JENNIFER HY LIN



■ DATA ANALYTICS PROJECTS

current 2022

Phenotypes associated with rare diseases¹

RStudio and Jupyter notebook in Anaconda

- · Exploratory data analysis stage
- · Used R mainly with some Python initially for data cleaning

current 2022

COVID-19 antivirals in phase 3 clinical trials

RStudio and/or Jupyter notebook in Anaconda

- · Data wrangling stage
- · Data collected from ClinicalTrials.gov

2022 2022 Drugs in rare diseases²

RStudio and Jupyter notebook in Anaconda

- Two R versions with one in base R and the other in Tidyverse

Natural history of rare diseases - malformation syndrome³

· Python version available

2022 2022

Jupyter notebook in Anaconda

- · Completed
- · All in Python

2022 2022 Long COVID data in SQL4

Dbeaver

- · Completed
- · Used mySQL server

2022 2022 Long COVID dashboard⁵

Tableau Public

- · Completed
- · Dataset from journal paper

View this CV online with links at jhylin.github.io/Portfolioprojects/Resume_JHYL.html

CONTACT

- **y** jenhylin
- **G**itHub
- **9** My data analytics and research portfolio in LinkedIn profile

ANALYTIC SKILLS

Made with the R package pagedown.

The source code is available on GitHub.

Acknowledgement to another helpful source on GitHub

Last updated on 2022-07-27.

