Data Management in the HPC

Tools and Workflows

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osf.io/98bzd/

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What you will learn

Part 1: learned about policies and basic tools/approaches for data management

Part 2: Learn about additional tools for transferring, organizing, and archiving HPC data.

Why: Setting up workflows means better collaboration, reproducibility => better research

Assume

- Familiarity with the Unix shell

Sample Transparent and Reproducible Research Pipeline

Basic workflow – less reproducible

Compute

Google Drive

Get data

Export

Google Drive

Data, Figures

Advanced workflow – More Reproducible

docker

Get

Manage

GitHub

Manage

Open Science Framework

Manage

MENDELEY

box

Dropbox

Collab w/ Co-authors (manuscripts, data)

ReDATA

arizona.figshare.com

Publish & archive

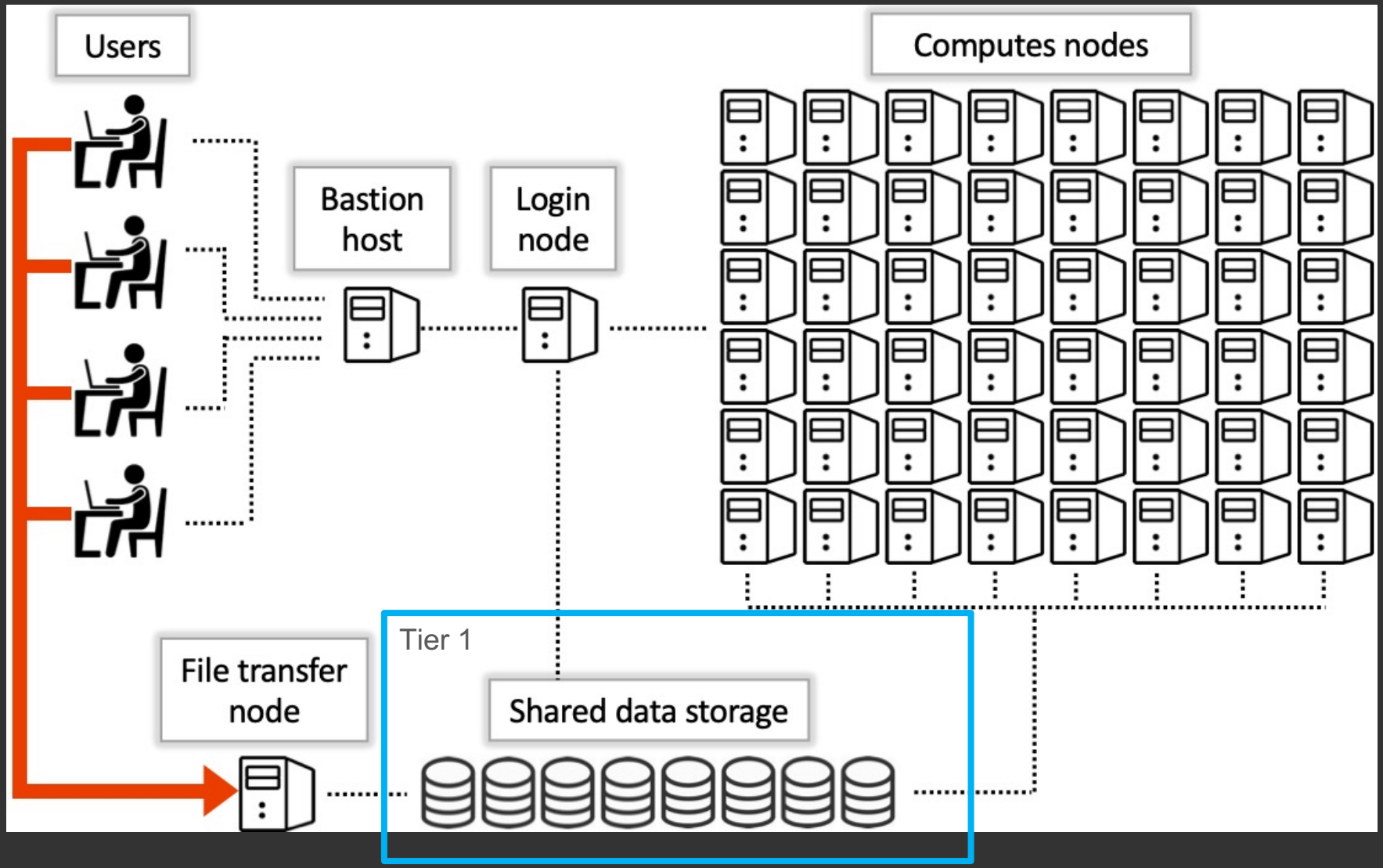
Get Data

THE UNIVERSITY OF ARIZONA

UA Storage Refresher

Tier	Uses	Examples
Tier 1 – HPC storage	High performance. Active research. **Limited permanent storage**.	Home directory PI allocations.
Tier 2 – General storage	Less frequent access Copy subsets as needed to tier 1. **Backups**	Google Drive Box Your computer
Tier 3 - Archival storage	Store data after project completion **Publish data that supports publications**	UA ReDATA Amazon Glacier

