



# RMACC New User Seminar

**Website:** [www.rc.colorado.edu](http://www.rc.colorado.edu)

**Documentation:** <https://curc.readthedocs.io>

**Helpdesk:** [rc-help@colorado.edu](mailto:rc-help@colorado.edu)



Research Computing  
UNIVERSITY OF COLORADO **BOULDER**

# RC Resources for RMACC

## Tech Support

- High Performance Computing
- Data Management

## Human Support

- Training Materials & Workshops
- Consultations & Office Hours
- Help Tickets

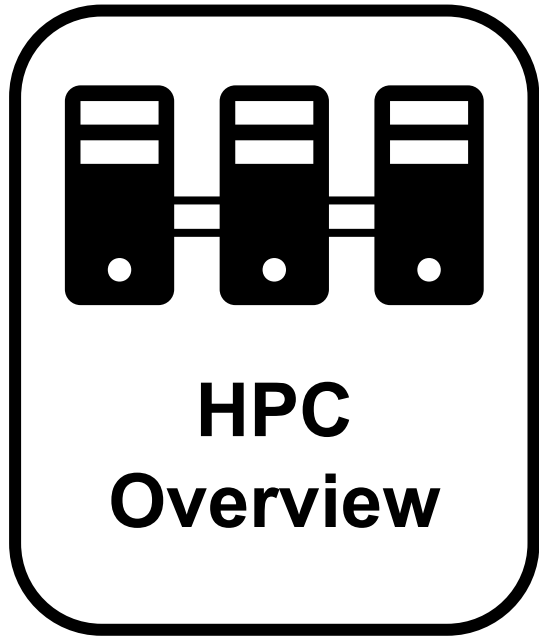


**Ask Questions**

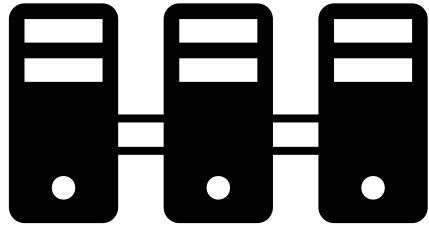


**Discuss Ideas**

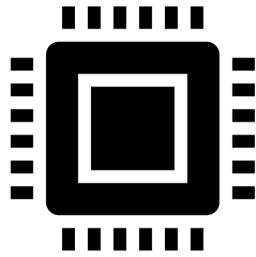
# Learning Goals



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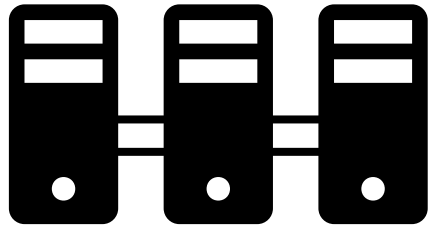


**HPC  
Overview**

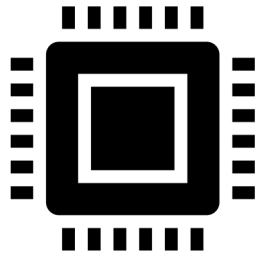


**Nodes &  
Partitions**

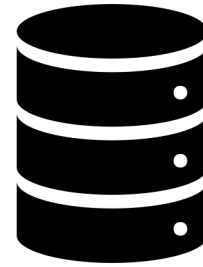
# Learning Goals



**HPC  
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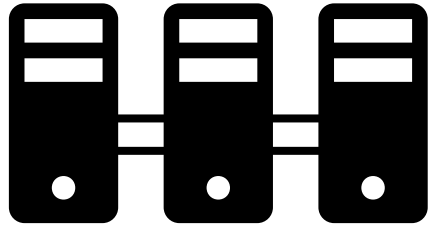
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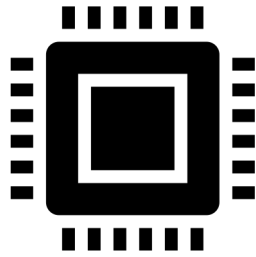
**Data  
Storage**



