menu: Jobs→Job Composer

- +New Job button
- Edit Files button
- Select 'myjob.sh'
- View button

```
#!/bin/bash
# JOB HEADERS HERE

echo "Hello World"
```

- Close browser tab
- Submit button

View results

Wait for “completed status”

- Edit Files button
- Select 'slurm-?????.out'
- View button

Hello World

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Open Ondemand Portal – First HPC job

Modify job script and resubmit

- Edit Files button
- Select 'myjob.sh'
- Edit button

```
#!/bin/bash
# JOB HEADERS HERE

echo "Hello World"
echo "running on host"
hostname
echo "on "
date
```

View Results

Wait for "completed status"

- Edit Files button
- Select 'slurm-?????.out'
- View button

```
Hello World
running on host
c1405
on
Sun Sep 25 19:21:58 CDT 2022
```

- Save button
- Close browser tab
- Submit button

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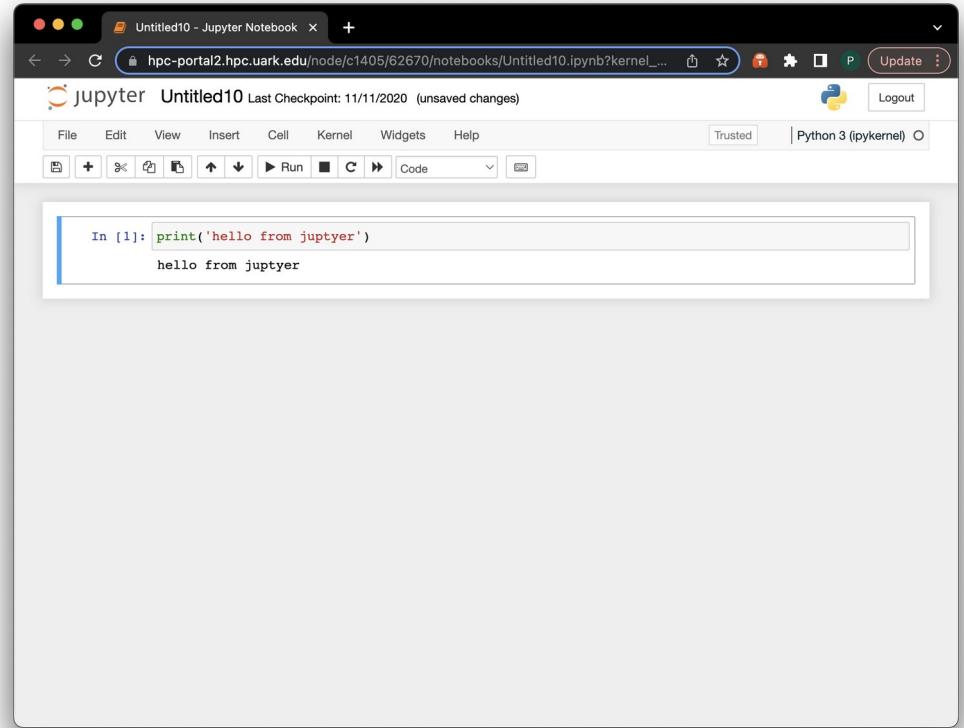
Open Ondemand Portal – interactive job

Interactive Jupyter Notebook job

- Top Menu: Interactive Apps → Jupyter Notebook
- Select “comp01, 1 hour, 1 node”
- Launch button
- Wait for the status “Running”
- Connect to Jupyter button

Interact with the Notebook

- New button → Python3 (ipykernel)
- Type in “print('hello')” into In [] field
- <CTRL>-Enter to run



The screenshot shows a Jupyter Notebook interface with a single cell containing the code "print('hello from jupyter')". The output of the cell is "hello from jupyter". The notebook has a title "Untitled10" and a subtitle "Last Checkpoint: 11/11/2020 (unsaved changes)". The toolbar includes File, Edit, View, Insert, Cell, Kernel, Widgets, Help, and a Python 3 (ipykernel) option.

```
In [1]: print('hello from jupyter')
hello from jupyter
```

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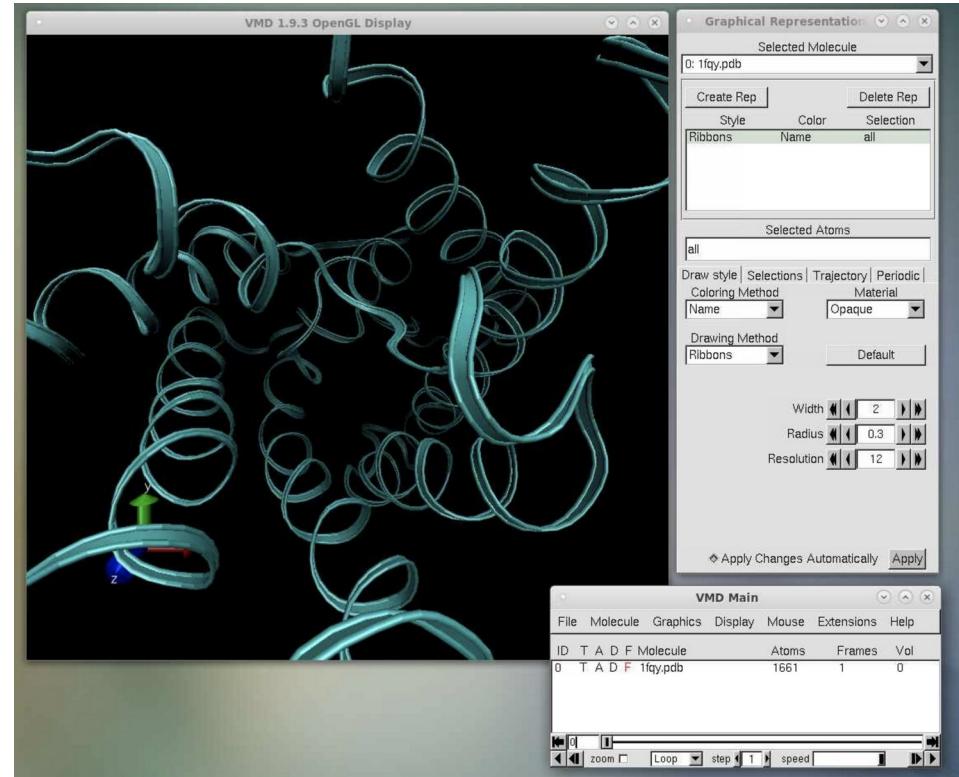
Open Ondemand Portal – interactive job

Interactive VMD job

- Top Menu: Interactive Apps → VMD
- Select “Queue: comp01, Number of hours: 1”
- Launch button
- Wait for the status “Running”
- Launch VMD button

Interact with VMD

- At the VMD prompt Type: “menu main on”
- In the menu “File→New Molecule”
- In the “Molecule File Browser”, “File Name”:
`/scrfs/ahpcc/ARP-workshop/1fqy.pdb`
- Load button
- In the menu
“Graphics→Representations:”
Drawing method: Ribbons



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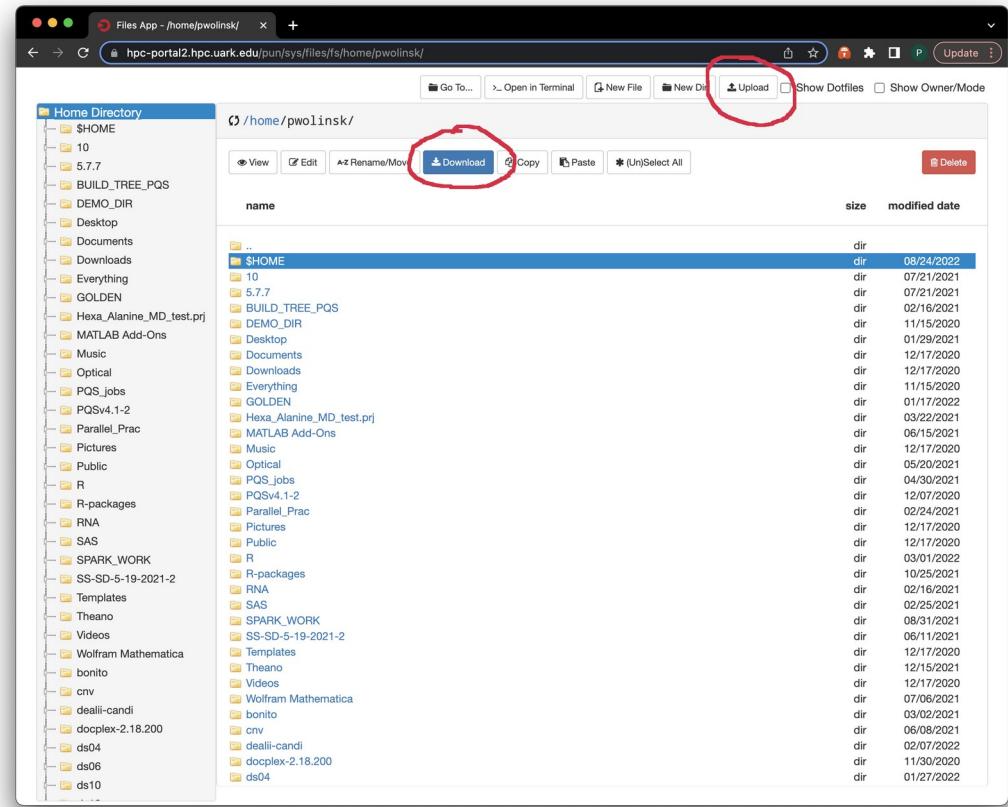
Open Ondemand Portal – Moving Data

File dialog window

- Top Menu: **Files** → **Home Directory**
New browser tab opens with the contents of your home directory
- **Upload** button – select file from your local machine and upload it to Pinnacle
- **Download** button – select file from your Pinnacle account and download it to your local machine

Using those two buttons:

- 1) download file etc/redhat-release to your laptop
- 2) Upload any text file to your home directory on Pinnacle



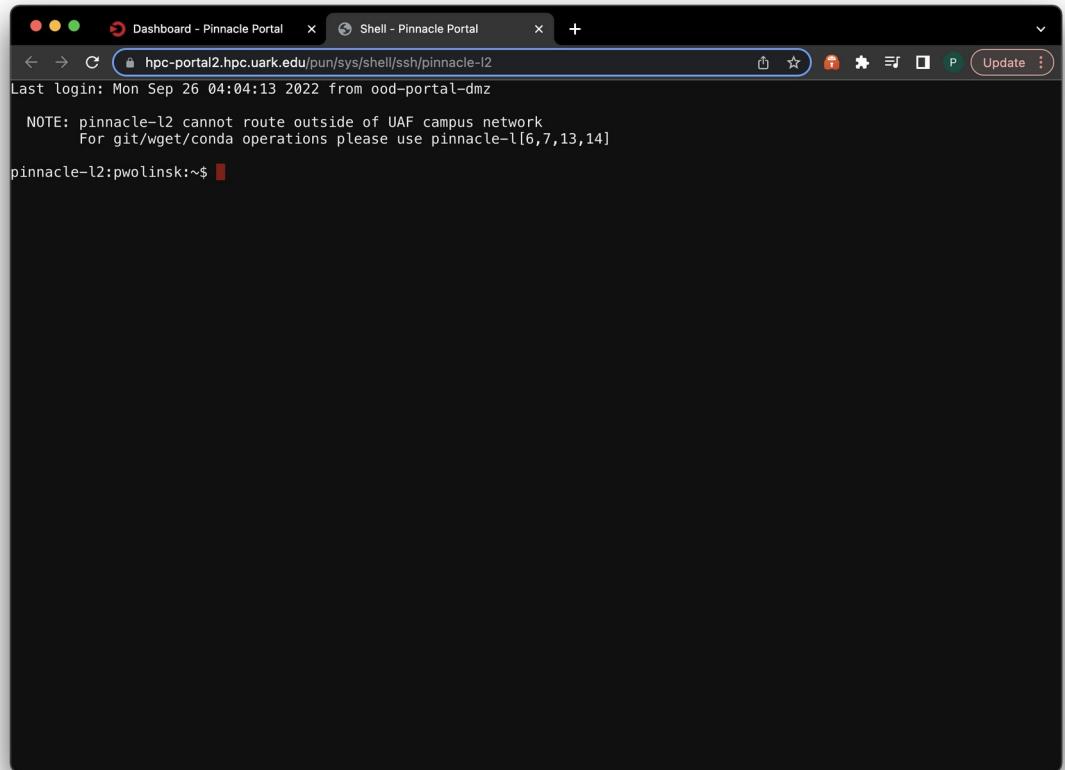
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Open Ondemand Pinnacle – Terminal

Terminal

- Top Menu: **Clusters** → **Pinnacle**
New browser tab opens with a terminal on a Pinnacle login node

OOD Portal on Pinnacle: <https://hpc-portal2.hpc.uark.edu>



The screenshot shows a web browser window with two tabs. The left tab is titled "Dashboard - Pinnacle Portal" and the right tab is titled "Shell - Pinnacle Portal". Both tabs have the URL "hpc-portal2.hpc.uark.edu/pun/sys/shell/ssh/pinnacle-l2".

