

CyVerse Data Store



Managing Your 'Big' Data



Cold
Spring
Harbor
Laboratory

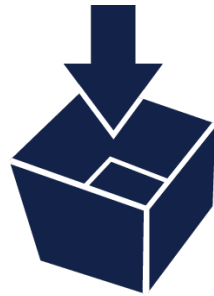


Download Slides and Follow Along

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



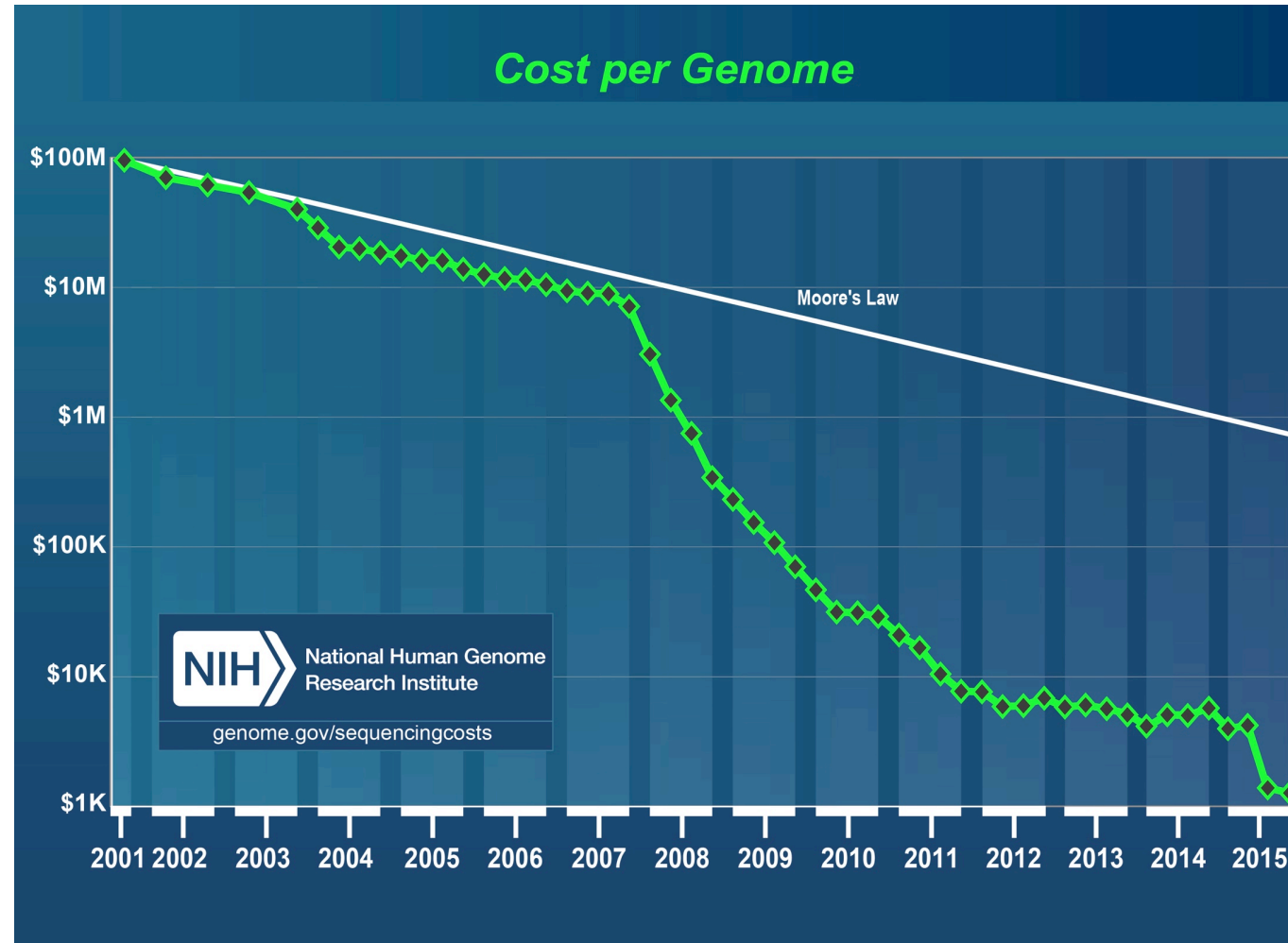
- Big data a term applied to data sets whose size is beyond the ability of commonly used software tools to capture, manage, and process the data within a **tolerable elapsed time**
- Big data sizes are a constantly moving target currently ranging from a **few dozen terabytes** (TB) to many petabytes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





