BigQuery: Qwik Start - Command Line | Google Cloud Skills Boost

Qwiklabs: 15-18 minutes

GSP071



Overview

Storing and querying massive datasets can be time consuming and expensive without the right hardware and infrastructure. BigQuery is a serverless, highly scalable cloud data warehouse that solves this problem by enabling super-fast SQL queries using the processing power of Google's infrastructure. Simply move your data into BigQuery and let us handle the hard work. You can control access to both the project and your data based on your business needs, such as giving others the ability to view or query your data.

You can access BigQuery by using the Console, Web UI or a command-line tool using a variety of client libraries such as Java, .NET, or Python. There are also a variety of solution providers that you can use to interact with BigQuery.

This hands-on lab shows you how to use bq, the python-based command line tool for BigQuery, to query public tables and load sample data into BigQuery.

Setup and requirements

Before you click the Start Lab button

Read these instructions. Labs are timed and you cannot pause them. The timer, which starts when you click **Start Lab**, shows how long Google Cloud resources will be made available to you.

This hands-on lab lets you do the lab activities yourself in a real cloud environment, not in a simulation or demo environment. It does so by giving you new, temporary credentials that you use to sign in and access Google Cloud for the duration of the lab.

To complete this lab, you need:

Access to a standard internet browser (Chrome browser recommended).

Note: Use an Incognito or private browser window to run this lab. This prevents any conflicts between your personal account and the Student account, which may cause extra charges incurred to your personal account.

• Time to complete the lab---remember, once you start, you cannot pause a lab.

Note: If you already have your own personal Google Cloud account or project, do not use it for this lab to avoid extra charges to your account.

How to start your lab and sign in to the Google Cloud Console

- 1. Click the **Start Lab** button. If you need to pay for the lab, a pop-up opens for you to select your payment method. On the left is the **Lab Details** panel with the following:
 - The Open Google Console button
 - Time remaining
 - o The temporary credentials that you must use for this lab
 - o Other information, if needed, to step through this lab
- 2. Click **Open Google Console**. The lab spins up resources, and then opens another tab that shows the **Sign in** page.

Tip: Arrange the tabs in separate windows, side-by-side.

Note: If you see the Choose an account dialog, click Use Another Account.

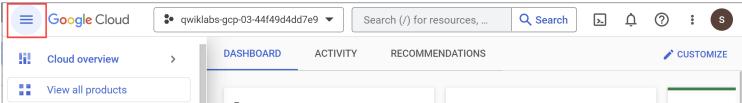
- 3. If necessary, copy the **Username** from the **Lab Details** panel and paste it into the **Sign in** dialog. Click **Next**.
- 4. Copy the Password from the Lab Details panel and paste it into the Welcome dialog. Click Next.

Important: You must use the credentials from the left panel. Do not use your Google Cloud Skills Boost credentials. **Note:** Using your own Google Cloud account for this lab may incur extra charges.

- 5. Click through the subsequent pages:
 - Accept the terms and conditions.
 - Do not add recovery options or two-factor authentication (because this is a temporary account).
 - Do not sign up for free trials.

After a few moments, the Cloud Console opens in this tab.

Note: You can view the menu with a list of Google Cloud Products and Services by clicking the **Navigation** menu at the top-left.



Activate Cloud Shell

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. Cloud Shell provides command-line access to your Google Cloud resources.

1. Click **Activate Cloud Shell 1.** at the top of the Google Cloud console.

When you are connected, you are already authenticated, and the project is set to your **PROJECT_ID**. The output contains a line that declares the **PROJECT_ID** for this session:

Your Cloud Platform project in this session is set to YOUR_PROJECT_ID

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

2. (Optional) You can list the active account name with this command:

gcloud auth list

- 3. Click Authorize.
- 4. Your output should now look like this:

Output:

ACTIVE: * ACCOUNT: student-01-xxxxxxxxxxx@qwiklabs.net To set the active account, run: \$ gcloud config set account `ACCOUNT`

5. (Optional) 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 **Note:** For full documentation of gcloud, in Google Cloud, refer to the gcloud CLI overview guide.

