Lab3 Display content of the files. Pipeline.

Read carefully the instructions before performing!!!

Submission in pairs : 02/12/24 for the groups of Monday 05/12/24 for the groups of Thursday

- To learn how to see (to display) the files.
- · Copy and rename the file.
- The first using of the pipeline concept.

Linux Commands

1. Commands cat, more = display the contain of the file. cat file_name more file1 file2 file3 2. Command cp = Copy file 1 into file 2 Example: >ls foo1.c > **cp** foo1.c foo2.c >ls foo1.c foo2.c > cp -r dir1 dir2 = Copy recursively **3.** Command **mv** = Copy file 1 into file 2 = Rename file1 Example: >ls

foo1.c

> **mv** foo1.c foo2.c

>ls

foo2.c

4. Wildcards

4.1 * = matches zero or more character

Examples:

>Is *.txt = show all the files which finish with .text

>Is a* = show all the files which start with the letter a

>Is a*.txt = files which start with a and end with .text

4.2 ? = matches <u>exactly one</u> character

Examples:

>Is ?.txt = show all the files which start with any character and finish with .text

>Is a???.txt = show all the files which start with a , then any 3 characters and finish with .txt

> Is a??.*txt = show all the files which start with a, then any 2 characters, then ., then any text and finally it finishes with txt

4.3 [] - A character class (exactly 1 character from the range)

Examples:

> Is *.[aeiou] = show all the files which end with a, e, i, o or u. Like foo.a or boo.u

>ls ca[nt] = show can or cat

4.4 Ranges

 $[a-g]^* = Matches all files that start with a,c,..g$

[3-6]* = Matches all files that start with 3,4,5 or 6

4.5 Special symbols – use the symbol \

> Is *\? = Show all the files which finish with ?

- **5.** head file_name prints the <u>first lines</u> (default 10) of the file file_name head -n file name prints 3 first lines of file name
- **6. tail** file_name prints the last lines (default 10) of the file file_name tail **-n** file_name prints **n** last lines of file_name

tail +n file_name- prints all the lines starting with the n-th line of the file_name

7. pipeline |

Pipeline is a mechanism for connecting the output of one command to the input of another command

Example1: > cat file_name | head -3 | tail +2

Example2:

>cat foo1.txt

line 1

line 2

line 3

line 4

> cat foo1.txt| head -3 | tail +2

line 2

line 3

Procedures

- 1. Create a new directory Lab3.
- 2. In Lab3 create a new directory TryDir.
- 3. In TryDir create 3 files foo1.dat, foo2.g, boo3.gt6, boo.
- 4. Open the file foo1.dat (use pico) and write here your name, and id.
- 5. Open the file foo2.g and write the name of your best friend.
- 6. Display foo1.dat using cat and more.
- 7. Display both foo1.dat and foo2.g (Use, for example **cat** with 2 parameters)
- 8. Show the names of the files starting with the letter **f.**
- 9. Show the names of the files which contain the letter g.
- 10. Show the names of the files which contain a number.

- 12. Make a **copy** of the file foo1.dat and call it newFoo.
- 13. Display the contain of the file newFoo. Is it identical to foo1.dat?
- 14. **Rename** the file newFoo to be Foo2.
- 15. **Move** the file Foo2 to the directory Lab3.
- 16. Write in **lab3.sh** all the above (1-15) commands.
- 17. Show the first 5 lines of lab3.sh.
- 18. Show the last 5 lines of lab3.sh.
- 19. Show the lines 5-7 of lab3.sh (use cat, head, tail and pipeline).
- 20. Display the contain of the files which start with the letter **f** and then show the second line only of the result.

Submission:

Submit the file **lab3.sh** and results of the runs of the commands 1-20 in file **answers.txt**.

Note:

Create one compressed file named Linux_Lab3_id1_id2.tar, where id1 and id2 must be changed to the id numbers of the two partners.

This file will include the lab3.sh file and addtional text file named **answers.txt** with the answers to the questions.

To create such a file in Lunix command line write:

\$ tar -cvf Linux Lab3 id1 id2.tar lab3.sh answers.txt

Submit only the file Linux_Lab3_id1_id2.tar in Moodle and only by one partners.