1. Steps to setup and run ETL pipeline.
   1. Download code from git - <https://github.com/medic/chis-visualizations/tree/ETL>
   2. Create and insert data to source tables in Postgres as mentioned in source\_data\_creation.sql
   3. Create target star schema tables and views using the queries present in star\_schema\_tables.sql.
   4. Create meta data table as provided in meta\_data\_details.sql
   5. Update the .dbenv file with the credentials of the data base on given environment.
   6. Execute main.py to trigger the ETL.
2. Documentation for super set dashboards can be found here - <https://docs.google.com/document/d/1kQ2UEOLFmPt6uFfcryq95dL9i-a6TyOX/edit>
3. Overview of the ETL, Datamodel, Superset dashboard can be found here

<https://docs.google.com/presentation/d/19yj8FSmQXwjRnJOUZfP3TZVk9YiNxdDH/edit#slide=id.p1>

1. Detail overview of the ETL can be found here –

<https://docs.google.com/presentation/d/1WItFFb7arPh67OTYvXKq0g06nCZpkJH0/edit>

1. Readme.md on git hub explains the details about below scenarios
   1. Capturing new column from source data.
   2. Adding new transformation rules in ETL pipeline
   3. On boarding new source and target to ETL
   4. Updating views involving x\_form\_ids