

Brian_Reppeto_DSC550_Week_6

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0.0.1 DSC 550 Week :

Activity 6.2

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0.0.2 Milestone 1: Predicting Patient Readmissions for Diabetic Patients

Project Narrative

Hospital readmissions within 30 days post-discharge are not only a significant financial burden on the healthcare system but also often reflect suboptimal patient outcomes and potentially preventable complications. This project addresses the critical challenge of predicting such readmissions among diabetic patients, a group particularly prone to frequent and costly hospitalizations.

The Problem

The specific problem this project targets is the prediction of unplanned readmissions within 30 days among patients diagnosed with diabetes. These readmissions may be due to a variety of factors including inadequate management of diabetes, complications arising from the condition, or insufficient patient education and follow-up care upon discharge. The goal is to provide a predictive tool that can identify at-risk patients before they leave the hospital. This tool will enable healthcare providers to initiate targeted interventions such as personalized discharge planning, enhanced patient education, and tailored follow-up care schedules.

Objectives

The primary objective of this project is to develop a predictive model that uses historical hospital data to forecast the likelihood of a diabetic patient being readmitted within 30 days of discharge. The insights gained from this model will assist healthcare professionals in making informed decisions about patient care strategies and resource allocation.

Data Utilization

The project utilizes data from the Diabetes 130-US hospitals dataset, which comprises information from over 100,000 hospital admissions from 1999 to 2008 and across 130 US hospitals. The dataset includes diverse variables such as patient demographics, admission and discharge statuses, diagnostic codes, number of inpatient visits, and medication details. This rich dataset provides a comprehensive foundation to explore and model the complexities associated with readmissions.

Potential Impact

By accurately predicting readmissions, the model can directly influence the development of personalized medicine approaches and proactive healthcare strategies. Hospitals can use these predictions to reduce readmission rates, thereby decreasing the penalties they face under healthcare regulations like the Hospital Readmissions Reduction Program (HRRP). Moreover, patients benefit from improved healthcare experiences and outcomes, contributing to overall patient satisfaction and health system sustainability.

In conclusion, this project aims to harness the power of machine learning and predictive analytics to tackle a pressing healthcare issue. By doing so, it not only addresses an immediate business need for hospitals but also plays a crucial role in advancing how data-driven strategies can be implemented in clinical settings to enhance patient care.

```
[82]: # import libraries

import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler, LabelEncoder
from sklearn.linear_model import LogisticRegression
from sklearn.metrics import confusion_matrix, classification_report, \
    accuracy_score
```

```
[83]: # load the dataset

data = pd.read_csv('diabetic_data.csv')
```

```
[84]: # data shape

data.shape
```

```
[84]: (101766, 50)
```

```
[85]: # head the data

data.head(15)
```

```
[85]:
```

	encounter_id	patient_nbr	race	gender	age	weight	\
0	2278392	8222157	Caucasian	Female	[0-10)	?	
1	149190	55629189	Caucasian	Female	[10-20)	?	
2	64410	86047875	AfricanAmerican	Female	[20-30)	?	
3	500364	82442376	Caucasian	Male	[30-40)	?	
4	16680	42519267	Caucasian	Male	[40-50)	?	
5	35754	82637451	Caucasian	Male	[50-60)	?	
6	55842	84259809	Caucasian	Male	[60-70)	?	
7	63768	114882984	Caucasian	Male	[70-80)	?	
8	12522	48330783	Caucasian	Female	[80-90)	?	

9	15738	63555939	Caucasian	Female	[90-100)	?
10	28236	89869032	AfricanAmerican	Female	[40-50)	?
11	36900	77391171	AfricanAmerican	Male	[60-70)	?
12	40926	85504905	Caucasian	Female	[40-50)	?
13	42570	77586282	Caucasian	Male	[80-90)	?
14	62256	49726791	AfricanAmerican	Female	[60-70)	?

