

1. Importing Libraries

```
import pandas as pd
import numpy as np
import seaborn as sns
from sklearn.preprocessing import LabelEncoder
import warnings
warnings.filterwarnings('ignore')
import matplotlib.pyplot as plt
import seaborn as sns
pd.set_option("display.max_rows", None, "display.max_columns", None)
```

2. Importing datasets

```
train = pd.read_csv("SPARC_S194_2016_SelectedHospitals.csv", parse_dates=["Discharge Year"])
```

User Defined Functions

```
def quality_report(df):
    """Description: Displays quality of data in terms of missing values,
    unique numbers, datatypes etc.

    Arguments: Dataframe

    """
    dtypes = df.dtypes
    nunique = df.apply(lambda x: x.nunique(), axis=1)
    total = df.isnull().sum().sort_values(ascending=False)
    percent = (df.isnull().sum()/df.isnull().count()*100).sort_values(ascending=False)
    quality_df = pd.concat([total, percent, nunique, dtypes], axis=1, keys=["Total NaN", "Percent of NaN", "Nunique", "display(quality_df)"])

def object_count_plot(df):
    """Description: Plot countplot for all categorical features present in the dataframe passed Argument : Dataframe
    for var in df.columns:
        if df[var].dtype == 'object':
            print(df[col].describe())
            plt.figure(figsize=(12,5))
            g = sns.countplot(x=var,data=df)
            ax = sns.distplot(df[col].dropna(), rotation=90, ha="right")
            plt.figure(figsize=(12,5))
            plt.show()

def numeric_distribution_plot(df):
    """Description: Gives distribution plot for all the numeric features in the dataframe passed Argument : Dataframe
    for col in df.columns:
        if df[col].dtype != 'object':
            print(df[col].describe())
            plt.figure(figsize=(12,5))
            plt.title("Distribution of "+col)
            ax = sns.distplot(df[col].dropna())
            plt.tight_layout()
            plt.show()
            value_counts = df[col].value_counts().rename_axis("unique_"+col).to_frame('counts').sort_values('counts', ascending=False)
            print(value_counts.head(5), "\n", value_counts.tail(5))
```

Visualising the data set

	Facility ID	Facility Name	Age Group	Zip Code - 3 digits	Gender	Race	Ethnicity	Length of Stay	Admit Day of Week	Type of Admission	Patient Disposition	Discharge	Discharge Year	Discharge Day of Week	CCS Diagnosis Code
0	66	Olean General Hospital	70 or Older	147	M	White	Span/Hispanic	Not	2	None	Emergency	Home or Self Care	2016-01-01	None	108
1	66	Olean General Hospital	50 to 69	147	F	White	Span/Hispanic	Not	5	None	Emergency	Inpatient Rehabilitation Facility	2016-01-01	None	108

3. Quick EDA

```
quality_report(train)
train.shape

#these are 5 missing values in the zip code field.
#dropping these
train.dropna(axis = 0,inplace=True)
train.shape

#dropping redundant columns i.e columns that give little new information
train.drop(['Facility Name', 'APR MDC Code', 'APR DRG Description', 'APR DRG Code', 'Discharge Day of Week', 'Discharge Year', 'Admit Day of Week', 'APR Medical Surgical Description', 'APR Risk of Mortality', 'Birth Weight'],axis=1, inplace = True)
train.shape

(4744, 22)
```

```
#the attributes reduced to 22 after dropping low information columns
```

```
train.columns
```

```
Index(['Facility ID', 'Age Group', 'Zip Code - 3 digits', 'Gender', 'Race', 'Ethnicity', 'Length of Stay', 'Type of Admission', 'Patient Disposition', 'CCS Diagnosis Code', 'CCS Procedure Code', 'APR DRG Description', 'APR DRG Code', 'APR MDC Description', 'APR Severity of Illness Code', 'APR Risk of Mortality', 'Source of Payment', 'Emergency Department Indicator', 'Total Charges', 'Total Costs'], dtype='object')
```

