



Supervised Learning Capstone

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Background

I've acquired a Diabetic Readmission Data Set from kaggle. The dataset is composed of integrated data from 130 U.S. hospitals over the course of 10 years. It includes over 50 features such as length of stay, medications, lab results and primary diagnosis.

In the healthcare field, determining the chances of a patient readmission is vital. If risks for readmission is known, better treatment plan can be created for patients.

Objective

To create various models to determine diabetic patient readmission prediction less than 30 days of previous admission.

Exploratory Data Analysis

- ❖ Categorical values includes gender, race, age (which were group into counts of 10), medications, diagnosis via ICD9 codes, medical specialty of admitting doctor, medication changes and more.
- ❖ Numerical columns included patient identifiers such as encounter id and patient number, admission type, number of times a patient was inpatient, outpatient, in emergency, procedures, time in hospital and number of medications.

Object Types

We have a mixture of 37 categorical values and 13 numerical/continuous values:

1 num_cols.dtypes

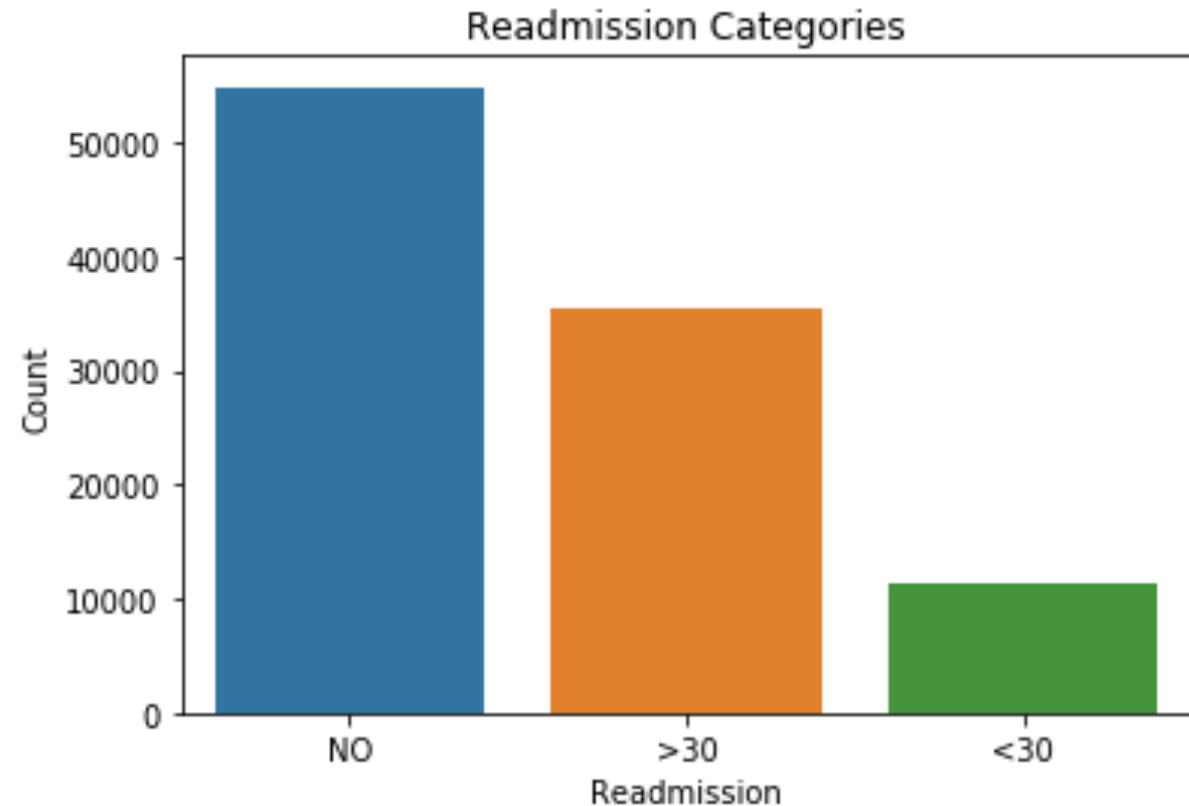
encounter_id	int64
patient_nbr	int64
admission_type_id	int64
discharge_disposition_id	int64
admission_source_id	int64
time_in_hospital	int64
num_lab_procedures	int64
num_procedures	int64
num_medications	int64
number_outpatient	int64
number_emergency	int64
number_inpatient	int64
number_diagnoses	int64
target	int64
age_group	int64
dtype:	object