The content area of the "Shell" tab shows a terminal session:
Last login: Mon Sep 26 04:04:13 2022 from ood-portal-dmz

NOTE: pinnacle-l2 cannot route outside of UAF campus network
For git/wget/conda operations please use pinnacle-l[6,7,13,14]
pinnacle-l2:pwolinsk:~\$ █

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Moving Data – Pinnacle/Grace - rsync

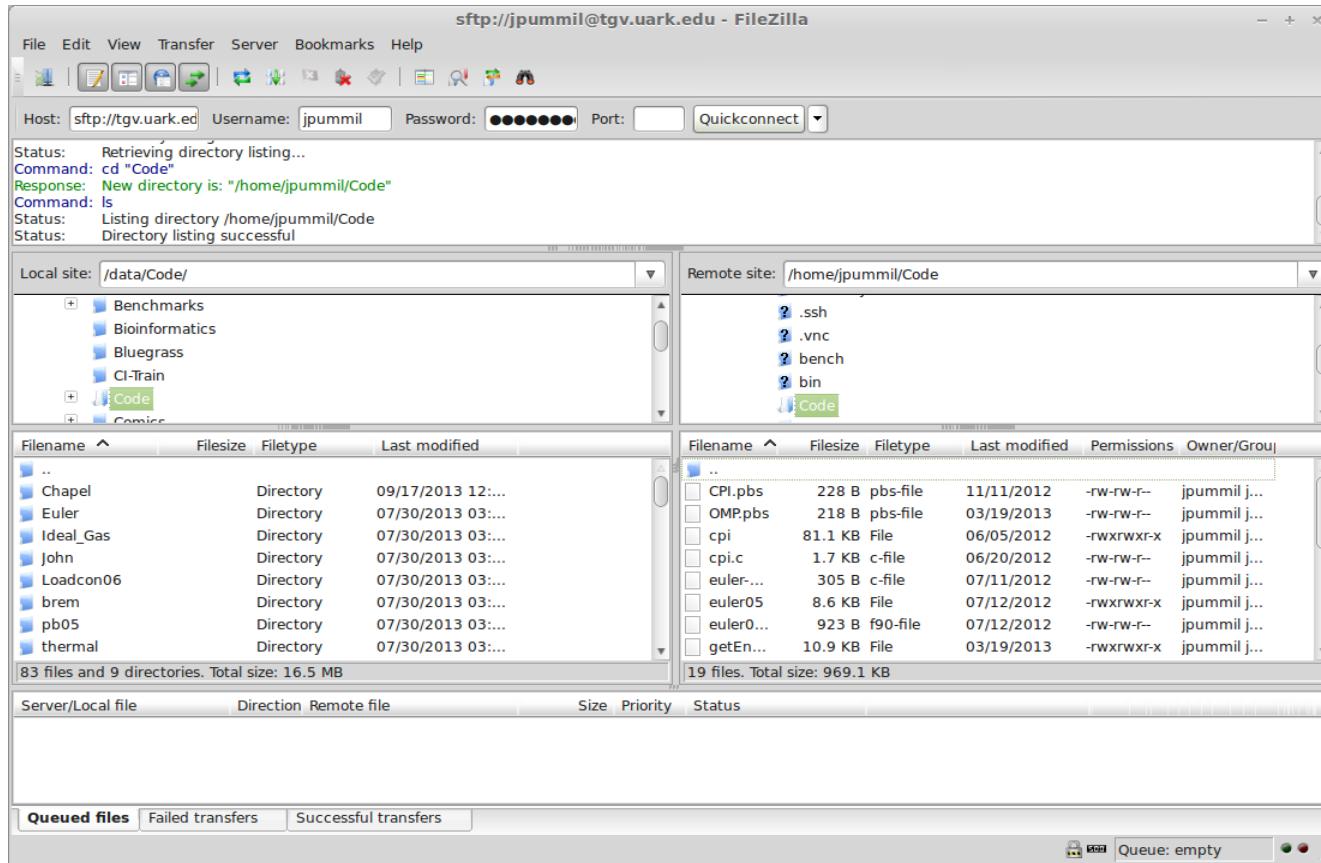
rsync - synchronize files between local and remote hosts. Only moves files which have been updated.

```
rsync -avz <source> <destination>
```

-avz options stand for archive, verbose, compress during transfer. Either source or destination can be of from
username@host:file_path

```
On Grace [login1]
[pwolinsk@login1 ~]$ scp -r slurm-pinnacle/ pwolinsk@hpc-portal12.hpc.uark.edu:
cgroup.conf.example
first-slurm-startup.sh
gres.conf
prolog
...
epilog
slurmdbd.conf.example
cgroup.conf
[pwolinsk@login1 ~]$ echo "some text" >> slurm-pinnacle/prolog
[pwolinsk@login1 ~]$ rsync -avz slurm-pinnacle pwolinsk@hpc-portal12.hpc.uark.edu:
sending incremental file list
slurm-pinnacle/
slurm-pinnacle/cgroup.conf
...
slurm-pinnacle/slurm.conf.example
slurm-pinnacle/slurmdbd.conf.example
slurm-pinnacle/layouts.d/
slurm-pinnacle/layouts.d/power.conf.example
slurm-pinnacle/layouts.d/power_cpufreq.conf.example
slurm-pinnacle/layouts.d/unit.conf.example
sent 1,058 bytes received 1,310 bytes 4,736.00 bytes/sec
total size is 118,072 speedup is 49.86
[pwolinsk@login1 ~]$
```

Moving Data – Pinnacle/Grace – GUI



<https://filezilla-project.org/download.php>

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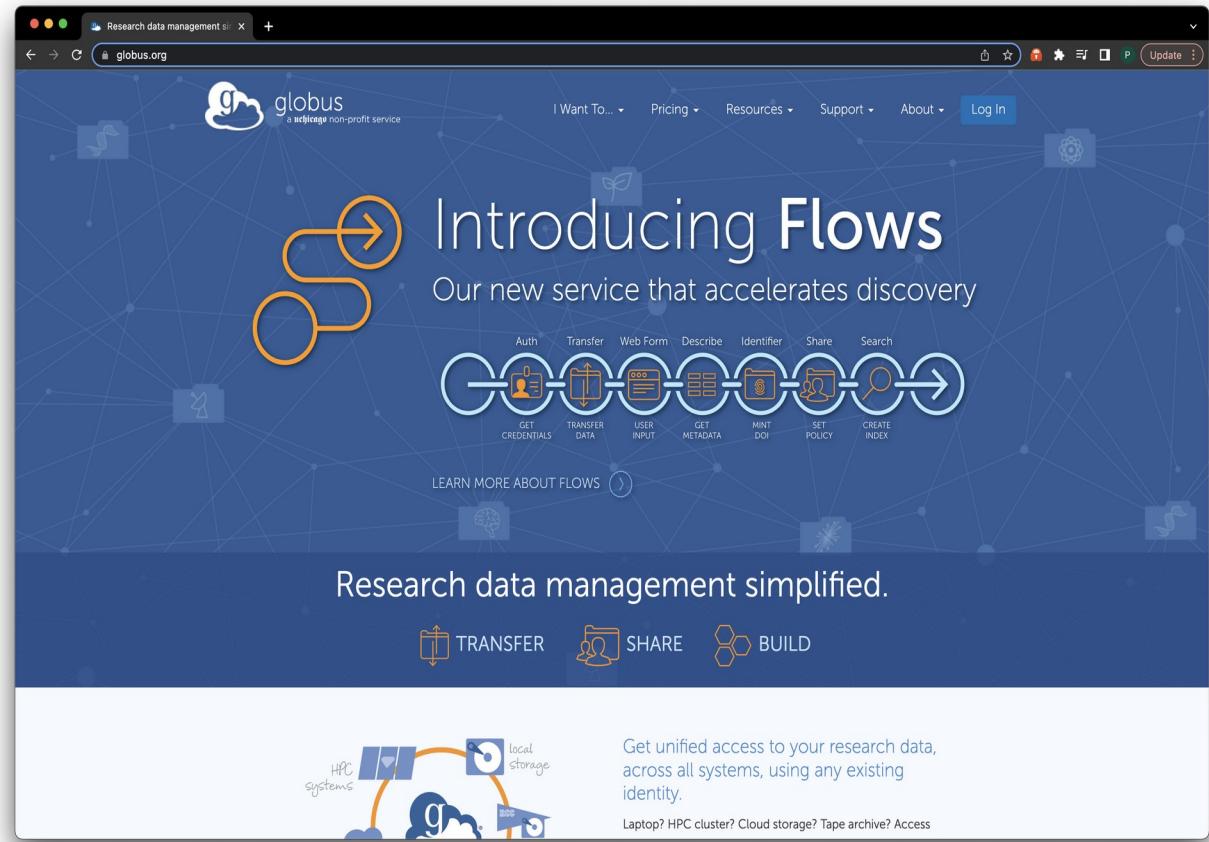
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Moving Data – Large Data Transfers - Globus

Globus – efficient, large data transfer and data sharing service.