Task 1. Examine a table

BigQuery offers a number of sample tables that you can run queries against. In this lab, you'll run queries against the shakespeare table, which contains an entry for every word in every play.

To examine the schema of the Shakespeare table in the samples dataset, run:

bq show bigquery-public-data:samples.shakespeare

In this command you're doing the following:

- bq to invoke the BigQuery command line tool
- · show is the action
- Then you're listing the name of the project:public dataset.table in BigQuery that you want to see.

Output:

Task 2. Run the help command

When you include a command name with the help commands, you get information about that specific command.

1. For example, the following call to be help retrieves information about the guery command:

bq help query

2. To see a list of all of the commands bq uses, run just bq help.

Task 3. Run a query

Now you'll run a query to see how many times the substring "raisin" appears in Shakespeare's works.

- 1. To run a query, run the command bq query "[SQL STATEMENT]":
- Escape any quotation marks inside the [SQL STATEMENT] with a \ mark, or
- Use a different quotation mark type than the surrounding marks ("versus").
- 2. Run the following standard SQL query in Cloud Shell to count the number of times that the substring "raisin" appears in all of Shakespeare's works:

bq query --use_legacy_sql=false \ 'SELECT word, SUM(word_count) AS count FROM `bigquery-public-data`.samples.shakespeare WHERE word LIKE "%raisin%" GROUP BY word'

In this command:

• --use_legacy_sql=false makes standard SQL the default query syntax.

Output:

Waiting on job_e19 ... (0s) Current status: DONE +-----+ | word | count | +-----+ | praising | 8 | | Praising | 4 | | raising | 5 | | dispraising | 2 | | dispraising | 1 | | raisins | 1 |

The table demonstrates that although the actual word **raisin** doesn't appear, the letters appear in order in several of Shakespeare's works.

Test completed task

Click **Check my progress** to verify your performed task. If you have successfully run a query against a public dataset, you will see an assessment score.

Run a query (dataset: samples, table: shakespeare, substring: raisin)

If you search for a word that isn't in Shakespeare's works, no results are returned.

• Run the following search for "huzzah", returns no matches:

bq query --use_legacy_sql=false \ 'SELECT word FROM `bigquery-public-data`.samples.shakespeare WHERE word = "huzzah"

Test completed task

Click **Check my progress** to verify your performed task. If you have successfully run a query against a public dataset, you will see an assessment score.

Run a query (dataset: samples, table: shakespeare, substring: huzzah)

Task 4. Create a new table

Now create your own table. Every table is stored inside a dataset. A *dataset* is a group of resources, such as tables and views.

Create a new dataset

1. Use the bg 1s command to list any existing datasets in your project:

bq Is

You will be brought back to the command line since there aren't any datasets in your project yet.

2. Run bq ls and the bigquery-public-data Project ID to list the datasets in that specific project, followed by a colon (:):

bq ls bigquery-public-data:

Output:

datasetId ------ austin_311 austin_bikeshare austin_crime austin_incidents austin_waste baseball bitcoin_blockchain bls census_bureau_construction census_bureau_international census_bureau_usa census utility chicago crime ...

Now create a dataset. A dataset name can be up to 1,024 characters long, and consist of A-Z, a-z, 0-9, and the underscore, but it cannot start with a number or underscore, or have spaces.

3. Use the bq mk command to create a new dataset named babynames in your project:

bq mk babynames

Sample output:

Dataset 'qwiklabs-gcp-ba3466847fe3cec0:babynames' successfully created.

Test completed task

Click **Check my progress** to verify your performed task. If you have successfully created a BigQuery dataset named babynames, you will see an assessment score.

Create a new dataset (name: babynames)

• Run bq 1s to confirm that the dataset now appears as part of your project:

bq Is

Sample output:

datasetId ----- babynames

Upload the dataset

Before you can build the table, you need to add the dataset to your project. The custom data file you'll use contains approximately 7 MB of data about popular baby names, provided by the US Social Security Administration.