# Learning Goals



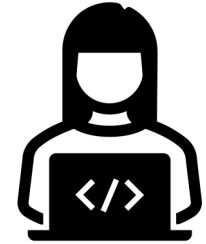
**HPC  
Overview**



**Nodes &  
Partitions**



**Data  
Storage**



**System  
Access**



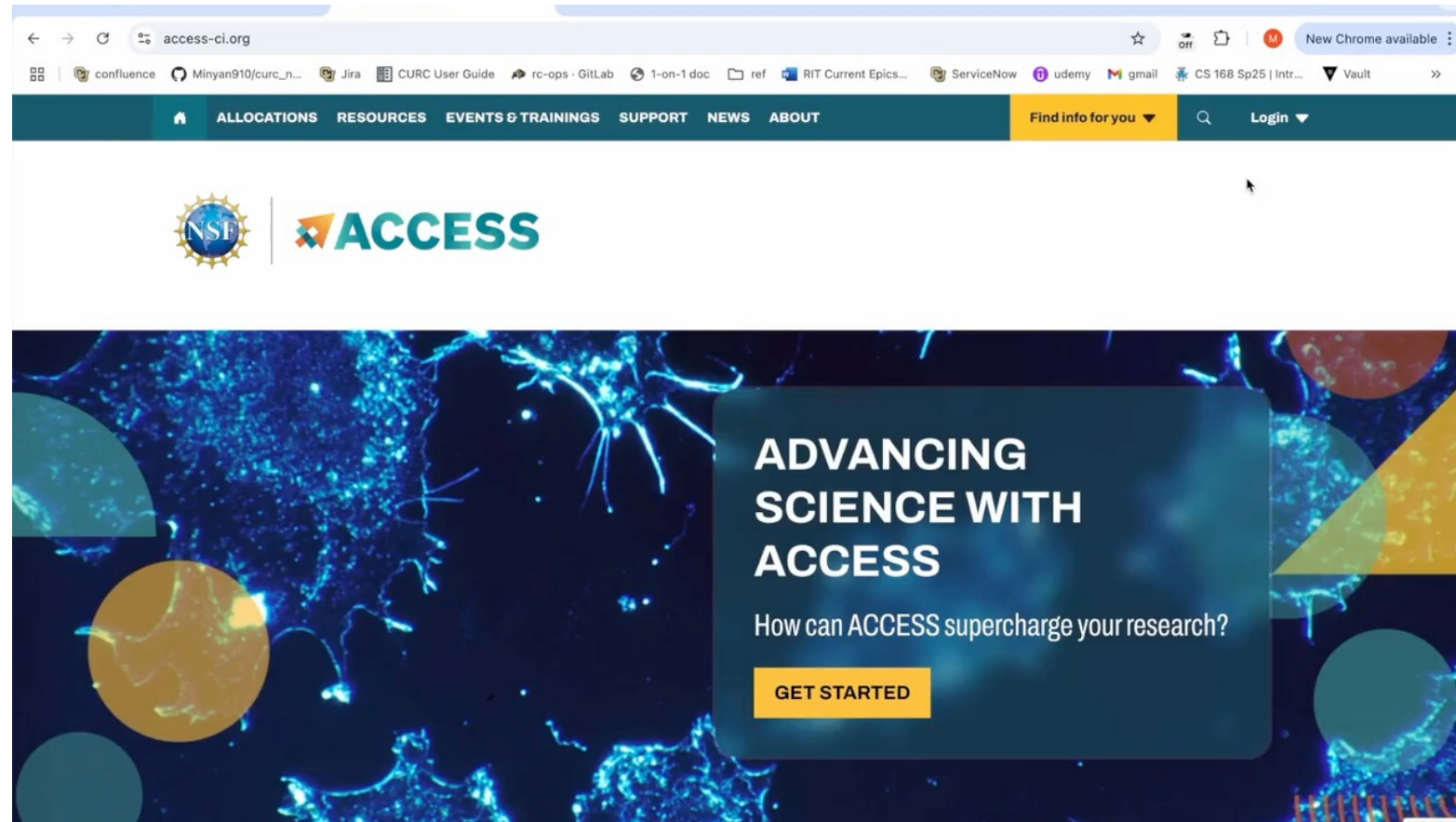
# How to Access RC Resources?