Process:

1. log into Globus
2. connect to two Globus end points
3. transfer data between end points



<https://globus.org>

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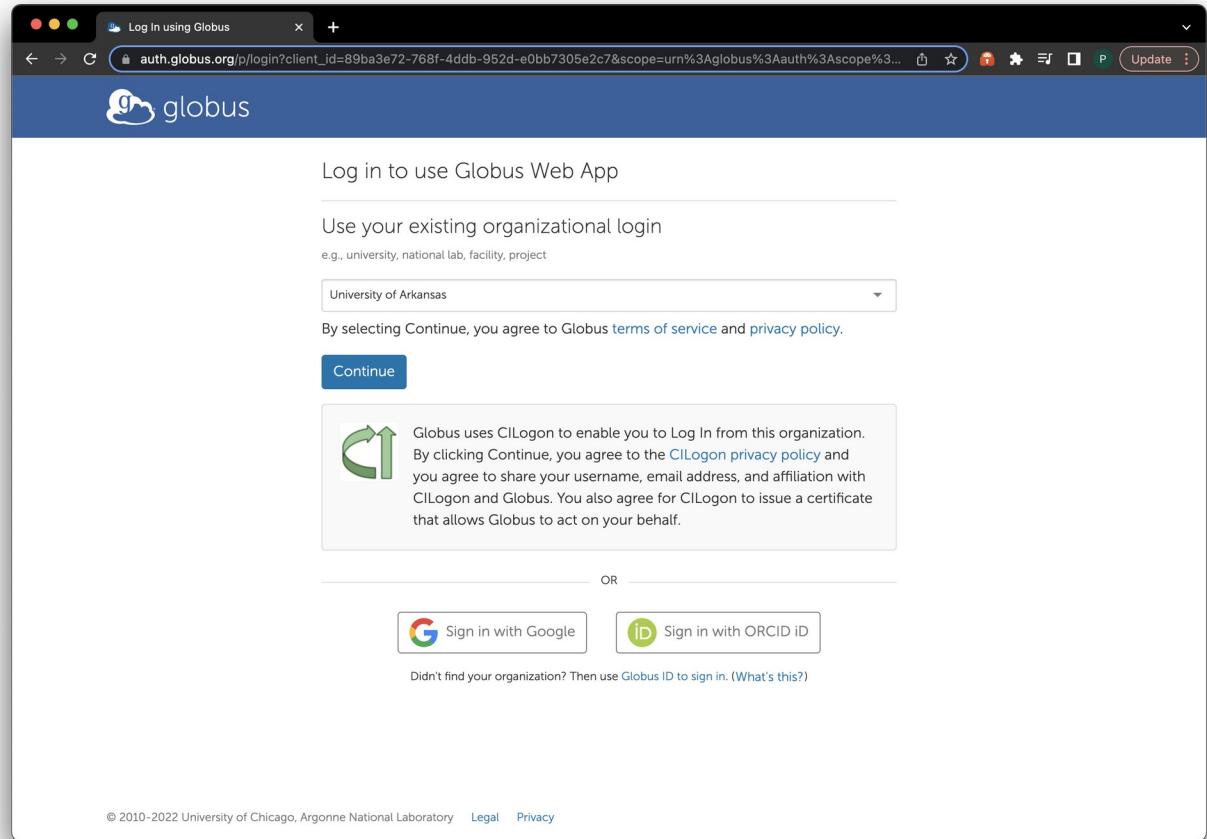
ARP

Moving Data – Large Data Transfers – Globus – Log in

If your institution is listed in the drop down menu, login with those credentials.

Google accounts and ORCID ID accounts are an option. Otherwise create a Globus ID account (link at the bottom of page).

In addition to other services, Globus is also an identity provider and maintains its own user accounts.



<https://globus.org>

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Moving Data – Large Data Transfers – Globus – Log in

Select Globus End Point for Pinnacle: UARK-Pinnacle

1. open File Manager
2. In the Collection text box: **UARK-Pinnacle**
3. log into the UARK-Pinnacle end point using your Pinnacle credentials

The image shows three overlapping browser windows:

- Top Window:** "File Manager | Globus" (app.globus.org/file-manager/collections). It displays a sidebar with "FILE MANAGER", "BOOKMARKS", "ACTIVITY", "COLLECTIONS", "GROUPS", "FLOWS", "ACCOUNT", and "LOGOUT". The main area shows a collection named "UARK-Pinnacle" with a description: "Owner: rwojinski@uark.edu Description: Pinnacle Cluster". Below it is another entry: "JEFF-UARK" with "Owner: pwojinski@uark.edu". A message at the bottom says "Please authenticate to access UARK-Pinnacle" with "Continue" and "Cancel" buttons.
- Middle Window:** "MyProxy Delegation Service" (globus.uark.edu/oauth/authorize?auth_token=myproxy%3aoa4mp%2C2012%3A%2FtempCred%2F0909a35bd8d3c8ff...). It features the AHPCC logo and the text "Arkansas High Performance Computing Center".
- Bottom Window:** "MyProxy Client Authorization" (globus.uark.edu/oauth/authorize?auth_token=myproxy%3aoa4mp%2C2012%3A%2FtempCred%2F0909a35bd8d3c8ff...). It contains the text "MyProxy Client Authorization" and "Welcome to the OAuth for MyProxy Client Authorization Page. The Client below is requesting access to your account. If you approve, please sign in with your username and password." It has fields for "Client Information" (Name: Globus, URL: https://www.globus.org), "Username" (pwojinski), "Password" (redacted), and "Sign In" and "Cancel" buttons.

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Moving Data – Large Data Transfers – Globus

Select another end point: Ambs_genomics

1. In the other Collection text box:

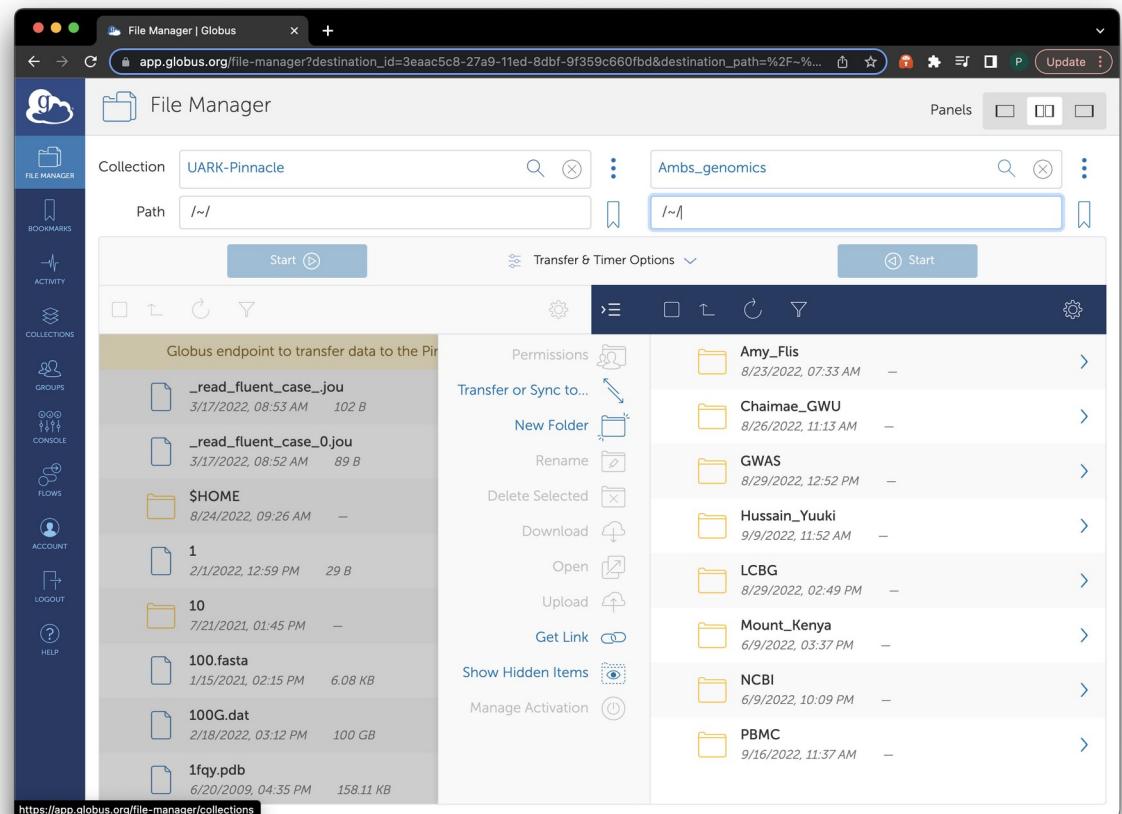
Ambs_genomics

2. double click on NCBI folder

3. select file “README”

4. Start button

When the data transfer is complete, you'll see the README file on the UARK-Pinnacle side (and in your HOME directory on Pinnacle)



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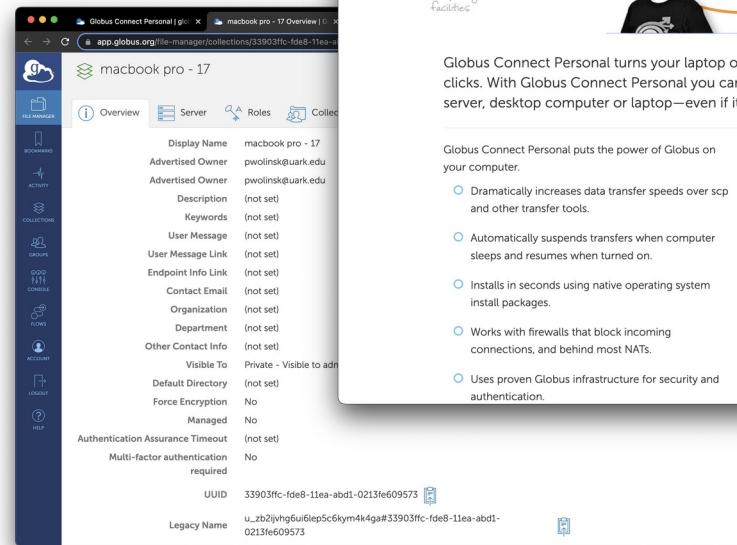
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Globus – Creating a Personal Endpoint

To move data using Globus between your personal device and any Globus end point you have access to, you have to install Globus Connect Personal server on your personal device.

It's a server that runs on your local machine and registers with your Globus account. Once set up and manually started, a new endpoint for your device will be available to you (only you) in the Globus collections.

A screenshot of the official Globus Connect Personal landing page at globus.org/globus-connect-personal. The page features a central image of a smiling woman with a laptop, surrounded by icons representing various cloud services like storage, sync, and scientific instruments. Text on the page explains the service: "Globus Connect Personal turns your laptop or other personal computer into a Globus endpoint with just a few clicks. With Globus Connect Personal you can share and transfer files to/from a local machine—campus server, desktop computer or laptop—even if it's behind a firewall and you don't have administrator privileges." Below this, there are two columns of bullet points and links for installation instructions.

<ul style="list-style-type: none">Dramatically increases data transfer speeds over scp and other transfer tools.Automatically suspends transfers when computer sleeps and resumes when turned on.Installs in seconds using native operating system install packages.Works with firewalls that block incoming connections, and behind most NATs.Uses proven Globus infrastructure for security and authentication.	<p>Globus Connect Personal is available for all major operating systems. Please click on the links below for installation instructions.</p> <p>Install Globus Connect Personal</p> <p>Globus Connect Personal for Mac for Mac OS X 10.9 or higher</p> <p>Globus Connect Personal for Linux for common x64-based distributions</p> <p>Globus Connect Personal for Windows for currently supported Windows versions</p>
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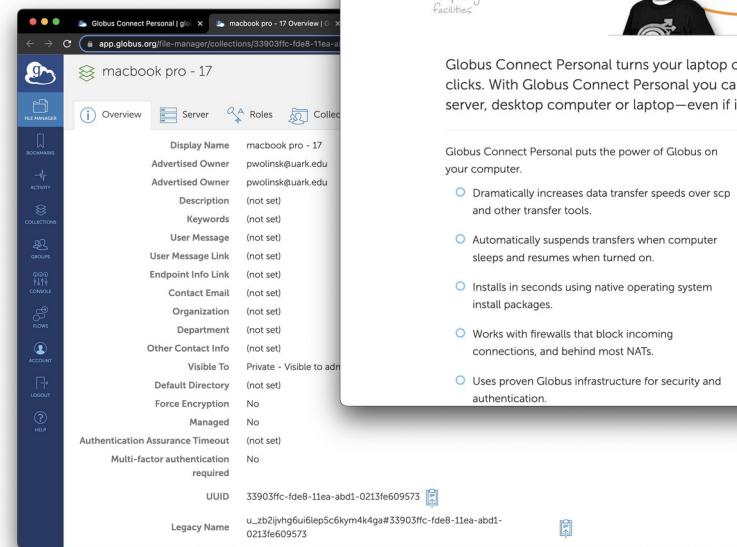
<http://globus.org/globus-connect-personal>

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Globus – Using a Personal Endpoint/Pinnacle

To move data using Globus between your personal device:

1. start your Globus Connect Personal server on your device
2. click on the Globus Connect Personal server icon and select **Web: Transfer Files**
3. A browser window with Globus File manager will appear with your local endpoint opened on one side
4. select another Globus end point and start moving data



The screenshot shows the Globus Connect Personal landing page. It features a central image of a smiling woman with the text "your computer becomes an endpoint in the Globus cloud". Above her are icons for "storage", "sync", and "globus connect". Below the image are sections for "Super computing facilities", "archives", "share", "scientific instruments", and "another personal computer". Arrows indicate data transfer between these components. A note at the bottom right states: "* File transfer and sharing between personal computers requires a paid subscription." The main text on the page reads: "Globus Connect Personal turns your laptop or other personal computer into a Globus endpoint with just a few clicks. With Globus Connect Personal you can share and transfer files to/from a local machine—campus server, desktop computer or laptop—even if it's behind a firewall and you don't have administrator privileges." To the right, there's a sidebar with links for "Install Globus Connect Personal" (Mac OS X 10.9 or higher, Linux, Windows), "Globus Connect Personal for Mac", "Globus Connect Personal for Linux", and "Globus Connect Personal for Windows".

