

Linux Commands Cheat Sheet

Command	Explanation
<code>cd <dir></code>	Change directories. When run without any arguments, go to your home directory.
<code>ls</code>	List contents of a directory. <code>ls -l</code> is a long list with more info.
<code>man <command></code>	Manual for the command. Space=forward, b=backwards, q=quit, /=search.
<code>pwd</code>	Print working directory. Shows you where you currently reside on the directory tree.
<code>gzip <files></code>	Compress files.
<code>gunzip <files.gz></code>	Decompress files.
<code>history</code>	Display command line history.
<code>cat <files></code>	Print files to the screen (conCATenate)
<code>less <files></code>	Open the file for viewing. Space=forward, b=backwards, q=quit, /=search.
<code>head <file></code>	Prints the first few lines of the file.
<code>tail <file></code>	Prints the last few lines of the file.
<code>grep <pattern> <file></code>	Pulls out lines from the file matching the pattern.
<code>wc <files></code>	Word count on files.
<code>cp <src> <dest></code>	Copies file from source to destination.
<code>mv <src> <dest></code>	Moves the file from source to destination. If dest is a directory, this will move the src file into the dest directory. Otherwise, this will rename the file called src to a file called dest.
<code>nano</code>	A basic text editor
<code>which</code>	Tells you where a program resides
<code>chmod +x <file></code>	Makes a file executable
<code>find .</code>	Finds all files in the current directory recursively. Add -type f to limit to files only. Add -name "pattern" to restrict files found to matching the pattern.
<code>parallel "cmd {}"</code>	Runs the command <code>cmd</code> on each file passed in on a pipe from find, which assumes the place of {}. E.g., running the command: <code>find . -name "*.fastq" parallel "somecommand {} > {}.results.txt"</code> Would go through all the fastq files found and run somecommand on that file, redirecting the output to a file ending in .results.txt. E.g. <code>somecommand sample1.fastq > sample1.fastq.results.txt</code> <code>somecommand sample2.fastq > sample2.fastq.results.txt</code> <code>somecommand sample3.fastq > sample3.fastq.results.txt</code>

Shortcut	Location
<code>.</code>	The current directory, "here".
<code>..</code>	The parent directory relative to here ("the directory above")
<code>~</code>	Your home directory. Same as <code>/home/<username></code> .