

\$ Welcome to the Matrix Unix/Linux I

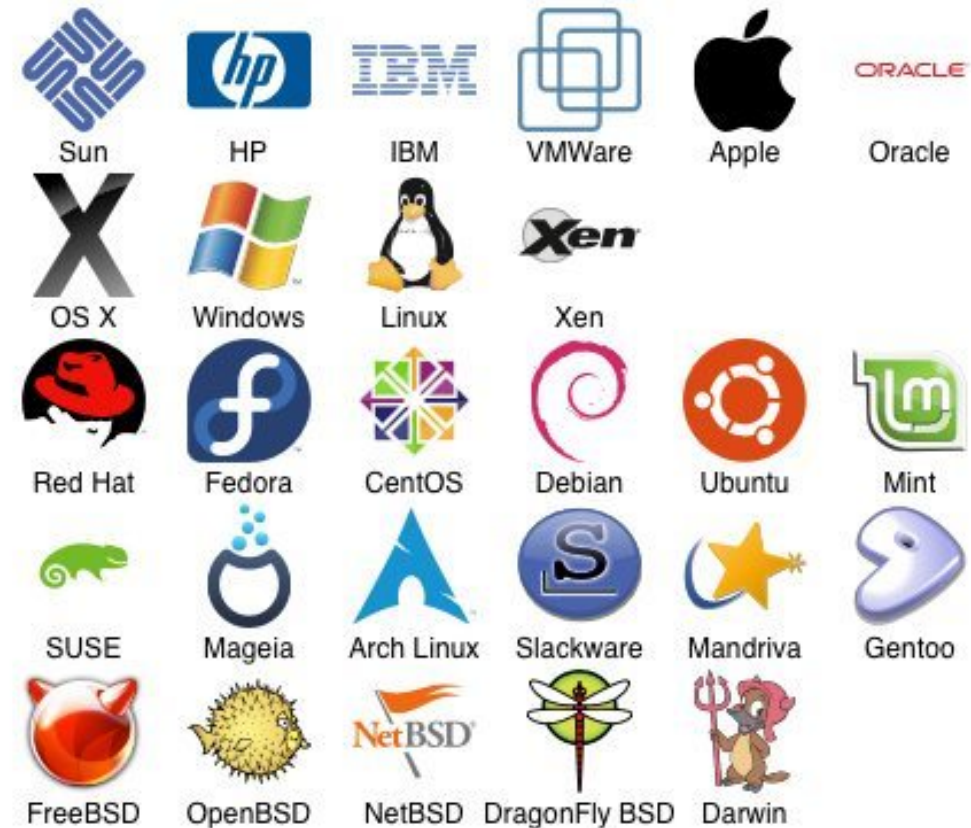
Computational Biology

Lecture 1

Dr. Chris Bird

LINUX is a Free & Open Source Version of the UNIX Operating System

- An **operating system** is the primary interface between you and the computer
- **Open source** is a decentralized development model where all aspects of a project are viewable and generally free to use
- Linux is free
 - Supercomputers
 - Useful text manipulation tools



2 Primary Methods of Interfacing with Computers



Graphical User Interface (GUI)



Command-line Interface (CLI)

```
checking lzma.h presence... yes
checking for lzma.h... yes
checking if lzma version >= 5.0.3... yes
checking for pcre_fullinfo in -lpcre... yes
checking pcre.h usability... yes
checking pcre.h presence... yes
checking for pcre.h... yes
checking pcre/pcre.h usability... no
checking pcre/pcre.h presence... no
checking for pcre/pcre.h... no
checking if PCRE version >= 8.20, < 10.0 and has UTF-8 support... yes
checking if PCRE version >= 8.32... yes
checking whether PCRE support suffices... yes
checking for pcre2-config... no
checking for curl-config... /home/cbird/anaconda3/bin/curl-config
checking libcurl version ... 7.64.0
checking curl/curl.h usability... yes
checking curl/curl.h presence... yes
checking for curl/curl.h... yes
checking if libcurl is version 7 and >= 7.22.0... yes
checking if libcurl supports https... no
configure: error: libcurl >= 7.22.0 library and headers are required with support for https
(base) cbird@LAPTOP-URS0LRPO:~/downloads/R-3.6.1$ ls
ChangeLog  configure  doc        m4         Makefile.in  Makefrag.cxx  README  SVN-REVISION  VERSION
config.log  configure.ac  etc       Makeconf.in  Makefrag.cc  Makefrag.m    share   tests         VERSION-NICK
config.site  COPYING      INSTALL  Makefile.fw  Makefrag.cc_lo  go            src     tools
(base) cbird@LAPTOP-URS0LRPO:~/downloads/R-3.6.1$ less -S config.log
(base) cbird@LAPTOP-URS0LRPO:~/downloads/R-3.6.1$ F
```



Why use CLI Linux?

- Free
- Automation
- Flexibility
- Powerful
- Designed for developers
- Supercomputers use it
- Many software tools for biologists
- Large body of support online



The UNIX Philosophy

- One program (**command**) does one thing
- All programs accept input as a text stream and output a modified **text** stream
- Programs can be linked together into serial **pipelines** to achieve complex results



The Unix philosophy (excerpt):

-Make each program do one thing well.

-Expect the output of every program to become the input to another, as yet unknown program.

