# FIT1043: Assignment 3 Report Processing large data set using BASH shell scripts and R

Name: Eu Jia Xin

Student ID: 30881676

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# Introduction

This aim of this assignment is to utilise BASH shell scripts and R programming language to read and process large data set. For this assignment, the data used is a pre-processed twitter dataset for 15<sup>th</sup> May 2020, filtered by COVID-19 related keywords. This dataset contains variety of information including user information, tweet content, follower counts and related information. The data will be read and processed using BASH shell scripts, and visualisations will be generated using R Studio. Instructions given from the assignment specifications will be included for reference. It should be noted that a \$ sign will be used to indicate the start of the shell commands; and the code in blue represents R codes inputted (same format as in the console). The relevant codes and outputs will be shown, followed by analysis or additional information where necessary in the following sections of the report.

# Part A: Inspecting the Data

1) Copy the downloaded file to your UNIX (Linux) terminal. State the size (in Bytes or MegaBytes) of the corona\_tweets.csv.gz file and provide the shell command that you used to determine the size.

shell command:

```
$ 1s -lh corona_tweets.csv.gz
output:
-rw-rw-r-- 1 jeuu0002 jeuu0002 118M May 31 13:33 corona_tweets.csv.gz
```

Analysis:

Based on the output, 118M represents 118 MegaBytes. Therefore, the file size is 118 MegaBytes.

2) The first line of the CSV file contains headers that are "tab" separated. What are the header names and provide the command you used to obtain it. Note that the command provided has to be in one line.

```
shell command:
```

```
$ cat corona_tweets.csv.gz | gunzip | head -1 | tr "\t" "\n"
output:
Created
Tweet_ID
Text
User_ID
User
User_Location
Followers_Count
Friends_Count
Geo
Place_Type
```

```
Place_Name
Place_Country
Language
```

## Analysis:

To allow a clearer view of the column headers, every column header is outputted in a new line and can be seen in the output. Based on the output, there are 13 headers in total, which are: Created, Tweet\_ID, Text, User\_ID, User, User\_Location, Followers\_Count, Friends\_Count, Geo, Place\_Type, Place\_Name, Place\_Country and Language.

3) How many lines are there in the dataset? Again, provide the single line code on how you obtained it.

shell command:

Analysis:

```
$ cat corona_tweets.csv.gz | gunzip | wc -1
output:
1143559
```

There are 1143559 lines in the dataset.

# Part B: Information from data

1) How many unique twitter users are there in the dataset. Provide the single line code that uses the "awk" and "uniq" command. You are also required to read the "man" pages of the "uniq" command to figure out if it is sufficient to answer the question. Provide explanation on the code you provided.

```
shell command:
```

```
$ cat corona_tweets.csv.gz | gunzip | awk -F '\t' '{print $4}' | head -10
output:
```

User ID

3269308680

255800212

803042908345696300

350289248

812218284640608300

3808461979

1161532339052261400

938616431259934700

1243052659944198100

First 10 rows are outputted to make sure we are using an appropriate column. Becuase User\_ID serves as an identifier for different users, this column will be used to find the number of unique twitter users.

shell command:

```
$ cat corona_tweets.csv.gz | gunzip | awk -F '\t' '{print $4}' | grep -v
'User_ID' | sort | uniq | wc -l
```

output:

641975

## Analysis:

The awk command is used to extract only the User\_ID column. Then, grep command is used to exclude the column header, as we only want to know the number of unique twitter users. In the manual of the uniq command, it says that the command "filters adjacent matching lines", which means that we are required to sort if first to retrieve the actual number of unique twitter users. The total number of unique twitter users is 641975.

