Programming For Biology 2025

programmingforbiology.org

Instructors

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Big Picture

Why?

Why is it important for **biologists** to learn to program?

You might already know the answer to this question since you are here.

We firmly believe that knowing how to program is just as essential as knowing how to run a gel or set up a PCR reaction. The data we now get from a single experiment can be overwhelming. This data often needs to be reformatted, filtered, and analyzed in unique ways. Programming allows you to perform these tasks in an **efficient** and **reproducible** way.

Helpful Tips

What are our tips for having a successful programming course?

- 1. Practice, practice, practice. Please spend as much time as possible actually coding.
- 2. Write only a line or two of code, then test it. If you write too many lines, it becomes more difficult to debug if there is an error.
- 3. Errors are not failures. Every error you get is a learning opportunity. Every single error you debug is a major success. Fixing errors is how you will cement what you have learned.
- 4. Don't spend too much time trying to figure out a problem. While it's a great learning experience to try to solve an issue on your own, it's not fun getting frustrated or spending a lot of time stuck. We are here to help you, so please ask us whenever you need help.
- 5. Lectures are important, but the practice is more important.
- 6. Review sessions are important, but practice is more important.
- 7. Our key goal is to slowly, but surely, teach you how to solve problems on your own.

Unix 1

Unix Overview

What is the Command Line?

Underlying the pretty Mac OSX Graphical User Interface (GUI) is the operating system (OS). It's based on BSD (Berkeley Standard Distribution), which is a version of Unix. Linux is pretty similar and also a very common OS in bioinformatics and you'll run into dialects by Red Hat, Ubuntu and others.

The command line gives you access to the internals of the OS, and is also a convenient way to write custom software and scripts.

Most bioinformatics tools are written to run on the command line and have no Graphical User Interface. In many cases, a command-line tool is more versatile than a graphical tool because you can easily combine command line tools into automated scripts that accomplish custom tasks sequentially without human intervention.

In this course, we will be writing Python scripts and running them from the command line.

The Unix OS is complicated and there are always more things to learn about it. We'll just cover enough to get you going and show you

The Basics

Logging into Your Workstation

Your workstation is an iMac. To log into it, provide your user name and password. Your username is 'pfb' and the password is 'pfb2025'

Bringing up the command line

To bring up the command line, use the Finder to navigate to *Applications->Utilities* and double-click on the *Terminal* application. This will bring up a window like the following:

```
student — -zsh — 80×24

Last login: Sun Oct 9 20:26:51 on console

student@INFO-01 ~ %
```

You can open many Terminal windows at once. This is often helpful. A common way to work is to open your code editor in one window, run the code from another and view data and files in another.

You will be using the Terminal application a lot, so I suggest that you drag its icon into the shortcuts bar at the bottom of your screen.

OK. I've Logged in. What Now?

The terminal window is running **shell** called "zsh". (Mac recently changed from bash) The shell is a loop that:

- 1. Prints a prompt
- 2. Reads a line of input from the keyboard
- 3. Parses the line into one or more commands.
- 4. Executes the commands (which usually print some output to the terminal)
- 5. Go back step 1.

There are many different shells with bizarre names like **bash**, **sh**, **csh**, **tcsh**, **ksh**, and **zsh**. The "sh" part means shell. Each shell has slightly different syntax and features. Your accounts are set up to use **zsh**. It's very similar to **bash** which is standard on linux systems.

Command Line Prompt

Most of bioinformatics is done by running command line software in a shell, so you should take some time to learn to use the shell effectively.

This is a command line prompt:

bush202>

This is another:

(~) 51%

This is another:

srobb@bush202 1:12PM>

What you see depends on how the system administrator has customized your login. You can customize it yourself (but we won't go into that here)

The prompt tells you the shell is ready for you to type a command. Most commands run almost instantly, but if you run a long command, the prompt will not reappear until it is finished and the system is ready to accept your next request.

Issuing Commands

You type in a command and press the <Enter> key to execute it. If the command has output, it will appear on the screen. Example:

```
(~) 53% ls -F
                          cool elegans.movies.txt man/
GNUstep/
                          docs/
                                                     mtv/
INBOX
TNBOX~
                          etc/
                                                     nsmail/
Mail@
                                                     pcod/
                          games/
News/
                          get_this_book.txt
                                                     projects/
                                                     public html/
axhome/
                          icod/
                          lib/
bin/
                                                     src/
build/
                          linux/
                                                     tmp/
ccod/
(~) 54%
```

The command here is 1s -F, which produces a listing of files and directories in the current directory (more on that later). Below its output, the command prompt appears again.

Some programs will take a long time to run. After you issue their command names, you won't recover the shell prompt until they're done. You can keep working in the meantime, either by launching a new shell (from Terminal's File menu), or running the command in the background by adding an ampersand at the end of the command like so

```
(~) 54% long_running_application &
(~) 55%
```

The command will now run in the background until it is finished. If it has any output, the output will be printed to the terminal window. You may wish to capture the output in a file (called redirection). We'll describe this later.

Command Line Editing

Most shells offer command line editing. Up until the moment you press <Enter>, you can go back over the command line and edit it using the keyboard. Here are the most useful keystrokes:

- Backspace: Delete the previous character and back up one.
- Left arrow, right arrow: Move the text insertion point (cursor) one character to the left or right.
- *control-a (^a)*: Move the cursor to the beginning of the line. (Mnemonic: A is first letter of alphabet)
- *control-e* (*^e*): Move the cursor to the end of the line. (Mnemonic: E for the End) (*^z* was already used to interrupt a command).
- *control-d* (^d): Delete the character currently under the cursor. D=Delete.
- *control-k (^k)*: Delete the entire line from the cursor to the end. k=kill. The line isn't actually deleted, but put into a temporary holding place called the "kill buffer". This is like cutting text

- *control-y* (*^y*): Paste the contents of the kill buffer onto the command line starting at the cursor. y=yank. This is like paste.
- *Up arrow, down arrow*: Move up and down in the command history. This lets you rerun previous commands, possibly modifying them before you do.

