

BGGN 213
Introduction to UNIX
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- ## Recap From Last Time:
- **Substitution matrices:** Where our alignment match and mis-match scores typically come from
 - **Comparing methods:** The trade-off between *sensitivity*, *selectivity* and *performance*
 - **Sequence motifs and patterns:** Finding functional cues from conservation patterns
 - **Sequence profiles** and **position specific scoring matrices (PSSMs)**, Building and searching with profiles, Their advantages and limitations
 - **PSI-BLAST algorithm:** Application of iterative PSSM searching to improve BLAST sensitivity
 - **Hidden Markov models (HMMs):** More versatile probabilistic model for detection of remote similarities

Today's Menu

	Time	Topics
I	9:00-9:15 AM	Setup and Motivation
II	9:15-10:00 AM	Beginning Unix
	10:00-10:40 AM	Hands on session
III	10:40-11:00 AM	Working with Unix
IV	11:00-11:20 PM	How to Get Working
	11:20-11:45 PM	Hands on session

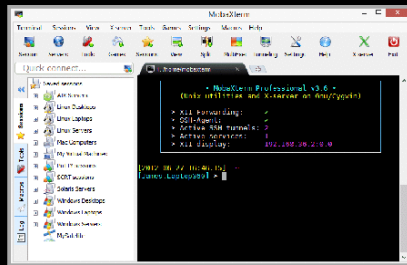
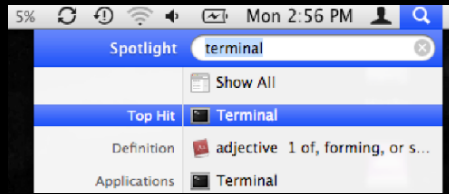
https://bioboot.github.io/bggn213_f17/setup/

- ## Setup Checklist
- https://bioboot.github.io/bggn213_f17/setup/
- ✓ **Mac:** Terminal or **PC:** MoblXterm
 - ✓ **PC:** MoblXterm **CygUtils** plugin (core UNIX tools)
 - ✓ Downloaded our class specific **jetstream keyfile** (See email: This is required for connecting to jetstream virtual machines with your terminal.)
 - ✓ Example data downloaded: [_bggn213_01_unix.zip](#)
- Problems?

Lets get started...

Do it Yourself!

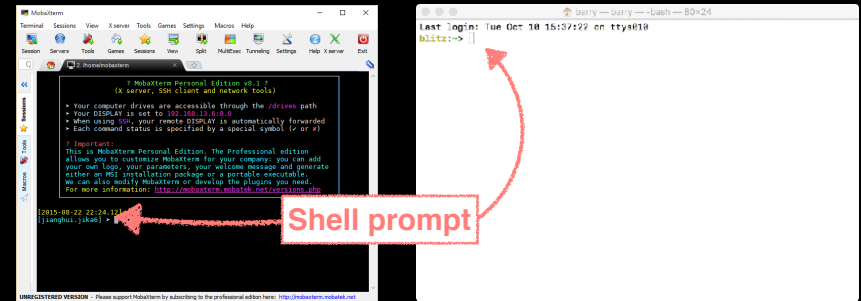
Mac
Terminal



PC
MobaXterm

SideNote: Terminal vs Shell

- **Shell**: A command-line interface that allows a user to interact with the operating system by typing commands.
- **Terminal [emulator]**: A graphical interface to the shell (i.e. the window you get when you launch MobaXterm).



Introduction To UNIX Shell



Motivation

Why do we use Unix?

Modularity	Core programs are modular and work well with others
Programmability	Best software development environment
Infrastructure	Access to existing tools and cutting-edge methods
Reliability	Unparalleled uptime and stability
Unix Philosophy	Encourages open standards

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Modularity

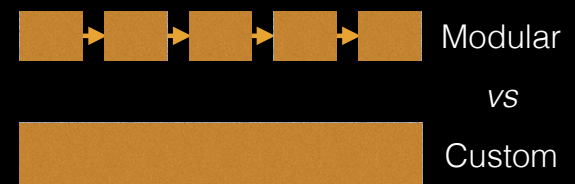
The Unix shell was designed to allow users to easily build complex workflows by interfacing smaller **modular programs** together.