- 2) For each of the sub-question below, provide the single line code (one each) and briefly explain your code.
- a) How many tweets mentioned the word "death" in any combination of uppercase or lowercase letters.

shell command:

```
$ cat corona_tweets.csv.gz | gunzip | cut -f 3 | grep -i "death" | wc -l
output:
```

50911

Analysis:

The cut command is used to extract the 3<sup>rd</sup> column of the dataset (Text). Then, grep command is to find tweets with the word 'death' in it. The -i flag is used to ignore case as we want the word 'death' in any combination of uppercase or lowercase letters. Then, wc -l is used to show how many lines there are, which is how many tweets there are. The output shows that there are 50911 tweets containing the word 'death'.

b) How many of those are not spelt exactly "death", "deaths", "Death" or "Deaths" but in other combination of uppercase and lowercase (e.g. DEath, deatHs)

shell command:

```
$ cat corona_tweets.csv.gz | gunzip | cut -f 3 | grep -i 'death' | grep -v
-E 'death|deaths|Death|Deaths' | wc -l
```

output:

450

Analysis:

To find the answer, we will modify the code from 2a. There will an additional grep command. For the second new grep command, the -v flag is used to filter out the unwanted words. The -E flag is used to make sure that the program searches and excludes 4 different words, instead of searching for a single long word. Based on the output, there are **450 tweets** that contain the word 'death' (case insensitive), but are not spelt exactly as "death", "deaths", "Death" or "Deaths".

c) Output the lines of sub-part 2 (b) into a file called myText.txt (not the number of lines but the specific lines).

shell command:

```
$ cat corona_tweets.csv.gz | gunzip | cut -f 3 | grep -i 'death' | grep -v
-E 'death|deaths|Death|Deaths' > myText.txt
```

To check if the .txt file is correct, it is opened in notepad. When scrolled to the bottom, the bottom right of the terminal (highlighted in a blue rectangle) shows that there are 450 lines in total, which matches the previous code in 2b.

# Part C: Data aggregation

1) Let's group the twitter user (ID) by the number of followers that they have into the following ranges. Provide the code for them. One line of code for each of them below (11 lines).

```
a) <= 1000
```

shell command:

```
$ cat corona_tweets.csv.gz | gunzip | awk -F'\t' '$7<=1000 {print $4,$7}'
| sort -u | less</pre>
```

The less command is used to show the user ID and follower count. A manual inspection was done to find possible repeating user IDs. A specific user ID 100003276 is shown to be outputted twice. And the follower count is 280 and 281. For a clearer output, the grep command is used to show only the specific user ID using the first attempted code.

#### shell command:

```
$ cat corona_tweets.csv.gz | gunzip | awk -F'\t' '$7<=1000 {print $4,$7}'
| sort -u | grep '100003276'
output:</pre>
```

100003276 280

100003276 281

This shows that the user\_id has an increase in follower by 1, which proved the problem cannot be solved by directly sorting with a -u flag as above.

From this, we can deduce that there are duplicated user IDs even though we have attempted to sort it with the -u flag. This is most likely because the same user might have tweeted multiple times a day, creating multiple records. And within the time intervals, they might have gained or lost followers, causing the follower count to be different. Therefore, we will obtain the most accurate follower count by taking the latest entry of the record.

We will try to rectify this using the following codes and checking if the specific user ID chosen earlier will only output follower count of 281.

# shell command:

```
$ cat corona_tweets.csv.gz | gunzip | awk -F'\t' '{print $4,$7}' | sort -r
-u | awk '!seen[$1]++' | awk -F' ' '$2<=1000 {print $1,$2}' | grep
'100003276'</pre>
```

output:

100003276 281

### Code breakdown:

The first awk command will select the User\_ID and Followers\_Count only. Then, we will sort using two flags, -r and -u to make sure that the latest record of unique users with duplicate records will be shown at the top. Then, awk '!seen[\$1]++ is used to only take the first record of any user IDs that are repeated. The final awk command will filter out user IDs based on the follower count range intended.

