Introduction to Linux – Part 2

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Linux Virtual Machine

Get a temporary account (or use your own CHPC account)

□ Virtual Machine FastX portal: <u>https://linuxclass.chpc.utah.edu:3300</u>

Open a XFCE Terminal

□ Adjust Font size: Edit → Preferences → Appearance → Click on Font → adjust Font Size

- □ Use Bash shell (quick check: echo \$SHELL); □How to change to bash: /bin/bash
- Copy and Paste issue on Mac;
- Prepare example data (if you missed Part 1)
 - □cd ~
 - Image: Contract of the second seco
 - Cd LinuxClass
 - □wget <u>https://home.chpc.utah.edu/~u0028729/CHPCPresentation/shell-lesson-data.zip</u>
 - unzip shell-lesson-data.zip
 - Cd shell-lesson-data

Editors

There are many choices – a few are:













Nano Editor

To start either

nano

OR

nano filename

-- if filename exists, it will open file in editor; if it does not, this will be the name used when you save the file.

if you start nano without a filename it will prompt you for a name when you "WriteOut" using ^O (Ctrl + O)

^? \rightarrow Ctrl + Letter Key; How to quit nano session: ^X \rightarrow Ctrl + x



Nano cheat sheet: https://www.nano-editor.org/dist/latest/cheatsheet.html

Vi editor

Another common choice

Start with the command vi or vi filename

- **vi** at CHPC is actually **vim**, which is an improved version of **vi**
- □ More feature rich, takes more time to learn
- Not going into detail but we do provide a vi cheat sheet and a vi graphical cheat sheet – linked on the presentation page <u>https://www.chpc.utah.edu/presentations/IntroLinux3parts.php</u>
- □ There is also a tutor program start with command **vimtutor** which is a great tool to learn to use the program
- □ How to quit Vi session (discard changes): ESC \rightarrow : \rightarrow q!

Loops

- Used when you want to preform the same action many times, such as on multiple files
- There are a number of ways you can do this
- One option
 - List multiple arguments for a command to act upon
 - DExample (go to the LinuxClass/shell-lesson-data/exercise-data/creatures directory):

Dhead –n 3 basilisk.dat minotaur.dat unicorn.dat

□Another option – do a loop with a for/do statement

Loop Terminology

□ in bash syntax a loop looks like:

Bash	
<pre>for thing in list_of_things do operation_using \$thing done</pre>	# Indentation within the loop is not required, but aids legibility

- In this loop thing is a variable. During execution \$thing is set to the first item in the list, the operation(s) is done, then it goes to the second and repeats the operation(s), etc until it reaches the last item in the list. Then the loop is exited.
- □You can choose anything for **thing** (eg mything, item, myfile...) however, your choice of the what to use for thing should help a person reading the file understand what the loop is doing and what it is acting upon
- Examples of list_of_thing: 2 4 6 8 10;

```
{2..10..2}; {start..end..step}
fileA fileB fileC fileD;
file*;
$variable or $(command) representing a list: $(ls)
```

Exercise

□ in bash syntax a loop looks like:

Bash	
<pre>for thing in list_of_things do</pre>	# Indentation within the loop is not required, but aids legibility

- Go to the directory shell-lesson-data/exercise-data/proteins and write a loop that lists all of the pdb files
- □Write a loop that prints all file names can use **echo** command
- Build upon this loop by adding another command
 - Copy each file to a new name based on the existing name, file.pdb to orig-file.pdb
- Add another command to the loop to make a single file that has the content of all of the pdb files in this directory
 - □ can use redirect >, >>

for file in *.pdb
do
 echo \$file
 cp \$file orig-\$file
 cat \$file >> all.pdb
done

Exercise – Nelle's Data

- To process her data files Nelle will need to run an analysis on each of her sample files in north-pacific-gyre. The files to be processed have the consistent names of NENExxxA.txt and NENExxxB.txt
- The analysis requires running a script more on this next time written by her supervisor is called goostats.sh; this script acts on one sample file at a time; it requires two arguments the input file name and the output file name.
 Dbash goostats.sh <INPUT> <OUTPUT>
- Nelle decides to call the output file stats-NENExxxA.txt prepending the filename with stats-
- □She is being careful so she wants to test (using echo instead of running the script
- □ Hint start with 1 file and run test to create the two arguments. You can prepend a variable with additional information: stats-\$variable

