Lab 01: HDFS, MapReduce, Pig, Hive, and Jaql

Hands-On Lab





Table of Contents

1	Intro	duction	3		
2	Abou	About this Lab			
3	Envi	ronment Setup Requirements	3		
	3.1	Getting Started	3		
4	Expl	oring Hadoop Distributed File System (HDFS)	6		
	4.1	Using the command line interface	6		
5	Mapl	Reduce	13		
	5.1	Running the WordCount program	. 13		
6	Work	king with Pig	15		
7	Work	king with Hive	18		
8	Work	king with Jaql	22		
9	Sum	mary	26		

1 Introduction

The overwhelming trend towards digital services, combined with cheap storage, has generated massive amounts of data that enterprises need to effectively gather, process, and analyze. Techniques from the data warehousing and high-performance computing communities are invaluable for many enterprises. However, often times their cost or complexity of scale-up discourages the accumulation of data without an immediate need. As valuable knowledge may nevertheless be buried in this data, related scaled-up technologies have been developed. Examples include Google's MapReduce, and the open-source implementation, Apache Hadoop.

Hadoop is an open-source project administered by the Apache Software Foundation. Hadoop's contributors work for some of the world's biggest technology companies. That diverse, motivated community has produced a collaborative platform for consolidating, combining and understanding data.

Technically, Hadoop consists of two key services: data storage using the Hadoop Distributed File System (HDFS) and large scale parallel data processing using a technique called MapReduce

2 About this Lab

After completing this hands-on lab, you'll be able to:

- Use Hadoop commands to explore the HDFS on the Hadoop system
- Use Hadoop commands to run a sample MapReduce program on the Hadoop system
- Explore Pig, Hive and Jaql

3 Environment Setup Requirements

To complete this lab you will need the following:

- 1. InfoSphere BigInsights Bootcamp VMware® image
- 2. VMware Player 2.x or VMware Workstation 5.x or later

For help on how to obtain these components please follow the instructions specified in VMware Basics and Introduction from module 1.

3.1 Getting Started

To prepare for the contents of this lab, you must go through the process of getting all of the Hadoop components started.

- 1. Start the VMware image by clicking the Power On button in VMware Workstation if it is not already on.
- 2. Log in to the VMware virtual machine using the following information:
 - User: biadmin
 - Password: password

Page 3 of 27



3. Open Gnome Command Prompt Window by right-clicking on the Desktop and selecting "Open in Terminal".

Figure 1 - Open a new terminal window

4. Change to the **\$BIGINSIGHTS_HOME** (which by default is set to /opt/ibm/biginsights).

cd \$BIGINSIGHTS_HOME/bin

or

cd /opt/ibm/biginsights/bin

5. Start the Hadoop components (daemons) on the BigInsights server. You can practice starting all components with these commands. Please note they will take a few minutes to run:

./start-all.sh

The following figure shows the different Hadoop components starting.

	biadmin@imtebi1:01_HadoopCore/HDFS 🔤 🗖 🗖	×
File E	idit View Terminal Help	
biadmi	n@imtebil:~/Desktop> start-all.sh	-
[INFO]	DeployCmdline - [IBM InfoSphere BigInsights QuickStart Edition]	
[INFO]	Progress - Start zookeeper	
[INFO]	@imtebil.imte.com - zookeeper started, pid 5450	
[INFO]	Deployer - zookeeper service started	
[INFO]	Progress - 9%	
[INFO]	Progress - Start hadoop	
[INFO]	@imtebil.imte.com - namenode started, pid 5643	-
[INFO]	@imtebil.imte.com - secondarynamenode started, pid 5890	-
[INFO]	@imtebil.imte.com - datanode started, pid 6059	
[INFO]	Progress - 14%	
[INFO]	Deployer - Waiting for Namenode to exit safe mode	
[INFO]	Deployer - Wait namenode to exit safemode for another 5 seconds, please check namenode log for details	
[INFO]	Deployer - Wait namenode to exit safemode for another 5 seconds, please check namenode log for details	
[INFO]	Deployer - Wait namenode to exit safemode for another 10 seconds, please check namenode log for details	
[INFO]	Deployer - Wait namenode to exit safemode for another 10 seconds, please check namenode log for details	
[INFO]	Deployer - HDFS cluster started successfully	
[INFO]	@imtebil.imte.com - jobtracker started, pid 6773	
[INFO]	@imtebil.imte.com - tasktracker started, pid 7057	
[INFO]	Progress - 18%	
[INFO]	Deployer - MapReduce cluster started successfully	
[INFO]	Progress - Start derby	
[INFO]	@imtebil.imte.com - derby started, pid 7231	
[INFO]	Progress - 27%	
[INFO]	Progress - Start hive	
[INFO]	@imtebil.imte.com - derby already running, pid 7231	
[INFO]	Progress - 28%	
[INFO]	@imtebil.imte.com - hive-web-interface started, pid 7408	
[INFO]	@imtebil.imte.com - hive-server started, pid 7733	
[INFO]	Progress - 36%	
[INFO]	Progress - Start hbase	
[INFO]	Deployer - check zookeeper services, make sure zookeeper service is started before start hbase service	
[INFO]	@imtebil.imte.com - hbase-master(active) started	
[INFO]	@imtebil.imte.com - hbase-regionserver started	
[INFO]	Deployer - hbase service started	
[INFO]	Progress - 45%	
[INFO]	Progress - Start bigsql	
[INFO]	@imtebil.imte.com - bigsql-server started, pid 8538	
[INFO]	Progress - 55%	
[INFO]	Progress - Start oozie	Y
	Figure 2 - Starting Hadoop components	

(1) Note: You may get an error that the server has not started, please be patient as it does take some time for the server to complete start.