<http://globus.org/globus-connect-personal>

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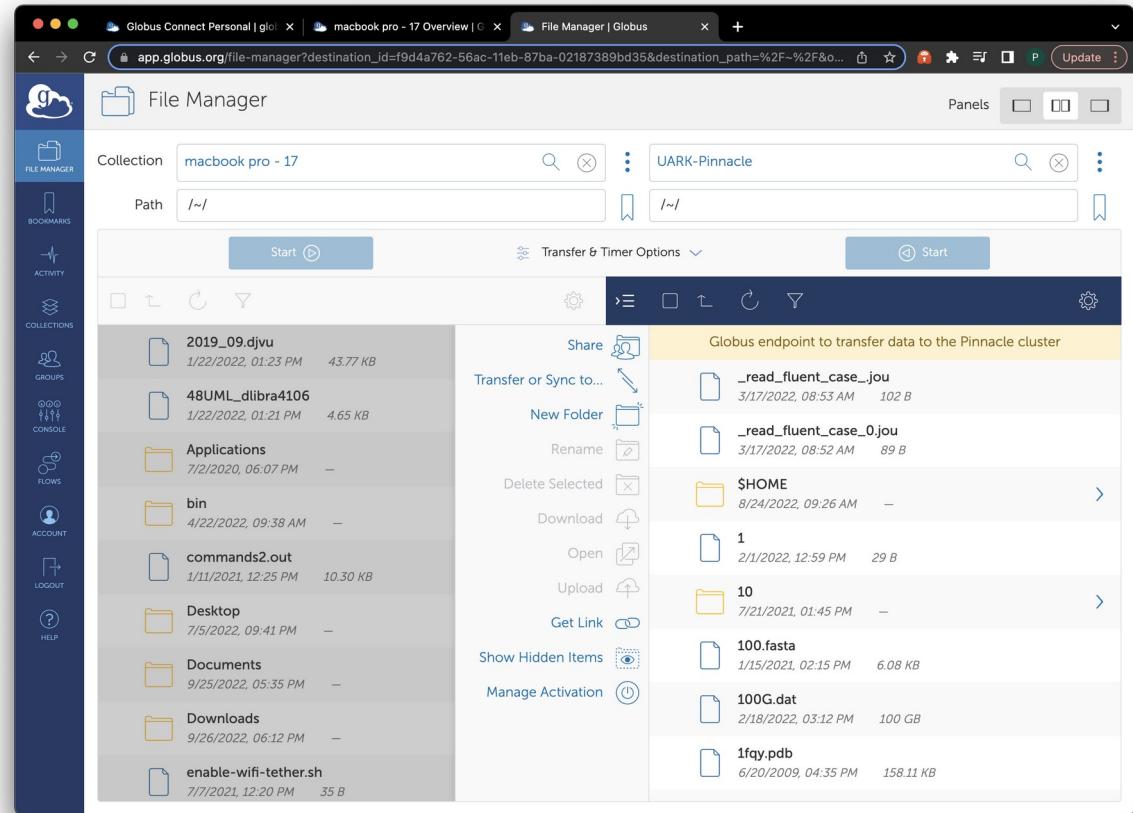
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Globus – Using a Personal Endpoint/Pinnacle

To move data using Globus between your personal device:

1. start your Globus Connect Personal server on your device
2. click on the Glonus Connect Personal server icon and select **Web: Transfer Files**
3. A browser window with Globus File manager will appear with your local endpoint opened on one side
4. select another Globus end point and start moving data



<http://globus.org/globus-connect-personal>

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Getting Started with the Arkansas Research Platform

SAU Campus
February 20

Day 2

supported by



ARKANSAS RESEARCH PLATFORM The logo for the Arkansas Research Platform, featuring the letters "ARP" in a large, blue, sans-serif font, with a stylized white swoosh graphic underneath the "P".

Day 2 Schedule

9:00 am	SSH access SSH clients (Windows/MacOS/Linux) Logging into Pinnacle and Grace Network Accessibility Pinnacle's 2-step login on the DMZ network
9:30 am	Intro to Command Line Interface (CLI) Bash shell Environment Basic linux commands File systems /home and /scratch
10:00 am	Queueing System Queues Running a sample job Interactive job Batch job Job arrays
10:30 am	Software modules Modules Conda environments Conda - GRACE deep dive
11:00 am	Moving and Storing Data, part 2 Basic object storage principles; What is ROSS? Request access to ROSS - identities and secret keys ecs-sync
12:00 pm	LUNCH
1:00 pm	Individual project help Questions, Tell me more, and Bring your own project (optional for attendees)

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Secure Shell Client

Secure Shell (SSH) is a protocol for exchanging information between two nodes on a network: an SSH server and an SSH client. SSH client requests a connection from and SSH server. Client runs only when a connection is established. Server runs continuously waiting for connection requests.

An SSH server is running on each of the login nodes on the ARP clusters (as well on each of the compute nodes).

Linux, Mac

SSH client is built in! (server too)

Windows

Windows Subsystem for Linux (WSL): Open PowerShell and run “wsl --install”

OR

1. please open this page in a browser:

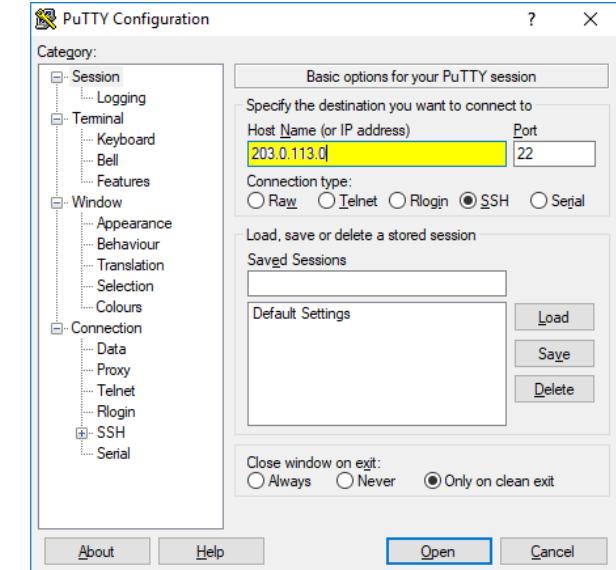
<http://www.chiark.greenend.org.uk/~sgtatham/putty/download.html>

(first link in google search with for keyword “putty”)

2. and right click and save to the desktop:

putty.exe, pscp.exe and plink.exe

<http://www.chiark.greenend.org.uk/~sgtatham/putty/download.html>



Secure Shell – log into Pinnacle and Grace

ssh user@hostname - log into a remote system

Linux

1. open a terminal and run
2. run:

*ssh <username>@hpc-portal2.hpc.uark.edu
ssh <username>@login.hpc.uams.edu*

*Pinnacle
Grace*

3. enter your supplied password when prompted.

Windows

1. double click on **putty.exe** on the desktop
2. specify

*hpc-portal2.hpc.uark.edu
login.hpc.uams.edu*

*Pinnacle
Grace*

as “Hostname” and click on “Open” button.

3. enter your username (w/o ‘@uark.edu’ or ‘@uams.edu’)
4. enter your supplied password

*ssh <username>@hpc-portal2.hpc.uark.edu
ssh <username>@login.hpc.uams.edu*

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Network Accessibility – Pinnacle & Grace

Pinnacle

Network Location	Access Type		Server DNS	Availability
UARK Campus/UARK VPN	SSH	1	pinnacle.uark.edu	Open
UARK Campus/UARK VPN	HTTPS	2	pinnacle-portal.uark.edu	Open
UARK Campus/UARK VPN	SSH/HTTPS	3	hpc-portal.uark.edu	Open
UARK Campus/UARK VPN	SSH/HTTPS	4	hpc-portal2.hpc.uark.edu	Open
Other UASYS Campus	SSH	1	pinnacle.uark.edu	Open
Other UASYS Campus	HTTPS	2	pinnacle-portal.uark.edu	Open
Other UASYS Campus	SSH/HTTPS	3	hpc-portal.uark.edu	Open
Other UASYS Campus	SSH/HTTPS	4	hpc-portal2.hpc.uark.edu	Open
Commodity Internet	SSH	1	pinnacle.uark.edu	Closed
Commodity Internet	HTTPS	2	pinnacle-portal.uark.edu	Closed
Commodity Internet	SSH/HTTPS	3	hpc-portal.uark.edu	Closed
Commodity Internet	SSH/HTTPS	4	hpc-portal2.hpc.uark.edu	Open (2-step login)

Grace

Network Location	Access Type		Server DNS	Availability
UAMS Campus/UAMS VPN	SSH	1	login.hpc.uams.edu	Open
UAMS Campus/UAMS VPN	HTTPS	2	portal.hpc.uams.edu	Open
Other UASYS Campus	SSH	1	login.hpc.uams.edu	Closed
Other UASYS Campus	HTTPS	2	portal.hpc.uams.edu	Open
Commodity Internet	SSH	1	portal.hpc.uams.edu	Closed
Commodity Internet	HTTPS	2	portal.hpc.uams.edu	Open

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Pinnacle – Demilitarized Zone (DMZ) access

hpc-portal2.hpc.uark.edu - login node for both SSH and HTTPS (OOD portal) accessible from anywhere on the internet. Access is granted using a 2-step process:

1. register your source IP and add it to the list of hosts allowed to access the Pinnacle SSH login nodes and Pinnacle OOD portal. Ssh into port 2022 of hpc-portal2.hpc.uark.edu.

```
ssh <username>@hpc-portal2.hpc.uark.edu -p 2022
```

```
[pawel@frontdesk ~]$ ssh testuser@hpc-portal2.hpc.uark.edu -p 2022
Password:
Last login: Tue Dec  7 16:58:10 2021 from 75.205.5.185
          Arkansas High Performance Computing Center
          off-campus access portal
=====
https://hpc-portal2.hpc.uark.edu      <- Open OnDemand Web Portal
ssh hpc-portal2.hpc.uark.edu         <- Standard AHPCC login

Access allowed from 75.205.5.185 until Wed Dec  8 06:02:43 CST 2021
=====
Connection to hpc-portal2.hpc.uark.edu closed.
[pawel@frontdesk ~]$
```

