Task - 3

Aim: Implement EDA -Exploratory data analysis Module using MapReduce. come up with proper problem architecture with algorithms and document well.

Theory:

- ➤ EDA stands for Exploratory data analysis. It is an approach to analyze data sets to summarize their main characteristics, often using statistical graphics and other data visualization methods.
- ➤ MapReduce is the tool which distributes the data and process it in parallel manner. It mainly consists of two phases Mapper And Reducer. The mapper takes the data generates the data in key-value pair then output of mapper considered as input for reduce which reduces the same key values and generates the output.

Algorithm:

- > Start
- ➤ I extract the data by web scraping using BeautifulSoup library in python.
- > Then I clean the data such as
 - a) Covert each word into lowercase
 - b) Tokenization
 - c) Remove Stopping words
 - d) Stemming/Lemmatization
 - e) Remove Special Symbols
- ➤ Then I pass the data into map reduce using Hadoop streaming, here I count the frequency of each word.
- ➤ For Visualization of data I use Elk Stalk pipeline, I create a index in Kibana and pass the output file using logstash configuration file into the kibana index.
- ➤ End.

Program Code:

> Mapper.py

```
#!/usr/bin/env python
   """mapper.py"""
   import sys
   for line in sys.stdin:
   line = line.strip()
   words = line.split()
   # increase counters
   for word in words:
           print '%s\t%s' % (word, 1)
> Reducer.py
#!/usr/bin/env python
"""reducer.py"""
from operator import itemgetter
import sys
current\_word = None
current\_count = 0
word = None
for line in sys.stdin:
   line = line.strip()
word, count = line.split('\t', 1)
try:
count = int(count)
except ValueError:
continue
if current_word == word:
```

```
current_count += count
else:
if current_word:
print '%s\t%s' % (current_word, current_count)
current_count = count
current_word = word

if current_word == word:
print '%s\t%s' % (current_word, current_count)
```

Outputs:

```
[cloudera@quickstart Desktop]$ hdfs dfs -put Task3.txt
[cloudera@quickstart Desktop]$ hdfs dfs -ls
Found 24 items
-rw-r--r-- 1 cloudera cloudera
                                              4705957 2020-11-27 02:02 Genome.fa
drwxr-xr-x - cloudera cloudera
                                                      0 2020-11-07 02:34 Outpul2
drwxr-xr-x - cloudera cloudera
drwxr-xr-x - cloudera cloudera
                                                      0 2020-11-01 22:58 Output12
                                                      0 2020-11-01 22:47 Output13
-rw-r--r-- 1 cloudera cloudera 57833 2021-05-28 01:41 Task3.txt
drwxr-xr-x - cloudera cloudera
drwxr-xr-x - cloudera cloudera
                                                      0 2020-12-04 01:12 assignment
                                                      0 2020-12-04 01:27 assignment1
drwxr-xr-x - cloudera cloudera drwxr-xr-x - cloudera cloudera
                                                      0 2020-11-27 02:10 checkoutput
                                                     0 2020-10-30 09:09 hadoops1
                                                    0 2020-10-30 09:16 hadoops2
                                                    0 2020-10-30 09:44 hadoops3
                                                     0 2020-10-31 10:10 hadoops4
                                                    0 2020-10-31 11:29 hadoops5
drwxr-xr-x - cloudera cloudera
                                                    0 2020-10-31 11:33 hadoops8
-rw-r--r-- 1 cloudera cloudera
drwxr-xr-x - cloudera cloudera
drwxr-xr-x - cloudera cloudera
                                                    61 2020-10-31 09:58 inputfile
                                                    0 2020-11-08 22:05 output100
                                                     0 2020-11-08 22:13 output101
drwxr-xr-x - cloudera cloudera
drwxr-xr-x - cloudera cloudera
drwxr-xr-x - cloudera cloudera
                                                     0 2020-11-08 22:17 output102
                                                    0 2020-12-04 08:39 output assignment
                                                    0 2020-12-04 06:24 outputass
drwxr-xr-x - cloudera cloudera
-rw-r--r-- 1 cloudera cloudera
                                                     0 2020-10-31 10:03 outputs1
                                               485246 2020-11-01 22:23 test_access_log
-rw-r--r-- 1 cloudera cloudera
                                                    59 2020-11-08 22:02 wordcount
 -rw-r--r--
                1 cloudera cloudera
                                                    59 2020-11-08 22:11 wordcount.txt
[cloudera@quickstart Desktop]$
```

Fig1: I put the data file into the mapreduce.