Terminal		×
<u>F</u> ile <u>E</u> dit <u>V</u> iew <u>T</u> erminal Ta <u>b</u> s <u>H</u> elp		
<pre>[INF0] Deployer - Starting Hadoop [INF0] @localhost.localdomain - Starting jaqlserver [INF0] @localhost.localdomain - Starting biginsightconsole [INF0] @localhost.localdomain - Starting bigsheets [INF0] Progress - 20%</pre>		
[INF0] @localhost.localdomain - namenode uncertain status, 9000 not open, 27 Retry the daemon command later, or tail log: ssh <node> tail /var/ibm/bigi</node>	pid 34 nsight	141 A
s/hadoop/log/hadoop-hdpadmin-namenode-localhost.log		

Figure 3 - Hadoop component error

6. Sometimes certain hadoop components may fail to start. You can start and stop the failed components one at a time by using **start.sh** or **stop.sh** respectively. For example, to start and stop Hadoop use:

./start.sh hadoop
./stop.sh hadoop

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Page 5 of 27

In the following example, the console component failed. The particular component was then started again using the **./start.sh console** command. It then succeeded without any problems. This approach can be used for any failed components.

🖬 biadmin@imtebi1:bm/biginsights/bin 💶 🗆 🗙
File Edit View Terminal Help
biadmin@imtebil:/opt/ibm/biginsights/bin> ./start.sh console
[INFO] DeployCmdline - [IBM InfoSphere BigInsights QuickStart Edition]
[INFO] Progress - Start console
[INFO] Deployer - /opt/ibm/biginsights/console/wlp
[INFO] Deployer - starting pigserver, logging to /var/ibm/biginsights/sheets/logs/bigsheets-biadmin-pigserve
r.out
[INFO] Deployer - Server waslp-server started with process ID 26854.
[INFO] Deployer - [AUDIT] CWWKF00111: The server waslp-server is ready to run a smarter planet.
[INFO] Deployer - BigInsights Management Console started, pid 26854
[INFO] Progress - 100%
[INFO] DeployManager - Start; SUCCEEDED components: [console]; Consumes : 23226ms
biadmin@imtebil:/opt/ibm/biginsights/bin>
biadmin@imtebil:/opt/ibm/biginsights/bin>

Figure 4 - Starting a specific component

Once all components have started successfully you can then move to the next section.

4 Exploring Hadoop Distributed File System (HDFS)

Hadoop Distributed File System (HDFS), allows user data to be organized in the form of files and directories. It provides a command line interface called *FS shell* that lets a user interact with the data in HDFS accessible to Hadoop MapReduce programs.

There are two methods to interact with HDFS:

- 1. You can use the command-line approach and invoke the FileSystem (fs) shell using the format: hadoop fs <args>. This is the method we will use in this lab..
- 2. You can also manipulate HDFS using the BigInsights Web Console. You will explore the BigInsights Web Console on another lab.

4.1 Using the command line interface

In this part, we will explore some basic HDFS commands. All HDFS commands start with *hadoop* followed by *dfs* (distributed file system) or *fs* (file system) followed by a dash, and the command. Many HDFS commands are similar to UNIX commands. For details, refer to the *Hadoop Command Guide* and *Hadoop FS Shell Guide*.

We will start with the hadoop fs -Is command which returns the list of files and directories with permission information.

Ensure the Hadoop components are all started, and from the same Gnome terminal window as before (and logged on as *biadmin*), follow these instructions:

1. List the contents of the root directory.

```
hadoop fs -ls /
```

	biadmin@imtebi1:bm/biginsights/bin	_ = ×
File Edit View Terminal Help	THE REPORT OF MALE AND THE PROPERTY AND THE CO	
biadmin@imtebil:/opt/ibm/biginsight	s/bin> hadoop fs -ls /	^
Found 5 items		
drwxr-xr-x - biadmin biadmgrp	0 2013-06-19 22:56 /biginsights	
drwxr-xr-x - biadmin supergroup	0 2013-07-09 13:48 /hadoop	
drwxr-xr-x - biadmin supergroup	0 2013-07-09 13:50 /hbase	
drwxrwxrwx - biadmin supergroup	0 2013-06-19 22:45 /tmp	
drwxrwxrwx - biadmin supergroup	0 2013-07-09 13:33 /user	
biadmin@imtebil:/opt/ibm/biginsight	s/bin>	

Figure 5 - List directory command

2. To list the contents of the /user/biadmin directory, execute:

hadoop fs -ls or

```
hadoop fs -ls /user/biadmin
```

Note that in the first command there was no directory referenced, but it is equivalent to the second command where /user/biadmin is explicitly specified. Each user will get its own home directory under /user. For example, in the case of user biadmin, his home directory is /user/biadmin. Any command where there is no explicit directory specified will be relative to the user's home directory.

