Carnegie Mellon University in Qatar

Distributed Systems

15-440 - Fall 2020

Project 4

Out: November 22, 2020 Due: December 3, 2020

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1 Summary & Intended Learning Outcomes

MapReduce is now a pervasive analytics engine on the cloud. Hadoop is an open source implementation of MapReduce and is currently enjoying wide popularity.

Hadoop offers a high-dimensional space of configuration parameters, which makes it difficult for practitioners to set for efficient and cost-effective execution. In this project, you will implement the K-Means clustering algorithm using Hadoop MapReduce and "characterize" the map phase parallelism accordingly. For the sake of this project, map phase parallelism is defined in terms of two Hadoop configurable parameters; the number of available map slots and the number of map tasks running over the slots. Within the confines of MapReduce (and this project), we define the characterization of the map phase parallelism as the process of observing, identifying and explaining various MapReduce runtime responses to different map task and map slot numbers. To this end, in addition to characterizing the map phase parallelism, you will compare and contrast the performance of your MPI K-Means implementation (from $\mathbf{P3}$) against your Hadoop MapReduce K-Means implementation from this project.

In short, the learning outcomes of this project are as follows:

- 1. Apply MapReduce to a popular real problem, namely cluster analysis using K-Means algorithm.
- 2. Characterize the map phase parallelism using the K-Means clustering algorithm.
- 3. Compare and contrast your MPI and Hadoop MapReduce implementations of *K-Means* in terms of performance and development effort.

2 **Project Objectives**

The overall goal of this project is to get a clear understanding on Hadoop map parallelism and how different parallel implementations of the same algorithm compare against each other. You will conduct and analyze some scalability studies on various degrees of parallelism for *K-Means*. The project will provide students with: (1) deep insights onto how parallelism affects performance in large-scale settings, and (2) a practical experience augmented with a methodology for solving clustering problems (and alike) on a distributed system using MapReduce.

3 Implementation Guidelines

In this project, you will provide a MapReduce implementation for K-Means with two types of data sets; a data set of data points and a data set of DNA strands (as was done in **P3**). Please use the datasets you generated in **P3** to run and test your MapReduce K-means implementation. For a complete explanation of the K-Means algorithm, please refer to the write-up of **P3**.

For this project, use the 4-VM virtual cluster (which can be deemed as a private cloud) provided to you in **P3**. We have already installed and tested Hadoop 0.20.2 for you. Hence, the clusters are ready to run your Hadoop MapReduce code.

4 Experimentation and Analysis

After implementing K-Means using Hadoop MapReduce, please conduct experiments and report on the following:

- A comparison between your 3 different *K-Means* implementations (the sequential and the MPI ones from **P3**, and the MapReduce one from this project) in terms of performance and development effort.
- Three scalability studies. In particular:
 - A scalability study on the number of map slots per each virtual machine (VM) for fixed cluster, data set and HDFS block sizes. Specifically, use 1, 2, and 4 map slots per each VM on your 4-VM cluster, with a data set size of your choice (use only the data set of the 2D data points) and 64MB HDFS block size.
 - A scalability study on the HDFS block size per fixed data set size, cluster size and map slot value. Specifically, use 32MB, 64MB and 128MB HDFS block sizes on your 4-VM cluster with 2 map slots per each VM and a data set size of your choice (again, use only the data set of the 2D data points).
 - A scalability study on the number of VMs per fixed data set size, HDFS block size, and map slot value. Specifically, use 1, 2, 3 and 4 VMs with 64MB HDFS block size, 2 map slots per a VM and a data set size of your choice (again, use only the data set of the 2D data points).

Note that these scalability studies are, in essence, the methodology we are using to characterize the map phase parallelism. You should report on your observations concerning the obtained parallelism (in each of the scalability studies) and the consequent overall influence on Hadoop MapReduce performance for *K-Means*. For instance, you should explain why a larger number of map tasks can sometimes expedite performance while at other times it might degrade performance.

