UNIX basics

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This slide set is called unix_basics.pdf and is located in the "15_unix_basics" folder of our Lectures repository.

• To learn basic UNIX commands

• To learn how to interact with running processes

Most of the content in this slide set is essential. You will need to become proficient with it to proceed in the course and to work at the command line in general.

Most of this is a review of what you read in the assigned Active Reading.

Warnings

- rm and rmdir are forever
- overwriting is forever (so be careful with cp, mv, etc.)
- a single whitespace matters a lot
 - rm -rf a*
 - ${\ensuremath{\, \circ }}$ recursively delete any file/folder starting with an ${\ensuremath{\rm a}}$
 - rm -rf a *
 - ${\ensuremath{\, \bullet }}$ recursively delete any file/folder starting with an ${\ensuremath{\rm a}}$
 - then do the same for *everything* (because * matches anything)
- these are all different, so be careful when copy-pasting
 - ",", and " (straight vs. curly double-quotes)
 - ',',', and ' (straight vs. curly single-quotes vs. back-tick)
 - -, -, and (hyphen, en dash, em dash)
- don't use spaces and special characters in file names
- don't work on the login node of htc (use srun -M teach -A hugen2071-2024f --pty bash)

- User interface for interacting with the UNIX operating system
- A program that accepts commands that are in turn other, more basic programs/utilities
- Important for scripting, making pipelines, calling other programs, organizing files, managing resource-intensive processes/jobs
- There are lots of versions
- We use bash here

- Very common
- Supports history (use the up arrow to save time!)
- Supports autocompletion (use tab to save time!)
- Supports wildcards (more on that later)
- Comments start with #

Files are organized hierarchically

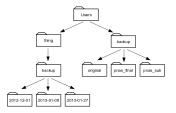


Figure 1: Example of a file tree (https://swcarpentry.github.io/shell-novice/02-filedir.html)

You can designate a file/folder with

- an absolute path (starting at the top of the file tree, root, /)
 - e.g., /Users/thing/backup/2013-01-08/
- a relative file path (starting at your working directory, .)
 - e.g., ../2013-01-08/ if you're currently in /Users/thing/backup/2012-12-01/

Shortcut	Meaning
~	home directory
	current/working directory
••	parent directory (up one from where you are now)
-	your previous working directory
/	• By itself: the root directory (top of the file tree)
	• Between names of folders: just a divider

Command	Meaning	Syntax	Useful options
pwd cd	print working directory change directory	pwd cd directory	
ls	list the files in a directory	ls directory	-lahFGpt

What does the output of 1s mean?



Figure 2: Columns of Is output

Commands for manipulating files

Command	Syntax	Meaning
ср	• cp file1 file2 file5 dir	 Copy one or more files into the folder
	• cp file1 file2	 Copy file1 as file2
mv	• mv file1 file2 file5 dir	 Move one or more files into the folder
rm -i	• mv file1 file2 rm file1 file2 file5	• Rename file1 as file2 Permanently delete one or more files
mkdir	mkdir dir1 dir2 dir5	Create one or more folders
rmdir	rmdir dir1 dir2 dir5	Permanently delete one or more empty folders

Who can do what with a file?

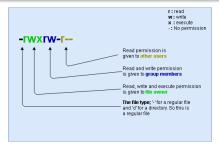


Figure 3: How to read file permissions (https://teaching.healthtech.dtu.dk/unix/index.php/File_permissions)

Permission code	Meaning
r	read file/list directory
W	edit file/make files in directory
х	execute file/cd into directory

Example	
command	Meaning
chmod -v a=rwx file	Give everyone (all) permission to read, write, and execute
chmod -v ug=rw file	Give yourself (the user/owner, u) and your group permission to read and write only
chmod -v o=r file	Give others read-only permission
chmod -v a+x file	Add (+) execute permission for all (without changing the ${\tt r}$ or w permissions)

-v means "verbose" and prints a useful message.

CommandMeaning		Syntax	Useful options
head	show first lines of file	head file	-n (first n lines)
tail	show last lines of file	tail file	-n (last n lines) or -n+ (start at line n)
more	read through file	more file	(Navigate with spacebar and q)
less	read through file	less file	(See documentation)
wc	get count of words, lines, and bytes	wc file	-1 (get only word count)

CommandUse		Example	Meaning	
cat	Prints contents of a file	cat file1.txt file2.txt	Print one or more files	
echo	Prints out whatever follows	echo "Hi there"	Print "Hi there"	
grep	Search for pattern in file • -i case- <u>i</u> nsensitive	grep gene file.txt	Print lines of file.txt containing "gene"	
	 -v invert (find non-matches) 			
	 color highlight matches 			

These are powerful - so be careful with them!