There are also some useful shell commands you can issue:

- history Show all the commands that you have issued recently, nicely numbered.
- !<number> Reissue an old command, based on its number (which you can get from history).
- !! Reissue the last command.
- !<partial command string>: Reissue the previous command that began with the indicated letters. For example, !1 (the letter el, not a number 1) would reissue the ls -F command from the earlier example.
- *control-r* (*^r*): enter string and search through history for commands that match it. This is really useful.

zsh offers automatic command completion and spelling correction. If you type part of a command and then the key, it will prompt you with all the possible completions of the command. For example:

```
(~) 51% fd<tab><tab>
(~) 51% fd

fd2ps    fdesign fdformat fdlist    fdmount fdmountd fdrawcmd fdumount
(~) 51%
```

If you hit tab after typing a command, but before pressing <Enter>, **zsh** will prompt you with a list of file names. This is because many commands operate on files.

Wildcards

You can use wildcards when referring to files. * stands for zero or more characters. ? stands for any single character. For example, to list all files with the extension ".txt", run 1s with the wildcard pattern "*.txt"

```
(~) 56% ls -F *.txt
final_exam_questions.txt genomics_problem.txt
genebridge.txt mapping_run.txt
```

There are several more advanced types of wildcard patterns that you can read about in the **zsh** manual page. For example, if you want to match files that begin with the characters "f" or "g" and end with ".txt", you can put both characters inside square brackets [fg] as part of the wildcard pattern. Here's an example

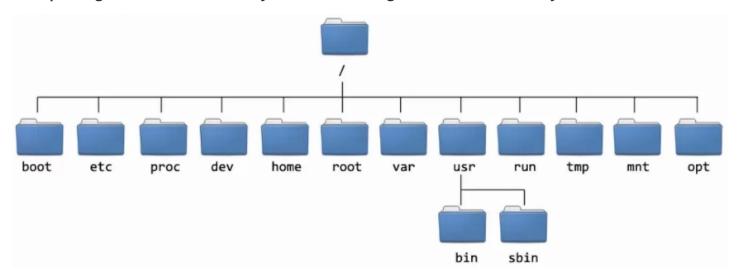
```
(~) 57% ls -F [fg]*.txt
final_exam_questions.txt genebridge.txt genomics_problem.txt
```

You can also search for a range of characters e.g. [a-e] or [1-5].

Directories and how they are organized

A computer comes with a place to store scripts, data, images, OS, and other files. It used to be floppy disks, then hard drives and these days it's often a solid state drive (SSD). Let's talk about how the storage is organized to help you find what you are working on. Directories or folders are created inside other directories. One directory is special. This is the **root directory** because it is not inside any other directories (it's written /). Files that go together are created inside a directory to keep them organized. This creates a structure that can be drawn like a branching tree. We find it clearer to turn it upside down to look like branching roots.

Example diagram of a Linux directory structure starting from the root directory



Home Sweet Home (your home directory ~)

When you first log in, you'll be in your personal directory (or folder), called the **home directory**. This directory has the same name as your login name, and on macOS is located inside the directory /users. (On Linux, it's typically in /home). If your username is dbrown, your home directory would (probably) be /users/dbrown. This is a filepath or a path. Unix if full of abbreviations to save on typing common things. The shell allows you to abbreviate it as ~username (where "username" is your user name or someone elses), or simply as ~. The weird character (called "tilde" or "twiddle") is at the upper left corner of your keyboard. The \$HOME variable is usually set to your home directory.

In your home directory, you have permission to save, delete, open and move files and other directories. In general, you can read but not write or modify files elsewhere in the system.

To see what is in your home directory, issue the command ls for list directory contents

```
Desktop Downloads Movies Pictures exptl.countdata.tsv
Documents Library Music Public notes.txt
```

You can modify the way a command works with switches. These are single letters typed after a dash -. The case is important. So ls -aF lists all (a) files including hidden filenames that start with a . with fancy formatting turned on (F) such that a / is added to the end of directory names.

```
./ .git/ Desktop/ Library/ Pictures/ notes.txt
../ .lesshst Documents/ Movies/ Public/
.CFUserTextEncoding .zshrc Downloads/ Music/ exptl.countdata.tsv
```

Don't go deleting the hidden files. Many of them are essential configuration files for commands and other programs. For example, the <code>lzshrc</code> file contains configuration information for the <code>zsh</code> shell. You can edit it (a bit later, when you know what you're doing) in order to change things like the command prompt and command search path.

Changing to a New Directory

You can move around from directory to directory using the cd command (for change directory). Give the name of the directory you want to move to, or give no name to move back to your home directory. Use the pwd command (print working directory) to see where you are (or, if configured, this information will appear in the prompt):

```
dbrown@WideIsLove ~ % cd Documents
dbrown@WideIsLove Documents % pwd
/Users/dbrown/Documents
dbrown@WideIsLove Documents % cd
dbrown@WideIsLove ~ % cd /
dbrown@WideIsLove / % ls -F
Applications/ System/ Volumes/ cores/ etc@ opt/ sbin/ usr/
Library/ Users/ bin/ dev/ home@ private/ tmp@ var@
```

Each directory contains two special hidden directories named . and ... The first, . refers always to the current directory. ... refers to the parent directory. This lets you move upward in the directory hierarchy like this:

```
(~/docs) 64% cd ..
```

and to do cool things like this:

```
(~/docs) 65% cd ../../dbrown/Documents
```

The latter command moves upward two levels, and then into a directory named <code>Documents</code> inside a directory called <code>dbrown</code>.

If you get lost, the pwd command prints out the full path to the current directory:

```
(~) 56% pwd
/Users/lstein
```

Absolute and relative paths

We can type a path in two ways, an absolute path always starts with a / character and reads from the root directory. A relative path starts with another character and reads from the directory you are currently in. Here are two ways to get to a directory called Music in the home directory of the user dbrown. From Documents we can go up a directory and down into Music (relative path) or go from / to Users to dbrown to Music (absolute path).

```
dbrown@WideIsLove Documents % cd ../Music
dbrown@WideIsLove Music % cd
dbrown@WideIsLove ~ % cd /Users/dbrown/Music
dbrown@WideIsLove Music %
```

Which of these paths would work for a different user as well as for dbrown?

Essential Unix Commands

With the exception of a few commands that are built directly into the shell (e.g. history, for, if), all Unix commands are actually executable programs. When you type the name of a command, the shell will search through all the directories listed in the \$PATH environment variable for an executable of the same name. If found, the shell will execute the command. Otherwise, it will give a "command not found" error.

Most commands live in /bin, /sbin, /usr/bin, /usr/sbin/ Or /usr/local/bin. You can use the which command to find a program's location, e.g., which ls may show /bin/ls.