McIlroy, Pinson & Tague, 1978

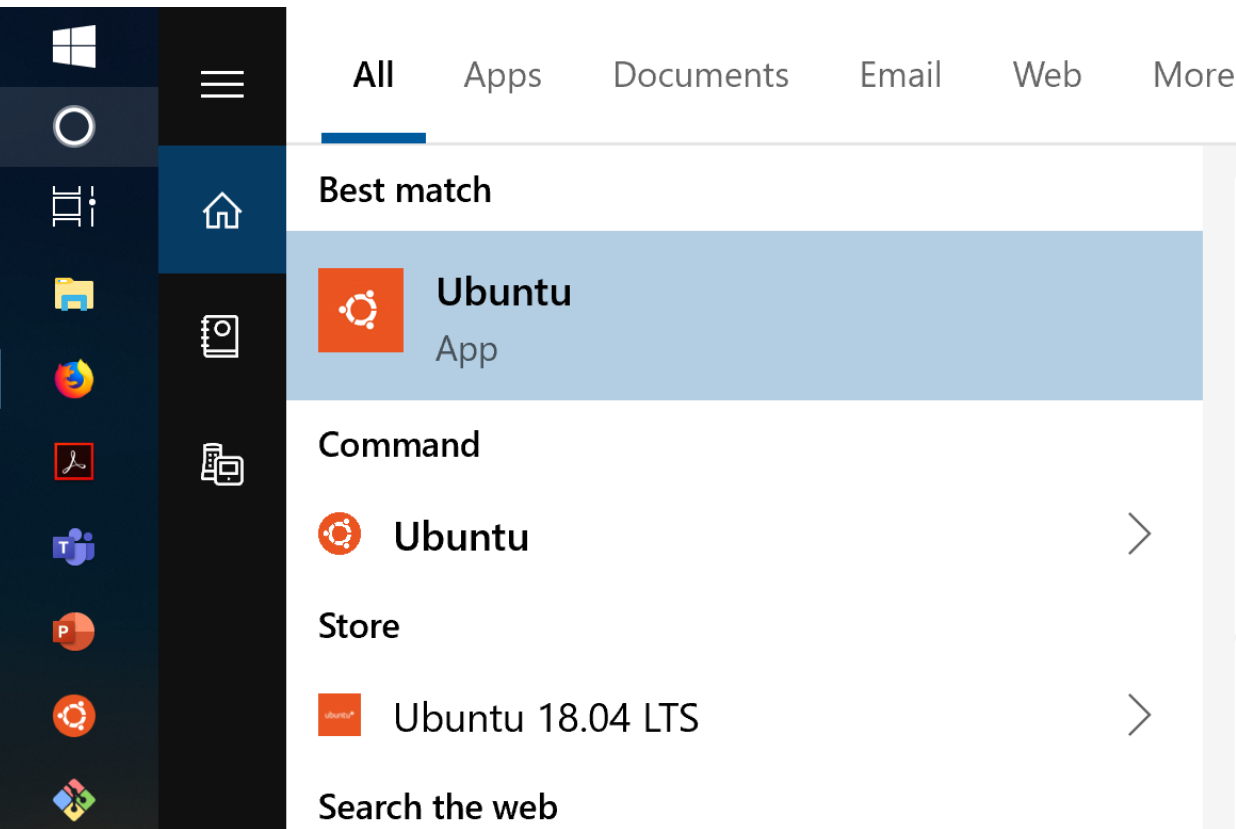
Linux CLI Pipelines Facilitate Scientific Reproducibility and Long-Term Efficiency

Comparison of GUI and CLI for manipulating data

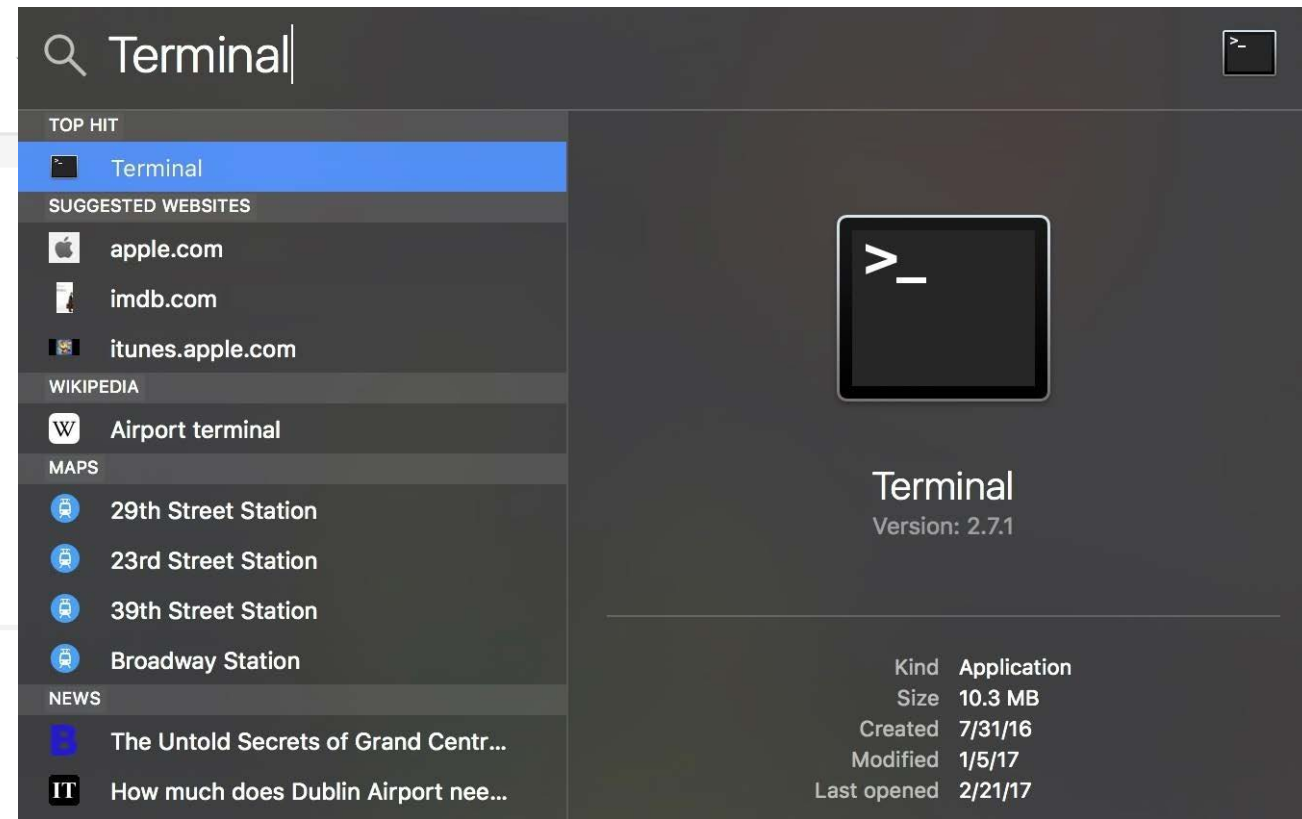
	GUI	CLI
Learning curve	Short, shallow	Long, steep
Amount of your time taken to process large amounts of data	Long	Short
Process Documented or Recorded	Often not, mouse clicks	Always
Ability to identify mistake	Poor	Excellent
Time to recover from mistake	Long	Short
Ease for another lab to reproduce	Difficult to impossible	Simple

