

The Linux Command Line

In this lecture we'll introduce you to a large number of new command line tools and show you how to connect them together via pipes.

The Command Line

In general, the best way to understand the command line is to learn by doing. We'll go through a number of useful commands with examples, and you should become reasonably skilled in the process, but it is nevertheless worth looking at a few books to consult as references. Here are some particularly good ones:

- [The Command Line Crash Course](#) is Zed Shaw's free online tutorial introduction to the command line. Much shorter than the others listed here and recommended.
- [Sobell's Linux Book](#): If you only get one book, get Sobell's. It is probably the single best overview, and much of the content is applicable to both Linux and OS X¹ systems.
- [Unix Power Tools](#): While this came out in 2002, much of it is still relevant to the present day. New options have been added to commands but old ones are very rarely deprecated.
- [Unix/Linux Sysadmin Handbook](#): Somewhat different focus than Sobell. Organized by purpose (e.g. shutdown), and a good starter guide for anyone who needs to administer machines. Will be handy if you want to do anything nontrivial with EC2.

With those as references, let's dive right in. As always, the commands below should be executed after `ssh`'ing into your EC2 virtual machine¹. If your machine is messed up for some reason, feel free to terminate the instance in the [AWS EC2 Dashboard](#) and boot up a new one. Right now we are doing all these commands in a "vanilla" environment without much in the way of user configuration; as we progress we will set up quite a lot.

The three streams: STDIN, STDOUT, STDERR

Most `bash` commands [accept input](#) as a single stream of bytes called `STDIN` or standard input and yields two output streams of bytes: `STDOUT`, which is the standard output, and `STDERR` which is the stream for errors.

- `STDIN`: this can be text or binary data streaming into the program, or keyboard input.
- `STDOUT`: this is the stream where the program writes its data. This is printed to the screen unless otherwise specified.

¹You can probably get away with running them on your local Mac, but beware: the built-in command line tools on OS X (like `mv`) are very old BSD versions. You want to install the new GNU tools using [homebrew](#). Then your local Mac environment will be very similar to your remote EC2 environment.

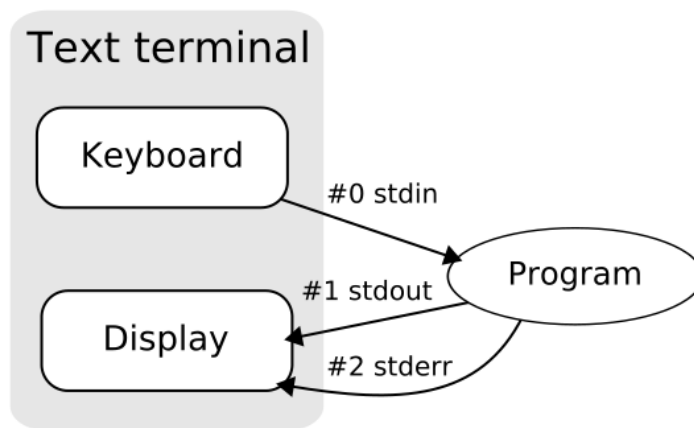


Figure 1: Visualizing the three standard streams. From [Wikipedia](#).

- **STDERR:** this is the stream where error messages are display. This is also printed to the screen unless otherwise specified.

Here are a few simple examples; execute these at your EC2 command line.

```

1  # Redirecting STDOUT with >
2  echo -e "line1\nline2"
3  echo -e "line1\nline2" > demo.txt
4
5  # Redirecting STDERR with 2>
6  curl fakeurl # print error message to screen
7  curl fakeurl 2> errs.txt
8  cat errs.txt
9
10 # Redirecting both STDIN and STDOUT to different files with 1> and 2>
11 curl google.com fakeurl 1> out1.txt 2> out2.txt
12
13 # Redirecting both to the same file with &>
14 curl google.com fakeurl &> out.txt
15
16 # Getting STDIN from a pipe
17 cat errs.txt | head
18
19 # Putting it all together
20 curl -s http://google.com | head -n 2 &> asdf.txt