Getting Information About Commands

The man command will give a brief synopsis of a command. Let's get information about the command we

Finding Out What Commands are on Your Computer

The apropos command will search for commands matching a keyword or phrase. Here's an example that looks for commands related to 'column'

On some systems, you can also use tldr (too long didn't read). On macOS, you can use Homebrew (https://breww.sh/) brew install tldr to see help documentation:

```
$ tldr wc

wc

Count lines, words, or bytes.
More information: <https://keith.github.io/xcode-man-pages/wc.l.html>.

- Count lines in file:
    wc -l path/to/file

- Count words in file:
    wc -w path/to/file

- Count characters (bytes) in file:
    wc -c path/to/file

- Count characters in file (taking multi-byte character sets into account):
    wc -m path/to/file

- Use `stdin` to count lines, words and characters (bytes) in that order:
    find . | wc
```

Arguments and Command Line Switches

Many commands take arguments. Arguments are often the names of one or more files to operate on. Most commands also take command line "switches" or "options", which fine-tune what the command does. Some commands recognize "short switches" that consist of a minus sign — followed by one or more single characters, while others recognize "long switches" consisting of two minus signs — followed by a whole word.

The wc (word count) program is an example of a command that recognizes both long and short options. You can pass it the single -c, -w and/or -1 options (or combinations of them) to count the characters, words, and lines in a text file, respectively. Or you can use the longer but more readable --chars, --words or --lines options. Both these examples count the number of characters and lines in the text file /var/log/messages:

```
(~) 102% wc -c -l /var/log/messages
     23     941 /var/log/messages
(~) 103% wc --chars --lines /var/log/messages
     23     941 /var/log/messages
```

You can cluster one-character switches by concatenating them together, as shown in this example:

```
(~) 104% wc -cl /var/log/messages
23 941 /var/log/messages
```

Many commands will give a brief usage summary when you call them with the -h or --help switch.

Spaces, tabs and newline Characters

The shell uses spaces to separate arguments. If you want to embed a space (see below for other whitespace or non-printing characters like a tab or newline etc) in an argument, put single quotes around it. For example:

```
mail -s 'An important message' 'Bob Ghost <bob@ghost.org>'
```

This will send an e-mail to the fictitious person Bob Ghost. The switch takes an argument, which is the subject line for the e-mail. Because the desired subject contains spaces, it has to have quotes around it. Likewise, my name and e-mail address, which contain embedded spaces, must also be quoted in this way.

Certain special non-printing characters have escape codes associated with them:

Escape Code	Description
\n	new line character
\t	tab character
\r	carriage return character
\a	bell character (ding! ding!)
\NNN	the character whose octal ASCII code is NNN e.g. printf '\101' prints 'A' <u>ASCII code tables</u>

Useful Commands

Here are some commands that are used extremely frequently. Use man to learn more about them. Some of these commands may be useful for solving the problem set;-)

Manipulating Directories

Command	Description
ls	Directory listing. Most frequently used as 1s -F (decorated listing), 1s -1 (long listing), 1s -a (list all files).
mv	Rename or move a file or directory.
ср	Copy a file. cp -r (recursively) to copy directories
rm	Remove (delete) a file. rm -rf olddata.tsv
mkdir	Make a directory
ln	Create a symbolic or hard link. ln -s makes a symbolic (sym) link
chmod	Change the permissions of a file or directory.

Reading files

Command	Description
cat	Concatenate program. Can be used to concatenate multiple files together into a single file, or, much more frequently, to view the contents of a file or files in the terminal.
more	Scroll through a file page by page. Very useful when viewing large files. Works even with files that are too big to be opened by a text editor.
less	A version of more with more features.
head	View the first few lines of a file. You can control how many lines to view head -3 prints the first three lines.
tail	View the end of a file. You can control how many lines to view. You can also use tail -f to view a file that you are writing to.

Analyzing, processing files

Command	Description
WC	Count words, lines and/or characters in one or more files.
sort	Sort the lines in a file alphabetically or numerically (-g or -n) in reverse order (-r).
uniq	Remove duplicated lines in a file.
cut	Remove columns from each line of a file or files.
grep	Filter a file for lines matching a specified pattern. Can also be reversed to print out lines that don't match the specified pattern.
gzip (gunzip)	Compress (uncompress) a file.
tar	Archive or unarchive an entire directory into a single file.

Editing files

Command	Description
tr	Substitute one character for another. Also useful for deleting characters.
nano	Very basic and easy to use text editor
emacs	Run the Emacs text editor (good for experts).
vi	Run the vi text editor (confusing even for experts).
echo	print text to the screen. E.g. echo 'Hello World!'

Connecting to other computers

Command	Description
ssh	A secure (encrypted) way to log into machines.
scp	A secure way to copy (cp) files to and from remote machines.

Standard I/O and Redirection

Unix commands print output to the terminal (screen) for you to see, and accept input from the keyboard (that is, from *you*!)

Every Unix program starts out with three connections to the outside world. These connections are called "streams", because they act like a stream of information (metaphorically speaking):

Stream Name	Description
standard input	This is initially attached to the keyboard. When the program reads from standard input, it reads whatever text you type in.
standard output	This stream is initially attached to the terminal. Anything the program prints to this stream appears in your terminal window.
standard error	This stream is also initially attached to the terminal. It is a separate stream intended for printing error messages.

The word "initially" might lead you to think that standard input, output, and error can somehow be detached from their starting places and reattached somewhere else. And you'd be right. You can attach one or more of these three streams to a file, a device, or even to another program. This sounds esoteric, but it is actually very useful.

A Simple Example

The wc program counts lines, characters, and words in data sent to its standard input. You can use it interactively like this:

In this example, I ran the wo program. It waited for me to type in a little poem. When I was done, I typed the END-OF-FILE character, control-d (^d for short). wo then printed out three numbers indicating the number of lines (6), words (20), and characters (107) in the input.