An alternative approach is to write a **single complex program** that takes raw data as input, and after hours of data processing, outputs publication figures and a final table of results.



Which would you prefer and why?



Advantages/Disadvantages

The 'monster approach' is **customized to a particular project** but results in **massive, fragile** and difficult to modify (therefore **inflexible, untransferable, and error prone**) code.

With **modular workflows**, it's easier to:

- **Spot errors** and figure out where they're occurring by inspecting intermediate results.
- **Experiment** with alternative methods by swapping out components.
- **Tackle novel problems** by remixing existing modular tools.

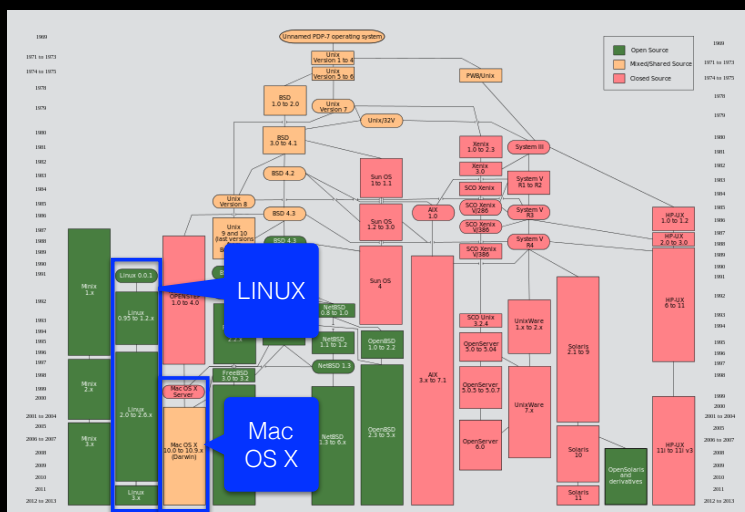
Unix 'Philosophy'

"Write programs that do one thing and do it well. Write programs to work together and that encourage open standards. Write programs to handle text streams, because that is a universal interface."



— Doug McIlroy

Unix family tree [1969-2010]



Source: https://commons.wikimedia.org/wiki/File:Unix_history-simple.svg

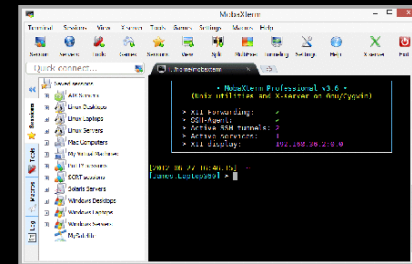
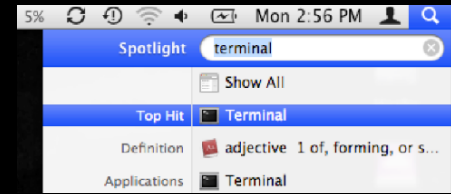
Basics	File Control	Viewing & Editing Files	Misc. useful	Power commands	Process related
ls	mv	less	chmod	grep	top
cd	cp	head	echo	find	ps
pwd	mkdir	tail	wc	sed	kill
man	rm	nano	curl	sudo	Ctrl-c
ssh	 (pipe)	touch	source	git	Ctrl-z
scp	> (write to file)		cat	R	bg
	< (read from file)		tmux	python	fg

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Lets get started...

Do it Yourself!

Mac Terminal



PC
MobaXterm

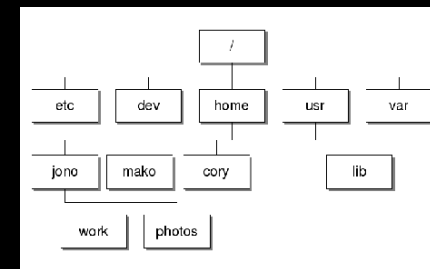
Beginning Unix

Getting started with basic Unix commands

Download the example data and mv to your Desktop
[bggn213_01_unix.zip](#)

File System Structure

- Information in the file system is stored in files, which are stored in directories (folders). Directories can also store other directories, which forms a directory tree.