Since this code takes the latest record for the specific user ID chosen, we assume this is correct for now, and will use the same approach to find the rest of the followers range. Instead of outputting the user IDs and follower count, we will directly count the number of records using wc -1. Afterwards, the sum of all 11 lines results is totalled up and compared to answer from b2 (641975 unique users) to double confirm if the code is correct.

```
a) <= 1000
shell command:
\ cat corona_tweets.csv.gz | gunzip | awk -F'\t' '{print $4,$7}' | sort -r -u | awk '!seen[$1]++' | awk -F' ' '$2<=1000 {print $1,$2}' | wc -1
output:
455751
b) 1001 to 2000
shell command:
-u | awk '!seen[$1]++' | awk -F' ' '$2>1000 && $2<=2000 {print $1,$2}' |
wc -1
output:
68527
c) 2001 to 3000
shell command:
$ cat corona_tweets.csv.gz | gunzip | awk -F'\t' '{print $4,$7}' | sort -r
-u | awk '!seen[$1]++' | awk -F' ' '$2>2000 && $2<=3000 {print $1,$2}' |
wc -1
output:
31264
d) 3001 to 4000
shell command:
\ cat corona_tweets.csv.gz | gunzip | awk -F'\t' '{print $4,$7}' | sort -r -u | awk '!seen[$1]++' | awk -F' ' '$2>3000 && $2<=4000 {print $1,$2}' |
wc -1
output:
18798
e) 4001 to 5000
shell command:
$ cat corona_tweets.csv.gz | gunzip | awk -F'\t' '{print $4,$7}' | sort -r
-u | awk '!seen[$1]++' | awk -F' ' '$2>4000 && $2<=5000 {print $1,$2}' |
wc -1
output:
```

```
f) 5001 to 6000
shell command:
$ cat corona_tweets.csv.gz | gunzip | awk -F'\t' '{print $4,$7}' | sort -r
-u | awk '!seen[$1]++' | awk -F' ' '$2>5000 && $2<=6000 {print $1,$2}' |
wc -1
output:
7975
g) 6001 to 7000
shell command:
$ cat corona_tweets.csv.gz | gunzip | awk -F'\t' '{print $4,$7}' | sort -r
-u | awk '!seen[$1]++' | awk -F' ' '$2>6000 && $2<=7000 {print $1,$2}' |
wc -1
output:
5874
h) 7001 to 8000
shell command:
-u | awk '!seen[$1]++' | awk -F' ' '$2>7000 && $2<=8000 {print $1,$2}' |
wc -1
output:
4367
i) 8001 to 9000
shell command:
$ cat corona_tweets.csv.gz | gunzip | awk -F'\t' '{print $4,$7}' | sort -r
-u | awk '!seen[$1]++' | awk -F' ' '$2>8000 && $2<=9000 {print $1,$2}' |
wc -1
output:
3522
j) 9001 to 10000
shell command:
$ cat corona_tweets.csv.gz | gunzip | awk -F'\t' '{print $4,$7}' | sort -r
-u | awk '!seen[$1]++' | awk -F' ' '$2>9000 && $2<=10000 {print $1,$2}' |
wc -1
```

output:

2738

#### k) More than 10000

shell command:

```
$ cat corona_tweets.csv.gz | gunzip | awk -F'\t' '{print $4,$7}' | sort -r
-u | awk '!seen[$1]++' | awk -F' ' '$2>10000 {print $1,$2}' | grep -v
'User_ID' | wc -1
```

output:

31470

#### Analysis:

By summing up all the outputs from a to k, we know that there is a total of 641975 users, which proves that the code is correct.

