## Data Storage and Sharing Best Practices



New User Training February 16, 2024 Lisa Gerhardt Data, AI, And Analytics Group



### File Systems Overview





### The System is a Sum of Many Parts!









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## Simplified NERSC File Systems





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- Users have ultimate responsibility for managing and backing up their data
   <a href="https://docs.nersc.gov/policies/data-policy/policy/">https://docs.nersc.gov/policies/data-policy/policy/</a>









### **Perlmutter File Systems**

#### **Global Home**

- Permanent, relatively small storage
- NOT tuned to perform well for parallel jobs
- Snapshot backups
- Perfect for storing data such as source codes, shell scripts
- cd \$HOME





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#### Global Community File System (CFS)

- Permanent, larger storage
- Medium performance for parallel jobs
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- Perfect for sharing data within research group
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#### **Local Scratch**

- Large, temporary storage
- Optimized for read/write operations, NOT storage
- Not backed up
- Purge policy (8 weeks)
- Perfect for staging data and performing computations
- cd \$SCRATCH









# Long-Term Storage System

#### HPSS

- High-Performance Storage System
- Archival storage of infrequently accessed data
- Hierarchical storage:
  - Data first ingested onto high-performance disk arrays
  - Migrated to large enterprise tape subsystem for long-term retention





### The System is a Sum of Many Parts!









## Perlmutter Scratch (>5TB bandwidth)

- Store data being actively read or written by jobs on computes
- Directories are user-readable and writable by default
- Purged! Back up any important data
- Quotas are 20TB (soft) and 30TB (hard).
   After you exceed the hard quota, you will not be able to write any more data to the file system

**How to:** move your run scripts into scratch and have your application set up to read/write data within scratch (e.g. file paths to locations within scratch)







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**How to:** move your run scripts into scratch and have your application set up to read/write data within scratch (e.g. file paths to locations within scratch)

Advanced users may want to optimize data striping:

	Single Shared-File I/O	File per Process		
File size (GB)	command			
< 1	keep default striping	keep default striping		
1 - 10	stripe_small	keep default striping		
10 - 100	stripe_medium	keep default striping		
> 100	stripe_large	keep default striping		
> 1000	stripe_large	stripe_large		
https://d	ocs.nersc.gov/perfo	rmance/io/lustre/		

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### **Global Common: Software Filesystem**

• For: software stacks - Why? Library load performance, and enhanced caching







## **Global Common: Software Filesystem**

• For: software stacks - Why? Library load performance, and enhanced caching



- Group writable directories similar to community, but with a smaller quota, /global/common/software/<projectname>
   Write from login pode: read only on compute pode
  - Write from login node; read-only on compute node
- Smaller block size for faster compiles than CFS





### **Community File System**

- For: large datasets that you need for a longer period
- Set up for sharing with group read permissions by default
- Not for intensive I/O use Scratch instead

How to: cd \$CFS/<projectNumber>

mkdir <name\_of\_choice>

https://docs.nersc.gov/filesystems/community/







## Community File System

- Use the "dvs\_ro" (/dvs\_ro/cfs) mount if you're reading from CFS during jobs
- Data is never purged
- Projects can split their space allocations between multiple directories and give separate working groups separate quotas

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#### **HPSS**





https://docs.nersc.gov/filesystems/archive/



#### HPSS

- For: data from your finished paper, raw data you might need in case of emergency, really hard to generate data
- HPSS is tape!
  - Data first hits a spinning disk cache and gets migrated to tapes, cache is sized for several days of retention
  - Files can end up spread all over, so use htar to aggregate into bundles of 100 GB - 2 TB
  - Archive the way you intend to retrieve the data
  - hsi and htar give the best performance within NERSC
- Quotas are controlled in Iris. If you're a member of multiple projects you can adjust the percentage you want charged to each



https://docs.nersc.gov/filesystems/archive/



#### **Home Directories**

- For: source files, scripts for **testing**, notes
- 40G quota
- Not intended for intensive I/O (e.g. application I/O) use Scratch instead
- Backed up monthly by HPSS
- Snapshots are also available e.g. my homedir is at /global/homes/.snapshots/2022-06-14/e/elvis



### General Advice for I/O

- I/O from batch jobs should go to Perlmutter's scratch file system (/pscratch, \$SCRATCH)
  - Input data
  - Configuration files
  - Output data







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  - Conda environments
  - Anything you install with config / make / cmake etc.