- The forward slash character '/' is used to represent the root directory of the whole file system, and is also used to separate directory names. E.g. **/home/jono/work/bggn213_notes.txt**

Basics: Using the filesystem

ls	List files and directories
cd	Change directory (i.e. move to a different 'folder')
pwd	Print working directory (which folder are you in)
mkdir	MaKe a new DIRectories
cp	CoPy a file or directory to somewhere else
mv	MoVe a file or directory (basically rename)
rm	ReMove a file or directory

Side Note: File Paths

- An **absolute path** specifies a location from the root of the file system. E.g. `/home/jono/work/bggn213_notes.txt`
- A **relative path** specifies a location starting from the current location. E.g. `../bggn213_notes.txt`

.	Single dot '.' (for current directory)
..	Double dot '..' (for parent directory)
~	Tilda '~' (for your home directory)
[Tab]	Pressing the tab key can autocomplete names

Finding the Right Hammer (**man** and **apropos**)

- You can access the manual (i.e. user documentation) on a command with **man**, e.g:
 - > `man pwd`
- The man page is only helpful if you know the name of the command you're looking for. **apropos** will search the man pages for keywords.
 - > `apropos "working directory"`

Inspecting text files

- **less** - visualize a text file:
 - use arrow keys
 - page down/page up with "space"/"b" keys
 - search by typing "/"
 - quit by typing "q"
- Also see: **head**, **tail**, **cat**, **more**

Creating text files

Creating files can be done in a few ways:

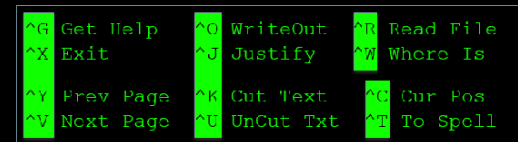
- With a **text editor** (such as **nano**, **emacs**, or **vi**)
- With the **touch** command (`> touch a_file`)
- From the command line with **cat** or **echo** and **redirection** (more on this later)
- **nano** is a simple text editor that is recommended for first-time users. Other text editors have more powerful features but also steep learning curves

Creating and editing text files with **nano**

Do it Yourself!

In the terminal type:

```
> nano yourfilename.txt
```



```
^G Get Help      ^O WriteOut     ^R Read File
^X Exit          ^J Justify     ^W Where Is
^Y Prev Page    ^K Cut Text    ^C Cur Pos
^V Next Page    ^U UnCut Txt   ^T To Spell
```

^ - Press Control

- There are many other text file editors (e.g. **vim**, **emacs** and **sublime text**, etc.)

Connecting to remote machines (with **ssh**)

- Most high-performance computing (HPC) resources can only be accessed by **ssh** (Secure Shell)

```
> ssh [user@host.address]
```

For example:

```
> ssh barry@bio3d.ucsd.edu
> ssh tb170077@IP_ADDRESS
```

Connecting to jetstream (with **ssh**)

- First we have to login online and fire-up a new “virtual machine”
- Full instructions here: https://bioboot.github.io/bggn213_f17/jetstream/boot/
- We will go through this process in the last section today!

Copying to and from remote machines (**scp**)

- The **scp** (Secure CoPy) command can be used to copy files and directories from one computer to another.

- > `scp [file] [user@host]:[destination]`
- > `scp localfile.txt bgrant@bigcomputer.net:/remotedir/.`

Basics	File Control	Viewing & Editing Files	Misc. useful	Power commands	Process related
ls	mv	less	chmod	grep	top
cd	cp	head	echo	find	ps
pwd	mkdir	tail	wc	sed	kill
man	rm	nano	curl	sudo	Crl-c
ssh	 (pipe)	touch	source	git	Crl-z
scp	> (write to file)		cat	R	bg
	< (read from file)		tmux	python	fg

Process refers to a running instance of a program

top	Provides a real-time view of all running processes
ps	Report a snapshot of the current processes
kill	Terminate a process (the "force quit" of the unix world)
Crl-c	Stop a job
Crl-z	Suspend a job
bg	Resume a suspended job in the background
fg	Resume a suspended job in the foreground
&	Start a job in the background

Hands-on time

Sections 1 to 3 of software carpentry UNIX lesson

<https://swcarpentry.github.io/shell-novice/>

~20 mins

Do it Yourself!

Working with Unix

How do we actually use Unix?

