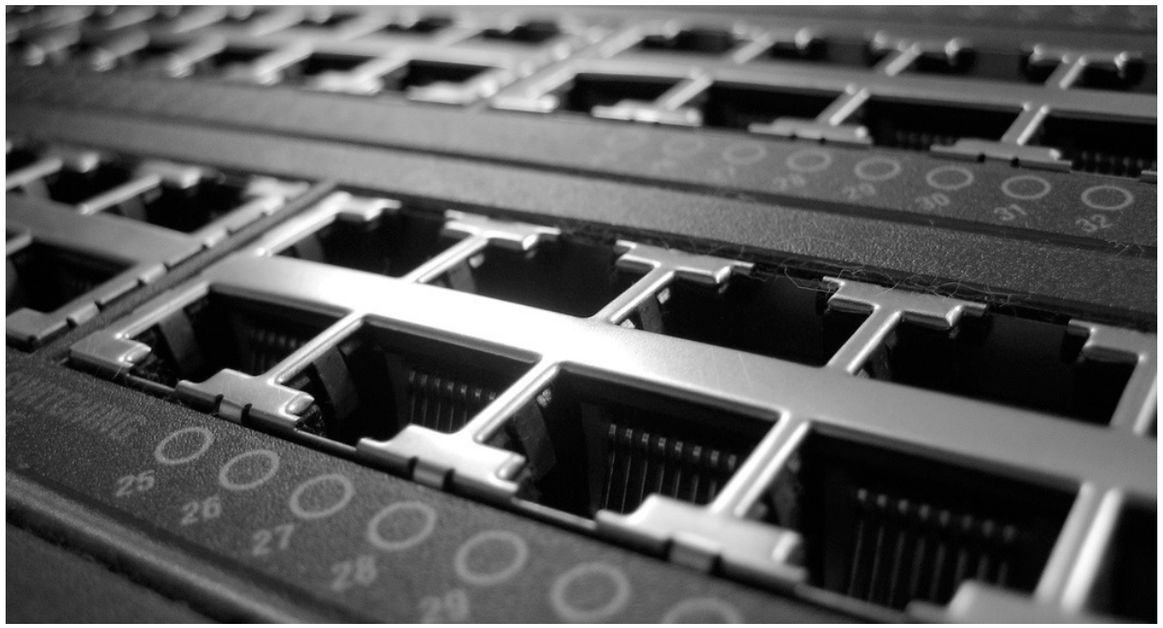


Introduction to the command-line interface

(shell)

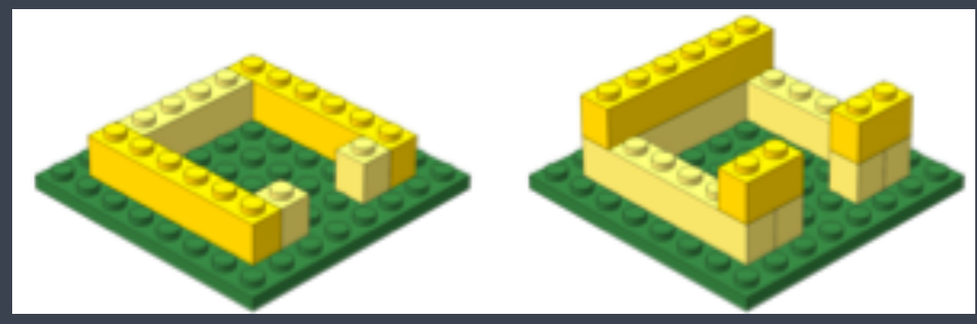
Harvard Chan Bioinformatics Core

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 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-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!

- FAS-RC crew
  - Daniel Caunt
  - Maggie McFee
  - Raminder Singh
  - Muneeba Syed

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

Twitter

@bioinfocore