Predicting Patient ReAdmissions with Classification

August 27, 2021

1 Research Question

Our goal is to predict whether or not a patient will be readmitted to a hospital within 30 days of their initial discharge. We will be using the K-Nearest Neighbor (KNN) method to examine the patient's pre-existing health conditions as well as demographics such as gender and age.

2 Method Justification

KNN works by grouping records together that have similar features. By doing this to our patient data we can see if a set of features is likely to contribute to readmissions. Predictions made by KNN assume that similar inputs produce similar outputs.

2.1 Tools, Packages, and Libraries

We'll be using python for this analysis. Python has many libraries that make the data manipulation and analysis tasks that we are going to perform easier. The particular libraries that we will make use of are:

- Numpy and pandas for standard dataframe and numerical operations
- sklearn for it's KNN implementation as well as it's preprocessing module for scalers, model_selection module for train/test splits, and metrics for the AUC score.
- matplotlib for plots

We will now install and import them:

```
[1]: %%capture
    # install packages
    !pip install numpy
! pip install pandas
    !pip install matplotlib
    !pip install sklearn

# import packages
import pandas as pd
import numpy as np

from sklearn.neighbors import KNeighborsClassifier
```

```
from sklearn.preprocessing import StandardScaler
from sklearn.model_selection import train_test_split
from sklearn.metrics import roc_auc_score
import matplotlib.pyplot as plt

# random seed
import random
random.seed(10)
```

3 Data Preparation

3.1 Selection

The variables that we will be including in our analysis are:

Variable	Type
Age	continuous
Gender	categorical
$VitD_levels$	continuous
$Soft_drink$	categorical
HighBlood	categorical
Stroke	categorical
Overweight	categorical
Arthritis	categorical
Diabetes	categorical
Hyperlipidemia	categorical
BackPain	categorical
Anxiety	categorical
Allergic_rhinitis	categorical
Reflux_esophagitis	categorical
Asthma	categorical
ReAdmis	categorical

```
'Diabetes',
    'Hyperlipidemia',
    'BackPain',
    'Anxiety',
    'Allergic_rhinitis',
    'Reflux_esophagitis',
    'Asthma',
    'ReAdmis'
]]
df.head()
# Define outcome and predictors
outcome = 'ReAdmis'
predictors = [
    'Age',
    'Gender',
    'VitD_levels',
    'Soft_drink',
    'HighBlood',
    'Stroke',
    'Complication_risk',
    'Overweight',
    'Arthritis',
    'Diabetes',
    'Hyperlipidemia',
    'BackPain',
    'Anxiety',
    'Allergic_rhinitis',
    'Reflux_esophagitis',
    'Asthma',
]
```

3.2 Cleaning and Scaling

There are many categorical variables in this set. KNN requires numeric variables, so we will need to convert them. We will also separate the dataset into outcome and predictors (X and y) and apply a scaler to the data.

The cleaned dataset can be found in medical_clean_numeric.csv

```
[3]: %%capture
    cat_columns = df.select_dtypes(exclude="number").columns

# Give categorical columns a numeric value
for col in cat_columns:
    df[col] = pd.Categorical(df[col])
    df[col] = df[col].cat.codes
```

```
# export
df.to_csv("data/medical_clean_numeric.csv", index=False)

# Separate into predictors and outcome

X = df.loc[1:, predictors]
y = df.loc[1:, outcome]

# apply scaler
scaler = StandardScaler()
scaler.fit(X * 1.0)
X = scaler.transform(X * 1.0)
```

3.3 Train test split

We'll also separating the data into training and testing sets on an 80/20 split. Copies of the train and test data can be found in train.csv and test.csv, respectively.

4 Analysis

Now that the data has been cleaned and split into train and test data we are ready to run a KNN Classifier!

We'll try the analysis using different values for k.

```
[5]: neighbors = range(2, 21)
    scores = np.empty(len(neighbors))

# Loop over different values of k
best_k = 0
best_score = -1
best_model = None
for i, k in enumerate(neighbors):
    #Create a k-NN classifier with k neighbors
    knn = KNeighborsClassifier(n_neighbors=k)

# Fit the classifier to the data
```

```
knn.fit(X_train, y_train)

# Score the classifier and add it the list of scores
score = knn.score(X_test, y_test)
scores[i] = score

if score > best_score:
    best_score = score
    best_k = k
    best_model = knn

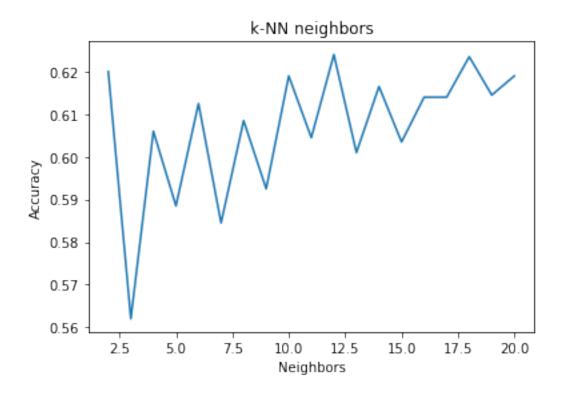
print('\nBest k-value is %i with a score of %.3f' % (best_k, best_score))

y_pred = best_model.predict(X_test)
auc_score = roc_auc_score(y_test, y_pred)
print('AUC score: %.3f' % auc_score)
```

Best k-value is 12 with a score of 0.624 AUC score: 0.505

We'll visualize the data with a graph as well.

```
[6]: plt.title('k-NN neighbors')
   plt.plot(neighbors, scores, label='Accuracy')
   plt.xlabel('Neighbors')
   plt.ylabel('Accuracy')
   plt.show()
```



5 Summary and Implications

5.1 Model effectiveness

The best version of the model we produces used a k-value of 12 and achieved an accuracy rating of 0.624 as well as an Area Under the Curve (AUC) score of .505. This means that the model is *slightly* better than random guessing.

The model's accuracy could use improvement. One limitation is the lack of complete patient medical records. Perhaps if the model could take into consideration more of the patient's medical history the accuracy could be increased.

At the moment the best course of action would be to gather more patient records to produce a better model.