	admission_type_id	discharge_disposition_id	admission_source_id	\
0	6	25	1	
1	1	1	7	
2	1	1	7	
3	1	1	7	
4	1	1	7	
5	2	1	2	
6	3	1	2	
7	1	1	7	
8	2	1	4	
9	3	3	4	
10	1	1	7	
11	2	1	4	
12	1	3	7	
13	1	6	7	
14	3	1	2	

	time_in_hospital	...	citoglipton	insulin	glyburide-metformin	\
0	1	...	No	No	No	
1	3	...	No	Up	No	
2	2	...	No	No	No	
3	2	...	No	Up	No	
4	1	...	No	Steady	No	
5	3	...	No	Steady	No	
6	4	...	No	Steady	No	
7	5	...	No	No	No	
8	13	...	No	Steady	No	
9	12	...	No	Steady	No	
10	9	...	No	Steady	No	
11	7	...	No	Steady	No	
12	7	...	No	Down	No	
13	10	...	No	Steady	No	
14	1	...	No	Steady	No	

	glipizide-metformin	glimepiride-pioglitazone	metformin-rosiglitazone	\
0	No	No	No	
1	No	No	No	
2	No	No	No	
3	No	No	No	
4	No	No	No	

5	No	No	No
6	No	No	No
7	No	No	No
8	No	No	No
9	No	No	No
10	No	No	No
11	No	No	No
12	No	No	No
13	No	No	No
14	No	No	No

	metformin-pioglitazone	change	diabetesMed	readmitted
0	No	No	No	NO
1	No	Ch	Yes	>30
2	No	No	Yes	NO
3	No	Ch	Yes	NO
4	No	Ch	Yes	NO
5	No	No	Yes	>30
6	No	Ch	Yes	NO
7	No	No	Yes	>30
8	No	Ch	Yes	NO
9	No	Ch	Yes	NO
10	No	No	Yes	>30
11	No	Ch	Yes	<30
12	No	Ch	Yes	<30
13	No	No	Yes	NO
14	No	No	Yes	>30

[15 rows x 50 columns]

0.0.3 Data Exploration and Cleaning

```
[86]: # explore missing values and clean data

data = data.replace('?', np.nan) # replace '?' with NaN for clarity
missing_data = data.isnull().sum()/len(data) * 100
print("Percentage of missing data per column:\n", missing_data[missing_data > 0])
```

```
Percentage of missing data per column:
race                2.233555
weight              96.858479
payer_code          39.557416
medical_specialty   49.082208
diag_1              0.020636
diag_2              0.351787
diag_3              1.398306
max_glu_serum       94.746772
```

```
A1Cresult          83.277322
dtype: float64
```

```
[87]: # drop columns with high percentage of missing values and those not relevant
      ↪for the analysis
```

```
columns_to_drop = ['weight', 'medical_specialty', 'payer_code']
data.drop(columns=columns_to_drop, inplace=True)
```

```
[89]: # head the data
```

```
data.head(15)
```

```
[89]:
```

	encounter_id	patient_nbr	race	gender	age	\
0	2278392	8222157	Caucasian	Female	[0-10)	
1	149190	55629189	Caucasian	Female	[10-20)	
2	64410	86047875	AfricanAmerican	Female	[20-30)	
3	500364	82442376	Caucasian	Male	[30-40)	
4	16680	42519267	Caucasian	Male	[40-50)	
5	35754	82637451	Caucasian	Male	[50-60)	
6	55842	84259809	Caucasian	Male	[60-70)	
7	63768	114882984	Caucasian	Male	[70-80)	
8	12522	48330783	Caucasian	Female	[80-90)	
9	15738	63555939	Caucasian	Female	[90-100)	
10	28236	89869032	AfricanAmerican	Female	[40-50)	
11	36900	77391171	AfricanAmerican	Male	[60-70)	
12	40926	85504905	Caucasian	Female	[40-50)	
13	42570	77586282	Caucasian	Male	[80-90)	
14	62256	49726791	AfricanAmerican	Female	[60-70)	

	admission_type_id	discharge_disposition_id	admission_source_id	\
0	6	25	1	
1	1	1	7	
2	1	1	7	
3	1	1	7	
4	1	1	7	
5	2	1	2	
6	3	1	2	
7	1	1	7	
8	2	1	4	
9	3	3	4	
10	1	1	7	
11	2	1	4	
12	1	3	7	
13	1	6	7	
14	3	1	2	

