Evolutionary Biology Metabarcoding Big Data wgs Next Generation Sequencing

BIOinformatics Working Group

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Objectives

- 1. Next-Generation technologies
- 2. Linux terminology
- 3. SSH into Pod —> Command Line basics
- 4. Best practices
- 5. Github

Quick Quiz

1. ___ command is used to 2. command is used to list contents of a directory. remove a directory. a. rdir a. tar b. Is b. remove c. rd c. dir d. rmdir d. lp 3. ____ command is used to 4. ___ command is used to grab specific strings in a file. see columns in a file. a. awk a. grep b. grab b. cut c. less c. sort d. head d. a and b

5. #!/bin/sh is called what?

Quick Quiz

- 1. __ command is used to 2. command is used to remove a directory. list contents of a directory. a. rdir a. tar b. Is b. remove c. rd c. dir d. rmdir d. lp
- 3. ____ command is used to 4. ___ command is used to grab specific strings in a file. see columns in a file.
 - a. grep
 - b. grab
 - c. less
 - d. head

- - a. awk
 - b. cut
 - c. sort
 - d. a and b

5. #!/bin/sh is called what?

sha-bang, hashbang, pound-bang

NGS technologies









- Single Molecule Real Time (SMRT) technology
- 35,000 70,000 good reads cell⁻¹
- ~11% error rate (as of 2013)
- Read length >45 kb and >190 kb
- More info: https://www.pacb.com/smrtscience/smrt-sequencing/





- Most common
- millions billions of reads lane⁻¹
- Low error rate
- Average read length ~300 bp
- More info: https://www.illumina.com/ techniques.html





- COOLEST sequencer ever!
- Long reads (e.g., 1 mbp)
- High error rate (~5-15%)
- More info: https://nanoporetech.com/how-it-works

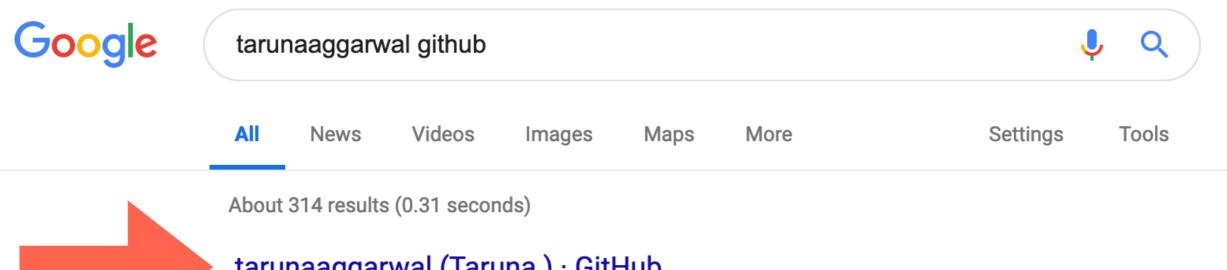
Key Linux Terms

- Linux a type of operating system
- Terminal a place where you type input and receive output
- Shell the program that interprets what you type in the terminal
- Bash a type of shell
- Directory another name for a folder
- Root the top level directory (denoted by /)
- Root the person with admin privileges
- Permissions describe who can do what to files and directories

SSH into Pod

SSH key

https://github.com/tarunaaggarwal/bwg-s19



tarunaaggarwal (Taruna) · GitHub

https://github.com/tarunaaggarwal ▼

tarunaaggarwal has 11 repositories available. Follow their code on GitHub.

GitHub - tarunaaggarwal/labsite

https://github.com/tarunaaggarwal/labsite ▼

Contribute to tarunaaggarwal/labsite development by creating an account on GitHub.