1. Get an RC account
2. Log in
3. Create greatness! (responsibly)

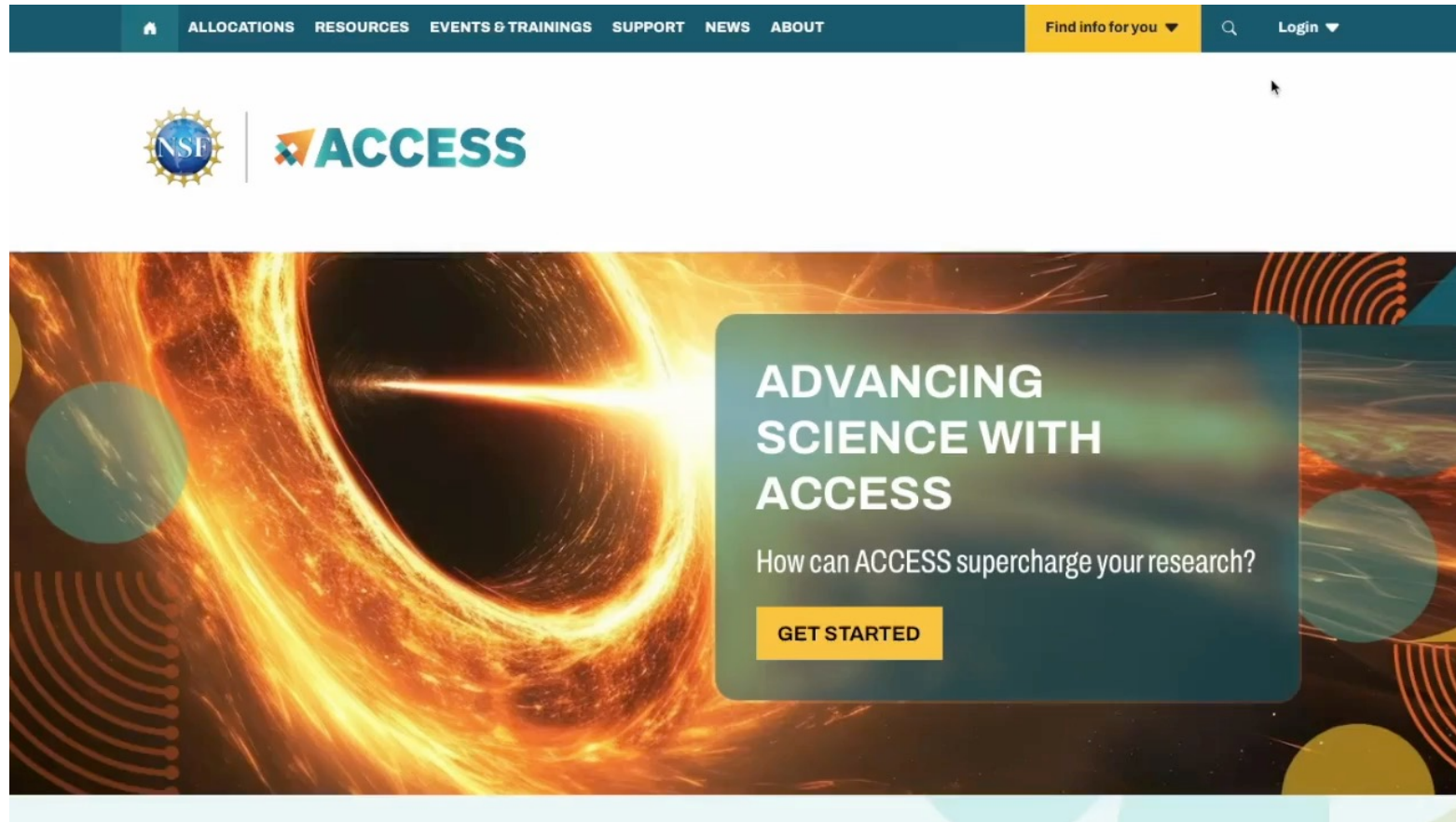
# Getting an Account

- CU Boulder, CSU users and affiliates:
  - Request an account through the RC Account request portal:  
<https://rcamp.rc.colorado.edu/accounts/account-request/create/organization>
- AMC, RMACC users and affiliates:
  - Request an account through the ACCESS-CI User Registration Portal:  
<https://identity.access-ci.org/new-user.html>

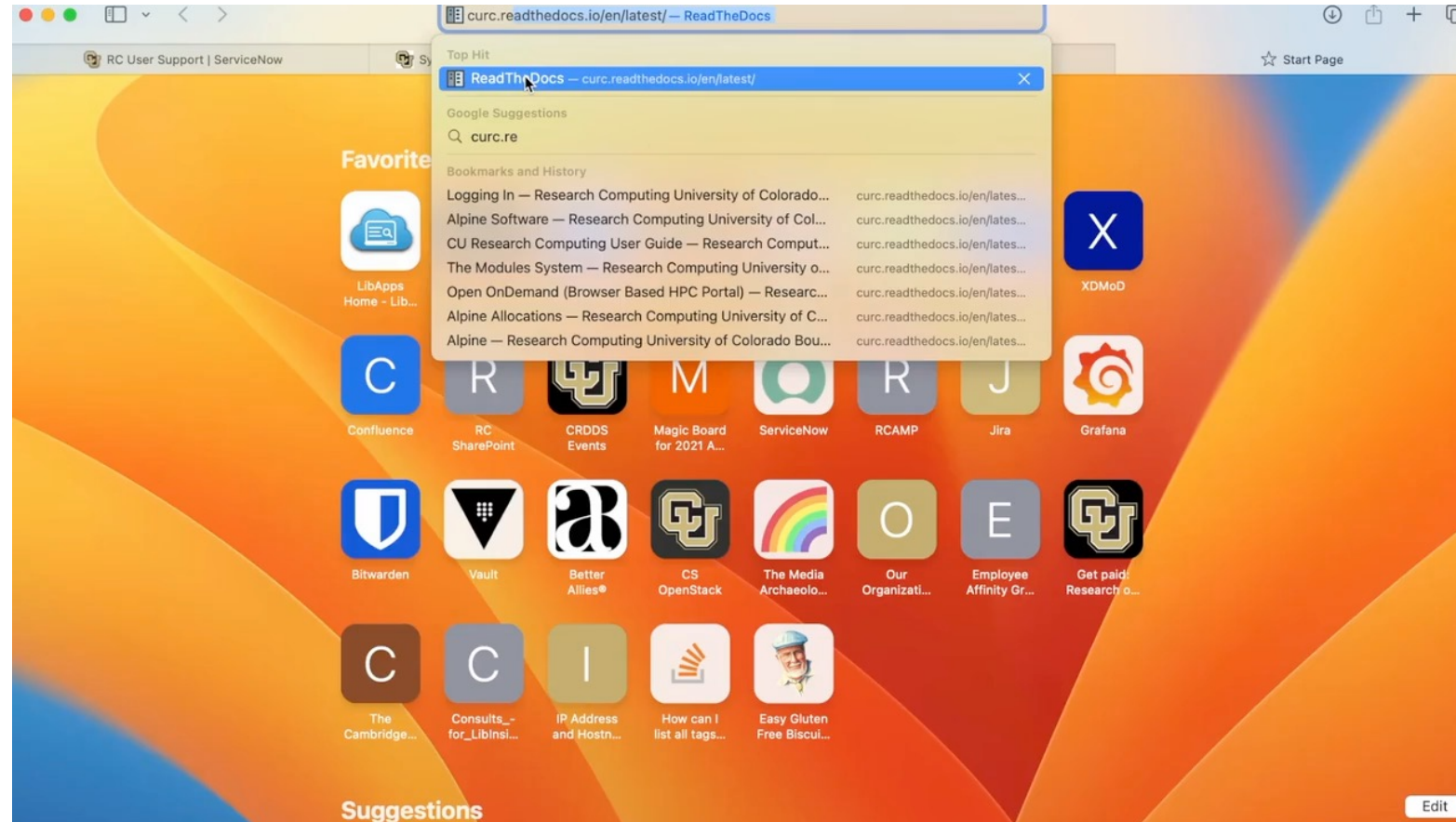
# ACCESS Account Creation (Part 1)



# ACCESS Account Creation (Part 2)



# RMACC Account Form



# Your RC Account

## Access to:

1. Alpine Cluster
2. Core Storage
3. PetaLibrary Storage\*
4. Open OnDemand
5. Approximately 2,000 Service Units (SUs) per month