Open A Terminal Window

WIN10: Search **Ubuntu**

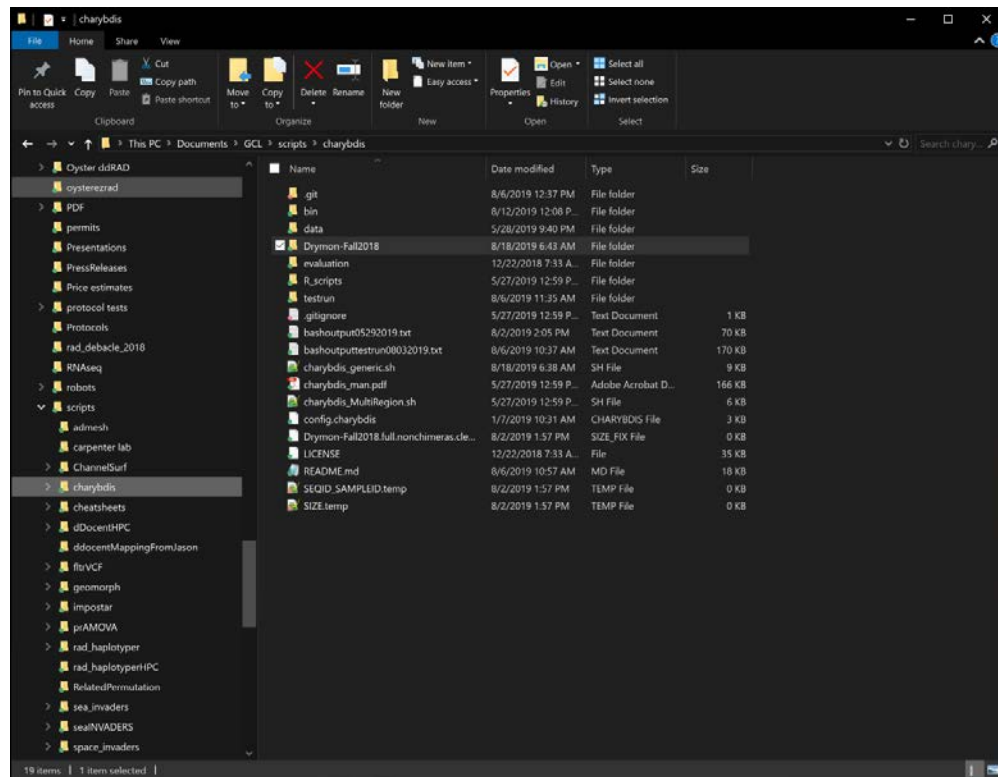


MacOS: Search **Terminal**

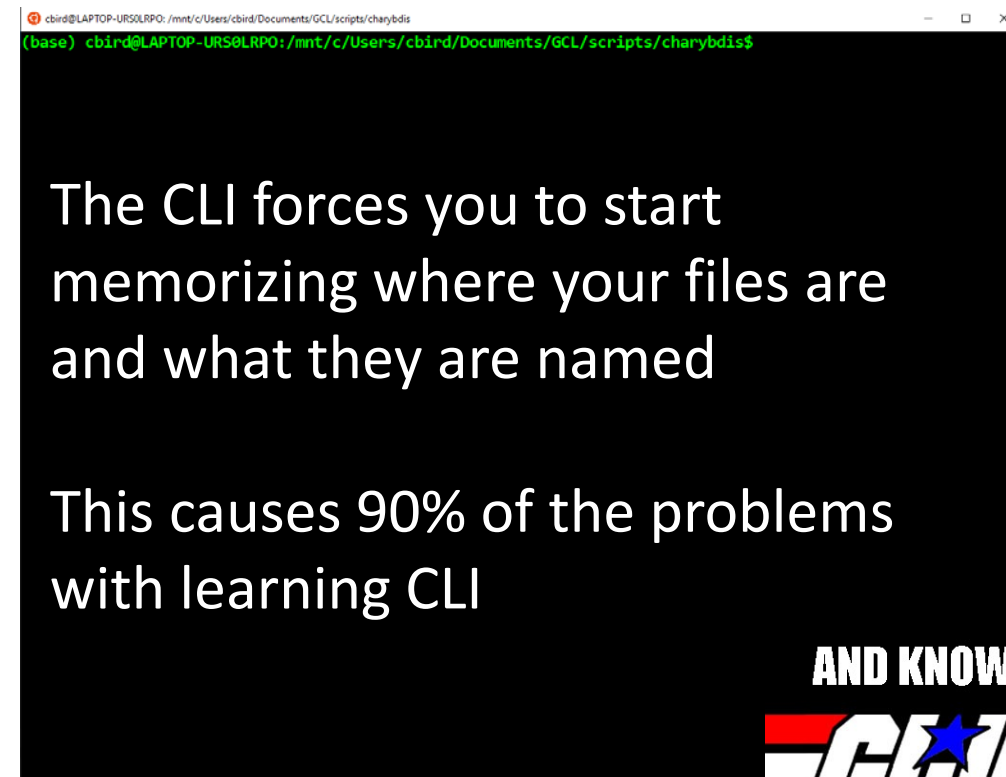


The **Directory Structure** is the Organization of Files and Folders (aka Directories) In Your Computer