More often, you'll want to count the number of lines in a big file and save the results in another file. We'll see this in two steps. First, we send the output of we to stdout. In the second command, we *redirect* stdout from we to a file with the > symbol.

```
(~) 63% wc big_file.fasta
2943 2998 419272
(~) 64% wc big_file.fasta > count.txt
```

Now if you cat the file *count.txt*, you'll see that the data has been recorded.

```
(~) 65% cat count.txt
2943 2998 419272
```

Redirection Meta-Characters

Here's the complete list of redirection commands for zsh:

Redirect command	Description
< myfile.txt	Redirect the contents of the file to standard input
> myfile.txt	Redirect standard output to file
>> logfile.txt	Append standard output to the end of the file
1> myfile.txt	Redirect just standard output to file (same as above)
2> myfile.txt	Redirect just standard error to file
&> myfile.txt	Redirect both stdout and stderr to file

These can be combined. For example, this command redirects standard input from the file named <code>/etc/passwd</code>, writes its results into the file <code>search.out</code>, and writes its error messages (if any) into a file named <code>search.err</code>. What does it do? It searches the password file for a user named "root" and returns all lines that refer to that user.

```
(~) 66% grep root /etc/passwd > search.out 2> search.err
```

For instance, the find command will look for files/directories/links.

When you try to look in a directory where you don't have permissions, you will get an error.

So, if I look from // for a file, I will get lots of error messages for directories that I can't read and that I'd like to ignore.

I can redirect to a file:

```
find / -name passwd.txt 2>err
```

That might actually produce a very large *err* file, and I don't actually care about the errors. Instead, I can redirect them to a special filehandle called <code>/dev/null</code>, which throws them away entirely:

```
find / -name passwd.txt 2>/dev/null
```

Filters, Filenames, and Standard Input

Many Unix commands act as filters, taking data from a file or standard input, transforming the data, and writing the results to standard output. Most filters are designed so that if they are called with one or more filenames on the command line, they will use those files as input. Otherwise they will act on standard input. For example, these two commands are equivalent:

```
(~) 67% grep 'gatttgc' big_file.fasta
(~) 68% grep 'gatttgc' < big_file.fasta</pre>
```

Both commands use the <code>grep</code> command to search for the string "gatttgc" in the file <code>big_file.fasta</code>. The first command is explicitly given the name of the file on the command line; the second one searches standard input, which is redirected from the file to stdin of <code>grep</code>.

Sometimes you want a filter to act on a series of files, one of which happens to be standard input. Many commands let you use — on the command line as an alias for standard input. Example:

```
(~) 70% grep 'gatttgc' big_file.fasta bigger_file.fasta -
```

This example searches for "gatttgc" in three places. First it looks in file big_file.fasta, then in bigger file.fasta, and lastly in standard input (which, since it isn't redirected, will come from the keyboard).

Standard I/O and Pipes

The coolest thing about the Unix shell is its ability to chain commands together into mini pipelines. Here's an example:

```
(~) 65% grep gatttgc big_file.fasta | wc -l 22
```

There are two commands here. grep searches a file or standard input for lines containing a particular string. Lines which contain the string are printed to standard output. wc -1 is the familiar word count program, which counts words, lines, and characters in a file or standard input. The -1 command line option instructs wc to print out just the line count. The | character, which is known as a "pipe", connects the two commands together so that the standard output of grep becomes the standard input of wc. Think of pipes connecting streams of flowing data.

What does this pipe do? It prints out the number of lines in which the string "gatttgc" appears in the file big_file.fasta.

More Pipe Idioms

Pipes are very powerful. Here are some common command line idioms.

Count the Number of Times a Pattern does NOT Appear in a File

The example at the top of this section showed you how to count the number of lines in which a particular string pattern appears in a file. What if you want to count the number of lines in which a pattern does **not** appear?

Simple. Reverse the test with the -v switch:

```
(~) 65% grep -v gatttgc big_file.fasta | wc -l
2921
```

Uniquify Lines in a File

If you have a long list of names in a text file, and you want to weed out the duplicates:

```
(~) 66% sort long_file.txt | uniq > unique.out
```

This works by sorting all the lines alphabetically and piping the result to the uniq program, which removes duplicate lines that occur one after another. That's why you need to sort first. The output is placed in a file named unique.out.

Concatenate Several Lists and Remove Duplicates

If you have several lists that might contain repeated entries among them, you can combine them into a single unique list by concatenating them together, then sorting and uniquifying them as before:

```
(~) 67% cat file1 file2 file3 file4 | sort | uniq
```

Count Unique Lines in a File

If you just want to know how many unique lines there are in the file, add a wc to the end of the pipe:

```
(~) 68% sort long_file.txt | uniq | wc -l
```

Page Through a Really Long Directory Listing

Pipe the output of ls to the more program, which shows a page at a time. If you have it, the less program is even better:

```
(~) 69% ls -1 | more
```

Monitor a Growing File for a Pattern

Pipe the output of tail -f (which monitors a growing file and prints out the new lines) to grep. For example, this will monitor the /var/log/syslog file for the appearance of e-mails addressed to 'mzhang':

```
(~) 70% tail -f /var/log/syslog | grep dbrown
```

Permissions

Often several different users work on the same computer. To help them work without disrupting each other, a set of permissions is attached to files and directories. These permissions allow or prevent reading, writing or executing a file. There are permissions for the user, group and other. You can view permissions with an ls -1 command.

```
dbrown@WideIsLove ~ % ls -1
total 16
drwx-----+ 3 dbrown staff 96 Oct 15 14:09 Desktop
drwx-----+ 3 dbrown staff 96 Oct 15 14:09 Documents
...
-rw-r--r-- 1 dbrown staff 9 Oct 15 14:12 exptl.countdata.tsv
-rw-r--r-- 1 dbrown staff 20 Oct 15 14:11 notes.txt
-rw-r--r-- 1 dbrown staff 23 Oct 16 13:37 print.py
```

The series of characters at the beginning of the line describe the permissions. The first character is

- d: directory
- 1: link
- -: file

Then there are three triples of characters (- means not allowed, a letter means you have read, write, exectute permission). x for a directory means you can access the directory

```
rwx
---
```

for user, group, other, like so

```
UsrGrpOth
drwxrwxrwx+
```

Users can belong to groups (who collaborate) and other is everyone else.

