

Chem 160 Module 19 Linux

Segment 1 Overview of the Linux Shell

Topics:

1. The Linux shell

Online references:

1. Bash shell
 - a. http://www.linuxtopia.org/online_books/bash_guide_for_beginners/index.html
 - b. List of Bash tutorials: <http://wiki.bash-hackers.org/scripting/tutoriallist>
 - c. Interactive Shell tutorial: <http://www.learnshell.org>
 - d. More tutorials on the shell: <http://swcarpentry.github.io/shell-novice/>
 - e. <http://www.ibm.com/developerworks/linux/library/l-bash/index.html>

I. The Shell

A. What is a shell?

- a. Program that interprets what you type
- b. Environment that keeps track of your previous commands and env. variables
- c. Programming language that lets you customize your environment and toolkit

B. How many shells are there?

- a. Many, but most people use one of a small group (most at least 30 years old):
 - i. **bash**—this is what we will learn, which is by far the most widely used
 - ii. **tcsh**—a more advanced descendent of the even older **csch**

tcsh

```
set file="/usr/local/prog.py"
```

```
echo $file
```

```
echo $file:t
```

```
echo $file:e
```

```
echo $file:h
```

```
echo $file:t:r
```

```
exit
```

- iii. **zsh**—expanded version of Bourne shell with great file completion

zsh

```
ls /u/lo/in[TAB]
```

```
cd [TAB]
```

```
ls -[TAB]
```

```
exit
```

- iv. **fish**—a shell designed to be user friendly (and have a sense of humor)

fish

```
ls /usr/local/bin
```

```
ls /var/**/*.log
```

```
exit
```

- b. For more info see: https://en.wikipedia.org/wiki/Unix_shell

C. You can switch between shells quite easily, just type the shell name to start a new shell and then **exit** to quit from a shell.

D. Look at how bash remembers what you've done

a. **history**

b. Recalling old commands

- i. Up arrow
- ii. Recovering commands by number !29
- iii. Recovering commands without executing them !29:p
- iv. Searching for old commands Ctrl-r text
- v. Try typing: Ctrl-r **nano**

Note that the search is *incremental*, updating with each new letter typed.

E. Let's look at some other features--download the GitHub repository for this module:
git clone https://github.com/mcolvinphd/chem160module19.git

F. Wildcards (aka "globbing")

- a. A wildcard is a pattern that can match multiple files (or directories)
- b. Many commands can take wildcards
 - i. "*" match 0 or more characters
 - ii. "?" match 1 character—can use several
 - iii. [ABC] match any character in the list
 - iv. [a-e] match in range
 - v. {a?,b??} match either a? or b??
- c. Let's use wildcards to sort college abbreviations:

```
cd colleges
```

```
ls ???
```

```
ls *U??
```

```
ls UC?
```

```
UC?[A-C]
```

```
ls UC{S?,?}
```

- d. Let's try it to collect output files

```
cd ../dna
```

```
ls DNA*.pdb
```

```
ls DNA*1.pdb
```

```
ls DNA?1.pdb
```

```
ls DNA?[135]? .pdb
```

```
ls DNA[0-1][3-5][13579].pdb
```

```
ls DNA[0-1][3-5][13579].pdb > dna.out
```

- e. You can use basic wildcards in the grep command (cd back to ~/class2a)

```
grep GL[YN] PK.pdb | grep CA
```

```
grep [A-Z]E[A-Z] PK.pdb | grep CA
```

```
grep 3[02468][13579][02468][7-9] PK.pdb | grep CA
```

```
grep 3[02468][13579][02468][7-9] PK.pdb | grep CA > CA.out
```

- f. A more sophisticated form of wildcards are "regular expressions" (Regex's) which are much more flexible (and complex) than wild cards
- g. Regex's are used in many languages and Linux commands, including Perl, Python, awk, egrep, and sed. If you've ever filling in an electronic form, you've probably used regex's.
- h. Regex's can be used in Bash scripts, as we'll see when we cover conditionals in Bash
- i. Example of using regular expression in Bash—match only sets of 8 digits [0-9]. Enter this at the Bash prompt:

```
date=11182019
```

```
[[ $date =~ ^[0-9]{8}$ ]] && echo "Date" || echo "Not date"
```

```
date=111819
```

(use up arrow to get this command back)

```
[[ $date =~ ^[0-9]{8}$ ]] && echo "Date" || echo "Not date"
```

- j. A more complicated date regex matcher is in the script: **datetest.bash** which will match dates of the format 11/18/19, 11-18-19, 11/18/2019, etc. and will check whether the ranges are correct for the days (01-31) and months (01-12) but not whether the dates actually exist in a given month (e.g. 02/31/2019 is okay)

```
datetest.bash 11/19/2019
```

```
datetest.bash 13/19/2019
```

```
datetest.bash 11/39/2019
```
- k. If you search for "date regex expressions" you'll see more sophisticated regex's that check if a date actually exists in a given year.

Chem 160 Module 19 Linux Segment 2 First Bash Script Example

Topics:

1. Examples of Shell scripting

I. A first immersion into shell scripts

- A. Shell script is a program written in the language of the shell
 - i. Bundles a set of commands in a useful way
 - ii. Can includes loops over sets of files or input values
 - iii. Can include conditionals (if statements) to do different things under different conditions
- B. cd into the **chem160module19** directory
- C. Edit a file called **files.bash** (using **nano**)
 - i. Go into insert mode and start the script with line **"#!/bin/bash"**
 - ii. With the exception of the first line, we use **"#"** to comment out a line (needed for debugging scripts or adding explanatory comments)
 - iii. Add the following 3 lines to the script (note quotes in echo optional)
echo "Directory contains this many files and dirs:"
pwd
ls | wc | colrm 10
- D. Make the script runnable by giving it "execute" permission:
chmod +x files.bash
- E. Run the script:
files.bash

II. Miniproject: Script to count how many of a particular amino acid is in a pdb file.

- A. This script will require 2 inputs, the amino acid abbreviation and the file name
- B. The Bash shell provide a special way to pass command line arguments in a script.
- C. If we have a script called **count_aa.bash** and run it as follows:
count_aa.bash GLY PK.pdb
Then inside the script:
 - i. **\$1** will contain "GLY"
 - ii. **\$2** will contain "PK.pdb"
- D. Writing the script:
 - i. Use nano to edit a file called **count_aa.bash**
 - ii. First line: **#!/bin/bash**
 - iii. Now echo what the result means:
echo -n "The number of \$1 amino acids in \$2 is:"
 - iv. Next use the grep, wc and colrm commands to get this information
grep \$1 \$2 | grep CA | wc | colrm 10
 - v. Now save the script and make it executable
 - vi. Try running it:
count_aa.bash GLY PK.pdb
count_aa.bash ALA PK.pdb
 - vii. What if you forget an argument or misspell the pdb file name?

- ```

count_aa.bash PK.pdb
count_aa.bash GLY PL.pdb

```
- viii. Let's improve the script by adding some *conditionals* to test for these issues
  - ix. Edit the script again and add the following right after the first line

```

if [$# -ne 2]; then
 echo "Script takes 2 arguments, an AA and a pdb file"
 exit 1
fi

```
  - x. Test running the script with the wrong number of arguments
  - xi. Edit the file again and add the following lines after the first conditional

```

if [! -f $2]; then
 echo "There is no file $2"
 exit 1
fi

```
  - xii. Test running the script with a misspelled pdb file name
  - xiii. Finally let's put your script into a loop (hint: this will help in Homework 2):

```

for i in GLU GLN GLY ALA PHE
do
count_aa.bash $i dynein.pdb
done

```