```

Note what we did in the last two examples, with the `|` symbol. That's called a pipe, and the technique of connecting one command's STDOUT to another command's STDIN is very powerful. It allows us to build up short-but-powerful programs by composing individual pieces. If you *can* write something this way, especially for text processing or data analysis, you almost always *should*...even if it's a complex ten step pipeline. The reason is that it will be incredibly fast and robust, due to the fact that the underlying GNU tools are written in C

and have received countless hours of optimization and bug fixes. It will also not be that hard to understand: it's still a one-liner, albeit a long one. See this piece on [Taco Bell programming](#) and [commentary](#). As a corollary, you should design your own programs to work in command line pipelines (we'll see how to do this as the course progresses). With these principles in mind, let's go through some interactive examples of Unix commands. All of these should work on your default Ubuntu EC2 box.

Navigation and Filesystem

list files: `ls`

The most basic and heavily used command. You will use this constantly to get your bearings and confirm what things do.

```
1 ls
2 ls --help
3 ls --help | less # 'pipe' the output of ls into less
4 ls -a
5 ls -alrth
6 alias ll='ls -alrth' # create an alias in bash
7 ll
```

modify/create empty files: `touch`

Used to change timestamps and quickly create zero byte files as placeholders. Useful for various kinds of tests.

```
1 touch file1
2 touch file2
3 ll
4 ll --full-time
5 touch file1
6 ll --full-time # note that the file1 modification time is updated
```

display text: `echo`

Useful for debugging and creating small files for tests.

```
1 echo "foo"
2 man echo
3 echo -e "foo\n"
4 echo -e "line1\nline2" > demo.txt # write directly to tempfile with '>'
```

copy files: `cp`

Powerful command with many options, works on everything from single files to entire archival backups of huge directories.

```

1 cp --help
2 cp demo.txt demo2.txt
3 cp -v demo.txt demo3.txt # verbose copy
4 cp -a demo.txt demo-archive.txt # archival exact copy
5 ll --full-time demo* # note timestamps of demo/demo-archive
6 echo "a new file" > asdf.txt
7 cat demo.txt
8 cp asdf.txt demo.txt
9 cat demo.txt # cp just clobbered the file
10 alias cp='cp -i' # set cp to be interactive by default

```

move/rename files: mv

Move or rename files. Also extremely powerful.

```

1 mv asdf.txt asdf-new.txt
2 mv -i asdf-new.txt demo.txt # prompt before clobbering
3 alias mv='mv -i'

```

remove files: rm

Powerful command that can be used to delete individual files or recursively delete entire trees. Use with caution.

```

1 rm demo.txt
2 rm -i demo2.txt
3 rm -i demo*txt
4 alias rm='rm -i'
5 alias # print all aliases

```

symbolic link: ln

Very useful command that allows a file to be symbolically linked, and therefore in two places at once. Extremely useful for large files, or for putting in one level of indirection.

```

1 yes | nl | head -1000 > data1.txt
2 head data1.txt
3 ll data1.txt
4 ln -s data1.txt latest.txt
5 head latest.txt
6 ll latest.txt # note the arrow. A symbolic link is a pointer.
7 yes | nl | head -2000 | tail -50 > data2.txt
8 head latest.txt
9 ln -s data2.txt latest.txt # update the pointer w/o changing the underlying file
10 head latest.txt

```

```
11 head data1.txt
12 head data2.txt
```

print working directory: pwd

Simple command to print current working directory. Used for orientation when you get lost.

```
1 pwd
```

create directories: mkdir

Create directories. Has a few useful options.

```
1 mkdir dir1
2 mkdir dir2/subdir    # error
3 mkdir -p dir2/subdir # ok
```

change current directory: cd

Change directories. Along with `ls`, one of the most frequent commands.

```
1 cd ~
2 cd dir1
3 pwd
4 cd ..
5 cd dir2/subdir
6 pwd
7 cd - # jump back
8 cd - # and forth
9 cd # home
10 alias ..='cd ..'
11 alias ...='cd ..; cd ..'
12 cd dir2/subdir
13 ...
14 pwd
```

remove directories: rmdir

Not used that frequently; usually recursive `rm` is used instead. Note that `rm -rf` is the most dangerous command in Unix and must be used with extreme caution, as it means “remove recursively without prompting” and can easily nuke huge amounts of data.

```
1 rmdir dir1
2 rmdir dir2 # error
3 rm -rf dir2 # works
```