			biadmir	n@imtebi1:	bm/biginsights	/bin		×
File Edit Vi	ew	Terminal	Help					
biadmin@imt	ebi	1:/opt/il	om/biginsigh	ts/bin> ha	doop fs -ls	1		~
Found 5 ite	ms	Ċ6			10			
drwxr-xr-x	14	biadmin	biadmgrp	0	2013-07-09	13:52	/biginsights	
drwxr-xr-x	12	biadmin	supergroup	0	2013-07-11	11:04	/hadoop	
drwxr-xr-x	8	biadmin	supergroup	0	2013-07-11	11:06	/hbase	
drwxrwxrwx	12	biadmin	supergroup	0	2013-06-19	22:45	/tmp	
drwxrwxrwx	10	biadmin	supergroup	0	2013-07-09	16:47	/user	
biadmin@imt	ebi	1:/opt/il	om/biginsigh	ts/bin> ha	doop fs -ls			
Found 3 ite	ms		Constantine for					
drwx	10	biadmin	supergroup	0	2013-07-05	10:31	/user/biadmin/.staging	
drwxxx	14	biadmin	supergroup	0	2013-07-09	13:56	/user/biadmin/credstore	
drwxr-xr-x	12	biadmin	supergroup	0	2013-07-05	10:31	/user/biadmin/oozie-biad	=
biadmin@imt	ebi	1:/opt/il	om/biginsigh	ts/bin>				-

Figure 6 - hadoop fs -ls command outputs

3. To create the directory *myTestDir* you can issue the following command:

```
hadoop fs -mkdir myTestDir
```

Where was this directory created? As mentioned in the previous step, any relative paths will be using the user's home directory.

4. Issue the Is command again to see the subdirectory myTestDir:

hadoop	fs	-ls
--------	----	-----

or

```
hadoop fs -ls /user/biadmin
```

biadmin@imtebi1:bm/biginsights/bin	_ 0	×
File Edit View Terminal Help		
biadmin@imtebil:/opt/ibm/biginsights/bin> hadoop fs -ls Found 4 items		^
drwx biadmin supergroup 0 2013-07-05 10:31 /user/biadmin/.stag	ing	
drwxxx - biadmin supergroup 0 2013-07-09 13:56 /user/biadmin/creds	tore	
rwxr-xr-x - biadmin supergroup 0 2013-07-26 02:23 /user/biadmin/myTes	tDır	
drwxr-xr-x - biadmin supergroup 0 2013-07-05 10:31 /user/biadmin/oozie	-biad	
biadmin@imtebil:/opt/ibm/biginsights/bin> hadoop fs -ls /user/biadmin		
Found 4 items		
drwx biadmin supergroup 0 2013-07-05 10:31 /user/biadmin/.stag	ing	
drwxxx - biadmin supergroup 0 2013-07-09 13:56 /user/biadmin/creds	tore	
drwxr-xr-x - biadmin supergroup 0 2013-07-26 02:23 /user/biadmin/myles	tDir	
drwxr-xr-x - biadmin supergroup 0 2013-07-05 10:31 /user/biadmin/oozie	-biad	Ξ
biadmin@imtebil:/opt/ibm/biginsights/bin>		9

Figure 7 - hadoop fs -ls command outputs

(i) Note: If you specify a relative path to hadoop fs commands, they will implicitly be relative to your user directory in HDFS. For example when you created the directory myTestDir, it was created in the */user/biadmin* directory.

To use HDFS commands recursively generally you add an "r" to the HDFS command (In the Linux shell this is generally done with the "-R" argument).

5. For example, to do a recursive listing we'll use the -lsr command rather than just -ls, like the examples below:

hadoop fs -ls /user hadoop fs -lsr /user

			biad	min@imte	bi1:0	1_Hado	opCore	/HDFS _ 🗆	×
File Edit	View	Terminal	Help						
bi admi n@i	mtehi	1.∼/hoot	camp/ipput/lab	01 Hadoor	Core	HDES>	hadoor	o fs als /user	^
Found 2 i	tome			- '					
drwxrwxrv	/x -	biadmin	supergroup	0	2013-	07-05	10:33	/user/applications	
drwxr-xr-	x -	biadmin	supergroup	0	2013-	07-22	13:10	/user/biadmin	
bi admi n@i	mtabi	1.~/hoot	amp/input/lab	al Hadoor	Core		hadoor	fe ler /user	
druwnuwn		biodmin	eupongnoup		2012	07 05	10.22	/user/enplications	
drwxr-xr-	x -	biadmin	supergroup	0	2013-	07-22	13:10	/user/biadmin	
drwx		biadmin	supergroup	0	2013-	07-05	10:31	/user/biadmin/.staging	
drwx		biadmin	superaroup	0	2013-	07-05	10:29	/user/biadmin/.staging/job 201307051019	
0002			1 5 1						-
drwxx	x -	biadmin	supergroup	0	2013-	07-09	13:56	/user/biadmin/credstore	
drwx		biadmin	supergroup	0	2013-	06-19	22:46	/user/biadmin/credstore/private	
- rw	- 3	biadmin	supergroup	99	2013-	06-19	22:46	/user/biadmin/credstore/private/hive key	v
store pwo	.prop		1 5 1						
- rw- r r -	- 1	biadmin	supergroup	109	2013-	07-09	13:56	/user/biadmin/credstore/properties.txt	
drwxr-xr-	x -	biadmin	superaroup	0	2013-	07-22	11:51	/user/biadmin/mvTestDir	
drwxr-xr-	x -	biadmin	supergroup	0	2013-	07-05	10:31	/user/biadmin/oozie-biad	
biadmin@i	mtebi	1:~/boot	camp/input/lab	01 Hadoor	Core	HDES>			Ξ
		,							~

