

Unix introduction

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What is Unix

- Unix is a family of operating systems and environments that exploits the power of linguistic abstractions to perform tasks
- Unix is not an acronym; it is a pun on “Multics”. Multics was a large multi-user operating system that was being developed at Bell Labs shortly before Unix was created in the early '70s. Brian Kernighan is credited with the name.
- All computational genomics is done in Unix



History of Unix

- Initial file system, command interpreter (shell), and process management started by Ken Thompson
- File system and further development from Dennis Ritchie, as well as Doug McIlroy and Joe Ossanna
- Vast array of simple, dependable tools that each do one simple task



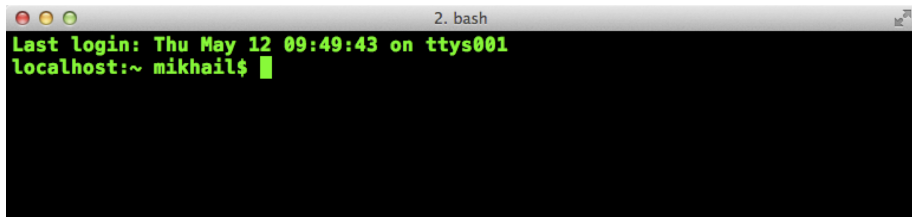
Ken Thompson (sitting) and Dennis Ritchie working together at a PDP-11

Philosophy of Unix

- Vast array of simple, dependable tools
- Each do one simple task, and do it really well
- By combining these tools, one can conduct rather sophisticated analyses

Know your Unix

- Unix users spend a lot of time at the **command line**
- In Unix, a word is worth a thousand mouse clicks

A screenshot of a terminal window titled "2. bash". The window has a standard macOS-style title bar with red, yellow, and green window control buttons on the left. The terminal content is displayed in green text on a black background. It shows the login message "Last login: Thu May 12 09:49:43 on ttys001" followed by the prompt "localhost:~ mikhail\$" and a green cursor block.

```
2. bash
Last login: Thu May 12 09:49:43 on ttys001
localhost:~ mikhail$
```

Unix systems

- Three common types of laptop/desktop operating systems: Windows, Mac, Linux.
- Mac and Linux are both Unix-like!
- What that means for us: Unix-like operating systems are equipped with “shells” that provide a command line user interface.



What is shell

- Shell is an interactive environment with a set of commands to initiate and direct computations
- Shell encloses the complexity of OS, hence the name
 - You type in commands
 - Shell executes them

https://en.wikipedia.org/wiki/Unix_shell

Command line, aka shell

- The Bourne shell (`sh`) is a shell, or command-line interpreter, for computer operating systems.
- Developed by Stephen Bourne at Bell Labs, 1976
- `bash` (the Bourne-Again shell) was later developed for the GNU project and incorporates features from the Bourne shell, `csh`, and `ksh`. It is meant to be POSIX-compliant.

https://en.wikipedia.org/wiki/Stephen_R._Bourne

Most popular types of shell

- `bash` - Bourne-Again shell
- `tcsh` - TENEX C shell
- `zsh` - Z shell
- Change shell: `chsh -s /bin/zsh`
- `$SHELL` environmental variable has path to the currently used shell

Getting to the command line

- Remote access, SSH, **PuTTY**
(<http://www.chiark.greenend.org.uk/~sgtatham/putty/>), **MobaXterm**
(<https://mobaxterm.mobatek.net/>)
- **Mac OS X + Xcode development suite** (free,
<https://developer.apple.com/xcode/>) + **X11 server** (free,
<https://www.xquartz.org/>) + **iTerm2** (optional, <https://iterm2.com/>)
- **Ubuntu Linux** (long-term support LTS version, XX.04,
<http://www.ubuntu.com/download/desktop>)

Getting to the command line | Windows users

- **Cygwin**, <http://www.cygwin.com/>
- **Git Bash**, <https://git-for-windows.github.io/>
- Boot from a CD or USB
- Install the whole Linux systems as a Virtual Machine in **VirtualBox** (<https://www.virtualbox.org/>)

