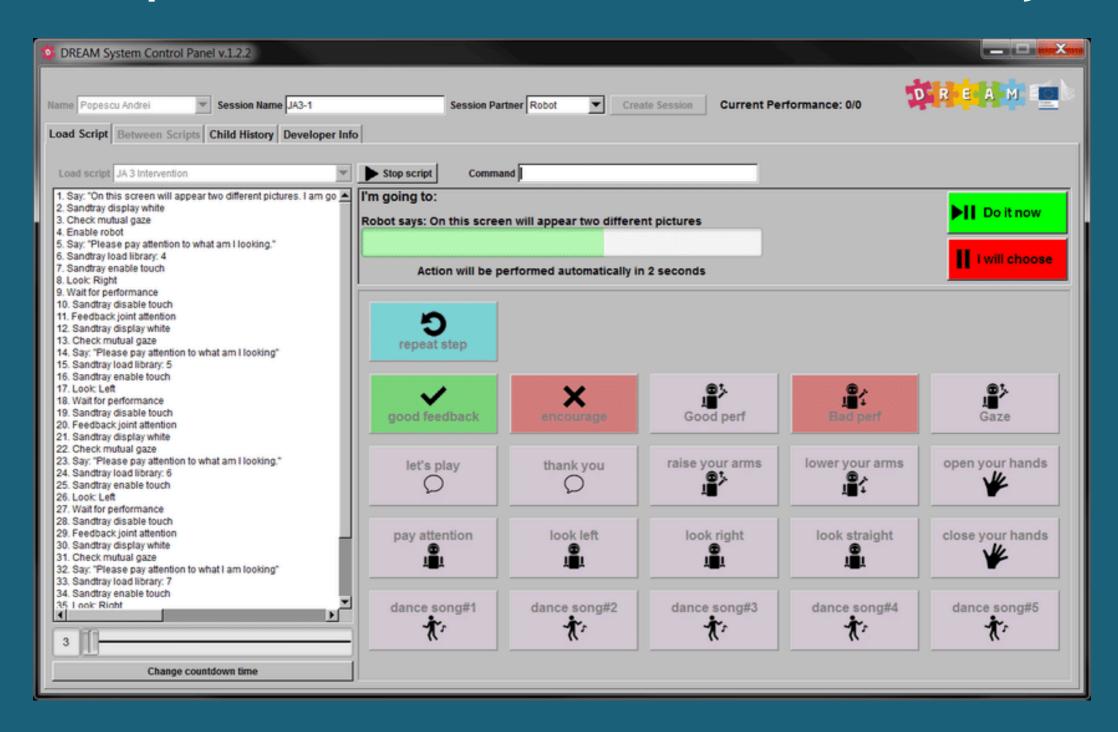
Unix tips and tricks

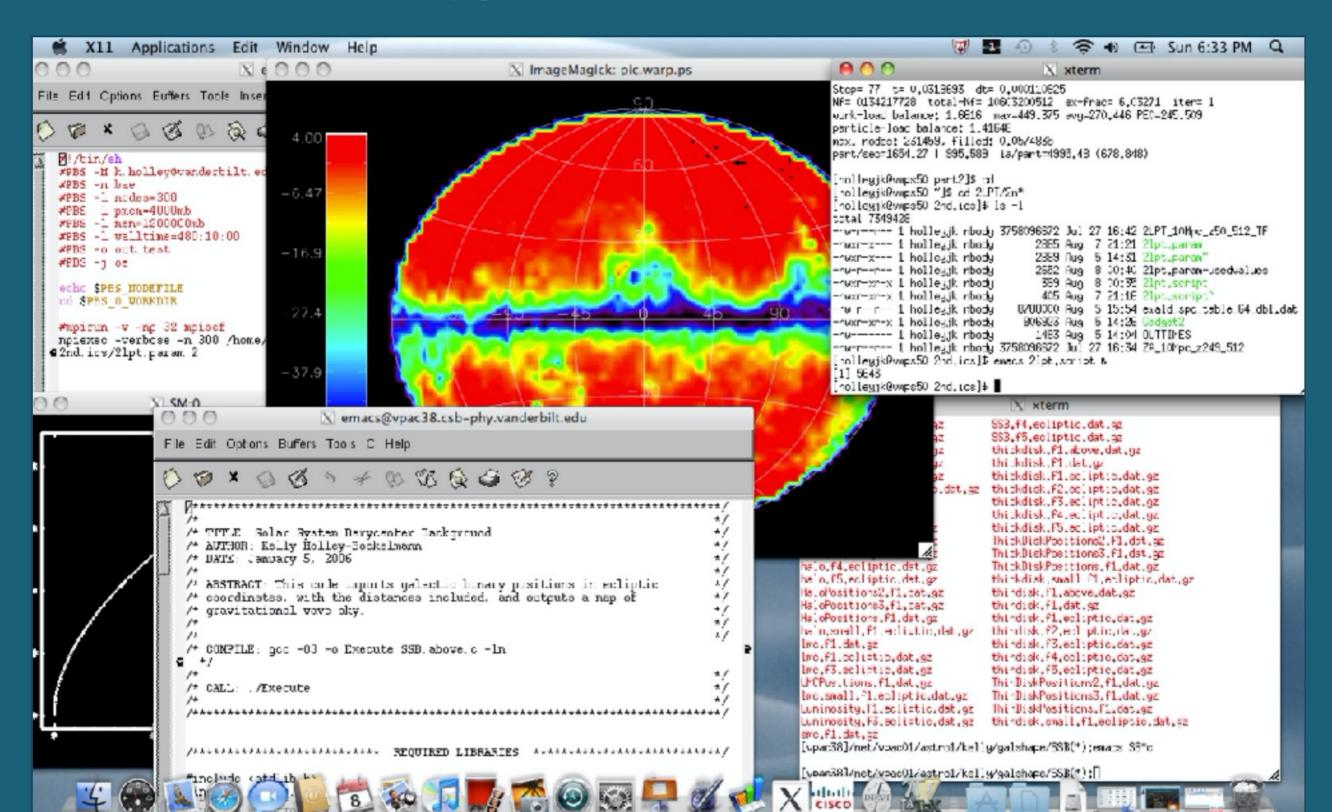


Most computational science is not done by GUI



GUI: Graphical User Interface

A typical science session



Scientific workstations use a UNIX-based operating system







Why UNIX?