2) Use (modified) the code above and include them into a shell script (with the extension .sh) to output a CSV file. The CSV file should contain two columns, the first column is the range and the second column is the number of twitter users.

```
$ nano c2shell.sh
```

Inside the nano editor:

#!/bin/bash

```
echo "<=1k,`cat corona_tweets.csv.gz | gunzip | awk -F'\t' '{print $4,$7}' | sort
-r -u | awk '!seen[$1]++' | awk -F' ' '$2<=1000 {print $1,$2}' | wc -l`" >
follower count.csv
echo "1-2k, `cat corona_tweets.csv.gz | gunzip | awk -F'\t' '{print $4,$7}' | sort
-r -u | awk '!seen[$1]++' | awk -F' ' '$2>1000 && $2<=2000 {print $1,$2}' | wc -
1`" >> follower count.csv
echo "2-3k,`cat corona_tweets.csv.gz | gunzip | awk -F'\t' '{print $4,$7}' | sort -r -u | awk '!seen[$1]++' | awk -F' ' '$2>2000 && $2<=3000 {print $1,$2}' | wc -
1`" >> follower_count.csv
echo "3-4k, `cat corona_tweets.csv.gz | gunzip | awk -F'\t' '{print $4,$7}' | sort -r -u | awk '!seen[$1]++' | awk -F' ' '$2>3000 && $2<=4000 {print $1,$2}' | wc -
1`" >> follower count.csv
echo "4-5k,`cat corona_tweets.csv.gz | gunzip | awk -F'\t' '{print $4,$7}' | sort -r -u | awk '!seen[$1]++' | awk -F' ' '$2>3000 && $2<=4000 {print $1,$2}' | wc -
1`" >> follower count.csv
echo "5-6k,`cat corona_tweets.csv.gz | gunzip | awk -F'\t' '{print $4,$7}' | sort -r -u | awk '!seen[$1]++' | awk -F' ' '$2>3000 && $2<=4000 {print $1,$2}' | wc -
1`" >> follower_count.csv
echo "6-7k,`cat corona_tweets.csv.gz | gunzip | awk -F'\t' '{print $4,$7}' | sort -r -u | awk '!seen[$1]++' | awk -F' ' '$2>3000 && $2<=4000 {print $1,$2}' | wc -
1`" >> follower count.csv
```

```
echo "7-8k,`cat corona_tweets.csv.gz | gunzip | awk -F'\t' '{print $4,$7}' | sort -r -u | awk '!seen[$1]++' | awk -F' ' '$2>3000 && $2<=4000 {print $1,$2}' | wc -1`" >> follower_count.csv

echo "8-9k,`cat corona_tweets.csv.gz | gunzip | awk -F'\t' '{print $4,$7}' | sort -r -u | awk '!seen[$1]++' | awk -F' ' '$2>3000 && $2<=4000 {print $1,$2}' | wc -1`" >> follower_count.csv

echo "9-10k,`cat corona_tweets.csv.gz | gunzip | awk -F'\t' '{print $4,$7}' | sort -r -u | awk '!seen[$1]++' | awk -F' ' '$2>9000 && $2<=10000 {print $1,$2}' | wc -1`" >> follower_count.csv

echo ">10k,`cat corona_tweets.csv.gz | gunzip | awk -F'\t' '{print $4,$7}' | sort -r -u | awk '!seen[$1]++' | awk -F' ' '$2>10000 {print $1,$2}' | grep -v 'User_ID' | wc -1`" >> follower_count.csv
```

The shell script is made using the nano editor. The line #!/bin/bash at the start is used to tell the program that the script should be interpreted and run by BASH shell. Then, there are 11 lines, each producing a record into the csv file. For every line, the echo command will be used to pipe the output into the csv file. The first column will be the follower count range, whereas the second column will be the number of users that are within the range. The second column is the code used to find the number of lines as in part C1. The code is enclosed with `` to make sure that the output of the code is piped into the csv. Aside from the first record, the remaining 10 lines uses >> instead of > to ensure that consecutive records are appended instead of overwriting previous lines.

#### shell command:

```
$ chmod a+x c2shell.sh
```

# \$ ./c2shell.sh

After executing the shell script, the csv file is created, we will check the output and open the file in Excel to ensure the shell script performed the tasks correctly.

