

CyVerse Overview

MCBIOS 2016 – University of Memphis, TN

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Download Slides and Follow Along

mcbios.readthedocs.org







iPlant 2008 Empowering a New Plant Biology *iPlant 2013* Cyberinfrastructure for Life Science **CyVerse 2016** Transforming Science Through Data-Driven Discovery





CyVerse Evolution



DBI-0735191 and DBI-1265383

We are funded by the National Science Foundation

- We are your colleagues and collaborators!
- \$100 Million in investment
- Freely available to the community
- Spur national/international collaboration
- Cite CyVerse:

CyVerse.org/acknowledge-cite-cyverse





CyVerse Evolution



Vision:

Transforming science through data-driven discovery

Mission:

Design, develop, deploy, and expand a national cyberinfrastructure for life science research, and train scientists in its use

CyVerse 2016 Transforming Science Through Data-Driven Discovery

More than 30K users, PB of data, and hundreds of publications, courses, and discoveries





What is Cyberinfrastructure?

- Data storage
- Software
- High-performance computing
- People

organized into systems that solve problems of size and scope that would not otherwise be solvable.





What is Cyberinfrastructure?

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Platforms, tools, datasets

Storage and compute

Training and support





CyVerse supports all domains of life science



Plant / Microbial

Animal

Biomedical Ecological/Climate



CyVerse is built for Data



CyVerse product stack





How was CyVerse built?



Published online <u>3 September 2008</u> | *Nature* **455**, 16-21 (2008) | doi:10.1038/455016a

News Feature Big data: Welcome to the petacentre

What does it take to store butos by the tops of theusands of the store 455, 30 (4 September 2008) | doi:10.1038/455030a; Published online 3 September 2008

which Big data: Distilling meaning from data

Felice Frankel¹ & Rosalind Reid²

 Buried in va:
 Nature 455, 28-29 (4 September 2008) | doi:10.1038/455028a; Published online 3 September 2008

 need to crafi
 Big data: How do your data grow?

Clifford Lynch¹

 Clifford Lynch is the executive director of the Coalition for Networked Information, 21 Dupont Circle, Washington DC 20036, USA, and an adjunct professor at the School of Information, University of California, Berkeley, California, 94720-4600, USA. Email: <u>cliff@cni.org</u>

Scientists need to ensure that their results will be managed for the long haul. Maintaining data takes big organization, says Clifford Lynch.

ature 455, 1 (4 September 2008) | doi:10.1038/4-01a; Published online 3 September 2008

Community cleverness required

Researches and provide the ansatz the second provide the second provid





CyVerse Community Priorities

Genomic data and analysis:

- Reference guided assembly
- De novo assembly
- RNA-Seq (expression; gene/isoform discovery)
- Variant calling
- Genome/Transcriptome annotation
- ChIP-Seq/Integration of epigenetic information
- Multiple sequencing platforms
- New and evolving technologies





CyVerse Community Priorities







CyVerse Community Priorities



Phylogenetic Tools for inference





CyVerse Collaborators





CyVerse collaborates to enable access to the solutions that work the best for you...



CyVerse Institutions



CyVerse is a collaborative virtual organization



CyVerse Products

- We strive to be the CI Lego blocks
- Danish 'leg godt' '**play well**'
- Also translates as 'I put together' in Latin
- If a solution is not available you can craft your own using CyVerse Cl components

Data Store

The resources you need to share and manage data with your lab, colleagues and community

- ✓ Initial 100 GB allocation TB allocations available
- ✓ Automatic data backup
- ✓ Easy upload /download and sharing

Discovery Environment

Hundreds of bioinformatics Apps in an easy-to-use interface

- \checkmark A <u>platform</u> that can run almost any bioinformatics application
- Seamlessly integrated with data and high performance computing
- ✓ User extensible add your own applications

- ✓ Simple: One-click access to more than 200 virtual machine images
- ✓ Flexible: Fully customize your software setup
- ✓ Powerful: Integrated with iPlant computing and data resources

Science APIs

Fully customize *iPlant* resources

- ✓ Science-as-a-service platform
- ✓ Define your own compute, and storage resources (local and *iPlant*)
- ✓ Build your own app store of scientific codes and workflows

DNA DNA Subway

SUBWAY Educational workflows for Genomes, DNA Barcoding, RNA-Seq

- Commonly used bioinformatics tools in streamlined workflows
- ✓ Teach important concepts in biology and bioinformatics
- ✓ Inquiry-based experiments for novel discovery and publication of data

- ✓ Secure image storage, analysis, and data management
- ✓ Integrate existing applications or create new ones
- ✓ Custom visualization and image handling routines and APIs

CyVerse Executive Team

Parker Antin Nirav Merchant Eric Lyons

TEXAS ADVANCED COMPUTING CENTER

Matt Vaughn

Doreen Ware Dave Micklos

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