HPC Storage Refresher



Getting data into the HPC

Many ways to do it…

Small, infrequent transfers

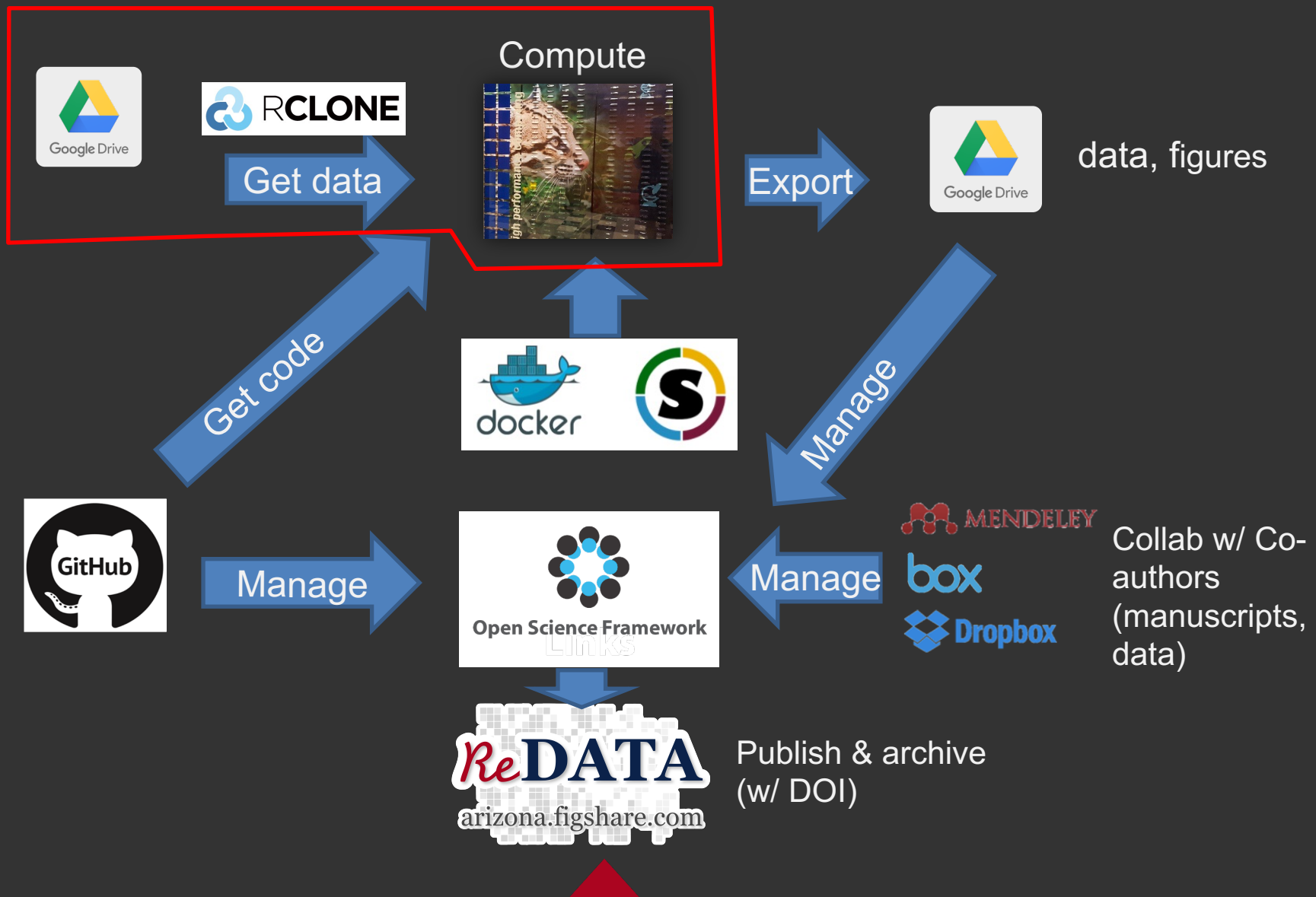
General purpose

Large transfer >100GB, scheduled transfers, transfer outside UA

Rclone

- Mature software for working with cloud storage
- Mirroring, syncing, encryption, union and more
- Supports 40+ providers
  - Gdrive
  - Box
  - Amazon S3
  - OneDrive
  - SFTP
  - ownCloud
  - …

Tier 2 to Tier 1 transfer



Basic Rclone workflow

The sample data is also available on the OSF osf.io/7rbpd

- Log in to HPC
  - ssh <netid>@filexfer.hpc.arizona.edu
- rclone config
  - Create the remote UA-gdrive – see rclone documentation
- rclone lsf UA-gdrive:'/OSF/HPC Demo'
  - Test the connection
- rclone copy UA-gdrive:'/OSF/HPC Demo' .

