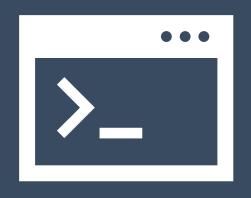


Introduction to the commandline interface (shell)

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

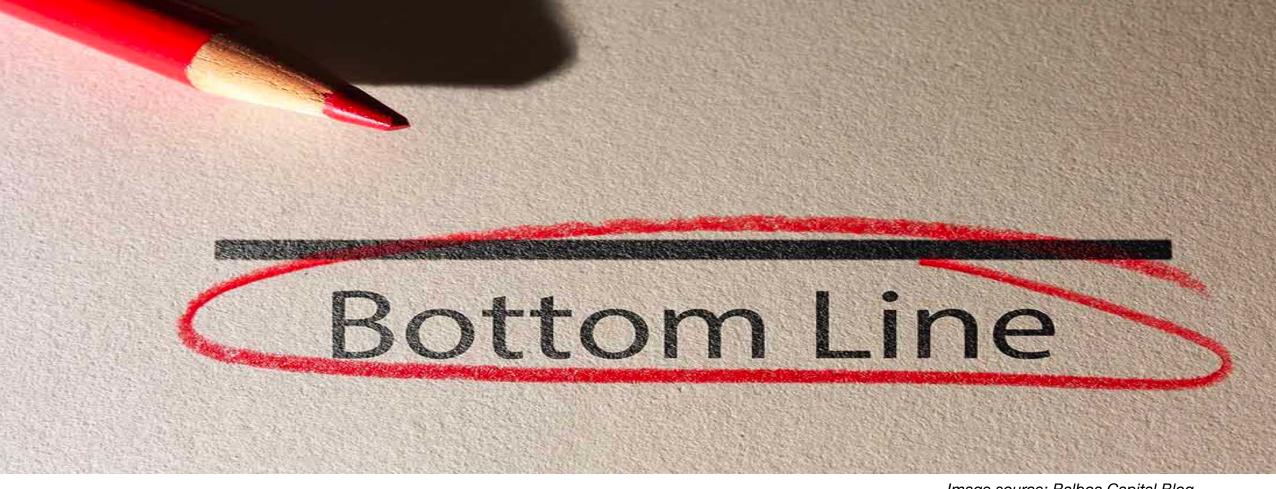


Image source: Balboa Capital Blog

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

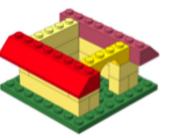
Exit survey

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

Keep building!



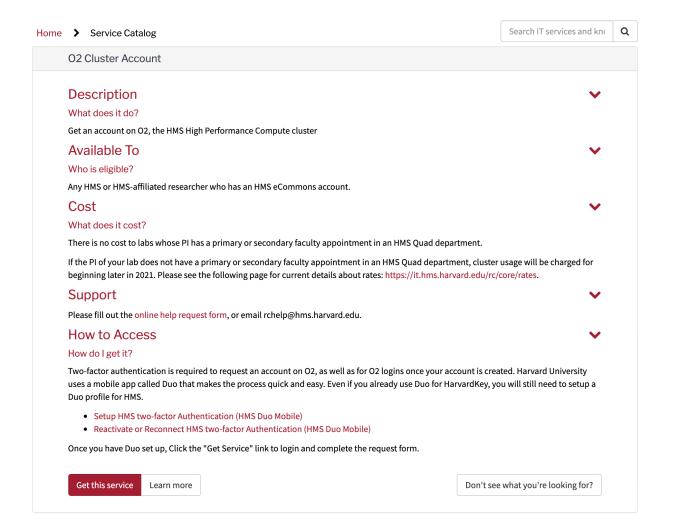






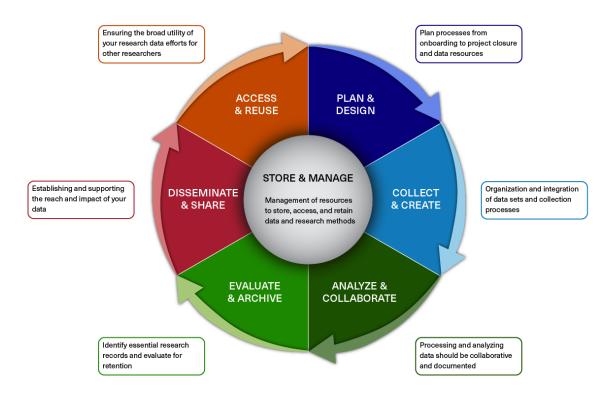
- Upcoming HBC advanced shell-based workshops:
 - Introduction to Variant Calling
 - May 28th, May 31st, June 4th, June 7th (in-person)
 - Using paired tumor-normal samples to analyze variants
 - Understanding chromatin biology using high throughput sequencing
 - Add dates here (online)
 - ChIP-seq, CUT&RUN, ATAC-seq

Get an O2 account!



Research Data Management (RDM)

BIOMEDICAL RESEARCH DATA LIFECYCLE



Date	Time	Event	Location
Jun 5	10am	RCBio: easy and quick HPC pipeline builder & runner	In-person
Jun 10	2pm	protocols.io webinar: Tips and Tricks	Zoom
Jun 14	12pm	Level Up Your GIS Skills: A Three- Part Series (Part 1)	Countway Library Classroom L2-025
Jun 21	12pm	Level Up Your GIS Skills: A Three- Part Series (Part 2)	Countway Library Classroom L2-025
Jun 26	1pm	R basics	Zoom
Jun 28	12pm	Level Up Your GIS Skills: A Three- Part Series (Part 3)	Countway Library Classroom L2-025

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



Thursday June 6th, 2024 9:00 to 10:30am

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



- HBC training team: hbctraining@hsph.harvard.edu
- HBC consulting: bioinformatics@hsph.harvard.edu
- O2 (HMS-RC): rchelp@hms.harvard.edu