Document clustering using K-means under Mahout

1. Edit system variables (can be ignored if you already successfully installed Hadoop)

Under your home directory, edit the file .bash_profile (for Mac) or .bashrc (for Linux) and add the following lines.

\$vi .bash_profile

export JAVA_HOME=\$(/usr/libexec/java_home)

export HADOOP_HOME=/Users/Double]/Software/hadoop-1.0.3 (replace with your hadoop directory)

export HADOOP_CONF_DIR=/Users/DoubleJ/Software/hadoop-1.0.3/conf

Then save and exit the file (press ESC and then : wq + Enter). To make the edition be effective, use source command. $source .bash_profile$

OR

\$source .bashrc

2. Download data

Download reuters.tar.gz file and store it in the folder of mahout-work (you can create it first if not exists)

 $\label{lem:condition} $$ \curl\ http://kdd.ics.uci.edu/databases/reuters 21578/reuters 21578.tar.gz -o/Users/DoubleJ/Workspace/mahout-work/reuters.tar.gz$

Note: The command line above is in one line.

3. Uncompress data

Create a new folder sgm under mahout-work first and then uncompress the original zip file into the sgm folder

\$mkdir /Users/DoubleJ/Workspace/mahout-work/sgm
\$tar zxvf /Users/DoubleJ/Workspace/mahout-work/reuters.tar.gz -C /Users/DoubleJ/Workspace/mahout-work/sgm

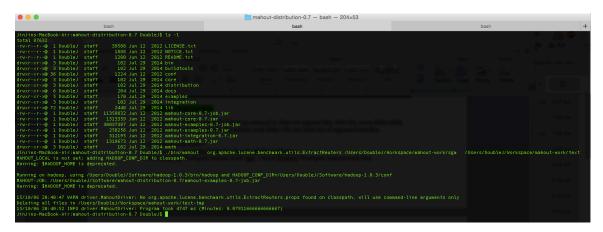
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4. Obtain the segment data

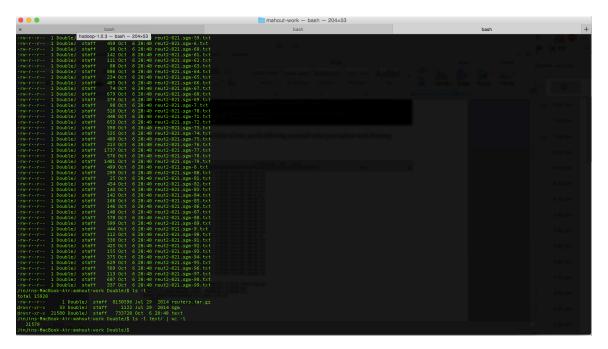
Go to the mahout folder and run the following command to obtain the segment data. After this, a new folder called text will be automatically created under mahout-work folder. The text folder has all segmented text files.

- \$ cd <your mahout folder>
- $\$./bin/mahout \quad org. apache.lucene. benchmark.utils. Extract Reuters$
- /Users/DoubleJ/Workspace/mahout-work/sgm /Users/DoubleJ/Workspace/mahout-work/text



To check how many files in the folder of text, use the following command under your mahout-work directory.

\$ls -l text/ | wc -l



5. Generate the sequence directory, which contains all texts

5.1 Send the segment data to the Hadoop file system

\$cd \$HADOOP_HOME

OR if you do not set $HADOOP_HOME$ environment variable in the .bashrc or .bash_profile

- \$cd <your hadoop folder>
- \$./bin/start-all.sh
- \$./bin/hadoop fs -mkdir hdfs-mahout (create a folder hdfs-mahout in HDFS)
- $\$./bin/hadoop\ fs\ -put\ /Users/DoubleJ/Workspace/mahout-work/text \ \ hdfs-mahout\ (takes\ for\ a\ while)$

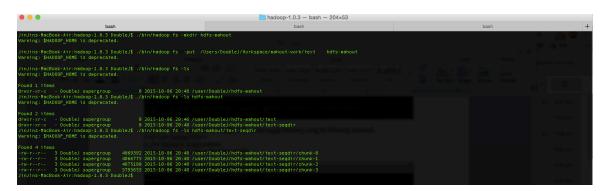
5.2 Create the sequence directory

\$cd <your mahout folder>

\$./bin/mahout seqdirectory -i hdfs-mahout/text -o hdfs-mahout/text-seqdir -c UTF-8 -chunk 5

To check new files generated in the hdfs-mahout/text-seqdir directory, using the following command.

