

# CyVerse Data Store



## Managing Your 'Big' Data

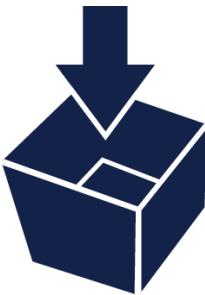


Download Slides and Follow Along

[mcbios.readthedocs.org](https://mcbios.readthedocs.org)



# Welcome to the Data Store



Manage and share your data across all CyVerse platforms





# Working with Big Data

Challenges: the scope and scale of life sciences data continue to grow

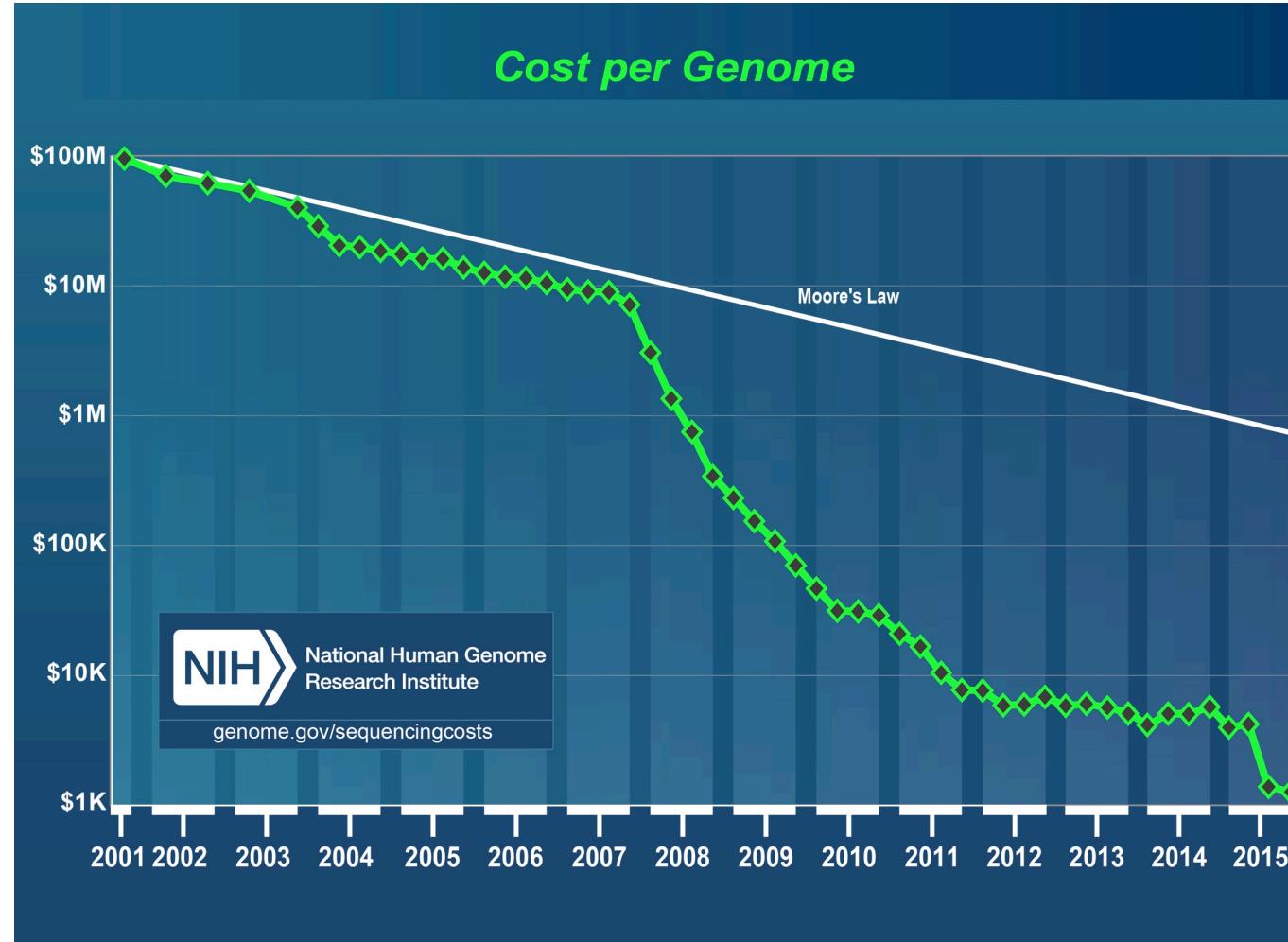


- Big data a term applied to data sets whose size is beyond the ability of commonly used software