

UNIX/Linux Operating System

Bash script exercises

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Write a bash script that computes the values of a function f(x) for all the triples of integer values stored in a file

```
    File content
    F(x) = 3 \cdot x^2 + 4 \cdot y + 5 \cdot z

Example
1    1    2    17
2    1    3    31
Output values
3    35

File content
```

Write two versions of the script using while and for statements, respectively

Using a for loop

Reads from file **one value at a time** because the output of the command goes in a list of strings

```
#!/bin/bash

iter=1
for val in `cat $1`
do
   if [ $iter -eq 1 ]
   then
   let f=3*val*val
```

```
elif [ $iter -eq 2 ]
   then
     let f=f+4*val
   elif [ $iter -eq 3 ]
   then
     let f=f+5*val
     iter=0
     echo -n "$f "
   fi
   let iter=iter+1
done
exit 0
```

Using the while loop (reads a line at a time)

Reads a line and the string is assigned to variable **line**

```
#!/bin/bash
```

```
while read line
do
   iter=1
   for val in $line
   do
```

Parsing the line

```
if [ $iter -eq 1 ]
     then
       let f=3*val*val
     elif [ $iter -eq 2
     then
       let f=f+4*val
     elif[ $iter -eq 3 ]
     then
       let f=f+5*val
     fi
     let iter=iter+1
   done
   echo -n "$f "
done < $1
```

Loop on file lines

Using the while loop (reads three values at a time)

```
#!/bin/bash

while read x y z
do
   f=3*x*x+4*y+5*z
   echo -n "$f "
done < $1
exit 0</pre>
```

Loop on file lines

- Write a bash script that
 - > Takes a filename from command line
 - Displays the file content
 - A line at a time, prepending the line number
 - A string at a time, prepending the string number

```
#!/bin/bash
n=1
while read line # read a line
do
 echo "$n: $line"
 let n=n+1
done < $1  # Redirection !</pre>
n=1
for str in `cat $1` # read a word
do
 echo "$n: $str"
 let n=n+1
done
```

Write a bash script that

- > Takes a filename from command line
- > Reads a sequence of integer number from the file
- > Each number represents a histogram bin value
- Displays a horizontal histogram using '*'
- > Example

| 1 | * | |
|---|------|--------|
| 3 | *** | |
| 5 | **** | |
| 4 | *** | Output |
| 2 | ** | Output |

File content

```
#!/bin/bash
                                      Reads a number at a time
for n in $(cat $1)
do
  i=1
  while [ $i -le $n ]
                                    Prints without
  do
                                      newline
    echo -n "*"
    let i=i+1
  done
                                   Prints a newline
  echo
done
exit 0
```

- Write a bash script that displays
 - > All files of the current directory
 - ➤ With ".c" extension
 - That include string "posix"
- Example of execution
 - > ./myScript .
 - ./myScript /home/bin

```
#!/bin/bash
for file in $(ls *.c); do
  grep --quiet "POSIX" $file
  if [ $? -eq 0 ]
  then
                                              grep
                             1) -q, --quiet, avoids printing the found line
    more $file
                             2) If a file is found, returns (echo $?) 0 i.e.,
  fi
                                the condition is TRUE
done
exit 0
# Alternative (single command):
                                                   grep -l
                                           displays only the filenames
# more $(grep -1 POSIX *.c)
                                           matching the string POSIX
# Notice the difference !!:
# grep -l POSIX *.c | more
```

- Write a bash script that
 - > Takes a filename from command line
 - > The file contains two columns of data
 - > Example

7 3

2 23

5 0

➤ The script must overwrite the file swapping the two columns

Uses a temporary file ...

```
#!/bin/bash
```

file="tmp"

while read var1 var2 do

echo \$var2 \$var1
done <\$1 >\$file

mv \$file \$1

... renamed at the end of the script

exit 0

- Write a bash script that
 - > Takes a set of strings from command line
 - > The first string is a directory name
 - > The others are filenames
 - \$myScript dir file1 file2 ... filen