Basics	File Control	Viewing & Editing Files	Misc. useful	Power commands	Process related
ls	mv	less	chmod	grep	top
cd	cp	head	echo	find	ps
pwd	mkdir	tail	wc	sed	kill
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	< (read from file)			python	fg

Combining Utilities with Redirection (>, <) and Pipes (|)

- The power of the shell lies in the ability to combine simple utilities (*i.e.* commands) into more complex algorithms very quickly.
- A key element of this is the ability to send the output from one command into a file or to pass it directly to another program.
- This is the job of >, < and |

Side-Note: Standard Input and Standard Output streams

Two very important concepts that unpin Unix workflows:

- Standard Output (**stdout**) - default destination of a program's output. It is generally the terminal screen.
- Standard Input (**stdin**) - default source of a program's input. It is generally the command line.

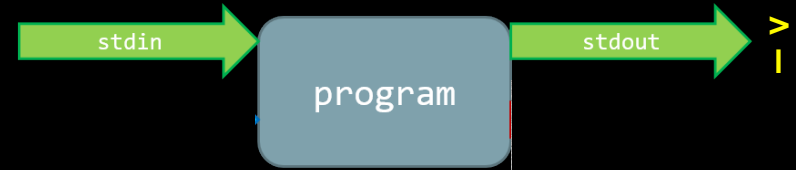
Output redirection and piping

Do it Yourself!



> ls /usr/bin # stdin is "/usr/bin"; stdout to screen

Output redirection and piping

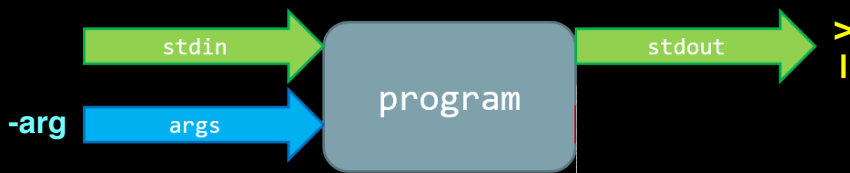


> ls /usr/bin # stdin is "/usr/bin"; stdout to screen

> ls /usr/bin > binlist.txt # stdout **redirected** to file

> ls /usr/bin | less # stdout **pipelined** to less (no file created)

Output redirection and piping



> ls /usr/bin # stdin is "/usr/bin"; stdout to screen

> ls /usr/bin > binlist.txt # stdout **redirected** to file

> ls /usr/bin | less # stdout **pipelined** to less (no file created)

> ls -l /usr/bin # extra optional input **argument** "-l"

Output redirection and piping



> ls /usr/bin # stdin is "/usr/bin"; stdout to screen

> ls /usr/bin > binlist.txt # stdout **redirected** to file

> ls /usr/bin | less # stdout **pipelined** to less (no file created)

> ls /nodirexists/ # stderr to screen

Output redirection and piping

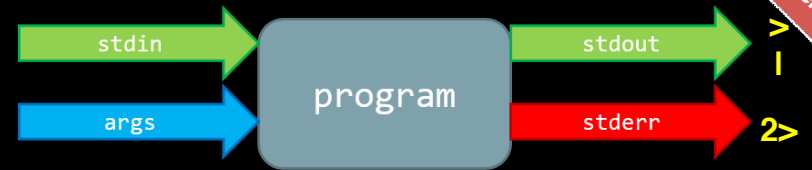
Do it Yourself!



- > ls /usr/bin # stdin is "/usr/bin"; stdout to screen
- > ls /usr/bin > binlist.txt # stdout **redirected** to file
- > ls /usr/bin | less # stdout **piped** to less (no file created)
- > ls /nodirexists/ > binlist.txt # stderr to **screen**

