# TDDE31/732A54-Big Data Analytics Lab Compendium

Notice: Please make sure you have read the whole lab compendium before you start to work on the server

from NSC.

April 5th, 2023

### 1 Description and Aim

In the lab you will work on the Sigma<sup>1</sup> set up which is a HPC cluster from the National Supercomputer Centre (NSC). You are supposed to work with the historical meteorological data from the Swedish Meteorological and Hydrological Institute (SMHI). Specifically, you will work with air temperature readings and precipitation readings from 812 meteorological stations in Sweden<sup>2</sup>. In these exercises, you will work with Spark 2.4.3<sup>3</sup>.

There are three lab topics which are BDA1-Spark, BDA2-Spark SQL and BDA3-Machine Learning with Spark. After completing the first two labs you will have basic knowledge of the programming environment, and PySpark API. You will work on exercises with Spark and Spark SQL and thus will be able to compare the differences between the two approaches. In the third lab, you are supposed to achieve a machine learning method with Spark. The overview of three labs is that you need to upload data to Hadoop Distributed File System (HDFS), read the data from HDFS in your code and then program with PySpark<sup>4</sup> to solve the assignments in each lab.

#### 2 SMHI Data

The data includes air temperature and precipitation readings from 812 stations in Sweden. The stations include both currently active stations and historical stations that have been closed down. The latest readings available for active stations are from October 10, 2016. The air temperature and precipitation records are hourly readings, however some stations provide only one reading every three hours.

The provided files<sup>5</sup> are prepared csv files with removed headers. Values are separated with semicolons. Some files are too big to be read using some text editors. Therefore, please use

<sup>&</sup>lt;sup>1</sup>Sigma at: https://www.nsc.liu.se/systems/sigma/

<sup>&</sup>lt;sup>2</sup>If interested in other readings please check: http://opendata-catalog.smhi.se/explore/

<sup>&</sup>lt;sup>3</sup>Spark 2.4.3 at: https://spark.apache.org/docs/2.4.3/index.html

<sup>&</sup>lt;sup>4</sup>PySpark 2.4.3 at: https://spark.apache.org/docs/2.4.3/api/python/index.html

<sup>&</sup>lt;sup>5</sup>Within the BDA\_demo: https://www.ida.liu.se/~732A54/lab/data.zip

either python to read the files or bash commands such as *tail* and *more* to get an overview of a file's content. Provided files:

- temperature-readings.csv ca 2 GB
- precipitation-readings.csv ca 660 MB
- stations.csv
- stations-Ostergotland.csv

All these files are available at /software/sse/manual/spark/BDA\_demo/input\_data folder on Sigma. The headers of these files are shown in Table 1, Table 2 and Table 3. If you notice any mistakes in the dataset or have any comments please contact the lab assistants.

Station number	Date	Time	Air temperature	(in °C)	Quality <sup>a</sup>
----------------	------	------	-----------------	---------	----------------------

<sup>a</sup> G - controlled and confirmed values, Y - suspected or aggregated values

Table 1: Headers for temperature-readings.csv

<sup>&</sup>lt;sup>a</sup> G - controlled and confirmed values, Y - suspected or aggregated values

Table 2: Headers for precipitation-readings.csv

Station	Station	Measurement	Latitude	Longitude	Readings from	Readings to	Elevation
number	name	height			(date and time)	(date and time)	

Table 3: Headers for stations.csv and stations-Ostergotland.csv

# 3 Running PySpark Program on Sigma

#### 3.1 Working on Sigma

The Sigma server is available at **sigma.nsc.liu.se** (log in using your NSC accounts). You can use **(Option 1:)** ssh forwarding connection or **(Option 2:)** Thinlinc<sup>6</sup> connection to log in Sigma. It's also fine to use regular ssh connection without forwarding option, in this case, you have to program locally and use *scp* command to upload your code to Sigma, and then check the history log from text file. **Note:** The forwarding function of ssh supports to run graphics applications (e.g., Atom editor) remotely.

- Option 1: when you are at a computer in an SU room at the university,
  - ssh -X username@sigma.nsc.liu.se where username is your NSC username (not the LiU one), -X indicates forwarding function of ssh which is used for running graphics applications remotely.