1. Run this command to add the baby names zip file to your project, using the URL for the data file:

curl -LO http://www.ssa.gov/OACT/babynames/names.zip

2. List the file:

ls

You can see the name of the file added to your project.

3. Now unzip the file:

unzip names.zip

4. That's a pretty big list of text files! List the files again:

ls

The bq load command creates or updates a table and loads data in a single step.

You will use the bq load command to load your source file into a new table called names2010 in the babynames dataset you just created. By default, this runs synchronously, and will take a few seconds to complete.

The bq load arguments you'll be running are:

datasetID: babynames tableID: names2010 source: yob2010.txt schema: name:string,gender:string,count:integer

5. Create your table:

bq load babynames.names2010 yob2010.txt name:string,gender:string,count:integer

Sample output:

Waiting on job_4f0c0878f6184119abfdae05f5194e65 ... (35s) Current status: DONE

Test completed task

Click **Check my progress** to verify your performed task. If you have successfully loaded data into a dataset table, you will see an assessment score.

Load the data into a new table

1. Run bq 1s and babynames to confirm that the table now appears in your dataset:

bq Is babynames

Output:

tableId Type ----- names2010 TABLE

2. Run bq show and your dataset.table to see the schema:

bq show babynames.names2010

Output:

Task 5. Run queries

Now you're ready to query the data and return some interesting results.

1. Run the following command to return the top 5 most popular girls names:

bq query "SELECT name,count FROM babynames.names2010 WHERE gender = 'F' ORDER BY count DESC LIMIT 5"

Output:

Waiting on job_58c0f5ca52764ef1902eba611b71c651 ... (0s) Current status: DONE +-----+ | name | count | +-----+ | Isabella | 22913 | | Sophia | 20643 | | Emma | 17345 | | Olivia | 17028 | | Ava | 15433 | +-----+

2. Run the following command to see the top 5 most unusual boys names:

bq query "SELECT name,count FROM babynames.names2010 WHERE gender = 'M' ORDER BY count ASC LIMIT 5" **Note:** The minimum count is 5 because the source data omits names with fewer than 5 occurrences.

Output:

Waiting on job_556ba2e5aad340a7b2818c3e3280b7a3 ... (1s) Current status: DONE +-----+ | name | count | +-----+ | Aaqib | 5 | | Aaidan | 5 | | Aadhavan | 5 | | Aarian | 5 | | Aamarion | 5 | +-----+

Test completed task

Click **Check my progress** to verify your performed task. If you have successfully a against a custom dataset, you will see an assessment score.

Run queries against your dataset table

Task 6. Test your Understanding

Below are multiple choice questions to reinforce your understanding of this lab's concepts. Answer them to the best of your abilities.

Task 7. Clean up

1. Run the bq rm command to remove the babynames dataset with the -r flag to delete all tables in the dataset:

bq rm -r babynames

2. Confirm the delete command by typing Y.

Test completed task

Click **Check my progress** to verify your performed task. If you have successfully removed babynames dataset, you will see an assessment score.

Remove the babynames dataset

Congratulations!

Now you can use the command line with BigQuery to manipulate data.

Finish your quest

This self-paced lab is part of the BigQuery for Data Warehousing, NCAA® March Madness®: Bracketology with Google Cloud, BigQuery Basics for Data Analysts, and Using the Cloud SDK Command Line quests. A quest is a series of related labs that form a learning path. Completing a quest earns you a badge to recognize your achievement. You can make your badge or badges public and link to them in your online resume or social media account. Enroll in any quest that contains this lab and get immediate completion credit. See the Google Cloud Skills Boost catalog to see all available quests.

Next steps / Learn more

This lab is also part of a series of labs called Qwik Starts. These labs are designed to give you a little taste of the many features available with Google Cloud. Search for "Qwik Starts" in the lab catalog to find the next lab you'd like to take!

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