Downloading and Syncing

synchronize local and remote files: `rsync`

Transfer files between two machines. Assuming you have set up your `~/.ssh/config` as in previous lectures, run this command on your *local* machine, making the appropriate substitutions:

```
1  # Run on local machine
2  $ yes | nl | head -500 > something.txt
3  $ rsync -avp something.txt awshost4:~/
4  building file list ... done
5  something.txt
6
7  sent 4632 bytes  received 42 bytes  1869.60 bytes/sec
8  total size is 4500  speedup is 0.96
9  $ rsync -avp something.txt awshost4:~/
10 building file list ... done
11
12 sent 86 bytes  received 20 bytes  42.40 bytes/sec
13 total size is 4500  speedup is 42.45
```

Notice that the first `rsync` invocation sent 4632 bytes but the second time sent much less data 86 bytes. That is because `rsync` can resume transfers, which is essential for working with large files and makes it generally preferable to `scp`. As an exercise, `ssh` in and run `cat something.txt` on the EC2 machine to see that it was uploaded.

retrieve files via `http/https/ftp`: `wget`

Very useful command to rapidly pull down files or webpages. Note: the distinction between `wget` and `rsync` is that the `wget` is generally used for publicly accessible files (accessible via a web browser, perhaps behind a login) while `rsync` is used for private files (usually accessible only via `ssh`).

```
1  wget http://startup-class.s3.amazonaws.com/simple.sh
2  less simple.sh # hit q to quit
3
4  # pull a single HTML page
5  wget https://github.com/joyent/node/wiki/modules
6  less modules
7
8  # recursively download an entire site, waiting 2 seconds between hits.
9  wget -w 2 -r -np -k -p http://www.stanford.edu/class/cs106b
```

interact with single URLs: `curl`

A bit different from `wget`, but has some overlap in functionality. `curl` is for interacting with single URLs. It doesn't have the spidering/recursive properties that `wget` has, but it supports

a much wider array of protocols. It is very commonly used as the building block for API calls.

```
1 curl https://install.meteor.com | less # an example install file
2 curl -i https://api.github.com/users/defunkt/orgs # a simple API call
3 GHUSER="defunkt"
4 GHVAR="orgs"
5 curl -i https://api.github.com/users/$GHUSER/$GHVAR # with variables
6 GHVAR="repos"
7 curl -i https://api.github.com/users/$GHUSER/$GHVAR # with diff vars
```

send test packets: ping

See if a remote host is up. Very useful for basic health checks or to see if your computer is online. Has a surprisingly large number of options.

```
1 ping google.com # see if you have internet connectivity
2 ping stanford.edu # see if stanford is up
```

Basic Text Processing

view files: less

Paging utility used to view large files. Can scroll both up and down. Use q to quit.

```
1 rsync --help | less
```

print/concatenate files: cat

Industrial strength file viewer. Use this rather than MS Word for looking at large files, or files with weird bytes.

```
1 # Basics
2 echo -e "line1\nline2" > demo.txt
3 cat demo.txt
4 cat demo.txt demo.txt demo.txt > demo2.txt
5 cat demo2.txt
6
7 # Piping
8 yes | head | cat - demo2.txt
9 yes | head | cat demo2.txt -
10
11 # Download chromosome 22 of the human genome
12 wget ftp://ftp.ncbi.nih.gov/genomes/Homo_sapiens/CHR_22/hs_ref_GRCh37.p10_chr22.gbk.gz
13 gunzip hs_ref_GRCh37.p10_chr22.gbk.gz
14 less hs_ref_GRCh37.p10_chr22.gbk
15 cat hs_ref_GRCh37.p10_chr22.gbk # hit control-c to interrupt
```