- Multitasking
- Multiuser
- Access data stored in another computer
- Able to store large data sets

Program whose primary purpose is to read commands and run other programs.

Pros: high action-to-keystroke ratio, automating repetitive tasks, accessing network machines

Cons: Cryptic commands

It all starts with a\$

If the shell can't find aprogram whose name is the command you typed the error message will read 'command not found'

What it looks like:

```
vivek@nixcraft:/tmp$ vi hello.sh
vivek@nixcraft:/tmp$
vivek@nixcraft:/tmp$ chmod +x hello.sh
vivek@nixcraft:/tmp$
vivek@nixcraft:/tmp$ ls -l hello.sh
-rwxr-xr-x 1 vivek vivek 31 Jan 21 15:08 hello.sh
vivek@nixcraft:/tmp$
vivek@nixcraft:/tmp$ ./hello.sh
Hello World
vivek@nixcraft:/tmp$
vivek@nixcraft:/tmp$ bash hello.sh
Hello World
vivek@nixcraft:/tmp$
vivek@nixcraft:/tmp$ sh hello.sh
Hello World
vivek@nixcraft:/tmp$ cat hello.sh
#!/bin/bash
echo "Hello World"
vivek@nixcraft:/tmp$
```

Bash shell online

https://tinyurl.com/terminalproject2021

Or

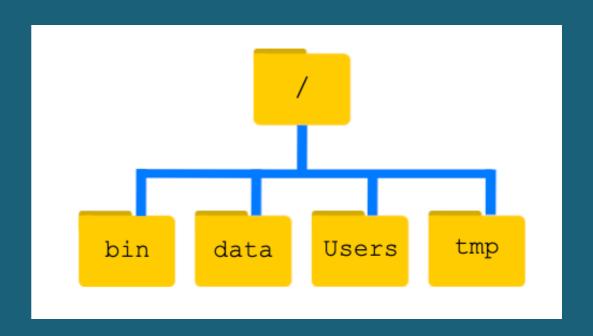
- Open Terminal on your Mac
- Download shell-lesson-data.zip
 - https://tinyurl.com/shelldata
- Save it on your Desktop

/ is the root dir

Otherwise it's apath separator (opposite of Windows)

Absolute (from root) vs relative path (from current location)

.. & . are special dirs.



Getting around the directory tree

- Is → lists content of directory
- pwd → print working directory
- · cd 'blah' change directory to 'blah'
- mkdir 'blah' → make directory 'blah'
- rmdir 'blah' → remove directory 'blah'

Practice: What directory are you in?
Practice: If on your computer, navigate to your Desktop directory
Practice: make a directory called 'Hello_World'
Practice: Go inside your new directory

*Hint use tab completion *

Feel the power: adding tags to your command can customize it.

ls → lists content of directory

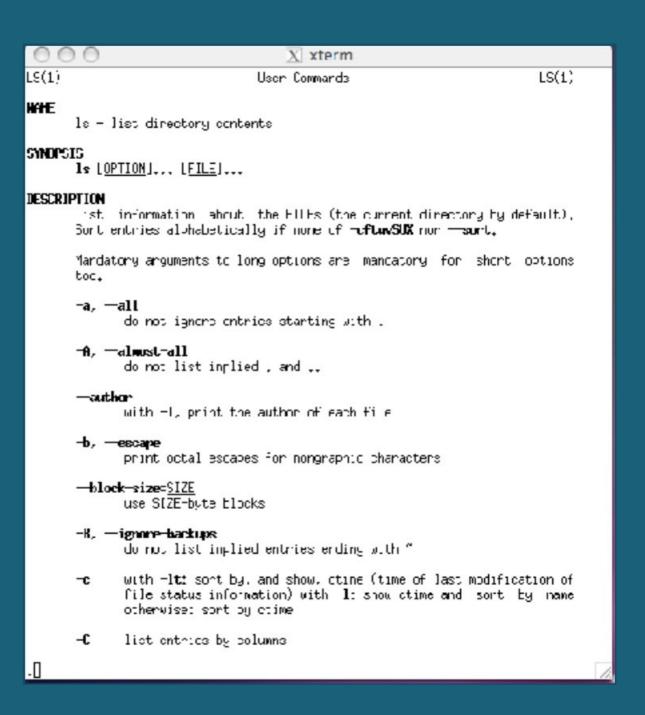
```
ls -l long format
ls -a
ls -lhS
ls -lhS
ls -F
```

```
total 101M
-rw-r--r-- 1 holleyjk astro 58M May 17 10:30 hdf5-1.8.4-patch1.tar
-rw------ 1 holleyjk astro 6.7M May 17 11:11 king_snap.dat
-rw------ 1 holleyjk astro 6.3M Aug 10 2009 mri.pdf
-rw------ 1 holleyjk astro 3.4M Jul 21 18:24 100719_Yaqiongcareerproposal.doc
-rw------ 1 holleyjk astro 2.6M Jan 1 2010 merger_44_20_2366.png
```

But how do I know the secret tags for each command?

man command

man intro is a good place to start!



Key Points

- •The file system is responsible for managing information on the disk.
- •Information is stored in files, which are stored in directories (folders).
- •Directories can also store other directories, which then form a directory tree.
- •cd [path] changes the current working directory.
- •Is [path] prints a listing of a specific file or directory; Is on its own lists the current working directory.
- •pwd prints the user's current working directory.
- on its own is the root directory of the whole file system.