<sup>&</sup>lt;sup>6</sup>https://www.cendio.com/thinlinc/download

- [username@sigma  $\sim$ ] \$ exit is used to logout Sigma. If it is hung on, please use ctrl-c to terminate the connection.
- You can use Atom editor for coding by running following two commands first in a terminal:
  - \* [username@sigma  $\sim$ ] \$ module load atom-editor/1.45.0-bdist
  - \* [username@sigma  $\sim$ ] \$ atom &
- [username@sigma ~] \$ emacs & You can also use Emacs for coding by running emacs & in the terminal after you connect to Sigma or program locally on your machine, then use scp command to copy you files to Sigma.
- scp LOCAL\_FILE\_PATH username@sigma.nsc.liu.se:Documents is used for uploading files from your local machine to Sigma. (Note: you are supposed to run scp command before you log in Sigma when you want to upload files to Sigma.)
- You can always first connect to the university server and from there connect to Sigma. Information regarding remote access to LiU's system can be found <u>here</u>.
- Option 2: Sigma's Thinlinc server is available same as *sigma.nsc.liu.se*. In this way, you can get a graphical environment on Sigma and given that you work directly on Sigma there is no need to use *ssh* or *scp*. Note: Please remember to log out when done working on the labs so that Sigma does not keep open Thinlinc sessions. Also, each pair of students, please use at most one Thinlinc connection during lab sessions, due to the limited number of Thinlinc licenses on Sigma.
  - When you are at a computer in an SU room at the university:
    - \* You need to open a terminal and run following two lines:
    - \* \$ module load courses/732A54 or \$ module load courses/TDDE31
    - \* \$ tlclient
  - If you use your own computers, you can just download Thinlinc first and then connect directly.
  - As mentioned above, on Sigma, you can use Atom editor for coding by running following two commands in a terminal.
    - \* [username@sigma  $\sim$ ]  $\$  module load atom-editor/1.45.0-bdist
    - \* [username@sigma  $\sim$ ] \$ atom &

NSC reserves nodes for each lab session of the course, which means other jobs on Sigma will not use these nodes during our lab sessions. Table 4 shows reservation names for our scheduled lab sessions as well as a reservation, **devel**, which you can use at any time (e.g., outside of lab sessions).

Each time when you submit a job, the options '-A liu-compute-2023-10', and '-reservation RESERVATION NAME' should be specified.

- [username@sigma ~] \$ listreservations, by running this command, you can see all the reservations.
- [username@sigma  $\sim$ ] \$ sbatch -A liu-compute-2023-10 -reservation bigdata-2023-04-13 run\_yarn\_with\_historyserver.q

Table 4: Time and Reservation Name

RESERVATION_NAME	Time
bigdata-2023-04-13	04-13, 08:15 to 10:15
bigdata-2023-04-14	04-14, 15:15 to 17:15
bigdata-2023-04-18	04-18, 13:15 to 17:15
bigdata-2023-04-20	04-20, 08:15 to 10:15
bigdata-2023-04-24	04-24, 10:15 to 12:15
bigdata-2023-04-25	04-25, 13:15 to 17:15
bigdata-2023-04-27	04-27, 08:15 to 10:15
bigdata-2023-04-28	04-28, 15:15 to 17:15
bigdata-2023-05-02	05-02, 13:15 to 17:15
bigdata-2023-05-04	05-04, 08:15 to 10:15
bigdata-2023-05-05	05-05, 15:15 to 17:15
bigdata-2023-05-11	05-11, 08:15 to 10:15
bigdata-2023-05-12	05-12, 15:15 to 17:15
bigdata-2023-05-16	05-16, 13:15 to 17:15
bigdata-2023-05-23	05-23, 13:15 to 17:15
devel	Anytime

#### 3.2 Submitting Jobs to Hadoop Cluster and Sigma

To submit your pyspark code to the Hadoop cluster, you will use:

• spark-submit --deploy-mode cluster --master yarn --num-executors 9 -driver-memory 2g --executor-memory 2g --executor-cores 4 CODE.py where CODE.py is the python script in the current folder. In this command, Yarn is used for resource management and the cluster-deploy mode is used.

