CS216: Introduction to Software Engineering Techniques (Summer, 2020) Lab Assignment 2 (20 points)

Today's Date: Friday, June 12

Due Date: Tuesday, June 16

The purpose of this lab assignment is:

- Getting familiar with Unix/Linux environment
- Practicing more Unix commands such as: sort and grep
- Learning the basic Regular Expression

Requirement:

1. Connect to your VM and change your working directory to CS216.

In the terminal window (either from "terminal" application in the GUI-based Linux or from connecting your Virtual Machine through SSH client), make the CS216 directory (which you created in Lab1) your current working directory.

2. Create the directory

Create a directory underneath the CS216 directory named Lab2 and make the Lab2 directory your current working directory.

3. Download the file.

Without opening a browser, use command curl to download a text file named imdb.txt from the link (http://www.cs.uky.edu/~yipike/CS216/imdb.txt) and save the file into your current working directory ~/CS216/Lab2 by typing the following command at the prompt \$:

\$curl -0 http://www.cs.uky.edu/~yipike/CS216/imdb.txt

Please note that the option at the above command is upper case letter O instead of number zero. You will use this text file to search for some movies so that you can generate the output for this lab assignment. You can type the following command at the prompt \$ to check the content of the file named imdb.txt:

\$ cat imdb.txt

It displays one movie title (with the year made) at each line, in the decreasing order of the number of votes (note that the number of votes for each movie is not included in the file though).

4. Read the manual.

You need to use commands **sort** and **grep** to finish this Lab assignment. Spend some time to read the manual pages of commands **sort** and **grep** respectively by typing the following commands at the prompt \$:

```
$ man sort
$ man grep
```

You can also read the reference part at the end of this document for **sort** command and lecture notes for **grep** command.

5. Solve problems.

The problem you need to solve for this lab assignment is:

- (1) to sort all the movies from imdb.txt you downloaded from step 3, in Lexicographical order (dictionary order) by title.
- (2) to search from the sorted file in step(1) and find all the movie titles, which start with "Star Wars" and which were made in between year 2002 and year 2009 (including 2002 and 2009), and print them to an output file, named Lab2output.txt.
- (3) to search from the sorted file in step(1) and find all the movies which contain "land" as part of the movie title (Ignore case distinctions) and which were made in either year 1979 or year 1999, and print (append) them at the end of the same output file, named Lab2output.txt.

The following is the sample output of Lab2output.txt:

```
Star Wars: A Musical Journey (2005) (V)
Star Wars: Episode II - Attack of the Clones (2002)
Star Wars: Episode III - Revenge of the Sith (2005)
Star Wars Episode V 1/2: The Han Solo Affair (2002)
Star Wars: Revelations (2005) (V)
Star Wars: The Clone Wars (2008)
Abendland (1999)
A Dog of Flanders (1999)
Backstreet Boys Homecoming: Live in Orlando (1999) (V)
Bear Island (1979)
Clubland (1999)
Drowning on Dry Land (1999)
Heartland (1979)
Hinterland (1999)
In a Savage Land (1999)
Jackson Pollock: Love and Death on Long Island (1999) (V)
```

```
Landspeed: CKY (1999) (V)
Sayehaye bolande bad (1979)
Schlaraffenland (1999)
The Adventures of Elmo in Grouchland (1999)
The Clandestine Marriage (1999)
The Lost Treasure of Sawtooth Island (1999)
The Wacky Adventures of Ronald McDonald: The Legend of Grimace Island (1999) (V)
Treasure Island (1999/I)
Treasure Island (1999/II)
Wonderland (1999)
```

Note the first **SIX** lines of the sample output is the result from matching the pattern (2): movies starting with "**Star Wars**" and made in between **year 2002 and year 2009** (including 2002 and 2009); and the rest part is the result from matching the pattern (3): movies made in **year 1979 or year 1999** and each containing "**land**" (ignore case distinctions) in the title.

6.type the following command at the prompt \$, then click "Enter" key.

```
$ history | tail -10 >> Lab2output.txt
```

The above command sends the output from the command "history" to the command tail as input through the piping operator(|), then append the output to the existing file named Lab2output.txt. It actually appends the latest ten commands you typed from the command line to the end of the file Lab2output.txt, make sure that it contains the commands you used to generate the output file at step 5. Check the content of your output file Lab2output.txt, if it does not contain the commands you used at step 5, then adjust the number 10 in the above command, so that the commands you used at step 5 are inside.