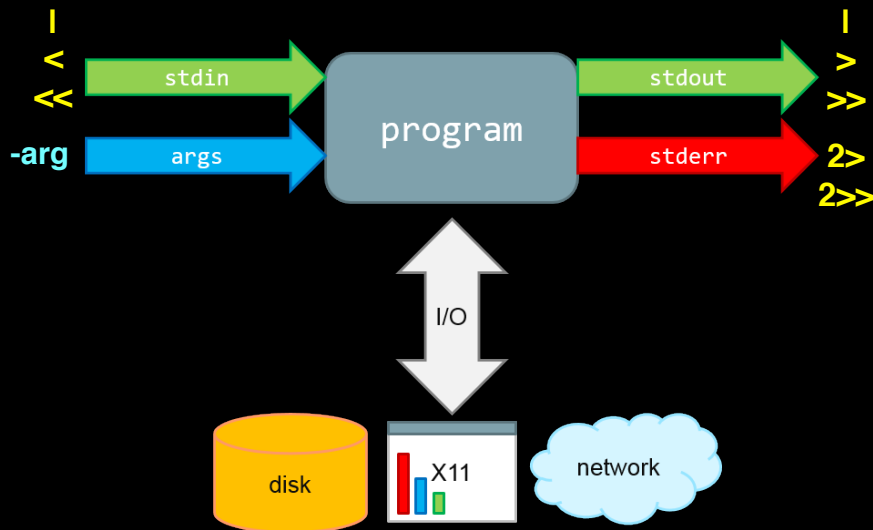
Output redirection and piping

Do it Yourself!

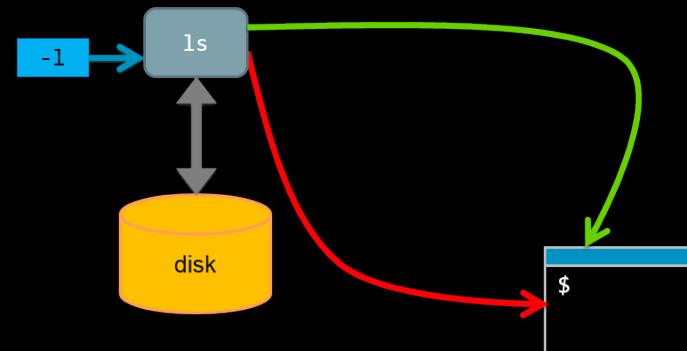


- > ls /usr/bin # stdin is "/usr/bin"; stdout to screen
- > ls /usr/bin > binlist.txt # stdout **redirected** to file
- > ls /usr/bin | less # stdout **piped** to less (no file created)
- > ls /nodirexists/ >2 binlist.txt # stderr to **file**

