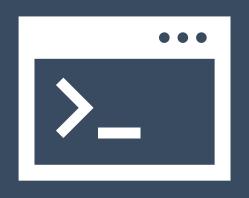


Introduction to the commandline interface (shell)

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



Harvard Chan Bioinformatics Core
in collaboration with
HMS Research Computing



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

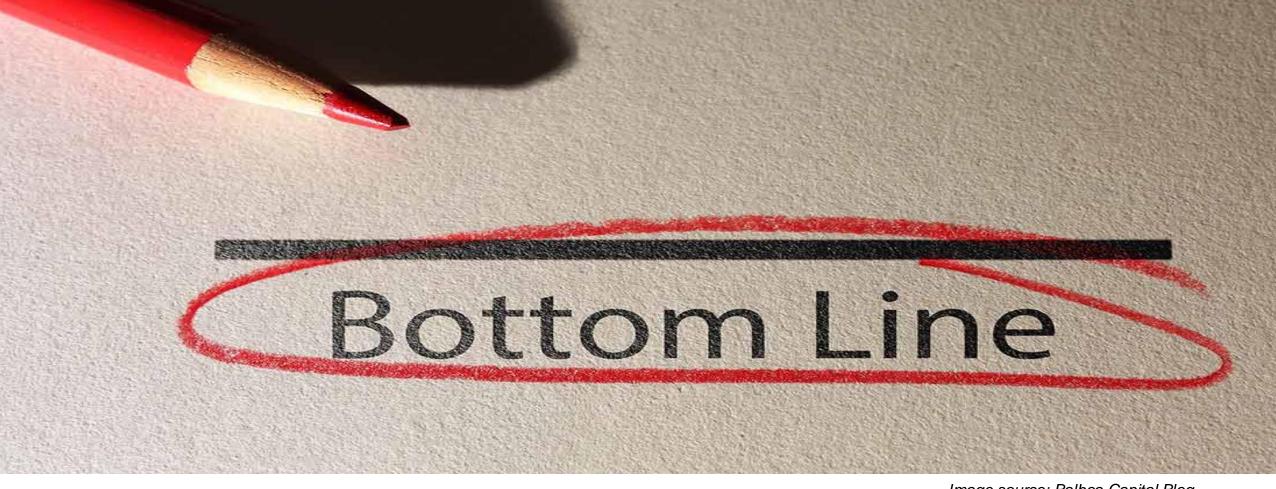


Image source: Balboa Capital Blog

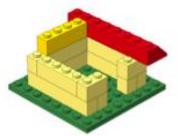
If you plan to process raw high throughput sequencing data yourself, you will need to learn shell.

Exit survey

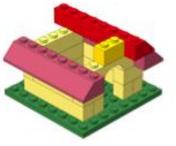
https://tinyurl.com/hbc-shell-exit

Keep building!



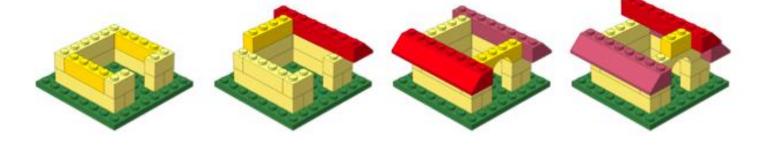






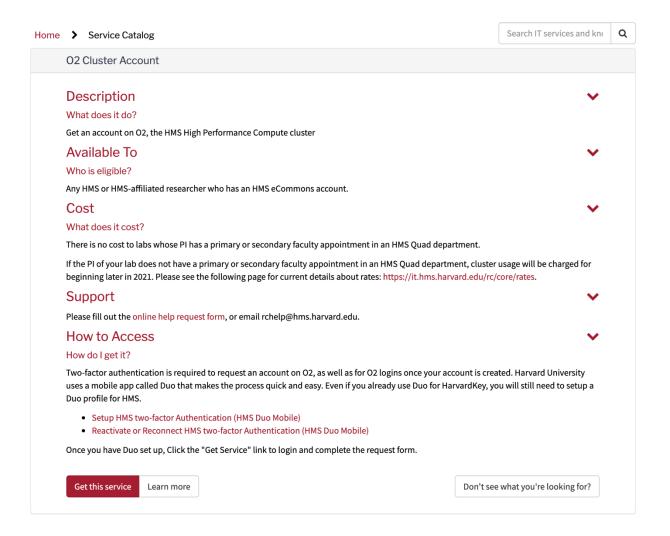
- Upcoming HBC advanced shell-based workshops:
 - Introduction to Variant Calling
 - September 17th, 20th, 24th and 27th (in-person)
 - Using paired tumor-normal samples to analyze variants
 - Introduction to R
 - October 8th, 11th, 15th and 18th
 - Basics of programming and figure creation in R

Keep building!



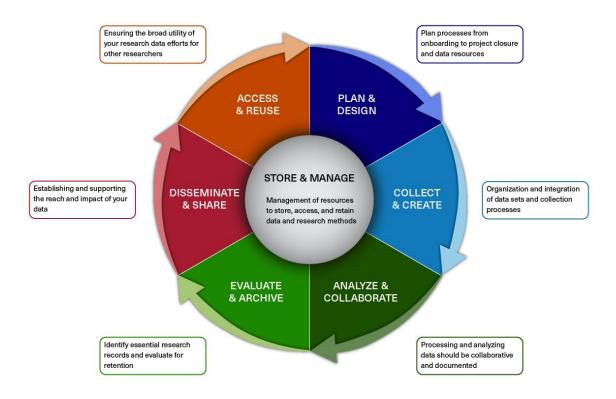
Topic	Pre-requisites	Date/Time	Time	Registration
RShiny	R basics	9/18/24	1 – 4pm	Register!

Get an O2 account!



Research Data Management (RDM)

BIOMEDICAL RESEARCH DATA LIFECYCLE



Fall 2024 Data Lifecycle Training

Plan & Design

September 24 🚇

Managing Research Data Efficiently

September 26

Project and Lab Onboarding

October 31 🙈

Data Horror Stories: Avoid the Nightmare

November 19 🙈

Writing a Data Management and Sharing Plan

Collect & Analyze

September 19 💂

Intro to MATLAB

October 10 💭

Research Computing: Intro to Python

October 16

scRNA pre-processing

November 21

Research Computing: Intro to O2

December 5

RCBio: easy and quick HPC pipeline builder & runner

Store & Evaluate

October 22 💷

Introduction to the General Records Schedule

October 24

Computing Strategies and Resources

November 22 💷

Managing Paper Records: Off-Site Records Storage

December 17

Managing Electronic Records: Shared Drives and Emails

Share & Publish

September 18 💭

Interact with your data using RShiny

November 14 💂

Principles of Finding and Citing Data

November 20 💂

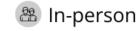
GitHub Part I

December 3 🙉

Research Management: Tools for Open Science

December 12 🖃

Data Sharing in Repositories







Better RDM practice benefits you

- HMS Data Management LMA
 - Webpage: https://datamanagement.hms.harvard.edu
 - Sign up for quarterly email updates
- Harvard-wide Research data Management
 - https://researchdatamanagement.harvard.edu/

Join us for HBC Community Breakfast!

- An opportunity to get to know others in the community
- Free food and beverages
- Great conversations



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 <u>Harvard Chan Bioinformatics</u> <u>Core (HBC)</u>. These are open access materials distributed under the terms of the <u>Creative Commons</u> <u>Attribution license (CC BY 4.0)</u>, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

Contact Us



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