



THE UNIVERSITY OF
CHICAGO

Center for
Research
Informatics

WORKSHOP SERIES

Using CRI's Computational Infrastructure for Bioinformatics Analysis

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CRI SYSTEMS RESOURCES OVERVIEW

HPC

Tarbell Cluster: High Performance Computation Cluster for Data Analysis

Large Memory Server: (SAS/STATA/X-Windows)

Web Portals

Galaxy: Integrates with the HPC & provides a suite of tools for data analysis

REDCap: for building online surveys and databases. Research data can be exported to statistical packages (SPSS, SAS, Stata, R)

Data Storage

Home Data: 10GB Limit

Lab Shares: From 100GB

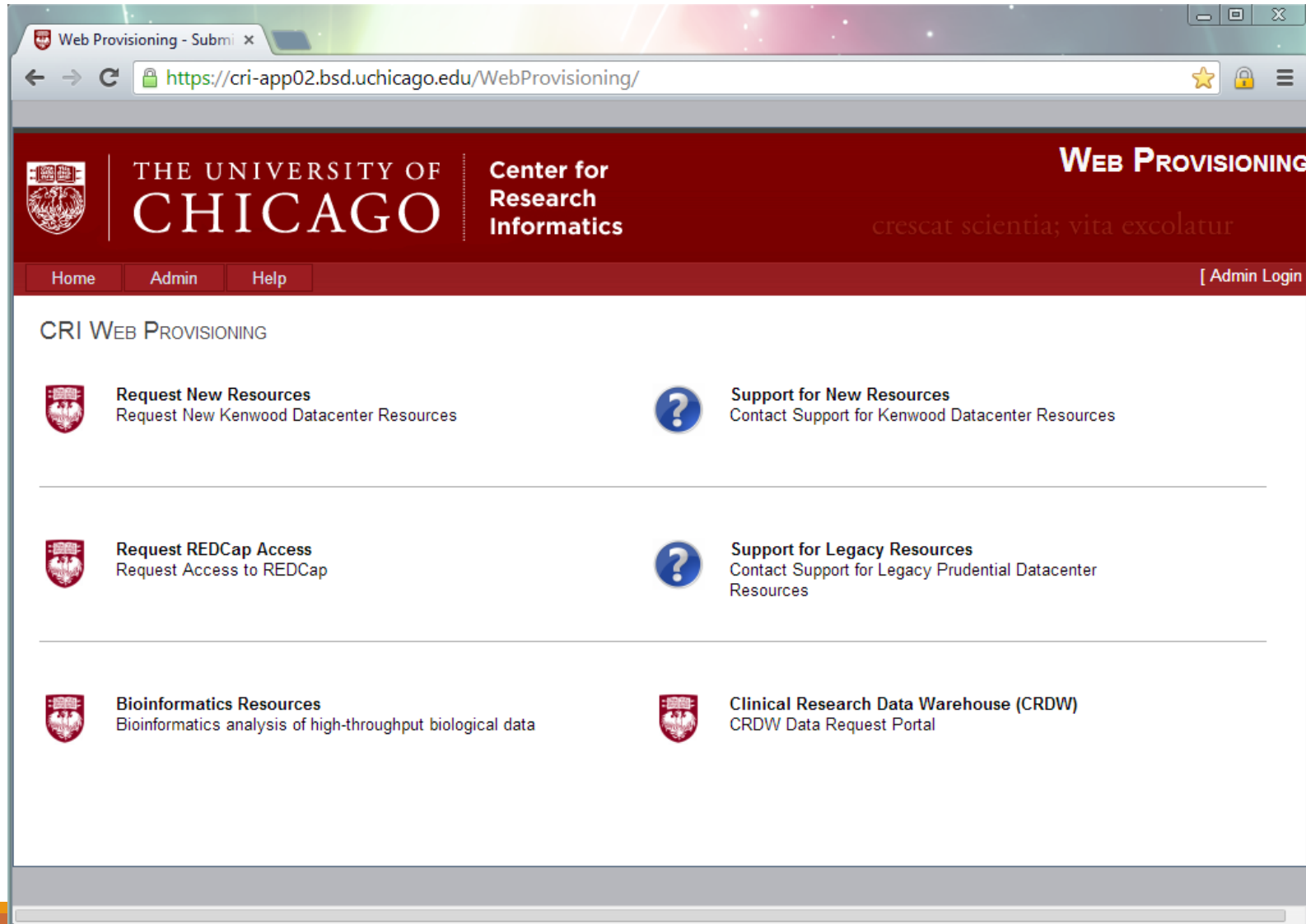
Fast Scratch Space: Shared 60TB

Virtualization

Virtual Machines: Hosted Windows & Linux Servers for building web applications and databases

HOW TO REQUEST CRI RESOURCES/SUPPORT


Visit: cri.uchicago.edu > Request Resources





ONLINE FORMS


[Home](#) [Admin](#) [Help](#)


CRI WEB PROVISIONING - NEW SERVICE REQUEST


**Request IBI User Account for REDCap**
Request IBI User Account (REDCap)


**Request IBI Collaborator Account for REDCap**
Request Collaborator Account (REDCap, IBICluster, Storage)


**Request Lab Share Creation**
Request creation of a Lab Share to store research data in a highly secure clustered storage system with tape library backup.


