

Serverless Data Analysis with Dataflow: MapReduce in Dataflow (Python) | Qwiklabs

Tuesday, November 17, 2020 3:41 PM

Clipped from:

https://googlecourses.qwiklabs.com/course_sessions/71782/labs/11648

Overview

In this lab, you will identify Map and Reduce operations, execute the pipeline, use command line parameters.

Objective

- Identify Map and Reduce operations
- Execute the pipeline
- Use command line parameters

Setup

For each lab, you get a new GCP project and set of resources for a fixed time at no cost.

1. Make sure you signed into Qwiklabs using an **incognito window**.
2. Note the lab's access time (for example,

02:00:00

and make sure you can finish in that time block.

3. When ready, click

START LAB

4. Note your lab credentials. You will use them to sign in to Cloud Platform Console.

[Open Google Console](#)

Caution: When you are in the console, do not deviate from the lab instructions. Doing so may cause your account to be blocked. [Learn more.](#)

Username

google2876526_student@qwiklabs.n

Password

TG959yrKDX

GCP Project ID

qwiklabs-gcp-0855e773352d3560



[New to labs? View our introductory video!](#)

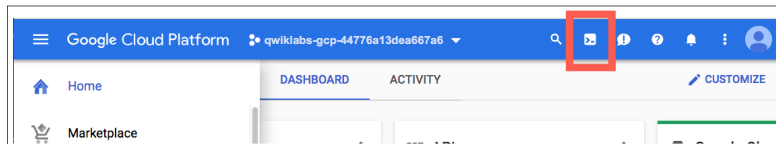
5. Click **Open Google Console**.
6. Click **Use another account** and copy/paste credentials for **this** lab into the prompts.

1. Accept the terms and skip the recovery resource page.

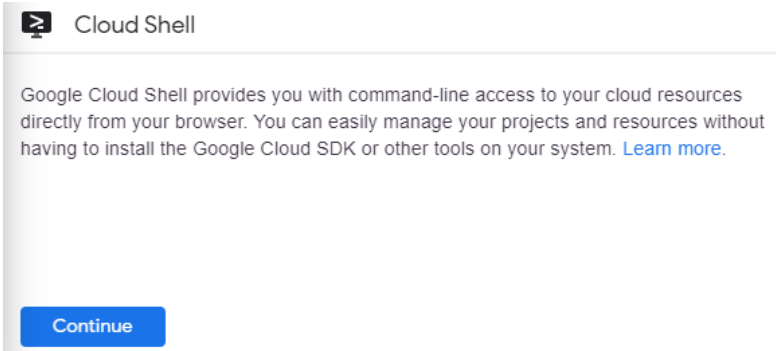
[Activate Google Cloud Shell](#)

Google Cloud Shell is a virtual machine that is loaded with development tools. It offers a persistent 5GB home directory and runs on the Google Cloud. Google Cloud Shell provides command-line access to your GCP resources.

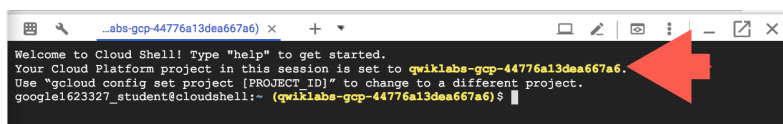
1. In GCP console, on the top right toolbar, click the Open Cloud Shell button.



2. Click **Continue**.



It takes a few moments to provision and connect to the environment. When you are connected, you are already authenticated, and the project is set to your *PROJECT_ID*. For example:



gcloud is the command-line tool for Google Cloud Platform. It comes pre-installed on Cloud Shell and supports tab-completion.