The dot means copy to the current folder

Advanced Rclone workflow

Mounting: Access Google as if it were a local folder (e.g., can use standard commands like ls, cat, cp).

- Configure the remote (if not configured already)
- Request an interactive session on a cluster
  - Mounting doesn't work on filexfer or login nodes
- rclone mount UA-gdrive:'/OSF/HPC Demo' ~/Desktop/mount/ &
  - Mount into a folder (e.g., mount)
- cd ~/Desktop/mount ; ls
  - Test the connection
- fusermount -uz ~/Desktop/temp/directory
- Note: lots of caveats!! See rclone docs

Now what?

File Management

Need a strategy before doing anything else

Data Management Best Practices

In Part 1 we covered versioning and file/folder organization

Better

```
Study001_Raw
└ BiopsyData
  ├ 20161101_Study001_Biopsy_visit1.xls
  ├ 20161101_Study001_Biopsy_visit1_v2.xls
  ├ 20161101_Study001_Biopsy_visit1_v3.xls
```

Document the naming scheme

Add version and/or date (ISO 8601)

Descriptive folder names

Bad

```
GW_model
  ├ elevation.mat
  ├ depth_wt.csv
  ├ well_loc.csv
  ├ flow_model.m
  ├ flow_model2.m
  ├ flow_model_final.m
  ├ flowlines1.png
  ├ flowlines2.png
  ├ contours.png
```

Better

```
GroundwaterModel
└ Code
  ├ 20170402_FlowModel_v1.m
  ├ 20170410_FlowModel_v2.m
  ├ 20170511_FlowModel_v3.m
└ Inputs
  ├ TerrainElevation.m
  ├ DepthToWaterTable.csv
  ├ WellLocations.csv
└ Outputs
  ├ 20170402_Flowlines_FlowModelv1.png
  ├ 20170402_Contours_FlowModelv1.png
  ├ 20170415_Flowlines_FlowModelv2.png
```

Project Setup

- Do you have a more complex project?

https://github.com/mkrapp/cookiecutter-reproducible-science

Basic Metadata

Each stage of data in it's own folder

Software in it's own folder

```
.
├── AUTHORS.md
├── LICENSE
├── README.md
├── bin                <- Your compiled model code can be stored here (not tracked by git)
├── config             <- Conf                                  doxygen or for your model if needed
├── data
│   ├── external       <- Data from third party sources.
│   ├── interim        <- Intermediate data that has been transformed.
│   ├── processed      <- The final, canonical data sets for modeling.
│   └── raw            <- The original, immutable data dump.
├── docs               <- Documentation, e.g., doxygen or scientific papers (not tracked by git)
├── notebooks          <- Ipython or R notebooks
├── reports            <- For                                   LaTeX, Markdown, etc., or any project reports
│   └── figures        <- Figu                                  reports
└── src                <- Sour
    ├── data           <- scripts and programs to process data
    ├── external       <- Any external source code, e.g., pull other git projects, or external libraries
    ├── models         <- Source code for your own model
    ├── tools          <- Any helper scripts go here
    └── visualization  <- Scripts for visualisation of your results, e.g., matplotlib, ggplot2 related.
```

Cookiecutter

- Install: See instructions on OSF (osf.io/9ceqd).
- cookiecutter gh:mkrapp/cookiecutter-reproducible-science

```
(puma) frios@r2u06n1 ~$ cookiecutter gh:mkrapp/cookiecutter-reproducible-science
You've downloaded /home/u17/frios/.cookiecutters/cookiecutter-reproducible-scienc
ownload it? [yes]: yes
full_name [Mario Krapp]: Fernando Rios
email [mariokrapp@gmail.com]: frios@arizona.edu
github_username [mkrapp]: zoidy
project_name [Name of your science project]: HPC Demo 2021
project_slug [hpc-demo-2021]:
project_short_description [A short description of your project]: Hello world!
release_date [2021-10-04]:
version [0.1.0]: 1.0
(puma) frios@r2u06n1 ~$
```