Key Points

- Most commands take options (flags) that begin with a -
- •A relative path specifies a location starting from the current location.
- An absolute path specifies a location from the root of the file system.
- Directory names in a path are separated with / on Unix, but \ on Windows.
- •.. means 'the directory above the current one'
- on its own means 'the current directory'.

Making and adding files

With Terminal you can write scripts, documents using vim*

Practice: On you bash shell type vim Hello.txt

You should have an empty text document

- To write using vim.
- Hit i on your keyboard (insert)
- Type your string Hello world
- Hit esc on your keyboard to stop writing
- Hit :wq on your keyboard to write and quite the text file
- Repeat the steps and make a empty text file
- Back in bash (\$) see if your text files are there with Is

*Vim is a text editor, people both love it and hate it.

If you have a prefer text editor feel free to use that one.

```
~$ pwd
/home/user Current directory
-$ mkdir 'Hello World' Make directory
-$ ls List things inside directory. Hello_World is bold because it's a directory
Hello_World 'Welcome to CoCalc.term'
-$ cd Hello World/ Change to the Hello_World directory
~/Hello World$ vim hello.txt Create hello.txt
~/Hello World$ vim hello.txt
~/Hello_World$ cd .. Go back one directory
~$ ls
Hello World 'Welcome to CoCalc.term'
```

Manipulating files

- cp file1 file2 → copy 'file1' to 'file2'
- mv file1 file2 → move 'file1' to 'file2'
- rm 'blah' → remove file 'blah'
 - *Terminal has no trash bin!

- means /home/usemame/
- means the current directory
- ... means go back 1 directory
 - * means wildcard

Copy a file

Practice: Copy a file

- To copy hello_world.
 - cp hello_world.txt hello_world_2.txt
 - or
 - cp hello_world.txt [path/filename]
- Are the files the same?

Move a file

Practice: Move a file

- To move hello_world up one directory
 - mv hello_world.txt ...
 - or
 - mv hello_world.txt [path/filename]
 - What is the new path for hello_world.txt?

Copy a directory

Practice: Copy a directory

- To copy Hello_world
 - cp –r Hello_world Hello_world_backup

We can also copy a directory and all its contents by using the <u>recursive</u> option -r

- List both directories
 - Is Hello_world Hello_world_backup

Move a directory

Practice: Move a directory

- Make a directory called Research_Test inside Hello_World folder
 - mkdir ~/shell-lessons-Crisel-1/Hello_world/Research_Test
 - Inside Hello_world > mkdir Research_Test
- To move Research Test to shell-lessons-Crisel-1
 - mv Research Test/ ~/shell-lessons-Crisel-1/

Remove a file

Practice: Remove a file/Delete a file (forever)

- Go to your Hello_World folder
- Remove hello world 2.txt
- rm hello_world_2.txt
- List all files
- Is hello_world_2.txt in your directory?
- What happened if you try:
 - rm -i Hello world/hello world.txt

Remove a directory

Practice: Remove a directory and everything inside it

- Remove your Hello_World directory safely
 - rm -ri Hello_World/
- List the items in your directory and make sure you don't have the Hello_World folder

Operations with multiple files and directories

Practice: Go to shell-lesson-data-YourName/molecules/

- * Wildcard matches characters
- List all the files inside the molecules/ directory

Try it out: Which command(s) will produce this output? ethane.pdb methane.pdb

```
Is *t*ane.pdb
Is *t?ne.*
Is *t??ne.pdb
Is ethane.*
```

Key Points

- •cp [old] [new] copies a file.
- •mkdir [path] creates a new directory.
- •mv [old] [new] moves (renames) a file or directory.
- •rm [path] removes (deletes) a file.
- •* matches zero or more characters in a filename, so *.txt matches all files ending in .txt.
- •? matches any single character in a filename, so ?.txt matches a.txt but not any.txt.
- The shell does not have a trash bin
- •Most files' names are something.extension. The extension is used to indicate the type of data in the file.

Practice: Go to shell-lesson-data-YourName/molecules/

- Word Count (wc)
- •\$ wc cubane.pdb
 - •20 156 1158 cubane.pdb
 - Number of lines, words, and characters in files
- •Practice:
- •wc *.pdb
- •wc -l *.pdb
- Save output
 - •wc -I *.pdb > lengths.txt

Check: Did you saved lengths.txt?

- Peek inside lengths.txt with:
 - cat -'concatenate' i.e. join together, and it prints the contents of files one after another.
 - head displays only the first 10 lines of the file
 - less screenful of the file, and then stops. You can go forward one screenful by pressing the spacebar, or back one by pressing b. Press q to quit.

Check: Sort

• Sort lengths.txt with:

•\$ sort -n lengths.txt > sorted-lengths.txt

• \$ head **-n** 1 sorted-lengths.txt

Check: echo prints statement

- \$ echo The echo command prints text
 - The echo command prints text

Check: > vs >>

- \$ echo hello > testfile01.txt
- \$ echo hello >> testfile02.txt
- > 'hello' is written to testfile01.txt, but the file gets overwritten each time we run the command.
- >> 'hello' is written to testfile01.txt, but the file gets overwritten each time we run the command.

Check: More commands in one line (pipe)

- \$ sort -n lengths.txt | head -n 1
- \$ wc -**I** *.pdb | sort -**n**
- \$ wc -I *.pdb | sort -n | head -n 1

```
$ wc -1 *.pdb our

Output in Shell

$ wc -1 *.pdb our

lengths
Output in File

$ wc -1 *.pdb our

lengths
Output in File

$ wc -1 *.pdb our

Note the second output in Shell

Output in Shell

Output in Shell
```

Key Points

- wc counts lines, words, and characters in its inputs.
- cat displays the contents of its inputs.
- sort sorts its inputs.
- head displays the first 10 lines of its input.
- tail displays the last 10 lines of its input.
- command > [file] redirects a command's output to a file (overwriting any existing content).