	time_in_hospital	num_lab_procedures	...	citoglipton	insulin	\
0	1	41	...	No	No	
1	3	59	...	No	Up	
2	2	11	...	No	No	
3	2	44	...	No	Up	
4	1	51	...	No	Steady	
5	3	31	...	No	Steady	
6	4	70	...	No	Steady	
7	5	73	...	No	No	
8	13	68	...	No	Steady	
9	12	33	...	No	Steady	
10	9	47	...	No	Steady	
11	7	62	...	No	Steady	
12	7	60	...	No	Down	
13	10	55	...	No	Steady	
14	1	49	...	No	Steady	

	glyburide-metformin	glipizide-metformin	glimepiride-pioglitazone	\
0	No	No	No	
1	No	No	No	
2	No	No	No	
3	No	No	No	
4	No	No	No	
5	No	No	No	
6	No	No	No	
7	No	No	No	
8	No	No	No	
9	No	No	No	
10	No	No	No	
11	No	No	No	
12	No	No	No	
13	No	No	No	
14	No	No	No	

	metformin-rosiglitazone	metformin-pioglitazone	change	diabetesMed	\
0	No	No	No	No	
1	No	No	Ch	Yes	
2	No	No	No	Yes	
3	No	No	Ch	Yes	
4	No	No	Ch	Yes	
5	No	No	No	Yes	
6	No	No	Ch	Yes	
7	No	No	No	Yes	
8	No	No	Ch	Yes	
9	No	No	Ch	Yes	
10	No	No	No	Yes	
11	No	No	Ch	Yes	

12	No	No	Ch	Yes
13	No	No	No	Yes
14	No	No	No	Yes

```

    readmitted
0          NO
1         >30
2          NO
3          NO
4          NO
5         >30
6          NO
7         >30
8          NO
9          NO
10         >30
11         <30
12         <30
13          NO
14         >30

```

[15 rows x 47 columns]

[90]: *# summary of cleaned data*

```
print(data.info())
```

```

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 101766 entries, 0 to 101765
Data columns (total 47 columns):
#   Column                                Non-Null Count  Dtype
---  -
0   encounter_id                          101766 non-null int64
1   patient_nbr                          101766 non-null int64
2   race                                 99493 non-null  object
3   gender                               101766 non-null object
4   age                                  101766 non-null object
5   admission_type_id                    101766 non-null int64
6   discharge_disposition_id             101766 non-null int64
7   admission_source_id                  101766 non-null int64
8   time_in_hospital                     101766 non-null int64
9   num_lab_procedures                   101766 non-null int64
10  num_procedures                        101766 non-null int64
11  num_medications                       101766 non-null int64
12  number_outpatient                     101766 non-null int64
13  number_emergency                      101766 non-null int64
14  number_inpatient                      101766 non-null int64
15  diag_1                                101745 non-null object

```

```

16 diag_2          101408 non-null object
17 diag_3          100343 non-null object
18 number_diagnoses 101766 non-null int64
19 max_glu_serum    5346 non-null object
20 A1Cresult        17018 non-null object
21 metformin        101766 non-null object
22 repaglinide      101766 non-null object
23 nateglinide      101766 non-null object
24 chlorpropamide   101766 non-null object
25 glimepiride      101766 non-null object
26 acetohexamide    101766 non-null object
27 glipizide        101766 non-null object
28 glyburide        101766 non-null object
29 tolbutamide      101766 non-null object
30 pioglitazone     101766 non-null object
31 rosiglitazone    101766 non-null object
32 acarbose         101766 non-null object
33 miglitol         101766 non-null object
34 troglitazone     101766 non-null object
35 tolazamide       101766 non-null object
36 examide          101766 non-null object
37 citoglipton      101766 non-null object
38 insulin          101766 non-null object
39 glyburide-metformin 101766 non-null object
40 glipizide-metformin 101766 non-null object
41 glimepiride-pioglitazone 101766 non-null object
42 metformin-rosiglitazone 101766 non-null object
43 metformin-pioglitazone 101766 non-null object
44 change           101766 non-null object
45 diabetesMed      101766 non-null object
46 readmitted       101766 non-null object
dtypes: int64(13), object(34)
memory usage: 36.5+ MB
None