WIN10 File Explorer

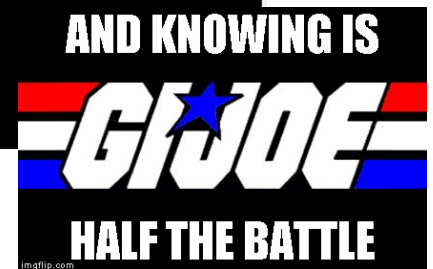


Ubuntu Terminal



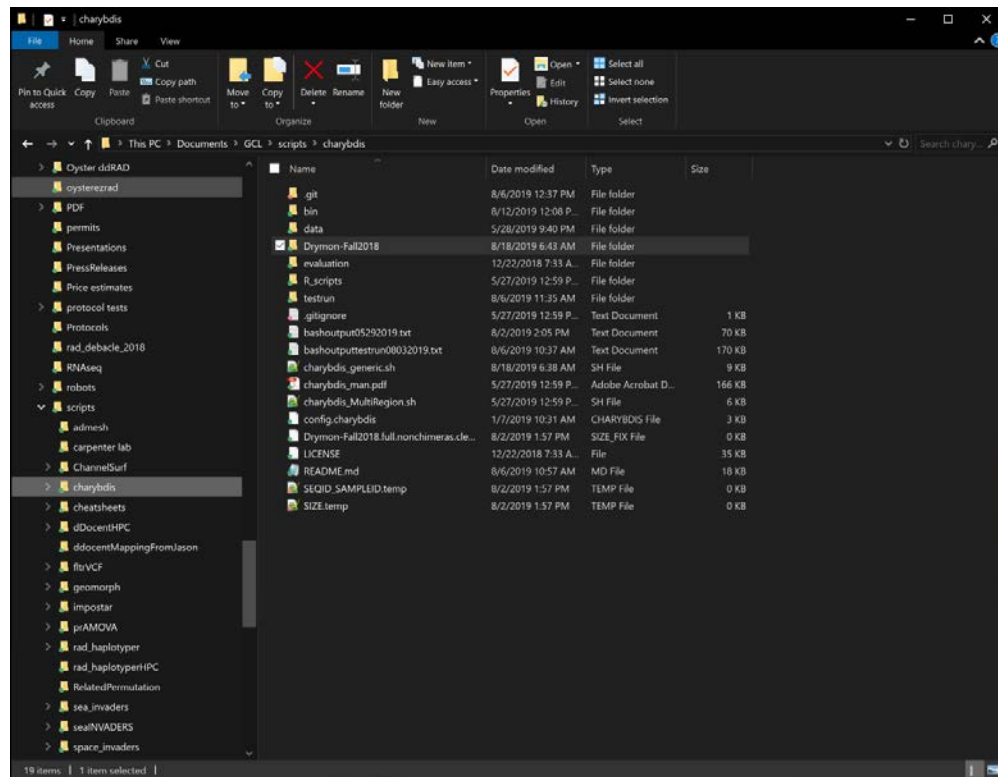
The CLI forces you to start memorizing where your files are and what they are named

This causes 90% of the problems with learning CLI

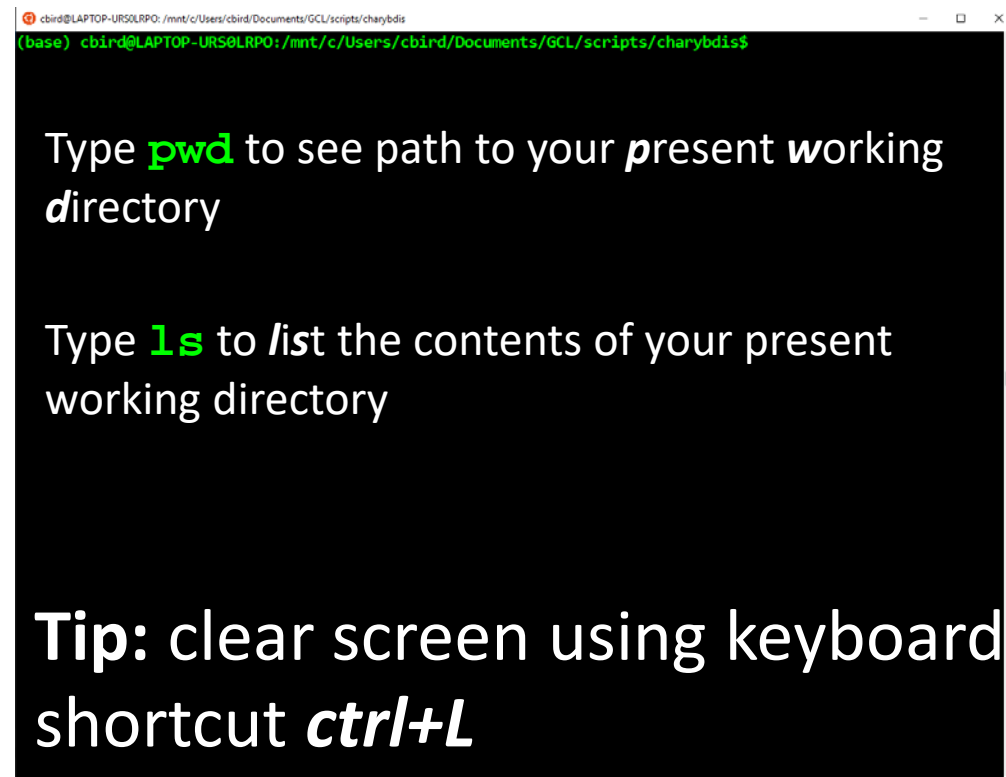


The **Directory Structure** is the Organization of Files and Folders (aka Directories) In Your Computer

WIN10 File Explorer



Ubuntu Terminal



Unix/Linux Command Line Terminology

The **path** is the address of a file or directory in the directory structure

Description	<u>Path</u> in Unix, Linux, Ubuntu, MacOS, Android	Path in Windows
Root , or top of the directory tree	/	c:\
A file named file.txt in the root dir	/file.txt	c:\file.txt
A directory named folder1 in the root dir	/folder1	c:\folder1
A file named dna.txt in folder1	/folder1/dna.txt	c:\folder1\dna.txt