Working with Big Data

Challenges: data generation is cheaper and faster



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



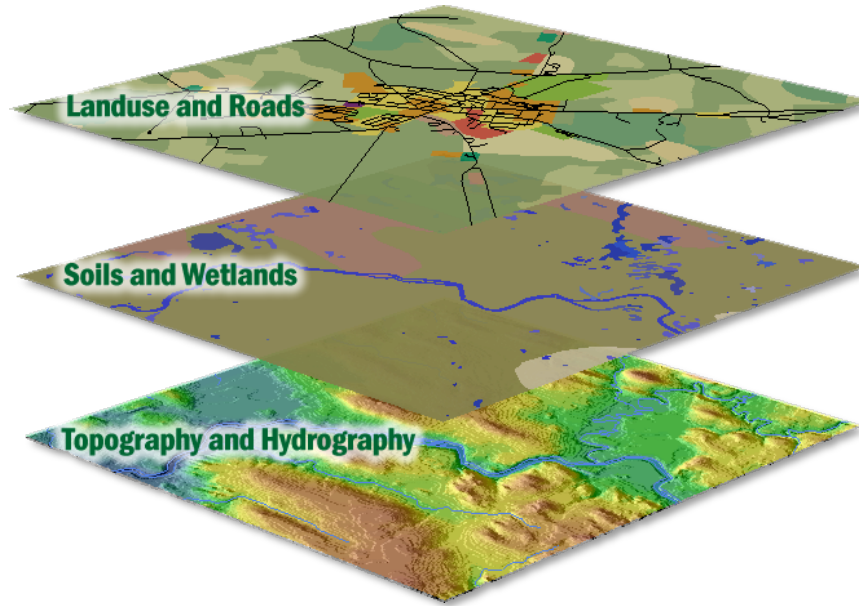


Working with Big Data

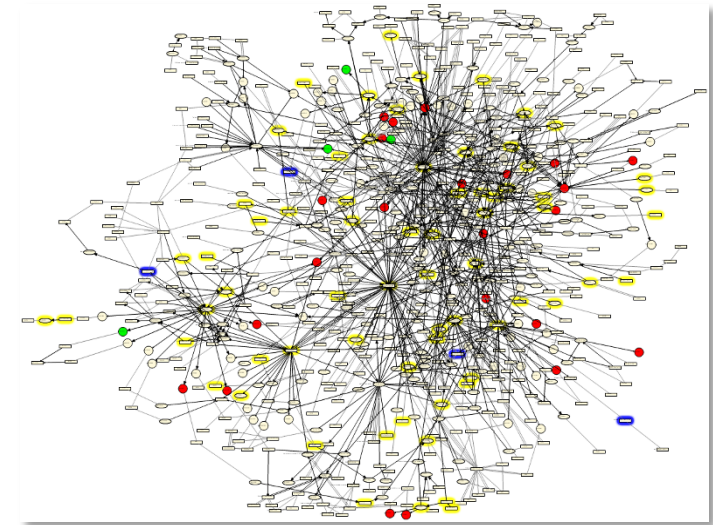
Challenge: biology encompasses more than sequence data



