CSc 3320: Systems Programming

Fall 2021 Homework

2: Total points 100

Submission instructions:

- 1. Create a Google doc for each homework assignment submission.
- 2. Start your responses from page 2 of the document and copy these instructions on page 1.
- 3. Fill in your name, campus ID and panther # in the fields provided. If this information is missing in your document TWO POINTS WILL BE DEDUCTED per submission.
- 4. Keep this page 1 intact on all your submissions. If this *submission's instructions* page is missing in your submission TWO POINTS WILL BE DEDUCTED per submission.
- 5. Each homework will typically have 2-3 PARTS, where each PART focuses on specific topic(s).
- 6. Start your responses to each PART on a new page.
- 7. If you are being asked to write code, copy the code into a separate txt file and submit that as well.
- 8. If you are being asked to test code or run specific commands or scripts, provide the evidence of your outputs through a screenshot and copy the same into the document.
- 9. Upon completion, download a .PDF version of the document and submit the same.

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PART I (2.5 points each): 10pts

1. What are the differences among *grep*, *egrep* and *fgrep*? Describe using an example.

grep: "Global Regular Expressions Print"

- grep is a program that scans a specific file or files line by line and returns lines with patterns. A pattern is an expression that identifies a set of strings by understanding characters as meta-characters.
- For instance, the asterisk meta character * is understood as "zero or more of the preceding element". This allows users to type characters and meta-characters into a grep command. This will have the computer show what lines in which file will match.
- standard grep command:
 - grep <flags> '<regular expression>' <filename>
- grep prints the search results to the screen (stdout) and returns this exit value:
 - O A match was found. 1 No match was found. >1 A syntax error was found or a file was inaccessible (even if matches were found)
- Common Flags:
 - -c: counts the number of matches but doesn't print actual matches
 - -i: makes searches case insensitive
 - -n: prints line number before each match printout
 - -l: prints file names of files with lines matching the expression

egrep: "Extended Global Regular Expressions Print"

- egrep handles patterns as regular expressions. It also handles +, ? , | , (, and) as meta-characters.
- Meta-characters lose their special meaning in grep. In order for grep to treat these characters as meta-characters, put a backslash before the character \+, \?, \|.
- grep using basic regular expression where the plus + is treated literally:
 - grep "+" file.txt
 - this returns any line with a plus +
- egrep treating the plus + as a meta-character and returns every line:
 - egrep "+" file.txt
- every line is returned since the plus + was treated by egrep as a meta-character; normal grep searches for only the lines with a plus +

fgrep: "Fixed-string Global Regular Expressions Print"

- fgrep, or *grep -F*, is a fixed/fast grep that behaves as a grep but doesn't recognize any regular expression meta-characters as anything special. This search completes faster because it only processes simple strings in comparison to complex patterns.
- When searching for a .bash_profile for a literal dot . using grep would be difficult. The dot . needs to be escaped because it's a meta-character meaning "wild-card, any single character".

```
- grep "." file.txt
```

- This command returns every line of the file.txt
 - fgrep "." file.txt
- Only the lines with a literal dot . in this command will be returned.
- fgrep helps by not bothering to escape the meta-characters.
- 2. Which utility can be used to compress and decompress files? And how to compress multiple files into a single file? Please provide one example for it.
 - tar:
 - tar: creates and extracts archives
 - tar archives without compression and then compresses with gzip. The file will then have an extension of .tar.gzip
 - Modes:
 - -c: creates an archive
 - -x: extracts an archive
 - Compress Single Directory/File:
 - Let there be a directory called *pictures* in the current directory that wants to save images. Below is the command:
 - \$ tar -czvf images.tar.gz pictures
 - -c: creates an archive
 - -z: compress the archive using gzip
 - -v: display progress in terminal
 - -f: specifying file name
 - Let there be a directory called *pictures* in /user/name/pictures that wants to save images. Below is the command:
 - \$ tar -czvf images.tar.gz /user/name/pictures

- <u>Compress Multiple Directories/Files:</u>
 - Let there be a directory called "pictures" at /user/name/pictures, "music" at /user/name/music, and "videos" at /user/name/videos and it would like to save as media. Below is the command:
 - \$ tar -czvf media.tar.gz /user/name/pictures
 /user/name/music /user/name/videos
 - Each path is separated by a space.