Figure 8 - hadoop fs-lsr command output

6. You can pipe (using the | character) any HDFS command to be used with the Linux shell. For example, you can easily use *grep* with HDFS by doing the following:

hadoop fs -mkdir /user/biadmin/myTestDir2
hadoop fs -ls /user/biadmin | grep Test

biadmin@imtebi1:bm/biginsights/bin	_ = ×
File Edit View Terminal Help	
biadmin@imtebil:/opt/ibm/biginsights/bin> hadoop fs -mkdir /user/biadmin/myTestDir2 biadmin@imtebil:/opt/ibm/biginsights/bin> hadoop fs -ls /user/biadmin grep Test	^
drwxr-xr-x - biadmin supergroup 0 2013-07-09 14:01 /user/biadmin/myTestDir	
drwxr-xr-x - biadmin supergroup 0 2013-07-09 14:21 /user/biadmin/myTestDir2 biadmin@imtebil:/opt/ibm/biginsights/bin>	



As you can see the grep command only returned the lines which had test in them (thus removing the "Found x items" line and the .staging and oozie-biad directories from the listing

7. To move files between your regular Linux filesystem and HDFS you can use the put and get commands. For example, move the text file README to the hadoop filesystem.

hadoop fs -put /home/biadmin/bootcamp/input/lab01_HadoopCore/HDFS/README README hadoop fs -ls /user/biadmin

	biadmin@imtebi1:01_HadoopCore/HDFS	×
File Edit ∨iew	Terminal Help	
biadmin@imteb:	l:~/bootcamp/input/lab01_HadoopCore/HDFS> hadoop fs -ls	^
Found 6 items		
drwx	biadmin supergroup 0 2013-07-05 10:31 /user/biadmin/.staging	
	biadmin supergroup 10 2010 07 22 10:17 /user/biadmin/PEADME	
rwxxx	biadmin supergroup 0 2013-07-09 13-56 (user/biadmin/credstore	
drwxr-xr-x	biadmin supergroup 0 2013-07-22 11:51 /user/biadmin/myTestDir	
drwxr-xr-x	biadmin supergroup 0 2013-07-22 13:13 /user/biadmin/myTestDir2	
drwxr-xr-x	biadmin supergroup 0 2013-07-05 10:31 /user/biadmin/oozie-biad	=
biadmin@imteb:	l:~/bootcamp/input/lab01_HadoopCore/HDFS>	~

Figure 10 - README file inside HDFS

You should now see a new file called /user/biadmin/README listed as shown above. Note there is a '1' highlighted in the figure. This represents the replication factor. By default, the replication factor in a BigInsights cluster is 3, but since this laboratory environment only has one node, the replication factor is 1.

8. In order to view the contents of this file use the –cat command as follows:

hadoop fs -cat README

You should see the output of the README file (that is stored in HDFS). We can also use the linux diff command to see if the file we put on HDFS is actually the same as the original on the local filesystem.

9. Execute the commands below to use the diff command:

cd /home/biadmin/bootcamp/input/lab01 HadoopCore/HDFS/

diff <(hadoop fs -cat README) README

Since the diff command produces no output we know that the files are the same (the diff command prints all the lines in the files that differ).

To find the size of files you need to use the -du or -dus commands. Keep in mind that these commands return the file size in bytes.

10. To find the size of the README file use the following command:

hadoop fs -du README



Figure 11 - Inspecting README file size

In this example, the README file has 18 bytes.

11. To find the size of all files individually in the /user/biadmin directory use the following command:

hadoop fs -du /user/biadmin

hadoop fs -dus /user/biadmin

	biadmin@imtebi1:01_HadoopCore/HDFS	_ 🗆 X
File	Edit ∨iew Terminal Help	
biadm	in@imtebil:~/bootcamp/input/lab01_HadoopCore/HDFS> hadoop fs -du /user/biadmin	<u>^</u>
Found	6 items	
0	hdfs://imtebil.imte.com:9000/user/biadmin/.staging	
18	hdfs://imtebil.imte.com:9000/user/biadmin/README	
208	hdfs://imtebil.imte.com:9000/user/biadmin/credstore	
0	hdfs://imtebil.imte.com:9000/user/biadmin/myTestDir	
0	hdfs://imtebil.imte.com:9000/user/biadmin/myTestDir2	
0	hdfs://imtebil.imte.com:9000/user/biadmin/oozie-biad	
biadm:	in@imtebil:~/bootcamp/input/lab01_HadoopCore/HDFS>	
	_	

Figure 12 - Inspecting files size in a specific directory

12. To find the size of all files in total of the /user/biadmin directory use the following command:

 biadmin@imtebi1:...01_HadoopCore/HDFS
 Image: Core/HDFS

 File Edit View Terminal Help

 biadmin@imtebi1:~/bootcamp/input/lab01_HadoopCore/HDFS> hadoop fs -dus /user/biadmin

 hdfs://imtebi1.imte.com:9000/user/biadmin
 226

 biadmin@imtebi1:~/bootcamp/input/lab01_HadoopCore/HDFS>
 Image: Core/HDFS>

Figure 13 - Inspecting the size of directories

13. If you would like to get more information about hadoop fs commands, invoke -help as follows:

hadoop fs -help



Figure 14 - Hadoop help command

14. For specific help on a command, add the command name after help. For example, to get help on the dus command you'd do the following:





Figure 15 - Help for specific Haoop commands

5 MapReduce

Now that we've seen how the FileSystem (fs) shell can be used to execute Hadoop commands to interact with HDFS, the same *fs* shell can be used to launch MapReduce jobs. In this section, we will walk through the steps required to run a MapReduce program. The source code for a MapReduce program is contained in a compiled .jar file. Hadoop will load the JAR into HDFS and distribute it to the data nodes, where the individual tasks of the MapReduce job will be executed. Hadoop ships with some example MapReduce programs to run. One of these is a distributed WordCount program which reads text files and counts how often words occur.