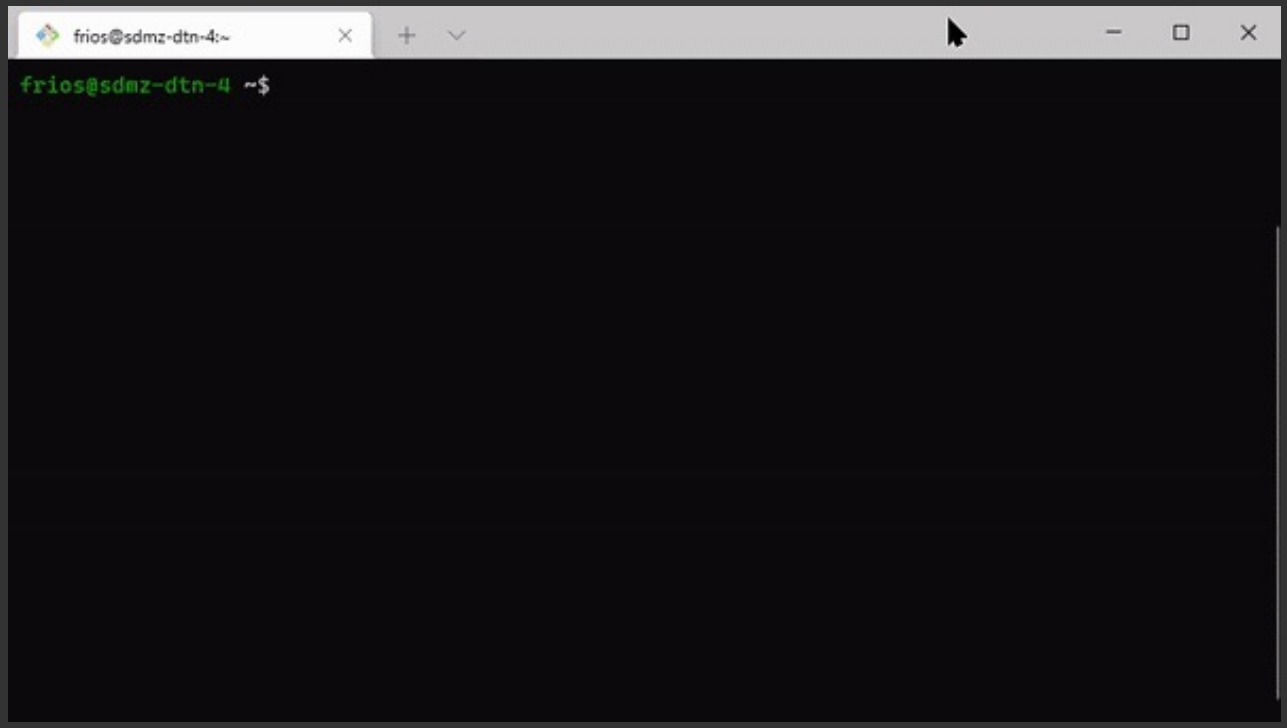
Other templates
https://github.com/luke-gregor/cookiecutter-science-pub
https://github.com/mnarayan/cookiecutter-data-science

File & Space Management Tools

- Checking your space & file limit: uquota

```
frios@login2 ~$ uquota
                                    used  soft limit  hard limit       files/limit
frios home & PBS                 46.84M         14G         15G        1044
/extra/frios                     56.72G        200G        200G           2/120000
```

