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

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CONERLY, OLIVER E	VETERANS AFFAIRS	BILOXI	NATIONAL CEMETERY	7	MISCELLANEOUS CLE.
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ederalEmployeeSalaries-2012 selected 86425 rows					1350593 rows

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"

May be a better way to do all this data manipulation

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!)