You can list the active account name with this command:

```
gcloud auth list
```

Output:

Credentialed accounts:

- [<myaccount>@<mydomain>.com](#) (active)

Example output:

Credentialed accounts:

- [google1623327_student@qwiklabs.net](#)

You can list the project ID with this command:

```
gcloud config list project
```

Output:

```
[core]
project = <project_ID>
```

Example output:

```
[core]
project = qwiklabs-gcp-44776a13dea667a6 Full
documentation of gcloud is available on Google
Cloud gcloud Overview.
```

[Launch Google Cloud Shell Code Editor](#)

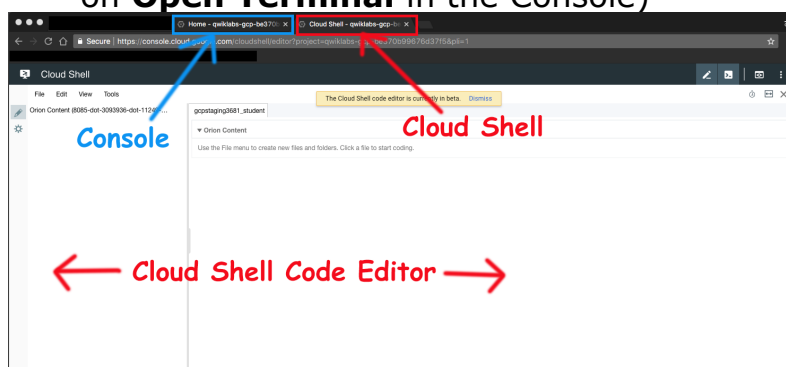
Use the Google Cloud Shell Code Editor to easily create and edit directories and files in the Cloud Shell instance.

Once you activate the Google Cloud Shell, click the **Open editor** button to open the Cloud Shell Code Editor.




You now have three interfaces available:

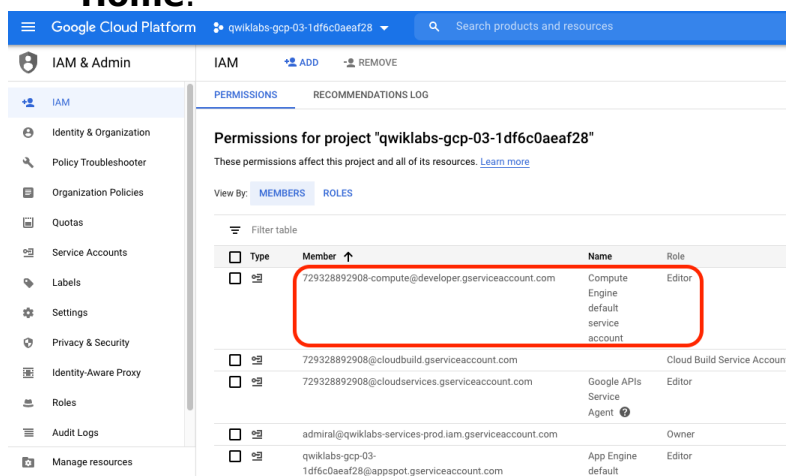
- The Cloud Shell Code Editor
- Console (By clicking on the tab). You can switch back and forth between the Console and Cloud Shell by clicking on the tab.
- The Cloud Shell Command Line (By clicking on **Open Terminal** in the Console)



Check project permissions

Before you begin your work on Google Cloud, you need to ensure that your project has the correct permissions within Identity and Access Management (IAM).

1. In the Google Cloud console, on the **Navigation menu** () , click **IAM & Admin** > **IAM**.
2. Confirm that the default compute Service Account [compute@developer.gserviceaccount.com](mailto:{project-number}-compute@developer.gserviceaccount.com) is present and has the editor role assigned. The account prefix is the project number, which you can find on **Navigation menu** > **Home**.



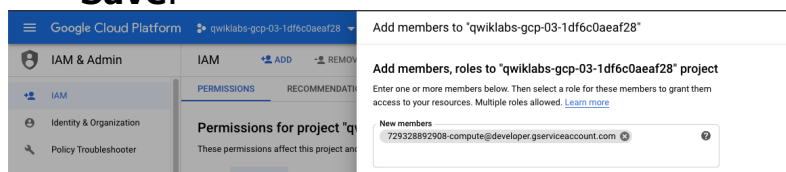
If the account is not present in IAM or does not have the editor role, follow the steps below to assign the required role.