5.1 Running the WordCount program

First we need to copy the data files from the local file system to HDFS.

1. Execute the commands below to copy the input files into HDFS.

hadoop fs -mkdir /user/biadmin/input

hadoop fs -put /home/biadmin/bootcamp/input/lab01_HadoopCore/MapReduce/*.csv
/user/biadmin/input

🖻 biadmin@	imtebi1:01_HadoopCore/HDFS _ 🗆 ×
File Edit View Terminal Help	
biadmin@imtebil:~/bootcamp/input/lab01_Hadoo biadmin@imtebil:~/bootcamp/input/lab01_Hadoo doopCore/MapReduce/*.csv /user/biadmin/inpu biadmin@imtebil:~/bootcamp/input/lab01_Hadoo	ppCore/HDFS> hadoop fs -mkdir /user/biadmin/input ppCore/HDFS> hadoop fs -put /home/biadmin/bootcamp/input/lab01_Ha c ppCore/HDFS>

Figure 16 - Copy input files into HDFS

2. Review the files have been copied with the following command:

```
hadoop fs -ls input
```



Figure 17 - List copied files into HDFS

3. Now we can run the wordcount job with the command below, where "/user/biadmin/input/" is where the input files are, and "output" is the directory where the output of the job will be stored. The "output" directory will be created automatically when executing the command below.

hadoop jar /opt/ibm/biginsights/IHC/hadoop-examples-1.1.1.jar wordcount /user/biadmin/input/ output

	biadm	nin@imtebi1:01_HadoopCore/HDFS	_ 🗆 🗙
File Edit	View Terminal Help		
biadmin@	imtebil:~/bootcamp/input/lab01 H	adoopCore/HDFS> hadoop jar /opt/ibm/biginsights/IHC/hadoop-	examples-
1.1.1.ja	r wordcount /user/biadmin/input/	output	22
13/07/09	14:40:04 INFO input.FileInputFo	rmat: Total input paths to process : 3	
13/07/09	14:40:04 INFO mapred.JobClient:	Running job: job_201307091355_0001	
13/07/09	14:40:05 INFO mapred.JobClient:	map 0% reduce 0%	
13/07/09	14:40:13 INFO mapred.JobClient:	map 33% reduce 0%	
13/07/09	14:40:15 INFO mapred.JobClient:	map 66% reduce 0%	
13/07/09	14:40:19 INFO mapred.JobClient:	map 100% reduce 0%	
13/07/09	14:40:26 INFO mapred.JobClient:	map 100% reduce 100%	
13/07/09	14:40:27 INFO mapred.JobClient:	Job complete: job_201307091355_0001	
13/07/09	14:40:27 INFO mapred.JobClient:	Counters: 29	
13/07/09	14:40:27 INFO mapred.JobClient:	Job Counters	
13/07/09	14:40:27 INFO mapred.JobClient:	Data-local map tasks=3	
13/07/09	14:40:27 INFO mapred.JobClient:	SLOTS_MILLIS_MAPS=12454	
13/07/09	14:40:27 INFO mapred.JobClient:	Launched map tasks=3	
13/07/09 ms)=0	14:40:27 INFO mapred.JobClient:	Total time spent by all reduces waiting after reservin	g slots (
13/07/09	14:40:27 INFO mapred.JobClient:	Total time spent by all maps waiting after reserving s	lots (ms)
=0			
13/07/09	14:40:27 IN=0 mapred.JobClient:	Launched reduce tasks=1	
13/07/09	14:40:27 INFO mapred.JobClient:	SLOIS_MILLIS_REDUCES=9452	
13/07/09	14:40:27 IN=0 mapred.JobClient:	File input Format Counters	
13/07/09	14:40:27 INFO mapred.JobClient:	Bytes Read=10954	
13/07/09	14:40:27 IN=0 mapred.JobClient:	File Output Format Counters	
13/07/09	14:40:27 INFO mapred.JobClient:	Bytes Written=9146	
13/07/09	14:40:27 INFO mapred.JobClient:	FILESystemcounters	
13/07/09	14:40:27 INFO mapred.JobClient:	HUFS_BITES_REAU=11480	
13/07/09	14:40:27 INFO mapred.JobClient:	FILE_BITES_WRITTEN=123009	
13/07/09	14:40:27 INFO mapred.JobClient:	HDEC DYTES WOTTEN-0146	
13/07/09	14:40:27 INFO mapred.JobClient:	HDFS_BTTES_WRITTEN=9140	
13/07/09	14:40:27 INFO mapred JobClient:	Vintual memory (butes) energhet=4761505004	
12/07/09	14:40:27 INFO mapred JobClient:	Peduce input appune=207	
12/07/09	14:40:27 INFO mapred JobClient:	Combine output pecende=200	=
12/07/09	14:40:27 INFO mapred lobClient:	Man output records=200	
13/07/09	14:40:27 INFO mapred JobClient.	CPU time spent (ms)=2390	
10,01,03	THE OWNER THE O HUP CONSODUCTION.	are crue sperie (mat-zase	

Figure 18 - WordCount MapReduce job running

4. Now review the output of step 3:

```
hadoop fs -ls output
```

	biadmin@imtebi1:01_HadoopCore/HDFS	_ = ×
File Edit View Terminal Help		
biadmin@imtebil:~/bootcamp/input/lab Found 3 items	01_HadoopCore/HDFS> hadoop fs -ls output	^
-rw-rr l biadmin supergroup drwxr-xr-x - biadmin supergroup -rw-rr l biadmin supergroup biadmin@imtebil:~/bootcamp/input/lab	0 2013-07-09 14:40 /user/biadmin/output/_SUCCESS 0 2013-07-09 14:40 /user/biadmin/output/_logs 9146 2013-07-09 14:40 /user/biadmin/output/part-r-00000 01_HadoopCore/HDFS>	

Figure 19 - MapReduce result files

In this case, the output was not split into multiple files.