first part of files: head

Look at the first few lines of a file (10 by default). Surprisingly useful for debugging and inspecting files.

```
1 head --help
2 head *gbk      # first 10 lines
3 head -50 *gbk  # first 50 lines
4 head -n50 *gbk # equivalent
5 head *txt *gbk # heads of multiple files
6 head -q *txt *gbk # heads of multiple files w/o delimiters
7 head -c50 *gbk # first 50 characters
```

last part of files: tail

Look at the bottom of files; default is last 10 lines. Useful for following logs, debugging, and in combination with head to pull out subsets of files.

```
1 tail --help
2 tail *gbk
3 head *gbk
4 tail -n+3 *gbk | head # start at third line
5 head -n 1000 *gbk | tail -n 20 # pull out intermediate section
6 # run process in background and follow end of file
7 yes | nl | head -n 10000000 > foo &
8 tail -F foo
```

extract columns: cut

Pull out columns of a file. In combination with head/tail, can pull out arbitrary rectangular subsets of a file. Extremely useful for working with any kind of tabular data (such as data headed for a database).

```
1 wget ftp://ftp.ncbi.nih.gov/genomes/Bacteria/\
2 Escherichia_coli_K_12_substr__W3110_uid161931/NC_007779.ptt
3 head *ptt
4 cut -f2 *ptt | head
5 cut -f2,5 *ptt | head
6 cut -f2,5 *ptt | head -30
7 cut -f1 *ptt | cut -f1 -d'.' | head
8 cut -c1-20 *ptt | head
```

numbering lines: nl

Number lines. Useful for debugging and creating quick datasets. Has many options for formatting as well.


```
1 nl *gbk | tail -1 # determine number of lines in file
2 nl *ptt | tail -2
```

concatenate columns: paste

Paste together data by columns.

```
1 tail -n+3 *ptt | cut -f1 > locs
2 tail -n+3 *ptt | cut -f5 > genes
3 paste genes locs genes | head
```

sort by lines: sort

Industrial strength sorting command. Very powerful standalone and in combination with others.

```
1 sort genes | less # default sort
2 sort -r genes | less # reverse
3 sort -R genes | less # randomize
4 cut -f2 *ptt | tail -n+4 | head
```

uniquify lines: uniq

Useful command for analyzing any data with repeated elements. Best used in pipelines with sort beforehand.

```
1 cut -f2 *ptt | tail -n+4 | sort | uniq -c | sort -k1 -rn #
2 cut -f3 *ptt | tail -n+4 | uniq -c | sort -k2 -rn # output number
3 cut -f9 *ptt > products
4 sort products | uniq -d
```

line, word, character count: wc

Determine file sizes. Useful for debugging and confirmation, faster than nl if no intermediate information is needed.

```
1 wc *ptt # lines, words, bytes
2 wc -l *ptt # only number of lines
3 wc -L *ptt # longest line length, useful for apps like style checking
```

split large files: split

Split large file into pieces. Useful as initial step before many parallel computing jobs.

```
1 split -d -l 1000 *ptt subset.ptt.  
2 ll subset.ptt*
```

Help

manuals: man

Single page manual files. Fairly useful once you know the basics. But Google or StackOverflow are often more helpful these days if you are really stuck.

```
1 man bash  
2 man ls  
3 man man
```

info: info

A bit more detail than `man`, for some applications.

```
1 info cut  
2 info info
```

System and Program information

computer information: uname

Quick diagnostics. Useful for install scripts.

```
1 uname -a
```

current computer name: hostname

Determine name of current machine. Useful for parallel computing, install scripts, config files.

```
1 hostname
```

current user: whoami

Show current user. Useful when using many different machines and for install scripts.

```
1 whoami
```

current processes: ps

List current processes. Usually a prelude to killing a process.

```
1 sleep 60 &
2 ps xw | grep sleep
3 # kill the process number
```

Here is how that looks in a session:

```
1 ubuntu@domU-12-31-39-16-1C-96:~$ sleep 20 &
2 [1] 5483
3 ubuntu@domU-12-31-39-16-1C-96:~$ ps xw | grep sleep
4  5483 pts/0    S      0:00 sleep 20
5  5485 pts/0    S+     0:00 grep --color=auto sleep
6 ubuntu@domU-12-31-39-16-1C-96:~$ kill 5483
7 [1]+  Terminated                  sleep 20
```

