Introduction to UNIX

Genome 559: Introduction to Statistical and Computational Genomics

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What is UNIX?

- Unix is a family of operating systems (like Linux)
- The language of the command-line (except on PCs, until Windows 10; see http://www.howtogeek.com/249966/how-to-install-and-use-the-linux-bash-shell-on-windows-10/)
- Set of programs/commands for:
  - Navigating file directories
  - Manipulating text files
  - (and much more we won't cover)

Why should you care about UNIX?

- Have you ever spent a long time systematically renaming files one by one, only to miss some?
- Do you ever have to work with files too large for your text editor?
- Will you have to work on a computer cluster?

You've already been using UNIX!

```
$ cd Documents
$ python hello.py
Hello, World!
```

Where are we?

Print working directory

```
$ pwd
/Users/seungsoo
```

(this is called a path)

List files and subdirectories

```
$ ls
Applications
Desktop
Documents
…
```

Let’s go somewhere …

Change directory

```
$ cd Documents
$ pwd
/Users/seungsoo/Documents
```

We’re now in a subdirectory.

Go up one level

```
$ cd ..
$ pwd
/Users/seungsoo
```

Let’s make a new folder (directory)

Make directory

```
$ mkdir GS559
```

Move into that directory

```
$ cd GS559
```

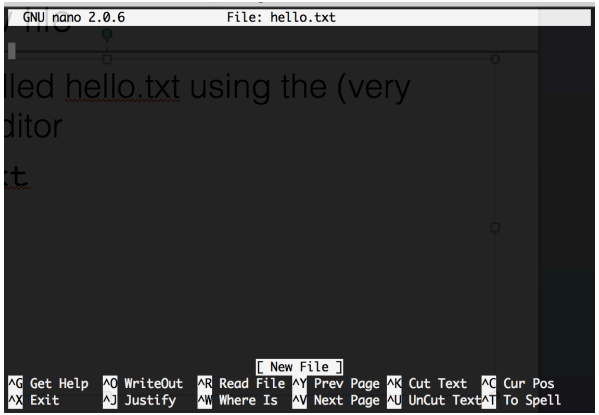
Go back home

```
$ cd
```

Let’s write a new file

Write a new file called hello.txt using the (very basic) nano text editor

```
$ nano hello.txt
```



nano text editor

- Instructions are at the bottom of the screen
- ^X means Control-X, etc.
- To save: ^X, then y, then Enter

Renaming/moving files

Rename hello.txt to greetings.txt

```
$ mv hello.txt greetings.txt
```

Move greetings.txt to the GS559 folder

```
$ mv greetings.txt GS559
```

Move greetings.txt back to the current folder

```
$ mv GS559/greetings.txt .
```

The single dot “.” is a shortcut for the current directory

Copying and deleting files

Make a copy of greetings.txt called greetings2.txt

```
$ cp greetings.txt greetings2.txt
```

Remove (delete) greetings2.txt

```
$ rm greetings2.txt
```

Be super careful with `rm` – unlike files put in the “Recycling Bin”, files deleted with `rm` are permanently gone.

Remember, you can use `ls` to check what files are in your current location.

Summary of part 1 (file navigation)

`pwd`	print working directory
`cd dir`	change directory to dir
`cd ..`	go up one level
`ls`	list directory contents
`nano file1`	edit file1 using text editor nano
`mv file1 file2`	move/rename file1 to file2
`cp file1 file2`	copy file1 and save as file2
`rm file1`	delete file1

The wildcards: * and ?

- UNIX is particularly powerful because of its wildcards
  - * indicates any string of characters (including none)
  - ? indicates any single character
- `ls` shows all files in the directory (except for some hidden files … check out ls -a)
- `ls *.py` shows all files in the directory that end in .py
- `ls D*` shows all files/directories that start with D (case-sensitive)
- Warning: be particularly careful using wildcards with `rm`! A good practice is to check which files you would delete with a command by first using `ls` in place of `rm`

More wildcard examples

If you had the following files in your directory:

- `PS1.txt`
- `PS1.py`
- `PS2.txt`
- `PS2.py`
- `PS3.txt`
- `PS3.py`
- `Lecture1.pptx`
- `Lecture1A.pptx`
- `Lecture1B.pptx`
- `Lecture2A.pptx`
- `Lecture2B.pptx`

How would you move all files ending in .txt to a new folder?

Which files would `rm Lecture1?.pptx` delete?