Extra (No credit)

If you still have time, please think about based on the matched pattern of step 5(3), how can you separate them into two groups: (i) all the movie titles containing "land" that form whole words; (ii) all the movie titles containing "land" as a sub-string but not the whole word.

```
Your matched pattern for this Extra part (i) group should be:

Drowning on Dry Land (1999)

In a Savage Land (1999)

Your matched pattern for this Extra part (ii) group should be:

Abendland (1999)

A Dog of Flanders (1999)

Backstreet Boys Homecoming: Live in Orlando (1999) (V)

Bear Island (1979)

Clubland (1999)

Heartland (1979)
```

Hinterland (1999)
Jackson Pollock: Love and Death on Long Island (1999) (V)
Landspeed: CKY (1999) (V)
Sayehaye bolande bad (1979)
Schlaraffenland (1999)
The Adventures of Elmo in Grouchland (1999)
The Clandestine Marriage (1999)
The Lost Treasure of Sawtooth Island (1999)
The Wacky Adventures of Ronald McDonald: The Legend of Grimace Island (1999) (V)
Treasure Island (1999/I)
Treasure Island (1999/II)
Wonderland (1999)

Submission

Open the link to Canvas LMS (https://uk.instructure.com/), and log in to your account using your linkblue user id and password. Please submit your file (Lab2output.txt) through the submission link for "Lab2". You can find your file Lab2output.txt from "Place", then the directory where you saved your file (if you are under the GUI-based Linux) or transferred from your VM to your local computer via Filezilla, or BitVise, or NoMachine, or scp command©

Grading (20 points + Bonus 3 points)

- 1. Attend the lab session or have a documented excused absence. (5 points)
- 2. Finish step 5 and generate the file named Lab2output.txt correctly. (8 points)

(if you just copy the output, you get 0 point)

- generate the matched text lines for pattern 5(2). (4 points)
- generate the matched text lines for pattern 5(3). (4 points)
- 3. At the end of the file Lab2output.txt, it contains the command you used to generate the sub-directory named Lab2 under CS216 directory. (1 point)
- 3. At the end of the file Lab2output.txt, it contains the commands you used to generate the list of matched text lines for patterns 5(2) and 5(3). (3*2=6 points)

Bonus: Demonstrate your program to your TA and answer TA's questions. (3 points)

(Late assignment will be reduced 10% for each day that is late. The assignment will not be graded (you will receive zero) if it is more than 3 days late. Note that a weekend counts just as regular days. For example, if an assignment is due Friday and is turned in Monday, it is 3 days late.

Reference

Unix sort command

NAME

sort - sort lines of text files

SYNOPSIS

sort [OPTION]... [FILE]...

OPTIONS

- -d, --dictionary-order
 consider only blanks and alphanumeric characters
- -f, --ignore-case
 fold lower case to upper case characters
- -g, --general-numeric-sort
 compare according to general numerical value
 - -r, --reverse reverse the result of comparisons
- - -r, --reverse
 reverse the result of comparisons
 - -o, --output=FILE
 write result to FILE instead of standard output

EXAMPLE

If you have an input file data.txt with the following data:

apple 55 orange 22 grape 11 pineapple 20 peach 39

...and you sort lines by dictionary order:

\$ sort -d data.txt

you will receive the following output:

apple	55
grape	11
orange	22
peach	39
pineapple	20

However, if you want to sort based on the second field in reverse numeric order, you can use the following command:

\$ sort -rg -k2 data.txt

It will produce the following g output:

apple	55
peach	39
orange	22
pineapple	20
grape	11

Redirection and Piping

Redirection is a mechanism for sending data to and from files.

- > redirects std-out to a file
- >> appends std-out to a file
- < redirects std-in from a file

Piping (|) is a mechanism for sending data from one command (or program) to another. The **piping operator**| feeds the output from the command (or program) on the left as input to the command (or program) on the right.

Example:

```
$ history | tail -10 >> Lab2output.txt
```