

Introduction to the command-line interface (shell)

Harvard Chan Bioinformatics Core

in collaboration with

HMS Research Computing

<https://tinyurl.com/hbc-shell-online>

Learning Objectives



- ✓ Learn what a “shell” is and become comfortable with the command-line interface
 - Find your way around a filesystem using written commands
 - Work with small and large data files
 - Become more efficient when performing repetitive tasks
- ✓ Understand what a computational cluster is and why we need it

Exit survey

<https://tinyurl.com/hbc-shell-exit>

Get an O2 account!

[Home](#) > [Service Catalog](#)

Search IT services and kno

Q

O2 Cluster Account

Description

What does it do?

Get an account on O2, the HMS High Performance Compute cluster

Available To

Who is eligible?

Any HMS or HMS-affiliated researcher who has an HMS eCommons account.

Cost

What does it cost?

There is no cost to labs whose PI has a primary or secondary faculty appointment in an HMS Quad department.

If the PI of your lab does not have a primary or secondary faculty appointment in an HMS Quad department, cluster usage will be charged for beginning later in 2021. Please see the following page for current details about rates: <https://it.hms.harvard.edu/rc/core/rates>.

Support

Please fill out the [online help request form](#), or email rchelp@hms.harvard.edu.

How to Access

How do I get it?

Two-factor authentication is required to request an account on O2, as well as for O2 logins once your account is created. Harvard University uses a mobile app called Duo that makes the process quick and easy. Even if you already use Duo for HarvardKey, you will still need to setup a Duo profile for HMS.

- [Setup HMS two-factor Authentication \(HMS Duo Mobile\)](#)
- [Reactivate or Reconnect HMS two-factor Authentication \(HMS Duo Mobile\)](#)

Once you have Duo set up, Click the "Get Service" link to login and complete the request form.

Get this service

Learn more

Don't see what you're looking for?

Data Management

- HMS Data management -

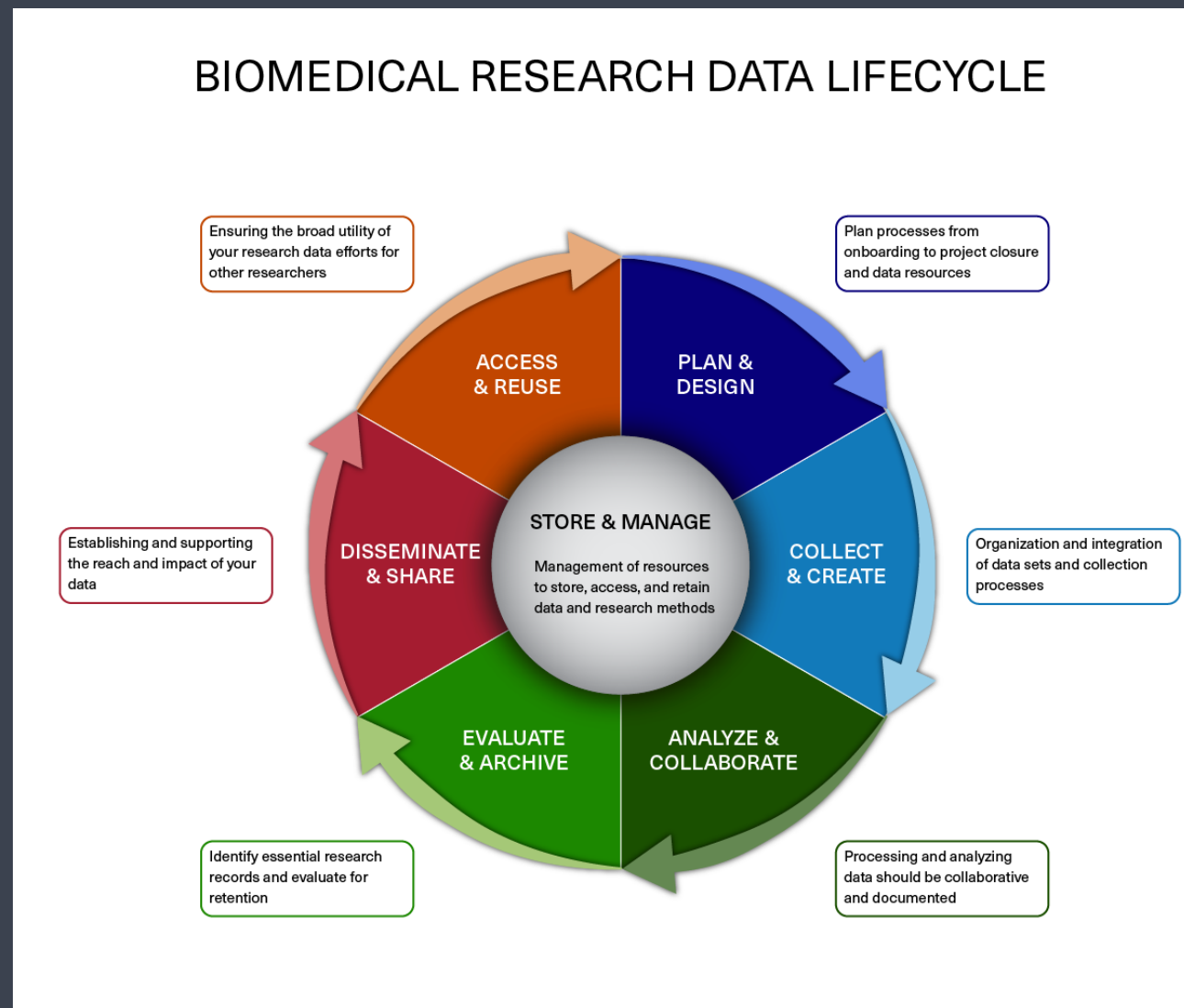
Webpage: <https://datamanagement.hms.harvard.edu/>