The chmod command (change mode) alters permissions. You add permissions with + and take them away with -

You can alter permissions for user u, group g, other o or all a

To make the print.py file executable (i.e. runnable like a program) you write

```
dbrown@WideIsLove ~ % ls -l print.py
-rw-r--r- 1 dbrown staff 23 Oct 16 13:37 print.py
dbrown@WideIsLove ~ % chmod a+x print.py
dbrown@WideIsLove ~ % ls -l print.py
-rwxr-xr-x 1 dbrown staff 23 Oct 16 13:37 print.py
dbrown@WideIsLove ~ %
```

To stop others from reading or executing the Python script, you write

```
dbrown@WideIsLove ~ % chmod o-rx print.py
dbrown@WideIsLove ~ % ls -1 print.py
-rwxr-x--- 1 dbrown staff 23 Oct 16 13:37 print.py
```

To convert a script into an exectuable program, you'll need to make it executable with chmod and then add the current directory to your \$PATH (we'll see how to do this later) for you, you can type the command in your terminal window.

NOTE: You must use the \$ when you want the shell to interpolate the value of the variable called PATH, but you do *not* use it when setting the value.

You can use echo to inspect your \$PATH variable, which is a colon-separated list of directories where the OS will search for executables.

You can pipe this output to tr to change the colon (:) to newlines to make it readable:

```
$ echo $PATH | tr : '\n'
/Library/Frameworks/Python.framework/Versions/3.12/bin
/opt/homebrew/bin
/opt/homebrew/sbin
/usr/local/bin
/usr/bin
/bin
/usr/sbin
/sbin
/Users/kyclark/.local/bin
```

More Unix

Here are a few more advanced Unix commands that are very useful, and when you have time you should investigate further. We list the page numbers for the Linux Phrasebook Second Edition by Scott Granneman or links to online tutorials.

- awk (Linux Phrasebook p.194-198) online tutorial
- sed (Linux Phrasebook p.188-194) online tutorial
- perl one-liners online tutoral
- for loops online tutorial

Link to Unix 1 Problem Set

Unix 2

Text Editors

It is often necessary to create and write to a file while using the terminal. This makes it essential to use a terminal text editor. There are many text editors out there. Some of our favorite are Emacs and vim. We are going to start you out with a simple text editor called vi

Introduction to vi

What is vi?

vi is a command line text editor. vi is included in every Linux installation. You don't have to install it, ever.

What is a command line text editor?

A command line text editor is an text editor that you use from the command line. In most command line text editors, don't expect to be able to point and click. You will need to naviage with keyboard key strokes. The two most popular text editors are **vi** and **emacs**. You are free to use either, but we will start with **vi** since the keystrokes are less complex than in **emacs**.

Why do I care about command line text editors?

If you are logged into a remote machine, a command line text editor is the fastest, easiest, most efficient way to write text files.

Getting Started with vi

Opening a file

On the command line, type vi followed by a file name.

```
srobb% vi <file>
```

Let's try it:

```
srobb% vi first_vi_file.txt
```

You will see this in your terminal.

```
"first_vi_file.txt" [New File]
```

Notice the file name at the bottom.

If you **do not** include a file name you will see something similar to this:

```
VIM - Vi IMproved

version 8.0.1283

by Bram Moolenaar et al.

Vim is open source and freely distributable

Become a registered Vim user!

type :help register<Enter> for information

type :q<Enter> to exit

type :help<Enter> or <Fl> for on-line help
type :help version8<Enter> for version info
```

Read what the message says and type :q<Enter> to Quit or exit.

vi has two modes.

- 1. Insert Mode
- 2. Command Mode

Insert Mode is for typing your file contents. All keyboard strokes will be interpreted as characters you want to see in your file.

Command Mode is for using commands. All keyboard stokes will be interpreted as commands and **not** as part of your file. Common commands are for deleting, copying, searching, replacing, and saving.

Creating, Writing, And Saving a File Walk through

Create

From the command line open a new file by typing

```
vi first_vi_file.txt
```

Write

Start typing content. To do this we need to enter **Insert Mode**.

To do this type i.

Your vi session will now look like this:

Notice the INSERT at the bottom of the screen.

Start typing your file contents. Remember that all keystrokes are ones you want to see in your file and that your mouse will not work.

Save

Now that the file contains some content let's enter **Command Mode** so that we can save our file.

- 1. Press the <ESC> key to enter **Command Mode**.
- 2. type w (colon followed by a w) to **Save (Write)**

If you want to type some more content, enter **Insert Mode** (i).

If, instead you want to exit, since you are already in Command Mode you can use the quit keystrokes [q]

Common Activities and vi Commands

Enter into **Command Mode** for all commands. If you are unsure that you are in **command mode**, just press the <esc> key. It will not hurt if you are already in **Command Mode**

Saving and Exiting

Remember to enter into **Command Mode** with <esc> key

key stroke	meaning
:wq	Save (W rite) and Q uit
:q!	Q uit without Saving!!!
:w	Save (W rite) Only

Most commands within vi are executed as soon as you press a sequence of keys. Any command beginning with a colon (:) requires you to hit <enter> to complete the command.

Getting around

Remember to enter into **Command Mode** with <esc> key

key stroke	meaning
Arrow keys	move the cursor around
j, k, h, 1	move the cursor down, up, left and right (similar to the arrow keys)
o (zero)	move cursor to beginning of current line
ˆ (caret)	move cursor to beginning of current line
\$	move cursor to end of the current line
:n	move to the nth line in the file
nG	move to the n th line (eg 5G moves to 5th line)
G	move to the last line
W	move to the beginning of the next word
nw	move forward n word (eg 2w moves two words forwards)
b	move to the beginning of the previous word
nb	move back n word

Deleting content

Remember to enter into **Command Mode** with <esc> key

key stroke	meaning
x	delete a single character
nx	delete n characters (eg 5x deletes five characters)
dd	delete the current line
dn	d followed by a movement command. Delete to where the movement command would have taken you. (eg d5w means delete 5 words)
уу	"yank" (copy) a line
p	"paste" a line that was deleted or yanked

Undoing

Remember to enter into **Command Mode** with <esc> key

key stroke	meaning
u	Undo the last action (you may keep pressing u to keep undoing)
U	(Note: capital)** - Undo all changes to the current line

Other Useful Tips

key stroke	meaning
:set number	display line numbers
:set nonumber	turn off line numbers
:/[search text]	find [this text] in your file
n	go to next occurance of your search result

Mug of vi



FILE COMMANDS

vi filename(s) edit a file or files vi -r filename retrieve saved file after crash ZZ, :wq, :x save and exit quit; quit without saving :q, :q! :w, :w fn save file, save file as fn edit filename :e filename :r filename insert filename :sh drop to shell : 1 cmd run command cmd :r !cmd execute cmd and insert output Imovement cmd pipe lines in movement through cmd