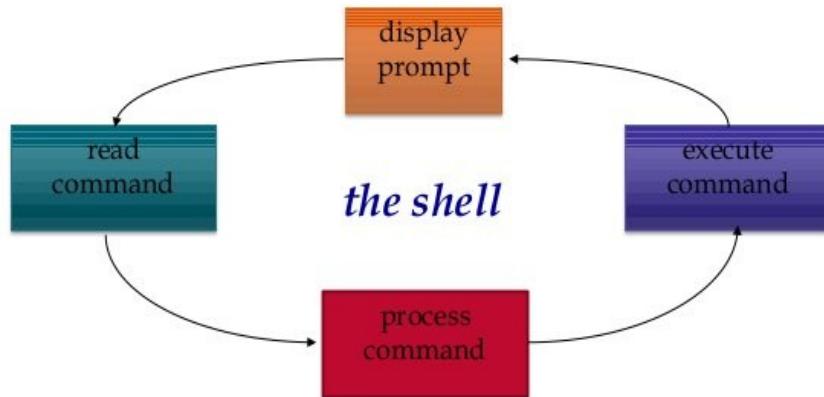
2. Log in as normal (without specifying port 2022):

```
ssh <username>@hpc-portal2.hpc.uark.edu
https://hpc-portal2.hpc.uark.edu
```

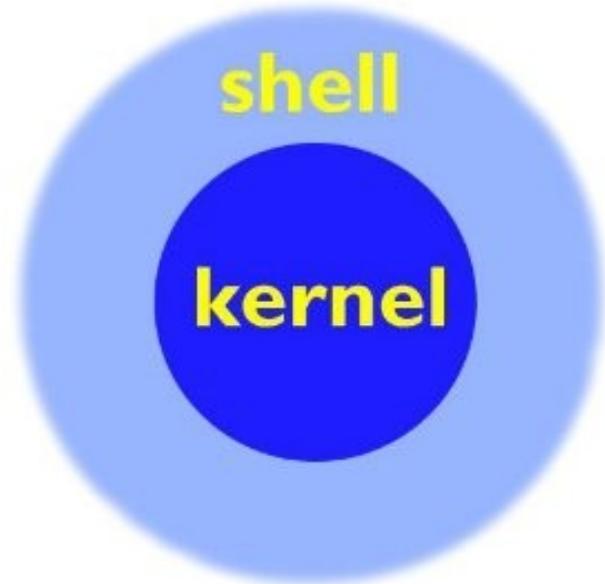
Bash Shell

Kernel – core part of the operating system. Contains all the drivers necessary to communicate with the underlying hardware. Manages the CPUs, memory, I/O operations from all peripheral devices (disk, keyboard, mouse, printers, scanners etc.)

Shell – provides the Command Line Interface (CLI) for interaction with the kernel and maintains the environment, a set of variables. Bash is the most commonly used shell on Linux.



Kernel receives commands from the shell and performs them. There may be multiple shells interacting with the kernel at one time (multiple users logged in at once, each with multiple shells).



Bash Shell - Environment

env – shows all of the environmental variables that are defined in your shell

- SHELL – the shell you are running
 - HOSTNAME - name of the host
 - PWD - present working directory
 - USER - user name
 - HOME - path of the user's home directory
 - PATH – list of directories to search when user types in a command
 - PS1 – configures prompt
- ...

[**pwolinsk@login1 ~]**

[**username@host working_directory**]

(~ is a shortcut notation for user's HOME directory)

```
[pwolinsk@login1 ~]$ echo $PS1  
[\u@\h \w]\$  
[pwolinsk@login1 ~]$ echo $SHELL  
/bin/bash  
[pwolinsk@login1 ~]$  
[pwolinsk@login1 ~]$ export PS1=" [hello \u@\h \w]\$"  
[hello pwolinsk@login1 ~]$
```

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Bash Shell - First Linux Commands

echo - display text string or contents of an environmental variable

who – list users currently logged on

whoami

hostname

date

pwd – show present working directory

man - show a manual page (help page) about any command

(in man type “/” followed by a search string, “n” to find next occurrence)

apropos - show any manual pages which reference a given string

history - show listing of your commands

nano – basic text editor

vi – powerful text editor

```
[pwolinsk@login1 ~]$ echo $PS1  
[\u@\h \w]\$  
[pwolinsk@login1 ~]$ echo $SHELL  
/bin/bash  
[pwolinsk@login1 ~]$  
[pwolinsk@login1 ~]$ export PS1=" [hello \u@\h \w]\$"  
[hello pwolinsk@login1 ~]$
```

File Systems – Pinnacle & Grace

df - disk free **mount** - show mounted file systems and mount points

```
pinnacle-11:pwolinsk:~$ df -h
Filesystem                                Size  Used Avail Use% Mounted on
devtmpfs                                     7.8G   0    7.8G  0% /dev
tmpfs                                         7.8G  24K  7.8G  1% /dev/shm
tmpfs                                         7.8G 153M  7.7G  2% /run
/dev/mapper/centos-root                      8.0G  6.7G  1.4G  84% /
/dev/vda1                                    1014M 173M  842M  18% /boot
172.17.27.1@o2ib,172.17.27.21@o2ib:172.17.27.2@o2ib,172.17.27.22@o2ib:/scrfs  2.3P  1.2P  1.2P  50% /scrfs
172.17.27.13@o2ib:/scr1                     11T   464G  9.7T  5% /scr1
172.17.27.14@o2ib:/scr2                     11T   964G  9.3T  10% /scr2
tmpfs                                         1.6G   0    1.6G  0%
/run/user/269030
pinnacle-11:pwolinsk:~$
```

```
[pwolinsk@xbt002 ~]$ df -h
Filesystem                                Size  Used Avail Use% Mounted on
/dev/mapper/sysvg-root                      88G  30G  59G  34% /
devtmpfs                                     63G   0    63G  0% /dev
tmpfs                                         63G   0    63G  0% /dev/shm
tmpfs                                         63G 490M  63G  1% /run
tmpfs                                         63G   0    63G  0% /sys/fs/cgroup
/dev/mapper/uservg-scratch                  465G  33M  465G  1% /local-scratch
/dev/sda2                                      997M 168M  829M  17% /boot
/dev/sda3                                      500M  16K  500M  1% /boot/efi
uams-gs                                       1.9P  1.4P  468T 75% /home
tmpfs                                         13G   0    13G  0% /run/user/0
tmpfs                                         13G   0    13G  0% /run/user/1692600016
[pwolinsk@xbt002 ~]$
```

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File Systems – Pinnacle & Grace

ls - list files **ls -l** - long listing **ls -al** - show all files (including hidden files, starting with ".")

```
pinnacle-11:guest01:~$ ls -l

drwxr-xr-x 2 guest01 guest01 32768 Aug 13 2009 mpiexamples
-rw-r--r-- 1 guest01 guest01 0 Mar 17 13:17 newfile
drwxrwxr-x 2 guest01 guest01 32768 Mar 17 14:55 tt
-----
|       |           |       |           |       |           |
| user/group/others user group   size modification   filename
| permissions             in      date
|                           bytes
file/directory/link
 r read    w write   x execute
```

```
pinnacle-11:guest01:~$ ls -al
drwx----- 6 guest01 guest01 32768 Mar 17 14:55 .
drwxr-xr-x 24 root     root      0 Mar 17 15:59 ..
-rw------- 1 guest01 guest01 4604 Mar 17 13:28 .bash_history
-rw-r--r-- 1 guest01 guest01 24 Aug 5 2009 .bash_logout
-rwxr-xr-x 1 guest01 guest01 132 Mar 17 13:20 .bash_profile
1rwxrwxrwx 1 guest01 guest01 13 Jan 27 2011 .bashrc -> .bash_profile
...
drwxr-xr-x 2 guest01 guest01 32768 Aug 13 2009 mpiexamples
-rwx----- 1 guest01 guest01 0 Mar 17 13:17 newfile
drwxrwxr-x 2 guest01 guest01 32768 Mar 17 14:55 tt
...
```

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File Systems – /home and /scratch

/home/\$USER – file storage for each user

/scratch/<jobid> - temporary directory for executing jobs

At the beginning of a job copy your input files to /scratch/<jobid> and at the end of job copy output files to back to /home/\$USER

```
c1601:pwolinsk:~$ df
Filesystem           1K-blocks      Used   Available  Use% Mounted on
devtmpfs
tmpfs
tmpfs
tmpfs
tmpfs
/sys/fs/cgroup
/dev/mapper/xcatvg-root          229595616  16333344  213262272  8% /
/dev/sda1                  517868    226032   291836  44% /boot
172.17.27.13@o2ib:/scr1          11468687248  486872436  10403223304  5% /scr1
172.17.27.14@o2ib:/scr2          11471001300  1011719572  9880567544  10% /scr2
172.17.27.10@o2ib,172.17.27.21@o2ib:172.17.27.20@o2ib,172.17.27.22@o2ib:/scrfs  2455426280448 1214971101936 1215600540940  50% /scrfs
tmpfs
/run/user/1006
tmpfs
/run/user/269030
c1601:pwolinsk:~$ pwd
/home/pwolinsk
c1601:pwolinsk:~$ ls -l /scratch |head
total 0
lrwxrwxrwx 1 root root 13 Feb 16 2022 1003741 -> /scr1/1003741
lrwxrwxrwx 1 root root 13 Feb 16 2022 1010024 -> /scr2/1010024
lrwxrwxrwx 1 root root 13 Feb 16 2022 1010044 -> /scr2/1010044
lrwxrwxrwx 1 root root 13 Feb 17 2022 1010047 -> /scr1/1010047
lrwxrwxrwx 1 root root 13 Feb 17 2022 1012608 -> /scr2/1012608
lrwxrwxrwx 1 root root 13 Feb 17 2022 1013345 -> /scr1/1013345
lrwxrwxrwx 1 root root 13 Feb 18 2022 1020146 -> /scr2/1020146
lrwxrwxrwx 1 root root 13 Feb 21 2022 1025910 -> /scr2/1025910
lrwxrwxrwx 1 root root 13 Feb 21 2022 1025911 -> /scr1/1025911
c1601:pwolinsk:~$
```