## shell command:

```
$ cat follower_count.csv
output:
```

<=1k,455751

1-2k,68527

2-3k,31264

3-4k,18798

4-5k,11689

5-6k,7975

6-7k,5874

7-8k,4367

8-9k,3522

9-10k,2738

>10k,31470

Screenshot of the file opened in Excel:

4	Α	В	С
1	<=1k	455751	
2	1-2k	68527	
3	2-3k	31264	
4	3-4k	18798	
5	4-5k	11689	
6	5-6k	7975	
7	6-7k	5874	
8	7-8k	4367	
9	8-9k	3522	
10	9-10k	2738	
11	>10k	31470	
12			
13			

As the output and Excel file seems alright, we will proceed to visualise the follower count range in a bar chart. This will be done in R studio.

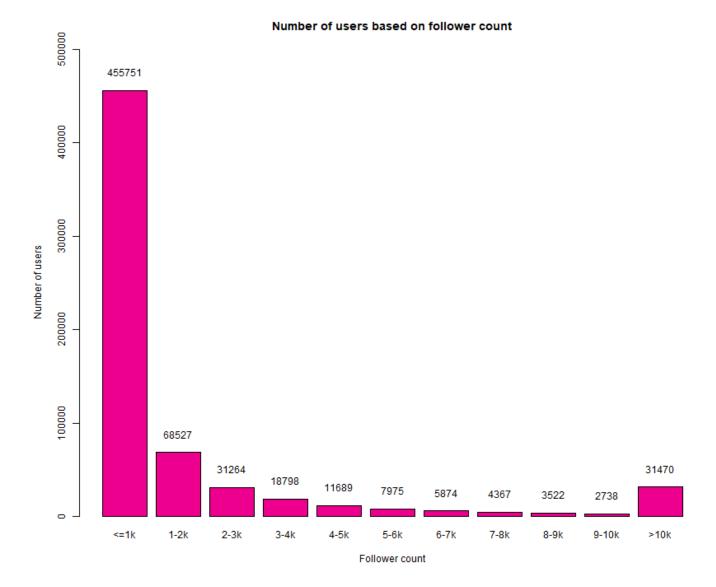
# **Visualisation**

Checking the file in R:

> options(scipen=100000)

To make sure the y-axis values are shown nicely without the scientific notations, scipen option is adjusted to a larger value.

```
> png("graph1.png",width= 850, height = 700)
> follower_graph = barplot(follower_count$V2,ylim = c(0,500000),main= "Number of users based on
follower count",names.arg = follower_count$V1,xlab="Follower count",ylab= "Number of users",col
= "#ED008E")
> text(x = follower_graph, y = follower_count$V2+20000, label=follower_count$V2)
> dev.off()
```



# Part D: Small challenge

Having done Part C: Data aggregation, let's assume that we want to compare against tweets that are not retweets. We can assume that this is indicated at the beginning of the text by the "RT @".

- 1) Provide a single line code that filters out the tweets that contain "RT @" and output the results into a compressed gz file.
- 2) Do the same process as Part C (1) (3) with this new file. Your R code can be a continuation from Part C.
- 3) Plot a side by side bar chart to visualise it. The bars should be coloured.

Similar to part c, a new user ID is randomly chosen from manual inspection beforehand to make sure that unique users are not double counted. The user ID selected is 100011309

```
shell command:
```

```
$ cat corona_tweets.csv.gz | gunzip | cut -f 3,4,7 | grep -v 'RT @' | awk
-F'\t' '{print $2,$3}' | grep '100011309'
output:
100011309 9417
100011309 9417
100011309 9417
100011309 9417
100011309 9419
```

To ensure that only the last line is outputted, the following code is used:

Shell command:

```
$ cat corona_tweets.csv.gz | gunzip | cut -f 3,4,7 | grep -v 'RT @' | awk
-F'\t' '{print $2,$3}' | sort -r -u | awk '!seen[$1]++' | grep '100011309'
Output:
```

```
100011309 9419
```