3 cat_cols.dtypes

race	object
gender	object
age	object
diag_1	object
diag_2	object
diag_3	object
max_glu_serum	object
AlCresult	object
metformin	object
repaglinide	object
nateglinide	object
chlorpropamide	object
glimepiride	object
acetohehexamide	object
glipizide	object
glyburide	object
tolbutamide	object
pioglitazone	object
rosiglitazone	object
acarbose	object
miglitol	object
troglitazone	object
tolazamide	object
examide	object
citoglipton	object
insulin	object
glyburide-metformin	object
glipizide-metformin	object
glimepiride-pioglitazone	object
metformin-rosiglitazone	object
metformin-pioglitazone	object
change	object
diabetesMed	object
readmitted	object
dtype:	object

Readmission Categories

- The target chosen for this project is to predict readmission in less than 30 days after previous admission
- It is worth noting that the readmission > 30 days does not specify exactly when patient was readmitted.



NO	52527
>30	35502
<30	11314

Looking for missing values

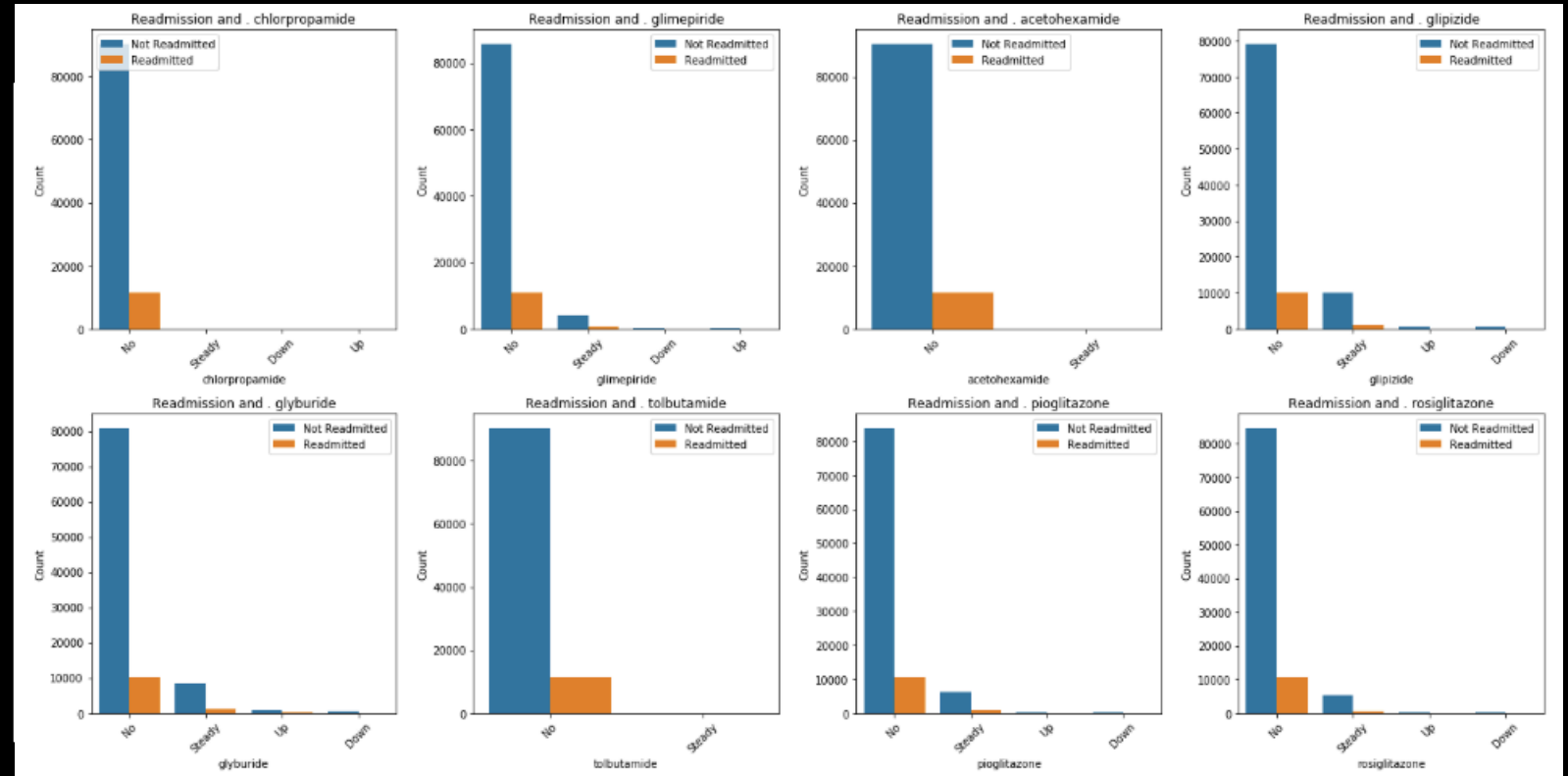
The dataset contained no null values upon first examination. However, it had many '?' values that represent missing values.