More wildcard examples - solutions

If you had the following files in your directory:

- `PS1.txt`
- `PS1.py`
- `PS2.txt`
- `PS2.py`
- `PS3.txt`
- `PS3.py`
- `Lecture1.pptx`
- `Lecture1A.pptx`
- `Lecture1B.pptx`
- `Lecture2A.pptx`
- `Lecture2B.pptx`

How would you move all files ending in .txt to a new folder?
`mv *.txt newfolder` (folder must already exist)

Which files would `rm Lecture1?.pptx` delete?
Lecture1A.pptx and Lecture 1B.pptx (not Lecture1.pptx)

A couple of handy shortcuts

- Tab-completion
  - if there's only one file/directory that starts with the set of characters you've typed, hitting Tab will complete it

```
$ ls gree<Tab>
$ ls greetings.txt
```

  - if there are multiple such files, hitting Tab twice will list them all
- Command history: use the up/down arrow keys to get your previously entered commands

Viewing/manipulating files

- UNIX is a text-based system – most files are flat (not fancy like Word) text files
- UNIX contains a lot of useful programs for working with text files
- UNIX programs read in files and write out to the standard out (and error) stream, unless redirected to a file
  - In general, they do not edit files in place

Print the beginning of the file

Print the top (head) of the file PS3_chr21.txt (by default, first 10 lines)

```
$ head PS3_chr21.txt
ctccaaagaaattgtagttttcttctggcttagaggtagatcatcttggt
ccaatcagactgaaatgccttgaggctagatttcagtctttgtggcagct
ggtgaatttctagtttgccttttcagctagggattagcttttaggggtc
ccaatgcctagggagatttctaggtcctctgttccttgctgacctccaat
tttgtctatccttttgctgagaggtctgcttaacttccttttagtcaggt
agctccattttatgctaagcttcttagttgctcaccttctgcagctaaag
aatcagaaaatgctgtgaaggaaaaacaaaatgaaattgcattgtttcta
ccggccctttatcaagccctggccaccatgatagtcatgaattccaattg
ttgtctatgcaggcctaccagatttctaacatctctgagctaccattttc
ttcttagctatctgctcagcaaatgtatccaaatgaaaggctgtggagaa
```

Print the first line in the file

```
$ head –n 1 PS3_chr21.txt
ctccaaagaaattgtagttttcttctggcttagaggtagatcatcttggt
```

Print the beginning of the file

Print the top (head) of the file PS3_chr21.txt (by default, first 10 lines)

```
$ head PS3_chr21.txt
ctccaaagaaattgtagttttcttctggcttagaggtagatcatcttggt
ccaatcagactgaaatgccttgaggctagatttcagtctttgtggcagct
ggtgaatttctagtttgccttttcagctagggattagcttttaggggtc
ccaatgcctagggagatttctaggtcctctgttccttgctgacctccaat
tttgtctatccttttgctgagaggtctgcttaacttccttttagtcaggt
agctccattttatgctaagcttcttagttgctcaccttctgcagctaaag
aatcagaaaatgctgtgaaggaaaaacaaaatgaaattgcattgtttcta
ccggccctttatcaagccctggccaccatgatagtcatgaattccaattg
ttgtctatgcaggcctaccagatttctaacatctctgagctaccattttc
ttcttagctatctgctcagcaaatgtatccaaatgaaaggctgtggagaa
```

Print the first line in the file

```
$ head -n 1 PS3_chr21.txt
ctccaaagaaattgtagttttcttctggcttagaggtagatcatcttggt
```

This is an option, specifying how many lines to print

Print the end of the file

Print the end (tail) of the file PS3_chr21.txt (by default, last 10 lines)

```
$ tail PS3_chr21.txt
NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN
NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN
NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN
NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN
NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN
NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN
NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN
NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN
NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN
NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN
```

Print the last line in the file

```
$ tail –n 1 PS3_chr21.txt
NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN
```

Print an entire file (or multiple, concatenated) to the screen

```
$ cat greetings.txt
hello

$ cat greetings.txt greetings.txt
hello
hello
```

Redirecting to standard in, standard out, and standard error

```
prog1 < file1
```

means feed file1 into the standard input of the program prog1

```
prog1 arg1 > file1
```

means run prog1 with argument arg1 and save the output to file1

```
prog1 arg1 | prog2
```

means run prog1 with argument arg1 and feed the output as the standard input to program prog2

Using the left arrow to replace an argument expecting a file with the output of a program

```
prog2 <(prog1 arg1)
```
is (mostly) equivalent to
```
prog1 arg1 > file1
prog2 file1
```

You can string these together!
```
prog3 <(prog1 arg1) <(prog2 arg2)
```
(mostly) equivalent to
```
prog1 arg1 > file1
prog2 arg2 > file2
prog2 file1 file2
```

Exercises

Create a new file twogreetings.txt that contains the contents of greetings.txt twice in a row.