- <u>Decompress Archive:</u>

- Let there be an existing archive *media.tar.gz* that wants to decompress to the current directory. Below is the command:
- \$ tar -cxvf media.tar.gz
 - Mode is set to x for extraction
- To decompress to /user/name/files directory, this is the command to decompress:
- \$ tar -cxvf media.tar.gz -C /user/name/files
 - - C allows extracting to specific paths
- 3. Which utility (or utilities) can break a line into multiple fields by defining a separator? What is the default separator? How to define a separator manually in the command line? Please provide one example for defining the separator for each utility.
 - awk
 - When using <u>awk</u>, the way to supply the delimiter is either through -F or a FS= postfix:
 - awk -F \t' { print \$2 }' file OR awk \t' { print \$2 } $FS='\t'$ file
 - Output in all cases:
 - "Testing one, two, three"
 - "Testing one, two, three"
 - "Testing one, two, three"
 - "Testing one, two three"
 - Quote Delimiter: if the double quote in the file are consistent, they can be used as a delimiter and can avoid the output using *cut*

4. What does the **sort** command do? What are the different possible fields? Explain using an example.

- sort command can sort the input file and arrange the records in a specific order. There are different options to sort the data in Linux. Sorting can be done numerically, alphabetically, in reverse, and many other ways.
 - sort -0: writes the output to an output file
 - *sort -r*: sorts the file in reverse
 - sort -n: sorts the file numerically
 - sort -nr: sorts the file numerically in reverse
- Other Options:

- cut

- sort -k: sorts the file with a table; k is used to denote the table
- sort -u: sorting and removing duplicated at the same time
- sort -M: sorting records by month
- Example:
- Let the text file *color.txt* have

```
red
yellow
green
```

```
blue
purple

Command: sort -r color.txt

Sorted Output:
yellow
red
purple
green
blue
```

Part II a (5 points each): 25pts

5. What is the output of the following sequence of bash commands:

echo 'Hello World' | sed 's/\$/!!!/g'

- Hello World!!!
- 6. What is the output for each of these awk script commands?

```
-- 1 <= NF { print $5 }
- awk `1 <= NF { print $5 }'</pre>
```

- prints the fifth field/column if 1 is less than or equal to the number of fields (NF). If the number of fields is less than 5, then nothing will print

```
-- NR >= 1 && NR >= 5 { print $1 }
- awk 'NR >= 1 && NR >= 5 { print $1 }'
```

- prints the first column for all the rows greater than or equal to 1 *AND* greater than or equal to 5

```
-- 1,5 { print $0 }
- awk `1,5 { print $0 }'
```

- prints all the lines in a file as 1,5; evaluates TRUE

```
-- {print $1 }
- awk `{ print $1 }'
```

- prints the first column in each line; using -F will provide a field separator; the default FS is space
- 7. What is the output of the following command line:

```
echo good | sed '/Good/d'
```

- good

8. Which **awk** script outputs all the lines where a plus sign + appears at the end of line?

```
- awk '$0 ~ /+$/ { print $0 } OR awk '/+$/ { print $0 }
```

- 9. What is the command to delete only the first 5 lines in a file "foo"? Which command deletes only the last 5 lines?
- Command to delete only the first 5 lines in a file "foo":
 sed -i '1,5d' foo

- tail -n +5 foo > foo.temp ; mv foo.temp foo
- awk 'NR > 5 { print \$5 } ' foo > foo.temp; mv foo.temp foo

- Command to delete only the last 5 lines:

- tac foo | sed "1,5d" | tac > foo.temp; mv foo.temp foo
- head -n -5 foo > foo.temp; mv foo foo.temp

Part II b (10pts each): 50pts

Describe the function (5pts) and output (5pts) of the following commands.

9. \$ cat float

Wish I was floating in blue across the sky, my imagination is strong, And I often visit the days

When everything seemed so clear.

Now I wonder what I'm doing here at all...

\$ cat h1.awk

NR>2 && NR<4{print NR ":" \$0

\$ awk '/.*ing/ {print NR ":" \$1}' float

- awk \'.*ing/ {print NR ":" \$1}' float
- prints the record number, a colon, and the first field for the lines finding "ing" in any of the strings
- Output:

1:Wish

2:When

3:Now

- Explanation: Anything inside the double backslashes // is used to find the string(s) in the file. The expression . *ing will find any string with "ing" anywhere in the file. The script will find three lines and print them with their record numbers.

10. As the next command following question 9,

\$ awk -f h1.awk float

- awk -f hl.awk float
- prints the record number, a colon and the third line form the file *float*
- Output:

3:When everything seemed so clear.

- Explanation: The command NR>2 && NR<4{print NR ":" \$0} makes awk act on record numbers greater than 2 and less than 4 (the number is 3). The record number, a colon and the entire line(\$0) will print.

