

Introduction

Introduction: What is Unix?

- An operating system
- Developed at AT&T Bell Labs in the 1960's
- Command Line Interpreter
- GUIs (Window systems) are now available

Introduction: Unix vs. Linux

- Unix was the predecessor of Linux
- Linux is a variant of Unix
 - So is Mac OS X, so much of this tutorial applies to Macs as well
- Linux is open source
- Most of the machines you'll use in the Bioinformatics program are running the Linux OS

Linux Distribution



Debian

....



Introduction: Why Unix/Linux?

- Linux is free
- It's fully customizable
- It's stable (i.e. it almost never crashes)

 These characteristics make it an ideal OS for programmers and scientists

Connecting to a Unix/Linux system

• Open up a terminal:

	×	● ● ●
zhome:~/linux_tutorial\$		Last login: Mon Jul 17 12:38:08 on ttys002
	•	

Putty

Mac

Mac : Terminal : ssh <u>your_username@bbcsrv3.biotech.uconn.edu</u> Windows : Putty



Open Putty it will open window1.

- 1. Provide host name e.g. ssh <u>your_username@bbcsrv3.biotech.uconn.edu</u>Expand SSH tab and select X11 (shown in window2)
- 2. Enable X11 forwarding by selecting it. (window2)
- 3. Scroll up the left panel and select Session.(window1)
- 4. Name your session e.g. BBC_cluster and click save tab to save.
- 5. Your session name should appear in saved sessions.

Double click on your session name to connect to server with SSH session.

Connecting to a Unix/Linux system

• Open up a terminal:



What exactly is a "shell"?

- After logging in, Linux/Unix starts another program called the **shell**
- The shell interprets commands the user types and manages their execution
 - The shell communicates with the internal part of the operating system called the **kernel**
 - The most popular shells are: tcsh, csh, korn, and bash
 - The differences are most times subtle
 - For this tutorial, we are using bash
- Shell commands are **CASE SENSITIVE!**

Unix/Linux File System

NOTE: Unix file names are CASE SENSITIVE!



Creating files in Unix/Linux

- Requires the use of an Editor
- Various Editors:
 - 1) nano
 - 2) vim
 - 3) emacs



Login into remote server

ssh -X <your_username>@<host_name>

ssh vsingh@bbcsrv3.biotech.uconn.edu

Basic Linux commands

pwd	# Get full path of the present working directory
	(same as "echo \$HOME")
ls	<pre># Content of pwd</pre>
ls -l	<pre># Similar as ls, but provides additional info on</pre>
	files and directories
ls –a	<pre># Includes all files including hidden files</pre>
	(.name) as well
ls -R	<pre># Lists subdirectories recursively</pre>
ls -t	# Lists files in chronological order

cd <dir_na< th=""><th>ame></th></dir_na<>	ame>		
cd	<pre># Brings you to the highest level of your</pre>		
	home directory.		
cd	# Moves one directory up		
cd//	<pre># Moves two directories up (and so on)</pre>		
cd –	# Go back to you were previously (before the last		
	directory change)		
echo ~	# View the full (complete) path of your home		
find ~	<pre># List all your files (including everything in</pre>		
	sub-directories)		
ls ~	# List the top level files of your home directory		

mkdir <dir_name></dir_name>	<pre># Creates specified directory</pre>
rmdir <dir_name></dir_name>	# Removes empty directory
rm <file_name></file_name>	# Removes file name
rm -r <dir_name></dir_name>	<pre># Removes directory including its</pre>
	content, but asks for confirmation, 'f'
	argument turns confirmation off
cp <name> <path></path></name>	<pre># Copy file/directory as specified in</pre>
	path (-r or -a to include content in
	directories)
<pre>mv <name1> <name2></name2></name1></pre>	# Renames directories or files
mv <name> <path></path></name>	<pre># Moves file/directory as specified in path</pre>

man <something> # general help (press the 'q' key to exit) # manual on program 'word count' wc man wc wc --help # short help on wc cat <file1> # Print contents of file1 cat <file1> <file2> > <cat.out> # concatenate files in output file 'cat.out' paste <file1> <file2> > <paste.out> # merges lines of files and separates them by tabs (useful for tables) cmp <file1> <file2> # tells you whether two files are identical diff <fileA> <fileB> # finds differences between two files head -<number> <file> # prints first lines of a file tail -<number> <file> # prints last lines of a file