Obtaining new command-line software

- Modern Unix systems have package managers to that download install (free) software for you
- On a Mac, **Homebrew** (<http://brew.sh/>) is a popular package-management system (alternatively, **MacPorts**, <https://www.macports.org/>)
- On Ubuntu, **apt** (https://en.wikipedia.org/wiki/Advanced_Packaging_Tool) is the standard package manager, with both a command-line and graphical interface available
- On Windows, **Cygwin** (<https://cygwin.com/install.html>) installs everything precompiled through its setup file. Do not delete `setup-x86_64.exe` file after installing Cygwin, explore what Linux tools are available (a lot)

Interacting with shell

- Most commands take additional arguments that fine tune their behavior
- If you don't know what a command does, use the command `man <command>`
- Some tools use an alternate documentation system called `info`
- Press `q` to quit the `man` page viewer
- Most often, you'll use `<command> -h` or `<command> --help`
- Some commands output help if executed without any arguments

Files

- Create an empty file: `touch filename.txt`
- Edit it: `nano filename.txt`

Move files over the network

scp: securely copy a file from one computer to another

- Use scp to securely transfer files between two Unix computers
- No need to start an FTP session or logging into the remote system explicitly
- The scp command uses SSH to transfer data, so it requires a password or passphrase for authentication
- scp encrypts both the file and any passwords exchanged
- Alternatively, use rsync

The syntax for the scp command is:

```
scp [options] username1@source_host:directory1/filename1 username2@target_host:directory2/filename2
```

scp examples

- Copy remote file locally (in the current folder)

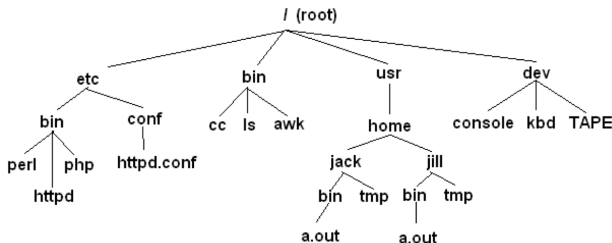
```
scp mdozmorov@merlot.bis.vcu.edu:sequencing/hg38.Ensembl.g
```

- Copy local file remotely

```
scp hg38.Ensembl.gtf mdozmorov@merlot.bis.vcu.edu:sequenci
```

- Use `-r` (recursive) option to copy a directory
- Explore `rsync` command as an alternative to `scp`

File system: Full vs. relative paths



- `cd /` - go to the root directory
- `cd /usr/home/jack/bin` - go to the user's sub-directory
- `cd ..` - go to the upper level directory
- `cd`, or `cd ~` - go to the user's home directory
- `cd --` - go to the last visited directory

Permissions: `chmod`, `chown` and `chgrp`

In Unix, every file and directory has an **owner** and a **group**

- Owner - is the one who created a file/directory
- Group - defines rules of file operations and/or permissions
- Every - user on a Unix machine can belong to one or more groups

Every file has **three permission levels**

- what the **u**ser can do
- what the **g**roup can do
- what the **a**ll can do

Permissions: `chmod`, `chown` and `chgrp`

- The first column tells you about the permissions on the file
 - The very first character in the permissions column tells you what kind of file it is. A `-` means it's a regular file. A `d` means it's a directory
 - The next nine characters come in three classes, each has three characters. The three classes are owner/group/world permissions
 - Inside a permission class, `r` means that class can **read** the file; `w` means that class can **write** the file; `x` means that class can **execute** the file
- The second column has the number of files (inside a directory)
- The third and fourth columns tell you the owner and group

Permissions: chmod, chown and chgrp

```
-rw-r--r--    1 mdozmorov  staff    205B Dec 19 11:01 BIOS692.2018.Rpt
-rw-r--r--    1 mdozmorov  staff    3.5K Dec 18 10:20 BUILD.md
-rw-r--r--    1 mdozmorov  staff    470B Dec 19 08:48 README.md
-rw-r--r--    1 mdozmorov  staff    2.1K Dec 19 07:51 _config.yml
drwxr-xr-x   10 mdozmorov  staff    340B Dec 18 10:20 _includes
drwxr-xr-x   10 mdozmorov  staff    340B Dec 18 10:20 _layouts
drwxr-xr-x    7 mdozmorov  staff    238B Dec 18 10:29 _posts
-rw-r--r--    1 mdozmorov  staff    1.0K Dec 20 15:54 acknowledgement.
```

Orienting in the filesystem

- `pwd` - print working directory
- `ls` - list all files in the current directory
- `ls -l` - list files in *one* column
- `ls -lah` - list files in long, human readable format, include all content, user, owner, permissions