- \$ cd <your hadoop folder>
- \$./bin/hadoop fs -ls hdfs-mahout
- \$./bin/hadoop fs -ls hdfs-mahout/text-seqdir



6. Create vector files to represent these documents

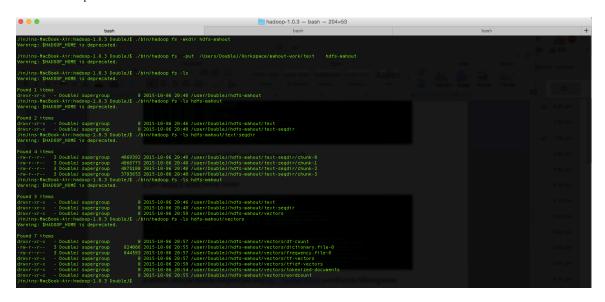
\$cd <your mahout folder>

\$./bin/mahout seq2sparse -i hdfs-mahout/text-seqdir/ -o hdfs-mahout/vectors --maxDFPercent 85 --namedVector

To check new files generated in the HDFS, use the following commands.

\$cd <your hadoop folder>

- \$./bin/hadoop fs -ls hdfs-mahout
- \$./bin/hadoop fs -ls hdfs-mahout/vectors



In addition, when you check the completed jobs using http://localhost:50030, you will see the following screen.

Scheduling Information

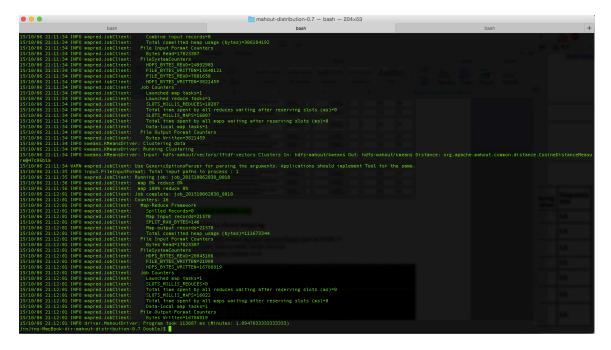
Queue Name	State	Scheduling Information
default	running	N/A
Filter (Jobid, Pr Example: 'user:smitl	iority, Us n 3200' will	ser, Name) [I filter by 'smith' only in the user fi
Running J	obs	
none		

Completed Jobs

Jobid	Priority	User	Name	Map % Complete		Maps Completed →	Reduce % Complete	Reduce Total	Reduces Completed	Job Scheduling Information	Diagnostic Info
job 201510062030 0004	NORMAL	DoubleJ	PartialVectorMerger::MergePartialVectors	100.00%	1	1	100.00%	1	1	NA	NA
job 201510062030 0005	NORMAL	DoubleJ	VectorTfldf Document Frequency Count running over input: hdfs- mahout/vectors/tf-vectors	100.00%	1	1	100.00%	1	1	NA	NA
job_201510062030_0006	NORMAL	DoubleJ	: MakePartialVectors: input-folder: hdfs-mahout/vectors/tf-vectors, dictionary-file: hdfs-mahout/vectors/frequency.file-0	100.00%	1	1	100.00%	1	1	NA	NA
job_201510062030_0007	NORMAL	DoubleJ	PartialVectorMerger::MergePartialVectors	100.00%	1	1	100.00%	1	1	NA	NA
job 201510062030 0001	NORMAL	DoubleJ	DocumentProcessor::DocumentTokenizer: input-folder: hdfs-mahout/text-seqdir	100.00%	4	4	100.00%	0	0	NA	NA
job 201510062030 0002	NORMAL	DoubleJ	DictionaryVectorizer::WordCount: input-folder: hdfs-mahout/vectors/tokenized-documents	100.00%	4	4	100.00%	1	1	NA	NA
job 201510062030 0003	NORMAL	DoubleJ	DictionaryVectorizer::MakePartialVectors: input-folder: hdfs-mahout/vectors/tokenized-documents, dictionary-file: hdfs-mahout/vectors/dictionary.file-0	100.00%	4	4	100.00%	1	1	NA	NA

7. Running K-means

\$cd <your mahout folder>

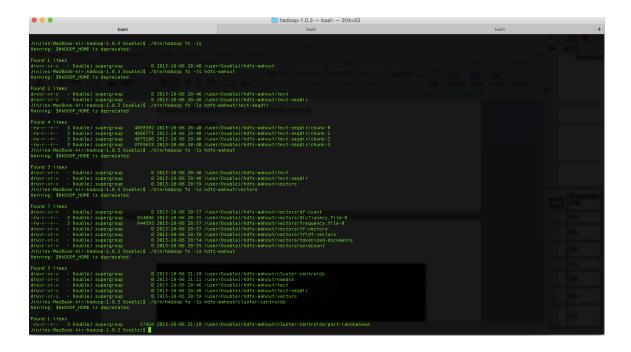


To check new files generated in HDFS, use the following commands.