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  - Anything you install with config / make / cmake etc.
- Don't generate a million small files, especially not in one directory
- Aggregating reads and writes into bigger pieces is generally better





#### **Best Practices for DVS**

- DVS is an I/O forwarder
- Uses a set of 24 nodes to forward I/O and offer high performance







### **Best Practices for DVS**

- DVS is an I/O forwarder
- Uses a set of 24 nodes to forward I/O and offer high performance
- Conda environments should be in a container or global common
  - By default they install to your home dir, which causes **A LOT** of problems at scale
  - Also, if you load a conda environment at login, ALL of the very large number of library paths are dragged along to your slurm job. Consider whether you want this or not
  - Python automatically adds your current working directory to the library load path
- Best choice for large scale I/O is always scratch!
- If your data is too large and you need to read it off of CFS, use "/dvs\_ro" instead of "/global"
  - "/global/cfs/cdirs/myproject/mega\_important\_config" ->
    - "/dvs\_ro/cfs/cdirs/myproject/mega\_important\_config"
- Avoid ACLs on files over DVS. These keep the system from using any caching and slows things down







#### Data Management Tools





### Adjusting Quotas in IRIS









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### PI Toolbox: <u>my.nersc.gov/pitools/</u>

PI To	oolb	ox							
Jump	Jump to das directory in cfs			Make entire proje	ect dir group-r	eadable	_		
Path: /global/cfs/cd		obal/cfs/cd All (But sele	Owning group can: Read (r) Write (w)				efresh file list		Show
Sel	ect	Name	Execute file, enter direct	ory			Size	Date	Permissions
Ē	Par	Make directory group openable and executable files group     executable (X)							
		.ipynb	<ul> <li>Execute binary file as member of owning group and force new items in directory to be owned by the group (s)</li> <li>Execute binary file normally, as member of user's default group (x)</li> <li>I want to apply these permissions recursively</li> </ul>				4096	Jul 18 13:14	drwxr-xr-x
	D	DaskE					6326	Sep 12 10:20	-rw-rr
		MODS					4096	Jan 18 15:32	drwxrwxr-x
		agrein					4096	Jul 22 18:23	drwxrwxrwx
		backu	Can	request		4096	Nov 18 13:35	drwxrwxr-x	
		canon			canon	das	4096	Aug 21 10:32	drwxrwx
		certs.ne	ersc.gov		dastest	das	4096	Sep 9 16:03	drwxrwxr-x
		dfulton			dfulton	das	4096	Aug 21 11:14	drwxrwxr-x







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#### Data Dashboard in my.nersc.gov

#### Data Dashboard

Showing disk space and inode usage for global directories at NERSC to which you have access as PI, PI proxy, or user (includes /cfs, /dna, and /projectb)





#### Data Sharing Best Practices





- Community File System: CFS
  - Every project has at least one directory that has permissions set up to be group writable and readable
  - PI Toolbox (<u>my.nersc.gov/pitools/</u>) can manage permissions
- HPSS Project Directories
  - Directories in HPSS with group writable and readable permissions

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- Scratch
  - User who desire to share data on scratch can do it by adjusting Linux permissions
    - Only share read access. If you want to allow writes, we recommend using a collaboration account instead
    - chgrp -R <project\_name> \$SCRATCH; chmod g+rX \$SCRATCH (read only)

