

NLP 100 Exercise - Chapter 2: Unix Commands



10. Line Count

Count the number of lines of the file. Confirm the result by using `wc` command.

```
$ wc popular-names.txt
2780 11120 55026 popular-names.txt
```

2780 lines
11120 words
55026 characters

More info about `wc` command here: <https://linuxize.com/post/linux-wc-command/>

11. Replace tabs into spaces

Replace every occurrence of a tab character into a space. Confirm the result by using `sed`, `tr`, or `expand` command.

Information for `sed` command: <https://www.geeksforgeeks.org/sed-command-in-linux-unix-with-examples/>

```
$ sed 's/\t/ /g' popular-names.txt
```

Information for `tr` command: <https://www.geeksforgeeks.org/tr-command-in-unix-linux-with-examples/>

```
$ cat popular-names.txt | tr '\t' ' '
```

12. col1.txt from the first column, col2.txt from the second column

Extract the value of the first column of each line, and store the output into `col1.txt` .
Extract the value of the second column of each line, and store the output into `col2.txt` .
Confirm the result by using `cut` command.

```
$ cut popular-names.txt -f 1 > col1.txt
$ cut popular-names.txt -f 2 > col2.txt
```

More info: <https://linuxize.com/post/linux-cut-command/>

13. Merging col1.txt and col2.txt

Join the contents of `col1.txt` and `col2.txt` , and create a text file whose each line contains the values of the first and second columns (separated by tab character) of the original file. Confirm the result by using `paste` command.

```
$ paste col1.txt col2.txt
```

More info: <https://www.geeksforgeeks.org/paste-command-in-linux-with-examples/>

14. First N lines

Receive a natural number N from a command-line argument, and output the first N lines of the file. Confirm the result by using `head` command.

```
$ head -n 15 popular-names.txt
Mary    F    7065    1880
Anna    F    2604    1880
Emma    F    2003    1880
Elizabeth F    1939    1880
Minnie  F    1746    1880
Margaret F    1578    1880
Ida F    1472    1880
Alice   F    1414    1880
Bertha  F    1320    1880
Sarah   F    1288    1880
John    M    9655    1880
William M    9532    1880
```

James	M	5927	1880
Charles	M	5348	1880
George	M	5126	1880

15. Last N lines

Receive a natural number N from a command-line argument, and output the last N lines of the file. Confirm the result by using `tail` command.

```
$ tail -n 15 popular-names.txt
Charlotte F 12940 2018
Mia F 12642 2018
Amelia F 12301 2018
Harper F 10582 2018
Evelyn F 10376 2018
Liam M 19837 2018
Noah M 18267 2018
William M 14516 2018
James M 13525 2018
Oliver M 13389 2018
Benjamin M 13381 2018
Elijah M 12886 2018
Lucas M 12585 2018
Mason M 12435 2018
Logan M 12352 2018
```

More info: <https://www.baeldung.com/linux/head-tail-commands>

16. Split a file into N pieces

Receive a natural number N from a command-line argument, and split the input file into N pieces at line boundaries. Confirm the result by using `split` command.

$N = 3$

```
$ split -n 1/3 popular-names.txt
```

(this command splits the file into 3 chunks xaa, xab, xac by line)

More info: <https://www.geeksforgeeks.org/split-command-in-linux-with-examples/>
(also `man split` is quite helpful for this one)

17. Distinct strings in the first column

Find distinct strings (a set of strings) of the first column of the file. Confirm the result by using `cut`, `sort`, and `uniq` commands.

[Sort command](#) sorts the lines by alphabetical order and also lowercase/uppercase.

```
$ sort col1.txt > col1_sorted.txt
```

[Uniq command](#) removes duplicate lines only if they are adjacent.

```
$ uniq col1_sorted.txt > col1_unique.txt
```

18. Sort lines in descending order of the third column

Sort the lines in descending numeric order of the third column (sort lines without changing the content of each line). Confirm the result by using `sort` command.

```
$ sort -nr -k 3 popular-names.txt
```

`-nr` : sort by numerical reverse order

`-k` : sort by column. `-k 3` means to sort by the 3rd column

19. Frequency of a string in the first column in descending order

Find the frequency of a string in the first column, and sort the strings by descending order of their frequencies. Confirm the result by using `cut`, `uniq`, and `sort` commands.

```
$ uniq -c col1_sorted.txt > col1_number_of_frequency.txt
```

```
$ sort -nr col1_number_of_frequency.txt > col1_number_of_frequency_sorted.txt
```

```
$ cat coll_number_of_frequency_sorted.txt
```

```
118 James
111 William
108 Robert
108 John
 92 Mary
 75 Charles
 74 Michael
 73 Elizabeth
 70 Joseph
 60 Margaret
 58 Thomas
 58 George
 57 David
 51 Richard
 45 Helen
 43 Frank
 43 Christopher
 41 Anna
 40 Edward
 39 Ruth
 38 Patricia
 37 Matthew
 36 Dorothy
 35 Emma
 32 Barbara
 31 Joshua
 31 Daniel
 26 Sarah
 26 Linda
 26 Jennifer
 26 Emily
 25 Jessica
 25 Jacob
 24 Susan
 24 Mildred
 24 Betty
 23 Henry
 23 Ashley
 22 Nancy
 21 Andrew
 20 Marie
 20 Florence
 20 Donald
```

20 Amanda
19 Samantha
18 Olivia
18 Melissa
18 Madison
18 Lisa
18 Karen
17 Stephanie
17 Abigail
16 Sandra
16 Mark
16 Ethel
15 Michelle
15 Isabella
15 Heather
15 Frances
15 Ethan
15 Carol
15 Angela
14 Shirley
14 Kimberly
14 Ava
14 Amy
13 Virginia
13 Sophia
13 Nicole
13 Jason
13 Hannah
13 Deborah
13 Brian
12 Minnie
12 Donna
12 Bertha
11 Cynthia
10 Ronald
10 Noah
10 Nicholas
10 Mia
10 Doris
10 Brittany
10 Alice
9 Tyler
9 Joan
9 Debra

8 Taylor
8 Mason
8 Judith
8 Ida
8 Clara
8 Alexis
8 Alexander
7 Tammy
7 Steven
7 Sharon
7 Liam
7 Harry
7 Brandon
6 Anthony
5 Jeffrey
5 Jayden
5 Gary
5 Charlotte
5 Annie
4 Lillian
4 Kathleen
4 Justin
4 Chloe
4 Benjamin
4 Austin
3 Megan
3 Harper
3 Evelyn
3 Elijah
3 Aiden
2 Rebecca
2 Oliver
2 Logan
2 Lauren
2 Larry
2 Bessie
2 Amelia
1 Walter
1 Tracy
1 Scott
1 Rachel
1 Pamela
1 Lucas
1 Lori

1 Laura
1 Kelly
1 Julie
1 Crystal
1 Carolyn

