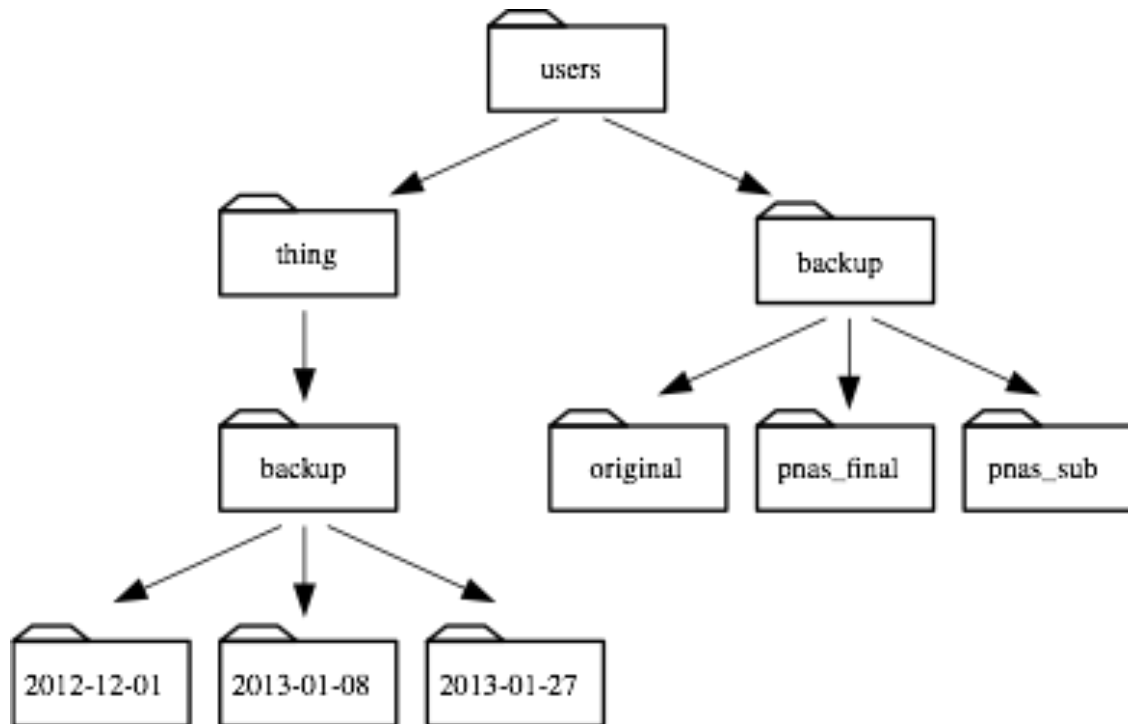


If `pwd` displays `/users/thing`, what will `ls ../backup` display?

- A. `../backup`: No such file or directory
- B. `2012-12-01 2013-01-08 2013-01-27`
- C. `2012-12-01/ 2013-01-08/ 2013-01-27/`
- D. `original pnas_final pnas_sub`



If `pwd` displays `/users/backup`, and `-r` tells `ls` to display things in reverse order, what command will display:  
`pnas-sub/ pnas-final/ original/`

- A. `ls pwd`
- B. `ls -r -F`
- C. `ls -r -F /users/backup`

What does the command `cd` without a directory name do?

- A. It has no effect.
- B. It changes the working directory to `/`.
- C. It changes the working directory to the user's home directory.
- D. It produces an error message.

What does the command `ls` do when used with `-s` and `-h` arguments?

- A. Lists only files that begin with `s` and `h`.
- B. Lists file sizes of files that begin with `h`.
- C. Lists ages of files in seconds .
- D. Lists file sizes in blocks.



# What is the output of the closing “ls”?

---

```
$ pwd
/home/jamie/data
$ ls
proteins.dat
$ mkdir recombine
$ mv proteins.dat recombine
$ cp recombine/proteins.dat ../proteins-saved.dat
$ ls
```

- A. proteins.dat
- B. proteins-saved.dat
- C. proteins.dat recombine
- D. recombine

Suppose that:

```
$ ls -F
analyzed/  fructose.dat  raw/  sucrose.dat
```

What command(s) could you run so that the commands below will produce the output shown?

```
$ ls
analyzed  raw
$ ls analyzed
fructose.dat  sucrose.dat
```

- A. `mv fructose.dat analyzed`  
`mv sucrose.dat analyzed`
- B. `cp fructose.dat analyzed`  
`cp sucrose.dat analyzed`
- C. `mv *.dat analyzed`

# What does **cp** do when given several filenames and a directory name, as in:

```
$ mkdir backup  
$ cp thesis/citations.txt thesis/quotations.txt backup
```