Outputting the results into a compressed gz file:

shell command:

```
\ cat corona_tweets.csv.gz | gunzip | cut -f 3,4,7 | grep -v 'RT @' | awk -F'\t' '{print $2,$3}' | sort -r -u | awk '!seen[$1]++' | gzip > no_rt.csv.gz
```

a) Less than or equal to 1000

shell command:

```
$ cat no_rt.csv.gz | gunzip | awk -F' ' '$2<=1000 {print $1,$2}' | wc -l
output:</pre>
```

142248

b) 1001 to 2000

shell command:

```
$ cat no_rt.csv.gz | gunzip | awk -F' ' '$2>1000 && $2<=2000 {print
$1,$2}' | wc -1
output:
```

24030

```
c) 2001 to 3000
shell command:
$ cat no_rt.csv.gz | gunzip | awk -F' ' '$2>2000 && $2<=3000 {print</pre>
$1,$2}' | wc -1
11770
d) 3001 to 4000
shell command:
$ cat no_rt.csv.gz | gunzip | awk -F' ' '$2>3000 && $2<=4000 {print</pre>
1,$2' | wc -1
output:
7348
e) 4001 to 5000
shell command:
$ cat no_rt.csv.gz | gunzip | awk -F' ' '$2>4000 && $2<=5000 {print</pre>
$1,$2}' | wc -1
output:
4724
f) 5001 to 6000
shell command:
$ cat no_rt.csv.gz | gunzip | awk -F' ' '$2>5000 && $2<=6000 {print</pre>
$1,$2}' | wc -1
output:
3459
g) 6001 to 7000
shell command:
$ cat no_rt.csv.gz | gunzip | awk -F' ' '$2>6000 && $2<=7000 {print</pre>
$1,$2}' | wc -1
output:
2498
h) 7001 to 8000
shell command:
```

```
$ cat no_rt.csv.gz | gunzip | awk -F' ' '$2>7000 && $2<=8000 {print</pre>
$1,$2}' | wc -1
output:
1908
i) 8001 to 9000
shell command:
$ cat no_rt.csv.gz | gunzip | awk -F' ' '$2>8000 && $2<=9000 {print</pre>
$1,$2}' | wc -1
output:
1611
j) 9001 to 10000
shell command:
$ cat no_rt.csv.gz | gunzip | awk -F' ' '$2>9000 && $2<=10000 {print</pre>
$1,$2}' | wc -1
output:
1279
k) More than 10000
shell command:
$ cat no_rt.csv.gz | gunzip | awk -F' ' '$2>10000 {print $1,$2}' | grep -
v 'User_ID' | wc -l
output:
17015
Now, by using the same approach as in part c, a new shell script to create a new csv file
containing the follower count range and number of users for non-retweets.
shell command:
$ nano d2shell.sh
Inside nano editor:
#!/bin/bash
echo "<=1k,`cat no_rt.csv.gz | gunzip | awk -F' ' '$2<=1000 {print $1,$2}' | wc -
1`" > no rt fcount.csv
echo "1-2k,`cat no_rt.csv.gz | gunzip | awk -F' ' '$2>1000 && $2<=2000 {print
$1,$2}' | wc -1`" >> no_rt_fcount.csv
echo "2-3k,`cat no_rt.csv.gz | gunzip | awk -F' ' '$2>2000 && $2<=3000 {print
$1,$2}' | wc -l`" >> no_rt_fcount.csv
```

```
echo "3-4k,`cat no_rt.csv.gz | gunzip | awk -F' ' '$2>3000 && $2<=4000 {print
$1,$2}' | wc -1`" >> no_rt_fcount.csv
echo "4-5k,`cat no_rt.csv.gz | gunzip | awk -F' ' '$2>4000 && $2<=5000 {print
$1,$2}' | wc -l`" >> no rt fcount.csv
echo "5-6k,`cat no_rt.csv.gz | gunzip | awk -F' ' '$2>5000 && $2<=6000 {print
$1,$2}' | wc -1`" >> no rt fcount.csv
echo "6-7k,`cat no_rt.csv.gz | gunzip | awk -F' ' '$2>6000 && $2<=7000 {print
$1,$2}' | wc -l`" >> no_rt_fcount.csv
echo "7-8k,`cat no_rt.csv.gz | gunzip | awk -F' ' '$2>7000 && $2<=8000 {print
$1,$2}' | wc -l`" >> no_rt_fcount.csv
echo "8-9k,`cat no_rt.csv.gz | gunzip | awk -F' ' '$2>8000 && $2<=9000 {print
$1,$2}' | wc -l`" >> no_rt_fcount.csv
echo "9-10k,`cat no_rt.csv.gz | gunzip | awk -F' ' '$2>9000 && $2<=10000 {print
$1,$2}' | wc -1`" >> no_rt_fcount.csv
echo ">10k,`cat no_rt.csv.gz | gunzip | awk -F' ' '$2>10000 {print $1,$2}' | grep
-v 'User_ID' | wc -1`" >> no_rt_fcount.csv
```