Columns with over 90% missing were dropped. Other columns that provide no insight such as insurance payer codes were dropped as well.

```
1 #looking for ? values.
2 for col in diabetes.columns:
3     if diabetes[col].dtype == object:
4         print(col, diabetes[col][diabetes[col] == '?'].count())

race 2273
gender 0
age 0
weight 98569
payer_code 40256
medical_specialty 49949
diag_1 21
diag_2 358
diag_3 1423
max_glu_serum 0
AlCresult 0
metformin 0
repaglinide 0
nateglinide 0
chlorpropamide 0
..
..
```

There were many medications (27) included in the dataset. Visualization helped determine relationship between target.



Feature Engineering

```
{ '[0-10)' : 0,  
  '[10-20)' : 10,  
  '[20-30)' : 20,  
  '[30-40)' : 30,  
  '[40-50)' : 40,  
  '[50-60)' : 50,  
  '[60-70)' : 60,  
  '[70-80)' : 70,  
  '[80-90)' : 80,  
  '[90-100)' : 90}  
'age_group'] = diabetes
```

- New feature was created for age since it was grouped by 10s. This in turn created a new numerical value for age.
- With discharge dispositions, patients who have expired or was sent to hospice were dropped because there will be no chance for readmission.

Grouped diagnosis by counts/occurrence

	Diagnosis	ICD_code	count
0	Heart Failure	428	6862
1	Other forms of Chronic Heart Disease	414	6581
2	Symptoms involving respiratory system and othe...	786	4016
3	Myocardial infarction	410	3614
4	Pneumonia	486	3508
5	Cardiac Dysrhythmias	427	2766
6	Emphysema	491	2275
7	Osteoarthritis	715	2151
8	Cellulitis	682	2042
9	General Symptoms	434	2028

	2nd_diagnosis	ICD_code	count
0	Disorders of fluid electrolyte and acid-base b...	276	6752
1	Heart Failure	428	6662
2	Diabetes mellitus without mention of complicat...	250	6071
3	Cardiac Dysrhythmias	427	5036
4	Essential Hypertension	401	3736
5	Chronic airway obstruction, not elsewhere clas...	496	3305
6	Other disorders of urethra and urinary tract	599	3288
7	Hypertensive chronic kidney disease	403	2823
8	Other forms of chronic ischemic heart disease	414	2650
9	Other acute and subacute forms of ischemic hea...	411	2566

	Other_diagnosis	ICD_code	count
0	Diabetes mellitus without mention of complicat...	250	11555
1	Essential Hypertension/t	401	8289
2	Disorders of fluid electrolyte and acid-base ...	276	5175
3	Heart Failure	428	4577
4	Cardiac Dysrhythmias	427	3955
5	Other forms of chronic ischemic heart disease	414	3664
6	Chronic airway obstruction, not elsewhere clas...	496	2605
7	Hypertensive chronic kidney disease	403	2357
8	Chronic Kidney Disease	585	1992
9	Disorders of lipid metabolism	272	1969