**Request User Access to Lab Share**
Request user access to an existing Lab Share


**Request Access to HPC Resources**
Request Access to Galaxy, the High Performance Computing Cluster or Large Memory Server


**Request Software Installation on HPC or LMEM**
Request installation of software on the 1024 Core High Performance Cluster or the Large Memory Super Computer

**Request Virtual Machine Creation**
Request creation of a virtual machine instance

**Request Collaborator Account Creation**
Request creation of an account for a collaborator

**Request Firewall Access**
Request firewall access

**Request Collaborator Account Deactivation**
Request deactivation of an account for a collaborator

**Revoke Resource Access**
Request removal of user access to the HPC Biocluster, Large Memory Server, Galaxy, or Lab Share

REQUESTING SUPPORT

- 1) For BSD Account issues, please contact CBIS help@bsd.uchicago.edu or call 773-702-3456
- 2) Use appropriate form on the Request Site to create a support ticket;
 Visit: cri.uchicago.edu, [Request Resources Link](#)
 -Trouble with installed software/ modules
 -Job submission issues
 -Questions
- 1) Call the CRI Help Desk to follow up on tickets - 773-834-8475

DATA MANAGEMENT

-Home Directories: /home/<username>

Limited to 10GB. Single user access to data

Check your current working directory with command #pwd

-Lab Shares: /group/<share-name>

Group access to backed-up data

Changing directory: #cd /group/share-name

-Scratch Space: /scratch/<share-name>

60TB Space for staging input used in analysis jobs and for temp files from job execution.

Data in scratch is purged periodically.

-Remote Access to Data

Login to the Campus VPN (cvpn.uchicago.edu)

DATA TRANSFER TO/FROM CRI STORAGE

Accessing Share From a Windows PC:

Option 1:

Click the Start Button > Type "run" in the Search Box and press "Enter" > In the run window, type \\bulkstorage.uchicago.edu\\<Lab-share-name>, press "Enter"

Login with your BSD Account, format "BSDAD\\<bsd-username>"

Option 2:

From "My Computer" > "Map Network Drive"

Folder name: \\bulkstorage.uchicago.edu\\<Lab-share-name>

*If you are not logged in to the BSDAD Domain, Check "Connect using different credentials"

Login with your BSDAD Account, format "BSDAD\\<bsd-username>"

Accessing Share From a Mac:

From "Finder" > "Go"

Server Address: smb://bulkstorage.uchicago.edu/<Lab-share-name>

Login with your BSDAD Account, format "BSDAD\\<bsd-username>"

DATA TRANSFER TO/FROM CRI STORAGE (Cont'd)

Using Rsync, SCP or SFTP to transfer large data:

Host: cri-syncmon.cri.uchicago.edu

Login: BSD Account/Password

Path: /group/<share-name>

Port: 22

Examples of Free SFTP Graphical Clients:

WinSCP, CoreFTP (Windows Only) or
FileZilla, Cyberduck (Windows & Mac)

HOW TO USE ANALYSIS RESOURCES

High Performance Computing (HPC) Cluster:

- Execute analysis in batch mode

- Execute analysis in interactive mode

- X Window System

Large Memory Server:

- Execute SAS or Stata analysis

- X Window System

SPECIFICATIONS

Interactive Nodes:

2 Nodes

2 AMD Opteron 6274 processors (32 cores)

128 GB RAM

Scientific Linux 6 Update 4

Compute Nodes:

34 Nodes

4 AMD Opteron 6274 processors (64 cores)

256 GB RAM

Scientific Linux 6 Update 4

SPECIFICATIONS (Cont'd)

Large Compute Nodes:

- 2 Nodes

- 4 Intel Xeon E7-4860 processors (40 cores)

- Hyper-threaded

- 1 TB RAM

- Scientific Linux 6 Update 4

Large Memory Server:

- 8 Intel Xeon E7-8870 processors (80 cores)

- Hyper-threaded

- 1 TB RAM

- Redhat Linux 6 Update 4

PRODUCTION ENVIRONMENT

Complete:

- TORQUE Resource Manager
- MOAB Scheduler
- Galaxy
- Scratch Space (60 TB)

Coming Soon:

- Web services
- Simplified Parametric Submission
- Job portability
- Trickled submission
- More Scratch Space

SOFTWARE ENVIRONMENT

Compilers:

GNU Compilers

Intel Compilers

PGI Compilers

Language Support:

C/C++

FORTRAN

Perl

Python

R

BEST PRACTICES

- DO NOT – Install Software into your home directory/lab share
- DO NOT – Write your job scripts using MS Word
- DO NOT – Copy/Paste from MS Word documents into your jobs
- DO – Contact ME (mjarsulic@bsd.uchicago.edu) if you are planning on running a large analysis
- DO – Understand that Java programs grow in memory
- DO NOT – Run analysis on the interactive nodes