\$cd <your hadoop folder>

\$./bin/hadoop fs -ls hdfs-mahout/

 $./bin/hadoop\ fs\ -ls\ hdfs-mahout/cluster-centroids$



In addition, when you check the completed jobs using http://localhost:50030, you will see the following screen.

Running Jobs

none

Completed Jobs

Jobid	Priority	User	Name	Map % Complete	Map Total	Maps Completed	Reduce % Complete	Reduce Total	Reduces Completed	Job Scheduling Information	Diagnostic Info
job 201510062030 0001	NORMAL	DoubleJ	DocumentProcessor::DocumentTokenizer: input-folder: hdfs-mahout/text-seqdir	100.00%	4	4	100.00%	0	0	NA	NA
job 201510062030 0002	NORMAL	DoubleJ	DictionaryVectorizer::WordCount: input-folder: hdfs-mahout/vectors/tokenized-documents	100.00%	4	4	100.00%	1	1	NA	NA
job_201510062030_0003	NORMAL	DoubleJ	Dictionary/Vectorizer::MakePartial/Vectors: input-folder: hdfs- mahout/vectors/tokenized-documents, dictionary-file: hdfs- mahout/vectors/dictionary.file-0	100.00%	4	4	100.00%	1	1	NA	NA
job 201510062030 0004	NORMAL	DoubleJ	PartialVectorMerger::MergePartialVectors	100.00%	1	1	100.00%	1	1	NA	NA
job 201510062030 0005	NORMAL	DoubleJ	VectorTfldf Document Frequency Count running over input: hdfs- mahout/vectors/tf-vectors	100.00%	1	1	100.00%	1	1	NA	NA
job 201510062030 0006	NORMAL	DoubleJ	: MakePartialVectors: input-folder: hdfs-mahout/vectors/tf-vectors, dictionary-file: hdfs-mahout/vectors/frequency.file-0	100.00%	1	1	100.00%	1	1	NA	NA
job_201510062030_0007	NORMAL	DoubleJ	PartialVectorMerger::MergePartialVectors	100.00%	1	1	100.00%	1	1	NA	NA
job 201510062030 0008	NORMAL	DoubleJ	Cluster Iterator running iteration 1 over priorPath: hdfs- mahout/kmeans/clusters-0	100.00%	1	1	100.00%	1	1	NA	NA
job 201510062030 0009	NORMAL	DoubleJ	Cluster Iterator running iteration 2 over priorPath: hdfs- mahout/kmeans/clusters-1	100.00%	1	1	100.00%	1	1	NA	NA
job_201510062030_0010	NORMAL	DoubleJ	Cluster Classification Driver running over input: hdfs-mahout/vectors/tfidf-vectors	100.00%	1	1	100.00%	0	0	NA	NA

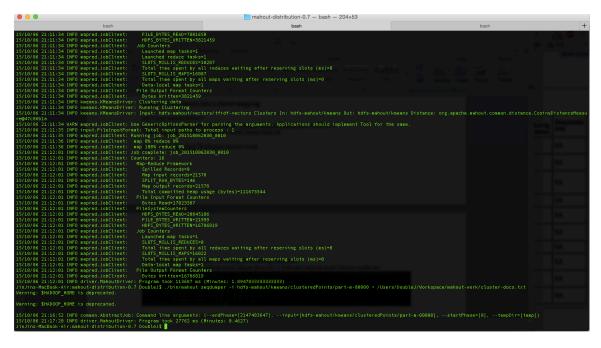
Retired Jobs

8. View results in a human readable way

8.1 Dump the documents into a cluster mapping

\$cd <your mahout folder>

 $./bin/mahout\ seqdumper\ -i\ hdfs-mahout/kmeans/clusteredPoints/part-m-00000 > /Users/DoubleJ/Workspace/mahout-work/cluster-docs.txt$



A new file (cluster-docs.txt) will be generated in your local system: /Users/DoubleJ/Workspace/cluster-docs.txt

\$ls -l /Users/DoubleJ/Workspace/mahout-work/

8.2 Download the perl view.pl and run it using the following command.

The cluster-results.txt is a text file where each line has two columns with a tab separated: clustered and its corresponding file name.