

```
#include <stdlib.h>
#include <string.h>
#include <ctype.h>
```

```
#define MAXPAROLA 30
#define MAXRIGA 80
```

```
int main(int argc, char *argv[])
```

```
{
    int freq[MAXPAROLA]; /* vettore di contatori
    delle frequenze delle lunghezze delle parole */
    char riga[MAXRIGA];
    int i, inizio, lunghezza;
    FILE *f;
```

```
for(i=0; i<MAXPAROLA; i++)
    freq[i]=0;
```

```
if(argc != 2)
{
    fprintf(stderr, "ERRORE, serve un parametro con il nome del file\n");
    exit(1);
}
```

```
f = fopen(argv[1], "r");
if(f==NULL)
{
    fprintf(stderr, "ERRORE, impossibile aprire il file %s\n", argv[1]);
    exit(1);
}
```

```
while( fgets( riga, MAXRIGA, f ) != NULL )
```



# UNIX/Linux Operating System

## Bash script exercises

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## Exercise

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

- $f(x) = 3 \cdot x^2 + 4 \cdot y + 5 \cdot z$

- Example

1	1	2	17
2	1	3	31
1	3	4	35

Output values

File content

- The name of the file must be passed from command line
- Write two versions of the script using **while** and **for** statements, respectively

# Solution 1

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
```

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

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

exit 0
```

## Solution 2

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
```

```
    flag=1
```

```
    for val in $line
```

```
    do
```

Parsing the line

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

Loop on file lines

# Solution 3

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

```
#!/bin/bash

while read x y z
do
    let f=3*x*x+4*y+5*z
    echo -n "$f "
done < $1

exit 0
```

Values read from  
file three at a time!

Loop on file lines

## Exercise

- ❖ Write a bash script that displays the content of
  - All files of the current directory
  - With ".c" extension
  - That include string "POSIX"



# Solution

```
#!/bin/bash
for file in $(ls *.c); do
    grep --quiet "POSIX" $file
    if [ $? -eq 0 ]
    then
        more $file
    fi
done
exit 0
```

```
# Alternative (single command):
# more $(grep  POSIX *.c -l)
# Notice the difference !!:
# grep -l POSIX *.c | more
```

grep

- 1) -q, --quiet, avoids printing the found line
- 2) If a file is found, returns (echo \$?) 0 i.e., the condition is TRUE

grep -l

displays only the filenames  
matching the string POSIX

## Exercise

❖ 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
- Note that output and input files are the same



# Solution

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

```
file="tmp"
```

```
while read var1 var2  
do
```

```
    echo $var2 $var1
```

```
done <$1 >$file
```

```
mv -f $file $1
```

```
exit 0
```

Uses a temporary file ...

... renamed at the end of the script

## Exercise

- ❖ 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

# Solution

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

# Exercise

## ❖ 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

File  
content

1	*
3	***
5	*****
4	****
2	**

Output

# Solution

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

```
for n in $(cat $1)
```

```
do
```

```
    i=1
```

```
    while [ $i -le $n ]
```

```
    do
```

```
        echo -n "*" 
```

```
        let i=i+1
```

```
    done
```

```
    echo
```

```
done
```

```
exit 0
```

Reads a number at a time

Prints without  
newline

Prints a newline

## Exercise

### ❖ 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 `dir`
- Copy only files confirmed by the user



# Solution

```
#!/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
```

# Solution 1

```
for i in $*
do
    if [ $i != $1 ]
    then
        echo -n "$i in $1 (y/n)? "
        read choice
        if [ $choice = "y" ] ; then
            cp $i $1
            if [ $? ]
            then
                echo "Copy done for $1/$i"
            else
                echo "Error for $1"
            fi
        fi
    fi
done
```

P.S.: \$\* does not include the name of the program

Skip the first parameter

```
fi
fi
done

exit 0
```

## Solution 2

```
dir=$1
shift
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
```

The command line arguments  
are shifted to the left

Exam Italian course: 2018/01/30

## Exercise: from exam

- ❖ The **df** file command shows the disk space available on the file system containing files
- ❖ Example

```
df /data/backup
```

```
Fifsystem 1K-blocks Used Avail Use% Mounted on  
/dev/sda7 41L1A492 5881472 33174616 16% /data
```

- The second, third and fourth fields show the total space used and available on the file system containing /data/backup
- Fields are separated with spaces
- Suppose no other separating character is used and spaces do not appear anywhere else

Exam Italian course: 2018/01/30

## Exercise: from exam

- ❖ Write a script that receives the path of a source file and a destination path, and
  - Check the correct passage of the parameters to the script
  - Make in background a copy of the source file in the destination path
  - Analyze the space occupied on the destination path at regular intervals of one second, displaying on the screen the percentage of progress of the copy operation

The command **sleep n** can be used to block the script for **n** seconds

# Solution

```
#!/bin/bash
if [ $# -ne 2 ]; then
    echo "Usage $0 <source> <destination>"
    exit 1
fi
if [ ! -f $1 ]; then
    echo "Source is not a valid file."
    exit 1
fi
if [ ! -d $2 ]; then
    echo "Destination is not a valid directory."
    exit 1
fi
source=$1
destination=$2
```

Check the number  
of parameters

Check the validity  
of the parameters



# Solution

```
size=$(ls -l $source | cut -d " " -f 5)
```

```
let "size=size/1024"
```

Calculate file size  
in 1KB blocks

```
startUsed = $(df $destination | \
               tail -n 1 | \
               tr -s " " | \
               cut -d " " -f 3)
```

Calculate  
destination file  
system size