```

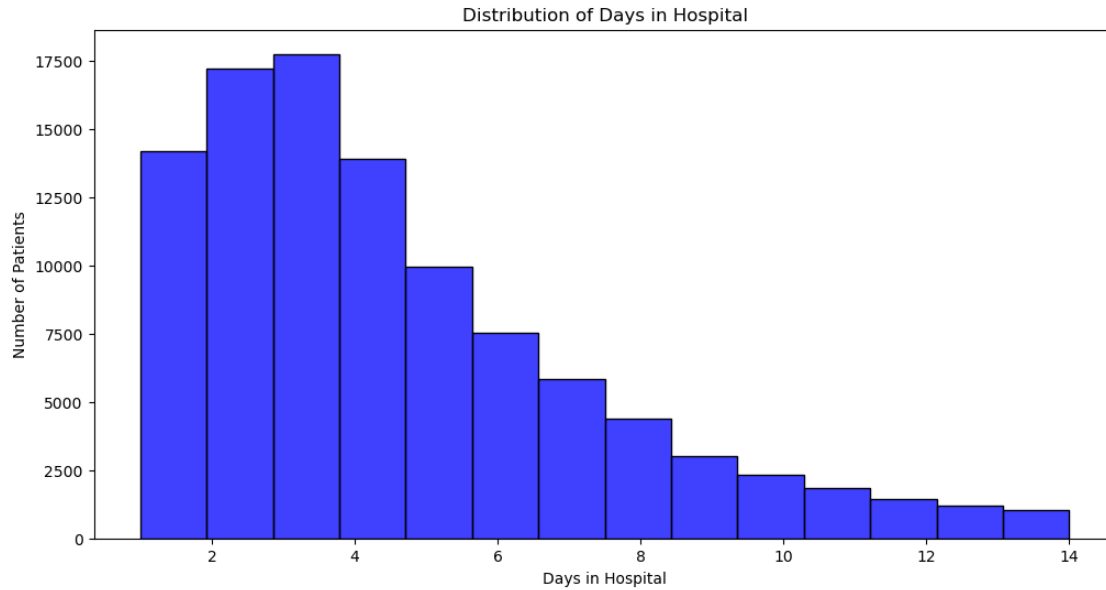
0.0.4 Graphical Analysis

```

[91]: # histogram of Distribution of the number of days in hospital

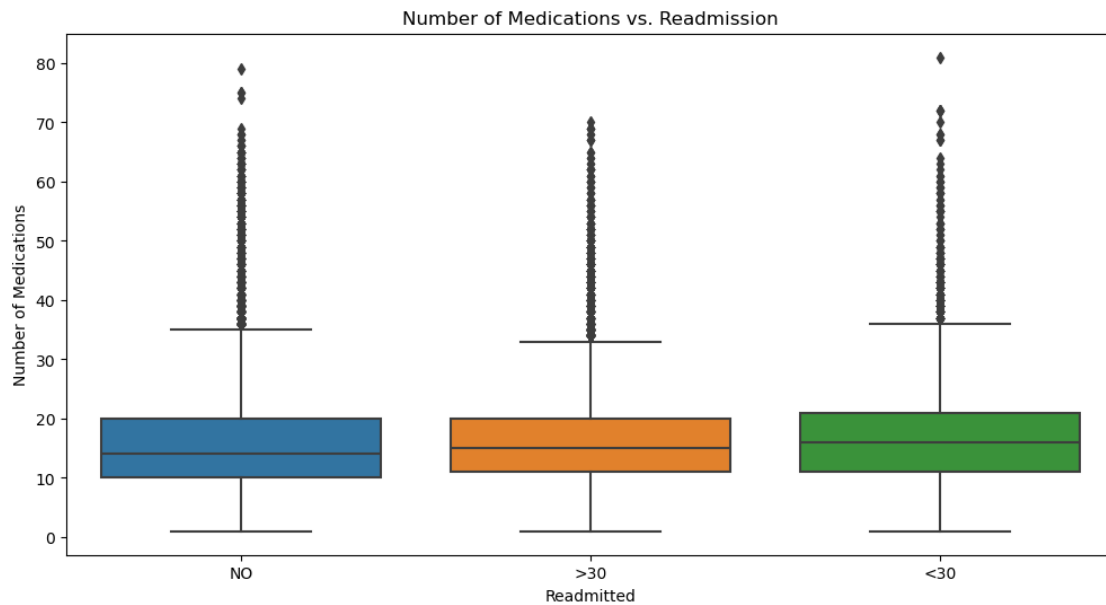
plt.figure(figsize=(12, 6))
sns.histplot(data['time_in_hospital'], bins=14, kde=False, color='blue')
plt.title('Distribution of Days in Hospital')
plt.xlabel('Days in Hospital')
plt.ylabel('Number of Patients')
plt.show()