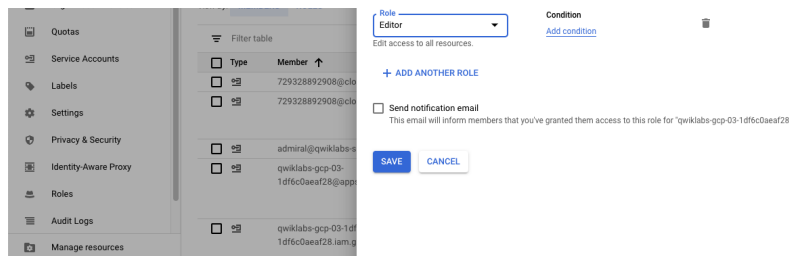
- In the Google Cloud console, on the **Navigation menu**, click **Home**.
- Copy the project number (e.g. 729328892908).
- On the **Navigation menu**, click **IAM & Admin** > **IAM**.
- At the top of the **IAM** page, click **Add**.
- For **New members**, type:

[compute@developer.gserviceaccount.com](mailto:{project-number}-compute@developer.gserviceaccount.com)

Replace {project-number} with your project number.

- For **Role**, select **Project** > **Editor**. Click **Save**.





Task 1. Review Preparations

These preparations should already have been done:

- Create Cloud Storage bucket
- Clone github [repository](https://github.com/GoogleCloudPlatform/training-data-analyst) to Cloud Shell

`git clone`

<https://github.com/GoogleCloudPlatform/training-data-analyst>

- Upgrade packages and install Apache Beam

```
cd training-data-analyst/courses/data_analysis/lab2/python
sudo ./install_packages.sh
```

Task 2. Identify Map and Reduce operations

1. In the Cloud Shell code editor navigate to the directory `/training-data-analyst/courses/data_analysis/lab2/python` and view the file `is_popular.py` in the Cloud Shell editor. **Do not make any changes to the code.**

Alternatively, you could view the file with `nano`.
Do not make any changes to the code.

```
cd ~/training-data-analyst/courses/data_analysis/lab2/python
nano is_popular.py
```

Can you answer these questions about the file `is_popular.py`?

- What custom arguments are defined?
- What is the default output prefix?
- How is the variable `output_prefix` in `main()` set?
- How are the pipeline arguments such as `--runner` set?
- What are the key steps in the pipeline?
- Which of these steps happen in parallel?
- Which of these steps are aggregations?

Task 3. Execute the pipeline

1. Run the pipeline locally:

```
cd ~/training-data-analyst/courses/data_analysis/lab2/python
python3 ./is_popular.py
```

1. Identify the output file. It should be **output**<suffix> and could be a sharded file.

```
ls -al /tmp
```

1. Examine the output file, replacing '-*' with the appropriate suffix.

```
cat /tmp/output-*
```

Task 4. Use command line parameters

1. Change the output prefix from the default value:

```
python3 ./is_popular.py --
output_prefix=/tmp/myoutput
```

1. What will be the name of the new file that is written out?
2. Note that we now have a new file in the **/tmp** directory:

```
ls -lrt /tmp/myoutput*
```

End your lab

When you have completed your lab, click **End Lab**. Qwiklabs removes the resources you've used and cleans the account for you.

You will be given an opportunity to rate the lab experience. Select the applicable number of stars, type a comment, and then click **Submit**.

The number of stars indicates the following:

- 1 star = Very dissatisfied
- 2 stars = Dissatisfied
- 3 stars = Neutral
- 4 stars = Satisfied
- 5 stars = Very satisfied

You can close the dialog box if you don't want to provide feedback.

For feedback, suggestions, or corrections,
please use the **Support** tab.

Last Updated Date: 2020-01-23

Last Tested Date: 2020-01-23