Chi-squared Feature Importance

Was able to reduce features that were not
Important using Chi-squared.

```
[ 'discharge_disposition_id_22',  
  'discharge_disposition_id_3',  
  'number_diagnoses',  
  'number_inpatient',  
  'number_emergency',
```

Chi-Squared Top 5 features

Have tried the following to address class imbalance:

Oversampling using SMOTE Technique

Undersampling

Class Imbalance

SVC

TRAIN

VALIDATION

AUC:0.672

AUC:0.668

accuracy:0.622

accuracy:0.685

recall:0.527

recall:0.542

precision:0.650

precision:0.189

specificity:0.693

specificity:0.682

F1:0.582

F1:0.281

UNDERSAMPLING

TRAIN

VALIDATION

AUC:0.94

AUC:0.56

accuracy:0.88

accuracy:0.689

recall:0.882

recall:0.345

precision:0.91

precision:0.112

specificity:0.77

specificity:0.712

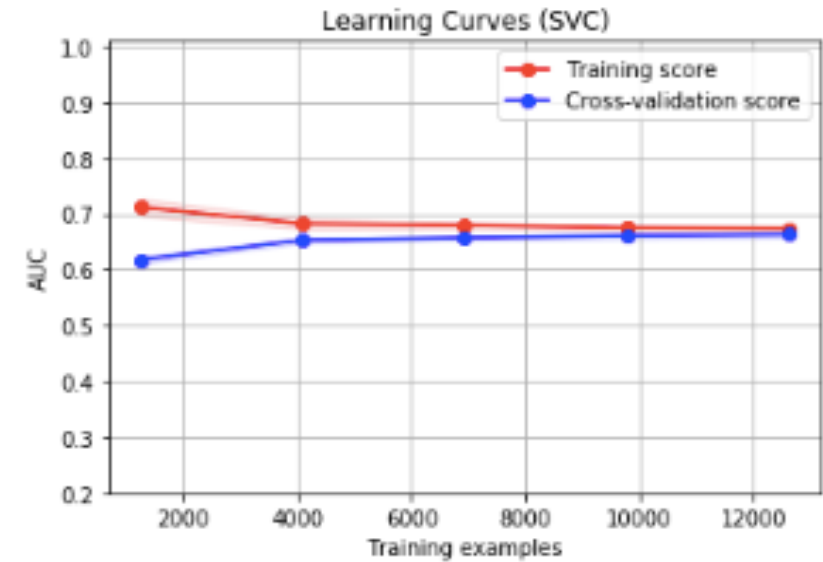
f1:0.89

f1:0.131

SMOTE

Support Vector Classifier

- Longest run times.
- Tuned C and gamma parameter without much improvement.



```
Baseline SVC
Training AUC:0.672
Validation AUC:0.668
Optimized SVC
Training AUC:0.672
Validation AUC:0.668
```