So you see that the process number 5483 was identified by printing the list of running processes with `ps xw`, finding the process ID for `sleep 20` via `grep`, and then running `kill 5483`. Note that the `grep` command itself showed up; that is a common occurrence and can be dealt with as follows:

```
1 ubuntu@domU-12-31-39-16-1C-96:~$ sleep 20 &
2 [1] 5486
3 ubuntu@domU-12-31-39-16-1C-96:~$ ps xw | grep sleep | grep -v "grep"
4  5486 pts/0    S      0:00 sleep 20
5 ubuntu@domU-12-31-39-16-1C-96:~$ kill 5486
```

Here, we used the `grep -v` flag to exclude any lines that contained the string `grep`. That will have some false positives (e.g. it will exclude a process named `foogrep`, so use this with some caution.

most important processes: top

Dashboard that shows which processes are doing what. Use `q` to quit.

```
1 top
```

stop a process: kill

Send a signal to a process. Usually this is used to terminate misbehaving processes that won't stop of their own accord.

```
1 sleep 60 &
2 # The following invocation uses backticks to kill the process we just
3 # started. Copy/paste it into a separate text file if you can't distinguish the
4 # backticks from the standard quote.
5 kill `ps xw | grep sleep | cut -f1 -d ' ' | head -1`
```

Superuser

become root temporarily: sudo

Act as the root user for just one or a few commands.

```
1 sudo apt-get install -y git-core
```

become root: su

Actually become the root user. Sometimes you do this when it's too much of a pain to type sudo a lot, but you usually don't want to do this.

```
1 touch normal
2 sudo su
3 touch rootfile
4 ls -alrth normal rootfile # notice owner of rootfile
```

Storage and Finding

create an archive: tar

Make an archive of files.

```
1 mkdir genome
2 mv *ptt* genome/
3 tar -cvf genome.tar genome
```

compress files: gzip

Compress files. Can also view compressed gzip files without fully uncompressing them with zcat.

```
1 gzip genome.tar
2 md temp
3 cp genome.tar.gz temp
4 cd temp
5 tar -xvzf genome.tar.gz
6 cd ..
```

```

7 gunzip genome.tar.gz
8 rm -rf genome.tar genome temp
9 wget ftp://ftp.ncbi.nih.gov/genomes/Homo_sapiens/CHR_22/hs_ref_GRCh37.p10_chr22.fa.gz
10 zcat hs_ref_GRCh37.p10_chr22.fa.gz | nl | head -1000 | tail -20

```

find a file (non-indexed): find

Non-indexed search. Can be useful for iterating over all files in a subdirectory.

```

1 find /etc | nl

```

find a file (indexed): locate

For locate to work, updatedb must be operational. This runs by default on Ubuntu but you need to manually configure it for a mac.

```

1 sudo apt-get install -y locate
2 sudo updatedb # takes a little bit of time
3 locate fstab

```

system disk space: df

Very useful when working with large data sets.

```

1 df -Th

```

directory utilization: du

Use to determine which subdirectories are taking up a lot of disk space.

```

1 cd ~ubuntu
2 du --max-depth=1 -b | sort -k1 -rn

```

Intermediate Text Processing

searching within files: grep

Powerful command which is worth learning in great detail. Go through `grep --help` and try out many more of the options.

```

1 wget ftp://ftp.ncbi.nih.gov/genomes/Bacteria/\
2 Escherichia_coli_K_12_substr__W3110_uid161931/NC_007779.ptt
3 grep protein *ptt | wc -l # lines containing protein
4 grep -l Metazoa *ptt *gbk # print filenames which contain 'Metazoa'
5 grep -B 5 -A 5 Metazoa *gbk
6 grep 'JOURNAL.*' *gbk | sort | uniq

