



Lab #1

Recommendations for a successful lab

BINF 2111, Fall 2023



Introductions

- Madeline (Madeline or Maddy)
- RAW Lab for 2 years
- Ph.D. in Bioinformatics and Computational Biology
 - M.S. in Bioinformatics (UNC Charlotte, 2022)
 - B.S. in Computer Science (UNC Charlotte, 2021)
 - B.A. in Anthropology (UNC Charlotte, 2021)
- Office: BINF 360
 - Office hours: T, R, F by appointment in person, M-F by appointment virtually

What is your name?

(I promise I will learn them all)

Recommendation #1 - Terminology

Know your terminology and commands, as this will be used in tests/quizzes.






- **Terminal:** a command line interface (CLI), where you can type commands, manipulate files, execute programs, and open documents
- **Directory:** folder or path to a folder/file
- **UNIX/bash:** language used in terminal
- **Print:** display information
- **Command:** a specific word or phrase that tells the computer what to do
- **Run:** execute a command or program
- **Options/flags:** an addition to a command that slightly changes the command in a specified manner



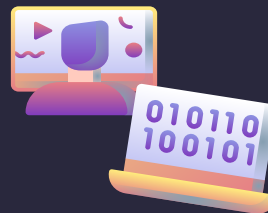


Command	Meaning	Usage
ls	lists everything in a directory	ls [options] [folder]
echo	prints text to a location	echo [phrase]
mkdir	create new directory	mkdir [folder name]
cd	change directory	cd [directory]
touch	make new file without any content	touch [file name]
more	view file one screen at a time	more [file]
cat	print full contents of file	cat [file]
mv	move file to a different location, rename file/folder	mv [file] [new location] mv [old name] [new name]
pwd	print current location (working directory)	pwd
wc	count the number of lines/words/bytes in a file	wc [options] [file]

 Command Options Input (like a file or folder)

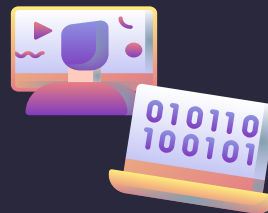
Recommendation #1 - Commands

- **ls**: lists everything in a directory
 - Useful options:
 - -a show all files (including ones that start with .)
 - -l use long listing format (file sizes, dates, permissions, etc)
 - -h use human readable format (1G, 27K, 736M)
 - -t sort by time with newest first
 - -o similar to -l, but without group permissions
 - -r reverse the order while sorting
 - Usage:
 - `ls -thor`
 - `ls -alh Desktop/`



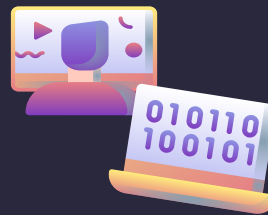
Recommendation #1 - Commands

- **wc**: count the number of lines/words/bytes/characters in a file
 - Default (no options) prints out:
line count word count byte count
 - Useful options:
 - `-c` print the byte count
 - `-l` print the line count
 - `-m` print the character count
 - `-w` print the word count
 - Usage:
 - `wc file.txt`
 - `wc -l file.txt`



Recommendation #1 - Commands

- **echo**: prints text to a location
 - Important notes:
 - Using **> file** after the command will put the text in the file
 - Using **>> file** after the command will append the text to the end of the file.
 - Usage:
 - `echo "hello"`
 - `echo "hello world" > file.txt`
 - `echo "hello student" >> file.txt`



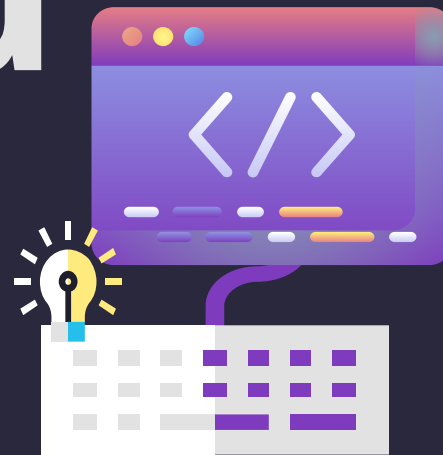
Recommendation #1 - Operators

Operator	Meaning	Usage
	Pipe, used as “and” or used to use the output of the first command as the input of the second command.	<code>echo “hello” echo “world”</code> <code>ls wc -l</code>
>	Output, put the output of the command into a file	<code>echo “hello” > file.txt</code>
>>	Append, add the output of a command to the end of a file	<code>echo “hello” >> file.txt</code>
*	Wildcard, used as a placeholder for any character for zero or more times	<code>cat file*</code>

```
((base) madelinebellanger@Madelines-MacBook-Air Lab1 % cat file1.txt
Hello Jose
((base) madelinebellanger@Madelines-MacBook-Air Lab1 % cat file2.txt
Hello World
((base) madelinebellanger@Madelines-MacBook-Air Lab1 % cat file3.txt
Hello Jose
Hello World
((base) madelinebellanger@Madelines-MacBook-Air Lab1 % cat file*
Hello Jose
Hello World
Hello Jose
Hello World
```