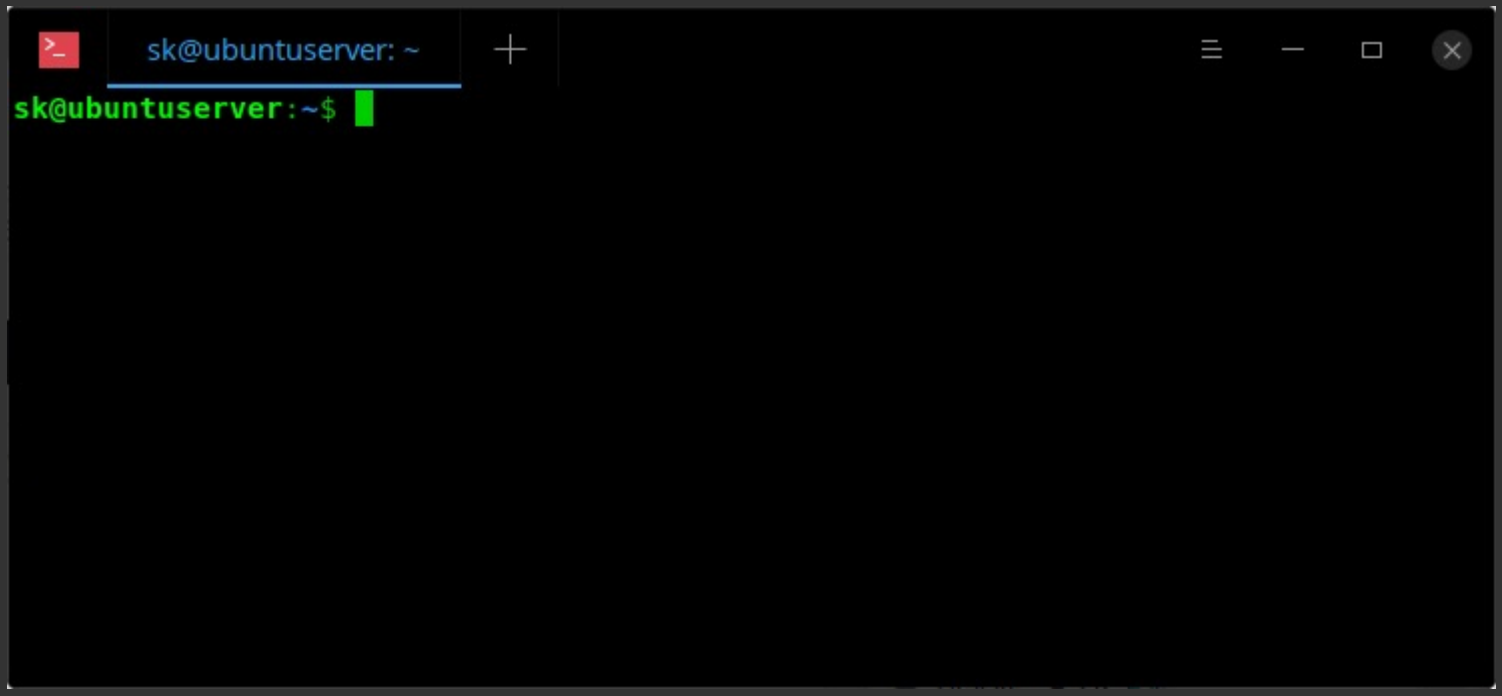
- Checking folder usage and count: NCDU



HPC installation instructions and pre-compiled binaries: https://osf.io/98bzd/

File & Space Management Tools

- Keeping file names tidy with renameutils



Pre-compiled version for HPC https://osf.io/98bzd/

Credit: ostechnix.com

Data Mgmt Best Practice: Storage & Backup

"I decide what data is important while I am working on it and typically save it in a single location"

Do

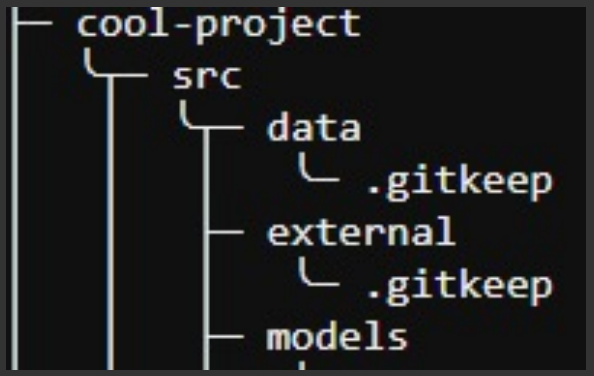
- 3-2-1: If possible, 3 copies, 2 different storage types, 1 copy offsite
- Keep offline backups if possible. Sync clients could propagate changes unintentionally

Avoid:

- Storing sensitive data on an unencrypted laptop or flash drive or insecure servers
- Relying on cloud storage for the only copy!
  http://www.cnet.com/news/dropbox-fixes-file-deletion-bug-offers-year-of-free-service/

For HIPAA compliance, use UA Box Health account.

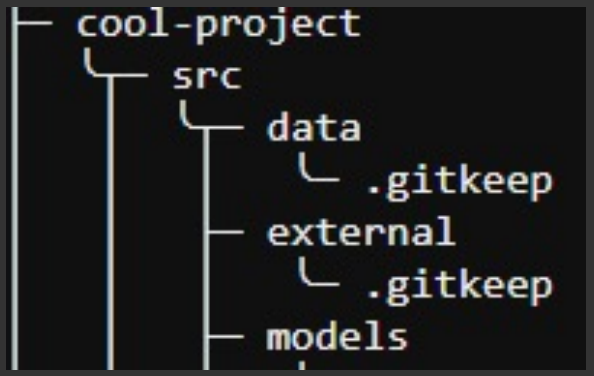
Backup and Restore to Google Drive (Tier 1 to Tier 2)