Important Directories

`/bin`

- Contains several basic programs

`/dev`

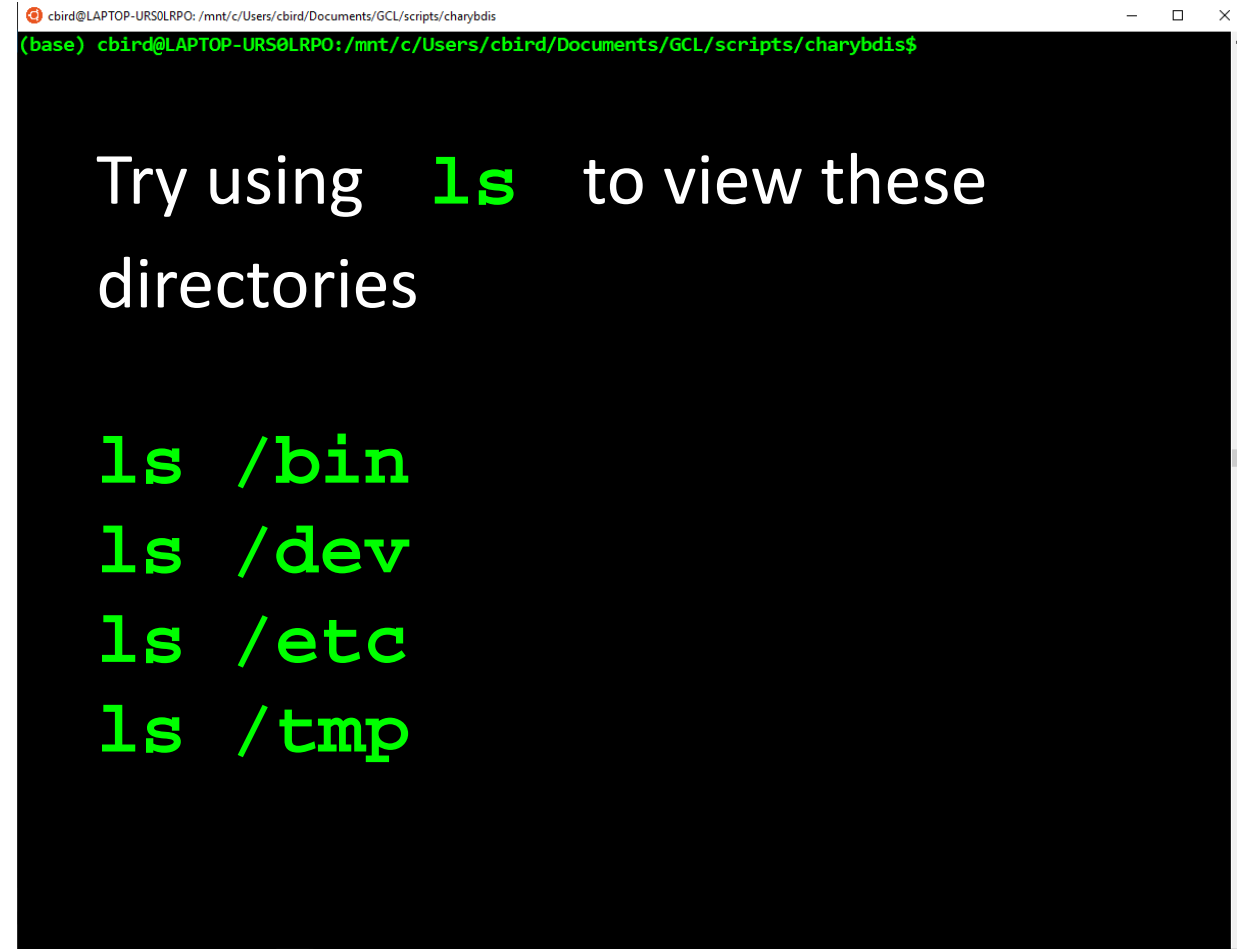
- Contains the files connecting to devices such as the keyboard, mouse, and screen

`/etc`

- Contains configuration files

`/tmp`

- Contains temporary files



```
cbird@LAPTOP-URS0LRPO: /mnt/c/Users/cbird/Documents/GCL/scripts/charybdis$  
(base) cbird@LAPTOP-URS0LRPO: /mnt/c/Users/cbird/Documents/GCL/scripts/charybdis$  
  
Try using ls to view these  
directories  
  
ls /bin  
ls /dev  
ls /etc  
ls /tmp
```

Your Home Directory

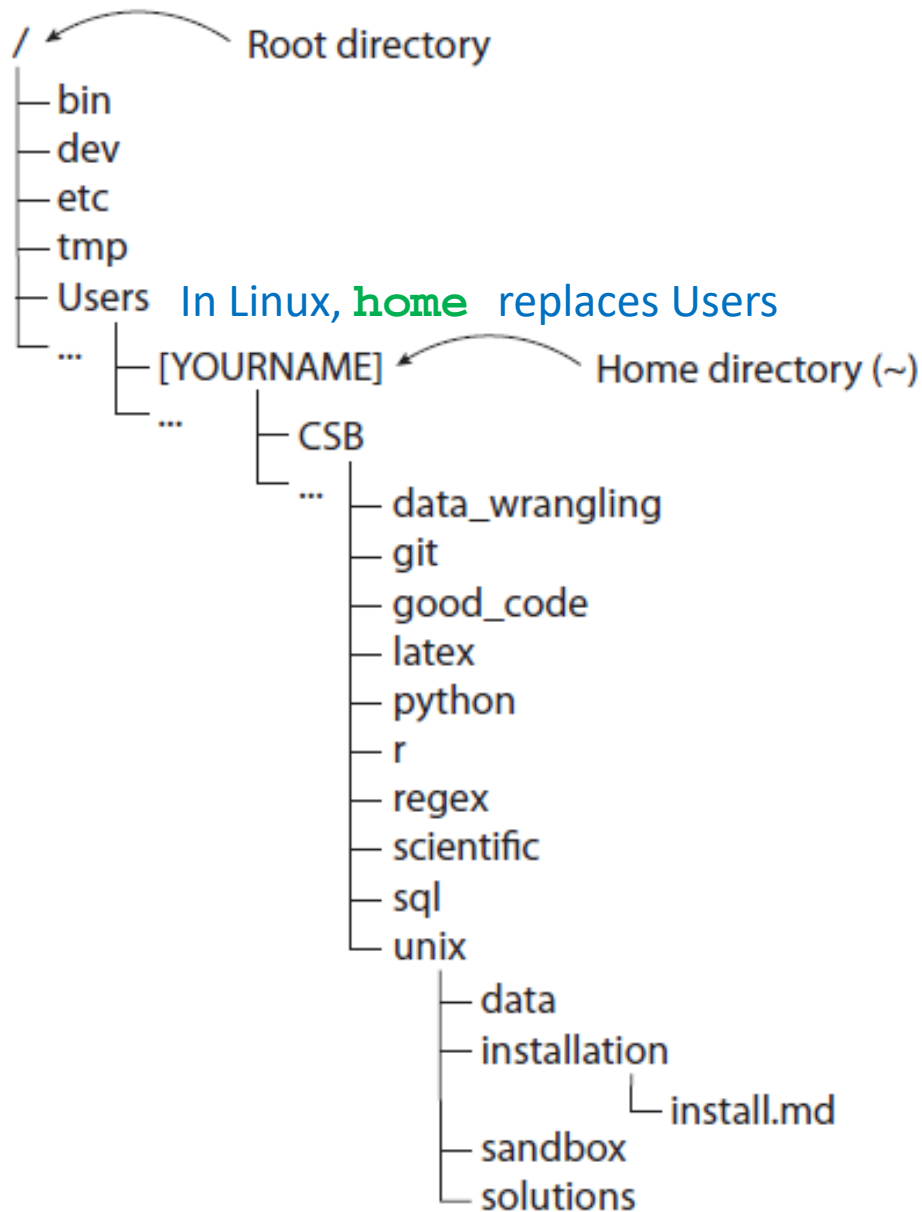
`/username/home`

- Starting or login directory
- Specific to user
- Place for personal files, dirs, programs, downloads etc