Advanced Imaging



Geospatial



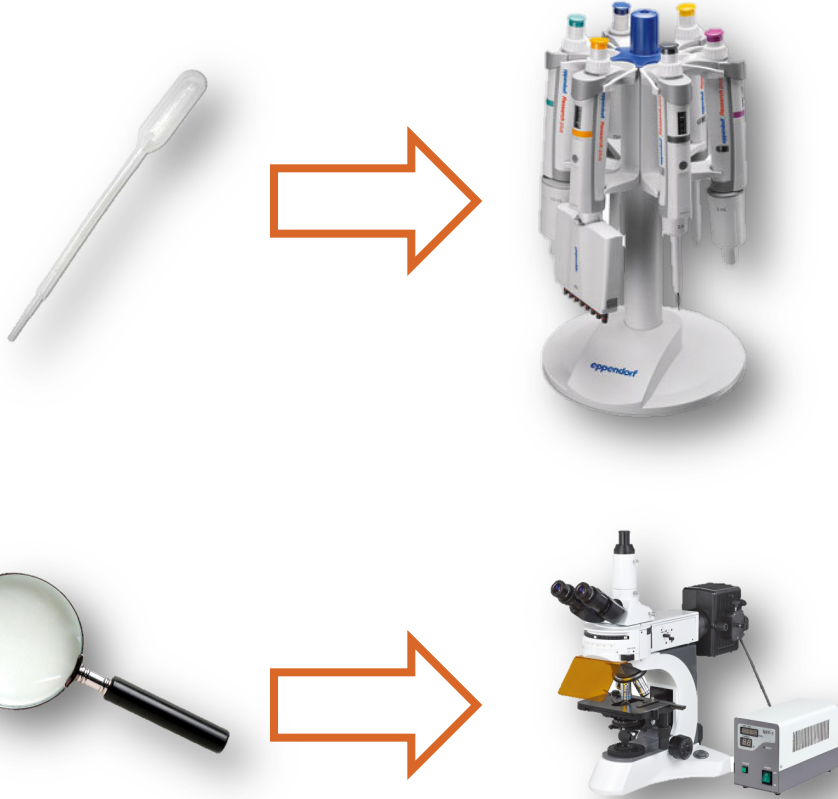
Network

Biologists work with and require **access** to diverse data types



Working with Big Data

Challenges: changes in data require changes in tools



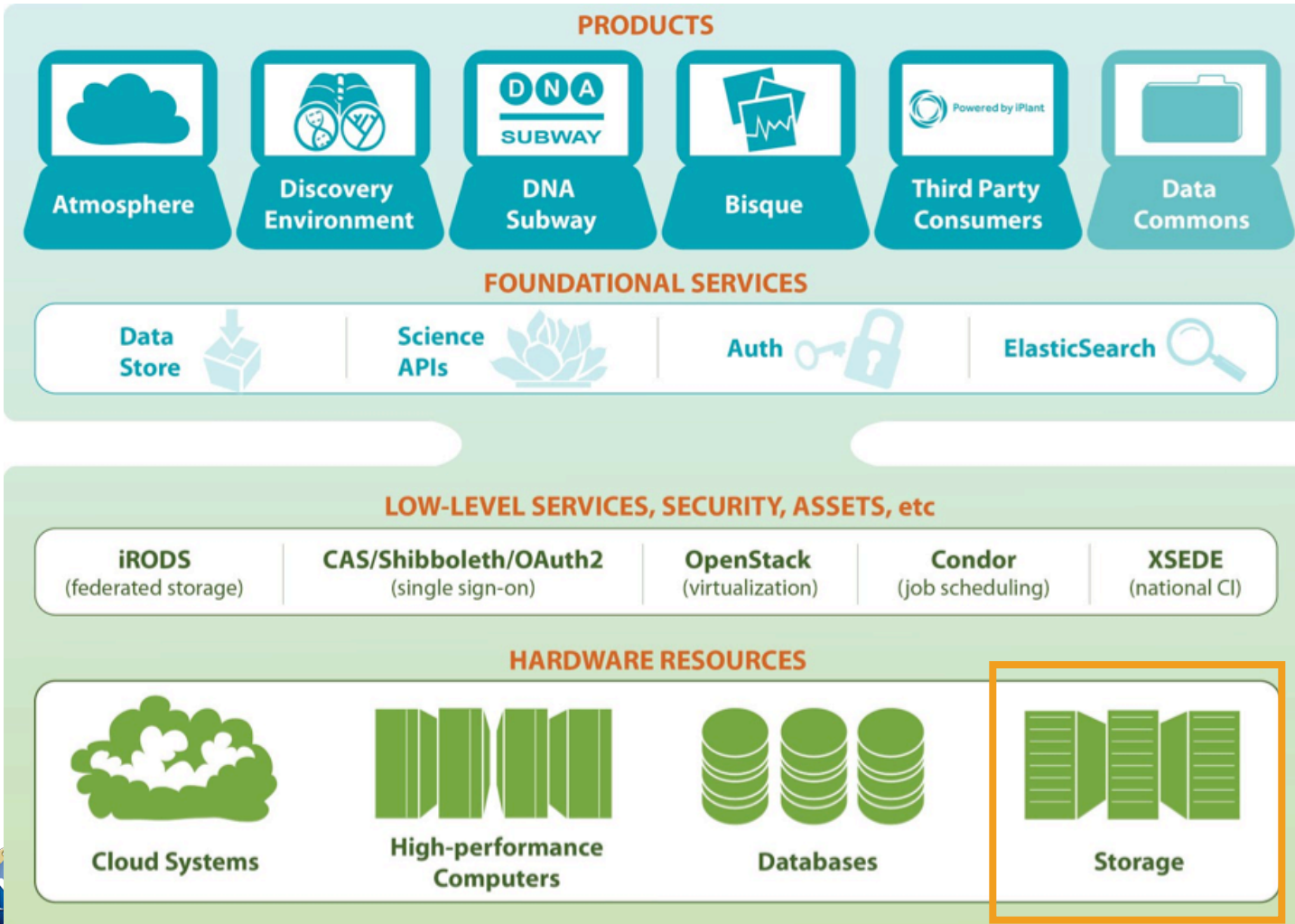
- Difficult / slow transfers
- Expense for storage / backup
- Difficult to share and publish
- Analysis
- Metadata (**What Is metadata?**)