GitHub - tarunaaggarwal/phylosift4NPRB-2017

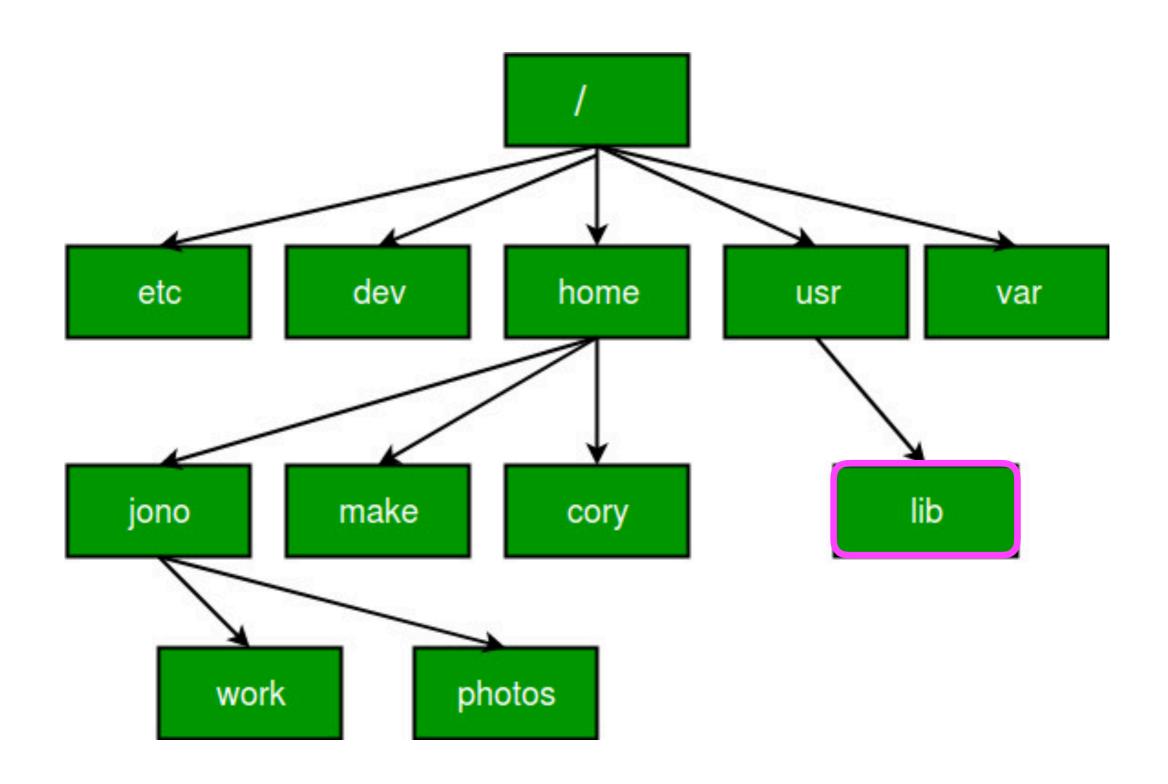
https://github.com/tarunaaggarwal/phylosift4NPRB-2017 ▼

Contribute to tarunaaggarwal/phylosift4NPRB-2017 development by creating an account on GitHub.

Command Line Basics

Command	Purpose
\$ 1s	shows a listing of dir contents
<pre>\$ pwd</pre>	print working dir
\$ cd	to navigate between dirs
\$	one dir above the current dir
\$.	current dir
\$~	home dir
<pre>\$ clear</pre>	to clear your screen
\$ rm	to remove a file
<pre>\$ rmdir</pre>	to remove a dir
<pre>\$ mkdir</pre>	to create a dir

PATH



Use a text editor to organize commands

Post your code on Github

Best Practices

Use a consistent directory structure

Case MaTTerS

Avoid spaces in file and dir names

Use a text editor to organize commands

Post your code on Github

Best Practices

Use a consistent directory structure

Case MaTTerS

Backup your work! Esp. scripts!!!

Avoid spaces in file and dir names

Review

1. Next-Generation technologies



- 2. Linux terminology (Linux is an OS)
- 3. SSH into Pod —> Command Line basics
- 4. Best practices
- 5. Github

Next Time

- 1. Next-Generation technologies —
- PACBIO®

 Illumina®

 NANOPORE
- 2. Linux terminology (Linux is an OS)
- 3. SSH into Pod —> Command Line basics
- 4. Best practices
- 5. Github
- 6. Files type and amplicon sequencing & analysis