The script

- Creates the directory if it does not exist
- For each file, ask the user if the file should be copied in the destination directory
- > Copy only files confirmed by the user

```
#!/bin/bash
if [ $# -le 1 ]
then
 echo "Run: $0 dir file1 file2 ..."
 exit 1
fi
if [ ! -d $1 ]
then
 echo "Create directory $1"
 mkdir $1
fi
```

```
dir=$1
                              The command line arguments
shift
                             are shifted to the left
for i in $*
do
  echo -n "$i in $dir (y/n)?"
  read choice
  if ["$choice" = "y" ]; then
    if cp $i $dir
    then
      echo "Copy done for $dir/$i"
    else
      echo "Error copying $i"
    fi
  fi
done
exit 0
```

Write a bash script that

- > Takes a filename (of a text file) from command line
- Copy the file with the same filename, but with extension xyx
- Modifies the original file
 - Adding at the beginning of each line the number of words in the line, and the total number of lines of the file
 - Sorting the lines according to their number of words

basename command

Syntax:

- basename NAME [SUFFIX]
- Prints NAME with any leading directory components removed. If specified, it will also remove a trailing suffix (typically a file extension)

```
> name='basename /home/user/current/file.txt txt'
> echo $name
file
```

```
#!/bin/bash
if [ $# -ne 1 ]
then
                                                     Filename without
  echo "usage $0 file.txt"
                                                        extension
  exit 1
fi
newfilename=$(basename $1 txt)
                                                      Better:
newfilename=$newfilename".xyz"
                                              nlines=\$(wc -l < \$1)
cp $1 $newfilename
nlines=$(cat $1 | wc -1)
rm -f tmp1.txt
                                                Add information on a
while read line
                                                   temporary file
do
  nwords=$(echo $line | wc -w)
  echo $nwords $nlines $line >> tmp1.txt
done < $1
cat tmp1.txt | sort -k 1 -n > $1
                                               Sort and overwrite the
rm tmp1.txt
                                                    original file
exit 0
                      Clean-up
```

Write a bash script that

- Takes 4 arguments (dir1, dir2 e dir3, directory names, and n, an integer number)
- Finds in dir1 and dir2 all files that have the same name, extension txt and more than n lines
- Creates in directory dir3 a version of these files with extension
 - eq save the lines that are equal in both files
 - dif save the lines that differ in the two files
 - cat concatenates the content of the two files

```
#!/bin/bash
if [ $# -ne 4 ]
then
  echo "usage: $0 dir1 dir2 dir3 n"
 exit 1
                                   find rather than Is
fi
                  `find $1 -maxdepth 1 -type f -name "*.txt"`
if [! -d $3]
then
 mkdir $3
fi
for file in $(ls $1/*.txt); do
  name=$(basename $file ".txt")
  if [ -f "$2/$name.txt" ]; then
    n1=\$(wc -l < \$file)
    n2=\$(wc -1 < "\$2/\$\{name\}.txt")
    if [ $n1 -gt $4 -a $n2 -gt $4 ]; then
```

Counts and controls the number of lines

```
Looks for this line
                                                   in the second file
      while read line; do
        grep -q -e "^$line$" "$2/$name.txt"
         if [ $? -eq 0 ]; then
           echo $line >> "$3/${name}.eq"
         else
                                                Looks for lines that are
           echo $line >> "$3/${name}.dif"
                                                      different
        fi
                                                  in the second file
      done < $file
      while read line; do
        grep -q -e "^$line$" "$3/${name}.eq"
        if [ $? -eq 1 ]; then
           echo $line >> "$3/${name}.dif"
        fi
      done < "$2/$name.txt"</pre>
      cat $file "$2/${name}.txt" > "$3/${name}.cat"
    fi
  fi
                                                  File concatenation
done
```