Key Points

- command >> [file] appends a command's output to a file.
- [first] | [second] is a pipeline: the output of the first command is used as the input to the second.
- The best way to use the shell is to use pipes to combine simple single-purpose programs (filters).

Loops

Check: More commands in one line (pipe)

- \$ sort -n lengths.txt | head -n 1
- \$ wc -**I** *.pdb | sort -**n**
- \$ wc -I *.pdb | sort -n | head -n 1

```
$ wc -1 *.pdb out

Output in Shell

$ wc -1 *.pdb > lengths

wc -1 *.pdb out

lengths
Output in File

$ wc -1 *.pdb | sort -n | head -n 1

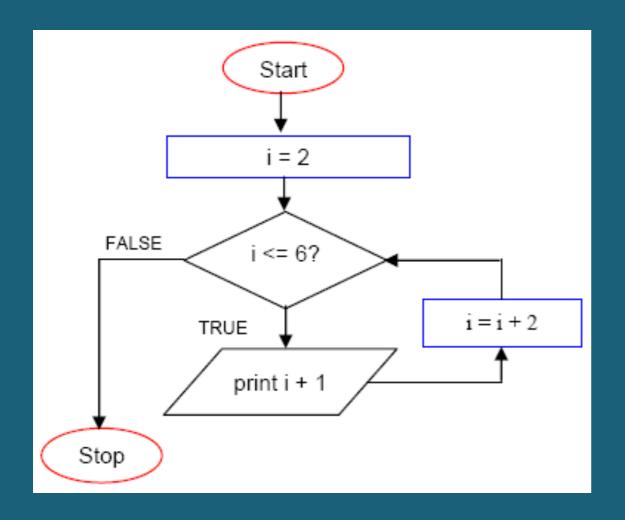
wc -1 *.pdb out | sort -n | out | head -n 1 out

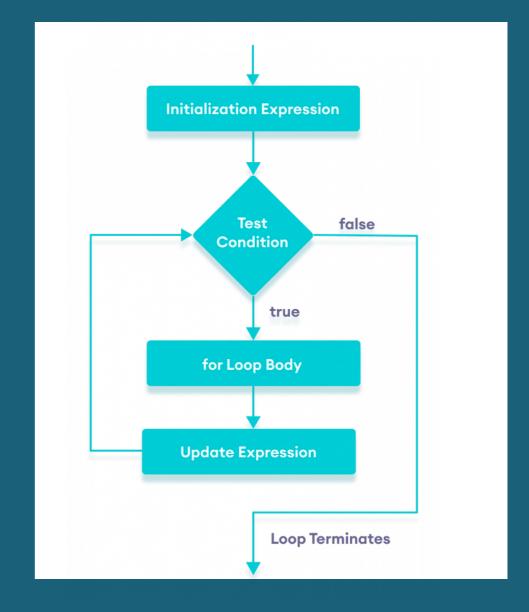
Output in Shell
```

Loops

Loops are a programming construct which allow us to repeat a command or set of commands for each item in a list. As such they are key to productivity improvements through

automation





Loops

```
$for thing in list_of_things
do
operation_using $thing
done
```

Practice: Go to shell-lesson-data-YourName/creatures/

```
$ for name in basilisk.dat minotaur.dat unicorn.dat
```

- >do
 - > head -n 2 \$name | tail -n 1
- > done

COMMON NAME: basilisk

COMMON NAME: minotaur

COMMON NAME: unicorn

Loops

```
$for thing in list_of_things
do
operation_using $thing
done
```

Practice: Go to shell-lesson-data-YourName/creatures/

```
$ for filename in *.dat
> do

cp $filename original-$filename
> done
```

Key Points

- A for loop repeats commands once for every thing in a list.
- Every for loop needs a variable to refer to the thing it is currently operating on.
- Use \$name to expand a variable (i.e., get its value).
- Do not use spaces, quotes, or wildcard characters such as '*' or '?' in filenames, as it complicates variable expansion.
- Give files consistent names that are easy to match with wildcard patterns to make it easy to select them for looping.

Bash scripts

How can I save and re-use commands?

Practice: Go to shell-lesson-data-YourName/molecules/

With Terminal you can write scripts, documents using vim*

Practice: Type vim middle.sh

You should have an empty text document

- To write using vim.
- Hit i on your keyboard (insert)
- Type your code head -n 15 octane.pdb | tail -n 5
- Hit esc on your keyboard to stop writing
- Hit :wq on your keyboard to write and quite the text file
- Back in bash (\$) bash middle.sh

Bash scripts

How can I save and re-use commands?

Practice: Type more general script vim middle.sh

- To write using vim.
- Hit i on your keyboard (insert)
- Type your code head -n 15 "\$1" | tail -n 5
 - Previously (head -n 15 octane.pdb | tail -n 5)
- Hit esc on your keyboard to stop writing
- Hit :wq on your keyboard to write and quite the text file
- Back in bash \$ bash middle.sh octane.pdb
- Back in bash \$ bash middle.sh pentane.pdb

Bash scripts

How can I save and re-use commands?

Practice: Type more general script vim script.sh

- To write using vim.
- Hit i on your keyboard (insert)
- Type your code
 - head -n \$2 \$1
 - tail -n \$3 \$1
- Hit esc on your keyboard to stop writing
- Hit :wq on your keyboard to write and quite the text file
- Back in bash \$ bash script.sh '*.pdb' 1 1

Key Points

- Save commands in files (usually called shell scripts) for re-use.
- bash [filename] runs the commands saved in a file.
- \$1, \$2, etc., refer to the first command-line argument, the second command-line argument, etc.