#### shell commands:

- \$ chmod a+x d2shell.sh
- \$ ./d2shell.sh

\$ cat no\_rt\_fcount.csv

## output:

- <=1k,142248
- 1-2k,24030
- 2-3k,11770
- 3-4k,7348
- 4-5k,4724
- 5-6k,3459
- 6-7k,2498
- 7-8k,1908
- 8-9k,1611
- 9-10k,1279
- >10k,17015

# Checking the Excel file:

010		
4	Α	В
1	<=1k	142248
2	1-2k	24030
3	2-3k	11770
4	3-4k	7348
5	4-5k	4724
6	5-6k	3459
7	6-7k	2498
8	7-8k	1908
9	8-9k	1611
10	9-10k	1279
11	>10k	17015
12		

The output and Excel file both looks alright, so we will proceed to visualising the results.

# Visualisation (side-by-side bar chart)

For this section, we will be using ggplot to plot the side by side bar chart.

First, we will import the new csv file made, and create a new dataframe based on the two csv files. The necessary libraries are then installed to help plot the bar chart. The following codes are used to plot the graph. Some outputs are shown to display the flow of code.

```
> no rt fcount = read.table("no rt fcount.csv",header=FALSE,sep=',')
> no_rt_fcount
  V1 V2
1 <=1k 142248
2 1-2k 24030
3 2-3k 11770
4 3-4k 7348
5 4-5k 4724
6 5-6k 3459
7 6-7k 2498
8 7-8k 1908
9 8-9k 1611
10 9-10k 1279
11 >10k 17015
> both_fcount = data.frame(follower_count,no_rt_fcount$V2)
> both_fcount
  V1 V2 no_rt_fcount.V2
1 <=1k 455751
                  142248
2 1-2k 68527
                  24030
3 2-3k 31264
                  11770
4 3-4k 18798
                  7348
5 4-5k 11689
                  4724
6 5-6k 7975
                 3459
7 6-7k 5874
                 2498
8 7-8k 4367
                 1908
9 8-9k 3522
                 1611
10 9-10k 2738
                  1279
```

```
> names(both fcount)[names(both fcount) == 'V1'] <- 'Follower Count'
> names(both_fcount)[names(both_fcount) == 'V2'] <- 'RT included'
> names(both_fcount)[names(both_fcount) == 'no_rt_fcount.V2'] <- 'RT excluded'
> both fcount
 Follower Count RT included RT excluded
              455751
1
       <=1k
                        142248
2
       1-2k
              68527
                        24030
3
       2-3k
              31264
                        11770
4
       3-4k
              18798
                        7348
5
       4-5k
                        4724
              11689
6
       5-6k
               7975
                        3459
7
       6-7k
               5874
                        2498
8
       7-8k
               4367
                        1908
9
       8-9k
               3522
                        1611
10
       9-10k
                2738
                         1279
11
       >10k
               31470
                         17015
> install.packages("tidyr")
> library(tidyr)
> library(ggplot2)
> df <- gather(both_fcount, Types of Tweet', 'Number of users', 'RT included':'RT excluded')
> df
 Follower Count Types of Tweet Number of users
1
       <=1k RT included
                             455751
2
       1-2k RT included
                             68527
3
       2-3k RT included
                             31264
4
       3-4k RT included
                             18798
5
       4-5k RT included
                             11689
6
       5-6k RT included
                              7975
7
       6-7k RT included
                              5874
8
       7-8k RT included
                              4367
9
       8-9k RT included
                              3522
10
       9-10k RT included
                               2738
11
       >10k RT included
                              31470
12
       <=1k RT excluded
                              142248
13
       1-2k RT excluded
                              24030
14
       2-3k RT excluded
                              11770
15
       3-4k RT excluded
                               7348
       4-5k RT excluded
16
                               4724
17
       5-6k RT excluded
                               3459
```