Changes in scale introduce quantitative and qualitative complications



Data Store Overview

The Data Store services all CyVerse platforms



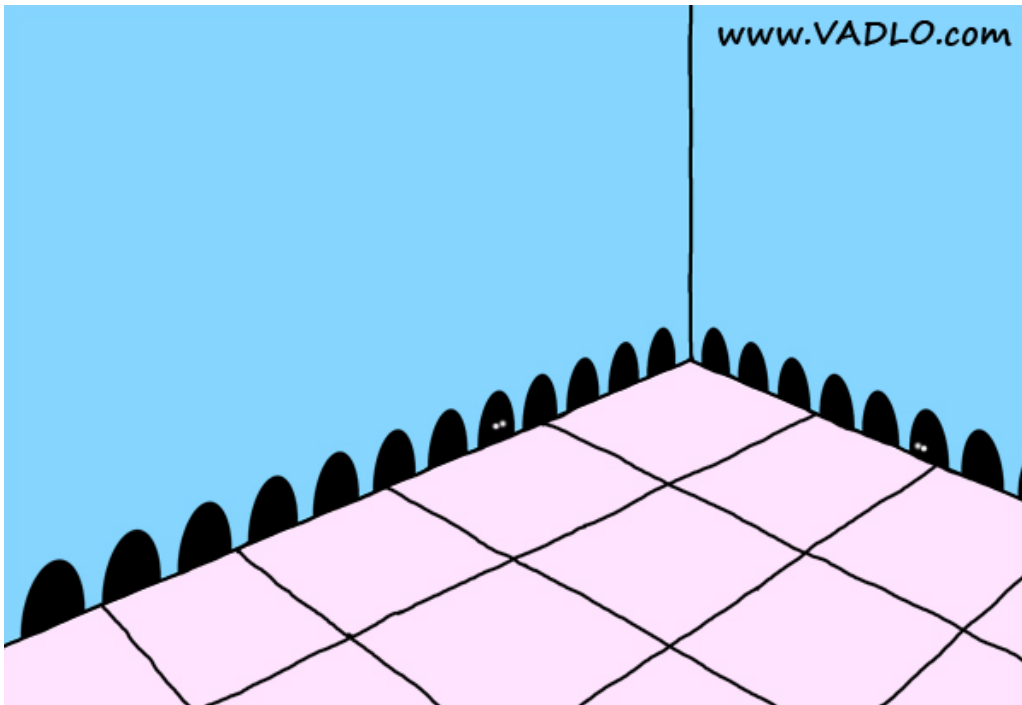
- Access your data from multiple CyVerse services
- Automatic backup (redundant between University of Arizona and University of Texas)
- Default 100 GB allocation, > 1 TB allocations available with justification





