# **Experiments protocol**

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#### 1. Overview

- The experiments aim to analyze data from the CSCR\_data dataset using various machine learning techniques.
- The experiments are implemented in Python scripts and a Jupyter notebook.

## 2. Directory Structure

Root Directory:

T:\Studies\CSCR\code\cscr\_stats\_ml

Gitlab:

papers / CSCR\_stats\_ML · GitLab (sib.swiss)

- Contains:
  - Data file: CSCR\_data.json
  - Code files:
    - correlation.py: Computes Pearson correlation between pairs of biomarkers.
    - stat\_test.py: Implements t-test and Kolmogorov-Smirnov (K-S) test.
    - stat\_model.py: Implements logistic regression statistical model.
    - utils.py: These are utilities functions used inside the three other py files
  - Notebook: experiment\_pipeline.ipynb
  - Environment file: ml\_paper\_env.yml

## 3. Environment Setup

- Create the environment using conda env create -f ml paper env.yml.
- Modify the prefix in the YAML file with the desired environment name.
- After setup, activate the environment using conda activate ml paper.

I had a problem with the seaborn library to display the correlation matrix. If it happens, try to reinstall it with pip (conda uninstall seaborn, then pip install seaborn)

# 4. Running Experiments

- Open and run the notebook experiment\_pipeline.ipynb.
- Execute all cells in the notebook to obtain results.

## 5. Results

- Upon running the notebook, the following results will be generated:
  - Correlation matrix
  - T-test results (for single visit data)
  - K-S test results (for longitudinal data)
  - Predictive modeling experiment

# 6. Figures

• Figures are generated within the notebook, allowing for easy reformatting for publication purposes.