

Course: Big Data

Lab 03

MapReduce

Fill answers of the questions below in the given tables.

Your screenshots must **contain commands** for required operations.

Question 1:

Given a tsv file [WHO-COVID-19-20210601-213841.tsv](#) which is corresponding to the [WHO Coronavirus \(COVID-19\) Dashboard](#).

Students are required to create a folder, named **lab03**, in HDFS and then copy the tsv to **lab03/input/**

Take a screenshot to show the content of **lab03/input/** in HDFS

```
hohuan@ubuntu:~/Desktop/hadoop$ bin/hdfs dfs -mkdir lab03
hohuan@ubuntu:~/Desktop/hadoop$ bin/hdfs dfs -mkdir lab03/input
hohuan@ubuntu:~/Desktop/hadoop$ bin/hdfs dfs -put /home/hohuan/Desktop/WHO-COVID-19-20210601-213841.tsv lab03/input/
2024-01-23 07:48:26,173 INFO sasl.SaslDataTransferClient: SASL encryption trust check: localhostTrusted = false, remoteHostTrusted = false
hohuan@ubuntu:~/Desktop/hadoop$ bin/hdfs dfs -ls lab03/input/
Found 1 items
-rw-r--r-- 2 hohuan supergroup 28907 2024-01-23 07:48 lab03/input/WHO-COVID-19-20210601-213841.tsv
hohuan@ubuntu:~/Desktop/hadoop$ bin/hdfs dfs -cat lab03/input/WHO-COVID-19-20210601-213841.tsv
2024-01-23 07:51:32,000 INFO sasl.SaslDataTransferClient: SASL encryption trust check: localhostTrusted = false, remoteHostTrusted = false
Name WHO Region Cases - cumulative total Cases - cumulative total per 100000 population Cases - newly reported in last 7 days Cases - newly reported in last 7 days per 100000 population Ca
ses - newly reported in last 24 hours Deaths - cumulative totalDeaths - cumulative total per 100000 population Deaths - newly reported in last 7 days Deaths - newly reported in last 7 days per 100000 population
Deaths - newly reported in last 24 hours Transmission Classification
Global 170,363,652.000 2,192,386 3,383,026.000 43,337 358,091.000 3,546,870.000 45,436 76,025.000 0.974 7,370.000
United States of America Americas 32,929,178.000 9,948,310 131,305.000 39,670 0.000 588,596.000 177,820 3,896.000 1.180 0.000 Community transmission
India South-East Asia 28,175,044.000 2,041,660 1,226,170.000 88,850 127,510.000 331,895.000 24,050 24,664.000 1.790 2,795.000 Clusters of cases
Brazil Americas 10,515,120.000 7,769,658 431,802.000 203,170 43,520.000 401,931.000 217,320 12,863.000 0.850 874.000 Community transmission
France Europe 5,566,214.000 9,559,220 59,437.000 91,390 9,541.000 108,558.000 106,910 756.000 1.100 15.000 Community transmission
Turkey Europe 5,242,911.000 6,216,470 56,424.000 66,900 6,933.000 47,405.000 50,210 1,137.000 1.350 134.000 Community transmission
Russian Federation Europe 5,071,917.000 3,475,480 62,000.000 42,490 8,475.000 121,501.000 83,200 2,700,0001.850 339.000 Clusters of cases
The United Kingdom Europe 4,484,000.000 6,805,280 21,510.000 31,700 3,111.000 127,781.000 188,230 60.000 0.000 6.000 Community transmission
Italy Europe 4,218,003.000 7,068,910 23,820.000 39,940 2,940.000 126,040.000 211,140 821.000 1.380 44.000 Clusters of cases
Argentina Americas 3,753,009.000 8,305,220 214,125.000 473,770 21,340.000 77,456.000 171,380 3,393,0007.510 348.000 Community transmission
Germany Europe 3,681,126.000 4,426,200 29,480.000 35,450 1,978.000 88,442.000 100,340 1,019.000 1.230 36.000 Community transmission
Spain Europe 3,643,176.000 7,739,220 13,542.000 28,530 0.000 79,888.000 168,780 54.000 0.110 0.000 Community transmission
Colombia Americas 3,383,279.000 6,649,150 150,823.000 296,410 20,218.000 88,282.000 173,580 3,558,0006.990 535.000 Community transmission
Iran (Islamic Republic of) Eastern Mediterranean 2,913,136.000 3,468,310 69,613.000 82,880 11,042.000 80,156.000 95,430 1,308.000 1.560 217.000 Community transmission
Poland Europe 2,872,283.000 7,566,900 6,102.000 16,080 233.000 73,745.000 194,208 800.000 2.110 7.000 Community transmission
Mexico Americas 2,412,810.000 1,871,370 16,286.000 12,570 1,387.000 223,587.000 173,350 1,809.000 1.440 52.000 Community transmission
Ukraine Europe 2,202,494.000 5,036,140 18,639.000 42,620 1,022.000 50,536.000 115,550 1,100.000 2.520 64.000 Community transmission
Peru Americas 1,955,469.000 5,930,720 30,188.000 91,530 3,818.000 69,342.000 210,310 1,289.000 3.910 140.000 Community transmission
Indonesia South-East Asia 1,921,703.000 666,810 40,576.000 14,830 5,662.000 50,578.000 18,490 1,123.000 0.410 174.000 Community transmission
South Africa Africa 1,662,825.000 2,803,680 27,360.000 46,130 3,755.000 50,439.000 95,160 637.000 1.070 76.000 Community transmission
Czechia Europe 1,661,272.000 15,534,710 3,188.000 29,740 113.000 30,108.000 281,540 80.000 0.750 4.000 Community transmission
Netherlands Europe 1,647,418.000 9,461,790 21,424.000 123,070 2,785.000 17,621.000 101,230 79.000 0.450 6.000 Community transmission
Chile Americas 1,384,346.000 7,241,740 49,005.000 250,770 6,839.000 29,300.000 153,270 752.000 3.930 132.000 Community transmission
Canada Americas 1,378,971.000 3,653,660 19,112.000 50,640 2,237.000 25,512.000 67,600 281.000 0.740 34.000 Community transmission
Philippines Western Pacific 1,230,301.000 1,122,730 45,601.000 41,610 6,684.000 20,966.000 19,130 983.000 0.900 100.000 Community transmission
```