Practice: Go to shell-lesson-data-YourName/writing/

- Look inside haiku.txt
- Which lines have contain 'not'?
- grep (find)
- \$ grep not haiku.txt
 - Is not the true Tao, until
 - "My Thesis" not found
 - Today it is not working

Practice: Go to shell-lesson-data-YourName/writing/

- \$ grep The haiku.txt
- \$ grep —w The haiku.txt
- \$ grep -w "is not" haiku.txt

- \$ grep -i => case insensitive
- \$ grep -n => number lines
- \$ grep -v => invert search
- \$ grep -r => recursive search
- \$ grep - help (no space)

Practice: Lets search recursively through the whole directory shell-lesson-data-YourName/writing/

\$ grep -r Yesterday -

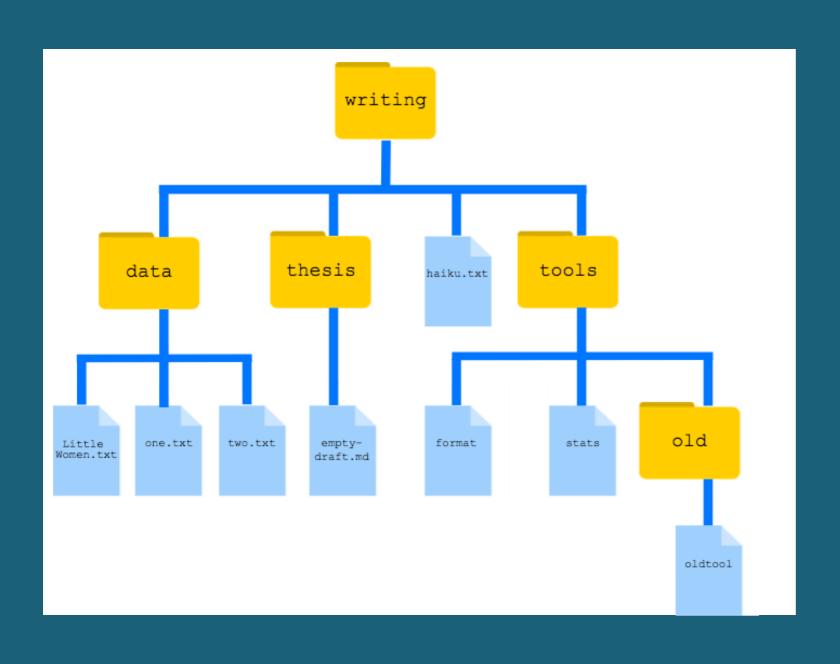
- data/LittleWomen.txt:"Yesterday, when Aunt was asleep and I was trying to be as still as a
- data/LittleWomen.txt:Yesterday at dinner, when an Austrian officer stared at us and then
- data/LittleWomen.txt:Yesterday was a quiet day spent in teaching, sewing, and writing in my
- haiku.txt:Yesterday it worked

Practice: Go to shell-lesson-data-YourName/writing/

- \$ grep -E "^.o" haiku.txt
- We use the -E option and put the pattern in quotes to prevent the shell from trying to interpret it.
- The ^ in the pattern anchors the match to the start of the line.
- The matches a single character
- o matches an actual 'o'.



Practice: Go to shell-lesson-data-YourName/writing/



\$ find.

\$ find . —type d

\$ find . —type f

\$ find . -name "*.txt"





Some more commands

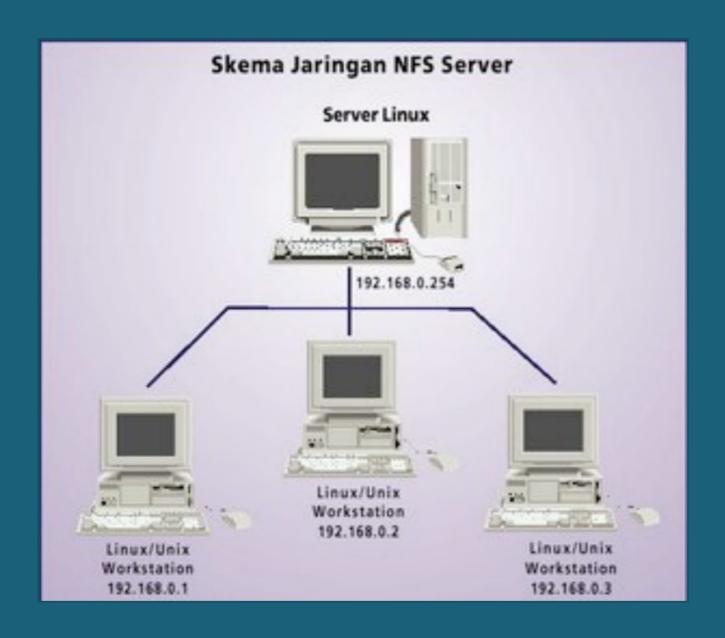
- rm -rf for directories (careful!)
- Maybe make rm interactive using rm -i in ~/.aliases
- history
- Unpack files from arXiv : tar -xvf <filename>

Translate this:

```
· cp ~/*.sm ../.
```

· mv ../density.sm ~/bootcamp/dense.s

• rm *

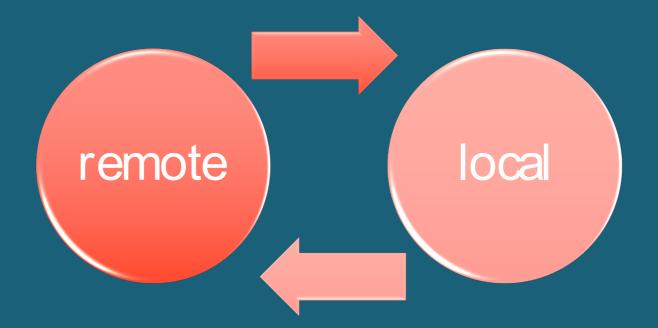


When you ssh into the vpac/accre network, you go to your home directory.

Your directory can be accessed through any of these network machines.