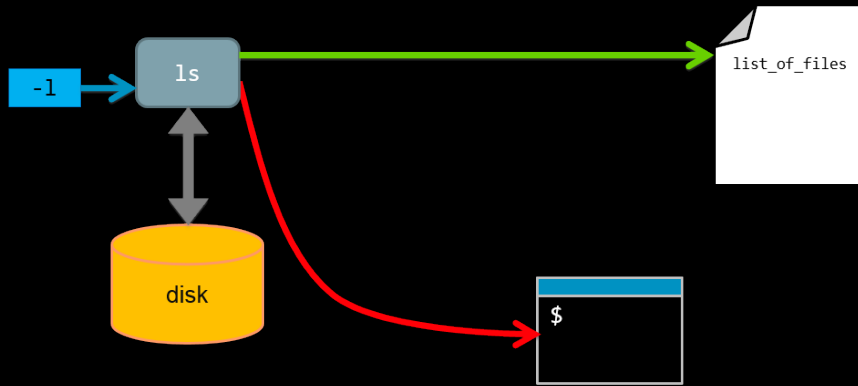
Output redirection summary



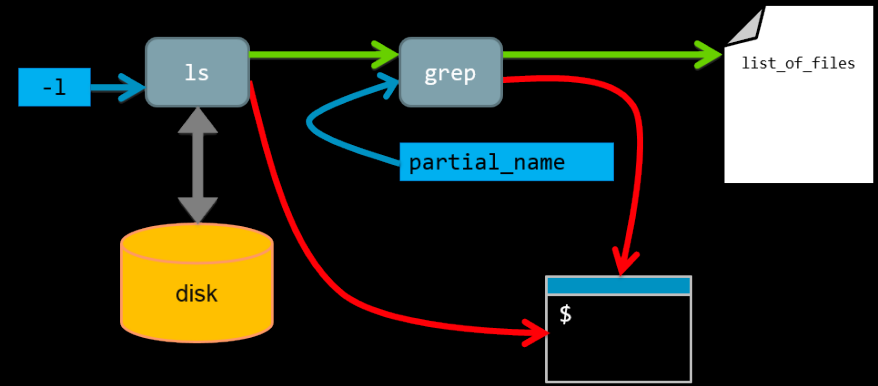
ls -l



ls -l > list_of_files

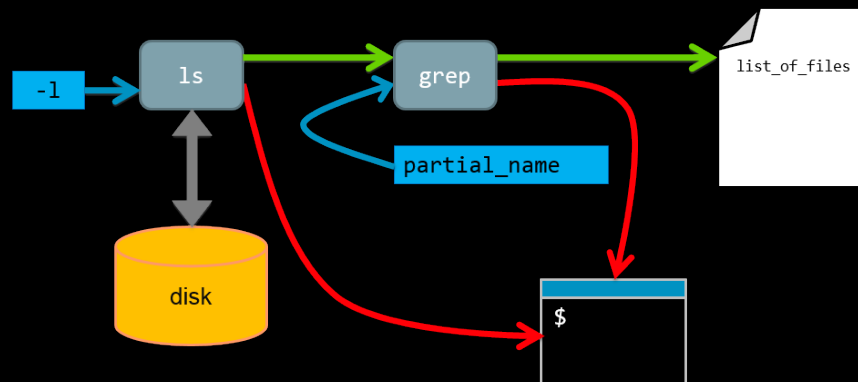


ls -l | grep partial_name > list_of_files



We have piped (|) the stdout of one command into the stdin of another command!

ls -l /usr/bin/ | grep "tree" > list_of_files



Do it Yourself!

grep: prints lines containing a string. Also searches for strings in text files.

Basics	File Control	Viewing & Editing Files	Misc. useful	Power commands	Process related
ls	mv	less	chmod	grep	top
cd	cp	head	echo	find	ps
pwd	mkdir	tail	wc	sed	kill
man	rm	nano	curl	sudo	Ctrl-c
ssh	 (pipe)	touch	source	git	Ctrl-z
	> (write to file)		cat	R	bg
	< (read from file)			python	fg

Side-Note: `grep` 'power command'

Do it Yourself!

- `grep` - prints lines containing a string pattern. Also searches for strings in text files, e.g.

```
> grep --color "GESGKS" sequences/data/seqdump.fasta
```

```
REVKLLLLGAGESGKSTIVKQMKIIEAGYSEEECKQYK
```

- `grep` is a 'power tool' that is often used with pipes as it accepts **regular expressions** as input (e.g. "`G..GK[ST]`") and has lots of useful options - see the [man page](#) for details.

`grep` example using regular expressions

Do it Yourself!

- Suppose a program that you are working with complains that your input sequence file contains non-nucleotide characters. You can eye-ball your file or ...

```
> grep -v "^>" seqdump.fasta | grep --color "[^ATGC]"
```

Exercises:

- (1). Use "man grep" to find out what the `-v` argument option is doing!
- (2). How could we also show line number for each match along with the output?
(tip you can grep the output of "man grep" for 'line number')

`grep` example using regular expressions

Do it Yourself!

- Suppose a program that you are working with complains that your input sequence file contains non-nucleotide characters. You can eye-ball your file or ...

```
> grep -v "^>" seqdump.fasta | grep --color -n "[^ATGC]"
```

- First we remove (with `-v` option) lines that start with a ">" character (these are sequence identifiers).
- Next we find characters that are *not* A, T, C or G. To do this we use `^` symbols second meaning: *match anything but* the pattern in square brackets. We also print line number (with `-n` option) and color output (with `--color` option).

Key Point: Pipes and redirects avoid unnecessary i/o

- Disc i/o is often a bottleneck in data processing!
- Pipes prevent unnecessary disc i/o operations by connecting the stdout of one process to the stdin of another (these are frequently called "**streams**")

```
> program1 input.txt 2> program1.stderr | \
program2 2> program2.stderr > results.txt
```

- Pipes and redirects allow us to build solutions from modular parts that work with **stdin** and **stdout streams**.

Unix 'Philosophy' Revisited

"Write programs that do one thing and do it well. Write programs to work together and that encourage open standards. **Write programs to handle text streams, because that is a universal interface.**"



— Doug McIlroy

Pipes provide speed, flexibility and sometimes simplicity...