[Click here to sign up for data management related emails](#)

- Harvard-wide Research Data Management -

<https://researchdatamanagement.harvard.edu/>

Data Management Short Workshops



<https://datamanagement.hms.harvard.edu/about/news-events/rdmwg-calendar>







Data Management Short Workshops

Spring 2024 Data Lifecycle Training






Plan & Design

- April 17**  Writing a Data Management and Sharing Plan for Grant Applications
- April 24**  Principles of Data Stewardship
- May 1**  Don't Leave Yet! Research Data Offboarding Done Right
- May 8**  Don't Leave Yet! Research Data Offboarding Done Right





Collect & Analyze



- February 21**  Basic Shell
- March 6**  Using Metadata to Find, Interpret & Share Your Data
- March 13**  Intro to MATLAB
- March 13**  Introduction to GIS
- April 17**  Intermediate Shell
- May 15**  Active Project Version Control with Git

Store & Evaluate

- February 7**  The When, Where, and How of Data Storage
- February 27**  Introduction to the General Records Schedule
- March 20**  Managing Data Transfer Between Collaborators
- March 26**  Managing Your Paper Records
- May 7**  Managing Your Electronic Records

Share & Publish

- January 16**  Data Sharing and Reuse on the Vivli Platform
- February 13**  Introduction to Harvard Dataverse
- February 15**  Research Management Using the Open Science Framework
- April 10**  Achieving FAIR Data: Selecting a Repository for Your Data

 In-person
 Virtual



Learn More & Register: bit.ly/rdmwg-calendar



<https://datamanagement.hms.harvard.edu/about/news-events/rdmwg-calendar>

Additional workshops on shell

Accelerate with Automation -
Making your code work for you
March 20th

Needle in a Haystack - Finding and
summarizing data from colossal files
April 17th

Shell Tips and Tricks on the
O2 cluster
May 15th

We hope
to see
you
soon!

Bioinformatics Community Networking Breakfast!



- *Free and open to the Harvard community*
- *Food is first-come-first-served*
- *No computers!*
- *Coming soon end of March!*

More info: <http://bioinformatics.sph.harvard.edu/breakfast/>

Thanks!

- Kathleen Chappell and Andy Bergman from HMS-RC
- [Data Carpentry](#)

These materials have been developed by members of the teaching team at the [Harvard Chan Bioinformatics Core \(HBC\)](#). These are open access materials distributed under the terms of the [Creative Commons Attribution license \(CC BY 4.0\)](#), which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.



Contact us!

HBC webpage: <http://bioinformatics.sph.harvard.edu>

HBC training materials: <http://hbctraining.github.io/main>

HBC workshop listserv: <https://tinyurl.com/hbc-mailing-list>

Training email: hbctraining@hsph.harvard.edu

Consulting email: bioinformatics@hsph.harvard.edu

O2 (HMS-RC): rhelp@hms.harvard.edu

Twitter

[@bioinfocore](https://twitter.com/bioinfocore)