Aims

This exercise aims to get you to:

- Compile, run, and debug MapReduce tasks via Hadoop Streaming
- Compile, run, and debug MapReduce tasks via MRJob
- Apply the design pattern "in-mapper combining" you have learned in Chapter 2.2 to MapReduce programming

One Tip on Hadoop File System Shell

Following are the three commands which appear same but have minute differences:

```
    hadoop fs {args}
    hadoop dfs {args}
    hdfs dfs {args}
```

The first command: fs relates to a generic file system that can point to any file system like local, HDFS etc. So this can be used when you are dealing with different file systems such as Local FS, HFTP FS, S3 FS, and others.

The second command: dfs is very specific to HDFS. It would work for operations relates to HDFS. This has been **deprecated** and we should use hdfs dfs instead.

The third command: It is the same as 2^{nd} . It would work for all the operations related to HDFS and is the recommended command instead of hadoop dfs.

Thus, in our labs, it is always recommended to use hdfs dfs {args}.

Hadoop Streaming

<u>Hadoop streaming</u> is a utility that comes with the Hadoop distribution. The utility allows you to create and run Map/Reduce jobs with any executable or script as the mapper/reducer.

NOTE: Please be careful when directly copying and pasting the commands in this instruction. It is better to enter the command by yourself.

1. In Lab1, you tested an application of word count. In Linux, we command can be used to find out the **number of lines**, word count, byte and character count in the files specified in the file arguments. Test we command:

NOTE: Please use the **whereis** command to check the location of the **cat** and **wc** commands. For example, \$whereis wc

```
$ wc $HADOOP HOME/etc/hadoop/*.xml
```

2. Run a streaming task with /bin/cat as the mapper and /bin/wc as the reducer:

```
$ hadoop jar $HADOOP_HOME/share/hadoop/tools/lib/hadoop-streaming-*.jar \
    -input input \
```

```
-output output2 \
-mapper /bin/cat \
-reducer /bin/wc
```

3. Check out the output:

```
$ hdfs dfs -cat output2/*

If you are using Mac OS, please add '\' before the '*' if the file/folder exists but you still see some errors. For example, in step 3 (check out the output)

$ hdfs dfs -cat output2/\*
```

4. To specify the number of reducers, for example two, use:

```
$ hadoop jar $HADOOP_HOME/share/hadoop/tools/lib/hadoop-streaming-*.jar \
-D mapreduce.job.reduces=2 \
-input input \
-output output3 \
-mapper /bin/cat \
-reducer /usr/bin/wc
```

Specifying Python Scripts as Mapper/Reducer

Next, you will learn to run Map/Reduce jobs with Python script as the mapper/reducer.

1. Create a folder "Lab2", and put all your codes written in this week's lab in this folder. Next, create a file named mapper.py and copy the codes below into the file.

```
#!/usr/bin/python3
import sys

for line in sys.stdin:
   line = line.strip()
   words = line.split()

  for word in words:
      print (word + "\t" + "1")
```

Make sure this file has the execution permission:

```
$ chmod +x mapper.py
```

2. Similarly, create a file named reducer.py and copy the codes below into the file.

```
#!/usr/bin/python3
import sys

results = {}
for line in sys.stdin:
    word, frequency = line.strip().split('\t', 1)
    results[word] = results.get(word, 0) + int(frequency)
words = list(results.keys())

for word in words:
    print(word, results[word])
```

Also, make sure this file has the execution permission:

```
$ chmod +x reducer.py
```

Compare the above code with that provided in slide 38 of Lecture 2.1. What are the differences? What problem the above approach may encounter?

3. Run the application (you need to enter the **Lab2** folder):

```
$ hadoop jar $HADOOP_HOME/share/hadoop/tools/lib/hadoop-streaming-*.jar \
    -input input \
    -output python_output \
    -mapper mapper.py \
    -reducer reducer.py \
    -file mapper.py \
    -file reducer.py
```

4. Check out the output (you can also download the file from HDFS):

```
$ hdfs dfs -cat python_output/*
```

Try to Write Your First Hadoop Streaming Job

1. Download the test file "pg100.txt" to your home folder from WebCMS3, and put it to HDFS:

```
$ hdfs dfs -rm input/*
$ hdfs dfs -put ~/pg100.txt input
```

2. Now please write your first MapReduce job with Hadoop Streaming to accomplish the following task:

Output the number of words that start with each letter. This means that for every letter we want to count the total number of words that start with that letter. In your implementation, please first convert all words to lowercase. You can ignore all non-alphabetic characters. Create "mapper. py", "reducer.py" and "combiner.py" scripts in the folder "LetterCount" to finish this task.

Hint: In the (key, value) output, each letter is the key, and its count is the value.

You can run the three scripts on Hadoop MapReduce by:

Compare your results with the answer provided at:

https://webcms3.cse.unsw.edu.au/COMP9313/23T3/resources/92020

Install MRJob

If you use Mac, you can skip installing pip3, since it is already in Mac OS.