supported by

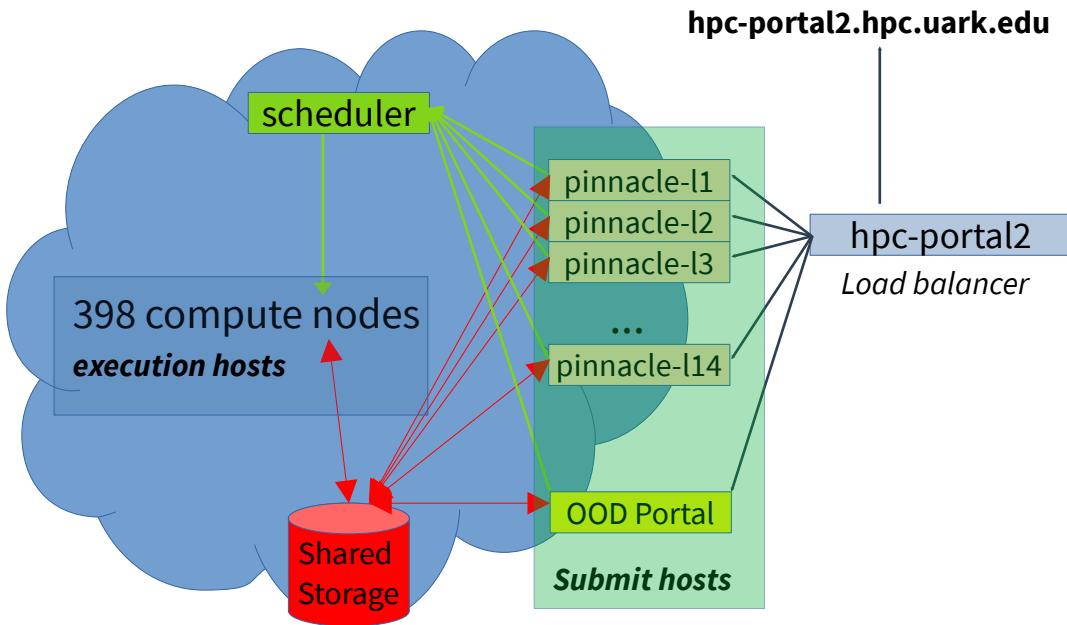


ARKANSAS RESEARCH PLATFORM

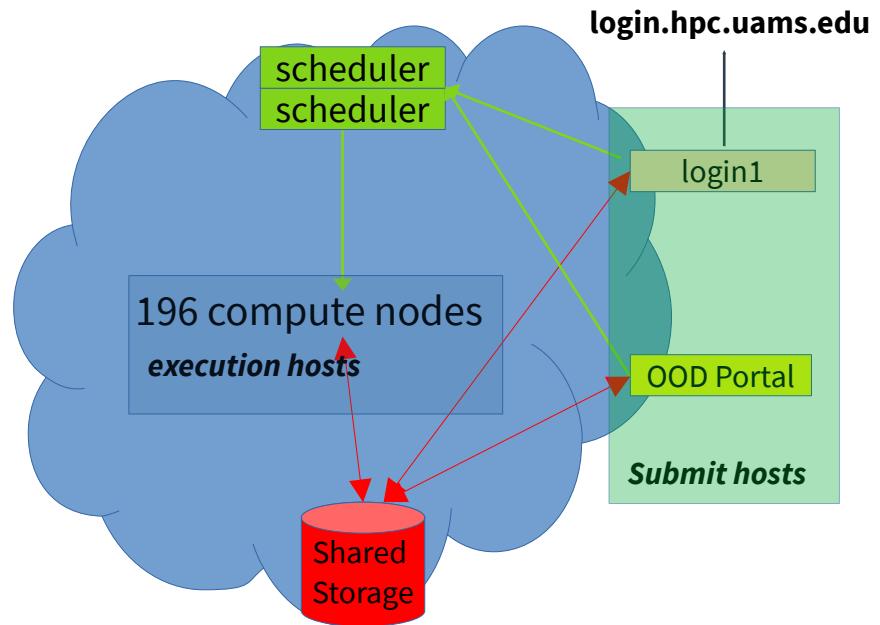


ARP Clusters – Queueing System

Pinnacle



Grace



scheduler – coordinates all of the nodes in the queue (SLURM queueing system)
submit hosts – nodes from which users submit jobs to the queues
login hosts – nodes accessible from the external network
compute nodes – nodes assigned to queues and run jobs

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High Performance Computing

AHPCC
Arkansas High Performance Computing Center

ARKANSAS RESEARCH PLATFORM **ARP**

Queueing System

sinfo - show node/queue assignment

squeue - jobs in the queues

Compute nodes are divided into queues (or partitions).

```
pinnacle-11:pwolinsk:~$ sinfo
PARTITION    AVAIL   TIMELIMIT  NODES  STATE NODELIST
comp01        up      1:00:00     3  drain c[1402-1404]
comp01        up      1:00:00    42  alloc c[1407-1420,1501-1520,1603-1610]
comp01        up      1:00:00     2  idle c[1405-1406]
comp01        up      1:00:00     1  down c1401
comp06        up      6:00:00     1  drain c1404
comp06        up      6:00:00    42  alloc c[1407-1420,1501-1520,1603-1610]
comp06        up      6:00:00     2  idle c[1405-1406]
comp72        up  3-00:00:00    42  alloc c[1407-1420,1501-1520,1603-1610]
gpu06         up      6:00:00     2  alloc c[1713,1715]
gpu06         up      6:00:00    17  idle c[1612-1615,1701-1712,1714]
gpu72         up  3-00:00:00     2  alloc c[1713,1715]
gpu72         up  3-00:00:00    17  idle c[1612-1615,1701-1712,1714]
condo          up  infinite     4  alloc c[1307,1310-1312]
condo          up  infinite    21  idle c[1308-1309,1313-1328,1611,1719-1720]
pubcondo06    up      6:00:00     4  alloc c[1307,1310-1312]
pubcondo06    up      6:00:00    21  idle c[1308-1309,1313-1328,1611,1719-1720]
pcon06*       up      6:00:00     4  alloc c[1307,1310-1312]
pcon06*       up      6:00:00    21  idle c[1308-1309,1313-1328,1611,1719-1720]
cloud72       up  3-00:00:00     2  idle c[1601-1602]
cloud72       up  3-00:00:00     1  down c1329
himem06       up      6:00:00     6  idle c[1301-1306]
himem72       up  3-00:00:00     6  idle c[1301-1306]
pinnacle-11:pwolinsk:~$
```

supported by



ARKANSAS RESEARCH PLATFORM The logo for the Arkansas Research Platform, featuring the letters "ARP" in a large, blue, sans-serif font, with three blue wavy lines underneath.

Queueing System

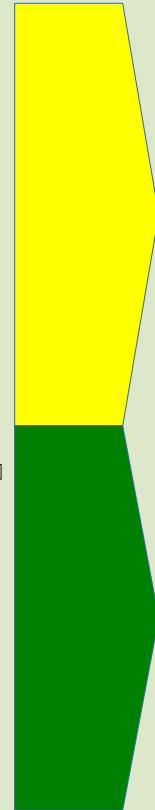
sinfo - show node/queue assignment

squeue - jobs in the queue

```
pinnacle-11:pwolinsk:~$ squeue
```

JOBID	PARTITION	NAME	USER	ST	TIME	NODES	NODELIST (REASON)
3633	condo	OG000050	aja	PD	0:00	1	(Resources)
3634	condo	OG000050	aja	PD	0:00	1	(Priority)
3635	condo	OG000050	aja	PD	0:00	1	(Priority)
3636	condo	OG000050	aja	PD	0:00	1	(Priority)
3637	condo	OG000049	aja	PD	0:00	1	(Priority)
3638	condo	OG000050	aja	PD	0:00	1	(Priority)
3639	condo	OG000050	aja	PD	0:00	1	(Priority)
3640	condo	OG000050	aja	PD	0:00	1	(Priority)
3641	condo	OG000049	aja	PD	0:00	1	(Priority)
3642	condo	OG000049	aja	PD	0:00	1	(Priority)
3643	condo	OG000049	aja	PD	0:00	1	(Priority)
3644	condo	OG000049	aja	PD	0:00	1	(Priority)
3645	condo	OG000048	aja	PD	0:00	1	(Priority)
3646	condo	OG000049	aja	PD	0:00	1	(Priority)
3650	condo	OG000048	aja	PD	0:00	1	(Priority)
3651	condo	OG000048	aja	PD	0:00	1	(Priority)
...							
6892	comp72	coalHMM_	tkchafin	PD	0:00	1	(Priority)
3632	condo	OG000051	aja	R	8:19:34	1	c1307
6551	comp72	Pmc21-MD	kypatel	R	6:09:20	8	c[1407,1409,1504,1506,1513,1605-1606,1608]
6749	comp72	coalHMM_	tkchafin	R	2-10:42:46	1	c1411
6761	comp72	coalHMM_	tkchafin	R	2-09:00:12	1	c1417
6775	comp72	coalHMM_	tkchafin	R	2-01:20:12	1	c1415
6776	comp72	coalHMM_	tkchafin	R	2-00:41:24	1	c1503
6777	comp72	coalHMM_	tkchafin	R	1-19:01:08	1	c1510
6779	comp72	coalHMM_	tkchafin	R	1-14:22:11	1	c1603
6780	comp72	coalHMM_	tkchafin	R	1-13:25:36	1	c1416
7147	gpu72	BO.bi_60	rpsander	R	1-21:56:14	1	c1715
7299	gpu72	BO.mo_80	rpsander	R	16:11:37	1	c1713
...							
7311	condo	fit_inte	yinyuan	R	3:16:02	1	c1310
7313	condo	fit_inte	yinyuan	R	3:08:58	1	c1311
7314	condo	afm	yinyuan	R	2:03:28	1	c1312

```
pinnacle-11:pwolinsk:~$
```



Pending jobs



Running jobs

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Queueing System – First job - command line

sbatch – submit a job to the queue

squeue - jobs in the queue

Copy a pre-made script into your home directory, examine it and run it on the login node.

```
pinnacle-16:pwolinsk:~$ mkdir EXAMPLES
pinnacle-16:pwolinsk:~$ cd EXAMPLES/
pinnacle-16:pwolinsk:~/EXAMPLES$ cp /scrfs/ahpcc/ARP-workshop/first.sh .
pinnacle-16:pwolinsk:~/EXAMPLES$ cat first.sh
#!/bin/bash

echo "Script is executing on: "
hostname
echo "My present working directory is $PWD"
echo "Time and date"
date
pinnacle-16:pwolinsk:~/EXAMPLES$
pinnacle-16:pwolinsk:~/EXAMPLES$ ./first.sh
Script is executing on:
pinnacle-16
My present working directory is /home/pwolinsk/EXAMPLES
Time and date
Tue Sep 27 19:04:21 CDT 2022
pinnacle-16:pwolinsk:~/EXAMPLES$
```

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ARKANSAS RESEARCH PLATFORM **ARP**

Queueing System – First interactive job - command line

sbatch – submit a batch job to the queue **srun** – submit an interactive job to the queue **squeue** - jobs in the queue

Submit the same script to the queueing system interactively – with **srun**.

```
pinnacle-16:pwolinsk:~/EXAMPLES$ srun -p comp01 --pty /bin/bash
c1405:pwolinsk:~/EXAMPLES$ ./first.sh
Script is executing on:
c1405
My present working directory is /home/pwolinsk/EXAMPLES
Time and date
Tue Sep 27 19:09:15 CDT 2022
c1405:pwolinsk:~/EXAMPLES$ squeue -u pwolinsk
      JOBID PARTITION      NAME      USER ST          TIME  NODES NODELIST(REASON)
      1315893    comp01      bash  pwolinsk  R      0:17      1  c1405
c1405:pwolinsk:~/EXAMPLES$ exit
exit
pinnacle-16:pwolinsk:~/EXAMPLES$
```

Queueing System – First batch job - command line

sbatch – submit a batch job to the queue **srun** – submit an interactive job to the queue **squeue** - jobs in the queue

Submit the same script to the queueing system as a batch job – with **sbatch**.

```
pinnacle-16:pwolinsk:~/EXAMPLES$ sbatch -p comp01 first.sh
Submitted batch job 1315894
pinnacle-16:pwolinsk:~/EXAMPLES$ squeue -u pwolinsk
      JOBID PARTITION      NAME      USER ST          TIME   NODES NODELIST(REASON)
pinnacle-16:pwolinsk:~/EXAMPLES$ ls -ltr
total 8
-rwxr-xr-x 1 pwolinsk pwolinsk 124 Sep 27 19:03 first.sh
-rw-rw-r-- 1 pwolinsk pwolinsk 130 Sep 27 19:13 slurm-1315894.out
pinnacle-16:pwolinsk:~/EXAMPLES$ cat slurm-1315894.out
Script is executing on:
c1405
My present working directory is /home/pwolinsk/EXAMPLES
Time and date
Tue Sep 27 19:13:02 CDT 2022
pinnacle-16:pwolinsk:~/EXAMPLES$
```

