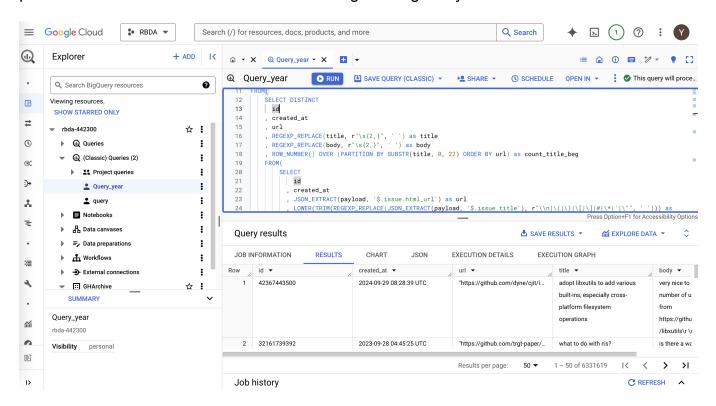
Data perparation and ingestion

In our project, I selected <u>CH Archive</u> of 2023 and 2024 as one of data source. CH Archive dataset includes all records of the public GitHub timeline, such as all kinds of issues. Therefore, we will use content of issues in recent two years to analyze the trend of technology that has been used with time.

Data Preparation

CH Archive dataset could be obtained from Google BigQuery with SQL query. The following picture shows the user interface of data fetching with BigQuery.



Parts of SQL query over CH Archive table of 2023 and 2024 is listed as below:

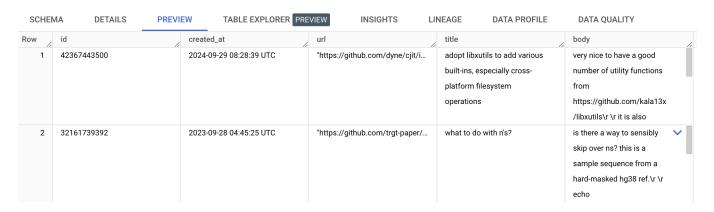
```
SELECT
   id,
   created_at,
   JSON_EXTRACT(payload, '$.issue.html_url') as url,
   LOWER(TRIM(REGEXP_REPLACE(JSON_EXTRACT(payload, '$.issue.title'),
r"\\n|\(|\)|\[|\]|#|\*|`|\"", ' '))) as title,
   LOWER(TRIM(REGEXP_REPLACE(JSON_EXTRACT(payload, '$.issue.body'), r"\\n|\(|\)|\[|\]|#|\*|`|\"", ' '))) as body
FROM `githubarchive.day.2024*`
WHERE
```

```
_TABLE_SUFFIX BETWEEN '0101' AND '1031'
AND type="IssuesEvent"
AND JSON_EXTRACT(payload, '$.action') = "\"opened\""
```

Finally we can get our raw dataset with following schema:

Field Name	Туре	Description
id	STRING	id of each issue
created_at	TIMESTAMP	create time of each issue
url	STRING	url for each issue
title	STRING	issue title
body	STRING	content of issue

Here is a preview of raw dataset:



The raw dataset consists of 6,331,619 rows, and it is 4.13 GB in size. Then the dataset is exported to bucket gs://nyu-dataproc-hdfs-ingest/gharchive for further ingestion.

Data ingestion

Data ingestion consists of several steps:

- 1. Total order sort for all data by id.
- Normalize the created_at timestamp to a consistent format as time field.
- Remove special characters or extra whitespaces from title and body, and combine them as text field.
- 4. Remove or replace non-printable characters, such as \r.
- 5. Add a source field with value GH Archive.

Design of Mapper and Reducer:

- Mapper:
 - For sampling phase, mapper should yield id from each line in JSON input file.
 - For total order sorting phase, only identity mapper is required.
- Reducer:
 - For sampling phase, we don't need any reducer.
 - For total order sorting phase, reducer should do data cleaning as mentioned above and then output values.

Execution:

Upload all DataIngest.java file to DataProc, compile source code and generate *.jar file.

```
javac -classpath `hadoop classpath` *.java
jar cvf dataIngest.jar *.class
```

Usage: DataIngest <input_path> <output_path>

Then we can run DataIngest job using jar file.

```
hadoop jar dataIngest.jar DataIngest gs://nyu-dataproc-hdfs-ingest/gharchive
gs://nyu-dataproc-hdfs-ingest/gharchive/output
```

Result:

After data ingestion with steps above, we can get the processed data, which is a snippet of the whole output data:

```
{
  "id": "26163418660",
  "time": "2023-01-01 00:00:00 UTC",
  "text": "chart editor glitched out and overwrote a completed chart with a
blank chart, describe your bug here....",
  "source": "GH Archive"
}

{
  "id": "26163419820",
  "time": "2023-01-01 00:00:08 UTC",
  "text":"not a problem but a request, is there any way of creating a column
for the numer of ratings.....",
  "source": "GH Archive"
}
```

```
{
  "id": "26163424178",
  "time": "2023-01-01 00:00:51 UTC",
  "text": "play button doesn't work after processing all files or changing
the playlist....",
  "source": "GH Archive"
}
```

There are 10 partitions of data file in total, each partition is about 260Mb in size. Data files are located at gs://nyu-dataproc-hdfs-ingest/gharchive/output directory.

nyu-dataproc-hdfs-ingest

