Structured Data Assignment – Akaike By Kavibarathi K

Problem statement:

The development of drugs is critical in providing therapeutic options for patients suffering from chronic and terminal illnesses. "Target Drug", in particular, is designed to enhance the patient's health and well-being without causing dependence on other medications that could potentially lead to severe and life-threatening side effects. These drugs are specifically tailored to treat a particular disease or condition, offering a more focused and effective approach to treatment, while minimising the risk of harmful reactions.

Objective:

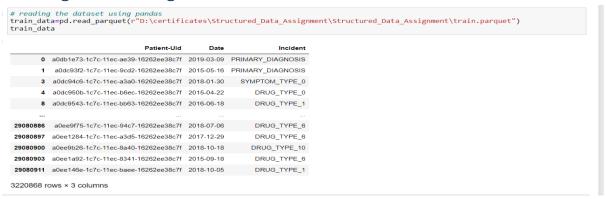
The objective in this assignment is to develop a predictive model which will predict whether a patient will be eligible*** for "Target Drug" or not in next 30 days. Knowing if the patient is eligible or not will help physician treating the patient make informed decision on the which treatments to give.

Necessary Libraries:

These are python libraries that are necessary to perform the functions for the model predictions.

```
import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.metrics import accuracy_score,f1_score,confusion_matrix,classification_report
from xgboost import XGBClassifier
from sklearn.svm import SVC
from sklearn.ensemble import RandomForestClassifier
```

Reading the Training Dataset:



Data Preprocessing:

This step includes detailed analysis of dataset and perform necessary process like checking for missing values, data type conversions, removing duplicate data's in the dataset. This is the most important process prior to model building in the predictions.

Creating a Positive and Negative data frames:

In this the positive data contains the set of data that are being test using the target drug and in the negative data incidents that are other than the "target drug" are taken into account .

```
# creating an object named positive result and storing the data that are tested with target drugs
positive_result_tr=train_data[train_data['Incident']=='TARGET DRUG']

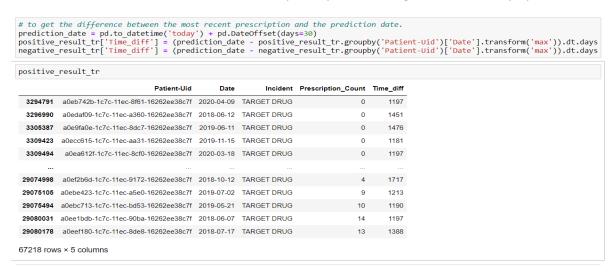
positive_result_tr

# creating an object named negative result and storing the data that are not tested with target drug
negative_data_tr=train_data['Patient-Uid'].isin(positive_result['Patient-Uid'])]
negative_result_tr = negative_data.groupby('Patient-Uid').tail(1)

negative_result_tr
```

Time difference column:

In this column, find out the time taken for the prescriptions of drug to find out the symptoms.



Concatenation the dataset:

The positive and negative dataset's are joined together using the pandas concat function to perform the model predictions.



Model building:

• The model building is the process in which we can use several classification algorithms to get the max accuracy from the dataset we have developed.

Train-Test-Split:

```
# creating train test split
x_train,x_test,y_train,y_test =train_test_split(new_data[['Prescription_Count','Time_diff']],new_data['Incident']=='TARGET_DRUG',
```

Model creation:

```
# model building for the predictions
model =XGBClassifier()
model.fit(x_train,y_train)
train_predict =model.predict(x_train)
test_predict =model.predict(x_test)
```

- Out of several model's that are tested for the best predictions XGBoost Classifier gave the highest F1-score.
- Model performance evaluated using several metrics.

```
# evaluating the model using accuracy score
print('Accuracy Score :',accuracy_score(y_test,test_predict))
Accuracy Score: 0.9634476916560779
# evaluating the model using the f1 score
print('F1 score :',f1_score(y_test,test_predict))
F1 score: 0.9784373984958649
# evaluating using confusion matrix
print('confusion Matrix :',confusion_matrix(y_test,test_predict))
confusion Matrix : [[ 3167 303]
 [ 560 19580]]
# evaluating using confusion matrix
print(classification_report(y_test,test_predict))
               precision recall f1-score support
                   0.85 0.91
0.98 0.97
        False
         True
                                        0.98
                                                     20140
                0.92 0.94 0.93
0.96 0.96 0.96
     accuracy
    macro avg
                                                     23610
weighted avg
                                                     23610
```

Tested model using test dataset:

In this data set all kind of necessary process as did in the training dataset.

```
# predicting the test data
test_data_predict =model.predict(new_data_ts[['Prescription_Count','Time_diff']])

test_data_predict
array([0, 0, 1, ..., 0, 0, 0])
```

Final submission:

```
: # final submission file after completing prediction
Final_submission = pd.DataFrame({'Patient-Uid': new_data_ts['Patient-Uid'], 'Prediction': test_data_predict})
Final_submission
```

	Patient-Uid	Prediction
57	a0f9e8a9-1c7c-11ec-8d25-16262ee38c7f	0
208	a0f9e9f9-1c7c-11ec-b565-16262ee38c7f	0
305	a0f9ea43-1c7c-11ec-aa10-16262ee38c7f	1
420	a0f9ea7c-1c7c-11ec-af15-16262ee38c7f	0
497	a0f9eab1-1c7c-11ec-a732-16262ee38c7f	0
1372381	a102720c-1c7c-11ec-bd9a-16262ee38c7f	1
1372432	a102723c-1c7c-11ec-9f80-16262ee38c7f	0
1372543	a102726b-1c7c-11ec-bfbf-16262ee38c7f	0
1372607	a102729b-1c7c-11ec-86ba-16262ee38c7f	0
1372859	a10272c9-1c7c-11ec-b3ce-16262ee38c7f	0

11482 rows × 2 columns