11.

```
$ cat h2.awk
BEGIN { print "Start to scan file" }
{print $1 ","$NF}
END {print "END-", FILENAME }
awk -f h2.awk float
```

- awk -f h2.awk float
- prints "Start to scan file" in the beginning, then for each record it would print the first column, followed by a comma, and another comma followed by the filename
- Output:

```
Start to scan file
```

Wish.is

Strong,,days

When, clear.

Now,all...

END- float

- Explanation: The *awk* script first executes the commands in the *BEGIN* section once per file, then executes the commands in the executable section once for each record, and then finally the commands in the *END* section.

12. sed ' $s/\$ ' float

- sed 's/\s/\t/g' float
- replaces space with tab for all occurrences of space.
- Output:

Wish	I was floati	ng	in	blue	across	the	sky,
my	imagination	is	strong	5,	And	I	often
visit	the days						
When	everything	seen	ned		so	clear.	
Now	I wonder		what	I'm	doing	here	at
all							

- Explanation: The *sed* will find space (\s) with tab (\t) and the flag *g* would replace them for all occurrences

13.

 $\$ ls *.awk| awk '{print "grep --color 'BEGIN' " $\$ }' |sh (*Notes: sh file runs file as a shell script . \$1 should be the output of 'ls *.awk' in this case, not the 1*st field)

- ls *awk | awk '{print "grep --color 'BEGIN' "\$1}' | sh
- Explanation: The command *ls *awk* lists all the .awk files in the current directory (h1.awk and h2.awk). The awk then builds a grep command to list all the lines with the BEGIN keyword. The sh then runs the generated commands.
- the generated commands:

```
grep --color BEGIN h1.awk
grep --color BEGIN h2.awk
```

- Output:

BEGIN { print "Start to scan file" }

- (the keyword BEGIN would be printed in red since we used *--color* option of grep)

14.

\$ mkdir test test/test1 test/test2 \$cat>test/testt.txt This is a test file ^D

- \$ cd test
- \$ ls -l. | grep '^d' | awk '{print "cp -r " \$NF " " \$NF ".bak"}' | sh
 - mkdir test test/test1 test/test2
 - This creates a directory named *test* with two more directories inside it named *test1* and *test2*
 - cat>test/testt.txt
 - This is a test file^D
 - This creates the *test.txt* file inside the test directory with the contents "This is a test file".
 - cd test (is entered into *test* directory)
 - ls -l . | grep '^d' | awk '{print "cp -r " \$NF " "\$NF".bak"}' | sh
 - Explanation: The *ls -l* lists the files in the *test* directory. The *grep* then finds the lines starting with *d* (the directories). *awk* then recursively copies the directories with the respective directories by adding *.bak* as the extensions to the directories.
 - Output:

ls -p

test1/ test1.bak. test2/ test2.bak/ test.txt

- (-p puts a slash after every directory listed)

Part III Programming: 15pts

- 15. Sort all the files in your class working directory (or your home directory) as per the following requirements:
- a. A copy of each file in that folder must be made. Append the string "_copy" to the name of the file
- b. The duplicate (copied) files must be in separate directories with each directory specifying the type of the file (e.g. txt files in directory named txtfiles, pdf files in directory named pdffiles etc).
- c. The files in each directory must be sorted in chronological order of months.
- d. An archive file (.tar) of each directory must be made. The .tar files must be sorted by name in ascending order.
- e. An archive file of all the .tar archive files must be made and be available in your home directory.

As an output, show your screen shots for each step or a single screenshot that will cover the outputs from all the steps.

```
[vdo10@gsuad.gsu.edu@snowball ~]$ pwd
/home/vdo10
[vdo10@gsuad.gsu.edu@snowball ~]$ ls
checkError.sh csc3320 hello.sh homeworks Lab3 Lab4 public Result simple.sh
[vdo10@gsuad.gsu.edu@snowball ~]$ [
```

I used the pwd and 1s commands to check all of the directories in my current working directory before I started programming this homework.

```
[vdol0@gsuad.gsu.edu@snowball ~]$ touch file1.txt file3.txt
[vdol0@gsuad.gsu.edu@snowball ~]$ touch file2.pdf file4.pdf
[vdol0@gsuad.gsu.edu@snowball ~]$ ls
checkError.sh file1.txt file3.txt hello.sh Lab3 public simple.sh
csc3320 file2.pdf file4.pdf homeworks Lab4 Result
[vdol0@gsuad.gsu.edu@snowball ~]$
```