To run your pyspark code on Hadoop cluster, you also need to first submit a job to Sigma. Some scripts in the demo (in next section) are provided to you so you can make the call of your *spark-submit* command. You will use the non-interactive way to submit a job on Sigma. Each time after you submit, the job will enter the scheduling queue. You can use **sbatch** command to submit the job, **squeue** command to monitor the submitted job and may use **scancel** command to cancel a job.

- [username@sigma ~] \$ sbatch -A liu-compute-2023-10 -reservation bigdata-2023-04-13 run\_yarn\_with\_historyserver.q (Note: Please remember to use '-A liu-compute-2023-10' if you are envolved in more than one project from NSC, it will guarantee to use the allocation reserved on Sigma for our course.)
- [username@sigma  $\sim$ ] \$ squeue -u username
- [username@sigma  $\sim$ ] \$ scancel JOB\_ID

Sigma uses Slurm for scheduling. Once you submit a job, the job will be assigned an ID. After the job is finished, you will see a slurm-JOB\_ID.out file returned, which includes the output information of the job script. The script run\_yarn\_with\_historyserver.q is supposed to contain commands for constructing the Hadoop cluster for your account, commands

for interacting with Hadoop Distributed File System (HDFS) and *spark-submit* command for running pyspark code. A detailed example (Figure 3) will be shown in following section which introduces the demo.

Another way to access the logs after the execution, you will need to set the spark.eventLog.enabled which is already contained in 'run\_yarn\_with\_historyserver.q'. After the job finishes, you can access the history server by running:

- module load spark/2.4.3-hadoop-2.7-nsc1
- spark\_browse\_historyserver A liu-compute-2023-10 --reservation RESER-VATION

Then a firefox session that points at the history server web interface will be opened. In this way, it only lists the latest finished job on the history server.

#### 3.3 Demo Code

You can find the demo from following folders after you log in Sigma.

• /software/sse/manual/spark/BDA\_demo/

The steps to run the BDA\_demo are shown as below, and in Figure 1 and Figure 2.

- Step 1: Login Sigma with 'ssh -X' connection or Thinlinc.
- Step 2: Copy the demo to your home folder on Sigma.
- Step 3: Use **sbatch** to submit the job. Then use *squeue* command to monitor your job. Once the job is finished, check the returned file named *slurm-ID.out*. Notice: when you want to run your own code, you need to make some changes in 'run\_yarn\_with\_historyserver.q' to make sure you create folders on HDFS and copy data to HDFS correctly (see Figure 3).
- Step 4: Check the history log from slurm-ID.out or run spark\_browse\_historyserver, or check the output folder.

In 'run\_yarn\_with\_historyserver.q' as shown in Figure 3, you can see a number of commands that are used to interact with HDFS.