\*Only Active tier available to non-CU users, if purchased



# High Performance Computing



**Scale**

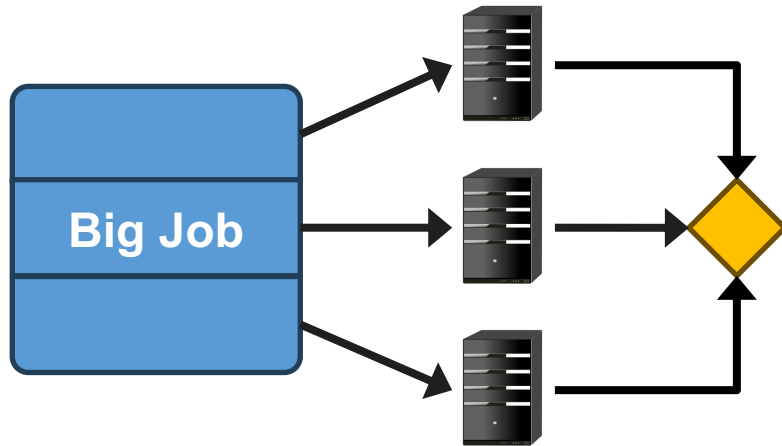
**vs**

**Speed**

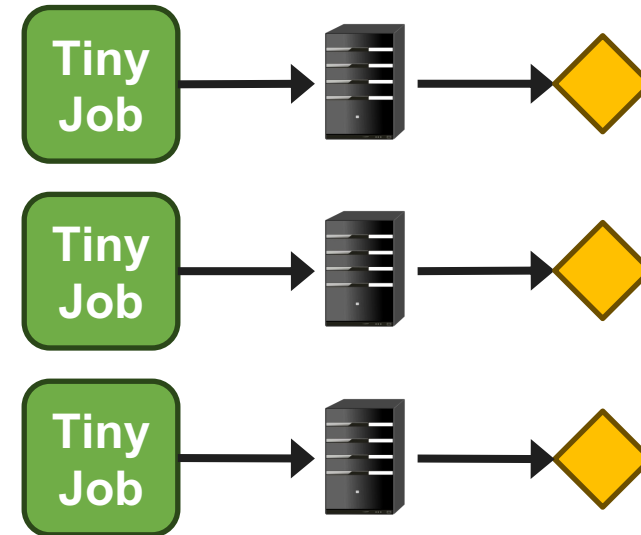




# What can / use HPC for?

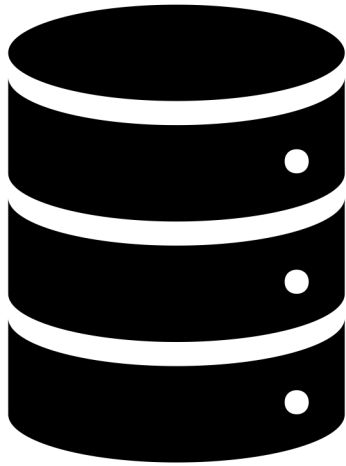


**Parallel Jobs**

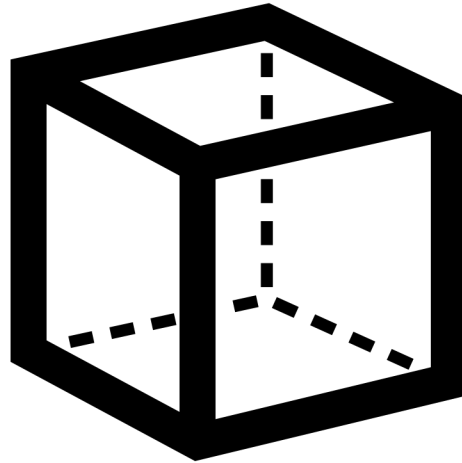


**Serial Jobs**

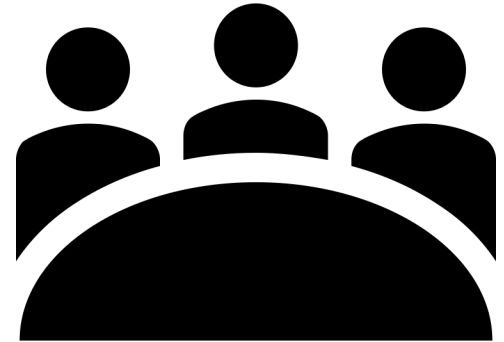
# What can / use HPC for?



**Big Data**



**Viz/Rendering**



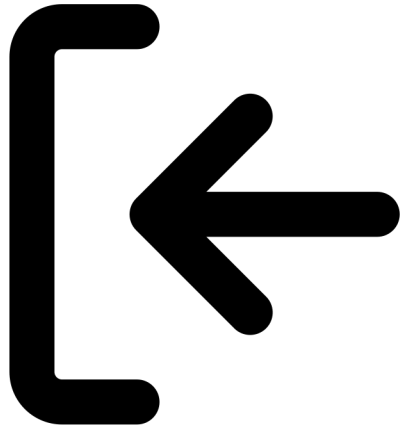
**Shared Work  
Environment**

# HPC Cluster: Alpine

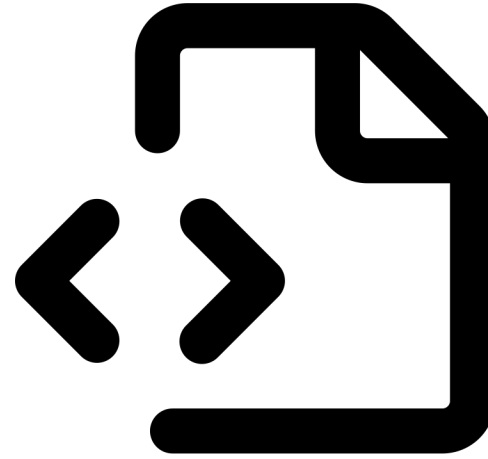


- Heterogeneous cluster
- Hardware provided by CU Boulder, CSU, and AMC
- Access available to CU Boulder, CSU, AMC and RMACC users