Fig2: Executing the Hadoop streaming command.

```
Found 25 items
-rw-r--r-- 1 cloudera cloudera
                                    4705957 2020-11-27 02:02 Genome.fa
drwxr-xr-x
            - cloudera cloudera
                                          0 2020-11-07 02:34 Outpul2
            - cloudera cloudera
                                          0 2020-11-01 22:58 Output12
drwxr-xr-x
                                          0 2020-11-01 22:47 Output13
drwxr-xr-x

    cloudera cloudera

            1 cloudera cloudera
                                      57833 2021-05-28 01:41 Task3.txt
drwxr-xr-x - cloudera cloudera
                                         0 2021-05-28 01:47 Task3 output
                                          0 2020-12-04 01:12 assignment
drwxr-xr-x - cloudera cloudera
drwxr-xr-x
            - cloudera cloudera
                                          0 2020-12-04 01:27 assignment1
                                          0 2020-11-27 02:10 checkoutput
drwxr-xr-x

    cloudera cloudera

drwxr-xr-x
            - cloudera cloudera
                                          0 2020-10-30 09:09 hadoops1
                                          0 2020-10-30 09:16 hadoops2
drwxr-xr-x

    cloudera cloudera

            - cloudera cloudera
                                          0 2020-10-30 09:44 hadoops3
drwxr-xr-x
drwxr-xr-x

    cloudera cloudera

                                          0 2020-10-31 10:10 hadoops4
drwxr-xr-x

    cloudera cloudera

                                          0 2020-10-31 11:29 hadoops5
drwxr-xr-x

    cloudera cloudera

                                          0 2020-10-31 11:33 hadoops8
-rw-r--r--
            1 cloudera cloudera
                                         61 2020-10-31 09:58 inputfile
drwxr-xr-x
            - cloudera cloudera
                                          0 2020-11-08 22:05 output100
drwxr-xr-x
            - cloudera cloudera
                                          0 2020-11-08 22:13 output101
            - cloudera cloudera
                                          0 2020-11-08 22:17 output102
drwxr-xr-x
            - cloudera cloudera
drwxr-xr-x
                                          0 2020-12-04 08:39 output assignment
drwxr-xr-x
            - cloudera cloudera
                                          0 2020-12-04 06:24 outputass
                                          0 2020-10-31 10:03 outputs1
drwxr-xr-x
            - cloudera cloudera
                                     485246 2020-11-01 22:23 test_access_log
-rw-r--r--
            1 cloudera cloudera
-rw-r--r--
                                         59 2020-11-08 22:02 wordcount
            1 cloudera cloudera
-rw-r--r--
            1 cloudera cloudera
                                         59 2020-11-08 22:11 wordcount.txt
[cloudera@quickstart Desktop]$ hdfs dfs -ls Task3 output
Found 2 items
            1 cloudera cloudera
                                          0 2021-05-28 01:47 Task3 output/ SUCCESS
-rw-r--r--
-rw-r--r--
                                       2195 2021-05-28 01:47 Task3_output/part-00000
            1 cloudera cloudera
```

Fig3: Generated Output Directory.

```
[cloudera@quickstart Desktop]$ hdfs dfs -cat Task3 output/part-00000
10specifies
                21
11
        22
23
        9
        16
a1
        55
agarwal 45
aggarwal
                45
agrawal 45
akshay 29
along 28
also
anything
                30
appearing
                29
arrowdropupsave 4
arrowdropupsave1
                         1
```