- hadoop fs -mkdir <FOLDER\_NAME> -make a folder on HDFS
- $hadoop\ fs\ -mkdir\ -p\ < FOLDER\_NAME> < FOLDER\_NAME>$ make multiple folders
- hadoop fs -test -d <FOLDER\_NAME> -if the path is a directory, return 0
- hadoop fs -rm -r <FOLDER\_NAME> -deletes the directory and any content under it recursively
- hadoop fs -cat <FOLDER\_ON\_HDFS> [local] -copy HDFS path to stdout
- hadoop fs -copyFromLocal <localsrc> ... <dst> -copy single src, or multiple srcs from local Sigma to HDFS
- hadoop fs -copyToLocal <dst> ... <localsrc> -copy single src, or multiple srcs from HDFS to local Sigma

```
(base) huali50@mac01048 ~ % ssh −X x_huali@sigma.nsc.liu.se 🚤
(x_huali@sigma.nsc.liu.se) Password:
(x_huali@sigma.nsc.liu.se) Verification code:
Last login: Mon Apr 3 17:13:23 2023 from 2001:6b0:17:fc08:c0d:92e5:b93c:2345
                                                                                       Step 1
Welcome to NSC and Sigma!
**≠ Project storage directories available to you:
/proj/theophys/users/x_huali
/proj/roarc/users/x_huali
/proj/liu-compute-2023-10/users/x_huali
**** Documentation and getting help:
https://www.nsc.liu.se/support/systems/sigma-getting-started/
https://www.nsc.liu.se/support
**** Useful commands
To see your active projects and CPU time usage: projinfo
To see available disk storage and usage: snicquota
To see your last jobs: lastjobs
Login to compute node to check running job: jobsh
To tweak job priorities, extend timelimits and reserve nodes: see
https://www.nsc.liu.se/support/batch-jobs/boost-tools/
(Run "nsc-mute-login" to not show this information)
[[x_huali@sigma ~]$ ls ◀─

    Check your current folder, you should have an empty one.

2_java_wordcount_1.0 732A54-te..
                                   Downloads MapReduceLab Pictures TDDD43LabNOSQL Videos
4_pyspark_wordcount Desktop
[x_huali@sigma ~]$ cp -r /software/sse/manual/spark/BDA_demo/ _/
[x_huali@sigma ~]$ ls
2_java_wordcount_1.0 732A54-test Desktop
                                              Step 2: Copy the demo code folder to you current folder.
   vspark wordcount
                      RDA demo
                                   Documents F1
[x_huali@sigma ~]$ cd BDA_demo/
                                                    Enter the folder and check the content
[x_huali@sigma BDA_demo]$ ls
Tx_huall@sigma_BuA_demoja_la_demo.py input_data_run_local_q run_local_with_historyserver.q Check reservations before submitting jobs.
[x_huali@sigma BDA_demo]$ listreservations 	
Reservations available to user:x_huali / project(s):liu-compute-2023-10,liu-2019-26
 bigdata-2023-04-13 from 2023-04-13T08:15:00 to 2023-04-13T10:15:00
                                                                       (project:liu-compute-2023-10)
 bigdata-2023-04-14
                    from 2023-04-14T15:15:00 to 2023-04-14T17:15:00
                                                                       (project:liu-compute-2023-10)
 bigdata-2023-04-18 from 2023-04-18T13:15:00 to 2023-04-18T17:15:00
                                                                       (project:liu-compute-2023-10)
 bigdata-2023-04-20
                    from 2023-04-20T08:15:00 to 2023-04-20T10:15:00
                                                                       (project:liu-compute-2023-10)
 bigdata-2023-04-24
                     from 2023-04-24T10:15:00 to 2023-04-24T12:15:00
                                                                       (project:liu-compute-2023-10)
 bigdata-2023-04-25
                     from 2023-04-25T13:15:00 to 2023-04-25T17:15:00
                                                                       (project:liu-compute-2023-10)
 bigdata-2023-04-27
                     from 2023-04-27T08:15:00 to 2023-04-27T10:15:00
                                                                       (project:liu-compute-2023-10)
                     from 2023-04-28T15:15:00 to 2023-04-28T17:15:00
                                                                       (project:liu-compute-2023-10)
 bigdata-2023-04-28
                     from 2023-05-02T13:15:00 to 2023-05-02T17:15:00
 bigdata-2023-05-02
                                                                       (project:liu-compute-2023-10)
 bigdata-2023-05-04
                     from 2023-05-04T08:15:00 to 2023-05-04T10:15:00
                                                                       (project:liu-compute-2023-10)
 bigdata-2023-05-05
                     from 2023-05-05T15:15:00 to 2023-05-05T17:15:00
                                                                       (project:liu-compute-2023-10)
 bigdata-2023-05-11
                     from 2023-05-11T08:15:00 to 2023-05-11T10:15:00
                                                                       (project:liu-compute-2023-10)
 bigdata-2023-05-12
                     from 2023-05-12T15:15:00 to 2023-05-12T17:15:00
                                                                       (project:liu-compute-2023-10)
 bigdata-2023-05-16
                     from 2023-05-16T13:15:00 to 2023-05-16T17:15:00
                                                                       (project:liu-compute-2023-10)
 bigdata-2023-05-23
                    from 2023-05-23T13:15:00 to 2023-05-23T17:15:00
                                                                       (project:liu-compute-2023-10)
 devel from NOW to INF (everyone)
Note: set one of the above as default by running:
 usereservation RESERVATIONNAME
Or without the usereservation alias:
 source /software/tools/bin/usereservation.sh RESERVATIONNAME
```

