

1. Imagine that we have just typed a shell script into the file `my_first_shell_script.sh` in the current directory. We then attempt to execute the script and observe the following:

```
$ my_first_shell_script.sh
my_first_shell_script.sh: command not found
```

Explain the possible causes for this, and describe how to rectify them.

2. Implement a shell script called `seq.sh` for writing sequences of integers onto its standard output, with one integer per line. The script can take up to three arguments, and behaves as follows:
- **seq.sh LAST** writes all numbers from 1 up to **LAST** inclusive, for example:

```
$ ./seq.sh 5
1
2
3
4
5
```

- **seq.sh FIRST LAST** writes all numbers from **FIRST** up to **LAST** inclusive, for example:

```
$ ./seq.sh 2 6
2
3
4
5
6
```

- **seq FIRST INCREMENT LAST** writes the sequence **FIRST**, **FIRST + INCREMENT**, **FIRST + 2 * INCREMENT**, up to **p** (where **p** is the largest integer in this sequence that is less than or equal to **LAST**), for example:

```
$ ./seq.sh 3 5 24
3
8
13
18
23
```

3. Write a shell script, **no_blinking.sh**, which removes all HTML files in the current directory which use the [blink element](#):

```
$ no_blinking.sh
Removing old.html because it uses the <blink> tag
Removing evil.html because it uses the <blink> tag
Removing bad.html because it uses the <blink> tag
```

Modify the shell script to instead take the HTML files to be checked as command line arguments and instead of removing them, add the suffix **.bad** to their name.

```
$ no_blinking.sh awful.html index.html terrible.html
Renaming awful.html to awful.html.bad because it uses the <blink> tag

Renaming terrible.html to terrible.html.bad because it uses the <blink> tag
```

4. Write a shell script, **list_include_files.sh**, which for all the C source files (**.c** files) in the current directory prints the names of the files they include (**.h** files), for example

```
$ list_include_files.sh
count_words.c includes:
    stdio.h
    stdlib.h
    ctype.h
    time.h
    get_word.h
    map.h
get_word.c includes:
    stdio.h
    stdlib.h
map.c includes:
    get_word.h
    stdio.h
    stdlib.h
    map.h
```

5. Consider the following columnated (space-delimited) data file containing contact information for various CSE academic staff:

G Heiser	Newtown	9381-1234
S Jha	Kingsford	9621-1234
C Sammut	Randwick	9663-1234
R Buckland	Randwick	9663-9876
J A Shepherd	Botany	9665-4321
A Taylor	Glebe	9692-1234
M Pagnucco	North Ryde	9868-6789

Note: This data is fictitious. Do not ring these phone numbers. I have no idea whether they are real or not, but they are certainly not the correct phone numbers for the academic staff mentioned.

The data is currently sorted in phone number order. Can we use the **sort** filter to re-arrange the data into "telephone-book" order? If not, how would we need to change the file in order to achieve this?

6. Consider the Unix password file (**/etc/passwd**):

```
root:ZHolHAHZw8As2:0:0:root:/root:/bin/bash
jas:iaiSHX49Jvs8.:100:100:John Shepherd:/home/jas:/bin/bash
andrewt:rX9KwSSPqkLyA:101:101:Andrew Taylor:/home/andrewt:/bin/cat
postgres:997:997:PostgreSQL Admin:/usr/local/pgsql:/bin/bash
oracle:999:998:Oracle Admin:/home/oracle:/bin/bash
cs2041:rX9KwSSPqkLyA:2041:2041:COMP2041 Material:/home/cs2041:/bin/bash
cs3311:mLRiCIvmtI902:3311:3311:COMP3311 Material:/home/cs3311:/bin/bash
cs9311:fIVLdSXYoVFai:9311:9311:COMP9311 Material:/home/cs9311:/bin/bash
cs9314:nTn.JwDgZE1Hs:9314:9314:COMP9314 Material:/home/cs9314:/bin/bash
cs9315:sOMXwkqmFbKlA:9315:9315:COMP9315 Material:/home/cs9315:/bin/bash
```

Provide a command that would produce each of the following results:

- display the first three lines of the file
- display lines belonging to class accounts
(assume that their login name starts with either "cs", "se", "bi" or "en", followed by a digit)
- display the user name of everyone whose shell is **/bin/bash**
- create a tab-separated file **passwords.txt** containing only login name and password for all users

7. The following shell script emulates the **cat** command using the built-in shell commands **read** and **echo**.

```
#!/bin/sh
while read line
do
    echo "$line"
done
```

- what are the differences between the above script and the real **cat** command?
 - modify the script so that it can concatenate multiple files from the command line, like the real **cat**
- (Hint: shell control structures (e.g. **if**, **while**, **for**) are commands in their own right and can form a component of a pipeline)

8. The **gzip** command compresses a text file and renames it to **oldName.gz**. The **zcat** command takes the name of a single compressed file as its argument and writes the original (non-compressed) text to its standard output.

Write a shell script called **zshow** that takes multiple **.gz** file names as its arguments, and displays the original text of each file, separated by the name of the file.