- A. Copies both .txt files into backup/
- B. Copies thesis/citations.txt onto thesis/quotations.txt, ignores backup
- C. Copies thesis/citations.txt onto thesis/quotations.txt and into backup /



# What does **cp** do when given three or more filenames, as in:

```
$ ls -F  
intro.txt      methods.txt    survey.txt  
$ cp intro.txt methods.txt survey.txt
```

- A. Copies intro.txt files over methods.txt and survey.txt
- B. Gives an error message and doesn't do anything
- C. Copies intro.txt files over methods.txt and copies methods.txt over survey.txt



How can we sort

```
10
2
19
22
6
```

and get

```
s 2
6
10
19
22
```

- A. `sort file.txt`
- B. `sort -n file.txt`
- C. `Sort -number file.txt`
- D. Either A or B

How many lines are in the output from  
running `uniq` on

```
coho  
coho  
steelhead  
coho  
steelhead  
steelhead
```

A. 2

B. 4

A file called `animals.txt` contains the following data:

```
2012-11-05,deer  
2012-11-05,rabbit  
2012-11-05,raccoon  
2012-11-06,rabbit  
2012-11-06,deer  
2012-11-06,fox  
2012-11-07,rabbit  
2012-11-07,bear
```

```
cat animals.txt | head -5 | tail -3 | sort -r > final.txt
```

**What's the first line in `final.txt`?**

- A. 2012-11-05,deer
- B. 2012-11-07,bear
- C. 2012-11-05-raccoon
- D. 2012-11-06-rabbit

The command:

```
$ cut -d , -f 2 animals.txt
```

produces the following output:

```
deer  
rabbit  
raccoon  
rabbit  
deer  
fox  
rabbit  
bear
```

What can we do with the output to get a list of animals with no duplicates?

- A. | sort
- B. | uniq | sort
- C. | uniq
- D. | sort | uniq



Suppose that `ls` initially displays:

```
fructose.dat    glucose.dat    sucrose.dat
```

What is the output of:

```
for datafile in *.dat
do
    ls *.dat
done
```

- A. fructose.dat glucose.dat sucrose.dat
- B. fructose.dat glucose.dat sucrose.dat  
fructose.dat glucose.dat sucrose.dat  
fructose.dat glucose.dat sucrose.dat
- C. fructose.dat  
glocuse.dat  
sucrose.dat



In the same directory, what does this loop do?

```
for sugar in *.dat
do
    echo $sugar
    cat $sugar > xylose.dat
done
```

- A. Prints fructose.dat, glucose.dat, and sucrose.dat, and copies sucrose.dat to create xylose.dat.
- B. Prints fructose.dat, glucose.dat, and sucrose.dat, and concatenates all three files to create xylose.dat.
- C. Prints fructose.dat, glucose.dat, sucrose.dat, and xylose.dat, and copies sucrose.dat to create xylose.dat.

# What is the output of

```
for left in 2 3
do
    for right in $left
    do
        expr $left + $right
    done
done
```

A. 2 4 3 6

B. 2 3

C. 4 6

D. 5 5

---

```
$ expr 3 + 5
```

```
8
```

```
$ expr 30 / 5 - 2
```

```
4
```



Suppose that `ls` initially displays:

```
fructose.dat    glucose.dat    sucrose.dat
```

What would **bash example.sh \*.dat** produce for `example.sh`

```
# Script 1  
echo *.*
```

- A. fructose.dat glucose.dat sucrose.dat
- B. fructose.dat glucose.dat sucrose.dat  
fructose.dat glucose.dat sucrose.dat  
fructose.dat glucose.dat sucrose.dat
- C. fructose.dat  
glucose.dat  
sucrose.dat

Suppose that `ls` initially displays:

```
fructose.dat    glucose.dat    sucrose.dat
```

What would **bash example.sh \*.dat** produce for **example.sh**

```
# Script 2
for filename in $1 $2 $3
do
    cat $filename
done
```

- A. Prints the contents of fructose.dat, glucose.dat and sucrose.dat
- B. Prints the contents of fructose.dat, glucose.dat and sucrose.dat, 3 times
- C. Combines fructose.dat and glucose.dat and overwrites them into sucrose.dat



**Grep -v junk file.txt** finds lines which don't contain "junk".

Which of the following commands will find all files in /data whose names end in ose.dat (e.g., sucrose.dat or maltose.dat), but do *not* contain the word temp?

A. `rm $(find ./ -name '*.tmp')`

How could you find and remove all of the .tmp files in your thesis/ directory and its subdirectories?

- A. `rm $(find ./ -name '*.tmp')`
- B. `rm *.tmp`
- C. `find ./ -name '*.tmp' | rm`
- D. `rm -i $(find ./ -name '*.tmp')`