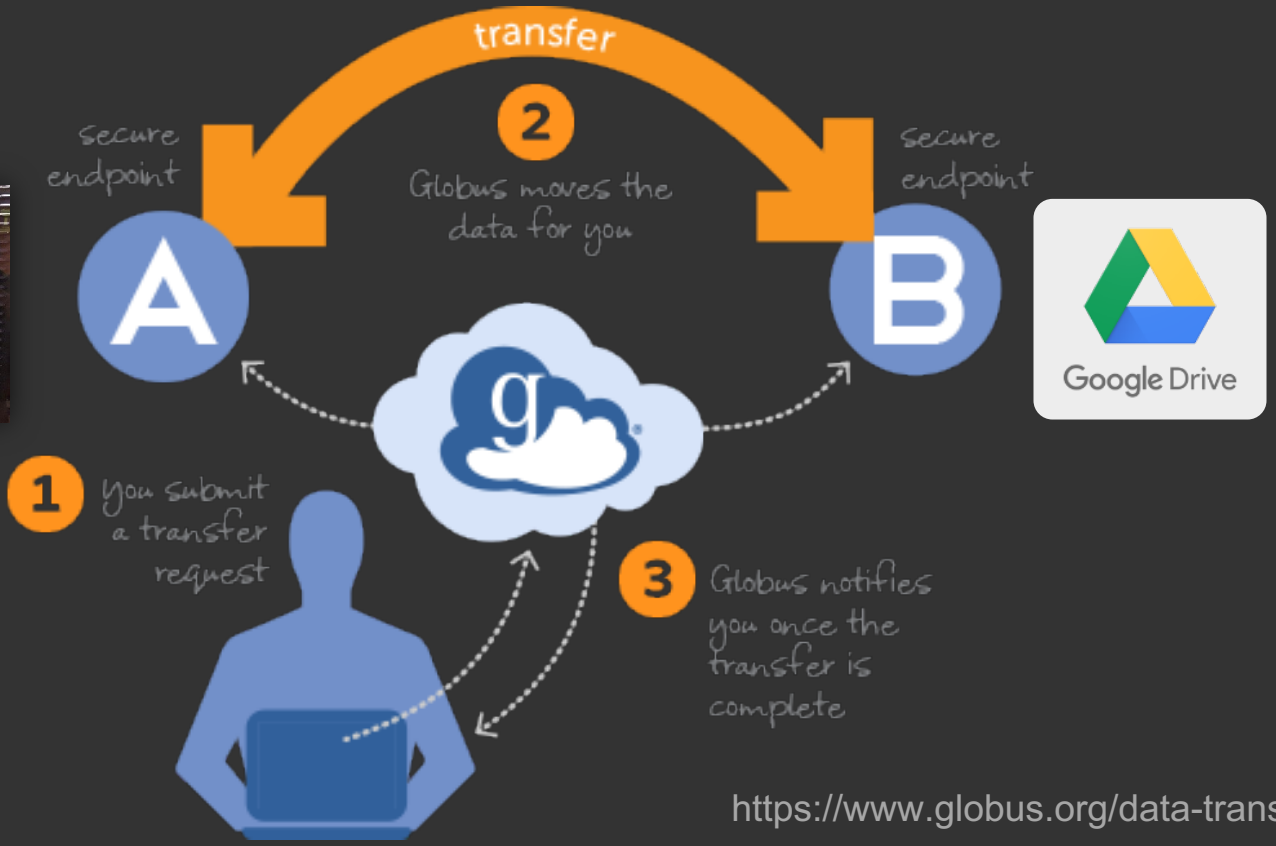
- Rclone to transfer directly to/from Google Drive, Dropbox, S3, Box… many more
- TIP: Transferring lots of little files is slow. Put everything in a tar archive

Finish Project and Export

- Export to Google Drive via Globus



Globus

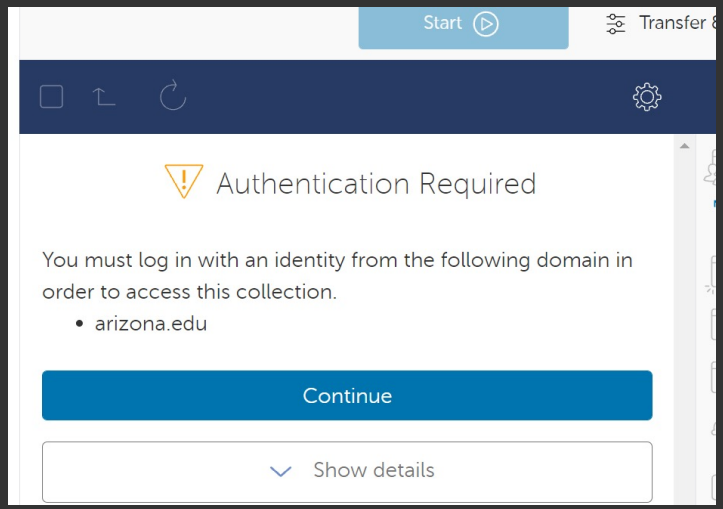


https://www.globus.org/data-transfer

- Why Globus?
  - Designed for large data
  - Reliable – supports resuming
  - Can initiate remotely, no babysitting transfers
    - Email on completion

Globus Demo

- Go to https://www.globus.org/ and log in
- In the File Manager tab,
  - Click the left hand Search box and search for the collection “UA HPC Filesystems”
  - In the box on the right, search for “UA Google Drive”
- You may have to authorize a bunch of requests



public.confluence.arizona.edu/display/UAHPC

Scheduled/recurring transfers using Globus

- Beta Globus Timer https://pypi.org/project/globus-timer-cli/
- pip3 install --user globus-timer-cli
- globus-timer session login

```
(elgato) frios@cpu1 ~$ globus-timer session login
Please log into Globus here:
--------------------------
https://auth.globus.org/v2/oauth2/authorize?client_id=bc7
th.globus.org%2Fv2%2Fweb%2Fauth-code&scope=profile+email+
wgYS81mcVAwZSrzsfgMhZEQH3qmOd5V-d-LM&code_challenge_metho
mand+Line+Interface+on+cpu1.elgato.hpc.arizona.edu
--------------------------

Enter the resulting Authorization Code here:
```