```
cp $source $destination &
```

```
transferred=0
```

```
percentage=0
```

Copy in  
background

# Solution

```
while [ $transferred -lt $size ]; do
    currentUsed = $(df $destination | \
                    tail -n 1 | \
                    tr -s " " | \
                    cut -d " " -f 3)

    let "transferred=currentUsed-startUsed"
    let "percentage=transferred*100/size"
    echo "Progress: $percentage%"

    sleep 1
done
```

Check the state of  
the copy in  
background

Exam Italian course: 2018/02/22

## Exercise: from exam

- ❖ A script receives the following parameters
  - the name of a file (fn) and three integers (n1, n2, and n3)
  - The file (fn) specifies a path on each line
- ❖ The script must
  - Verify that the 4 parameters are correct, i.e., integer numbers must be positive, and  $n1 \leq n2$
  - For each row of the file (fn)
    - Check that each string refers to a regular file

Exam Italian course: 2018/02/22

**Exercise: from exam**

- If the dimension of the file (fn) is
- Smaller than n1 bytes, delete it
  - Between n1 and n2 bytes, ignore it
  - Greater than n2 bytes, compress it. Compress a file means
    - Make a copy in a file with the same path but with the additional extension (e.g., .compressed)
    - Modify the contents by copying only one string every n3 strings (ie just copy strings in position 0, 1 \* n3, 2 \* n3, etc.). Consider strings separated by spaces or by "newline" characters

# Solution

```
#!/bin/bash
if [ $# -ne 4 ]; then
    echo "Usage $0 <list> <n1> <n2> <n3>"
    exit 1
fi
if [ ! -f $1 ]; then
    echo "List is not a valid file."
    exit 1
fi
if [ $2 -lt 0 ] || [ $3 -lt 0 ] || [ $4 -lt 0 ]; then
    echo "Values n1, n2 and n3 should be non-negative integers."
    exit 1
fi
if [ $2 -gt $3 ]; then
    echo "Values n1 should be non-greater than n2."
    exit 1
fi
```

Check the number  
of parameters

Check the validity  
of the parameters

# Solution

```
while read file; do
```

For each path  
read from file

```
    if [ ! -f "$file" ]; then  
        echo "Invalid file: $file"  
        continue  
    fi
```

Skips paths not  
associated to  
regular files

```
fi
```

```
size=$(cat $file | wc -c)
```

Computes the  
dimension of the file

```
if [ $size -lt $2 ]; then  
    rm -f $file
```

Removes small files



# Solution

```
elif [ $size -gt $3 ]; then
    i=1
    for word in $(cat $file); do
        let "i--"
        if [ $i -eq 0 ]; then
            echo $word >> $file".compressed"
            i=$4
        fi
    done
fi
done < $1
```

Compresses big  
files

Exam Italian course: 2013/01/28

## Exercise

- ❖ 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)
```

```
> echo $name
```

```
file.txt
```

```
> name=$(basename /home/user/current/file.txt ".txt")
```

```
> echo $name
```

```
file
```

# Solution

```
#!/bin/bash
if [ $# -ne 1 ]
then
    echo "usage $0 file.txt"
    exit 1
fi
newfilename=$(basename $1 ".txt")
newfilename=$newfilename".xyz"
cat $1 > $newfilename
nlines=$(cat $1 | wc -l)
rm -f tmp1.txt
while read line
do
    nwords=$(echo $line | wc -w)
    echo $nwords $nlines $line >> tmp1.txt
done < $1
cat tmp1.txt | sort -k 1 -n > $1
rm tmp1.txt
exit 0
```

Filename without extension

".txt"="\*.txt"=.txt

Copy file. Also:  
cp \$1 \$newfilename

Also:  
nlines=\$(wc -l < \$1)

Add information on a temporary file

Sort and overwrite the original file

Clean-up

Exam Italian course: 2014/02/03

# Exercise

## ❖ 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
  - **diff** save the lines that differ in the two files
  - **cat** concatenates the content of the two files

Control the number of parameters.  
Create directory **dir3** if it does not exist

# Solution

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

```
if [ $# -ne 4 ]  
then
```

```
    echo "usage: $0 dir1 dir2 dir3 n"  
    exit 1  
fi
```

```
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=$(cat $file | wc -l)
```

```
        n2=$(wc -l < "$2/${name}.txt")
```

```
        if [ $n1 -gt $4 -a $n2 -gt $4 ]; then
```

find rather than ls

```
`find $1 -maxdepth 1 -type f -name "*.txt"`
```

It was enough to remove the path

For each .txt file in the first directory, generate the corresponding name in the second directory

Counts and controls the number of lines



# Solution

```
while read line; do
    grep -q -e "^$line$" "$2/$name.txt"
    if [ $? -eq 0 ]; then
        echo $line >> "$3/${name}.eq"
    else
        echo $line >> "$3/${name}.dif"
    fi
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"
cat $file "$2/${name}.txt" > "$3/${name}.cat"
fi
fi
done
```

Control  
on the  
result of  
grep  
\$?=0  
(true or  
find)

Lines in file1 and file2 go in eq,  
lines not in file2 go in dif

Inverse control for lines  
potentially in dif

File concatenation

## More exercises...

- ❖ Other examples and small exercises about bash
  - [https://www.skenz.it/cs/bash\\_language](https://www.skenz.it/cs/bash_language)