

CSE291E-lab3

Running Hadoop map reduce in docker

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Input file (pseudo distributed mode)

Input.txt:

```
line1. line1. line1!  
line2, line2, line2  
line3 line3..
```

Input file (cluster mode)

Input1.txt:

```
Hello Hadoop! xixixi  
Hello Docker! hehehe
```

Input2.txt

```
Hello Docker!  
Hello World!
```

Input3.txt

```
Hello Hadoop!  
Hello World!
```

How to run word count in a single docker

container using Hadoop pseudo distributed mode

- Change current directory: `cd /XXX.../lab3/single/` (change `XXX...` according to your machine)
- Build docker image and run the task: `single/build.sh`, wait until finish, then you will see the result.
- To clean up, execute `./clean.sh`

How to run Hadoop in cluster mode using docker

- Note: this section is based on the work of Kai LIU:
<https://github.com/kiwenlau/hadoop-cluster-docker>
- Configure the environment
 - Change current directory `cd cluster`
 - Pull the dependent docker image `sudo ./boost.sh`
 - Build the data volume image `sudo ./build-image.sh data-volume`, which includes most our code.
 - WARN: this script will remove all running / waiting containers, in order to work properly.
 - Start cluster containers `sudo ./bootstrap.sh`, now you will be in a session with the master node of a Hadoop cluster with 4 slave nodes.
- Run the same word count program as last section in cluster mode
 - `/data/wordcount.sh`
- Run the bigram count program in cluster mode
 - `/data/bigrams.sh`

How to run the demo in a pre-configured server (For CSE291E staff, available until June 9)

- log in the server: `ssh course_staff@291elab3.philosopherwang.me`, password is `cse291e`.
- `cd lab3`
- Run the demo in single mode:
 - `cd single`

- `sudo ./build.sh`, and see the result, compared with input file (pseudo distributed mode) above
- `cd ..`
- Run the demo in docker cluster:
 - `cd cluster`
 - `sudo ./bootstrap.sh`, you'll login to the master node of the cluster
 - To see all cluster members `serf members`
 - To run word count demo `/data/wordcount.sh`, compared with input file (cluster mode) above.
 - To run bigram demo `/data/bigrams.sh`, compared with input file (cluster mode) above.
 - Exit session with master node `exit`
- Exit the session with the server: `exit`

Reference

- Single node pseudo distributed mode is based on: <https://hub.docker.com/r/sequenceiq/hadoop-docker/>
- Cluster mode is based on: <https://github.com/kiwenlau/hadoop-cluster-docker>
- Word count example: https://hadoop.apache.org/docs/current/hadoop-mapreduce-client/hadoop-mapreduce-client-core/MapReduceTutorial.html#MapReduce_Tutorial