Basic Linux commands cont..... Files and permission

```
"r" means "read only" permission
"w" means "write" permission
"x" means "execute" permission
In case of directory, "x" grants
permission to list directory contents
```

```
xanadu-submit-ext ~ $ ls -l
total 892
rw-r-rr-
drwxr-xr-x 7 vsingh domain users 257 Jun 2 09:18 .
drwxr-xr-x 7 vsingh domain users 1601 May 2 21:26 aening_masurca
-rw-r--r- 1 vsingh domain users 499 Jun 2 00:39 arrayjob.sh
drwxr-xr-x 2 vsingh domain users 1431 Jun 2 00:39 arrayout
-rwxr-xr-x 1 vsingh domain users 19384 Jun 16 12:07 assemble_xandu2.sh
-rw-r--r- 1 vsingh domain users 649 Jun 14 13:30 basemount_cp.sh
    Other/world (safe: r-x) : o
    Group (safe: r-x) : g
    User : u
File(-) or directory (d)
```

Basic Linux commands cont..... Change permission

If you own the file, you can change it's permissions with "chmod"
 Syntax: chmod [user/group/others/all]+[permission] [file(s)]
 Below we grant execute permission to all:

```
xanadu-submit-ext ~ $ clear
xanadu-submit-ext ~ $ ls -l test.sh
-r-x---- 1 vsingh domain users 484 Jun 16 12:06 test.sh
xanadu-submit-ext ~ $ chmod u+w test.sh
xanadu-submit-ext ~ $ ls -l test.sh
-rwx----- 1 vsingh domain users 484 Jun 16 12:06 test.sh
xanadu-submit-ext ~ $ chmod g+rwx test.sh
xanadu-submit-ext ~ $ ls -1 test.sh
-rwxrwx--- 1 vsingh domain users 484 Jun 16 12:06 test.sh
xanadu-submit-ext ~ $ chmod a+rwx test.sh
xanadu-submit-ext ~ $ ls -l test.sh
-rwxrwxrwx 1 vsingh domain users 484 Jun 16 12:06 test.sh
xanadu-submit-ext ~ $ chmod go+rx test.sh
xanadu-submit-ext ~ $ ls -l test.sh
-rwxrwxrwx 1 vsingh domain users 484 Jun 16 12:06 test.sh
xanadu-submit-ext ~ $ chmod go-w test.sh
xanadu-submit-ext ~ $ ls -l test.sh
-rwxr-xr-x 1 vsingh domain users 484 Jun 16 12:06 test.sh
xanadu-submit-ext ~ $
```

```
Running a program (a.k.a a job)
Make sure the program has executable permissions
chmod u+x program
Use "./" to run the program
./ program
```

INPUT/OUTPUT: STDIN STDOUT and redirecting (Pipes)

```
program_a > file.txt
    program_a's output is written to a file called
    "file.txt"
program_a < input.txt
    program_a gets its input from a file called
    "input.txt"
Programs can output to other programs
Called "piping"
program_a | program_b
program_a's output becomes program_b's input</pre>
```

Exercise 1:

Create a file myscript.sh : touch myscript.sh Check for file : 1s Open file to edit : vi myscript.sh : i : #!/bin/bash Write commands: interpreter : echo "Hello world" : echo "This is my first script in Linux." save and exit : esc-:-w-q-! Check permissions of file : ls -l myscript.sh Change permission if required : chmod u+x myscript.sh Run script/Program : ./myscript.sh : sh myscript.sh Store output to a file : sh myscript.sh > myscript output.txt Check the output (limited) : less myscript output.txt print the contents : cat myscript output.txt Count the number of lines : cat myscript output.txt | wc -1