5. To view the contents of the part-r-0000 file issue the command below:





Figure 20 - MapReduce output

(i) Note: You can use the BigInsights Web Console to run applications such as WordCount. This same application (though with different Input files) will be run again in the lab describing the BigInsights Web Console. More detail about the job will also be described then.

6 Working with Pig

In this tutorial, we are going to use Apache Pig to process the 1988 subset of the Google Books 1-gram records to produce a histogram of the frequencies of words of each length. A subset of this database (0.5 million records) has been stored in the file **googlebooks-1988.csv** under */home/biadmin/bootcamp/input/lab01_HadoopCore/PigHiveJaql* directory.

Let us examine the format of the Google Books 1-gram records.

1. Execute the commands below to examine the format of the records:

```
cd /home/biadmin/bootcamp/input/lab01 HadoopCore/PigHiveJaql
```

```
head -5 googlebooks-1988.csv
```

				biadmin@imtebi1:opCore/PigHiveJaql	_ = ×	
File Edi	t ∨iew	Terminal	Help			
biadmin(@imtebil	l:~/booto	amp/inpu	t/lab01_HadoopCore/HDFS> cd /home/biadmin/bootcamp/input/lab01_HadoopCo	re/Pig 🔨	
HiveJaq	/					
biadmin(@imtebi]	l:~/booto	amp/inpu	ıt/lab01_HadoopCore/PigHiveJaql> head -5 googlebooks-1988.csv		
#	1988	94000	45770	9585		
\$0.000	1988	4	4	2		
\$0.0006	1988	3	3	3		
\$0.0027	1988	1	1	1		
\$0.003	1988	4	4	2		
biadmin(jiadmin@imtebil:~/bootcamp/input/lab01_HadoopCore/PigHiveJaql>					

Figure 21 – Googlebooks-1988.csv file

The columns these data represent are the word, the year, the number of occurrences of that word in the corpus, the number of pages on which that word appeared, and the number of books in which that word appeared.

2. Copy the data file into HDFS.

hadoop fs -put googlebooks-1988.csv pighivejaql/googlebooks-1988.csv

Note that directory /user/biadmin/pighivejaql is created automatically for you when the above command is executed.

3. Start pig. If it has not been added to the PATH, you can add it, or switch to the \$PIG_HOME/bin directory

cd \$PIG HOME/bin

./pig

🗉 biadmin@ii	ntebi1:iginsights/pig/bin _ 🗆 🛪
File Edit View Terminal Help	
biadmin@imtebil:~/bootcamp/input/labOl_HadoopCo biadmin@imtebil:/opt/ibm/biginsights/pig/bin> . grunt>	re/PigHiveJaql> cd \$PIG_HOME/bin ∕pig

Figure 22 – Pig command line

4. We are going to use a Pig UDF to compute the absolute value of each integer. The UDF is located inside the piggybank.jar file (This jar file was created from the source, following the instructions in https://cwiki.apache.org/confluence/display/PIG/PiggyBank, and copied to the piggybank directory). We use the REGISTER command to load this jar file:

REGISTER /opt/ibm/biginsights/pig/contrib/piggybank/java/piggybank.jar;

The first step in processing the data is to LOAD it.

5. Execute the step below to load data.

```
records = LOAD 'pighivejaql/googlebooks-1988.csv' AS (word:chararray,
year:int, wordcount:int, pagecount:int, bookcount:int);
```

This returns instantly. The processing is delayed until the data needs to be reported.

6. To produce a histogram, we want to group by the length of the word:

```
grouped = GROUP records by
org.apache.pig.piggybank.evaluation.string.LENGTH(word);
```

7. Sum the word counts for each word length using the SUM function with the FOREACH GENERATE command.

```
final = FOREACH grouped GENERATE group, SUM(records.wordcount);
```

8. Use the DUMP command to print the result to the console. This will cause all the previous steps to be executed.

DUMP final;

This should produce output like the following:

	biadmin@imtebi1:iginsights/pig/bin	
File Edit View Terminal H	leip	
grunt> grouped = GROUP r grunt> final = FOREACH g grunt> DLMP final:	records by org.apache.pig.piggybank.evaluation.s grouped GENERATE group, SUM(records.wordcount);	string.LENGTH(word);
INFO [Thread-17] org.ap	pache.hadoop.mapreduce.lib.input.FileInputFormat	t - Total input paths to process :
INFO [main] org.apache. (1,12225611)	.hadoop.mapreduce.lib.input.FileInputFormat	- Total input paths to process : 1
(2,222545981)		
(4,20200542)		
(5, 20412451)		
(6 24985568)		
(7 28833898)		
(8,19880788)		
(9,16946072)		
(10,10711870)		
(11,7580500)		
(12,3472760)		
(13,1876286)		
(14,1275906)		
(15,502004)		
(16,181439)		
(17,80973)		
(18,22877)		
(19,15419)		
(20,13832)		
(21,6765)		
(22,2125)		
(23,1179)		
(24,1435)		
(25,290)		
(26,130)		
(27,118)		-
(28,28)		=
(29,61)		
(30,125)		~