Exercise – Nelle's Data Answer

To make sure getting all of the files for datafile in NENE*A.txt NENE*B.txt do echo \$datafile

done

Test getting the output file name for datafile in NENE*A.txt NENE*B.txt do

echo \$datafile stats-\$datafile done

Executing the script across all files for datafile in NENE*A.txt NENE*B.txt do echo \$datafile bash goostats.sh \$datafile stats-\$datafile done

Some other useful Linux commands

□cut-e.g. cut -d , -f 2,3 animals.csv

□Example file: shell-lesson-data/exercise-data/animal-counts
□-d (delimiter) -f (column ids)

□ Prints selected parts of lines from file to standard output (screen)

□du-e.g. du -h or du -sh

□Scan a given file/directory (and subdirs) and report space usage; -s give summary of total usage, -h gives it in "human readable" format of K, M, G

$\Box df - e.g. df - h$

Overview of file system disk space usage (-h: human readable)

Dln-e.g. ln -s ~/bin/prog.exe progl.exe

□ create a link between files (In -s FILE LINK)

On your own – Use and explore options of these commands

Linux File Permissions



Can also use a for all to set u, g, and o to same settings

Chmod – to change permissions of file or directory, can set

Examples:

```
□chmod g=rwx file
□chmod g+x file
□chmod o-rwx *.c
```

Executable files (programs and scripts) must have executable permissions; directories must be executable in order to be able to cd into them

 \Box chmod +x *.sh

Login Scripts & Environment Variables

In your home directory are a number of dot files - .bashrc and .custom.sh, .tcshrc and .custom.csh Depending on your shell choice, the appropriate pair of these are executed during login.

- These set the environment (as environment variables) needed for you to work on CHPC resources
- **Commands to check your environment:** env **or** printenv
- □Some important variables
 - □\$USER
 - □\$HOME
 - □\$PATH paths to search for commands

□\$LD_LIBRARY_PATH – paths to search for libraries when linking a program (more on that later)

Processes

- A Process is a running Linux program
 Each process has a PID (Process ID)
- top displays processes and resource usage in real time (Ctrl + C to quit)
 Itop –u <user>
 Ctrl + C to quit
- **ps** reports a snapshot of current processes
 - Display ALL of your processes
 - Display ALL processes
 - Display ALL processes (more detailed)
- □ kill PID kills the process with the specified PID
- **killall processname** kills all process with the processname
- □ kill -9 PID kills the process with the specified PID if a kill does not work

Monitoring processes/usage

- **uptime** how long the system has been running
- □ **free** free –h, memory and swap usage
- enhanced top
 - **atop** (available on CHPC clusters)
 - □ htop (available on CHPC clusters)
- □ sar historical system usage report (cpu, memory, I/O...)

Other Job Controls

Ctrl+C (^C) terminate the currently running process

- Ctrl-Z (^Z) suspends the currently running process
 Example: ping www.byu.edu
- **jobs**: lists all jobs, with their number
- □ **fg** %**n**: bring a program back to the foreground (and continue/resume execution)
- □ **bg** %**n**: puts current or specified job (%n) in the background (and continue/resume execution)
- □ & (ampersand) runs the job in the background
 - Example: sleep 10 VS sleep 10 &

Moving files to/from CHPC

https://www.chpc.utah.edu/documentation/data_services.php

- Can mount CHPC file systems on your local machine (Windows, Mac or Linux), must be on campus or using the campus VPN
- □Windows there are graphical tools such as WinSCP
- □ Mac, Windows, cloud options cyberduck, another graphical tool
- 🖵 Linux
 - **scp** command (secure shell copy) to copy files between linux systems
- □ wget to download from web with URL
 - **Curl** is another option
- For larger data sets look into the Data Transfer Nodes (DTNs) and transfer tools such as Globus, see
 - <u>https://www.globus.org/quickstart</u>
 - <u>https://www.chpc.utah.edu/documentation/data_services.php</u>

Have Questions?

helpdesk@chpc.utah.edu