Consider the following example execution of **zshow**:

```
$ zshow a.gz b.gz bad.gz c.gz
===== a =====
... original contents of file "a" ...
===== b =====
... original contents of file "b" ...
===== bad =====
No such file: bad.gz
===== c =====
... original contents of file "c" ...
```

9. Consider the marks data file from last week's tutorial, and assume that it is stored in a file called **Marks**.

```
2111321 37 FL
2166258 67 CR
2168678 84 DN
2186565 77 DN
2190546 78 DN
2210109 50 PS
2223455 95 HD
2266365 55 PS
...
```

Assume also that we have a file called **Students** that contains the names and student ids of for all students in the class, e.g.

```
2166258 Chen, X
2186565 Davis, PA
2168678 Hussein, M
2223455 Jain, S
2190546 Phan, DN
2111321 Smith, JA
2266365 Smith, JD
2210109 Wong, QH
...
```

Write a shell script that produces a list of names and their associated marks, sorted by name, e.g.

```
67 Chen, X
77 Davis, PA
84 Hussein, M
95 Jain, S
78 Phan, DN
37 Smith, JA
55 Smith, JD
50 Wong, QH
```

Note: there are many ways to do this, generally involving combinations of filters such as **cut**, **egrep**, **sort**, **join**, etc. Try to think of more than one solution and discuss the merits of each.

10. Implement a shell script called **grades.sh** that reads a sequence of (studentID, mark) pairs from its standard input and writes (studentID, grade) pairs to its standard output. The input pairs are written on a single line, separated by spaces, and the output should use a similar format. The script should also check whether the second value on each line looks like a valid grade, and output an appropriate message if it doesn't. The script can ignore any extra data occurring after the mark on each line.

Consider the following input and corresponding output to the program:

Input

```
2212345 65
2198765 74
2199999 48
2234567 50 ok
2265432 99
2121212 hello
2222111 120
2524232 -1
```

Output

```
2212345 CR
2198765 CR
2199999 FL
2234567 PS
2265432 HD
2121212 ?? (hello)
2222111 ?? (120)
2524232 ?? (-1)
```

To get you started, here is a framework for the script:

```
#!/bin/sh
while read id mark
do
    # insert mark/grade checking here
done
```

Note that the shell's **read** operation assumes that the components on each input line are separated by spaces. How could we use this script if the data was supplied in a file that used commas to separate the (studentID, mark) components, rather than spaces?

11. Write a shell script **time_date.sh** that prints the time and date once an hour. It should do this until a new month is reached. Reminder the **date** command produces output like this:

```
Friday 5 August 17:37:01 AEST 2016
```

12. Consider a scenario where we have a directory containing two LaTeX files, **a.tex** and **b.tex**. The file **a.tex** is 20 lines long, and **b.tex** is 30 lines long. What is the effect of each of the commands below? How will their output differ?

```
$ wc -l *.tex
$ echo `wc -l *.tex`
```

13. Write a shell script that displays the name and size of all files in the current directory that are bigger than (say) 100,000 bytes.

(Hint: use **wc** to do the counting, and capture its output using back-ticks. How do you get rid of the file name and/or line and word counts?)

14. What is the output of each of the following pipelines if the text

```
this is big Big BIG
but this is not so big
```

is supplied as the initial input to the pipeline?

- `tr -d ' ' | wc -w`
- `tr -cs 'a-zA-Z0-9' '\n' | wc -l`
- `tr -cs 'a-zA-Z0-9' '\n' | tr 'a-z' 'A-Z' | sort | uniq -c`

15. Consider the standard "split-into-words" technique from the previous question:

```
tr -c -s 'a-zA-Z0-9' '\n' < someFile
```

Explain how this command works (i.e. what does each argument do)

16. Assume that we are in a shell where the following shell variable assignments have been performed, and **ls** gives the following result:

```
$ x=2 y='Y Y' z=ls
$ ls
      a          b          c
```

What will be displayed as a result of the following **echo** commands:

- `$ echo a b c`
- `$ echo "a b c"`
- `$ echo $y`
- `$ echo x$x`
- `$ echo $xx`
- `$ echo "$y"`
- `$ echo '$y'`
- `$ echo ` $y ``
- `$ echo ` $z ``
- `$ echo `echo a b c``

17. The following C program and its equivalent in Java both aim to give precise information about their command-line arguments.

C:

```
// Display command line arguments, one per line
#include <stdio.h>
int main(int argc, char *argv[]) {
    int i;
    printf("#args = %d\n", argc-1);
    for (i = 1; i < argc; i++)
        printf("arg[%d] = \"%s\"\n", i, argv[i]);
    return 0;
}
```

Java:

```
public class args {
    public static void main(String args[]) {
        System.out.println("#args = " + args.length);
        for (int i = 0; i < args.length; i++)
            System.out.println("arg[" + (i+1) + "] = \"" + args[i] + "\"");
    }
}
```

Assuming that the C program is compiled into a command called **args**, consider the following examples of how it operates:

```
$ args a b c
#args = 3
arg[1] = "a"
arg[2] = "b"
arg[3] = "c"
$ args "Hello there"
#args = 1
arg[1] = "Hello there"
```

Assume that we are in a shell where the following shell variable assignments have been performed.

```
$ x=2 y='Y Y' z=ls
```

Assume that we are in a shell with the same variable assignments and the same current directory as the previous question. What will be the output of the following:

a.

b.

c.

d.

e.

f.

g.

h.

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