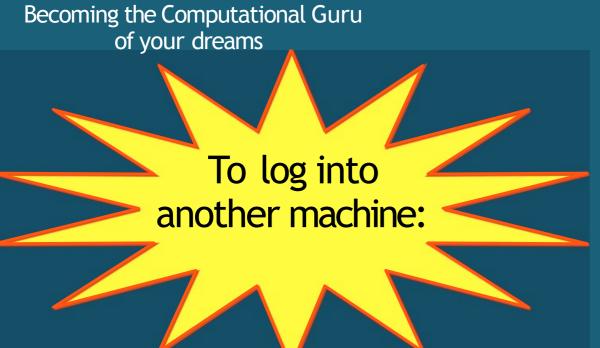
And, you can (theoretically) access data on other user's accounts, too.

SCP secure copy protocol



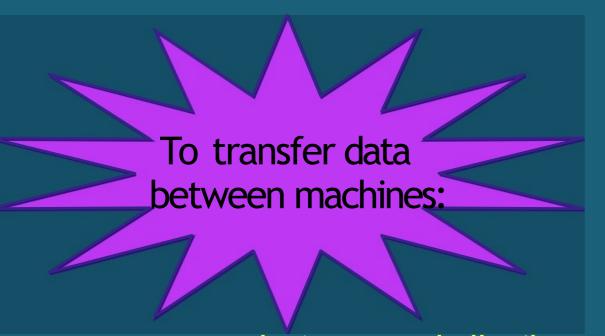
~ stands for home dir (make sure you're in the right place!)

Use * wildcard to grab multiple files that meet some condition



ssh -YC username@computer's whole address

example: ssh -YC holleyjk@vpac03.phy.vanderbilt.edu



scp -pr complete path of file you want to move complete path describing where you want to put it

example 1: scp -pr holleyjk@vpac03.phy.vanderbilt.edu:/home/holleyjk/superfile/Users/kelly/Desktop/superfile.moo

example 2: scp -pr /Users/kelly/awesomefile holleyjk@vmplogin.accre.vanderbilt.edu:/home/holleyjk/.

Your data is protected by file permissions

Aside: How did I get this list?

```
X xterm
             1 holleyjk astro
                                 28672 Oct 14 2009 sesaps 09 program.xls
             3 holleyjk astro
                                 4096 May 17 10:42 share
                                 14176 Jul 6 16:42 sig.khb.jpeg
            1 holleyjk astro
            1 holleyjk astro
                                13717 Jul 6 16:46 sig.khb.pdf
                                21470 Jun 10 14:56 sinha_69.pdf
            1 holleyjk astro
            1 hollewik astro
                                230880 May 28 15:08 SMD-10-1678_Holley-Bokelmann
.pdf
             4 holleuik astro
                                              2009 smooth, 2,01
                                 4096 May 13
            1 holleyjk astro
                                407473 Jan 25 2010 starcounts.dat.gz
            1 holleyjk astro
                                407478 Jan 23 2010 starcounts.orig.dat.gz
            1 holleyjk astro
                                 79428 Jun 15 13:30 stassun_career_report_2009.p
            1 holleyjk astro
                                     8 Jun 1 14:16 TAR.proposal.aux
                                  7436 Jun 1 14:16 TAR.proposal.dvi
             1 holleyjk astro
             1 holleyjk astro
                                  3674 Jun 1 14:16 TAR.proposal.log
            1 holleyjk astro
                                 26545 Jun 1 14:16 TAR.proposal.pdf
                                 72587 Jun 1 14:16 TAR.proposal.ps
             1 holleyjk astro
            1 holleyjk astro
                                 6086 Jun 1 14:16 TAR.proposal.tex
            6 holleyjk astro
                                 4096 Sep 21 2007 tibs2107
             5 holleyjk astro
                                 4096 Feb 2 2009 tipsy-2.2.3c
             2 holleyjk astro
                                 4096 Mar 27
                                              2008 TRS
                                 4096 Har 5 2008 web
             3 holleyjk astro
                                 33280 Jul 21 18:23 YaqiongSummary.doc
            1 holleyjk astro
           15 holleyjk astro
                                 4096 May 17 16:48 yt
drwxr-xr-x
                                798679 Jun 14 11:03 Zampieri, 2009, MNRAS, 400, 677,
            1 holleyjk astro
-rw-----
pdf
drwxr-xr-x 12 holleyjk astro
                                  4096 May 17 10:42 zlib-1.2.5
```

user
permissions
r=Read
w=Write
x=eXecute

Your data is protected by file permissions

```
X xterm
             1 holleyjk astro
                                 28672 Oct 14 2009 sesaps 09 program.xls
             3 holleyjk astro
                                 4096 May 17 10:42 share
                                 14176 Jul 6 16:42 sig.khb.jpeg
            1 holleyjk astro
            1 holleyjk astro
                                13717 Jul 6 16:46 sig.khb.pdf
                                 21470 Jun 10 14:56 sinha_69.pdf
            1 holleyjk astro
            1 hollewik astro
                                230880 May 28 15:08 SMD-10-1678_Holley-Bokelmann
.pdf
             4 holleuik astro
                                              2009 smooth, 2,01
                                  4096 May 13
            1 holleyjk astro
                                407473 Jan 25
                                              2010 starcounts.dat.gz
            1 holleyjk astro
                                407478 Jan 23 2010 starcounts.orig.dat.gz
            1 holleyjk astro
                                 79428 Jun 15 13:30 stassun_career_report_2009.p
             1 holleyjk astro
                                     8 Jun 1 14:16 TAR.proposal.aux
                                  7436 Jun 1 14:16 TAR.proposal.dvi
             1 holleyjk astro
             1 holleyjk astro
                                  3674 Jun 1 14:16 TAR.proposal.log
             1 holleyjk astro
                                 26545 Jun 1 14:16 TAR.proposal.pdf
             1 holleyjk astro
                                 72587 Jun 1 14:16 TAR.proposal.ps
            1 holleyjk astro
                                          1 14:16 TAR.proposal.tex
                                  6086 Jun
             6 holleyjk astro
                                  4096 Sep 21 2007 tibs2107
             5 holleyjk astro
                                  4096 Feb 2 2009 tipsy-2.2.3c
             2 holleyjk astro
                                  4096 Mar 27
                                               2008 TRS
                                  4096 Mar 5
                                              2008 web
             3 holleyjk astro
                                 33280 Jul 21 18:23 YaqiongSummary.doc
            1 holleyjk astro
                                  4096 May 17 16:48 yt
           15 holleyjk astro
drwxr-xr-x
                                798679 Jun 14 11:03 Zampieri, 2009, MNRAS, 400, 677,
            1 holleyjk astro
-rw-----
pdf
drwxr-xr-x 12 holleyjk astro
                                  4096 May 17 10:42 zlib-1.2.5
```