Question 2:

Create one and only one java file, named **ASEANCaseCount.java**, to run a MapReduce job that counts the number of cumulative total cases among ASEAN countries (*South-East Asia Region in the given data table*).

The output of the MapReduce job is located in **lab03/output-java/**.

Submit the source code file following the instructions in Submission Notice.

Question 3 (optional):

Create a pair of Python files, named **ASEANDeathCountMapper.py** and **ASEANDeathCountReducer.py**, to run a MapReduce job that counts the number of cumulative total deaths among ASEAN countries (*South-East Asia Region in the given data table*).

The output of the MapReduce job is located in **lab03/output-python/**.

Submit the source code files following the instructions in Submission Notice.

Submission Notice

- Export your answer file as pdf
- Rename the pdf following the format:
lab03_<student number>_HoTen.pdf
E.g. lab03_123456_NguyenThanhAn.pdf
If you have not been assigned a student number yet, then use 123456 instead.
- Create a folder with the name as **<student number>_HoTen**, which contains
 - **<student number>_HoTen.pdf** → your answer
 - **java/** | → Java source code folder
| **ASEANCaseCount.java**
 - **python/** | → Python source code folder
| **ASEANDeathCountMapper.py**
| **ASEANDeathCountReducer.py**
- Compress the folder **<student number>_HoTen** in zip format and finally submit to the given form.
E.g. 123456_HoTen.zip
- Careless mistakes in filename, format, question order, etc. are not accepted (0 pts).