6-7k RT excluded

7-8k RT excluded

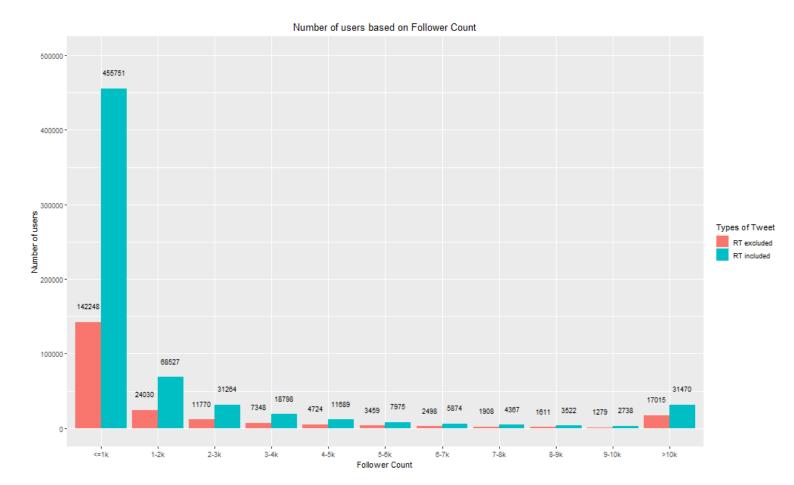
8-9k RT excluded

9-10k RT excluded

>10k RT excluded

<sup>&</sup>gt; df\$`Follower Count`<- as.character(df\$`Follower Count`)

```
> df$`Follower Count`<- factor(df$`Follower Count`,levels=unique(df$`Follower Count`))
> png("both_follower_graph.png",width= 1000, height = 600)
> plot <- ggplot(df, aes(x=`Follower Count`,y=`Number of users`,fill=`Types of Tweet`),width=10) +
coord_cartesian(ylim = c(0,500000))
> plot <- plot + geom_col(position = 'dodge') + ggtitle('Number of users based on Follower Count') +
theme(plot.title = element_text(hjust = 0.5))
> plot <- plot + geom_text(aes(label=`Number of users`, y= `Number of users`+
20000),position=position_dodge(width=0.9),vjust=0.25,size = 3)
> plot
> dev.off()
RStudioGD
2
```



# Conclusion

By using BASH shell scripts, large datasets can be processed in a short amount of time. The data can further be aggregated and manipulated efficiently, moreover we are able to generate filtered data for further processing. For example, a new compressed file which contains far less redundant data is created quickly, and the data is extracted into a csv file efficiently. This shows that the inter-process communication of piping in UNIX is extremely convenient for the manipulation of large data sets. Combined with R programming language, we are able to visualise the findings we obtained through visualisations like bar charts.