SEARCH AND REPLACE

/txt, ?txt find txt forward or backward find next line that starts with txt

n, N repeat last search forward, backward

R replace text from current character

DELETING/INSERTING TEXT

dw, dd, x delete word, line, character ndd, nx delete n lines, n characters delete character forward, backward x, X delete to end of line D. ds dmotion delete from cursor to motion (\$, 0, etc.) indent, outdent line :>, :< replace text with blank line 0, 0 insert new line below, above current line undo last change repeat last change

CUT/COPY/PASTE

nyy, nY copy n lines
yw, yy copy word, line
p, P paste text after, before cursor
a, i insert text after, before cursor
A, I insert text end, beginning of line

MOVING AROUND

nG move to line n left, right, up, down one character h, 1, k, j nb, nw left or right n words CTRL-B, F backward, forward one screen CTRL-U, D up, down one screen \$, G go to end of line, end of file 0 go to beginning of line (zero)), (}, { move to next, previous sentence move to next, previous paragraph w, b move forward, back one word go to end of current or next word e

WICKED COOL STUFF

change case

xp transpose characters

combine current line with next

mp create a mark called p

p return to p

d'x, y'x del, copy text from mark to cursor

indent n lines



Visual Studio Code

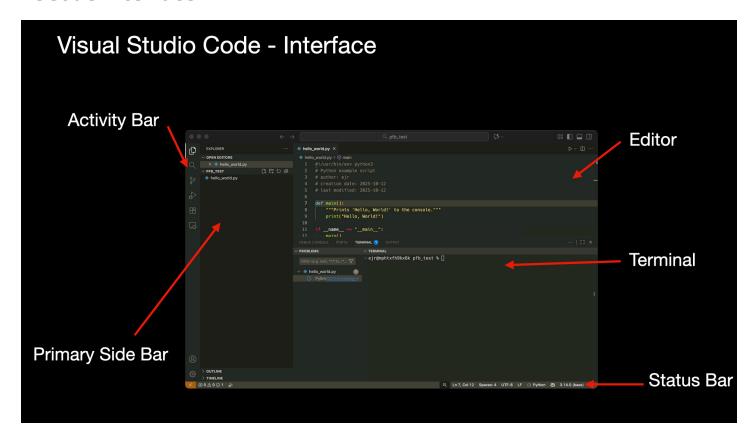
What is VSCode

Where vi is a text only editor with keystroke only commands, VSCode allows for on screen mouse interactions. vi is installed by default on all linux servers, VSCode needs to be installed and set up on each of your computers and can be used to edit files remotely.

- IDE (Integrated Development Environment).
- Owned by Microsoft, but "free".

- Can be used for many programming languages.
- Many official and community extensions.
 - Integration with ssh.
 - Integration with git.
 - Integration with AI coding tools.

VSCode interface



How Does VSCode Work

- Works like Word or other Text editor (copy, paste, undo etc).
- Formats and colors code in a language specific way.
- Saves text files that can be opened as any other text file.

Documentation

Documentation

Git for Beginners

Git is a tool for managing files and versions of files. It is a *Version Control System*. It allows you to keep track of changes. You are going to be using Git to manage your course work and keep your copy of the lecture notes and files up to date. Git can help you do very complex task with files. We are going to keep it simple.

The Big Picture

A Version Control System is good for Collaborations, Storing Versions, Restoring Previous Versions, and Managing Backups.

Collaboration

Using a Version Control System makes it possible to edit a document with others without the fear of overwriting someone's changes, even if more than one person is working on the same part of the document. All the changes can be merged into one document. These documents are all stored one place.

Storing Versions

A Version Control System allows you to save versions of your files and to attach notes to each version. Each save will contain information about the lines that were added or altered.

Restoring Previous Versions

Since you are keeping track of versions, it is possible to revert all the files in a project or just one file to a previous version.

Backup

A Version Control System makes it so that you work locally and sync your work remotely. This means you will have a copy of your project on your computer and the Version Control System Server you are using.

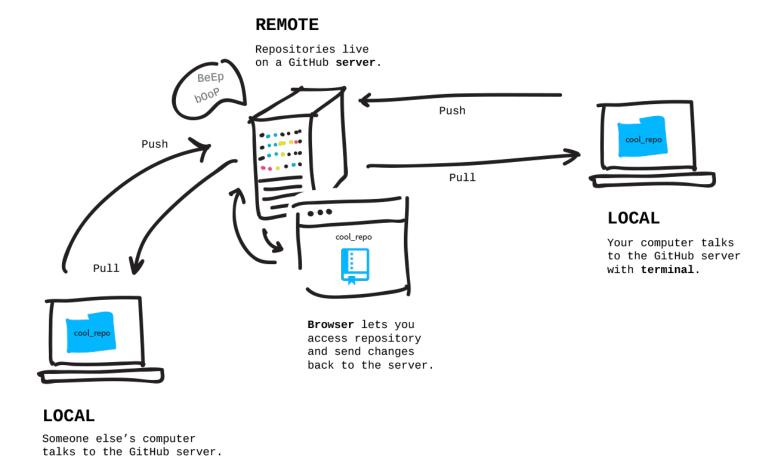
The Details

git is the Version Control System we will be using for tracking changes in our files.

GitHub is the Version Control System Server we will be using. They provide free account for all public projects.

The Basics

Usually you have a local copy of your project/repository and a remote copy. The **local** repository is stored on your computer and the **remote** is on a online service like GitHub.



You can use a web browser to interact with the remote server (gitHub) and the terminal to interact with the local repository.

SSH Keys

SSH Keys are needed to write to a personal repository. Lets set this up first

Github requires authentication with the use of ssh keys. Essentially, our github repos are LOCKED, we need a KEY to write to them. We will generate the a key (private) and a lock (public). Then We tell github about our lock and keep our key to ourselves.

Here is a great GitHub Tutoral.

