# Code and dataset description

A comprehensive evaluation of drug responsiveness prediction models, using cell viability inhibitory concentrations (IC50 values)

## Code repository



This repository contains python code for the construction of drug response prediction models in each dataset

### Code repository (e.g. scenario 1: MDG-160K)



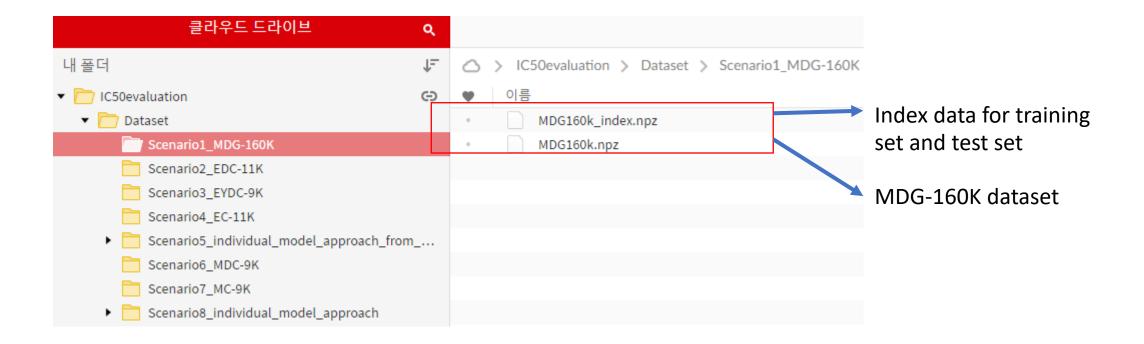
This repository contains python code for the construction of drug response prediction models in each dataset

## Code repository (e.g. scenario 2: EDC-11K)

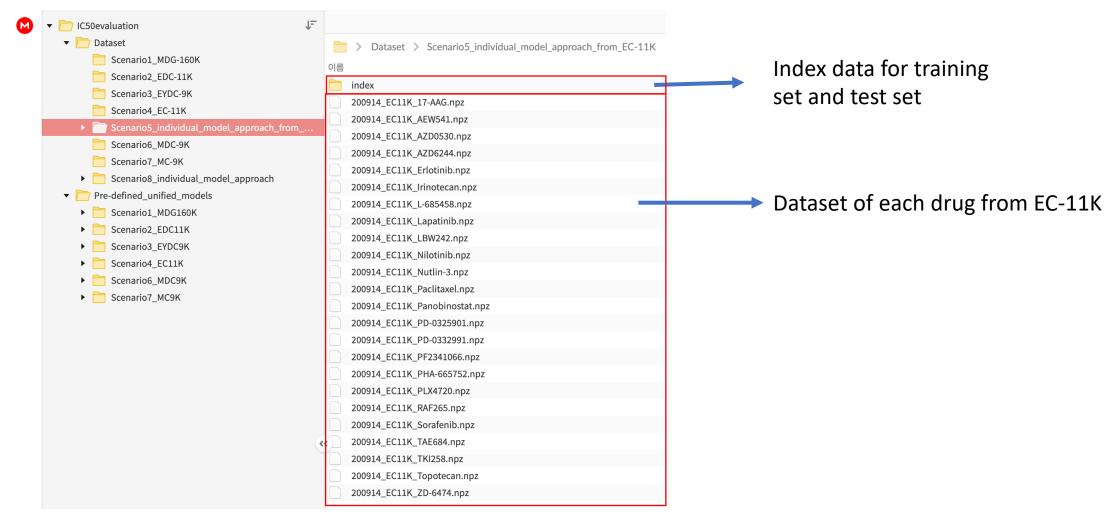
labnams IC50evaluation		785edfd on 12 Mar 🖰 <b>History</b>
ipynb_checkpoints	IC50evaluation	6 months ago
191205_EDC11K_znorm_minmax_CNN.ipynb	IC50evaluation	6 months ago
191205_EDC11K_znorm_minmax_Lasso.ipynb	IC50evaluation	6 months ago
191205_EDC11K_znorm_minmax_RF.ipynb	IC50evaluation	6 months ago
191205_EDC11K_znorm_minmax_ResNet.ipynb	IC50evaluation	6 months ago
191205_EDC11K_znorm_minmax_Ridge.ipynb	IC50evaluation	6 months ago
191205_EDC11K_znorm_minmax_SVR.ipynb	IC50evaluation	6 months ago
191205_EDC11K_znorm_minmax_XGB.ipynb	IC50evaluation	6 months ago

# Code repository (e.g. scenario 5: individual model approach)

#### Dataset repository – dataset for unified model



# Dataset repository – dataset for individual model approach



https://mega.nz/folder/CeYGDKyS#uqkmWJ4E2XSGJp C2VO2gg/folder/nWZVGYjZ

#### Dataset repository