tools to capture, manage, and process the data within a **tolerable elapsed time**
- Big data sizes are a constantly moving target currently ranging from a **few dozen terabytes (TB)** to many petabytes of data in a single data set.



# Working with Big Data

Challenges: data generation is cheaper and faster



<http://www.genome.gov/sequencingcosts/>



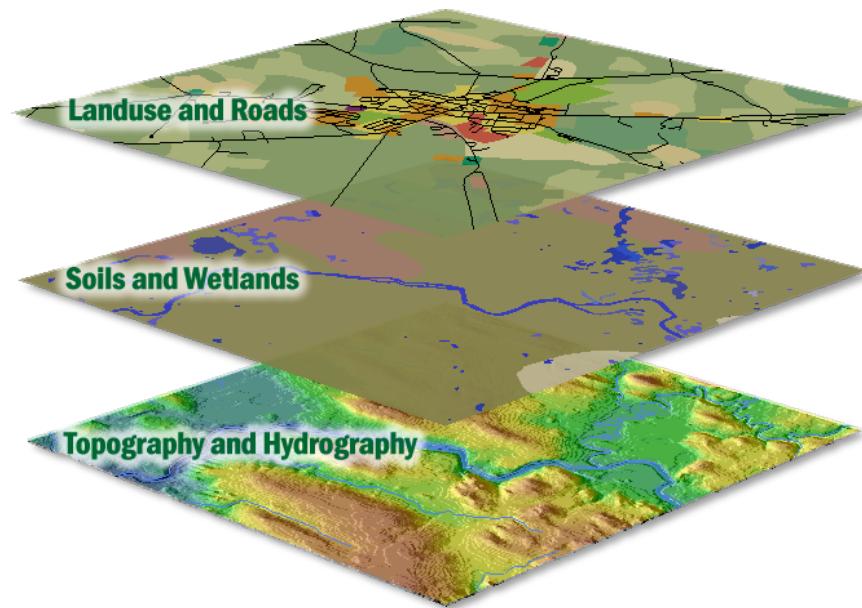


# Working with Big Data

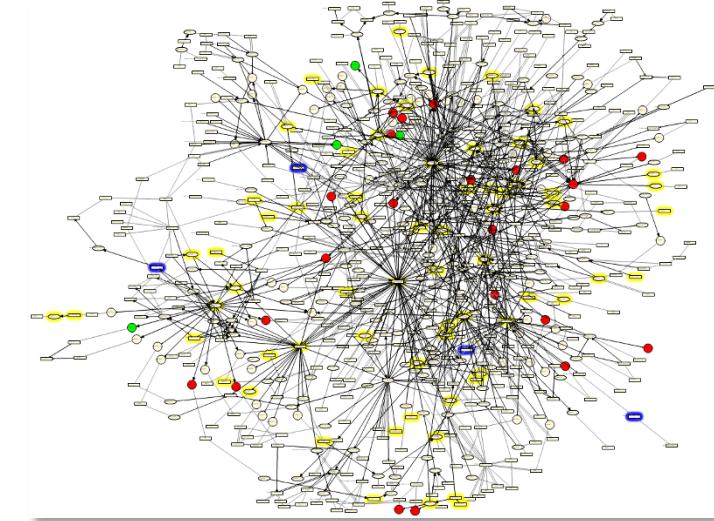
Challenge: biology encompasses more than sequence data