- A discussion on:
 - Your experience in applying MapReduce to the K-Means clustering algorithms.
 - Your insights concerning the performance trade-offs of MPI and MapReduce with K-Means.
 - Your thoughts on the applicability of *K*-Means to MapReduce.
 - Your recommendations regarding the usage of MapReduce for algorithms similar to K-Means.

5 Final Deliverables

As final deliverables, you should submit:

- 1. An archive containing a fully tested and debugged code for your MapReduce *K-Means* implementation. Specifically, you must submit <u>two</u> programs for the *K-Means* implementation:
 - (1) points.java
 - (2) dna.java

They must follow the specifications below:

- Input: (*in this order*)
 - The **Input directory** containing the data
 - The **Output directory** containing the data
 - Total number of **points**/strands provided
 - Number of **clusters**
 - Number of iterations (if it's given as 0, your program should stop as necessary)
 - 1: length of DNA strand (only applies for the DNA strands K-Means implementations)
 - * Sample program execution:

- Output: a file with the following information for each cluster c_i (all separated by new-lines):
 - The final centroid value for c_i
 - The number of points/strands assigned to the cluster c_i
 - * Sample output for 2D points (the above, separated by commas on each line):

// x-coordinate, y-coordinate, number of points
5.212, 9.880, 400
1.511, 3.201, 320
...

- * <u>Note</u>: The output file(s) must be in the output directory specified above, NOT locally nor in any other HDFS path
- 2. An **article** with a maximum of 5 pages (*similar to research articles*) that presents your solution, findings, observations and analysis.

6 Rubric

MapReduce K-Means (50 Points)

- K-Means using MapReduce for 2D data points
- K-Means using MapReduce for DNA strands.

Details (these are same for DNA and 2D data points versions):

- $\,\circ\,$ The code compiles and runs correctly
- $\circ\,$ Create map and reduce classes, extend MapReduceBase and implement Mapper and Reducer
- Configuring jobs correctly including setting the mapper and reducer classes, the input and output formats, the file input and output formats, launching a job, and looping over for new rounds
- $\circ~{\rm Read}$ the value of centroids at each mapper
- Create the map function with correct input and output key-value pairs (i.e., correct input and output formats) (3%)
- $\circ\,$ Apply K-Means at the mapper
- Create the reduce function with correct input and output key-value pairs (i.e., correct input and output formats)
- $\circ~$ Compute new means at the reducer
- $\circ~$ Write the value of centroids at each reducer
- Outputting final results and aborting cleanly

Write-up (47.5 Points)

- Performance comparison between sequential, MPI, and MapReduce with a fixed dataset size
- Development effort comparison between sequential, MPI and MapReduce
- MapReduce scalability w.r.t 1, 2, and, 4 map slots
- MapReduce scalability w.r.t 32MB, 64MB, and 128MB block sizes
- MapReduce scalability w.r.t 1, 2, 3 VMs
- Performance trade-offs between MPI and MapReduce
- Thoughts on the applicability of K-Means to MapReduce
- Recommendations regarding the usage of MapReduce for algorithms similar to K-Means
- Paper structure, level of writing, and language

Code Style (2.5 Points)

• Method Comments, Block comments, Readability, Dead code, Code Design

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 $25 \mathrm{pts}$





3pts

| 3pts | |
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| 3pts | |
| 3pts | |

| 10pts |
|--------|
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| 1.5pts |
| 10pts |
| 10pts |
| 10pts |
| 1.5pts |
| 1.5pts |
| 1.5pts |
| 1.5pts |

7 Late Policy

- If you hand in on time, there is no penalty.
- 0-24 hours late = 25% penalty.
- 24-48 hours late = 50% penalty.
- More than 48 hours late = you lose all the points for this project.

 ${\bf NOTE}:$ You ${\bf CANNOT}$ use your grace-days quota. For details about the quota, please refer to the syllabus.