`$HOME`

- The **path** to your home dir is stored in this **variable**
- A variable stores information
- Always preceded by a **\$** after it is created
- **\$HOME** is an **environmental variable** created by the operating system and bash

```
cbird@LAPTOP-URS0LRPO: /mnt/c/Users/cbird/Documents/GCL/scripts/charybdis$  
(base) cbird@LAPTOP-URS0LRPO:/mnt/c/Users/cbird/Documents/GCL/scripts/charybdis$  
  
echo $HOME  
pwd  
ls  
ls $HOME  
ls ~  
  
If you followed install instructions,  
you should have a CSB dir
```



Full path of the file install.md:

/Users/[YOURNAME]/CSB/unix/installation/install.md

Directory Tree

Showing Contents of \$HOME/CSB/unix/installation

```
cbird@LAPTOP-URS0LRPO: /mnt/c/Users/cbird/Documents/GCL/scripts/charybdis$
(base) cbird@LAPTOP-URS0LRPO: /mnt/c/Users/cbird/Documents/GCL/scripts/charybdis$

ls $HOME/CSB/unix/installation

On your own time, if you install tree,
you can view the directory tree on screen

sudo apt-get install tree
cd $HOME
tree CSB
tree -L 1 CSB
tree -L 2 CSB
man tree

On mac:
brew install tree
```


CSB/unix Repository

CSB/unix/data

- Contains data for examples and exercises

CSB/unix/installation

- Contains instructions for installing software for this chapter

CSB/unix/sandbox

- Dir where we work and experiment

CSB/unix/solutions

- Solutions in code (bash) pseudocode (plain English) for your consultation when you get stuck with an exercise.

cbird@LAPTOP-URS0LRPO: /mnt/c/Users/cbird/Documents/GCL/scripts/charybdis

(base) cbird@LAPTOP-URS0LRPO:/mnt/c/Users/cbird/Documents/GCL/scripts/charybdis\$

```
cd $HOME
```

```
ls CSB/unix
```

```
ls CSB/unix/data
```

```
ls CSB/unix/installation
```

```
ls CSB/unix/sandbox
```

```
ls CSB/unix/solutions
```

Tip: use the ↑ key to recall last command

\$ Welcome to the Matrix

1.4 The Shell

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The Shell

- The **shell** is software that controls the operating system **kernel** and is accessed through a **terminal** window
- The shell we are using in Ubuntu and MacOS is **BASH**, or **Born Again Shell**
- The **commands** we've been using are BASH commands which allow us to control the operating system

~

- Indicates where I am, the home dir

\$

- Indicates the terminal is ready to accept commands
- From here forward, \$<space><command> indicates you should type the command into the terminal

#

- A hash symbol means that everything that follows is a comment, usually in English

cbird@LAPTOP-URS0LRPO: ~

```
(base) cbird@LAPTOP-URS0LRPO:~$
```

Below, I've indicated that I want you to do what follows the # by typing the command that follows the \$ and you expect your output to be similar to the line(s) not preceded by # or \$

```
# display the date and time
```

```
$ date
```

```
Sat Aug 24 12:18:24 DST 2019
```

Bash Keyboard Shortcuts

- ↑ Scroll through previous commands
- Tab** autocomplete command, dir, or file name
 - if you hit tab and nothing happens there's either multiple matches or 0 matches
- Tab,Tab** show matches
- Ctrl+A** Go to the beginning of the line.
- Ctrl+E** Go to the end of the line.
- Ctrl+L** Clear the screen.
- Ctrl+U** Clear the line before the cursor position.
- Ctrl+K** Clear the line after the cursor.
- Ctrl+C** Kill the command that is currently running.
- Ctrl+D** Exit the current shell.
- Alt+F** Move cursor forward one word (in OS X, Esc+F).
- Alt+B** Move cursor backward one word (in OS X, Esc+B).

cbird@LAPTOP-URS0LRPO: ~

(base) cbird@LAPTOP-URS0LRPO:~\$

```
# try some of the shortcuts
$
```

Bash Commands

`cal 2020 -j`

- **Commands** like `cal` are programs that follow the UNIX philosophy
- **Arguments** like `2020` are essentially options, order usually matters and some commands require particular arguments
 - `cp` or copy requires at least which file to copy and where to copy it, in that order
- `-j` is an **option**, in this case it means Julian calendar
 - `--julian` is the same as `-j`, options that are words are always preceded by two dashes

```
# print calendar
```

```
$ cal
```

```
August 2019
```

Su	Mo	Tu	We	Th	Fr	Sa
				1	2	3
4	5	6	7	8	9	10
11	12	13	14	15	16	17
18	19	20	21	22	23	24
25	26	27	28	29	30	31

```
$ cal 2020
```

```
$ cal -j
```

```
$ cal --julian
```

```
$ cal -j 2020
```

If you want to stop a command, ***ctrl+c***

Getting Bash Help

- It's impossible to remember all command and arguments
- If you know what you want to do, but you don't know the command
 - Google search "bash <English description of what you want to do>"
- If you know the command, but you don't know the arguments
 - `man <CommandName>`
 - All manuals have same format

```
# view calendar manual
```

```
$ man cal
```

```
NAME
```

```
<name and brief descrip>
```

```
SYNOPSIS
```

```
<examples of how to run>
```

```
DESCRIPTION
```

```
<detailed description>
```

```
<list of arguments/options>
```

Tip: scroll with arrow keys and close manual with *q* key

Changing Directories

`cd ..`

- Move up to parent directory

`cd /`

- Move to root directory

`cd ~`

- Move to home directory

`cd -`

- Move to last directory

`pwd`

- Path to present working dir

`ls`

- Show contents of present directory

move around dir system

\$ `cd ..`

\$ `pwd`

\$ `cd /`

\$ `pwd`

\$ `cd -`

\$ `pwd`

\$ `cd ~`

\$ `pwd`

show dir contents

\$ `ls`

\$ `ls -l`

\$ `ls -ltrh`

Note: single letter *options* can typically be combined together, `-l -t -r -h`

Interpreting Output of `ls -l`

Dirs are highlighted below, files are not

```
(base) cbird@LAPTOP-URS0LRPO:~$ ls -ltrh
total 1.0K
-rwxrwxrwx 1 cbird cbird 515 Jul 10 2018 hosts
-rw-rw-rw- 1 cbird cbird 146 Jul 10 2018 initialize.bash
-rw-rw-rw- 1 cbird cbird 39 Aug 2 2018 tamucchpcmlogin.bash
-rw-rw-rw- 1 cbird cbird 42 Jan 11 2019 oduhpcmlogin.bash
-rw-rw-rw- 1 cbird cbird 61 Feb 15 2019 mntUSB.bash
-rw-rw-rw- 1 cbird cbird 93 Jun 21 06:46 onedrive.bash
drwxrwxrwx 1 cbird cbird 512 Aug 24 10:57 downloads
drwxrwxrwx 1 cbird cbird 512 Aug 24 11:25 CSB
(base) cbird@LAPTOP-URS0LRPO:~$
```