As you will install mrjob with pip3, you should first install pip3:

```
$ sudo apt install python3-pip
```

Then, you will be asked to enter the sudo password: comp9313

When pip3 is successfully installed, install mrjob with pip3:

```
$ pip3 install mrjob
```

Warning: You also need to configure and start YARN since running mrjob on Hadoop requires YARN. Please edit the mapred-site.xml and yarn-site.xml by following Lab 1 instructions. Then, start HDFS and YARN by running "start-dfs.sh" and "start-yarn.sh".

An Example of MRJob

Create a file called mr_word_count.py in the folder Lab2 and type this into it:

```
from mrjob.job import MRJob

class MRWordFrequencyCount(MRJob):

    def mapper(self, _, line):
        yield "chars", len(line)
        yield "words", len(line.split())
        yield "lines", 1

    def reducer(self, key, values):
        yield key, sum(values)

if __name__ == '__main__':
    MRWordFrequencyCount.run()
```

We still use the downloaded file "pg100.txt" as input (make sure it exists in /home/comp9313). Run the codes locally with:

```
$ python3 mr_word_count.py ~/pg100.txt
```

You will see the result:

```
Running step 1 of 1...
job output is in /tmp/mr_word_count.comp9313.20210927.075446.090653/output
Streaming final output from /tmp/mr_word_count.comp9313.20210927.075446.090653/output...
"chars" 5340313
"lines" 124787
"words" 904061
Removing temp directory /tmp/mr_word_count.comp9313.20210927.075446.090653...
```

If you want to run in Hadoop, use the "-r hadoop" option, and then you can check the result in the file "output":

```
$ python3 mr_word_count.py -r hadoop pg100.txt > output
```

If your files are in HDFS, you can run like:

```
$ python3 mr word count.py -r hadoop hdfs://localhost:9000/user/comp9313/input
```

If you want to store your results in HDFS (e.g., in the "output" folder) as well, you can run:

```
$ python3 mr_word_count.py -r hadoop hdfs://localhost:9000/user/comp9313/input -o
hdfs://localhost:9000/user/comp9313/output
```

There are different ways to run your job, see more details <u>here</u>.

Try to Write Your First MRJob Program

Please write your first mrjob program to complete the above "letter count" task, and compare it with the Hadoop streaming approach.

Improve WordCount by In-Mapper Combining

A better tokenization method:

Use the following codes to tokenize a line of document:

```
import re
words = re.split("[ *$&#/\t\n\f\"\'\\,.:;?!\[\](){}<>~\-_]", line.lower())
```

Documents will be split according to all characters specified (i.e., " $*$&#/\t^"'\.:;? {}<>~-_")$, and higher quality terms will be generated.

Convert all terms to lower case as well (by using lower() method).

Apply this to mapper.py of WordCount we have used. Note that you need to check if the word is an empty string now.

- a) Put the input file to HDFS by:
 - \$ hdfs dfs -rm input/*
 \$ hdfs dfs -put ~/pg100.txt input
- b) Go into the folder Lab2 and use the existing mapper.py and reducer.py scripts.
- c) Use the new method to tokenize a line of document
- d) Run the application with Hadoop Streaming/mrjob
- e) Remember to delete to output folder if it exists in HDFS
- f) If you forget the details, please refer to the previous instructions.

Type the following command in the terminal:

```
$ hdfs dfs -cat output/* | head
```

You should see results:

```
comp9313@comp9313-VirtualBox:~/9313_files$ hdfs dfs -cat python_output/* | head
0 2
00 1
000 1
01 1
02 1
03 1
04 1
05 1
08 1
1 362
```

Use the following command:

```
$ hdfs dfs -cat output/* | tail
```

You should see:

```
comp9313@comp9313-VirtualBox:~/9313_files$ hdfs dfs -cat python_output/* | tail
zephyrs 1
zip 1
zipped 1
zir 2
zo 1
zodiac 1
zodiacs 1
zone 1
zounds 24
zwagger 1
```

Apply the in-mapper combing approach:

Based on the previous python scripts, you are required to write an improved version using the "in-mapper combining" approach.

Hadoop Streaming: Create a new script "mapper2.py" in the folder Lab2 and solve this problem. Your results should be the same as generated by mapper.py.

mrjob: Create a new script "mr_word_count2.py" in the folder Lab2 and solve this problem. Your results should be the same as generated by mapper.py.

Hints:

- 1. Refer to the pseudo-code shown in slide 23 of Chapter 2.2.
- 2. You can use a dictionary in the mapper script to buffer the partial results for Hadoop streaming.
- 3. You need to override the methods mapper init() and mapper final() in mrjob
- 4. Do you need to change the reducer?

Solutions of these Problems

I hope that you are able to finish all problems by yourself, since the hints are already given. All the source codes will be published in the course homepage on Friday.