- In 1986 "*Communications of the ACM magazine*" asked famous computer scientist Donald Knuth to write a simple program to count and print the k most common words in a file alongside their counts, in descending order.
- Knuth wrote a literate programming solution that was 7 pages long, and also highly customized to this problem (e.g. Knuth implemented a custom data structure for counting English words).

- Doug McIlroy replied with one line:

```
> cat input.txt | tr A-Z a-z | sort | uniq -c | sort -rn | sed 10q
```

Key Point:

You can chain any number of programs together to achieve your goal!



This allows you to build up fairly complex workflows within one command-line.

Do it Yourself!

Hands-on time

Section 4 of software carpentry UNIX lesson

<https://swcarpentry.github.io/shell-novice/>

~15 mins

Shell scripting

Do it Yourself!

```
#!/bin/bash
# This is a very simple hello world script.
echo "Hello, world!"
```

Exercise:

- Create a "Hello world"-like script using command line tools and execute it.
- Copy and alter your script to redirect output to a file using `>` along with a list of files in your home directory.
- Alter your script to use `>>` instead of `>`. What effect does this have on its behavior?

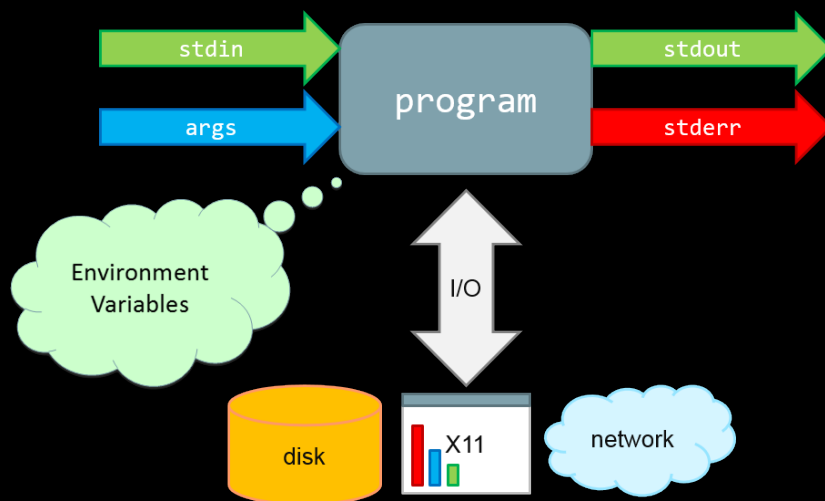
Variables in shell scripts

Do it Yourself!

```
#!/bin/bash
# Another simple hello world script
message='Hello World!'
echo $message
```

- "message" - is a **variable** to which the string 'Hello World!' is assigned
- echo - prints to screen the contents of the variable "\$message"

Side-Note: Environment Variables



\$PATH 'special' environment variable

- What is the output of this command?
 - > echo \$PATH
- Note the structure: <path1>:<path2>:<path3>
- **PATH** is an environmental variable which Bash uses to search for commands typed on the command line without a full path.
- *Exercise:* Use the command **env** to discover more.

Summary

- Built-in unix shell commands allow for easy data manipulation (e.g. sort, grep, etc.)
- Commands can be easily combined to generate flexible solutions to data manipulation tasks.
- The unix shell allows users to automate repetitive tasks through the use of shell scripts that promote reproducibility and easy troubleshooting
- Introduced the 21 key unix commands that you will use during ~95% of your future unix work...

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ls	mv	less	chmod	grep	top
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man	rm	nano	curl	sudo	Ctrl-c
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	> (write to file)		cat	R	bg
	< (read from file)			python	fg

How to Get Working

1) Connecting to jetstream

2) Best practices for organizing your computational biology projects

Read: Noble *PLoS Comp Biol* (2009)

- "A Quick Guide to Organizing Computational Biology Projects"

<http://journals.plos.org/plocompbiol/article?id=10.1371/journal.pcbi.1000424>

All files and directories used in your project should live in a single project directory.

- Use sub-directories to divide your project into sub-projects.
- Do not use spaces in file and directory names!

Document your methods and workflows with plain text README files

- Also document the origin of all data in your project directory
- Also document the versions of the software that you ran and the options you used.
- Consider using Markdown for your documentation.

Use version control and backup to multiple destinations!

Be reproducible:

<http://ropensci.github.io/reproducibility-guide/sections/introduction/>