```

simple substitution: sed

Quick find/replace within a file. You should review this outstanding set of [useful sed one-liners](#). Here is a simple example of using `sed` to replace all instances of `kinase` with `STANFORD` in the first 10 lines of a `ptt` file, printing the results to STDOUT:

```
1 head *ptt | sed 's/kinase/STANFORD/g'
```

`sed` is also useful for quick cleanups of files. Suppose you have a file which was saved on Windows:

```
1 # Convert windows newlines into unix; use if you have such a file
2 wget http://startup-class.s3.amazonaws.com/windows-newline-file.csv
```

Viewing this file in `less` shows us some `^M` characters, which are Windows-format newlines (`\r`), different from Unix newlines (`\n`).

```
1 $ less windows-newline-file.csv
2 30,johnny@gmail.com,Johnny Walker,,^M1456,jim@gmail.com,Jim Beam,,^M2076,...
3 windows-newline-file.csv (END)
```

The `\r` is interpreted in Unix as a “carriage return”, so you can’t just `cat` these files. If you do, then the cursor moves back to the beginning of the line everytime it hits a `\r`, which means you don’t see much. Try it out:

```
1 cat windows-newline-file.csv
```

You won’t see anything except perhaps a blur. To understand what is going on in more detail, read [this post](#) and [this one](#). We can fix this issue with `sed` as follows:

```
1 ubuntu@domU-12-31-39-16-1C-96:~$ sed 's/\r/\n/g' windows-newline-file.csv
2 30,johnny@gmail.com,Johnny Walker,,
3 1456,jim@gmail.com,Jim Beam,,
4 2076,jack@stanford.edu,Jack Daniels,,
```

Here `s/\r/\n/g` means “replace the `\r` with `\n` globally” in the file. Without the trailing `g` the `sed` command would just replace the first match. Note that by default `sed` just writes its output to STDOUT and does not modify the underlying file; if we actually want to modify the file in place, we would do this instead:

```
1 sed -i 's/\r/\n/g' windows-newline-file.csv
```

Note the `-i` flag for in-place replacement. Again, it is worth reviewing this list of [useful sed one-liners](#).

advanced substitution/short scripts: awk

Useful scripting language for working with tab-delimited text. Very fast for such purposes, intermediate size tool.

```
1 tail -n+4 *ptt | awk -F"\t" '{print $2, $3, $3 + 5}' | head
```

To get a feel for `awk`, review this list of [useful awk one liners](#). It is often the fastest way to operate on tab-delimited data before importation into a database.

Intermediate bash

Keyboard shortcuts

Given how much time you will spend using `bash`, it's important to learn all the [keyboard shortcuts](#).

- *Ctrl + A*: Go to the beginning of the line you are currently typing on
- *Ctrl + E*: Go to the end of the line you are currently typing on
- *Ctrl + F*: Forward one character.
- *Ctrl + B*: Backward one character.
- *Meta + F*: Move cursor forward one word on the current line
- *Meta + B*: Move cursor backward one word on the current line
- *Ctrl + P*: Previous command entered in history
- *Ctrl + N*: Next command entered in history
- *Ctrl + L*: Clears the screen, similar to the `clear` command
- *Ctrl + U*: Clears the line before the cursor position. If you are at the end of the line, clears the entire line.
- *Ctrl + H*: Same as backspace
- *Ctrl + R*: Lets you search through previously used commands
- *Ctrl + C*: Kill whatever you are running
- *Ctrl + D*: Exit the current shell
- *Ctrl + Z*: Puts whatever you are running into a suspended background process. `fg` restores it.
- *Ctrl + W*: Delete the word before the cursor
- *Ctrl + K*: Kill the line after the cursor
- *Ctrl + Y*: Yank from the kill ring

- *Ctrl + _*: Undo the last bash action (e.g. a yank or kill)
- *Ctrl + T*: Swap the last two characters before the cursor
- *Meta + T*: Swap the last two words before the cursor
- *Tab*: Auto-complete files and folder names

Let's go through a brief screencast of the most important shortcuts.