Figure 23 - Wordcount application output

9. Quit pig.

grunt> quit

7 Working with Hive

In this tutorial, we are going to use Hive to process the 1988 subset of the Google Books 1-gram records to produce a histogram of the frequencies of words of each length. A subset of this database (0.5 million records) has been stored in the file googlebooks-1988.csv under */home/biadmin/bootcamp/input/lab01_HadoopCore/PigHiveJaql* directory.

1. Ensure the Apache Derby component is started. Apache Derby is the default database used as metastore in Hive. A quick way to verify if it is started, is to try to start it using:

start.sh derby

 Biadmin@imtebil:...bm/biginsights/bin

 File Edit View Terminal Help

 biadmin@imtebil:/opt/ibm/biginsights/bin> ./start.sh derby

 [INFO] DeployCmdline - [IBM InfoSphere BigInsights QuickStart Edition]

 [INFO] Progress - Start derby

 [INFO] Ogimtebil.imte.com - derby started, pid 5770

 [INFO] DeployManager - Start; SUCCEEDED components: [derby]; Consumes : 6179ms

 biadmin@imtebil:/opt/ibm/biginsights/bin>

Figure 24 - Start Apache Derby

2. Start hive interactively. Change the directory to the \$HIVE_HOME/bin first, and execute from there using ./hive

cd \$HIVE_HOME/bin ./hive

biadmin@imtebi1:ginsights/hive/bin	_		×
File Edit View Terminal Help			
biadmin@imtebil:/opt/ibm/biginsights/bin> cd \$HIVE_HOME/bin biadmin@imtebil:/opt/ibm/biginsights/hive/bin> ./hive Tue Jul 09 14:50:49 EDT 2013 : Connection obtained for host: imtebil.imte.com, port number 152 Logging initialized using configuration in file:/opt/ibm/biginsights/hive/conf/hive-log4j.prop Hive history file=/var/ibm/biginsights/hive/query/biadmin/hive_job_log_biadmin_201307091450_61 hive> ■	28. perties 14262047.txt	:	<

Figure 25 - Start Apache Hive

3. Create a table called wordlist.

CREATE TABLE wordlist (word STRING, year INT, wordcount INT, pagecount INT, bookcount INT) ROW FORMAT DELIMITED FIELDS TERMINATED BY '\t';

biadmin@imtebi1:ginsights/hive/bin	_		×
File Edit View Terminal Help			
biadmin@imtebil:/opt/ibm/biginsights/bin> cd \$HIVE_HOME/bin biadmin@imtebil:/opt/ibm/biginsights/hive/bin> ./hive Tue Jul 09 14:50:49 EDT 2013 : Connection obtained for host: imtebil.imte.com, port number 1528. Logging initialized using configuration in file:/opt/ibm/biginsights/hive/conf/hive-log4j.properties Hive history file=/var/ibm/biginsights/hive/query/biadmin/hive_job_log_biadmin_201307091450_61426204 hive> CREATE TABLE wordlist (word STRING, year INT, wordcount INT, pagecount INT, bookcount INT) ROW DELIMITED FIELDS TERMINATED BY '\t'; OK Time taken: 6.391 seconds hive>	7.txt FORM	IAT	



4. Load the data from the googlebooks-1988.csv file into the wordlist table.

LOAD DATA LOCAL INPATH

```
'/home/biadmin/bootcamp/input/lab01_HadoopCore/PigHiveJaql/googlebooks-
1988.csv' OVERWRITE INTO TABLE wordlist;
```

	biadmin@imtebi1:ginsights/hive/bin	- 🗆 X
File Edit View Terminal	Help	
hive> LOAD DATA LOCAL ' OVERWRITE INTO TABLE	<pre>INPATH '/home/biadmin/bootcamp/input/lab01_HadoopCore/PigHiveJaql/googlebooks-190 wordlist;</pre>	88.csv 🛆
Copying data from file	:/home/biadmin/bootcamp/input/lab01_HadoopCore/PigHiveJaql/googlebooks-1988.csv	
Copying file: file:/ho	me/biadmin/bootcamp/input/lab01_HadoopCore/PigHiveJaql/googlebooks-1988.csv	
Loading data to table	default.wordlist	
OK	inte.com.sodo/biginsignts/nive/warenouse/wordtist	
Time taken: 1.011 seco	Inds	
hive>		
- Contraction of the second seco		

Figure 27 - Load data into wordcount table

5. Create a table named wordlengths to store the counts for each word length for our histogram.

CREATE TABLE wordlengths (wordlength INT, wordcount INT);



Figure 28 - Create wordlength table

6. Fill the wordlengths table with word length data from the wordlist table calculated with the length function.

INSERT OVERWRITE TABLE wordlengths SELECT length(word), wordcount FROM wordlist;

