```
In [14]:
          import pandas as pd
          import numpy as np
          import pickle
          from tensorflow.keras.models import load_model
          import warnings
          from sklearn.metrics import log_loss
          from tqdm import tqdm
          warnings.filterwarnings("ignore")
In [224..
          def final_fun_1(X):
              #encoding categorical features
              X.iloc[:,1] = X.iloc[:,1].map({'trt_cp':0, 'ctl_vehicle':1})
              X.iloc[:,2] = X.iloc[:,2].map({24:0, 48:1, 72:2})
              X.iloc[:,3] = X.iloc[:,3].map({'D1':0, 'D2':1})
              #normalizing gene and cell columns
              transformer = pickle.load(open('transform.pkl', 'rb'))
              X.iloc[:,4:] = transformer.transform(X.iloc[:,4:])
              #getting more features using auto-encoder
              encoder = load_model('encoder.h5')
              encoded_features = encoder.predict(X.iloc[:,1:])
              #adding encoded features with original features
              total_X = pd.concat([X,pd.DataFrame(encoded_features)], axis=1)
              #loading pre-trained model
              model = pickle.load(open('final_model.pkl','rb'))
              #loading column names of target columns
              columns = pickle.load(open('target_columns.pkl', 'rb'))
              #predictions
              pred = model.predict(total_X.iloc[:,1:])
              pred_data = pd.DataFrame(pred, columns = columns)
              pred_data.insert(loc=0, column='sig_id', value = X['sig_id'])
              pred_prob = model.predict_proba(total_X.iloc[:,1:])
              pred_prob_data = pd.DataFrame(pred_prob,columns=columns)
              pred_prob_data.insert(loc=0, column='sig_id', value = X['sig_id'])
              return pred_data, pred_prob_data
In [37]:
          def final_fun_2(X,y):
              #encoding categorical features
              X.iloc[:,1] = X.iloc[:,1].map({'trt_cp':0, 'ctl_vehicle':1})
              X.iloc[:,2] = X.iloc[:,2].map({24:0, 48:1, 72:2})
              X.iloc[:,3] = X.iloc[:,3].map({'D1':0, 'D2':1})
              #normalizing gene and cell columns
              transformer = pickle.load(open('transform.pkl','rb'))
              X.iloc[:,4:] = transformer.transform(X.iloc[:,4:])
              #getting more features using auto-encoder
              encoder = load_model('encoder.h5')
              encoded_features = encoder.predict(X.iloc[:,1:])
              #adding encoded features with original features
              total_X = pd.concat([X,pd.DataFrame(encoded_features)], axis=1)
              #loading pre-trained model
              model = pickle.load(open('final_model.pkl','rb'))
              #prediction
              pred_prob = model.predict_proba(total_X.iloc[:,1:])
              metric_value = [] #list to store log-loss of each target features
              for i in tqdm(range(y.shape[1]-1)): #iterating over each target columns
                  loss = log_loss(y.iloc[:,i],pred_prob[:,i], labels=[0, 1]) #computing log-loss
                  metric_value.append(loss)
              return np.mean(metric_value)
```