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#### • Scratch

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  - Only share read a collaboration acco
  - chgrp -R <proj</pre>
- **How to:** give -u <receiving\_username> <file or directory> take -u <sending\_username> <filename>

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only)

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### Sharing with External Collaborators

#### • Public HTML access

- Project specific area can be created:
  - /global/cfs/cdirs/<yourproject>/www
- These are available for public access under the URL:
  - https://portal.nersc.gov/project/<yourproject>/
- Science Gateways (<u>docs.nersc.gov/services/science-gateways/</u>)
  - Web portals allow you to interface with your data and computation at NERSC
  - For more sophisticated web applications: **Spin** (<u>docs.nersc.gov/services/spin/</u>)
- Globus Sharing (<u>docs.nersc.gov/services/globus/#globus-sharing</u>)
  - Projects can set up read-only endpoints for sharing data with certain Globus users
  - Excellent way to share large volumes of data, can be incorporated into web pages

#### NERSC's Dedicated Data Transfer Nodes

- Data Transfer Nodes (DTNs, https://docs.nersc.gov/systems/dtn/)
  - Dedicated servers for moving data at NERSC (dtnXX.nersc.gov)
  - Servers include high-bandwidth network interfaces & are tuned for efficient data transfers
    - Monitored bandwidth capacity between NERSC & other major facilities such as ORNL, ANL, BNL, SLAC...
  - Direct access to Community, HPSS Archive

#### How to (for small-ish files):

(logged onto the system you want to move the files to)

```
scp <username>@dtn0[1-4].nersc.gov:<path/to/file> <local_path>
```

scp lgupta@dtn01.nersc.gov:\$HOME/script.sh .







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  - Direct access to Community, HPSS Archive
- Use NERSC DTNs to move large volumes of data in and out of NERSC or between NERSC systems
- User Perlmutter Login nodes for data transfers to Perlmutter Scratch





# General Tips for Transferring Data: Globus

The **recommended** tool for moving data in, out & within NERSC

- Reliable & easy-to-use web-based service:
  - Automatic retries
  - Email notification of success or failure
- Accessible to all NERSC users



- NERSC-managed endpoints on DTNs for optimized data transfers
- Web based GUI for drag-and-drop transfers
- NERSC Globus scripts for command line transfers
- REST/API for scripted interactions with service
- Globus Connect Personal for setting up endpoints on your laptop

#### https://docs.nersc.gov/services/globus/





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- Globus Copy full Personal for

#### https://docs.nersc.gov/servi ces/globus/

Examples

Transfer files from NERSC's Community file system to NERSC's Perlmutter Scratch file system

This can be used to stage data on Perlmutter scratch before using it in a running job. See the script stage\_data.script (included in the globus-tools module) for an example of how to do this.

First, generate a list of files and directories you wish to transfer. If a directory is included in this Email notification of succes list, its contents will be recursively transferred to the target directory.

> fglobal/cfs/cdirs/<myrepo>/<my\_dataset\_directory> > transfer.txt global/cfs/cdirs/<myrepo>/<my\_other\_dataset\_directory>/data01.dat >> transfer.

transfer\_files.py -s dtn -t perlmutter -d /pscratch/sd/<letter>/<your\_username>/in

transfer information:

Please go to this URL and login: https://auth.globus.org/v2/oauth2/authorize?client Please enter the code you get after login here: <snipped> 

You can check on the status of the transfer with the check\_transfer.py script

nersc> module load globus-tools 



### **Performance Considerations**

- Performance is often **limited by the remote endpoint** 
  - Not tuned for WAN transfers or have limited network link
  - These can lower performance <100 MB/sec.
- File system contention may be an issue
  - Try the transfer at a different time or on a different FS.
- Don't use your \$HOME directory for I/O!

Instead use CFS, \$SCRATCH ...

 If you think you are not getting the transfer rates you expect, let us know: <u>help.nersc.gov</u>







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Thank You and Welcome to NERSC!