group permissions r=Read w=Write x=eXecute

Your data is protected by file permissions

```
X xterm
             1 holleyjk astro
                                 28672 Oct 14 2009 sesaps 09 program.xls
             3 holleyjk astro
                                 4096 May 17 10:42 share
                                 14176 Jul 6 16:42 sig.khb.jpeg
             1 holleyjk astro
             1 holleyjk astro
                                 13717 Jul 6 16:46 sig.khb.pdf
                                 21470 Jun 10 14:56 sinha_69.pdf
             1 holleyjk astro
             1 hollewik astro
                                230880 May 28 15:08 SMD-10-1678_Holley-Bokelmann
.pdf
             4 holleuik astro
                                               2009 smooth, 2,01
                                  4096 May 13
            1 holleyjk astro
                                407473 Jan 25
                                               2010 starcounts.dat.gz
             1 holleyjk astro
                                407478 Jan 23 2010 starcounts.orig.dat.gz
             1 holleyjk astro
                                 79428 Jun 15 13:30 stassun_career_report_2009.p
             1 holleyjk astro
                                     8 Jun 1 14:16 TAR.proposal.aux
                                  7436 Jun 1 14:16 TAR.proposal.dvi
             1 holleyjk astro
             1 holleyjk astro
                                  3674 Jun 1 14:16 TAR.proposal.log
             1 holleyjk astro
                                 26545 Jun 1 14:16 TAR.proposal.pdf
                                 72587 Jun 1 14:16 TAR.proposal.ps
             1 holleyjk astro
             1 holleyjk astro
                                           1 14:16 TAR.proposal.tex
                                  6086 Jun
             6 holleyjk astro
                                  4096 Sep 21 2007 tibs2107
             5 holleyjk astro
                                  4096 Feb 2
                                              2009 tipsy-2.2.3c
             2 holleyjk astro
                                               2008 TRS
                                  4096 Mar 27
                                               2008 web
             3 holleyjk astro
                                  4096 Mar
                                 33280 Jul 21 18:23 YaqiongSummary.doc
             1 holleyjk astro
           15 holleyjk astro
                                  4096 May 17 16:48 yt
drwxr-xr-x
                                798679 Jun 14 11:03 Zampieri, 2009, MNRAS, 400, 677,
            1 holleyjk astro
-rw-----
pdf
drwxr-xr-x 12 holleyjk astro
                                  4096 May 17 10:42 zlib-1.2.5
```

other
permissions
r=Read
w=Write
x=eXecute

To change permissions:

chmod ugo+/-rwx filename

Example: chmod go+r pro*.dat

Advanced: chgrp -R <group> <directory>

changes the group ownership of a directory (R=recursively). Have to be admin or be member of both groups

Taking a peek at a file:

- head -n50 file → displays first 50 lines of file
- tail -n12 file → displays last 12 lines of file
- more file →scrolls through file page by page Enter to scroll line by line or Spacebar to scroll "page by page"
- less file →a better version of more

Practice: figure out what is the last line in ~/alias.holleyjk

Is the file compressed? gunzip file

Looking for a file? Try find .-iname file

What jobs are running in your shell? ps ...on your machine? htop

Need to kill a job? kill -9 PID

Need to count lines in a file? wc -l file

Smoosh files into 1 cat file1 file2 > smooshfile

Split file into many of N bytes split -b N file

String commands together with pipes:

try these:

ls -l /home/holleyjk |less

ps aux |grep username

the possibilities are huge!

- 1. list all the files in the directory
- 2. list them by modification time, most recent first
- 3. display that list in human readable format
- 4. display only the top 5 in that list

- 1. list all the files in the directory ls -al
- 2. list them by modification time, most recent first
- 3. display that list in human readable format
- 4. display only the top 5 in that list

- 1. list all the files in the directory ls -al
- 2. list them by modification time, most recent first ls -alt
- 3. display that list in human readable format
- 4. display only the top 5 in that list

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Q. List the first 10 most space-hogging files in a directory in human readable format

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ls -halS |head -n11 |tail -n10

- last tells you the people who logged in, arranged chronologically
- w shows you who are logged in currently
- date gives you the current date (time-stamp)
- du shows you the disk usage of given directory (and sub-dirs)
- sort sorts data file/piped output based on specified column
- diff finds differences between two files
- > redirects result to a file
- >> appends data to a file
- history returns history of commands executed in terminal ping attempts to open aline of communication with anetwork host
- wget download afile from the Internet

Exercises:

- 1. Who are the last 5 people that logged in to vpac38 today?
- 2. How many times did 'hoffmare' login to vpac01?
- 3. Copy wang.merritt.dat from the 2015 bootcamp directory to your laptop and sort on the black hole mass. Dump results to a file.
- 4. How many people are on vpac02 right now?
- 5. How many bytes are you using on your home directory?
- 6. How many files does Kelly have in her \$HOME/bootcamp dir?