Queueing System – Second batch job – job array

sbatch – submit a batch job to the queue **srun** – submit an interactive job to the queue **squeue** - jobs in the queue

sbatch command with the **--array** parameter will run the same job script a specified amount of times with **\$SLURM_JOB_ARRAY_ID** variable storing a unique array element number.

```
pinnacle-16:pwolinsk:~/EXAMPLES$ cp /scrfs/ahpcc/ARP-workshop/second.sh .
pinnacle-16:pwolinsk:~/EXAMPLES$ cat second.sh
#!/bin/bash

echo "My array task id is: $SLURM_ARRAY_TASK_ID"
pinnacle-16:pwolinsk:~/EXAMPLES$ sbatch -p comp01 --array [1-5] second.sh
Submitted batch job 1315912
pinnacle-16:pwolinsk:~/EXAMPLES$ ls -ltr
total 188
-rwxr-xr-x 1 pwolinsk pwolinsk 124 Sep 27 19:03 first.sh
-rwxrwxr-x 1 pwolinsk pwolinsk 62 Sep 27 19:32 second.sh
-rw-rw-r-- 1 pwolinsk pwolinsk 23 Sep 27 19:33 slurm-1315912_5.out
-rw-rw-r-- 1 pwolinsk pwolinsk 23 Sep 27 19:33 slurm-1315912_4.out
-rw-rw-r-- 1 pwolinsk pwolinsk 23 Sep 27 19:33 slurm-1315912_3.out
-rw-rw-r-- 1 pwolinsk pwolinsk 23 Sep 27 19:33 slurm-1315912_2.out
-rw-rw-r-- 1 pwolinsk pwolinsk 23 Sep 27 19:33 slurm-1315912_1.out
pinnacle-16:pwolinsk:~/EXAMPLES$ cat slurm-1315912_5.out
My array task id is: 5
pinnacle-16:pwolinsk:~/EXAMPLES$ cat slurm-1315912_4.out
My array task id is: 4
pinnacle-16:pwolinsk:~/EXAMPLES$
```

Queueing System – Third batch job – job array

sbatch – submit a batch job to the queue **srun** – submit an interactive job to the queue **squeue** - jobs in the queue

sbatch command with the **--array** parameter will run the same job script a specified amount of times with **\$SLURM_JOB_ARRAY_ID** variable storing a unique array element number.

```
pinnacle-16:pwolinsk:~/EXAMPLES$ cp /scrfs/ahpcc/ARP-workshop/1fqy.pdb .
pinnacle-16:pwolinsk:~/EXAMPLES$ head 1fqy.pdb
HEADER      MEMBRANE PROTEIN          07-SEP-00    1FQY
TITLE       STRUCTURE OF AQUAPORIN-1 AT 3.8 A RESOLUTION BY ELECTRON
TITLE       2 CRYSTALLOGRAPHY
COMPND     MOL_ID: 1;
COMPND     2 MOLECULE: AQUAPORIN-1;
COMPND     3 CHAIN: A;
COMPND     4 SYNONYM: AQP1
SOURCE     MOL_ID: 1;
SOURCE     2 ORGANISM_SCIENTIFIC: HOMO SAPIENS;
SOURCE     3 ORGANISM_COMMON: HUMAN;
pinnacle-16:pwolinsk:~/EXAMPLES$ cp /scrfs/ahpcc/ARP-workshop/third.sh .
pinnacle-16:pwolinsk:~/EXAMPLES$ cat third.sh
#!/bin/bash

head -$SLURM_ARRAY_TASK_ID 1fqy.pdb |tail -1
pinnacle-16:pwolinsk:~/EXAMPLES$ sbatch -p comp01 --array [100,102] third.sh
Submitted batch job 1315918
pinnacle-16:pwolinsk:~/EXAMPLES$ ls -ltr
total 176
-rwxr-xr-x 1 pwolinsk pwolinsk   124 Sep 27 19:03 first.sh
-rwxrwxr-x 1 pwolinsk pwolinsk    62 Sep 27 19:32 second.sh
-rw-r--r-- 1 pwolinsk pwolinsk 158112 Sep 27 19:38 1fqy.pdb
-rwxrwxr-x 1 pwolinsk pwolinsk    58 Sep 27 19:38 third.sh
-rw-rw-r-- 1 pwolinsk pwolinsk    81 Sep 27 19:39 slurm-1315918_102.out
-rw-rw-r-- 1 pwolinsk pwolinsk    81 Sep 27 19:39 slurm-1315918_100.out
pinnacle-16:pwolinsk:~/EXAMPLES$ cat slurm-1315918_102.out
REMARK 3
pinnacle-16:pwolinsk:~/EXAMPLES$ cat slurm-1315918_100.out
REMARK 3 SIDE-CHAIN BOND          (A**2) : NULL ; NULL
pinnacle-16:pwolinsk:~/EXAMPLES$
```

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ARKANSAS RESEARCH PLATFORM



Software Modules

Hunders of applications, compilers and scripting languages, some with multiple versions, are installed on Grace and Pinnacle. They cannot all be available to the users at once because they interfere with each other.

The Environment Modules system is a tool to help users manage the shell environment and provide access to specific applications and versions on demand. It does this by manipulating the values of shell environmental variables. The most important of those are **\$PATH** and **\$LD_LIBRARY_PATH**.

```
pinnacle-16:pwolinsk:~$ ls /share/apps/
abinit      cplex          ganglia    idl           mathematics   OpenFOAM        qt           synopsys
accumulo    CPMD          gappa     iftop         Matjes       openmpi        Qt            szip
acml         crabz         Gaussian  iftop-0.17   matlab       opt          QTNetwork    TauDEM
alamode     cuda          gcc       igb-5.3.5.4  maui        oracle_developer_studio R  tcl
amber       curl           gd        imp          mauiversions orca        rclone      texinfo
amber16tools17 dar           gdal     intel       maven       orca_3_0_3_linux_x86-64 relion
amdappsdk   ddscat        gdrcopy   intel18     meeple     osu-micro-bench rh6
ansys       dealii        geomview intel2020-2 MKDAT      ovito        RITHM
...
pinnacle-16:pwolinsk:~$ ls /share/apps/bioinformatics/
abyss       bismark        DataAnalysis GCTA      last        MUMmer      picard      sate-tools-linux swarm
admixture   blas           datamash    Gctf      lib         muscle      pilon      sepp
aegean      blasr          dDocent    gemma    libBigWig  mutect      pindel      sepp-tools
agalma      blast          deepTools GeneMark lordec     MutMap      pitchfork seqtk
allpaths-lg blat           diamond   genemark_es lorma     ncbi        plink      shapeit
angsd       BLINK          dock      genometools lsc       necat      PopLDdecay sickle
annovar     blobtools     edirect   gtdbtk    MACSE      newbler    popoolation2 simple
astalavista bonito        eigen     GTDBTK   mafft      newick-utils pplacer    smrtlink
...

```

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Software Modules

module avail - list all available modules

module list – list modules currently loaded

module load – load a module

module remove – remove a loaded module

```
pinnacle-16:pwolinsk:~$ module avail
```

/share/apps/modulefiles				
BayesAss3-SNPs/1.0		edirect/20200815	(D)	java_ibmjdk_1.8.0
BerkeleyGW/2.1		elmer/1.0		java_openjdk_1.7.0
BerkeleyGW/3.0.1	(D)	elpa/2021.11.002		java_openjdk_1.8.0_20
CAFE/4.2.1		emboss/6.6.0		java_openjdk_1.8.0
CERN/6.24		epa-ng/0.3.6		java_openjdk_11.0.10
EnTAP/0.8.1		exabayes/1.5.1		java_openjdk_14.0.1
LIGGGHTS/1		exonerate/2.4.0		java_openjdk_16.0.0
METABOLIC/4.0		express/1.5.1		java_sunjdk_1.7.0_80
MUMmer/3.23		fastANI/1.32		java_sunjdk_1.7.0
MaSuRCA/3.4.2		fastepistasis/2.07		java_sunjdk_1.8.0_72
MetaGeneMark/3.38		fastqc/0.11.5		java_sunjdk_1.8.0_92
PAML/4.6		fasttree/2.1.10		java_sunjdk_1.8.0
PAML/4.8		fastx/0.0.14		java_sunjdk_11.0.10
PAML/4.9e	(D)	fftw/2.1.5	(D)	jellyfish/2.2.6
PASTA/1.8.3		fftw/3.2.2		jellyfish/2.3.0
PASTA/1.8.6	(D)	fftw/3.3.4-double-openmp-old		julia/1.7.0-rc1
PBSuite/15.8.24		fftw/3.3.4-double-openmp		kallisto/0.43.1
PGI/18.4		fftw/3.3.4-double		ktrim/1.1.0
PGI/18.10		fftw/3.3.4-openmpi		last/719
PGI/19.4		fftw/3.3.4		libxc/5.2.3
PGI/2016		fftw/3.3.6		libxsmm/1.17.0
PGI/2016.5		fftw/3.3.8		mafft/7.304b
PGI/2017		flash/1.2.11		mafft/7.490
PGI/2019	(D)	fluent/2020R2		maker/2.31.8
PopLDdecay/3.4.1		fluent/2022R1	(D)	maker/2.31.9
R/MR3.3.1		flye/2.3.3		mathematica/12.3
R/MR3.4.2		flye/2.6		mathematica/13.0
R/MR4.0.2		flye/2.8.3	(D)	mathematica/13.1
R/3.3.2		gadget/2.0.7-SF		matlab/r2017a
R/3.4.3-deprecated		gadget/2.03-SF	(D)	matlab/r2019a
...				

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Software Modules

module avail - list all available modules

module list – list modules currently loaded

module load – load a module

module remove – remove a loaded module

```
pinnacle-16:pwolinsk:~$ which python
/usr/bin/python
pinnacle-16:pwolinsk:~$ python
Python 2.7.5 (default, Nov 16 2020, 22:23:17)
[GCC 4.8.5 20150623 (Red Hat 4.8.5-44)] on linux2
Type "help", "copyright", "credits" or "license" for more information.
>>> quit()
pinnacle-16:pwolinsk:~$ module avail python
----- /share/apps/modulefiles -----
python/anaconda-2.7.13    python/anaconda-3.10    python/2.7.5          python/2.7.16-anaconda   python/3.7.3-anaconda
python/anaconda-2.7.15    python/intel-2.7.16   python/2.7.9          python/2.7.16-intel     python/3.7.7-intel
python/anaconda-2.7.16    python/intel-3.6.8   python/2.7.11         python/2.7.18-anaconda   python/3.8-anaconda
python/anaconda-3.6.0      python/intel-3.7.7   python/2.7.13-anaconda python/3.5.1           python/3.9-anaconda
python/anaconda-3.7.3      python/rh27        python/2.7.15-anaconda python/3.6.0-anaconda   python/3.10-anaconda (D)
python/anaconda-3.8        python/2.7-rh       python/2.7.15b        python/3.6.0           python/3.6.0
python/anaconda-3.9        python/2.7.3        python/2.7.15          python/3.6.8-intel     python/3.6.8-intel

Where:
D: Default Module
pinnacle-16:pwolinsk:~$ module load python/3.10-anaconda
pinnacle-16:pwolinsk:~$ module list
Currently Loaded Modules:
 1) os/el7  2) python/3.10-anaconda
pinnacle-16:pwolinsk:~$ which python
/share/apps/python/anaconda-3.10/bin/python
pinnacle-16:pwolinsk:~$ python
Python 3.10.4 | packaged by conda-forge | (main, Mar 24 2022, 17:38:57) [GCC 10.3.0] on linux
Type "help", "copyright", "credits" or "license" for more information.
>>> quit()
pinnacle-16:pwolinsk:~$ module remove python
pinnacle-16:pwolinsk:~$ which python
/usr/bin/python
pinnacle-16:pwolinsk:~$
```