Backticks

Sometimes you want to use the results of a bash command as input to another command, but not via STDIN. In this case you can use [backticks](#):

```
1 echo `hostname`
2 echo "The date is "`date`"
```

This is a useful technique to compose command line invocations when the results of one is the argument for another. For example, one command might return a hostname like `ec2-50-17-88-215.compute-1.amazonaws.com` which can then be passed in as the `-h` argument for another command via backticks.

Running processes in the background: `foo &`

Sometimes we have a long running process, and we want to execute other commands while it completes. We can put processes into the background with the ampersand symbol as follows.

```
1 sleep 50 &
2 # do other things
```

Here is how that actually looks at the command line:

```
1 ubuntu@domU-12-31-39-16-1C-96:~$ sleep 60 &
2 [1] 5472
3 ubuntu@domU-12-31-39-16-1C-96:~$ head -2 *ptt
4 Escherichia coli str. K-12 substr. W3110, complete genome - 1..4646332
5 4217 proteins
6 ubuntu@domU-12-31-39-16-1C-96:~$ ps xw | grep 5472
7   5472 pts/0    S      0:00 sleep 60
8   5475 pts/0    S+     0:00 grep --color=auto 5472
9 ubuntu@domU-12-31-39-16-1C-96:~$
10 ubuntu@domU-12-31-39-16-1C-96:~$
11 [1]+  Done                  sleep 60
```

Note that we see `[1] 5472` appear. That means there is one background process currently running (`[1]`) and that the ID of the background process we just spawned is `5472`. Note also at the end that when the process is done, this line appears:


```
1 [1]+  Done                  sleep 60
```

That indicates which process is done. For completeness, here is what it would look like if you had multiple background processes running simultaneously, and then waited for them all to end.

```
1 ubuntu@domU-12-31-39-16-1C-96:~$ sleep 10 &
2 [1] 5479
3 ubuntu@domU-12-31-39-16-1C-96:~$ sleep 20 &
4 [2] 5480
5 ubuntu@domU-12-31-39-16-1C-96:~$ sleep 30 &
6 [3] 5481
7 ubuntu@domU-12-31-39-16-1C-96:~$
8 [1]  Done                  sleep 10
9 [2]-  Done                  sleep 20
10 [3]+  Done                  sleep 30
```

Execute commands from STDIN: `xargs`

This is a very powerful command but a bit confusing. It allows you to programmatically build up command lines, and can be useful for spawning parallel processes. It is commonly used in combination with the `find` or `ls` commands. Here is an example which lists all files under `/etc` ending in `.sh`, and then invokes `head -2` on each of them.

```
1 find /etc -name '*.sh' | xargs head -2
```

Here is another example which invokes the `file` command on everything in `/etc/profile.d`, to print the file type:

```
1 ls /etc/profile.d | xargs file
```

A savvy reader will ask why one couldn't just do this:

```
1 file /etc/profile.d/*
```

Indeed, that will produce the same results in this case, and is feasible because only one directory is being listed. However, if there are a huge number of files in the directory, then `bash` will be unable to expand the `*`. This is common in many large-scale applications in search, social, genomics, etc and will give the dreaded **Argument list too long** error. In this case `xargs` is the perfect tool; see [here](#) for more.

Pipe and redirect: `tee`

Enables you to save intermediate stages in a pipeline. Here's an example:

```
1 ls | tee list.txt
```

The reason it is called `tee` is that it is like a “T-junction”, where it passes the data through while also serializing it to a file. It can be useful for backing up a file while simultaneously modifying it; see some [examples](#).

time any command: `time`

This is a `bash` built in useful for benchmarking commands.

```
1 time sleep 5
```

That should echo the fact that the command took (almost) exactly 5 seconds to complete. It is also useful to know about a more sophisticated GNU `time` command, which can output a ton of useful information, including maximum memory consumption; extremely useful to know before putting a job on a large compute cluster. You can invoke it with `/usr/bin/time`; see documentation for Ubuntu 12 LTS [here](#).

Summary

We covered quite a few command line programs here, but nothing substitutes for running them on your own and looking at the online help and flags. For example, with `head` you can do `head --help`, `man head`, and `info head` to get more information.