Figure 1: Login Sigma and Copy the demo code (Steps 1 and 2).

```
| huali50 - x_huali@sigma:~/BDA_demo/output - ssh -X x_huali@sigma.nsc.liu.se - 124×84
[x_huali@sigma BDA_demo]$ sbatch -A liu-compute-2023-10 --reservation devel run_yarn_with_historyserver.q
Submitted batch job 3450199
                                                               Step 3
[x_huali@sigma BDA_demo]$ squeue -u x_huali
                           NAME USER ST
                                                 TIME NODES NODELIST(REASON)
           JOBID PARTITION
                   sigma run_yarn x_huali R
          3450199
                                                0:12
                                                         2 n[1206,1209]
[[x_huali@sigma BDA_demo]$ squeue -u x_huali
           JOBID PARTITION
                            NAME
                                   USER ST
                                               TIME NODES NODELIST(REASON)
[[x_huali@sigma BDA_demo]$ ls
demo.py
[x_huali@sigma BDA_demo]$ cd output/
[x_huali@sigma output]$ ls
 SUCCESS part-00000 part-00001 ← Partition files copied back from
                                                                    Step 4: check this file
[x_hualicsigma output]$
                                  HDFS.
    This means the run of your program is successful (no syntax
    errors).
```

Figure 2: Submit a job and check the output (Steps 3 and 4).

```
#SBATCH --time=10:00
#SBATCH --nodes=2
#SBATCH --exclusive
module add spark/2.4.3-hadoop-2.7-nsc1
# Cleanup and start from scratch
rm -rf spark
echo "START AT: $(date)"
hadoop setup
echo "Prepare output and input directories and files..."
# The following command will make folders on your home folder on HDFS, the input and output folders should be correspondi
hadoop fs -mkdir -p "BDA" "BDA/input"

if [ "$?" == "0" ]; then

| "BDA/ | "BDA/| | 
                                                                                                                                                                                          Please read the
        hadoop fs -rm -r "BDA/output"
                                                                                                                                                                                           comments, you will need
                                                                                                                                                                                          to change this part.
hadoop fs -copyFromLocal ./input_data/temperature-readings-small.csv "BDA/input/"
# Remove the comment when you need specifc file below
#hadoop fs -copyFromLocal ./input_data/temperature-readings.csv "BDA/input/"
#hadoop fs -copyFromLocal ./input_data/precipitation-readings.csv "BDA/input/"
#hadoop fs -copyFromLocal ./input_data/stations.csv "BDA/input/"
"hadoop fs -copyFromLocal ./input_data/stations-Ostergotland.csv "BDA/input/"
                                                                                                                                                                            Here, you need to replace with
# Run your program
                                                                                                                                                                           your own code.
echo "Running Your program..."
exec 5>&1
APPLICATION_ID=$(spark-submit --conf spark.eventLog.enabled=true deploy-mode cluster --master yarn --num-executors 9 --
driver-memory 2g --executor-memory 2g --executor-cores 4 demo.py 2>&1 | tee >(cat - >&5) | awk '!found && /INFO.*Yarn.*Su
bmitted application/ {tmp=gensub(/^.*Submitted application (.*)$/,"\\1","g");print tmp; found=1}')
hadoop fs -cat "BDA/output"/*
echo
echo "Applicaton id: $APPLICATION_ID"
echo "=
yarn logs -applicationId "$APPLICATION_ID" | awk -F: '/^LogType/ {if($2=="stderr") {output=1} else {output=0}} output==1
{print}
echo "-------
echo "=
                                                                              result
echo "============
yarn logs -applicationId "$APPLICATION_ID" | awk -F: '/^LogType/ {if($2=="stdout") {output=1} else {output=0}} output==1
{print}' | grep -v "WARN\|INFO"
rm -rf output
hadoop fs -copyToLocal 'BDA/output' ./
hadoop_stop
echo "END AT: $(date)"
```

Figure 3: 'run\_yarn\_with\_historyserver.q' for running the code

# 4 Hand In

You are supposed to use GitLab<sup>7</sup> to submit your report and code. For each lab, please submit the code and a report that contains your results (a snippet of the results is enough if the results contain many rows) and answers to the questions. In cases where a plot of your results is asked, you can include the figure directly in the report. You can use a tool of your preference to produce the plots (R, Excel, matplotlib in Python, etc.). Comment each step in your code to provide a clear picture of your reasoning when solving the problem.

<sup>&</sup>lt;sup>7</sup>https://gitlab.liu.se/olaha93/bigdata