Creating, moving, copying, and removing files

- `touch <file>` - creates an empty file
- `mkdir <dirname>` - creates a directory
- `cp <source_file> <target_file>` - copy a file to another location/file
- `mv <source_file> <target_file>` - move a file
- `rm <file>` - remove a file. If multiple files provided, removes all of them
- `rm -r <dirname>` - recursive removal (deletes a directory)

Finding your files

- `find` - lists all files under the working directory (and its subdirectories) based on arbitrary criteria
- `find .` - prints the name of every file or directory, recursively. Starts from the current directory
- `find . -type f` - finds files only
- `find . -type d -maxdepth 1` - finds directories only, at most 1 level down
- `find . -type f -name "*.mp3"` - finds only *.mp3 files
- `find . -type f -name "README.md" -exec wc -l {} \;` - find files and execute a command on them

Wildcards and patterns

- `*` - matches any character
- `?` - matches a single character
- `[chars]` - matches any character in chars
- `[a-zA-Z]` - matches any character between a and z, including capital letters
- `ls *.md`
- `ls [Rt]*`

Looking inside files

- `cat <file>` - prints out content of a file. If multiple files, consequently prints out all of them (concatenates)
- `zcat` - prints out content of gzipped files
- `more/less <file>` - shows the content of the file one screen at a time

Keyboard shortcuts for more command

- space - forward
- b - backward
- g - go to the beginning
- G - go to the end
- `/<text>` - starts forward search, enter to find next instance
- q - quit

Chaining commands: pipes

One of the most useful capabilities of Unix is the ability to redirect the STDOUT of one command into the STDIN of another

The | (pipe) character feeds output from the first program (to the left of the |) as input to the second program on the right. Therefore, you can string all sorts of commands together using the pipe

```
find . | wc -l  
cat names.txt | sort | uniq -c
```

Executing one command AFTER another completed successfully:
<command> && <command>

```
mkdir music && mv *.mp3 music/
```

Chaining commands: redirections

- Nearly every command in Unix makes use of a convention to have a “standard input” (also called stdin or STDIN, or channel 0) and “standard output” (also called stdout or STDOUT, or channel 1)
- There is also a “standard error” (stderr or STDERR, or channel 2) output that is, by convention, reserved for error messages
- `find / 2> error.log` - capture STDERR into a file
- `find / 2> /dev/null` - suppress STDERR messages
- `find / 2>&1` - add STDERR to STDOUT

Chaining commands: redirections

- If you want to dump the standard output into a file, use `command > file` (overwrites the file). `command >> file` (appends to the file)
- Redirection example: `ls > README.md` - save file list in the current directory into README.md file
- Redirection works in another direction: `grep CC0 < License.md`
- Or, the content of a command into another command: `join <(sort file1) <(sort file2)`

Other essential commands

head/tail	cut
for	comm
sort	echo
uniq	basename
wc	dirname
tr	history
grep	which
join	who
kill	grep

Shell conveniences

- Tab completion
- Ctrl-c - cancel the command you are writing
- Ctrl-r - reverse search through your command line history
- history - shows your previous commands
- !<history number> - repeats specific command
- !! - repeats the last command

Statistical command line goodies

- **data_hacks**, https://github.com/bitly/data_hacks
 - Command line tools for data analysis
 - `histogram.py`
 - `bar_chart.py`
 - `sample.py`
- **datamash**, <https://www.gnu.org/software/datamash/>
 - summary statistics
 - transposing matrixes

Unix for high-performance cluster computing

- Allow you to submit multiple jobs at once
- Depending on the system, can schedule jobs for you
- Are optimized for high-throughput performance

Unix at VCU

- VCU Biostatistics cluster information,
<https://wiki.vcu.edu/display/biosit/Home>
- Contact Helen Wang (huwang at vcu.edu) to establish an account
- Google if you run into problems using Unix!

Learn more

- Heng Li's "A Bioinformatician's UNIX Toolbox",
<http://lh3lh3.users.sourceforge.net/biounix.shtml>
- Bioinformatics one-liners by Stephen Turner,
<https://github.com/stephenturner/oneliners>
- Collection of bioinformatics-genomics bash one liners, using awk, sed etc. <https://github.com/crazyhottommy/bioinformatics-one-liners>
- Links and references to many genomics and bioinformatics resources,
<https://github.com/crazyhottommy/getting-started-with-genomics-tools-and-resources>