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



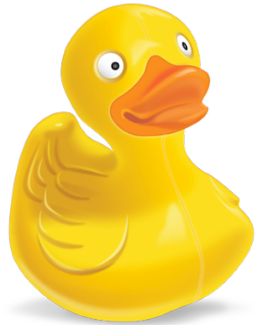
- iRODS makes **high-speed** transfers possible (100 GB in ~30 min)
- Share data **instantly** with collaborators within CyVerse



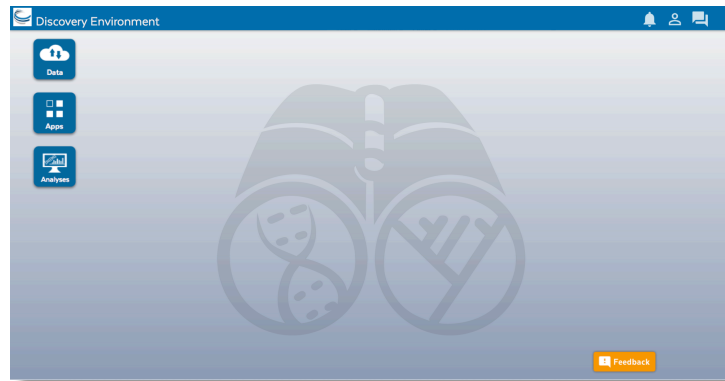
Data Store Overview

Multiple ways to access

Point-and-click



Cyberduck



Discovery Environment

Command line

```
jasonwilliams — bash — 44x11
C- /iplant/home/williams/analyses/Soapdeno
vo_2.04b_analysis1-2013-09-18-13-35-36.219
C- /iplant/home/williams/analyses/Soapdeno
vo_2.04b_analysis1_47-2013-09-18-22-50-52.01
6
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
```