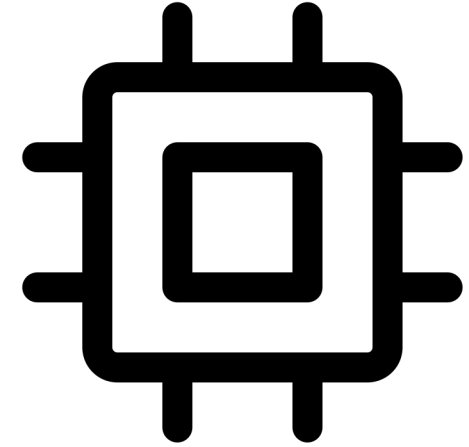
# Node Types



**LOGIN**

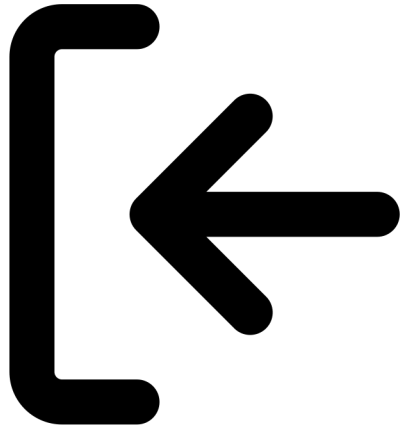


**COMPILE**



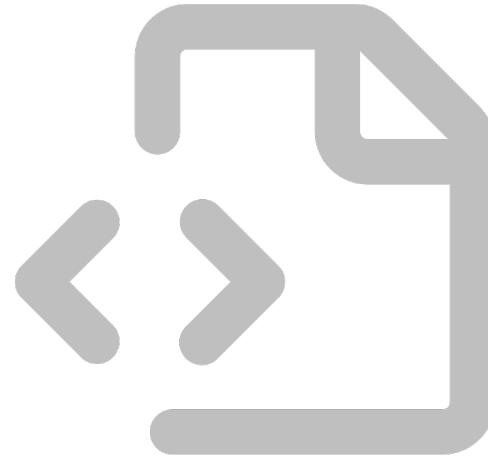
**COMPUTE**

# Node Types

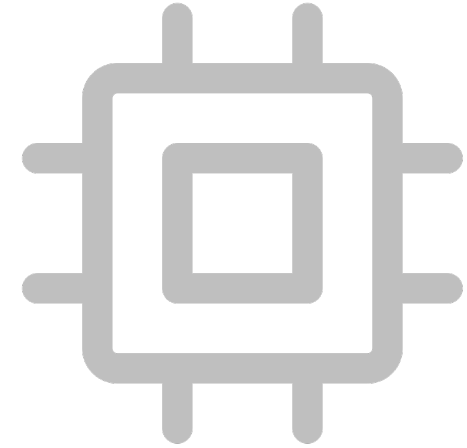


## LOGIN

- Entry to system
- View or edit files
- Submit jobs

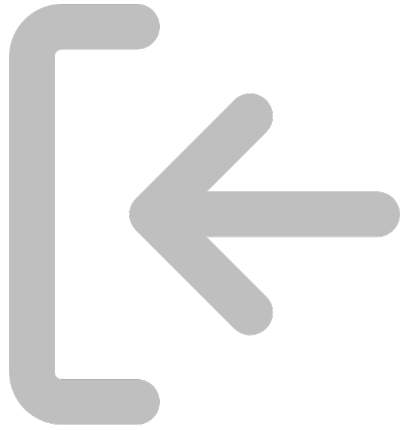


## COMPILE

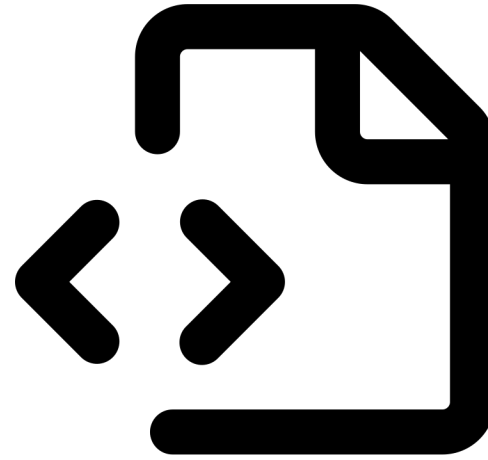


## COMPUTE

# Node Types

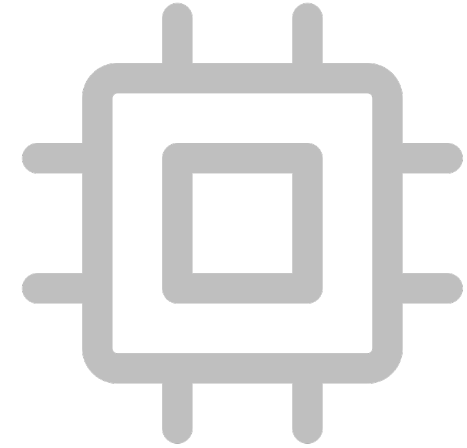


LOGIN



COMPILE

- View or edit files
- Submit jobs
- **Compile code**
- **Install software**

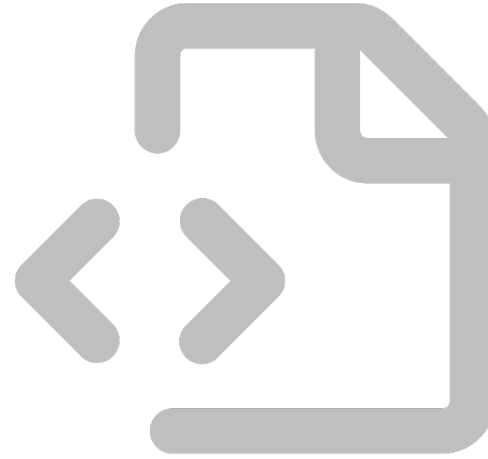


COMPUTE

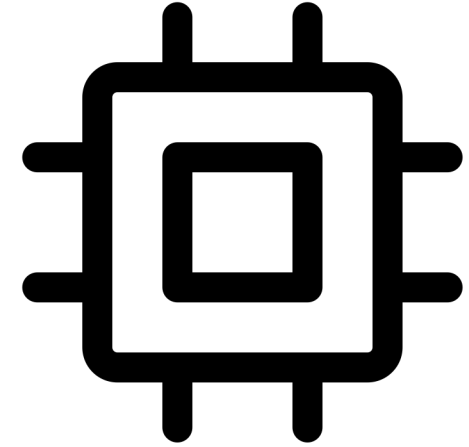
# Node Types



LOGIN



COMPILE

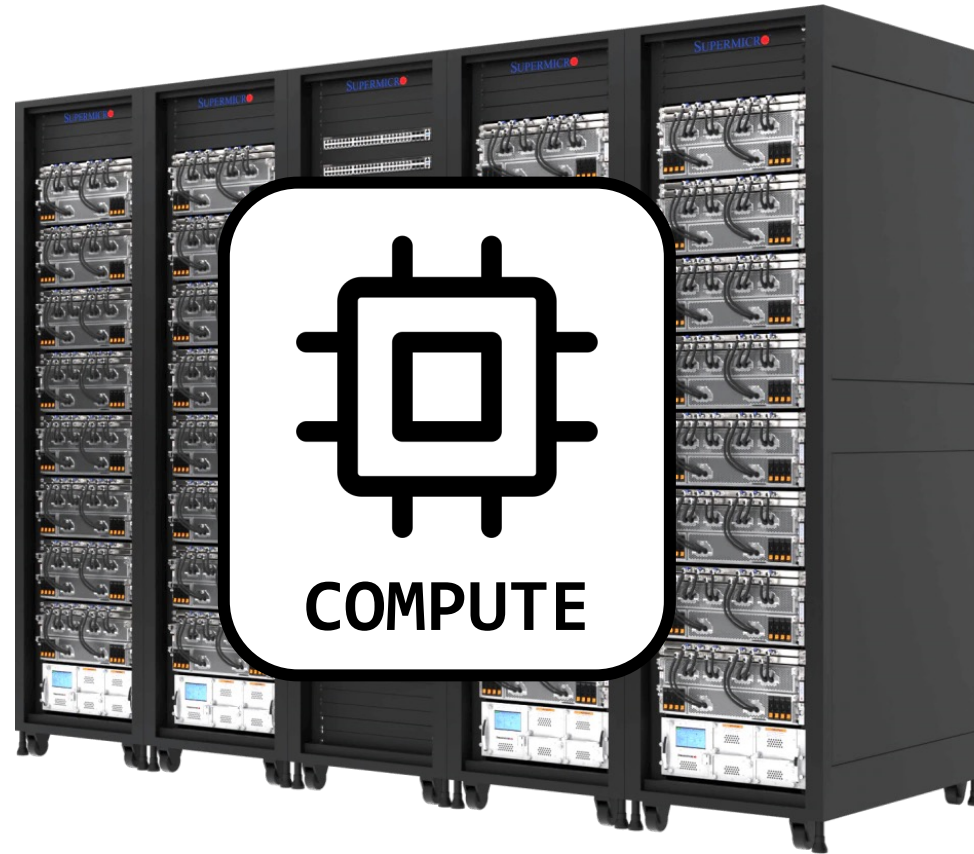


**COMPUTE**

- Run scheduled jobs
- Handle calculations



# Alpine Partitions



# Alpine Partitions

**amilan**  
General Usage



# Alpine Partitions

**amilan**

General Usage

**amem**

High Memory



# Alpine Partitions

**amilan**

General Usage

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**aa100**

Nvidia GPU's

# Alpine Partitions

**amilan**

General Usage

**amem**

High Memory



**aa100**

Nvidia GPU's

**ami100**

AMD GPU's

# Data Storage

## Core

- Personal Storage
- Includes 3 Directories
  - /home (2 GB)
  - /projects (250 GB)
  - /scratch (10 TB)

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## PL

- PetaLibrary
- Tiered Storage
  - Active, Archive
- Requires Funding
- Starts at 1 TB



# Core

## Home

- Personal data
- Config files
- OnDemand data
- Do not share!

**2 GB**

# Core

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- Personal data
- Config files
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- Do not share!

**2 GB**

## Projects

- Code, job scripts
- Installed software
- Shared data

**250 GB**

# Core

## Home

- Personal data
- Config files
- OnDemand data
- Do not share!

**2 GB**

## Projects

- Code, job scripts
- Installed software
- Shared data

**250 GB**

## Scratch

- Job data (in/out)
- Shared data
- Deleted after 90 days

**10 TB**

# PetaLibrary

## Active

- Performance tier
- Accessible by all nodes
- No file limit

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- Integrity tier
- Accessible by data transfer (DTN) and login nodes only
- 10,000 File Limit

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- Same features as Active
- Copy is synced to separate on-campus datacenter every 15 mins.