KNN

TRAIN

VALIDATION

AUC:0.652

AUC:0.624

accuracy:0.605

accuracy:0.650

recall:0.518

recall:0.504

precision:0.627

precision:0.163 s

specificity:0.658

specificity:0.637

F1:0.567

F1:0.246

UNDERSAMPLING

TRAIN

VALIDATION

AUC:0.923

AUC:0.551

accuracy:0.850

accuracy:0.672

recall:0.882

recall:0.333

precision:0.901

precision:0.131

specificity:0.768

specificity:0.706

f1:0.88

f1:0.08

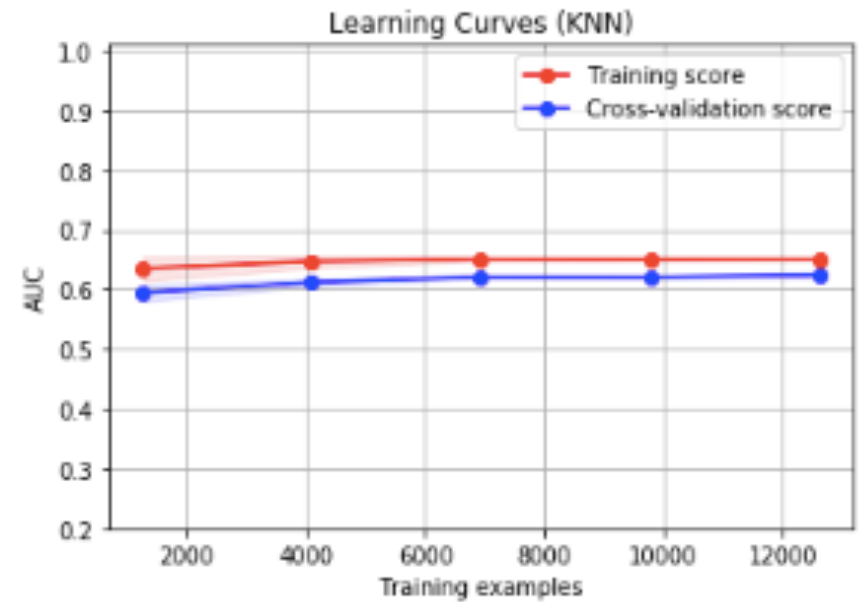
SMOTE

KNN

Moderate running times.

Tuned by increasing `n_neighbors`, using the minkowski metric, and adding uniform weight.

```
Baseline KNN
Training AUC:0.652
Validation AUC:0.624
Optimized KNN
Training AUC:0.649
Validation AUC:0.629
```



Logistic Regression

TRAIN

VALIDATION

AUC:0.675

AUC:0.667

accuracy:0.623

accuracy:0.668

recall:0.550

recall:0.572

precision:0.644

precision:0.186

specificity:0.696

specificity:0.681

F1: 0.59

F1: 0.28

UNDERSAMPLING

TRAIN

VALIDATION

AUC:0.908

AUC:0.543

accuracy:0.844

accuracy:0.773

recall:0.833

recall:0.188

precision:0.851

precision:0.138

specificity:0.855

specificity:0.849

F1:0.891

F1:0.188

SMOTE

Logistic Regression

An advantage is model is interpretability. Has feature importance

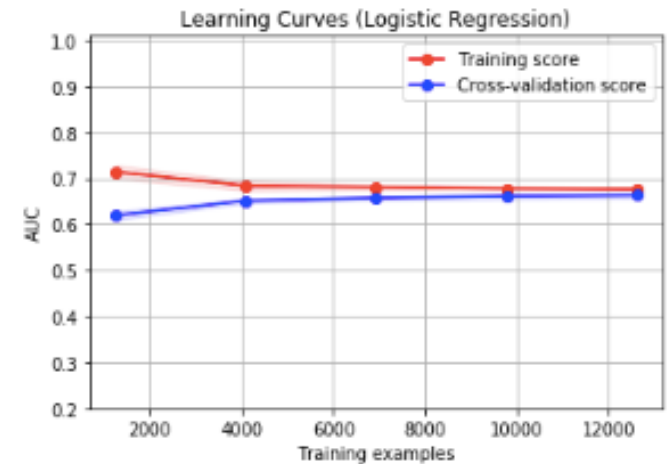
Fast training times.

Tuned by reducing C to 0.1, created l1 penalty, class weight was balanced.

Scores were not much different.

```
1 feature_importances.head(10)
```

	importance
number_inpatient	0.357715
discharge_disposition_id_22	0.188475
rosiglitazone_No	0.180301
repaglinide_No	0.173352
repaglinide_Steady	0.156140
rosiglitazone_Steady	0.134689
diabetesMed_Yes	0.120780
discharge_disposition_id_3	0.119310
discharge_disposition_id_28	0.110808
discharge_disposition_id_5	0.109521