I made up some random .txt and .pdf files using touch command and checked to see if they were actually there by using ls.

```
[vdol0@gsuad.gsu.edu@snowball ~]$ mkdir {txtfiles,pdffiles}
[vdol0@gsuad.gsu.edu@snowball ~]$ ls
checkError.sh filel.txt file3.txt hello.sh Lab3 pdffiles Result txtfiles
csc3320 file2.pdf file4.pdf homeworks Lab4 public simple.sh
[vdol0@gsuad.gsu.edu@snowball ~]$
```

I made two separate directories named *txtfiles* and *pdffiles* using the mkdir command and checked back at the home directory using 1s.

```
[vdo10@gsuad.gsu.edu@snowball ~]$ cp filel.txt txtfiles/filel_copy.txt
[vdo10@gsuad.gsu.edu@snowball ~]$ cp file3.txt txtfiles/file3_copy.txt
[vdo10@gsuad.gsu.edu@snowball ~]$ cp file2.pdf pdffiles/file2_copy.pdf
[vdo10@gsuad.gsu.edu@snowball ~]$ cp file4.pdf pdffiles/file4_copy.pdf
[vdo10@gsuad.gsu.edu@snowball ~]$ cd ~/txtfiles
[vdo10@gsuad.gsu.edu@snowball txtfiles]$ ls
file1_copy.txt file3_copy.txt
[vdo10@gsuad.gsu.edu@snowball txtfiles]$ cd ~/pdffiles
[vdo10@gsuad.gsu.edu@snowball pdffiles]$ ls
file2_copy.pdf file4_copy.pdf
[vdo10@gsuad.gsu.edu@snowball pdffiles]$ ls
file2_copy.pdf file4_copy.pdf
[vdo10@gsuad.gsu.edu@snowball pdffiles]$
```

I copied the files to their respective directories using the cp command (i.e *file1.txt* would be copied into *txtfiles* as *file1_copy.txt*), went into the directories using the cd ~/txtfiles and cd ~/pdffiles commands, and checked for the duplicate copies using 1s.

```
[vdol0@gsuad.gsu.edu@snowball ~]$ cd ~/txtfiles
[vdol0@gsuad.gsu.edu@snowball txtfiles]$ sort -M filel_copy.txt
[vdol0@gsuad.gsu.edu@snowball txtfiles]$ cd
[vdol0@gsuad.gsu.edu@snowball ~]$ tar -czvpf txtfiles.tar.gz txtfiles
txtfiles/
txtfiles/file3_copy.txt
txtfiles/file1_copy.txt
[vdol0@gsuad.gsu.edu@snowball ~]$ ls -lrth | grep -i tar
-rw-rw-r--. l vdol0@gsuad.gsu.edu vdol0@gsuad.gsu.edu 187 Sep 30 22:43 txtfiles.tar.gz
[vdol0@gsuad.gsu.edu@snowball ~]$ [
```

I sorted the *txtfiles* in chronological order of months by using the sort -M command and then archived this file of directory using the tar command.

```
[vdo10@gsuad.gsu.edu@snowball ~]$ cd ~/pdffiles
[vdo10@gsuad.gsu.edu@snowball pdffiles]$ sort -M file2_copy.pdf
[vdo10@gsuad.gsu.edu@snowball pdffiles]$ cd
[vdo10@gsuad.gsu.edu@snowball ~]$ tar -czvpf pdffiles.tar.gz pdffiles
pdffiles/
pdffiles/file4_copy.pdf
pdffiles/file4_copy.pdf
[vdo10@gsuad.gsu.edu@snowball ~]$ ls -lrth | grep -i tar
-rw-rw-r--. l vdo10@gsuad.gsu.edu vdo10@gsuad.gsu.edu 187 Sep 30 22:43 txtfiles.tar.gz
-rw-rw-r--. l vdo10@gsuad.gsu.edu vdo10@gsuad.gsu.edu 187 Sep 30 22:48 pdffiles.tar.gz
[vdo10@gsuad.gsu.edu@snowball ~]$
```

I did the exact same thing as the previous screenshot step above, but with the *pdffiles* instead.

```
[vdo10@gsuad.gsu.edu@snowball ~]$ ls
checkError.sh filel.txt file3.txt hello.sh Lab3 pdffiles public simple.sh txtfiles.tar.gz
csc3320 file2.pdf file4.pdf homeworks Lab4 pdffiles.tar.gz Result txtfiles
[vdo10@gsuad.gsu.edu@snowball ~]$
```

I used 1s to check for the *tar* files in the home directory.

Here is the history of my commands.