

# UNIX COMMANDS

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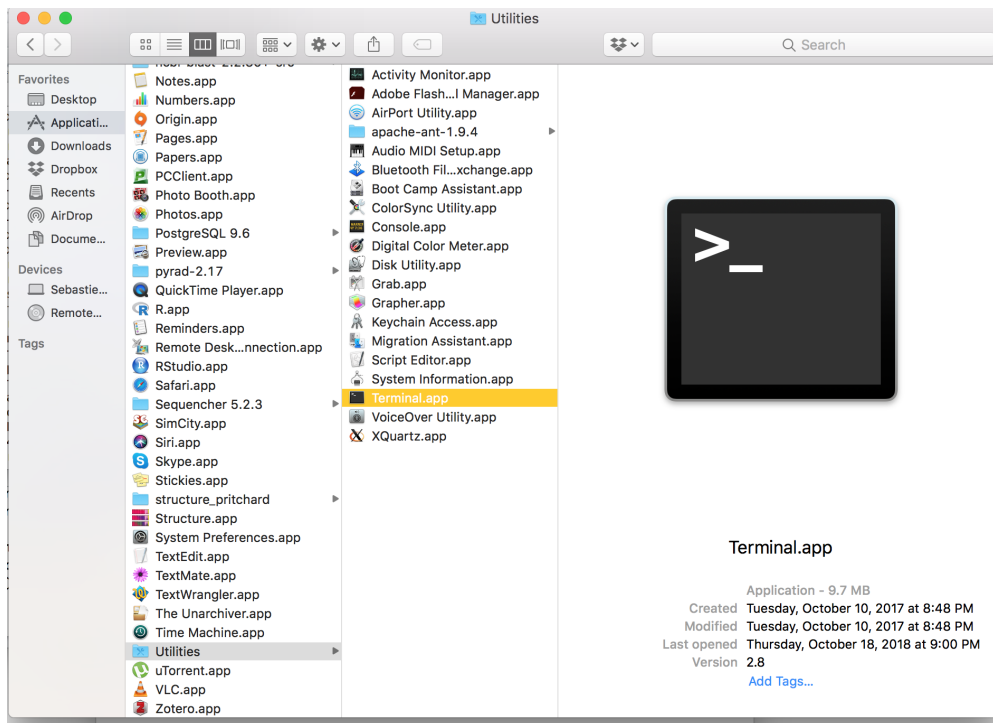
*October 22, 2018*

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## Get started

- Open terminal



- Download github repo (`curl -L https://github.com/seb951/basic_unix/archive/master.zip`  
>master.zip)
- Decompress (`unzip master.zip`)

## General principles

- All commands are short, perform simple tasks & are highly optimized (fast)
- Nearly all commands and options (~99%) work across Unix-based OS
- Commands can take (many) options
- Commands can be combined (see more later)
- Many commands don't require any argument, some do
- `command -option1 -option2 -optionN... parameter object`
- Many commands output directly to terminal

## Exploration

Command	Meaning
<code>ls</code>	what's there
<code>ls -l</code>	with an option
<code>ls -t -h</code>	with more options
<code>ls -thor</code>	combining even more options
<code>man ls</code>	how to run command (q = quit; d = next page; ; arrows & fn arrows = scrolling)
<code>pwd</code>	where are we (full path)
<code>cd</code>	move somewhere else (default to home)
<code>cd ..</code>	move back
<code>cd</code>	move back, then forward
<code>../Applications</code>	

Command	Meaning
<code>cd ~</code>	move home
<code>cd /</code>	move to root

## Moving / removing / creating files

Command	Meaning
<code>mv</code>	move
<code>cp</code>	copy
<code>rm</code>	<b>careful</b> remove a file
<code>rmdir</code>	<b>careful</b> remove an empty directory
<code>rm -r</code>	<b>VERY careful</b> remove directory recursively
<code>mkdir</code>	create empty directory
<code>touch</code>	create empty file
<code>echo</code>	print message to screen

## Examining files

Command	Meaning
<code>head</code>	print first 10 lines to screen
<code>tail</code>	print last 10 lines to screen
<code>less</code>	print file line by line (use arrows + fn arrows to move up/down)
<code>cat</code>	print file content to screen
<code>sort</code>	sort a file
<code>uniq</code>	unique lines in a file
<code>clear</code>	clear terminal
<code>wc</code>	wordcount
<code>du</code>	disk usage

## Tricks

Command	Meaning
<code>&gt;</code>	redirect
<code>&gt;&gt;</code>	append to file
<code> </code>	pipe
<code>*</code>	anything (special character: for literal *, use \*)
<code>?</code>	any character (special character)
<code>Ctrl+A</code>	go to beginning of line in terminal
<code>Ctrl+E</code>	end of line
<code>tab key</code>	autocomplete

## Working remotely

Command	Meaning
<code>ssh</code>	secure shell (remote connection)
<code>scp</code>	secure copy (remote connection)

## Other useful commands

Command	Meaning
<code>chmod</code>	change mode of file
<code>sudo</code>	superuser do
<code>top</code>	what's running
<code>kill</code>	kill a process
<code>curl</code>	get stuff from web (MACOS native) or <code>wget</code> (MACOS non-native)
<code>history</code>	see recent history
<code>open</code>	open a file
<code>ln</code>	symbolic link
<code>which</code>	where is a command
<code>alias</code>	create own commands
<code>nano</code>	edit text (whole file loaded in memory)
<code>nano +123</code>	specify location in file

## Profile

- Let's look at `.bash_profile` to modify options (e.g. `nano ~/.bash_profile`)
- Usefull options are here: `examples/bash_example_commands`

## Compression

Command	Meaning
<code>gzip</code>	compression/decompression of files
<code>gzip -cd file.gz</code>	decompress to standard out (use a <code>&gt;</code> redirect)
<code>gunzip</code>	compression/decompression of files
<code>tar</code>	compression/decompression of archives
<code>tar -xvzf file tar.gz</code>	decompress a tar.gz (tarball)

## Regular expressions

Command	Meaning
<code>grep</code>	regular expression
<code>grep -options 'pattern' file</code>	regular expression

## Examples

### Combining commands:

- Let's look at the file `examples/sequence.fasta`
- print line 17 by piping `head` and `tail`
- Count lines starting with '`>`' (pipe `grep` and `wc`)
- Count unique lines starting with '`>`' (pipe `grep`, `sort`, `uniq` and `wc`)
- How many files is there in the directory (pipe `ls` & `wc`)

### Install and run command line programs

- Let's look at this example: <https://github.com/rec3141/rec-genome-tools/blob/master/bin/fastagrep.pl>
- Notice the `#!/` (shebang) on the first line
- It is a short `perl` script that *greps* sequences in a *.fasta* formatted file
- Copy-paste the text in a new file (e.g. `touch fastagrep.pl`, `nano fastagrep.pl`, paste and save)
- Make it executable (`chmod +x fastagrep.pl`)
- Let's look at it (`./fastagrep.pl`)
- Let's try it (`./fastagrep.pl 'deserticola' examples/sequence.fasta`)
- Let's symbolic link it so the computer knows where to find it (`ln source target`)

### bash

- The terminal runs (by default) the `bash` command language
- We can write or use more complex executable `bash` code
- Let's look at an example `less examples/bash_cmd2`