Connection to CHEMBL database and get molecules for Zika virus Apuntes de clase

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IMPORT LIBRARIES

```
(Adam, Lee, y Wang 2023)

# PATH TO CUSTOM MODULES
import sys

sys.path.append("../src")

# IMPORT LIBRARIES
from chembl_webresource_client.new_client import new_client
import pandas as pd

# IMPORT CUSTOM MODULES
import repository.chembl_api_connection as chembl
```

PATHS

PATHS TO SAVE DATA
data_path = "../data/raw/"

GET MOLECULES FOR ZIKA VIRUS

```
# GET THE TARGET ID WITH MAXIMUM NUMBER OF IC50 MOLECULES FOR A GIVEN ORGANISM
organism = "Zika virus"
activity = "IC50"
target_id = chembl.get_target_id_maximum_activity(organism, activity)

# GET MOLECULES FOR TARGET ID AND ACTIVITY TYPE
molecules = chembl.get_molecules_from_target_activity(target_id, activity)
```

SAVE MOLECULES TO CSV

```
# SAVE MOLECULES TO CSV
molecules.to_csv(data_path + target_id + "_raw.csv", index=False)
```

Adam, Awadalkareem, Christy Lee, y Tian Wang. 2023. «Rational Development of Live-Attenuated Zika Virus Vaccines». *Pathogens* 12 (2): 194. https://doi.org/10.3390/pathogens12020194.