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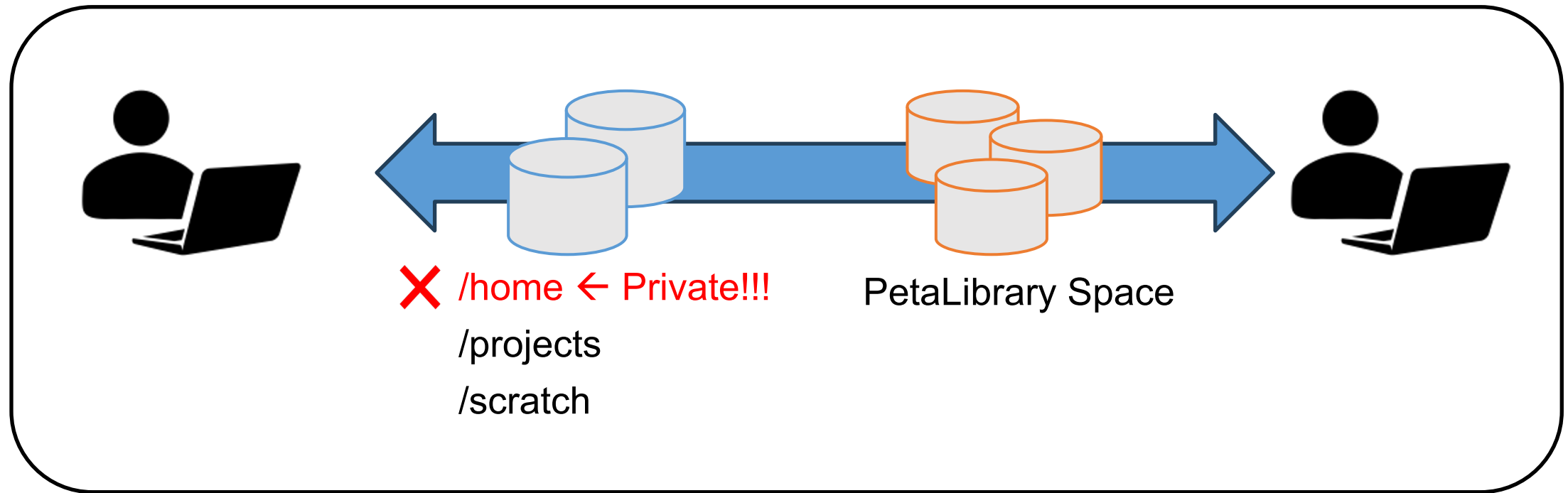
## Archive + DR\*

- Same features as Archive
- Copy is synced to offsite datacenter monthly

\*only available to CU System users



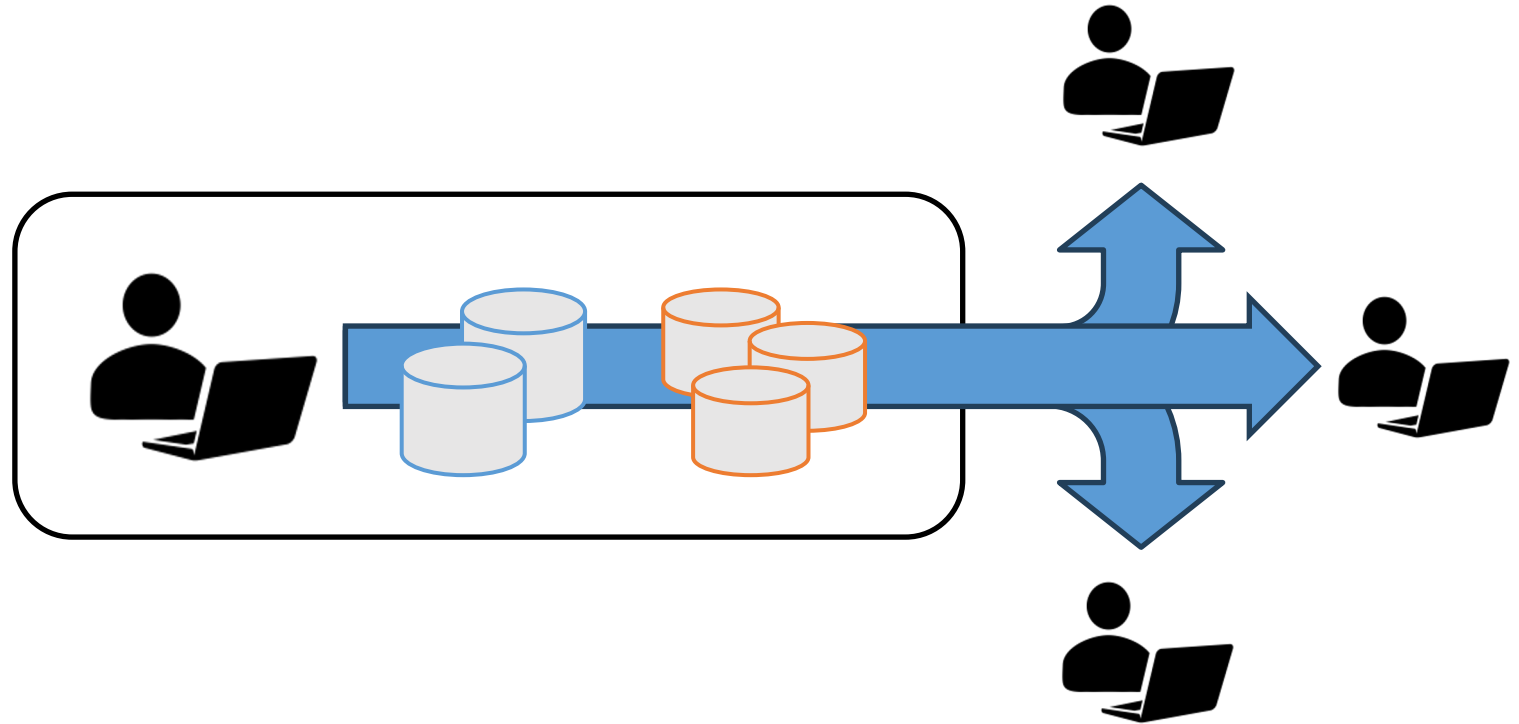
# Data Sharing: Within RC



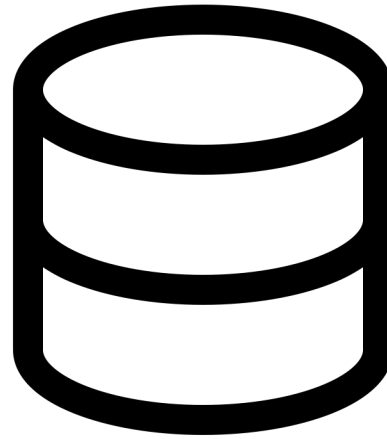
# Data Sharing: Outside RC

## Large Data Transfers:

- Globus (Recommended)
- Data Transfer Nodes (DTN)
- Terminal/Command Line:
  - rsync
  - rclone
  - sftp
  - scp



# Data Transfer Nodes



## DATA (DTN)

- Support data transfers
- Can be selected when using scp, sftp, or ssh transfers

# Acceptable data storage and use

CURC systems and services **should not be used to store** any data that is US government Classified, nor any Controlled Unclassified Information.