Transferring files between systems

nload Ailla Client			
•••		FileZilla	
	. 🜔 🎼 🙁 🗽 🐌 🔳 🖉 🔗 🛝		
Host: U	sername: Password: Port:	Quickconnect	
Software_tool_ Tools_articles	comparison		
Filename	Filesize V Filetype Last modified		
	4.905.256 HTML docum 01/10/2017 12:1		
 Turtle_analysis.nb.html 	776,304 HTML docum 04/25/2017 12:1		
results.nb.html	775,856 HTML docum 01/16/2017 15:3	Filename	Filesize Filetype
report.pdf	64,649 pdf-file 01/19/2017 12:1		
RSEM_output.pdf	4/,160 pdf-file 01/16/2017 11:5 46.404 pdf-file 01/05/2017 16:2	Not connected to	any server
.DS_Store	10,244 File 05/04/2017 11:5		
JellyFish.sh	8,808 Shell script 12/23/2016 05:3		
JellyFish.py	8,808 Python Sour 12/23/2016 05:3		
tst.sh	3,675 Shell script 02/01/2017 14:5		
help with R R	2 967 R Source File 01/12/2017 16:3		
results.Rmd	2,819 R Markdown 01/16/2017 15:3		
Turtle_analysis.Rmd	2,819 R Markdown 04/25/2017 12:1		
filter trinity sh	2,216 Shell script 01/16/2017 13:3		
Intel_chinty.sh	1,732 Python Sour 04/21/2017 17:0		
cluster_resource_request	1,616 txt-file 01/19/2017 17:0		
cluster_resource_request report.txt	1.4.47 Bython Sour 01/05/2017 15:2		
cluster_resource_request report.txt viz_trinitymapResults.py	1,447 Python Sour 01/05/2017 15:3 1.122 R Source File 01/12/2017 14:0		
cluster_resource_request report.txt viz_trinitymapResults.py Unitiled.R STAR_map.sh	1,447 Python Sour 01/05/2017 15:3 1,122 R Source File 01/12/2017 14:0 761 Shell script 01/19/2017 17:4		
cluster_resource_request report.txt viz_trinitymapResults.py Untitled.R STAR_map.sh STAR_map	1,447 Python Sour 01/05/2017 15:3 1,122 R Source File 01/12/2017 14:0 761 Shell script 01/19/2017 17:4 761 File 01/19/2017 17:41		
cluster_resource_request report.txt viz_trinitymapResults.py Untitled.R STAR_map.sh STAR_map 25 files and 5 directories. Total size	1,447 Python Sour 01/05/2017 15:3 1,122 R Source File 01/12/2017 14:0 761 Shell script 01/19/2017 17:4 761 File 01/19/2017 17:41 6,570,995 bytes	Not connected.	