Wildcard	Matches	Example pattern	Example matches
*	Anything (zero or more characters)	*.txt	Any .txt file
?	Any single character	chr1?.txt	chr11.txt, chr12.txt, chr1a.txt (but not chr1.txt, chr111.txt, chr.txt)
[]	Anything in the set of characters in the brackets	chr[0-9][0-9]. chr[XY].txt	txtbrMN.txt where M and N are any integers 0-9 chrX.txt or chrY.txt

cat, echo, grep, wildcard examples

echo example echo "Echoing a comment to the screen. Then using a cat command!" Echoing a comment to the screen. Then using a cat command!

cat example
print out some short text files
echo "Below are the contents of my chr*.txt files"
Below are the contents of my chr*.txt files
cat data/chr*.txt
chr1:36926582
chr1:6782904
chr1:77840389
chr2:66318540
chr2:85739014

grep example echo "Here are all the lines of chr1.txt without a 3" Here are all the lines of chr1.txt without a 3 grep -v 3 data/chr1.txt chr1:66782904

Some more advanced pattern-matching options:

Regular expres- sion	Use	Example	Meaning
		- 1	8
^	Matches beginning of line	grep "^chr1" file.txt	Print lines of file that start with "chr1" (this would also match "chr12", etc.)
\$	Matches end of line	grep "0\$" file.txt	Print lines of file that end with the digit 0
[]	Matches lines containing characters listed inside the brackets	grep "[ACGT]" file.txt	Print lines of file that contain A, C, G or T $$
[^]	Matches lines containing characters not listed inside the brackets	grep "[^ACGT]" file.txt	Print any lines of the file containing characters other than A, C, G, or T

Look at top of the file head -n3 data3/tb1.fasta >gi|385663969|gb|JQ90005a1| Zea mays subsp. mexicana isolate IS9 teosinte branched 1 (tb1) gene, complete GCCAGGACCTAGAGAGGGGAGCGTGGAGAGGGGCATCAGGGGGCCTTGGAGTCCCATCAGTAAGCACATG TTTCCTTTGTGTGATTCCTCAAGCCCCATGGACTTACCGCTTTACCGACACAGCGCAGCTAAGCCCCGTCTT

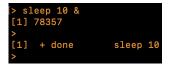
Pull out just the header grep ">" data3/tb1.fasta >xi|385663969|gb]JQ900508.1| Zea mays subsp. mexicana isolate IS9 teosinte branched 1 (tb1) gene, complet.

- A command you run is run as a "process" and assigned a process ID number (pid)
- By default a process runs in the **foreground** (you wait for it to finish before starting another)
- To keep interacting with the terminal, you can run a process in (or move it to) the **background**
 - Run it in the background by putting & after the command (this prints job id and pid)
 - To move a process from foreground to background
 - ctrl + z suspends the process
 - then bg to restart it in the background
 - Use jobs to see what jobs are running in the background (shows the commands)
 - Use ps to see process status (including pid)
 - Terminating a process
 - \bullet ctrl + z for a foreground process
 - kill followed by the pid for a background process

The sleep command suspends operations for a specified time. Here it runs for 10 seconds in the foreground:

> date; sleep 10; date Tue Oct 17 13:45:25 EDT 2023 Tue Oct 17 13:45:35 EDT 2023

Here it runs in the background, getting a job id (1) in addition to a process id (78357). Then a message when it finishes 10 seconds later:



Or we can start in the foreground and then background it:



To see what jobs are running in the background, use jobs. You can foreground a job with fg followed by its job id. (If you don't specify a job id, fg operates on the job with the +; and the job with - would be next after the + job finishes.)

> sleep 60 &	
[1] 78646	
> sleep 120 &	
[2] 78647	
> sleep 180 &	
[3] 78649	
> jobs	
<pre>[1] Running</pre>	sleep 60 &
[2] – Running	sleep 120 &
[3]+ Running	sleep 180 &

To check the status of all processes, use ps:

[> ps				
PID	TTY	TIME	CMD	
78552	ttys000	0:00.24	-bash	
78646	ttys000	0:00.00	sleep	60
78647	ttys000	0:00.00	sleep	120
78649	ttys000	0:00.00	sleep	180

To terminate a process, use kill:





- When it ends, a process returns an **exit status** stored in the variable \$?
- Exit code 0 means no error
- Any other exit code means error/failure for some reason