For users requiring storage for sensitive data types, please see the secure research computing resources:

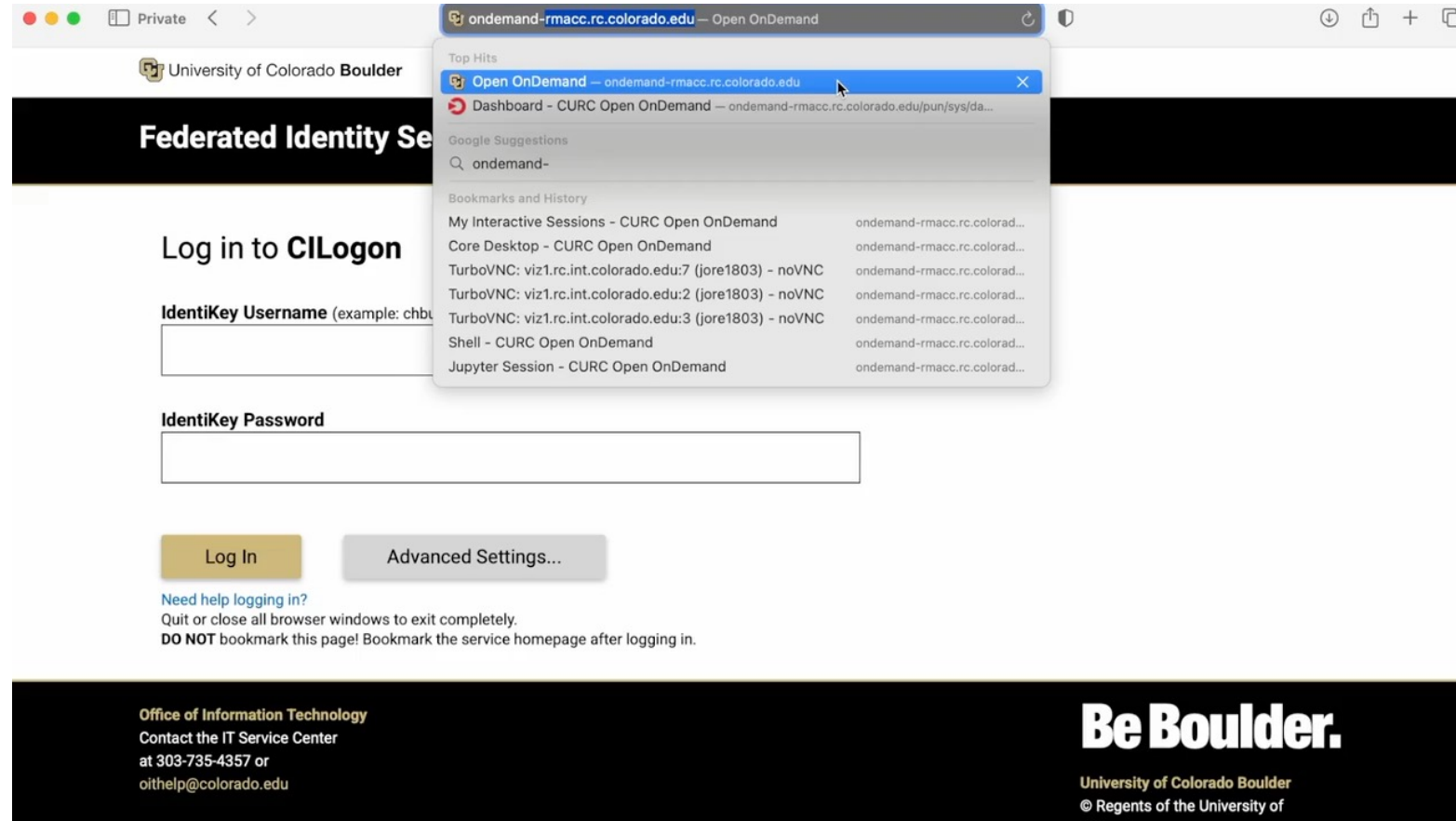
<https://www.colorado.edu/rc/secure-research-computing-resources>

# Demo: logging in with OnDemand

CURC Open OnDemand is a browser based, integrated, single access point for all of your HPC resources at CU Research Computing.

- CU Boulder: Visit <https://ondemand.rc.colorado.edu>.
- Other RMACC Institutions: Visit <https://ondemand-rmacc.rc.colorado.edu/>

# Demo: logging in with OnDemand



# Help! I'm stuck, where do I go?

- **Documentation**: [curc.readthedocs.io/](https://curc.readthedocs.io/)
- **Trainings with Center for Research Data and Digital Scholarship (CRDDS)**: <https://www.colorado.edu/crdds/>
- **Helpdesk**: [rc-help@colorado.edu](mailto:rc-help@colorado.edu)
- **Consult Hours** (Tuesday 12:00-1:00 in-person or virtually, Thursday 1:00-2:00 virtually)



# Help Desk Tickets

To: [rc-help@colorado.edu](mailto:rc-help@colorado.edu)

Dear Research Computing,

Help! My code won't run!  
Help!

Help please,  
John

To: [rc-help@colorado.edu](mailto:rc-help@colorado.edu)

Dear Research Computing,

I am running into issues running my Python script. I am using a conda environment called `my_python_env` with the pytorch software, and I am receiving the following error. I am not sure how to troubleshoot. My job ID is 1620350. Let me know what I can try!

`sbatch: error: Batch job submission failed: Invalid partition name specified.`

Thanks,  
John