```

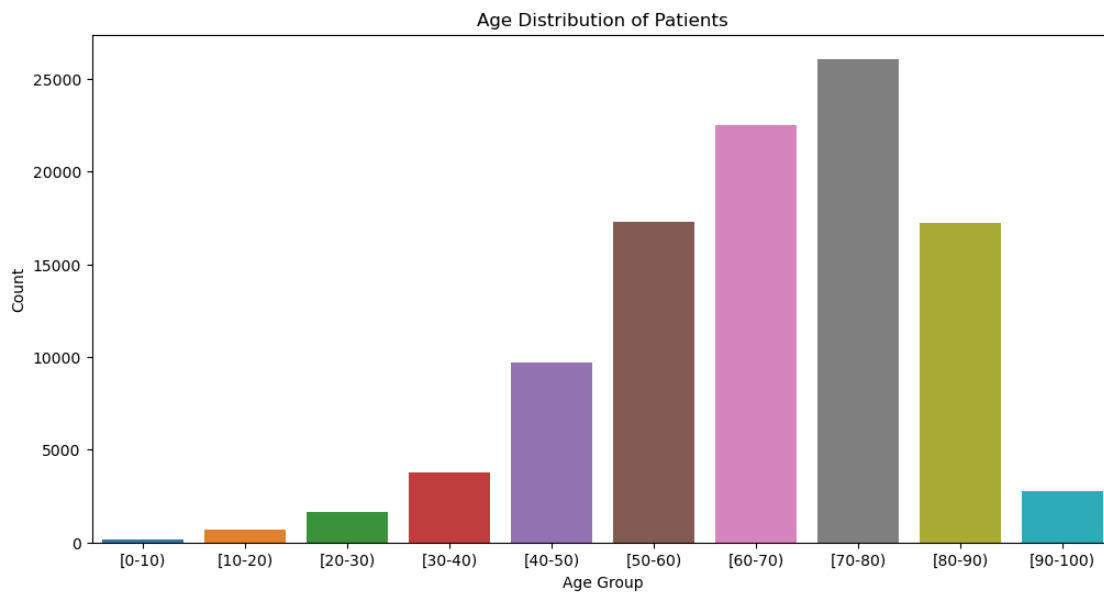
[92]: *# number of medications vs. readmissions*

```
plt.figure(figsize=(12, 6))
sns.boxplot(x='readmitted', y='num_medications', data=data)
plt.title('Number of Medications vs. Readmission')
plt.xlabel('Readmitted')
plt.ylabel('Number of Medications')
plt.show()
```