🗧 biadmin@imtebi1:ginsights/hive/bin _ 🗆 🗙
File Edit View Terminal Help
hive> INSERT OVERWRITE TABLE wordlengths SELECT length(word), wordcount FROM wordlist;
Total MapReduce jobs = 2
Launching Job 1 out of 2
Number of reduce tasks is set to 0 since there's no reduce operator
Starting Job = job_201307091355_0003, Tracking URL = http://imtebi1.imte.com:50030/jobdetails.jsp?jobid=job_ 201307091355_0003
Kill Command = /opt/ibm/biginsights/IHC/libexec//bin/hadoop job -Dmapred.job.tracker=imtebi1.imte.com:900 1 -kill job 201307091355 0003
Hadoop job information for Stage-1: number of mappers: 1; number of reducers: 0
2013-07-09 14:55:07,460 Stage-1 map = 0%, reduce = 0%
2013-07-09 14:55:12,507 Stage-1 map = 100%, reduce = 0%, Cumulative CPU 2.04 sec
2013-07-09 14:55:14,781 Stage-1 map = 100%, reduce = 0%, Cumulative CPU 2.04 sec
2013-07-09 14:55:15,993 Stage-1 map = 100%, reduce = 100%, Cumulative CPU 2.04 sec
MapReduce Total cumulative CPU time: 2 seconds 40 msec
Ended Job = job_201307091355_0003
Ended Job = -420676167, job is filtered out (removed at runtime).
Moving data to: hdfs://imtebil.imte.com:9000/tmp/hive-biadmin/hive_2013-07-09_14-54-58_095_21059989715052616 47/-ext-10000
Loading data to table default.wordlengths
Deleted hdfs://imtebil.imte.com:9000/biginsights/hive/warehouse/wordlengths
Table default.wordlengths stats: [num_partitions: 0, num_files: 1, num_rows: 0, total_size: 2109626, raw_dat a size: 0]
454574 Rows loaded to wordlengths
MapReduce Jobs Launched:
Job 0: Map: 1 Cumulative CPU: 2.04 sec HDFS Read: 9466046 HDFS Write: 2109626 SUCCESS
Total MapReduce CPU Time Spent: 2 seconds 40 msec
OK
Time taken: 18.594 seconds
hive>

Figure 29 - Fill wordlengths table

7. Produce the histogram by summing the word counts grouped by word length.

SELECT wordlength, sum(wordcount) FROM wordlengths group by wordlength;



Figure 30 - Executing MapReduce job

8. Quit hive.

quit;

	biadmin@imtebi1:ginsights/hive/bin	_ 🗆 ×
File Edit ∨iew Terminal	Help	
Time taken: 22.421 sec hive> quit;	onds	▲
biadmin@imtebil:/opt/i	bm/biginsights/hive/bin>	$\overline{\checkmark}$



8 Working with Jaql

In this tutorial, we are going to use Jaql to process the 1988 subset of the Google Books 1-gram records to produce a histogram of the frequencies of words of each length. A subset of this database (0.5 million records) has been stored in the file googlebooks-1988.csv under */home/biadmin/bootcamp/input/lab01_HadoopCore/PigHiveJaql* directory.

1. Let us examine the format of the Google Books 1-gram records:

cd /home/biadmin/bootcamp/input/lab01 HadoopCore/PigHiveJaql





Figure 32 - googlebooks-1998.csv file format

The columns these data represent are the word, the year, the number of occurrences of that word in the corpus, the number of pages on which that word appeared, and the number of books in which that word appeared.

2. Copy the googlebooks-1988.del file to HDFS.

```
hadoop fs -put googlebooks-1988.del googlebooks-1988.del
```



Figure 33 - Copy googlebooks-1988.csv file to HDFS

3. Change directory to \$JAQL_HOME\bin, and then execute ./jaqlshell to start the JaqlShell.

cd \$JAQL_HOME/bin ./jaqlshell



Figure 34 - Start Jaqlsell

4. Read the comma delimited file from HDFS. Note that this operation might take a few minutes to complete.

```
$wordlist = read(del("googlebooks-1988.del", { schema: schema { word:
string, year: long, wordcount: long, pagecount: long, bookcount: long } }));
```

```
      biadmin@imtebi1:...ginsights/jaql/bin
      _ _ _ ×

      File Edit View Terminal Help
      _ _ _ ×

      jaql> $wordlist = read(del("googlebooks-1988.del", { schema: schema { word: string, year: long, wordcount: long, page count: long, bookcount: long } ));
```

Figure 35 - Read googlebooks-1988.del from HDFS

5. Transform each word into its length by applying the strLen function.

```
$wordlengths = $wordlist -> transform { wordlength: strLen($.word),
wordcount: $.wordcount };
```

2	biadmin@imtebi1:ginsights/jaql/bin	_ 🗆 X
File	Edit View Terminal Help	
jaql>	• \$wordlengths = \$wordlist -> transform { wordlength: strLen(\$.word), wordcount: \$.wordcount };	<u>^</u>
jaql>		

Figure 36 - Applying strLen() function

6. Produce the histogram by summing the word counts grouped by word length.

```
$wordlengths -> group by $word = {$.wordlength} into { $word.wordlength,
counts: sum($[*].wordcount) };
```

This should produce output like the following:





7. Quit Jaql.

quit;





9 Summary

You have just completed Lab 1 which focused on the basics of the Hadoop platform, including HDFS, MapReduce, Pig, Hive, and Jaql. You should now know how to perform the following basic tasks on the platform:

- Start/Stop the Hadoop components
- Interact with the data in the Hadoop Distributed File System (HDFS)
- Navigate within HDFS
- Run MapReduce programs
- Use Pig, Hive, and Jaql languages to interact with Hadoop



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