Fig4: Frequency of each word

```
filename
                 4
filter
find
        38
found
        40
free
        60
geek
        29
geekfiletxt
                 130
geekfiletxt12
                 24
geekfiletxt4
                 11
                         33
geekfiletxtoutput
geeksforgeeks
                 58
geeksforgeeksorg
                         29
generate
give
        70
given
globally
                 1
great
        60
grep
        389
help
        29
idegeeksforgeeksorg
                         1
ignores 2
incorrect
                 30
                 30
information
input
        5
insensitive
insensitively
inverting
                 17
1
        22
learn
        78
                 124
learnunix
```

SnapShots of Visualizing the data using Kibana.

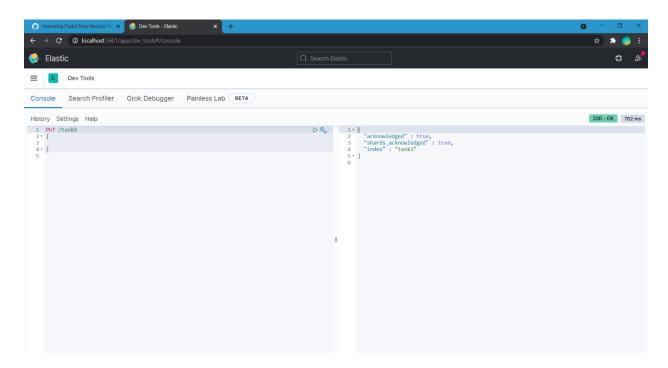
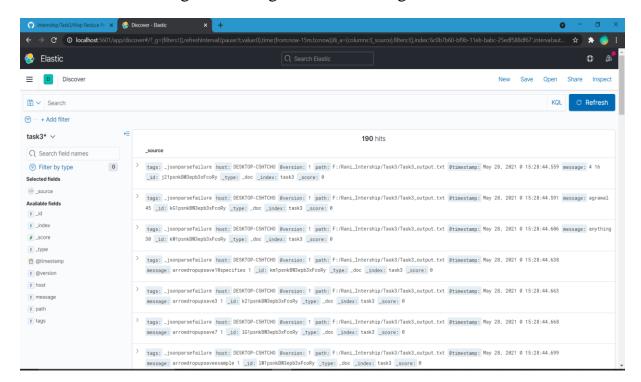


Fig 5: Creating index for storing the data.



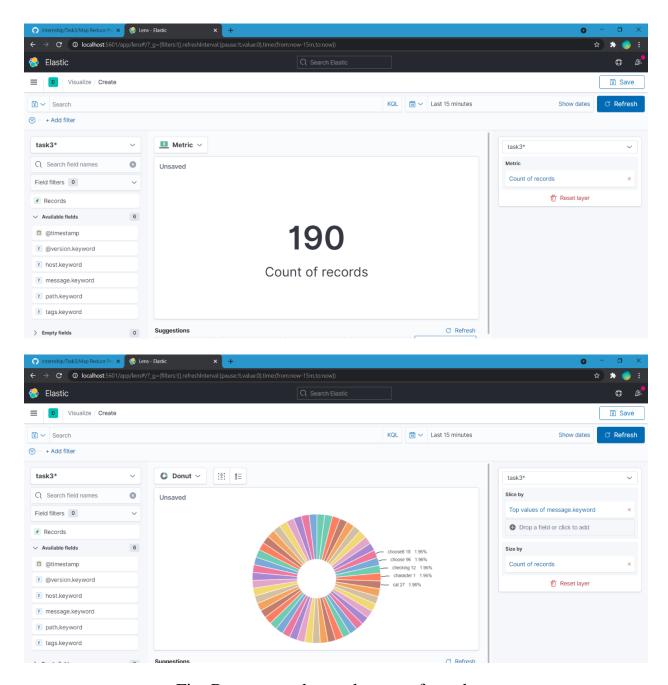


Fig: Represents the total count of word.