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ARKANSAS RESEARCH PLATFORM



Conda (Python Modules)

Package, dependency and environment management for Python (and other languages)

On Pinnacle, load the Python 3.10 module and activate the base environment.

```
· pinnacle-15:pwolinsk:~$ module load python/3.10-anaconda
pinnacle-15:pwolinsk:~$ module list

Currently Loaded Modules:
 1) os/el7   2) python/3.10-anaconda

pinnacle-15:pwolinsk:~$ which python
/share/apps/python/anaconda-3.10/bin/python
pinnacle-15:pwolinsk:~$ source /share/apps/bin/conda-3.10.sh
(base) pinnacle-15:pwolinsk:~$ conda env list
# conda environments:
#
BoltzTraP-env          /home/pwolinsk/.conda/envs/BoltzTraP-env
bwa-env                 /home/pwolinsk/.conda/envs/bwa-env
circlator               /home/pwolinsk/.conda/envs/circlator
delly-env                /home/pwolinsk/.conda/envs/delly-env
...
htseq-3.10              /share/apps/python/anaconda-3.10/envs/htseq-3.10
mpi4py-mvapich-3.10    /share/apps/python/anaconda-3.10/envs/mpi4py-mvapich-3.10
mpi4py-openmpi-3.10     /share/apps/python/anaconda-3.10/envs/mpi4py-openmpi-3.10
phonopy-3.10             /share/apps/python/anaconda-3.10/envs/phonopy-3.10
pymor-3.10               /share/apps/python/anaconda-3.10/envs/pymor-3.10
qiime2-2022.2-3.10      /share/apps/python/anaconda-3.10/envs/qiime2-2022.2-3.10

(base) pinnacle-15:pwolinsk:~$
```

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ARKANSAS RESEARCH PLATFORM



Conda (Python Modules) – existing environments

Package, dependency and environment management for Python (and other languages)

conda env list – list installed environments **conda activate <env_name>** - activate env **conda deactivate** - deactivate

```
· (base) pinnacle-15:pwolinsk:~$ conda env list
bwa-env                               /home/pwolinsk/.conda/envs/bwa-env
circlator                             /home/pwolinsk/.conda/envs/circlator
delly-env                             /home/pwolinsk/.conda/envs/delly-env
...
htseq-3.10                            /share/apps/python/anaconda-3.10/envs/htseq-3.10
mpi4py-mvapich-3.10                  /share/apps/python/anaconda-3.10/envs/mpi4py-mvapich-3.10
mpi4py-openmpi-3.10                  /share/apps/python/anaconda-3.10/envs/mpi4py-openmpi-3.10
...
· (base) pinnacle-15:pwolinsk:~$ which htseq-qa
/usr/bin/which: no htseq-qa in (/share/apps/python/anaconda-3.10/bin:/share/apps/python/anaconda-3.10/condabin:/share/
apps/python/anaconda-3.10/condabin:/share/apps/python/anaconda-3.10/bin:/usr/lib64/qt-3.3/bin:/usr/local/bin:/usr/bin:/
usr/local/sbin:/usr/sbin:/opt/ibutils/bin:/scrfs/ahpcc/libvirt/bin:/share/apps/bin:/home/pwolinsk/bin:/home/pwolinsk/
Leyh/t)
(base) pinnacle-15:pwolinsk:~$ conda activate htseq-3.10
(htseq-3.10) pinnacle-15:pwolinsk:~$ which htseq-qa
/share/apps/python/anaconda-3.10/envs/htseq-3.10/bin/htseq-qa
(htseq-3.10) pinnacle-15:pwolinsk:~$ ls /share/apps/python/anaconda-3.10/envs/htseq-3.10
bin  compiler_compat  conda-meta  etc  include  lib  man  sbin  share  ssl  x86_64-conda_cos7-linux-gnu  x86_64-conda-
linux-gnu
(htseq-3.10) pinnacle-15:pwolinsk:~$ conda deactivate
(base) pinnacle-15:pwolinsk:~$ which htseq-qa
/usr/bin/which: no htseq-qa in (/share/apps/python/anaconda-3.10/bin:/share/apps/python/anaconda-3.10/condabin:/share/
apps/python/anaconda-3.10/condabin:/share/apps/python/anaconda-3.10/bin:/usr/lib64/qt-3.3/bin:/usr/local/bin:/usr/bin:/
usr/local/sbin:/usr/sbin:/opt/ibutils/bin:/scrfs/ahpcc/libvirt/bin:/share/apps/bin:/home/pwolinsk/bin:/home/pwolinsk/
Leyh/t)
(base) pinnacle-15:pwolinsk:~$
```

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ARKANSAS RESEARCH PLATFORM



Conda (Python Modules) – install your own package

Package, dependency and environment management for Python (and other languages)

conda config -add channels <channel> – add channel **conda search <package>**

```
(base) pinnacle-15:pwolinsk:~$ conda config --add channels defaults
(base) pinnacle-15:pwolinsk:~$ conda config --add channels bioconda
(base) pinnacle-15:pwolinsk:~$ conda config --add channels conda-forge
(base) pinnacle-15:pwolinsk:~$ cat .condarc
channels:
  - conda-forge
  - bioconda
  - defaults
(base) pinnacle-15:pwolinsk:~$ conda search tensorflow
Loading channels: done
# Name          Version      Build Channel
tensorflow      0.7.1        py27_0  conda-forge
tensorflow      0.7.1        py34_0  conda-forge
...
tensorflow      2.9.1        mkl_py37h58a621a_0  pkgs/main
tensorflow      2.9.1        mkl_py38h96f9fba_0  pkgs/main
...
tensorflow      2.10.0       cpu_py39h4655687_0  conda-forge
tensorflow      2.10.0       cuda112py310he87a039_0  conda-forge
tensorflow      2.10.0       cuda112py37h01c6645_0  conda-forge
tensorflow      2.10.0       cuda112py38hded6998_0  conda-forge
tensorflow      2.10.0       cuda112py39h01bd6f0_0  conda-forge
...
```

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ARKANSAS RESEARCH PLATFORM



Conda (Python Modules) – install your own packages

Package, dependency and environment management for Python (and other languages)

conda install -c <channel> package=<version>=<build>

```
. (base) pinnacle-15:pwolinsk:~$ conda install -c conda-forge tensorflow=2.10.0=cpu_py310hd1aba9c_0
Collecting package metadata (current_repodata.json): done
Solving environment: -
The environment is inconsistent, please check the package plan carefully
The following packages are causing the inconsistency:

- conda-forge/noarch::ipywidgets==7.7.1=pyhd8ed1ab_0
- conda-forge/noarch::jupyterlab==3.4.4=pyhd8ed1ab_0
- conda-forge/noarch::notebook-shim==0.1.0=pyhd8ed1ab_0
- conda-forge/noarch::jupyterlab_server==2.15.0=pyhd8ed1ab_0
- conda-forge/noarch::nbclassic==0.4.3=pyhd8ed1ab_0
- conda-forge/noarch::nbconvert-core==6.5.0=pyhd8ed1ab_0
- conda-forge/noarch::notebook==6.4.12=pyha770c72_0
- conda-forge/noarch::nbconvert==6.5.0=pyhd8ed1ab_0
- conda-forge/noarch::nbconvert-pandoc==6.5.0=pyhd8ed1ab_0
- conda-forge/noarch::jupyter_server==1.18.1=pyhd8ed1ab_0
- conda-forge/noarch::widgetsnbextension==3.6.1=pyha770c72_0
done
```

```
## Package Plan ##
```

```
environment location: /share/apps/python/anaconda-3.10
```

```
added / updated specs:
- tensorflow==2.10.0=cpu_py310hd1aba9c_0
...
```

Users can't install into base environment

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Conda (Python Modules) – create a private environment

Package, dependency and environment management for Python (and other languages)

conda create -n <environment>

```
. (base) pinnacle-15:pwolinsk:~$ conda create -n tensorflow-pawel
Collecting package metadata (current_repodata.json): done
Solving environment: done

## Package Plan ##

environment location: /home/pwolinsk/.conda/envs/tensorflow-pawel

Proceed ([y]/n)? y

Preparing transaction: done
Verifying transaction: done
Executing transaction: done
#
# To activate this environment, use
#
#     $ conda activate tensorflow-pawel
#
# To deactivate an active environment, use
#
#     $ conda deactivate

Retrieving notices: ...working... done
(base) pinnacle-15:pwolinsk:~$ conda activate tensorflow-pawel
(tensorflow-pawel) pinnacle-15:pwolinsk:~$
```

Users can install into private environments

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ARKANSAS RESEARCH PLATFORM **ARP**

Conda (Python Modules) – install into a private environment

Package, dependency and environment management for Python (and other languages)

conda install -c <channel> package=<version>=<build>

```
(tensorflow-pawel) pinnacle-15:pwolinsk:~$ conda install -c conda-forge tensorflow=2.10.0=cpu_py310hd1aba9c_0
Collecting package metadata (current_repodata.json): done
Solving environment: done

## Package Plan ##

environment location: /home/pwolinsk/.conda/envs/tensorflow-pawel
added / updated specs:
- tensorflow==2.10.0=cpu_py310hd1aba9c_0

The following packages will be downloaded:
package          | build
-----|-----
_openmp_mutex-4.5 | 2_gnu      | 23 KB  conda-forge
absl-py-1.2.0    | pyhd8ed1ab_0 | 94 KB  conda-forge
aiohttp-3.8.3    | py310h5764c6d_0 | 449 KB  conda-forge
aiosignal-1.2.0  | pyhd8ed1ab_0 | 12 KB   conda-forge
...
wheel            | conda-forge/noarch::wheel-0.37.1-pyhd8ed1ab_0
wrapt             | conda-forge/linux-64::wrapt-1.14.1-py310h5764c6d_0
xz                | conda-forge/linux-64::xz-5.2.6-h166bdaf_0
yarl              | conda-forge/linux-64::yarl-1.7.2-py310h5764c6d_2
zipp              | conda-forge/noarch::zipp-3.8.1-pyhd8ed1ab_0
zlib              | conda-forge/linux-64::zlib-1.2.12-h166bdaf_3

Proceed ([y]/n)? y

Downloading and Extracting Packages
markdown-3.4.1    | 65 KB      |
#####
##### | 100%
```

Users can install into private environments

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ARKANSAS RESEARCH PLATFORM



Conda (Python Modules) – install into a private environment

Package, dependency and environment management for Python (and other languages)

conda install -c <channel> package=<version>=<build>

```
libzlib-1.2.12      | 65 KB      |
#####
grpc-cpp-1.47.1    | 5.1 MB      |
#####
wrapt-1.14.1       | 51 KB      |
#####
readline-8.1.2     | 291 KB      |
#####
libedit-3.1.20191231 | 121 KB      |
#####
zlib-1.2.12         | 92 KB      |
#####
Preparing transaction: done
Verifying transaction: done
Executing transaction: done
Retrieving notices: ...working... done
(tensorflow-pawel) pinnacle-15:pwolinsk:~$ ls .conda/envs/tensorflow-pawel/
bin/                  etc/                  man/                  ssl/
compiler_compat/      include/              sbin/              x86_64-conda_cos6-linux-gnu/
conda-meta/           lib/                  share/              x86_64-conda-linux-gnu/
(tensorflow-pawel) pinnacle-15:pwolinsk:~$ ls .conda/envs/tensorflow-pawel/bin/
2to3                 genbrk                h5dump               kpasswd            pip3
tensorboard
...

```

Users can install into private environments

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