Baseline Logistic Regression
Training AUC:0.675
Validation AUC:0.667

Optimized Logistic Regression
Training AUC:0.675
Validation AUC:0.668

Decision Trees

TRAIN

VALIDATION

AUC:0.729

AUC:0.637

accuracy:0.665

accuracy:0.664

recall:0.590

recall:0.539

precision:0.693

precision:0.177

specificity:0.738

specificity:0.679

F1: 0.63

F1: 0.26

UNDERSAMPLING

TRAIN

VALIDATION

AUC:0.848

AUC:0.538

accuracy:0.768

accuracy:0.757

recall:0.703

recall:0.216

precision:0.809

precision:0.138

specificity:0.833

specificity:0.826

F1:0.76

F1:0.11

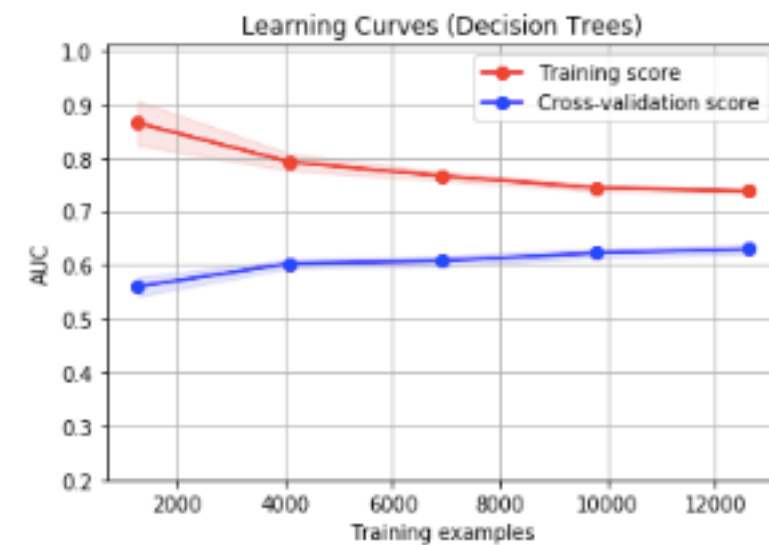
SMOTE

Decision Trees

Has tendency to overfit.

Decided not to tune this model.

Moderate run time.



Random Forest

TRAIN

VALIDATION

AUC:0.671

AUC:0.635

accuracy:0.624

accuracy:0.618

recall:0.586

recall:0.587

precision:0.634

precision:0.165

specificity:0.661

specificity:0.622

F1: 0.61

F1: 0.26

UNDERSAMPLING

TRAIN

VALIDATION

AUC:0.839

AUC:0.536

accuracy:0.760

accuracy:0.698

recall:0.761

recall:0.279

precision:0.759

precision:0.126

specificity:0.759

specificity:0.752

F1:0.891

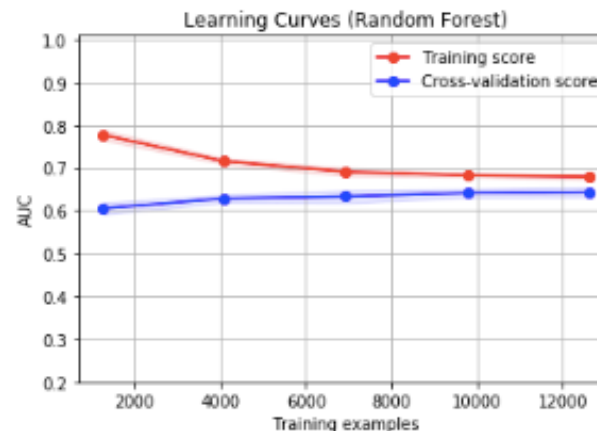
F1:0.188

SMOTE

Random Forest

- Has feature importance.
- Tuned by increasing number of estimators and depth.
- Moderate time training.
- Optimized score improved in training, but seems overfitted in validation set.

Baseline Random Forest
Training AUC:0.671
Validation AUC:0.635
Optimized Random Forest
Training AUC:0.713
Validation AUC:0.662



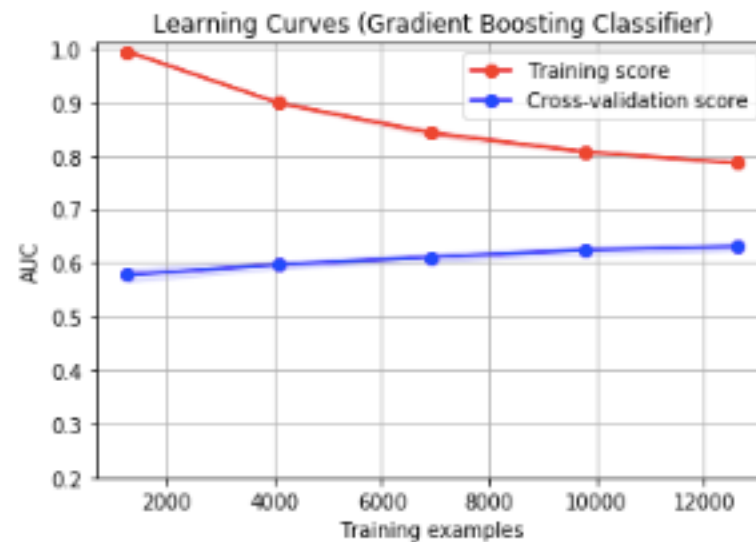
	importance
number_inpatient	0.183701
time_in_hospital	0.098817
number_emergency	0.093810
discharge_disposition_id_22	0.077494
num_medications	0.057466
num_lab_procedures	0.052790
number_diagnoses	0.045715
number_outpatient	0.028754
number_outpatient	0.023830
insulin_No	0.023613