Advanced Imaging



Geospatial



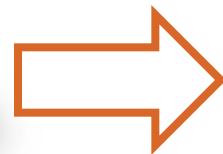
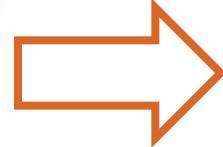
Network

Biologists work with and require access to diverse data types



# Working with Big Data

Challenges: changes in data require changes in tools



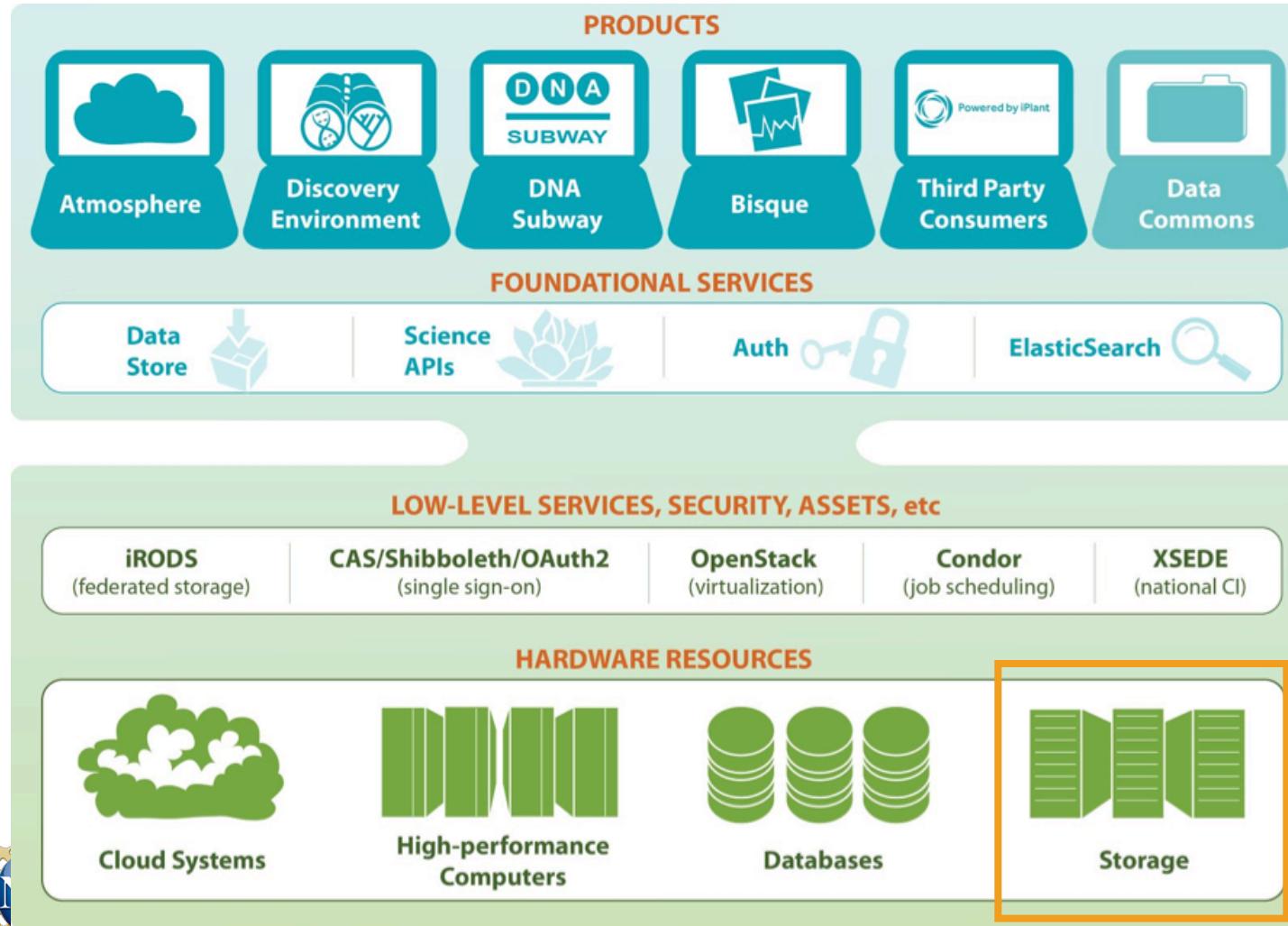
- Difficult / slow transfers
- Expense for storage / backup
- Difficult to share and publish
- Analysis
- Metadata (*What Is metadata?*)