Exit status example

cat data/chr1.txt chr1:36926582 chr1:66782904 chr1:77840389 echo \$? 0 ehco "Hi" bash: line 6: ehco: command not found echo \$? 127 grep "369" data/chr1.txt chr1:36926582 echo \$? 0 grep "CHR" data/chr1.txt echo \$? grep "CHR" data/chr1.txt | cat echo \$? 0

- htc (high throughout computing) is the computing cluster we'll use for this class
- the CRC (Center for Research Computing) administers it
- don't work on the login node (always 'srun --pty bash' when you first log in via the terminal)
- you can also **submit** larger jobs using the workload manager, Slurm (we'll cover that later)

First, log into the VPN with GlobalProtect.

There are two ways to log on to the cluster

- Via the web: ondemand.htc.crc.pitt.edu
- Via a terminal window
 - ssh <your_user_name>@htc.crc.pitt.edu

For details, see

https://crc.pitt.edu/getting-started/accessing-cluster.

Next, always start an interactive job: srun --pty bash .

- When you first ssh onto the cluster, you're on the **login node**, which is only for logging into and not for working
- Always do srun --pty bash to start an "interactive job" when you log in via the terminal (default: 1 hour)
- If you work on the login node
 - you will slow down the cluster and inconvenience the many people using it
 - you will get yourself and me in trouble

Two main options

- the in-browser interface at ondemand.htc.crc.pitt.edu in the Files menu (recommended)
- a free FTP program like Cyberduck (instructions: https://crc.pitt.edu/managingdata)

You can also use the more cumbersome scp command (again, see https://crc.pitt.edu/managingdata).

- nano is a simple text command-line text editor we'll use
- others include vim and pico
- In an interactive job on htc, first load nano with module load nano
- Entering nano will open the editor
 - Just nano opens a new blank file
 - nano fileame opens a new/existing file (called filename)
- Along the bottom are commands, where ^X means CTRL + x and so on
- To exit: ^X, then type y to save, and hit RETURN

Use man to learn how commands work

Suppose you want to quickly see how a command (e.g., 1s) works. You can do a web search, use an LLM like ChatGPT, or consult the man page. I entered man 1s and paged down with spacebar:



Figure 4: Part of the man page for Is

total 1736						
drwxrwxrwx	4 jonathanchernus	staff	128B O	lct 17	2023	data
drwxr-xr-x	3 jonathanchernus	staff	96B O	lct 17	2023	data2
drwxr-xr-x	3 jonathanchernus	staff	96B O	lct 17	2023	data3
drwxr-xr-x	4 jonathanchernus	staff	128B O	lct 17	2023	data4
drwxr-xr-x	15 jonathanchernus	staff	480B O	lct 17	2023	figures
-rw-rr-@	1 jonathanchernus	staff	357B D	lec 9	2022	header_pagenrs.tex
drwxr-xr-x@	5 jonathanchernus	staff	160B S	lep 5	2023	images
-rw-rr-@	1 jonathanchernus	staff	25K S	ep 23	11:09	unix_basics.Rmd
-rw-rr-@	1 jonathanchernus	staff	803K S	ep 23	11:09	unix_basics.pdf
		D	A 4 1		LINUNZ I	

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Using git at the command line - make a personal access token first

In order to clone your repository (and push changes) to your home directory on htc, you will need to create a personal access token.

- log in to your GitHub account
- click on your profile picture (top right) and select Settings
- click Developer Settings at the bottom left and then Personal access tokens - Tokens (classic)
- click Generate new token Generate new token (classic)
- enter something like hugen2071_htc_token for the Note
- set the expiration date as 90 days
- under Select scopes click the repo box to automatically check the 5 boxes below it
- now click the Generate token button
- copy the token and paste/store it somewhere safe
- you won't be able to look it up again without resetting it

You only need to this step once:

- copy the ssh link for the repository you want to clone
- cd to the directory you want
- enter git clone link (paste your link in place of the word link)
- provide your username and personal access token at the prompt

Using git at the command line - adding, committing, pushing

- First, if you create or change any files, enter git add -A
 - make sure all of the changes are staged for the next commit (otherwise git won't keep track of the changes)
 - or do git add file1 file2 if there are files you don't want under git control
 - do this every time before you commit/push
- Second, commit
 - enter git commit -m "message" (where "message" is a short, useful description of what changes you've made
 - do this every time before you push
- Finally, push
 - enter git push
 - this ensures your commit is saved remotely on GitHub