Concatenate the first 10 lines of PS3_chr21.txt with the last 10 lines of PS3_chr21.txt and print to the screen.

Exercises - solutions

Create a new file twogreetings.txt that contains the contents of greetings.txt twice in a row.

```
$ cat greetings.txt greetings.txt > twogreetings.txt
```

Concatenate the first 10 lines of PS3_chr21.txt with the last 10 lines of PS3_chr21.txt and print to the screen.

```
$ cat <(head PS3_chr21.txt) <(tail PS3_chr21.txt)
```

(How many lines of Python would this take?)

How big is the file?

`wc` counts the number of lines, words, and characters (bytes) in a file

```
$ wc PS3_chr21.txt
 774374  774374 40267443 PS3_chr21.txt
```

Just print the number of lines

```
$ wc -l PS3_chr21.txt
774374
```

less: a better viewer for looking at big files

less works with files one screen at a time
Try `less PS3_chr21.txt`

You can search for strings in the file:
type: `/GATT` to search “GATT” and highlight all matches
then hit “n” to go to the next hit

Hit arrow keys to navigate
Hit Space to go a page down
Hit “q” to exit

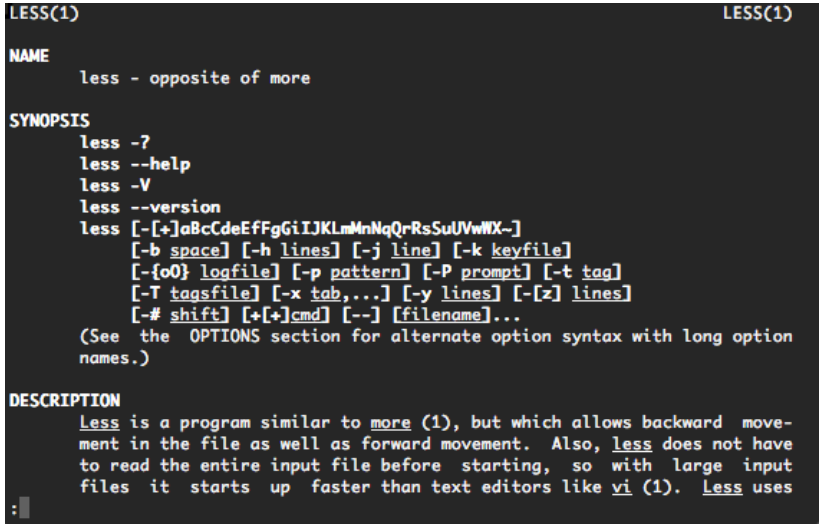
How do I remember all those options?

Every command has a manual page. Access it with the command **`man`**

```
$ man less
```

Read through manuals

using the **`less`** commands!



Working with big files - grep

Print all lines in PS3_chr21.txt that contain a string of interest, here GATT

```
$ grep GATT PS3_chr21.txt
```

Print all lines in PS3_chr21.txt that do NOT contain “N”

```
$ grep –v N PS3_chr21.txt
```

Some options:

-f: instead of just a string, take a file with a list of query sequences

-w: require the match to be a word (have whitespace on either side)

Working with big files - cut

We often work with tables, with columns separated by tabs (or spaces, commas, etc.)

Print the 3rd, 4th, 5th, and 9th columns (fields) of file1.txt

$cut –f 3-5,9 file1.txt

some options:

-d: specify delimiter - comma, space, tab (default)

-c: get characters rather than fields

How can we keep a record of these kinds of complex commands, and rerun them later?

- Shell scripts are programs that can be run by the UNIX interpreter, as if you had typed each line directly on the command-line.
- They can run other programs (e.g. Python programs), so they're useful for building complex programs (or analysis pipelines) that use programs other people have written (like BLAST)
- Like Python programs, they can take arguments, use loops and conditional statements, etc.
- They end in .sh and are executable pieces of text

Shell scripts

Suppose you had a Python program called hello-n.py (what does it do?)

```
import sys
for i in range(int(sys.argv[1])):
   print “Hello!”
```

And a shell script called five.sh

```
python hello.py 5
```

Then running five.sh would print “Hello!” five times by running hello-n.py with the argument 5.

How do we run a shell script?

We first have to make the script *executable*, with the command **`chmod`**

```
$ chmod +x five.sh
```

Then we can enter the name of the program, five.sh, preceded by "./" (strictly speaking, needs to be a path – e.g. could be myfolder/five.sh)

```
$ ./five.sh
Hello!
Hello!
Hello!
Hello!
Hello!
```