Exercises:

- 1. Who are the last 5 people that logged in to vpac38 today? first, ssh into vpac38 then... last head -n5
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- 2. How many times did 'hoffmare' login to vpac01? first, ssh into vpac01, then...

last |grep hoffmare > dummy; wc -l dummy

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- 3. Copy wang.merritt.dat from the 2015 bootcamp directory to your laptop and sort on the black hole mass. Dump results to a file.

sort -n -k2,2 < wang.merritt.dat > sorted.wang.merritt.dat

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w sort

- 5. How many bytes are you using on your home directory?
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du -h

6. How many files does Kelly have in her \$HOME/bootcamp dir?

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ls -a |wc -l

Setting your shell (how you interface with your operating system)

--More--(91%)

do a less on cshrc.holleyjk

execute an alias file that nicknames unix commands

tells your system where to look for external commands, like graphics programs

```
X xterm
 .cshrc
set listjobs
limit coredumpsize 0
umask 022
unset autologout
 if ( -e "/_alias )
                        source "/.alias
if( $?REMOTEHOST && ! $?DISPLAY ) then
    setenv DISPLAY ${REMOTEHOST}:0
endif
#set path=(/usr/local/products/mpich2/bin $path)
# pattans 20080123
set osv = `uname -a | awk '{print $1 substr ($3,1,1)}'`
set arch = `uname -m
setenv MPICH_HOME /usr/local/mpich2/${osv}/1.0.6p1/${arch}
setenv MPI_HOME /usr/local/mpich2/${osv}/1.0.6p1/${arch}
#setenv MPICH_HOME /usr/local/products/mpich2
#setenv MPI_HOME /usr/local/products/mpich2
setenv PATH /usr/local/bin:/usr/bin:/home/holleyjk/bin:/usr/local/python2.6/x86_64/
bin:/usr/local/jdk:/usr/local/lib/jdk/jdk1.6.0_01/bin:/usr/local/mpich2/Linux2/1.0.
6p1/x86_64/bin:/usr/lib64/qt-3.3/bin:/usr/kerberos/bin:/bin:/usr/X11R6/bin
setenv CLASSPATH /home/holley.jk/lstore/accre-loci-1.0-1802.jar:/home/holley.jk/lstor
e/aspectjrt.jar:/home/holleyjk/lstore/jargs.jar:/home/holleyjk/lstore/log4j-1.2.12.
setenv LD_LIBRARY_PATH /usr/lib:${HOME}/lib:/usr/local/lib:/usr/local/python2.6/x86
_64/lib:/usr/local/jdk/lib:/usr/local/lib/jdk/jdk1.6.0_01/lib:/usr/local/mpich2/Lin
ux2/1.0.6p1/x86_64/lib:/usr/include/kde/arts/gsl:/usr/lib64/qt-3.3/lib:/usr/kerbero
s/lib:/lib64:/usr/X11R6/lib:/lib
setenv IDL_DIR /usr/local/itt/idl
```

Setting your shell (how you interface with your operating system)

Open a web browser to github.com/disissom

Go to bootcamp repository

Click "Clone or Download" button, and choose "Download Zip"

Scp the zip file to your vpac home directory

Unzip the file on your laptop (Mac) and on VPAC

Setting your shell (how you interface with your operating system)

What does 'echo \$SHELL' return?

My bashrc to ~/.bashrc (if it doesn't exist yet...)

Less .bashrc to read it

source .bashrc will run the file and install your commands

To change your .bashrc , you need a text editor

emacs <filename> &

- Ctl-v -- page up
- Esc-v -- page down
- · Ctl-n -- next line
- Ctl-p -- previous line
- Esc-< -- top of file
- Esc -> -- bottom of file
- Esc-Esc-Esc -- get out of hotkey
- · Ctl-X-S -- save file
- · Ctl-X-C -- quit

- Ctl-k -- kill a line
- Ctl-y -- paste line
- Esc -% -- query replace
- Ctl-x (-- start macro
- Ctl-x) -- end macro
- Example: Ctl-x (Ctl-n Ctl-x
- Ctl-x e -- execute macro
- Esc-num -- do num times
- Ctl-/ -- undo

Practice: Open a new file called 'motivation' and write 'I love emacs with my whole heart!' 216 times. Then, replace 'heart'

with 'mind' on every other line.

Editing text files with Vim

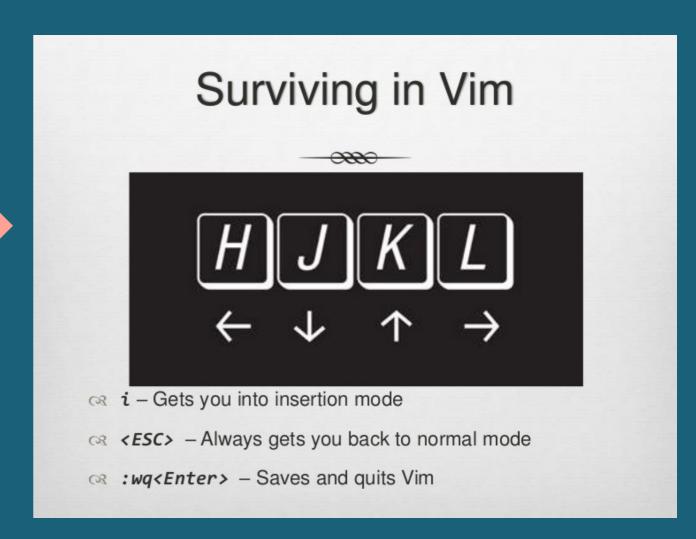
Set up vim files (dark terminal recommended)

\$ vim .bashrc

More slides at https://www.slideshare.net/

brandonliu/introduction-to-vim

/- to search and n/N to navigate search results



Vim walkthrough

https://www.openvim.com/

Remember the cheat sheet

http://cheatsheetworld.com/programming/unix-linux-cheat-sheet/

For homework:

Add your own nicknames for unix commands -- most useful aliases win a prize!

Add aliases to make rm, mv, and cp safer (check the man pages for the interactive option for each)

Practice with your text editor of choice



-- Find a file with cereal in its name

(hint: it's in a vpac directory involving /home/holleyjk...)

-- copy it to your home directory, add your name to the filename, and change permissions so that you can read and write to it, but group/other can only read it

-- write the name of your favorite cereal in the file

-- copy it to Kelly's bootcamp/2017 directory and tell me when you've done it. First one wins a prize!!