1. Install Postgressql if you do not have it on your machine yet (<https://www.postgresql.org/download/>)
2. Run pgadmin
   1. Go to PostgreSQL15 tab
   2. Right Click on Database and create database named “glycogenes”
   3. Click on database “glycogenes”
   4. Go to “Tools/Query Tool”
   5. Open and run “create\_schema\_glyco.sql” in the subfolder db-queries (this should create a table called “tf\_glyco”)
   6. Check that you can find “tf\_glyco” table under “Schemas/public/tables”
3. Go to “data” subfolder. You should see the folder “glyco” containing txt files of the GRNs for different tissues.
4. While in “data” subfolder, run: “python ../data-loader/load\_data\_glyco.py glyco/\*.txt”
5. Run ui.