Gradient Boosting Classifier

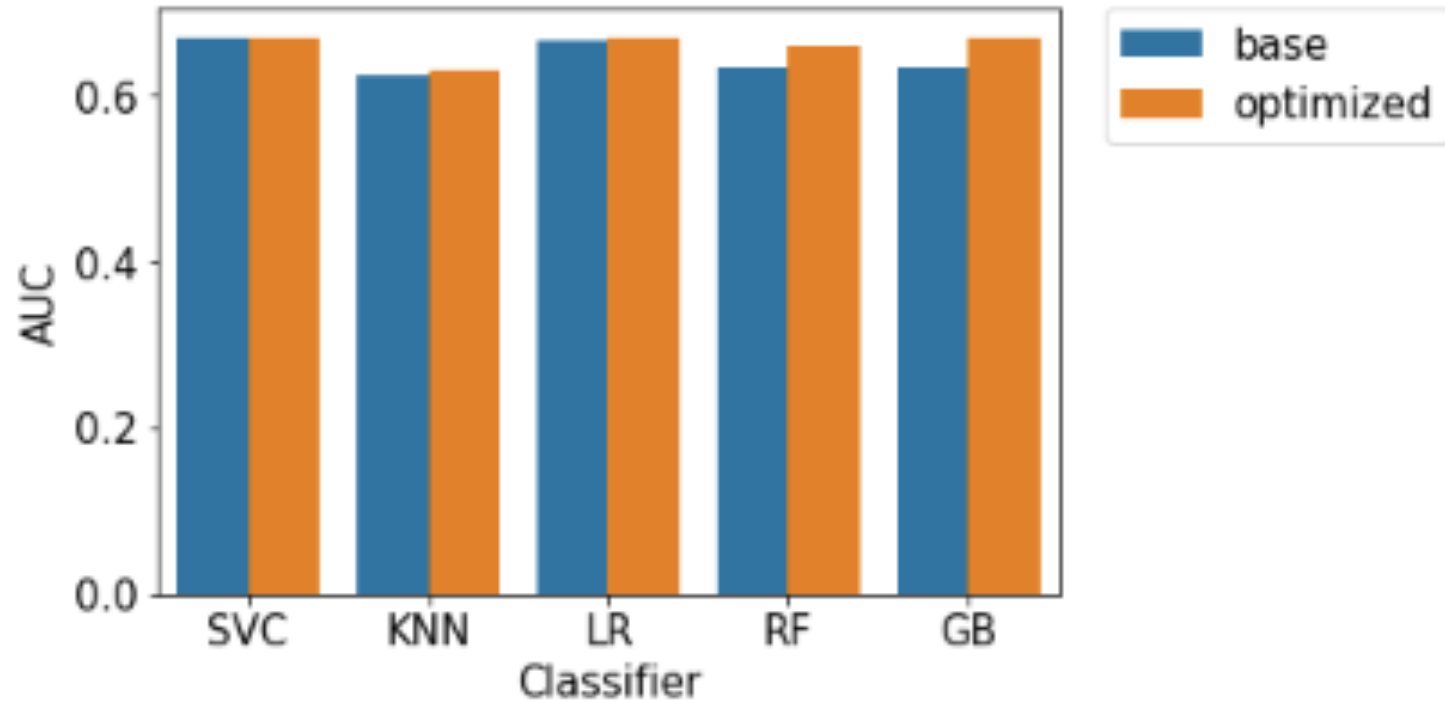
TRAIN	VALIDATION	TRAIN	VALIDATION
AUC:0.770	AUC:0.632	AUC:0.926	AUC:0.553
accuracy:0.694	accuracy:0.614	accuracy:0.861	accuracy:0.793
recall:0.670	recall:0.586	recall:0.839	recall:0.186
precision:0.704	precision:0.163	precision:0.878	precision:0.157
specificity:0.718	specificity:0.618	specificity:0.884	specificity:0.871
F1: 0.68	F1: 0.25	F1: 0.86	F1. 0.116
UNDERSAMPLING		SMOTE	