Interpreting Output of `ls -l`

Next

Slide

Usr

Grp

Size

Date

Names

-rwxrwxrwx	1	cbird	cbird	515	Jul 10	2018	hosts
-rw-rw-rw-	1	cbird	cbird	146	Jul 10	2018	initialize.bash
-rw-rw-rw-	1	cbird	cbird	39	Aug 2	2018	tamucchpcmlogin.bash
-rw-rw-rw-	1	cbird	cbird	42	Jan 11	2019	oduhpcmlogin.bash
-rw-rw-rw-	1	cbird	cbird	61	Feb 15	2019	mntUSB.bash
-rw-rw-rw-	1	cbird	cbird	93	Jun 21	06:46	onedrive.bash
drwxrwxrwx	1	cbird	cbird	512	Aug 24	10:57	downloads
drwxrwxrwx	1	cbird	cbird	512	Aug 24	11:25	CSB

(base) cbird@LAPTOP-URS0LRPO:~\$

Interpreting Output of `ls -l`

Permissions

	User	Group	Global
File	<div><div>r</div><div>w</div><div>-</div></div>	<div><div>r</div><div>w</div><div>-</div></div>	<div><div>r</div><div>w</div><div>-</div></div>
Dir	<div><div>r</div><div>w</div><div>x</div></div>	<div><div>r</div><div>w</div><div>x</div></div>	<div><div>r</div><div>w</div><div>x</div></div>

Paths

- A path is the address of file or directory
- An **absolute path** is complete and starts with root `/` or a variable that starts with root
 - These return the same result regardless of pwd
 - `/home/<username>/CSB`
 - `~/CSB`
 - `$HOME/CSBB`
- **Relative paths** start from the present location
 - These only work if you are in the right dir
 - `.` Means present directory
 - `..` means parent directory
 - `./CSB`
 - `CSB`
- It's best not to use spaces in dir and file names
 - See pg 21 for dealing w/ spaces

```
$ cd ~
```

```
# show contents of CSB dir
```

```
# absolute paths
```

```
$ ls /home/<username>/CSB
```

```
$ ls ~/CSB
```

```
$ ls $HOME/CSB
```

```
# relative paths
```

```
$ ls ./CSB
```

```
$ ls CSB
```

Note: if a path includes a space, either wrap path in quotes or precede each space with `\`

Mind Expander 1.1

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1.5 Commands to Remember

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Copy with **cp** <from> <to>

```
# goto sandbox
$ cd ~/CSB/unix/sandbox

# copy the following file to the present directory
$ cp ../data/Buzzard2015_about.txt .

# copy file and rename it in present dir
$ cp ../data/Buzzard2015_about.txt ./Buzzard2015_about2.txt

# copy whole data dir to present dir, then view present dir
$ cp -rf ../data .
$ ls
```

Note: -r means recursive, -f means force

Move or rename with **mv** **<from>** **<to>**

```
# make sure you are still in sandbox, if not then cd ~/CSB/unix/sandbox
$ pwd

# move the file to the data directory
$ mv Buzzard2015_about2.txt ../data

# rename a file that isn't in your pwd
$ mv ../data/Buzzard2015_about2.txt ../data/Buzzard2015_about_new.txt

# check your work
$ ls ../data
```

Note: bash gives no positive feedback, only negative if something is wrong

Create file with **touch** <filename>

```
# make sure you are still in sandbox, if not then cd ~/CSB/unix/sandbox
$ pwd

# inspect the current contents of the directory
$ ls -l

# create a new file (you can list multiple files)
$ touch new_file.txt

# inspect the contents of the directory again
$ ls -l

# if you touch the file a second time, the time of last access will change
$ touch new_file.txt
$ ls -l
```

Note: bash gives no positive feedback, only negative if something is wrong

Remove file(s) or dir(s) with **rm <name>**

Make dir with **mkdir <name>**

```
# make sure you are still in sandbox, if not then cd ~/CSB/unix/sandbox
```

```
$ pwd
```

```
# delete new_file.txt in sandbox, the -i requests confirmation
```

```
$ rm -i new_file.txt
```

```
# make dir d1 in present dir, d2 in d1, and d3 in d2; if you have tree try it
```

```
$ mkdir -p d1/d2/d3
```

```
$ tree d1
```

```
d1
├── d2
│   └── d3
```

```
# remove the d1,d2,& d3 dirs recursively
```

```
$ rm -rf d1
```

be careful with **rm**, you could delete your whole computer and there is no undo

View large files with	<code>less -S <filename></code>
Print and concatenate files	<code>cat <filename></code>
Print and sort files	<code>sort <filename></code>

```
# move to the data dir
```

```
$ cd ~/CSB/unix/data
```

```
# look at DNA alignment file, try duckduckgo search on "bash less commands"
```

```
$ less -S Marra2014_data.fasta
```

```
# type /ATCG inside of less to search; u=up, d=down, G=end, g=begin, q=exit
```

```
# concatenate files and/or print to screen
```

```
$ cat Marra2014_about.txt Gesquiere2011_about.txt Buzzard2015_about.txt
```

```
# print the sorted lines of a file
```

```
$ sort Gesquiere2011_data.csv
```

```
# sort numerically by column 2 in reverse order and view in less
```

```
$ sort -n -k2 -r Gesquiere2011_data.csv | less
```

Count words with
Determine file type

```
wc <filename>  
file <filename>
```

```
# count lines, words, and characters
```

```
$ wc Gesquiere2011_about.txt
```

```
# count lines only
```

```
$ wc -l Marra2014_about.txt
```

```
# determine file type, ASCII is a type of human-readable text file
```

```
$ file Marra2014_about.txt
```

```
Marra2014_about.txt: ASCII English text
```

Don't forget to use *Tab* key to autocomplete names, prevents spelling mistakes

Get beginning of file **head -n # <filename>**
Get end of file **tail -n # <filename>**

```
# display first two lines of a file
```

```
$ head -n 2 Gesquiere2011_data.csv
```

```
# display last two lines of file
```

```
$ tail -n 2 Gesquiere2011_data.csv
```

```
# display from line 2 onward
```

```
# (i.e., removing the header of the file)
```

```
$ tail -n +2 Gesquiere2011_data.csv
```

```
# display all but the last line
```

```
$ head -n -1 Gesquiere2011_data.csv
```