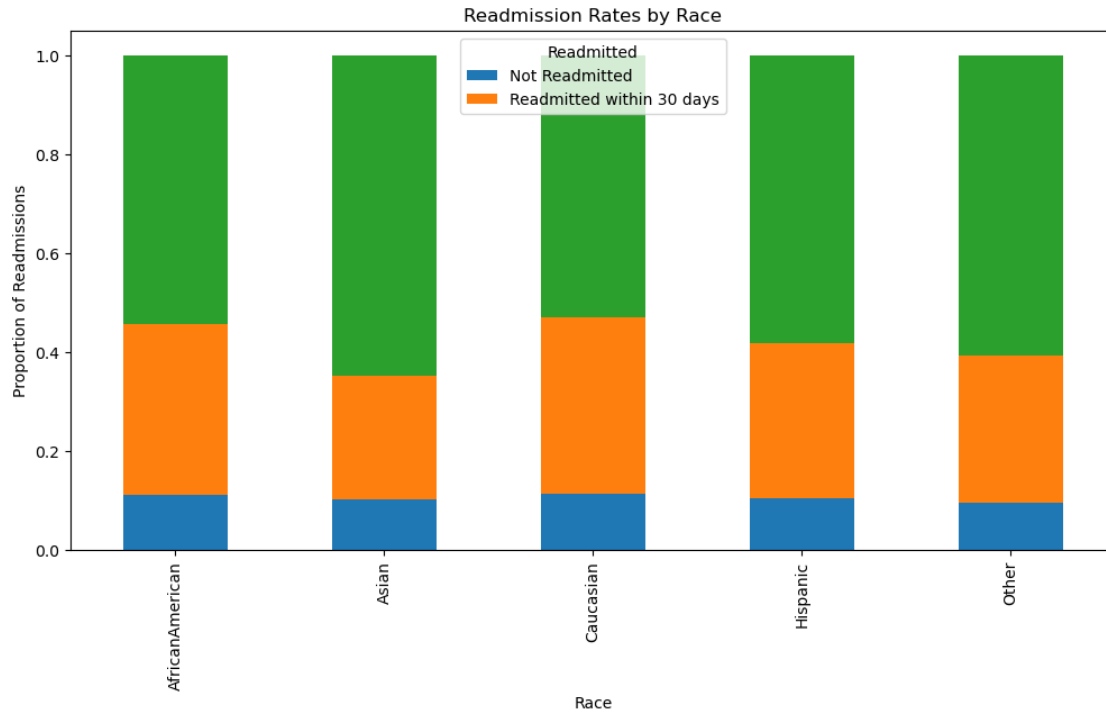
```
[93]: # age distribution of the patients
```

```
plt.figure(figsize=(12, 6))
sns.countplot(x='age', data=data, order=sorted(data['age'].unique()))
plt.title('Age Distribution of Patients')
plt.xlabel('Age Group')
plt.ylabel('Count')
plt.show()
```



```
[94]: # readmission rates by race
```

```
readmission_by_race = data.groupby('race')['readmitted'].  
    ↳value_counts(normalize=True).unstack().fillna(0)  
readmission_by_race.plot(kind='bar', stacked=True, figsize=(12, 6))  
plt.title('Readmission Rates by Race')  
plt.xlabel('Race')  
plt.ylabel('Proportion of Readmissions')  
plt.legend(title='Readmitted', labels=['Not Readmitted', 'Readmitted within 30_↳  
    ↳days'])  
plt.show()
```



0.0.5 Analysis of Graphs

Distribution of Days in Hospital:

The histogram shows that the most common duration of hospital stays is between 2 to 4 days. The distribution is right-skewed, indicating that longer stays are less frequent but not uncommon. This suggests that most diabetic patients have relatively short hospital stays, but a subset requires extended care.

Number of Medications vs. Readmission:

From the boxplot comparing the number of medications between readmitted and not readmitted groups, there is a noticeable overlap, but it seems that patients who were readmitted tend to be on slightly more medications. This could imply that patients with more complex medication schedules are at a higher risk of readmission, possibly due to more severe underlying conditions.

Age Distribution of Patients:

The age distribution shows that the majority of the patients fall into the 60-80 age range, with fewer younger patients. This is typical for diabetic cohorts where prevalence increases with age.

Readmission Rates by Race:

The bar chart demonstrates that readmission rates vary somewhat by race. The proportions show that certain racial groups might have higher or lower rates of readmission, which could be important for targeted interventions or understanding disparities in healthcare outcomes.

0.0.6 Conclusion

The graphical analysis provided insights into factors that might influence hospital readmission among diabetic patients. The analysis suggests that duration of hospital stay, complexity of medication regimens, patient age, and race could be significant predictors of readmission risk.

[]: