### Do more and type less on the command line

Helen Cook February 2018



### We are not here because we like typing

Rather we use the command line for its flexibility and power

# We are not here because I want to teach you about the shell

(although, that will happen too)

Rather because I want to inspire you to customize and improve your tools

#### New stuff first!

Ask questions about anything!

Shout if you have a better way!

### 3 different ways to extend your command toolbox

```
sam (alias/function)
newest (function)
tab (script)
```

A new, perhaps better, way to do all this stuff

### How to always invoke Is with -I argument?

alias II='Is -I'

alias II='Is -I -color=auto'

### How to randomly sample lines from a file?

Usage: cat file | sam



create alias for this part

There is a better way to do this, we'll do that next

#### sam — outline

- 1.
- 2. Randomize the lines in the file
- 3. Select the top 10 lines
- 4.
- 5.

#### sam — outline

- 1. Number the lines in the file
- 2. Randomize them
- 3. Select the top 10 lines
- 4. Sort them back into their original order
- 5. Remove the line numbers

### How to number the lines in a file?

#### nl

```
(helen@helens-mbp[ttys000])-(~/projects/type-less/examples)
(0)-(0)% head -5 chemicals
             carnitine
CID100000085
               gamma-aminobutyric acid (GABA
CID100000119
               5-aminolevulinic acid
CID100000137
               leucovorin
CID100000143
CID100000158
               prostaglandin E2
(helen@helens-mbp[ttys000])-(~/projects/type-less/examples)
(0)-(o)\% head -5 chemicals | nl
    1 CID100000085 carnitine
    2
       CID100000119
                       gamma-aminobutyric acid (GABA
    3
       CID100000137
                       5-aminolevulinic acid
                     leucovorin
       CID100000143
    4
                       prostaglandin E2
       CID100000158
```

### How to randomize the lines in a file?

#### shuf

```
(helen@helens-mbp[ttys000])-(~/projects/type-less/examples)
(0)-(o)% head -n 5 chemicals|shuf
CID100000158    prostaglandin E2
CID100000137    5-aminolevulinic acid
CID100000143    leucovorin
CID100000085    carnitine
CID100000119    gamma-aminobutyric acid (GABA)
```

### How to get the first N lines in a file?

#### head

```
(helen@helens-mbp[ttys000])-(~/projects/type-less/examples)
(0)-(o)% cat chemicals|nl|shuf|head -5
    49 CID100000853    thyroxine
1065 CID100119830    tenofovir disoproxil fumarate
    782 CID100006058    cysteamine
173 CID100002554    carbamazepine
1125 CID100159247    sevelamer hydrochloride
```

### How to sort the lines in a file?

#### sort

### How to remove columns from a file?

#### cut (or awk)

#### sam — outline

- 1. (nl) Number the lines in the file
- 2. (shuf) Randomize them
- 3. (head) Select the top 10 lines
- 4. (sort) Sort them back into their original order
- 5. (cut) Remove the line numbers

alias sam='nl | shuf | head | sort -n | cut -f2-'

#### As an alias, we can't do

sam file

### How to define a function?

```
sam () {
```

### How to get the arguments to a function?

# First argument is in \$1 Second is in \$2 etc...

### How to test if a file exists?

[-f \$file]

## [-f\$file] \spaces!!

### What is the syntax for if?

```
sam () {
  if [
               ]; then
  else
  fi
```

```
sam () {
  if [[ -f "$1" ]]; then
  else
```

```
sam () {
  if [[ -f "$1" ]]; then
     input="$1"
  else
```

# How do you read from STDIN?

cat -

```
sam () {
  if [[ -f "$1" ]]; then
     input="$1"
  else
     input="-"
```

```
sam () {
  if [[ -f "$1" ]]; then
     input="$1"
  else
     input="-"
  cat $input | nl | shuf | head | sort -n | cut -f2-
```

```
sam () {
  if [[ -f "$1" ]]; then
     input="$1"
  else
     input="-"
  cat $input | nl | shuf | head | sort -n | cut -f2-
```

```
(helen@sund29736[ttys009])-(~/projects/type-less/examples)
(0)-(0)% sam chemicals
                cladribine
CID100001546
                amlodipine
CID100002162
                cyproterone acetate
CID100002914
CID100003100
                diphenhydramine
                ethotoin
CID100003292
CID100003462
                gemeprost
                isoflurane
CID100003763
                lorazepam
CID100003958
                ceftobiprole
CID106918430
                canagliflozin
CID124812758
(helen@sund29736[ttys009])-(~/projects/type-less/examples)
(0)-(o)% cat chemicals|sam
                dihydrocodeine
CID100003063
                fluphenazine
CID100003372
CID100003696
                imipramine
                nisoldipine
CID100004499
                Milli
CID100004536
                strontium
CID100083606
                pyocin S3
CID100092725
                eslicarbazepine
CID100114709
                Humalog
CID116132438
                Cephalosporins
CID118609542
```

### For fun: adapt sam so it will work with very large files

### For fun: adapt sam so it will work with very large files

Hint: seek

# How to list the N newest files in a directory tree?

Usage: newest N

#### newest — outline

- 1. Find all the files in the directory tree
- 2. For each file ...
- 3. Get the last modified date
- 4. Sort files into the right order
- 5. Print only the top 10 lines

# Use a shell function to define "newest" that takes an optional argument

```
newest(){
```

# How to specify an optional argument?

```
newest () {
    lines=10
    if [[ $1 != "" ]]; then
        lines=$1
    fi
```

# How to list all files in a directory tree?

#### Is -R

Will list files recursively, but will order them by directory

#### find

find. -type f

find. -type f

find. -type f

```
(helen@sund29736[ttys009])-(~/projects/type-less/examples)
(0)-(o)% find . -type f
./chemicals
./solutions/sam.py
```

## How to get the last modified date for a file?

#### find . -type f -printf '%T@ %p \n'

```
(helen@Helens-MacBook-Pro[ttys007])-(~/Google Drive/Human-Virus-PPI/data)
(0)-(o)% find . -type f -printf '%T@ %p \n'|head
1520072281.00000000000 ./.DS_Store
1519736844.00000000000 ./.STRING-VIRUSES-INSTALL.swp
1515412339.0000000000 ./.swp
1464360694.0000000000 ./A
1504086376.00000000000 ./actions_inconsistencies
1418833586.0000000000 ./alberto-payload/create_payload.py
496841761.00000000000 ./alberto-payload/Icon
1418833739.0000000000 ./alberto-payload/stringrnautils.py
1418833742.00000000000 ./alberto-payload/stringrnautils.py
1464360700.00000000000 ./B
```

#### find . -type f -printf '%T@ %p \n'

UNIX time: Seconds since Jan 1 1970

```
newest () {
    lines=10
    if [[ $1 != "" ]]; then
        lines=$1
    fi
    find . -type f -printf '%T@ %p \n' |
```

## How to sort on the dates?

sort -n

```
newest(){
  lines=10
  if [[$1!=""]]; then
     lines=$1
  find. -type f -printf '%T@ %p \n' |
  sort -n |
```

# How to show the last \$lines lines?

```
newest(){
  lines=10
  if [[$1!=""]]; then
     lines=$1
  find. -type f -printf '%T@ %p \n' |
  sort -n
  tail -n $lines |
```

# How to show only the filenames?

cut

Field delimiter is a space

Print field 2 and onwards

```
newest(){
  lines=10
  if [[$1!=""]]; then
     lines=$1
  find. -type f -printf '%T@ %p \n'
  sort -n
  tail -n $lines |
  cut -f2- -d " "
```

```
newest(){
  local lines=10
  if [[ $1 != "" ]]; then
     lines=$1
  find . -type f -printf '%T@ %p \n' |
  sort -n
  tail -n $lines |
  cut -f2- -d " "
```

### But it doesn't work on my Mac

#### Gnu find has -printf BSD find doesn't



The missing package manager for macOS

http://brew.sh brew install coreutils

#### How to get the last modified date for a file?

stat -f "%m %N"

#### How to run a command on each file found?

#### xargs

```
find ... | xargs < cmd>
```

```
(helen@sund29736[ttys009])-(~/projects/type-less/examples)
(0)-(o)% find . -type f -print0 | xargs -0 ls -lt
-rwxr-xr-x  1 helen staff  1723 Mar  8 11:10 ./solutions/sam.py
-rw-r--r-  1 helen staff  35785 Feb  1 12:05 ./chemicals
```

```
newest(){
  lines=10
  if [[$1!=""]]; then
     lines=$1
  find . -type f -print0
  xargs -0 stat -f "%m %N"
  sort -n
  tail -n $lines
  cut -f2 -d " "
```

```
newest(){
  local lines=10
  if [[ $1 != "" ]]; then
     lines=$1
  find . -type f -print0 |
  xargs -0 stat -f "%m %N" |
  sort -n
  tail -n $lines |
  cut -f2 -d " "
```

#### How to avoid awk for tab delimited files?

awk -F"\t" '{print \$3 "\t" \$5 "\t" \$3 "\t"}'

awk -F"\t" '{print \$3 "\t" \$5 "\t" \$3 "\t"}'

This is terrible

Let's replace:

awk -F"\t" '{print \$3 "\t" \$5 "\t" \$3 "\t"}'

With:

tab 3 5 2

#### tab — outline

- 1. Get columns to print from cmd line
- 2. Build up awk print string
- 3. Run awk

#### Use a script to define "tab"

#### Pass the column numbers as arguments

#!/bin/sh

### How to get the arguments to a script?

# First argument is in \$1 Second is in \$2 etc...

## How to iterate through all arguments?

#### while \$1 is not empty: shift the arg array to move the next arg into \$1

#### What is the syntax for while condition in sh?

```
while [ ... ]; do
...
done
```

#### MANDATORY SPACES!! while [ ]; do done

#!/bin/sh

while [ ]; do

done;

#!/bin/sh

```
while [ "$1" != "" ]; do
```

shift; done;

### How to construct the awk string?

```
1
1 \t 2
1 \t 2 \t 3
```

... and get the quoting correct

```
#!/bin/sh
```

```
first=1
while [ "$1" != "" ]; do
  if [\$first == 1]; then
     first=0
  else
  shift;
done;
```

### How to concatenate strings in sh?

```
str = "some"
str = "$str"more
```

```
#!/bin/sh
printstring=""
first=1
while [ "$1" != "" ]; do
  if [\$first == 1]; then
     first=0
   else
  shift;
done;
```

```
#!/bin/sh
printstring=""
first=1
while [ "$1" != "" ]; do
  if [\$first == 1]; then
     printstring=
     first=0
  else
     printstring=
  shift;
done;
```

```
#!/bin/sh
printstring=""
first=1
while [ "$1" != "" ]; do
  if [\$first == 1]; then
     printstring="$"$1
     first=0
  else
     printstring=
  shift;
done;
```

```
#!/bin/sh
printstring=""
first=1
while [ "$1" != "" ]; do
  if [\$first == 1]; then
     printstring="$"$1
     first=0
   else
     printstring=$printstring'"\t"
  shift;
done;
```

```
#!/bin/sh
printstring=""
first=1
while [ "$1" != "" ]; do
  if [\$first == 1]; then
     printstring="$"$1
     first=0
   else
     printstring=$printstring'"\t""$"$1
  shift;
done;
```

```
#!/bin/sh
printstring=""
first=1
while [ "$1" != "" ]; do
  if [\$first == 1]; then
     printstring="$"$1
     first=0
   else
     printstring=$printstring'"\t""$"$1
  shift;
done;
awk -F"\t" '{print '$printstring'}'
```

```
#!/bin/sh
printstring=""
first=1
while [ "$1" != "" ]; do
   if [\$first == 1]; then
     printstring="$"$1
     first=0
   else
     printstring=$printstring'"\t""$"$1
   shift;
done;
awk -F"\t" '{print '$printstring'}'
```

#### Alias, function or script?

#### Alias

#### Just replaces text Best for changing defaults of commands

#### **Function**

Takes arguments
Is in memory
Executes quickly
(autoload in some shells)

Script

Spawns new process
Can write in any language





about docs contributing releases support

name	department	duty_station	division	grade #	_job_title>
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CONEENEY, CHRISTOPHL	GENERAL SERVICES AL	WASHINGTON	OFFICE OF GOVERNME	14	REALTY
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CONERLY, CAROLYN E	VETERANS AFFAIRS	JOHNSON CITY	VETERANS HEALTH AD	0	SOCIAL WORK
CONERLY, EVELYN SUE		LAS VEGAS	ENVIRONMENTAL PROT	12	ENVIRONMENTAL PRO_
CONERLY, JOYCE A	HEALTH AND HUMAN S	BETHESDA	NATIONAL INSTITUTE	14	HUMAN RESOURCES ML
CONERLY, KATINA R	VETERANS AFFAIRS	NEW ORLEANS	VETERANS HEALTH AD	6	DENTAL ASSISTANT
CONERLY, LACI L	VETERANS AFFAIRS	ALEXANDRIA	VETERANS HEALTH AD	6	PRACTICAL NURSE
CONERLY, MARY C	VETERANS AFFAIRS	DALLAS	VETERANS HEALTH AD	2	NURSE
CONERLY, OCTAVIA D.		WASHINGTON	ENVIRONMENTAL PROT	13	GENERAL PHYSICAL _
CONERLY, OLIVER E	VETERANS AFFAIRS	BILOXI	NATIONAL CEMETERY	7	MISCELLANEOUS CLE.
CONERLY, YVETTE FRA	AGRICULTURE	NEW ORLEANS	OFFICE OF THE CHIE	12	INFORMATION TECHN
CONERLY, ZENA P	AGRICULTURE	WASHINGTON	FOREST SERVICE	12	INFORMATION TECHN_
CONERLY-SMITH, CATH	VETERANS AFFAIRS	NEW ORLEANS	VETERANS HEALTH AD	8	MEDICAL INSTRUMEN
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VisiData (vd on the command-line) is an <u>open-source</u> multitool for interacting with tabular data from within the terminal. It's great for investigative journalists, data scientists, unix command-line users, and anyone else who wants to quickly look at or manipulate data.

http://visidata.org/

(vi for spreadsheets)

#### Visidata tutorial:

https://jsvine.github.io/intro-to-visidata/

#### Exercises

Implement an alias, function or script for something that you do frequently (and share it!)

# A short explanation of the conventions used for the following exercises

### Questions are on blue slides

### Answers are on yellow slides

## Hints and explanations are on green slides

### Pitfalls/traps/avoid are on red slides

# Exercises are on purple slides

#### Code is on white slides

### How to not print output on STDERR?

2>/dev/null