Don't forget to use *Tab* key to autocomplete names, prevents spelling mistakes

Mind Expander 1.2

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1.6 Advanced Commands

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Redirection of output (stdout) to file	<code>[command] > filename</code>
Append stdout to file	<code>[command] >> filename</code>
Redirect contents of file to stdin	<code>[command] < filename</code>

```
# let's start by moving to our sandbox
```

```
$ cd ~/CSB/unix/sandbox
```

```
# print text to screen, then print to file, then print file to screen
```

```
$ echo "My first line"
```

```
$ echo "My first line" > test.txt
```

```
$ cat test.txt
```

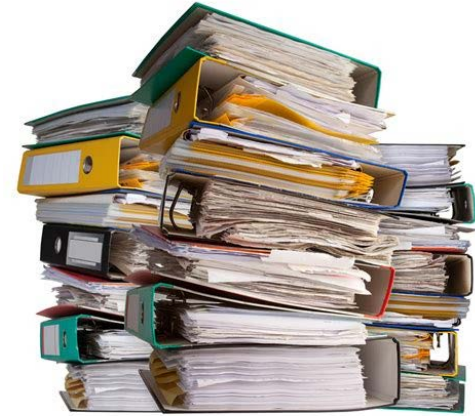
```
# append file with additional text, then print file to screen
```

```
$ echo "My second line" >> test.txt
```

```
$ cat test.txt
```

Don't forget to use *Tab* key to autocomplete names, prevents spelling mistakes

Problem Solving Scenario



- A machine provides you with thousands of data files
- There's so many, it's breaking your file browser
- How many files are there?
- We will use `unix/data/Saavedra2013` as an example of a directory with many files

```
# save file names to file in pwd
$ ls ../data/Saavedra2013 > filelist.txt

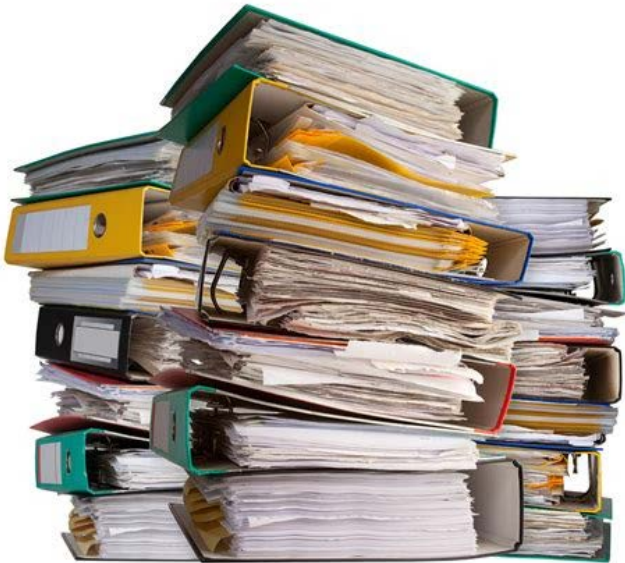
# look at the file
$ cat filelist.txt

# count lines in a file
$ wc -l filelist.txt

# remove the file
$ rm filelist.txt
```

Problem Solving Scenario – Application of pipe |

- A pipe passes the stdout from one command to the stdin of another
- How many files are there?



```
# list file names
$ ls ../data/Saavedra2013

# list file names and pipe into wc
$ ls ../data/Saavedra2013 | wc -l
59
```

TSV and CSV Data Files

	Ozone	Solar.R	Wind	Temp	Month	Day
128	32.0	92.0	15.5	84	9	6
78	61.0	285.0	6.3	84	7	18
105	65.0	157.0	9.7	80	8	14
64	NaN	101.0	10.9	84	7	4
98	122.0	255.0	4.0	89	8	7
145	36.0	139.0	10.3	81	9	23
27	23.0	13.0	12.0	67	5	28
28	45.0	252.0	14.9	81	5	29
113	9.0	36.0	14.3	72	8	22
132	24.0	259.0	9.7	73	9	10

- Tab Separated Values (TSV)
 - Tabs denote columns
- Comma Separated Values (CSV)
 - Commas denote columns
- Tidy data
 - Each row is one unit of observation
 - Each column is one dimension or aspect of the units of observation
- File extensions not always accurate

It's Easy to Convert Among Formats Using **tr**

```
# view contents of csv
```

```
$ less -S ../data/Pacifici2013_data.csv
```

```
# replace semicolons with commas using tr [find] [replace]
```

```
$ cat ../data/Pacifici2013_data.csv | tr ";" "," | less -S
```

```
# view as tsv
```

```
# \t is the nearly universal symbol for tab
```

```
$ cat ../data/Pacifici2013_data.csv | tr ";" "\t" | less -S
```

tr is short for translate

Using **cut** to grab columns and **head** to grab rows

```
# change directory
```

```
$ cd ~/CSB/unix/data
```

```
# display first line of file (i.e., header of CSV file)
```

```
$ head -n 1 Pacifici2013_data.csv
```

```
# display first column of file
```

```
$ cut -d ";" -f 1 Pacifici2013_data.csv
```

```
# display second through fourth columns
```

```
$ cut -d ";" -f 2-4 Pacifici2013_data.csv
```

```
# display first "cell" of data
```

```
$ head -n 1 Pacifici2013_data.csv | cut -d ";" -f 1
```

Connecting **cut** **head** **tail** **sort** **uniq**

```
# select 2nd column, display first 5 elements
```

```
$ cut -d ";" -f 2 Pacifici2013_data.csv | head -n 5
```

```
# select 2nd and 8th columns, display first 3 elements
```

```
$ cut -d ";" -f 2,8 Pacifici2013_data.csv | head -n 3
```

```
# select 2nd column without header, show 5 first elements
```

```
$ cut -d ";" -f 2 Pacifici2013_data.csv | tail -n +2 | head -n 5
```

```
# identify the orders in csv
```

```
# select 2nd column without header, unique sorted elements
```

```
$ cut -d ";" -f 2 Pacifici2013_data.csv | tail -n +2 | sort |\n> uniq
```

```
# count how many records per order in csv
```

```
$ cut -d ";" -f 2 Pacifici2013_data.csv | tail -n +2 | sort |\n> uniq -c
```


Welcome to the Matrix Questions?

Computational Biology

Lecture 1

Dr. Chris Bird

Mind Expander 1.3

Computational Biology

Lecture 1

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