Adding a new SSH key to your GitHub account

Here is a summary of the steps:

Generating a new SSH KEY

Make your key and add your email address.

ssh-keygen -t ed25519 -C "your_email@example.com"

Save your key and lock in your ssh directory. It will prompt you for a path, the default is usually a good place to store your SSH key files

```
> Enter a file in which to save the key (/Users/YOU/.ssh/id_ALGORITHM: [Press enter]
```

Next, pick a passphrase, something easy. It is a password, so that not just anyone on your physical computer can access your GitHub account. You CAN leave it blank.

```
> Enter passphrase (empty for no passphrase): [Type a passphrase]
> Enter same passphrase again: [Type passphrase again]
```

Adding your SSH key to the ssh-agent

```
$ eval "$(ssh-agent -s)"
> Agent pid 59566
```

First, check to see if your ~/.ssh/config file exists in the default location.

```
$ open ~/.ssh/config
> The file /Users/YOU/.ssh/config does not exist.
```

If the file doesn't exist, create the file.

```
touch ~/.ssh/config
```

Open your ~/.ssh/config file using vi and add the following content:

```
Host *

IdentityFile ~/.ssh/id_ed25519
```

Adding a new SSH key to your GitHub account

Print the contents of your PUBLIC ssh key file (our lock) and paste them into your github account

```
cat ~/.ssh/id_ed25519.pub
```

Paste into your GitHub account

1. Go to Settings



Signed in as octocat



Set status

Your profile

Your repositories

Your organizations

Your projects

Your stars

Your gists

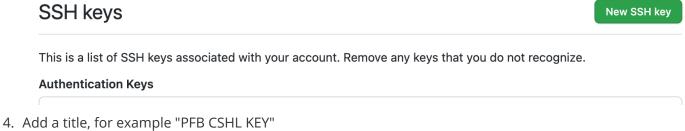
Feature preview

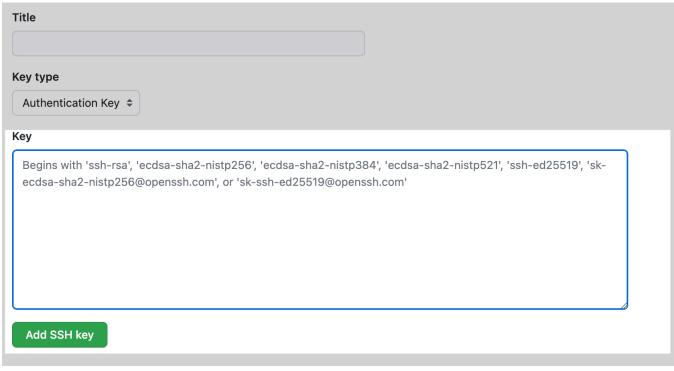
Help

Settings

Sign out

- 2. In the "Access" section of the sidebar, click "SSH and GPG keys".
- 3. Click New SSH key or Add SSH key. https://github.com/settings/keys





5. Click "Add SSH Key"



Do this to ensure that you do not encounter errors by attemping to add files that are too large

cd ~/PFB_problemsets/.git/hooks/ curl -0 https://raw.githubusercontent.com/prog4biol/pfb2025/master/setup/pre-commit

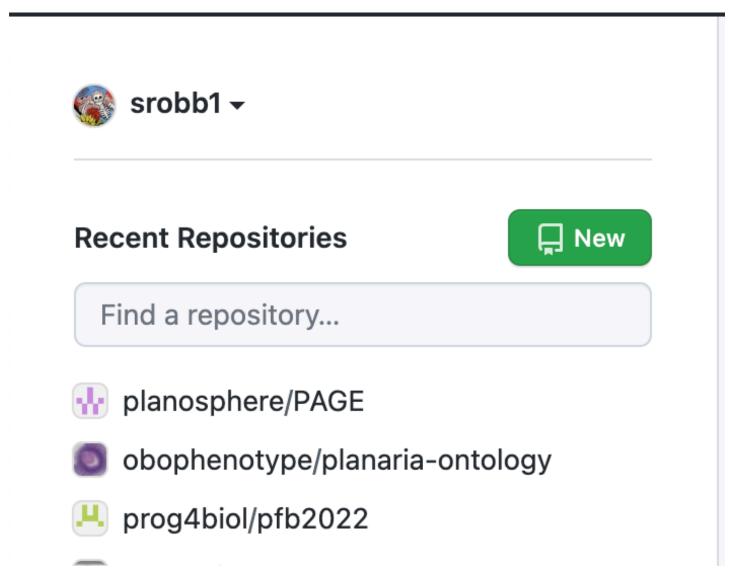
Creating a new repository

A repository is a project that contains all of the project files, and stores each file's revision history. Repositories can have multiple collaborators. Repositories usually have two components, one **remote** and one **local**.

Use Steps 1 and 2 below to create the **remote repository**.

Then, use Step 3 to create your **local repository** and **link it** to the **remote repository**.

1. Navigate to GitHub --> Create Account / Log In --> Click on your icon --> Click "Your Repositories" --> Click 'New'



2. Add a name (i.e., PFB_problemsets) and a description (i.e., Solutions for PFB Problem Sets) and click "Create Repository"

Create a new repository

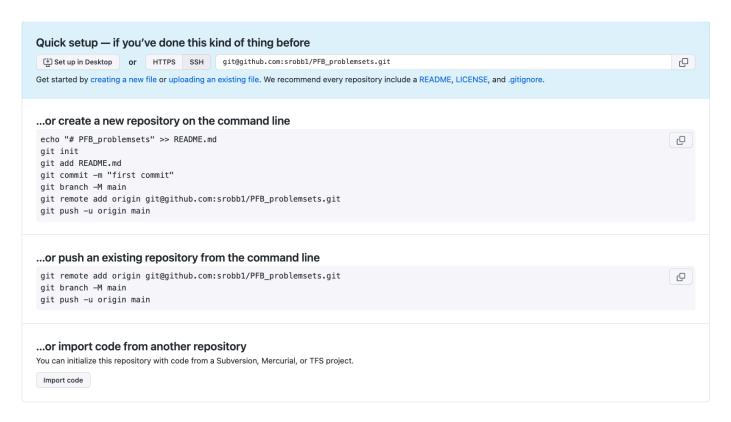
A repository contains all project files, including the revision history. Already have a project repository elsewhere? Import a repository.

Repository template Start your repository with a template repository's contents. No template -Owner * Repository name * srobb1 ▼ PFB_problemsets Great repository names are short and memorable. Need inspiration? How about effective-octo-waffle? Description (optional) Solutions for PFB Problemsets **Public** Anyone on the internet can see this repository. You choose who can commit. **Private** You choose who can see and commit to this repository. Initialize this repository with: Skip this step if you're importing an existing repository. □ Add a README file This is where you can write a long description for your project. Learn more. Add .gitignore Choose which files not to track from a list of templates. Learn more. .gitignore template: None -Choose a license A license tells others what they can and can't do with your code. Learn more. License: None ▼ i You are creating a public repository in your personal account. Create repository

3. Create a directory on your computer in your home directory

```
cd ~
mkdir PFB_problemsets
```

- 4. Make sure to select SSH in the GitHub new repository window.
- 5. Now follow the directions provided by github.