1: Host: sftp://bbcsrv3.biotech.uconn.edu

- 2: Username:
- 3. Password
- 4.Quickconnect

sftp://vsingh@bbcsrv3.biotech.uconn.edu - FileZilla					
Host: sftp://bbcsrv3.biote Us	sername: vsingh	Password: •••••• Port: Quickco	onnect 🗨		
Status: Connecting to bbcsrv3.biotech.uconn.edu Status: Connected to bbcsrv3.biotech.uconn.edu Status: Retrieving directory listing Status: Listing directory listing of "/home/vsingh Status: Directory listing of "/home/vsingh" successful					
Local site: /Users/vijendersing	Local site: //Users/vijendersingh/Documents/Scripts/				<u> </u>
Software_tool_comparison		▼ 12 / ▼ 12 home ▶ Singh			
Filename	Filesize V Filetype	Last modified			
···	4 805 356 HTML docum	01/19/2017 12:1			
 Turtle_analysis.nb.html 	776,304 HTML docum	04/25/2017 12:1			
results.nb.html	775,856 HTML docum	01/16/2017 15:3	Filename	Filesize Filetype 🗸	Last modified
report.pdf	64,649 pdf-file	01/19/2017 12:1		774 xhol-filo	04/25/2017
RSEM_output.pdf	47,160 pdf-file	01/16/2017 11:5	sickle latmier report tyt	1 498 tyt-file	12/08/2016 :
RSEM_output.gene.pdf	46,404 pdf-file	01/05/2017 16:2		2 2 2 2 tyt-file	06/12/2017 1
.DS_Store	10,244 File	05/04/2017 11:5	merred fasta file state tyt	925 tyt-file	01/17/2017 1
JellyFish.sh	8,808 Shell script	12/23/2016 05:3		2.216 txt-file	01/17/2017 1
JellyFish.py	8,808 Python Sour	12/23/2016 05:3	Tripity filtered fasta tyt	2,510 txt-file	04/18/2017
tst.sh	3,675 Shell script	02/01/2017 14:5	Trinity_intered_lasta.txt	0 txt-file	01/16/2017 1
master_script.sh	3,503 Shell script	07/14/2017 16:3	README test 2307/11 tyt	225 tyt-file	02/22/2017
help_with_R.R	2,967 R Source File	01/12/2017 16:1	Alignment to Green see Turtle result tyt	7466 txt-file	02/16/2017 1
results.Rmd	2,819 R Markdown	01/16/2017 15:3	array job script sh	1078 Shell script	04/12/2017 1
Turtle_analysis.Rmd	2,819 R Markdown	04/25/2017 12:1	test r Bout	865 Pout-file	07/13/2017 1
filter_trinity.sh	2,216 Shell script	01/16/2017 13:3	o test r	80 P Source	07/13/2017 1
cluster_resource_request	1,732 Python Sour	04/21/2017 17:0	graph py	93 Python S	03/10/2017 1
report.txt	1,616 txt-file	01/19/2017 17:0	RSEM output odf	46.404 pdf-file	01/05/2017 1
viz_trinitymapResults.py	1,447 Python Sour	01/05/2017 15:3	BBC test 272507 out	12 out-file	07/10/2017 1
Untitled.R	1,122 R Source File	01/12/2017 14:0	BBC test 272506 out	12 out-file	07/10/2017 1
STAR_map.sh	761 Shell script	01/19/2017 17:4	BBC test 272505 out	12 out-file	07/10/2017 1
STAR_map	761 File	01/19/2017 17:41		iz out ino	07710720171
25 files and 5 directories. Total size:	6,570,995 bytes		38 files and 41 directories. Total size: 1,998,121 bytes		
Server/Local file Direction Remote file Size Priority Status					
Queued files Failed transfers Successful transfers					

STDIN, STDOUT, STDERR, Redirections

<beginning-of-filename>*</beginning-of-filename>	<pre># * is wildcard to specify many files</pre>
ls > file	<pre># prints ls output into specified file</pre>
command < my_file	# uses file after '<' as STDIN
<pre>command >> my_file</pre>	<pre># appends output of one command to file</pre>
command tee my_file	# writes STDOUT to file and prints it to
	screen
<pre>command > my_file; cat my_file</pre>	# writes STDOUT to file and prints it to
	screen
<pre>command > /dev/null</pre>	<pre># turns off progress info of applications by redirecting their output to /dev/pull</pre>
gren my nattern my file wc	# Pipes (1) output of 'grep' into 'wg'
grep my_pattern my_iiie we	π ripes () output of grep fillo we file
grep my_pactern my_non_existing_	IIIe Z > my_stderr # prints SIDERR to IIIe
> file redirects stdout to fil	.e
1> file redirects stdout to fi	le

- 2> file redirects stderr to file
- &> file redirects stdout and stderr to file

/dev/null is the null device it takes any input you want and throws it away. It can be used to suppress any output.