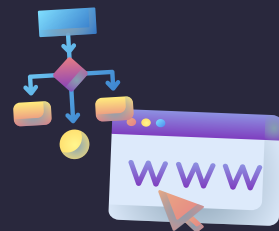
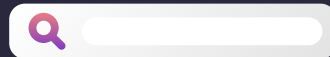



EVERYTHING IS TYPED



Recommendation #2

- Use your resources:
 - Lectures
 - Me, Dr. White, Andra, and your peers
 - Websites: GeeksForGeeks, Explainshell, Bash scripting cheatsheet
 - AI: ChatGPT, DALL-E, DeepMind, etc.
 - Your terminal
 - Test commands out!
 - **man** command
 - Pulls up the manuals for each command
 - Press Q to exit



Recommendation #3

- Use your tab button to autofill file/folder names.

```
Lab1 — -zsh — 68x24
((base) madelinebellanger@Madelines-Air Lab1 % ls
file1.txt      file2.txt      file3.txt
(base) madelinebellanger@Madelines-Air Lab1 % cat f
```

tab

```
Lab1 — -zsh — 79x24
((base) madelinebellanger@Madelines-Air Lab1 % ls
file1.txt      file2.txt      file3.txt
(base) madelinebellanger@Madelines-Air Lab1 % cat file
```

tab

```
Lab1 — -zsh — 79x24
((base) madelinebellanger@Madelines-Air Lab1 % ls
file1.txt      file2.txt      file3.txt
(base) madelinebellanger@Madelines-Air Lab1 % cat file
file1.txt file2.txt file3.txt
```

```
Lab1 — -zsh — 79x24
((base) madelinebellanger@Madelines-Air Lab1 % ls
file1.txt      file2.txt      file3.txt
(base) madelinebellanger@Madelines-Air Lab1 % cat file1
```

tab

```
Lab1 — -zsh — 79x24
((base) madelinebellanger@Madelines-Air Lab1 % ls
file1.txt      file2.txt      file3.txt
(base) madelinebellanger@Madelines-Air Lab1 % cat file1.txt
```

Recommendation #4

Tip: Do not use spaces in your file and folder names!

Make a directory specifically for this class.

```
[(base) madelinebellanger@Madelines-Air ~ % cd Desktop Move to my Desktop
[(base) madelinebellanger@Madelines-Air Desktop % ls List everything on my Desktop
Flow Data NFixDB
GLK_assembly_HQ_large-only.fasta PhD
GLO_best.fasta RYT200
GLO_plasmid_best.fasta Screenshot 2023-08-10 at 1.45.30 PM.png
MicroscopyImages cyclops-main
[(base) madelinebellanger@Madelines-Air Desktop % mkdir BINF2111 Create a new directory for this class called BINF2111
[(base) madelinebellanger@Madelines-Air Desktop % ls List everything on my Desktop again to confirm the new directory was
BINF2111 NFixDB created
Flow Data PhD
GLK_assembly_HQ_large-only.fasta RYT200
GLO_best.fasta Screenshot 2023-08-10 at 1.45.30 PM.png
GLO_plasmid_best.fasta cyclops-main
MicroscopyImages
```

Make a subdirectory for each lab.

```
[(base) madelinebellanger@Madelines-Air Desktop % cd BINF2111 Move to the BINF2111 folder
[(base) madelinebellanger@Madelines-Air BINF2111 % mkdir F23 Create a new directory for Fall 2023 (not necessary)
[(base) madelinebellanger@Madelines-Air BINF2111 % cd F23 Move to the F23 folder
[(base) madelinebellanger@Madelines-Air F23 % mkdir Lab1 Create a new directory for Lab 1 called Lab1
[(base) madelinebellanger@Madelines-Air F23 % ls List everything in F23 to confirm the new directory was created
Lab1
```

Helpful Hints

- Copy and paste does not always work :(
 - When in doubt, type it out!
- Press the up and down arrows to cycle through previous commands.
- Use the **history** command to see a list of your past commands.

```
[(base) madelinebellanger@Madelines-MacBook-Air Lab1 % history -9
702 cd Desktop/BINF2111/F23/Lab1
703 echo "Hello Jose" > file1.txt
704 echo "Hello World" >> file2.txt
705 cat file1.txt file2.txt > file3.txt
706 cat file1.txt
707 cat file2.txt
708 cat file3.txt
709 cat file*
710 history
```

Lab Notes

- Question 10:

```
echo "Hello Jose" > file1.txt | echo "Hello World" >> file2.txt
```

- Underlined portion will **not** work for **Mac** users
- Run the following for the last command (cat...) to work:
 - `echo "Hello Jose" > file1.txt`
 - `echo "Hello World" >> file2.txt`
- You still need to know what the command does!
 - You can run it on Replit to see it working.