Scheduled/recurring transfers using Globus

- Set up the transfer. Transfer some files from xdisk to Google Drive (may need to authorize up to 3x)

```
globus-timer job transfer \
    --name my-job \
    --label "Timer Transfer Job" \
    --interval 120 \
    --start '2021-10-04T12:57:00' \
    --source-endpoint 7c4462b2-7ca4-4f44-820a-xxxxxxxxxxxx \
    --dest-endpoint 26b96369-5f03-4742-9ab8-xxxxxxxxxxxx \
    --verify-checksum \
    --sync-level 2 \
    --item '/home/u17/frios/xdisk/coca/Shared Files/' '/xdisk/coca/Shared Files/folder1' true
```

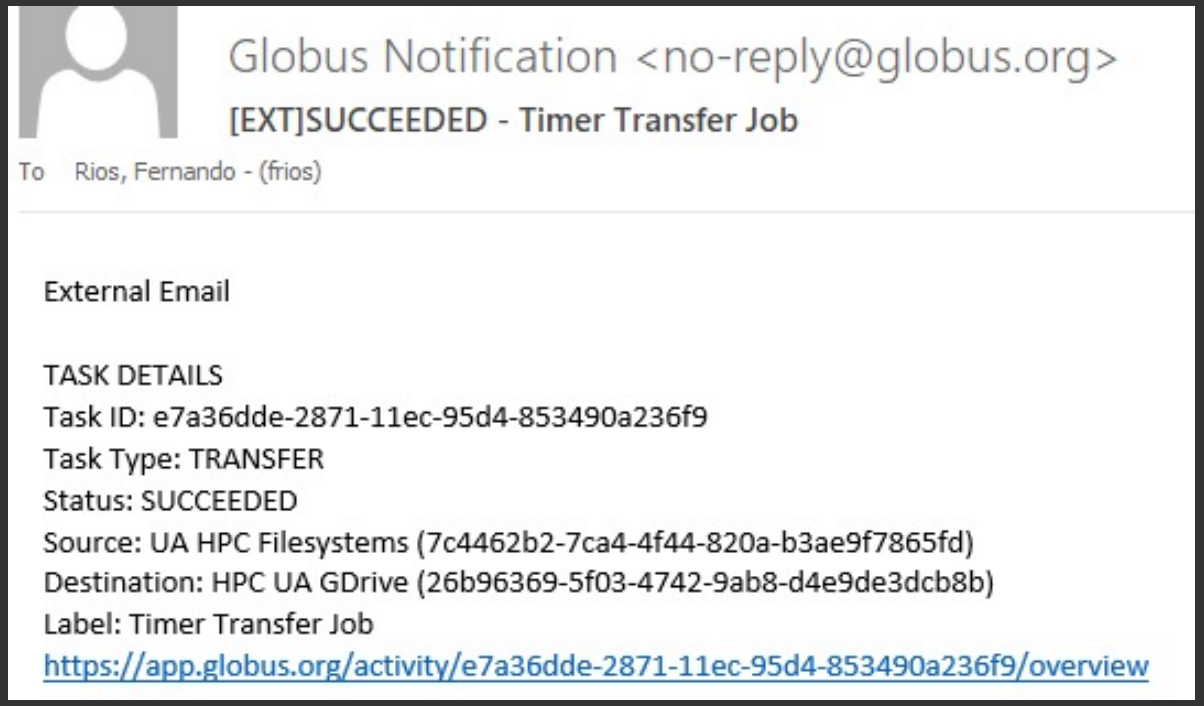
Don't have to be logged in to the HPC. Get email notification + status in Globus web

Source and dest endpoint UUIDs from Globus web globus.org

```
(puma) frios@junonia ~$ globus-timer job status  1ced207e-29d9-4d98-a4aa-e6368f3f4369
Name:             my-job
Job ID:           1ced207e-29d9-4d98-a4aa-e6368f3f4369
Status:           loaded
Start:            2021-10-04T19:57:00+00:00
Interval:         0:02:00
Next Run At:      2021-10-04T20:03:00+00:00
Last Run Result: RUN COMPLETE
```

Scheduled/recurring transfers using Globus

- Job runs automatically
- Check status
  - globus-timer job status <job_id>



Project is Done – publish an article

Work done, data analyzed, paper ready to submit

Now what?