Changes in scale introduce quantitative and qualitative complications



# Data Store Overview

The Data Store services all CyVerse platforms



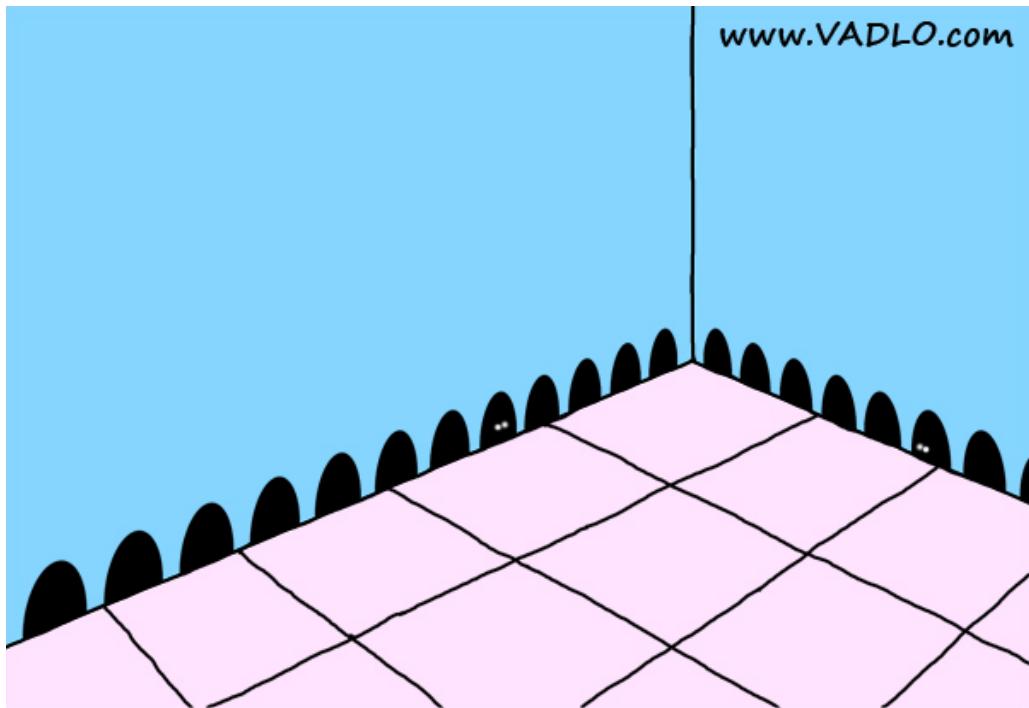
- Access your data from multiple CyVerse services
- Automatic backup (redundant between University of Arizona and University of Texas)
- Default 100 GB allocation, > 1 TB allocations available with justification





# Data Store Overview

Avoid reinventing the wheel



Mouse house that did not receive infrastructure funding.

