

# Introduction to the command-line interface (shell)

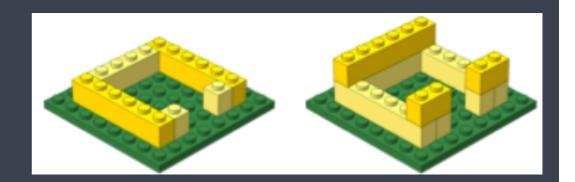
Harvard Chan Bioinformatics Core

in collaboration with

FAS Research Computing

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

### Learning Objectives



- ✓ Learn what a "shell" is and become comfortable with the commandline 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-hsph-shell-exit

### 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/

#### Thanks!

- Daniel Caunt and Maggie McFee from FAS-RC
- Data Carpentry

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#### 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

FAS-RC: create a ticket

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