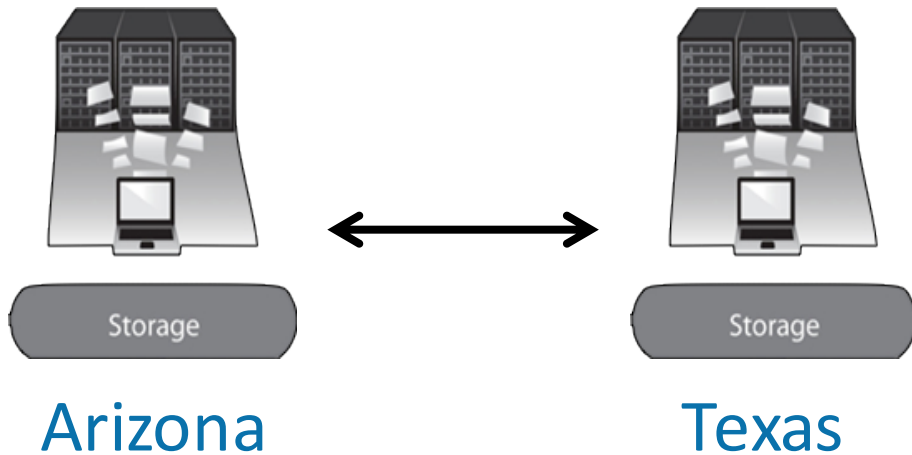
iCommands



Data Store Overview

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

Data Backups



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

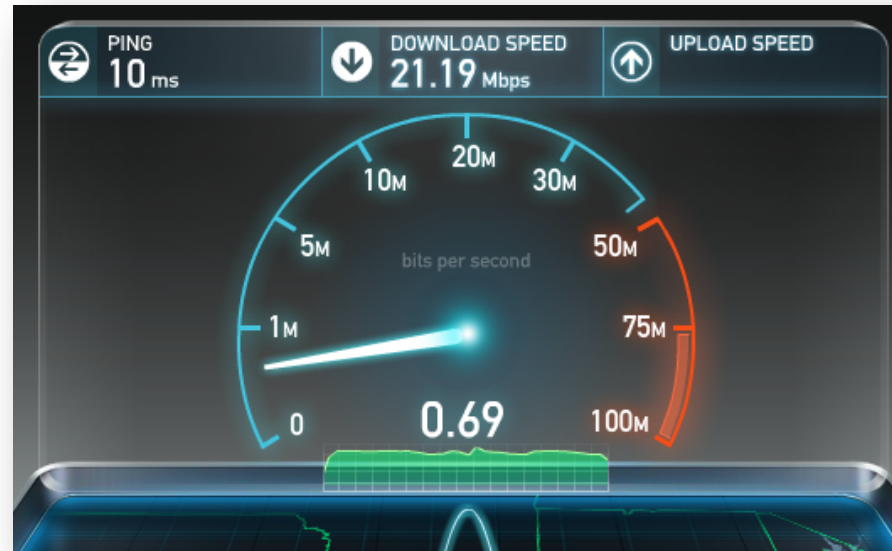
100 GB: 26m15s, 1 GB 17.5s





Data Store Overview

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

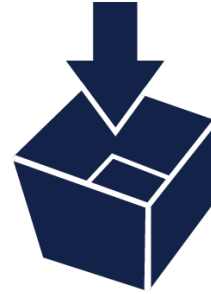


<http://www.speedtest.net/>

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

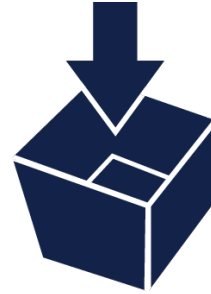
Core Facilities



- 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 files 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



Keep asking: ask.iplantcollaborative.org

Detailed instructions with videos, manuals, documentation in Learning Center

The screenshot shows the homepage of the iPlant Collaborative Ask iPlant forum. The header includes the iPlant Collaborative logo, navigation links for Tags, People, and Badges, and options to login, register, or get help. Below the header, there are filters for 'ALL' and 'UNANSWERED', a search bar, and an 'ASK YOUR QUESTION' button. The main content area displays a list of questions with their titles, tags, and statistics (votes, answers, views). The questions shown are:

- Broken link in AGAVE Beginners Guide** (Tag: Agave_API, 1 answer, 6 views, 3 hours ago by jdebarry)
- I'm having proplems 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.** (Tags: soapdenovo-trans, DiscoveryEnvironment, 2 answers, 12 views, 3 hours ago by matthewterry)
- Where is shared data?** (Tag: DiscoveryEnvironment, 2 answers, 23 views, Jan 25 '16 by kycklark)
- Output from Scythe-Adapter-Trimming_analysis is not showing up...** (Tags: DiscoveryEnvironment, scythe, 1 answer, 11 views, Jan 22 '16 by jdebarry)

On the right side, there are sections for 'Contributors' (a grid of user avatars) and 'Tags' (a list of popular tags such as Agave_API, Assembly, Atmosphere, Bisque, CoGe, DNASubway, DataStore, DiscoveryEnvironment, EST, General, HPC, Metadata, blast, bwa, concatenate, cuffdiff, error, genome, icommands, instance).