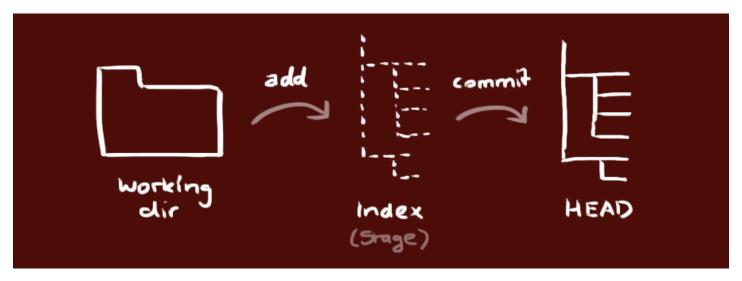
6. If you did not select SSH and use the SSH URL, do a git remote set-url 'git@blalabbalalal' with the SSH URL

The new local repository consists of three "trees" maintained by git. The first one is your "Working Directory" which holds the actual files. the second one is the "Index" which acts as a staging area and finally the "HEAD" which points to the last commit you've made.

Every git local repository has three main elements called trees:

- 1. The Working Directory contains your files
- 2. The *Index* is the staging area
- 3. The *HEAD* points to the last commit you made.

There are a few new words here. We will explain them as we go



Command Review

command	description
git init	Creates your new local repository with the three trees (local machine)
git remote add remote-	Links your local repository to a remote repository that is often named origin and is found at the given URL
git add filename	Propose changes and add file(s) with changes to the index or staging area (local machine)
git commit -m 'message'	Confirm or commit that you really want to add your changes to the HEAD (local machine)
git push -u remote-name remote-branch	Upload your committed changes in the HEAD to the specified remote repository to the specified branch

Using your new repository

You created a **local** repository above. It is linked to a **remote**. The <code>git remote</code> add command connects your local to the remote. Before this command the local will not know anything about your remote and vice versa.

Review:

- 1. You created a new remote repository on the github website.
- 2. You followed all the instructions given to you on github when you created your repository.

Now we will add some files to your new repository:

- 1. Change directory to your local repository
- 2. Create a new file with vi: vi git_exercises.txt
- 3. Add a line of text to the new file.

- 4. Save :w and Exit :q
- 5. (Add) Stage your changes. git add git_exercises.txt
- 6. (Commit) Become sure you want your changes. git commit -m 'added a line of text'
- 7. (Push) Sync/Upload your changes to the **remote** repository. git push origin main

That is all there is to it! There are more complicated things you can do, but we won't get into those. You will know when you are ready to learn more about git when you figure out there is something you want to do but don't know how. There are thousands of online tutorials for you to search and follow.

Keeping track of differences between local and remote repositories

If you are ever wondering what do you need to add to your remote repository use the <code>git status</code> command. This will provide you with a list of files that have been modified, deleted, and those that are untracked.

Untracked files are those that have never been added to the staging area with <code>git add</code>

command	description
git status	To see a list of files that have been modified, deleted, and those that are untracked

Deleting and moving files

command	description
git rm	Remove files from the index, or from the working tree and from the index
git mv	Move or rename a file, a directory, or a symlink

these two commands will update your index as well as change your local files. If you use just rm or mv you will have to update the index with add/commit.

Get a copy of file on your remote

Sometimes you really really mess up a file, or you delete it by mistake. You have a small heart attack then you remember that you have a good copy in your remote github repo. How do you get it in your local repo?

```
git checkout <filename>
```

Whew, what a life saver!

Tips

- 1. Adding files over 50M will break your git repo. Don't add large files. Don't blindly use git add -A when there might be large files present. You will be very sad if you do.
- 2. Don't clone a git repository into another git repository. This makes git really unhappy.

3. Don't be afraid to ask your questions on Google. git can be complicated and a lot of people ask a lot of questions that get answered in online forums, or GitHub will have a tutorial

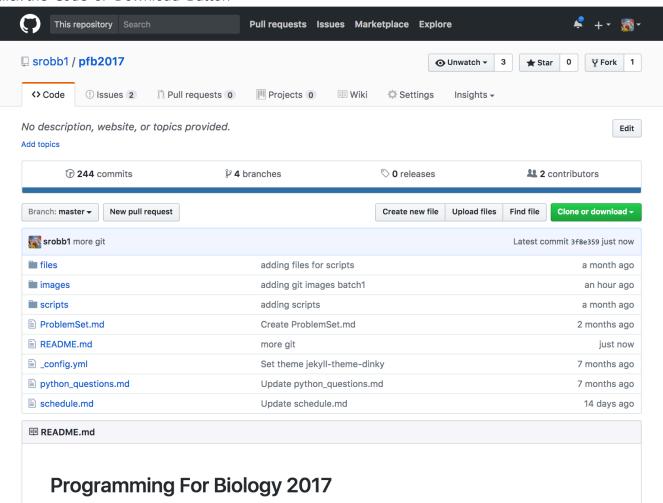
Cloning a Repository

Sometimes you want to download and use someone else's repository. This is different from above where we created our own repository. This is just a copy of someone else's repository

Let's clone the course material.

Let's do it!

- 1. Go to our PFB GitHub Repository
- 2. Click the 'Code' or 'Download' Button



- 3. Copy the URL: git@github.com:prog4biol/pfb2025.git
- 4. *Clone* the repository to your local machine. On the command line use this command: git clone git@github.com:prog4biol/pfb2025.git

Now you have a copy of the course material on your computer!

Bringing Changes in from the Remote Repository to your Local Repository

If changes are made to any of these files in the online, remote repository, and you want to update your local copy, you can *pull* the changes.

git pull

command	description
git pull	To get changes from the remote into your local copy

Links to slightly less basic topics

You will KNOW if you need to use these features of git.

- 1. <u>View Commit History</u>
- 2. Resolving Merge Conflicts
- 3. <u>Undoing Previous Commits</u>

Link To Unix 2 Problem Set