- Archive the “final” data
- Cite it in the paper => get credit
- Publish it in a data repository

UA Research Data Repository (ReDATA)

- Long-term archival repository for “final” data
  - Get a DOI
  - Comply with funder, journal policies, UA retention policy
  - Get help improving the data for reuse
  - You don’t have to worry about keeping data around, even if you leave UA

Tier 1 Tier 2 Tier 3

Google Drive

Open Science Framework

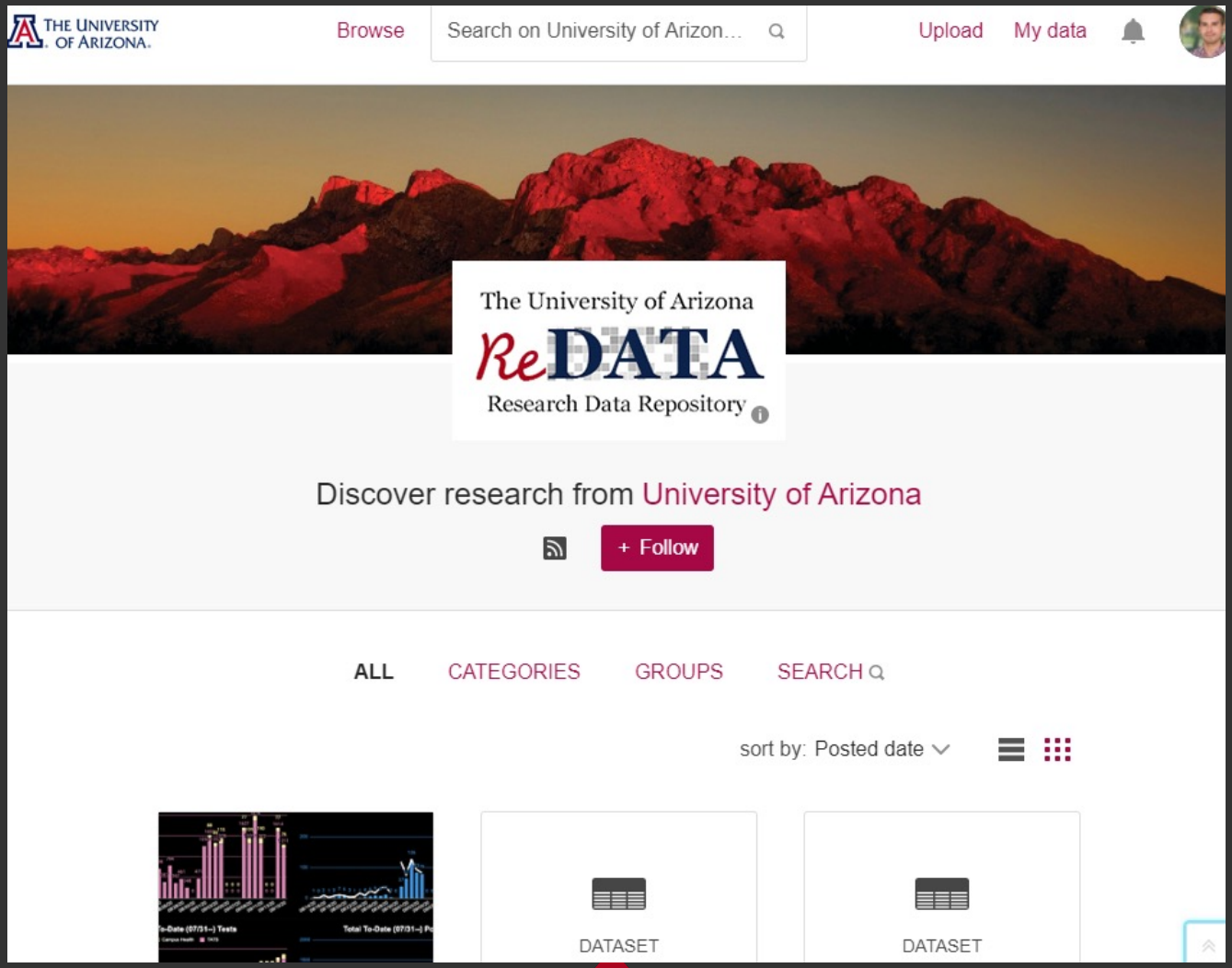
ReDATA

arizona.figshare.com

Private Private/Public Public

ReDATA

- Go to arizona.figshare.com, log in



Takeaways

- Setting up data management workflows increase efficiency and support doing good research
- By linking together both UA-provided and 3rd party tools and resources, you can build a solid workflow at no cost
- Refer to documentation and guides
  - public.confluence.arizona.edu/display/UAHPC
  - data.library.arizona.edu/osf
  - data.library.arizona.edu/redata

UNIVERSITY LIBRARIES

Data Cooperative

The data cooperative is a group of library-based data services providers
https://data.library.arizona.edu

Data Management

Consulting, data management plans, support for DMPTool, OSF, data archiving via ReDATA

Data Science & Visualization

Data analysis & data visualization suppor through consulting and instruction

Geospatial Support

Data management consulting, data management plans, data archiving via ReDATA

data.library.arizona.edu/data-management/events-schedule-current
Pipelines: Data news, events for UA
https://redata.tiny.us/dm-news