Introduction to Unix

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For the sake of clarity, the concepts mentioned in these slides have been simplified significantly.

Most of the command line tools and programs we explain have many more features that can't be found here. Unix command line tools are very well documented and their manual pages are freely available.

These slides are prepared for the users of MGHPCC. MGHPCC system has features that don't exist in standard Unix systems such as loading modules.

A rigorous treatment of topics on Unix and Bash can be found in various books in several levels.

Unix:

Unix in a Nutshell , Arnold Robbins, ISBN-13: 978-0596100292 Learning the bash Shell: Unix Shell Programming, Cameron Newham, ISBN-13: 978-0596009656

Umass High Perfotmance Computing:

http://wiki.umassrc.org/

Bioinformatics:

UCSC genome bioinformatics: https://genome.ucsc.edu/ Sam file format: samtools.github.io/hts-specs/SAMv1.pdf Samtools: http://samtools.sourceforge.net/ Bedtools: http://bedtools.readthedocs.org/en/latest/ bowtie2: http://bowtie-bio.sourceforge.net/bowtie2/manual.shtml tophat: http://ccb.jhu.edu/software/tophat/index.shtml An operating system is a collection of software that manages system resources (such as memory, cpu(s), display and etc.) and provide applications a simpler interface to the system hardware.





Developed in the 1960's by Ken Thompson, Dennis Ritchie. It became very popular in the 70's and 80's. "Contrary to popular belief, Unix is user friendly. It just happens to be very selective about who it decides to make friends with." –anonymous *Unix-like* operating systems *"behave"* like the original Unix operating system and comply (at least partially) with POSIX (portable operating system interface) standards.

Examples: Linux, OS X, Free BSD, Solaris, Hurd.

 $\text{Unix} \equiv \text{Unix-like} \equiv \text{Unix-clone}$



What is a Terminal?



A terminal is a software that emulates a teletype writer terminal used in the early days of Unix.

A shell is a software that runs inside a terminal and interprets and executes user commands.

One of the most popular shells being used today is called **BASH** (Bourne Again Shell). It is also the default shell in the UMass HPC cluster, OS X and many Linux distributions.



\$ denotes the Bash command line prompt. It is not meant to be typed. All Bash commands will appear in a gray box.

\$ bashcommand

We connect to MGHPCC server.

\$ ssh username@ghpcc06.umassrc.org

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Let's verify that we are at the right place.

\$ hostname

To print something on the screen, we use echo

\$ echo Hello World

Navigating the Unix File System



Bash always has a working directory. To see the working directory,

\$ pwd

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To see the contents of the working directory,

\$ ls

To see the contents of a particular directory, provide the path of the directory.

\$ ls /bin

displays th contents of the directory /bin

A program's execution and / or output format can be modified, according to our needs, by providing command line arguments.



We can get more information about the files or directories by running ls with additional arguments.

Compare

with

A closer look at Is

ls list the given directory contents. To see the complete list of options,

\$ man ls

This will display the manual page of ls.

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Some commonly used options are:

- -I : list contents in more detail
- -A : list all files in the direcory
- -h : when used with the -1 option, prints file sizes in KB, MB, GB, and etc. to reduce the number of digits on the screen.

These options can be grouped

Let's first have some files and folders to experiment with.

\$ /tmp/bootcamp.pl

Upon success, you will see /home/user_name/bootcamp

First, we see the working directory

\$ pwd

Now, lets change the directory to the bootcamp directory.

\$ cd ~/bootcamp

We can verify the change by

\$ pwd

\$ mkdir directory_path

Note that

\$ mkdir /home/username/bootcamp/my_new_directory

is the same as

\$ mkdir ~/bootcamp/my_new_directory

\$ cp sourcefile(s) destination_path

Let's copy our sample files into the home folder.

\$ cp ~/bootcamp/transcriptomics.tar.gz ~/transcriptomics.tar.gz



Copying Directories



\$ cp source_directory destination_directory

will not work in general. You need to provide the -r parameter.

\$ cp -r source_directory destination_directory

Another useful option is -v which prints the files being copied.

\$ cp -rv source_directory destination_directory



\$ mv source_file destination_file



rm : Remove comand

\$ rm file_path

To remove non-empty directories, use rm with the -r option.

\$ rm -r directory

There is no undoing of rm! You may want to use it in the ineractive mode not to delete accidentally files

File Path

- ~ : Home directory
 - : current directory
- .. : parent directory
 - If a file path begins with /, then it is an *absolute path*.
 It doesn't depend on the current working directory.
 - If a file path begins WITHOUT /, then it is a *relative path*. The path is defined according to the current directory. So, the path depends on the current working directory.

/tmp/bootcamp/bootcamp.pl : Absolute path.

bootcamp.pl : Relative path. This is the botcamp.pl file in the working directory.

../bootcamp.pl : Relative path. This is the botcamp.pl file in the parent directory of the working directory.

~/bootcamp.pl : This is the file in the **home** directory of the current user.

Technically, every file is a binary file.

Yet, it is common practice to group the files into two:

Text Files	:	Contains only printable characters.
Binary Files	:	Files that are not text files.

Binary files are generally more compact and therefore smaller in size. They are read and processed by specific applications. Text files are of extreme importance in bioinformatics. There are many tools to view and edit text files. **less** is a very useful program.

\$ less text_file

For details read the manual page

\$ man less

For a very simple text editor:

\$ nano

vi and **emacs** are very powerful text editors with sophisticated features.

Let's try to run a bash script.

\$./hello.sh

Let's try another one

\$./other_hello.sh

Why didn't this work?

Let's try to spy on Manuel and see what he is up to.

\$ ls /home/mg44w

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\$ ls /home/mg44w

In Unix, users can only access (read, write or execute) files that they have permission to.

A typical output of

is

-rw-r–r–	1	ho86w	moore	11M	Nov 5	00:41	RPKM.csv
-rwxrwxrwx	1	ho86w	moore	143B	Oct 27	02:53	my_script.sh
drwxr-xr-x	21	ho86w	moore	714B	Jan 24	13:52	sam_files

- r : read
- w : write
- x : execute



- User : Has read, write and execute access.
- Group : Has read and execute access but can't write to the file
- Others : Has only read access.

chmod: Change file mode bits.

\$ chmod filemode file

To give only read access to other users,

\$ chmod o=r myscript.sh

To give read and execution access together,

\$ chmod o=rx myscript.sh

A computer process is an instance of a program. It takes up space in the memory and its instructions are executed in the CPU.

You can list **your** running processes

\$ ps -u username

where $\tt username$ is your username. You can see them in more detail with the -F option

You can terminate a process by

\$ kill -9 processID

where $\verb"processID"$ is the process id that can be obtained by the $\verb"ps"$ command.

To send a process to the background, first put the process on hold by pressing ctrl and z, and then send it to the background by

\$ bg

You can check your processes by

\$ jobs

To put a back to the foreground, first get the process id by

\$ jobs

and then

\$ fg % processNumber

To start a process on the background, put & at the end of your command

\$ command &

A common type of binary files is zipped files. We can save space by zipping text files.

We use

gzip to compress / decompress files

tar to pack files into one file for archiving purposes.

To compress a file

\$ gzip file

To decompress, reads.fastq.gz,

\$ gzip -d reads.fastq.gz

Say we collect our results on a project in one directory. We want to send this directory in an email attachment to a collaborator. How do we do that?



To pack a directory

To pack a directory in a zipped tar file

To get our directory back from the tar file

To get our directory back from the tar.gz file

\$ tar -xvzf archive.tar.gz