Gradient Boosting Classifier

- Has the best AUC scores for training and validation set.
- Moderate run time.



```
Baseline Gradient Boosting Classifier
Training AUC:0.770
Validation AUC:0.632
Optimized GBC
Training AUC:0.702
Validation AUC:0.670
```

Hyperparameter Tuning Results

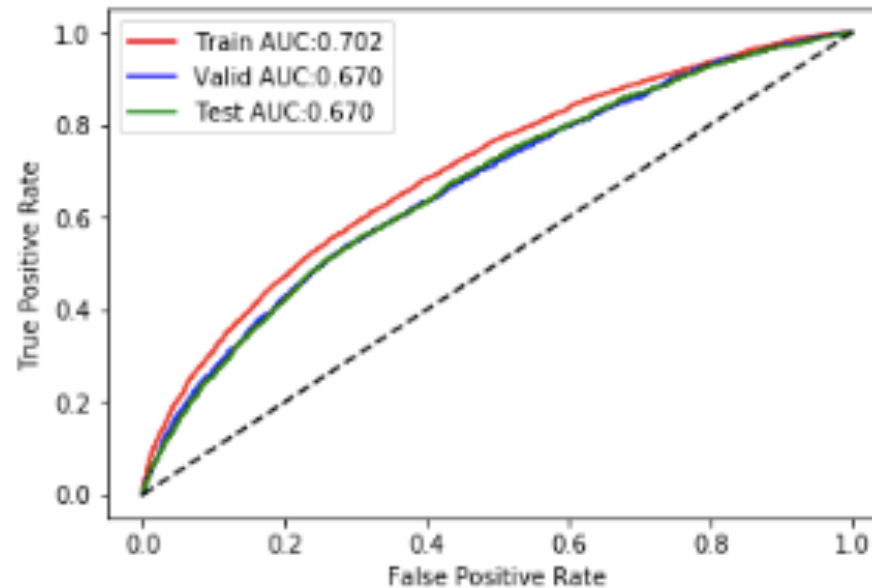
Training:
AUC:0.702
accuracy:0.644
recall:0.593
precision:0.660
specificity:0.694
prevalence:0.500

Validation:
AUC:0.670
accuracy:0.651
recall:0.584
precision:0.179
specificity:0.660
prevalence:0.113

Test:
AUC:0.670
accuracy:0.644
recall:0.590
precision:0.183
specificity:0.651
prevalence:0.117

gbc f1:0.280

Model Selection: GBC



Conclusion

What have we learned from exploring this dataset?

1. There is a correlation between number of inpatient visits and being readmitted less than 30 days.
2. A patient being discharged to the rehab or a subacute facility has a higher chance of readmission in less than 30 days.
3. Since many patients have a primary diagnosis of heart related conditions, it is worth looking at studying readmission rates for this population. Diabetes and heart disease have a known correlation, how does this affect readmission rates?
4. If the intention is to truly predict readmission for diabetic patients, it may be helpful to look at diabetes as a primary diagnosis. According to the ICD 10, primary diagnosis requires the most serious attention and is resource intensive while secondary and tertiary diagnosis could be diseases that co-exist during admission or develop thereafter admission.
5. Additional information may be needed for this dataset. Information such as procedures and certain blood work could provide more insight into readmission.

Reference:

Dataset acquired from: <https://www.kaggle.com/brandao/diabetes>

ICD 10: <https://www.icd10watch.com/blog/clearing-confusion-between-principal-and-primary-diagnoses>

ICD 9 Codes : <http://www.icd9data.com/>