- iRODS (integrated Rule-Oriented Data System) is an established, scalable, open-source data management system
- iRODS supports many data intensive projects
- iRODS abstracts data services from data storage to facilitate executing services across heterogeneous, distributed storage systems



# Data Store Overview

## Benefits

### Get Science Done



- Store **any type of files** related to your research
- An evolving “Data Commons” lets you access important datasets

### Reproducibility



- Metadata captures information needed for **reproducibility**
- Automatic backup and accessibility support your **data management plan**

### Productivity



- iRODS makes **high-speed** transfers possible (100 GB in ~30 min)
- Share data **instantly** with collaborators within CyVerse

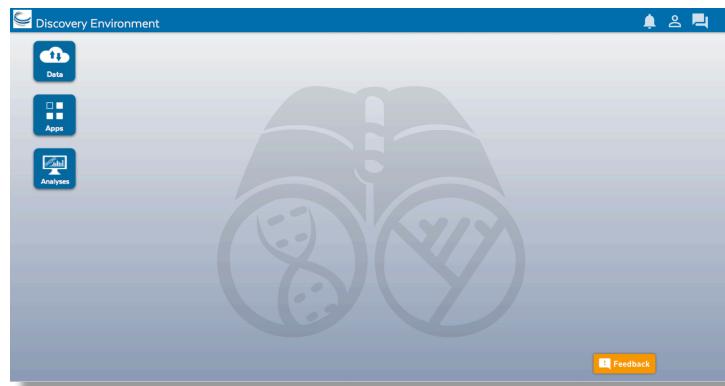


# Data Store Overview

Multiple ways to access



Cyberduck



Discovery Environment

## Point-and-click

## Command line

```
C- /iplant/home/williams/analyses/Soapdenovo_2.04b_analysis1-2013-09-18-13-35-36.219
C- /iplant/home/williams/analyses/Soapdenovo_2.04b_analysis1_47-2013-09-18-22-50-52.016
C- /iplant/home/williams/analyses/TASSEL_4.3.0 MLM analysis1-2013-09-11-20-17-30.232
C- /iplant/home/williams/analyses/TASSEL_4.3.0 MLM analysis1-2013-09-12-14-52-35.844
C- /iplant/home/williams/analyses/Test_of_New_App_analysis1-2013-10-25-14-40-49.857
```

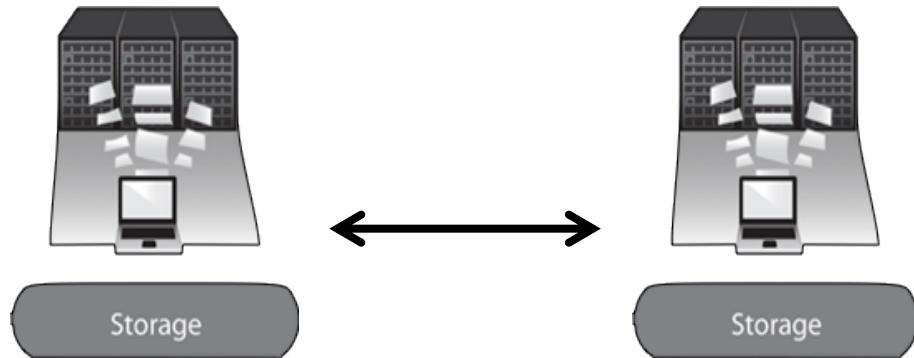
iCommands



# Data Store Overview

Some important things we will not “see” in the demo

## Data Backups



Arizona

Texas

Key component of your data management

Worry-free

## Data Transfer

Source	Destination	Copy Method	Time (seconds)
CD	My Computer	cp	320
Berkeley Server	My Computer	scp	150
External Drive	My Computer	cp	36
USB 2.0 Flash	My Computer	cp	30
Data Store	My Computer	iget	18
My Computer	My Computer	cp	15

Closer to optimum conditions:  
transfers between University of  
Arizona and UC Berkeley

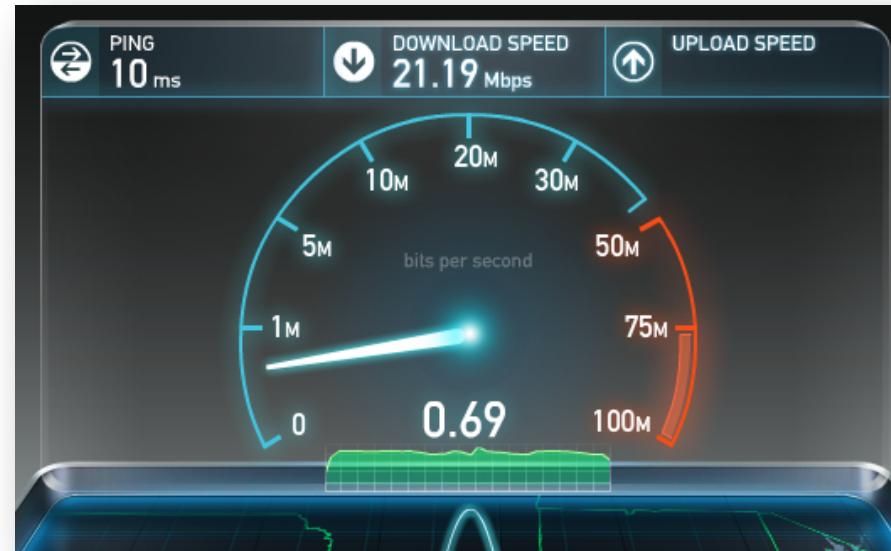
100 GB: 26m15s, 1 GB 17.5s





# Data Store Overview

Some important things we will not “see” in the demo



<http://www.speedtest.net/>

Local connections and institutional policies limit data transfer





# Data Store Overview



## Hands-on demo



# Data Store Overview

## User perspectives and potential applications

Bench Scientist



- Uploads all of his .fastq files along with 50GB of root growth videos
- Shares all his analyses results with his thesis advisor

Bioinformatician



- Created a metadata template for assembled genomes her students and collaborators will place in a shared folder
- Uses public links in the supplemental materials of her publications

Core Facilities

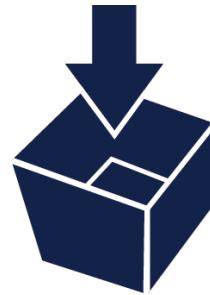


- Developed a script to automate transfer of data to core users
- Uses a shared folder to make large datasets accessible





# Data Store Overview



Time for Summaries and Tips



# Tips for any transfer method

## Spaces / Special Characters

- Many software packages are sensitive to spaces in file names and/or the special characters below
- Rename uploaded files before using them in an analysis

**~ ! @ # \$ % ^ & \* ( ) + =**

**{ } [ ] | \ : ; " ' < > , ? /**



# Tips

When sharing, use this chart to decide appropriate permissions

Permission	Read	Download	Metadata	Rename	Move	Delete
Read	x	x				
Write	x	x	x			
Own	x	x	x	x	x	x



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454 Agave\_API Assembly  
Atmosphere Bisque CoGe  
DNASubway DataStore  
DiscoveryEnvironment  
EST General HPC Metadata blast  
bwa concatenate cuffdiff error  
genome icommands instance

**Broken link in AGAVE Beginners Guide**

Agave\_API

no votes 1 answer 6 views  
3 hours ago jdebarry

I'm having problems with SoapdeNovo runs failing. I think there might be an error in how the program is getting the manual entries for insert size. I'll paste my error file below.

soapdenovo-trans DiscoveryEnvironment

no votes 2 answers 12 views  
3 hours ago matthewterry

**Where is shared data?**

DiscoveryEnvironment

no votes 2 answers 23 views  
Jan 25 '16 kyclark

**Output from Scythe-Adapter-Trimming\_analysis is not showing up...**

DiscoveryEnvironment scythe

no votes 1 answer 11 views  
Jan 22 '16 jdebarry