Univariate Analysis

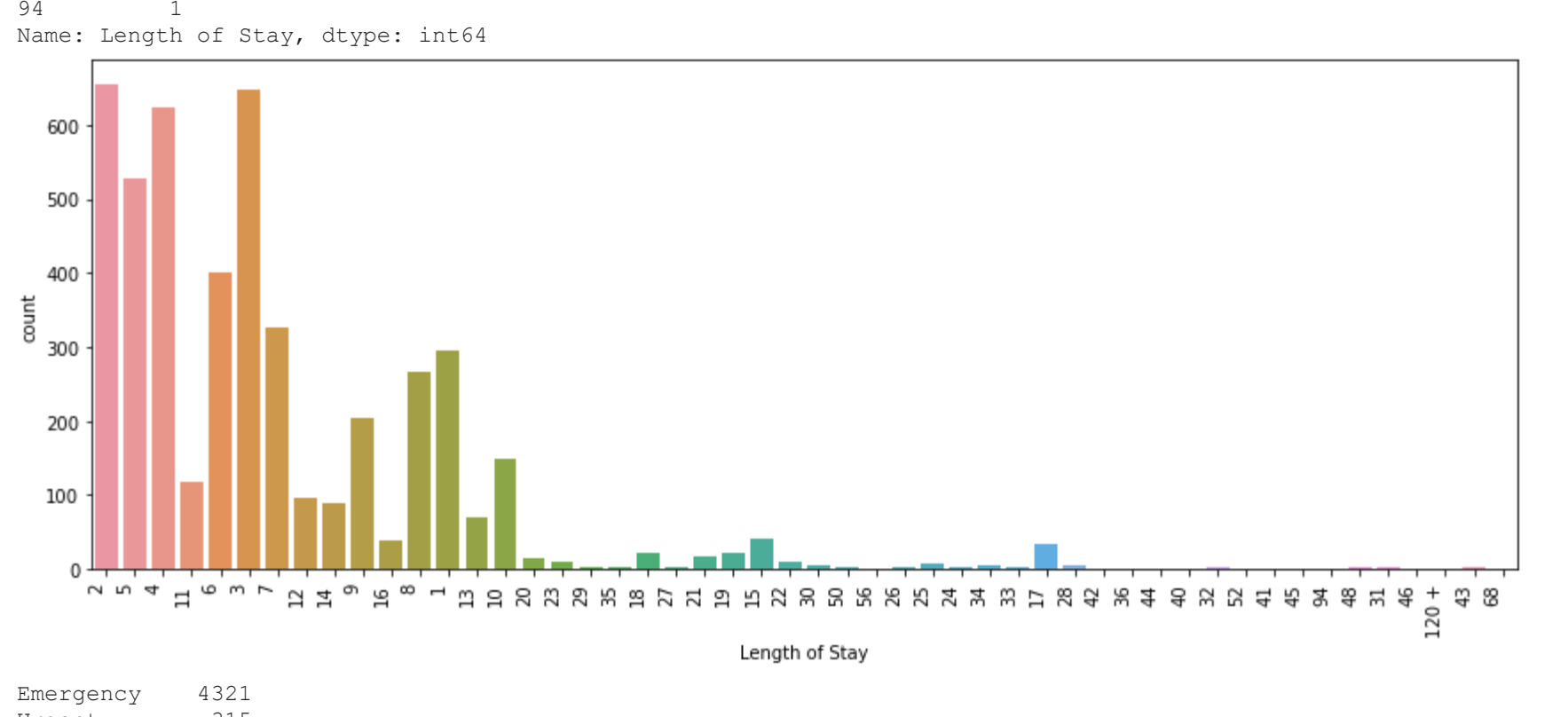
Categorical data

```
using the user defined function to generate plots for the object variables
object_count_plot(train.drop(['Emergency Department Indicator', 'APR Risk of Mortality', 'APR Severity of Illness Code'], axis=1))
```

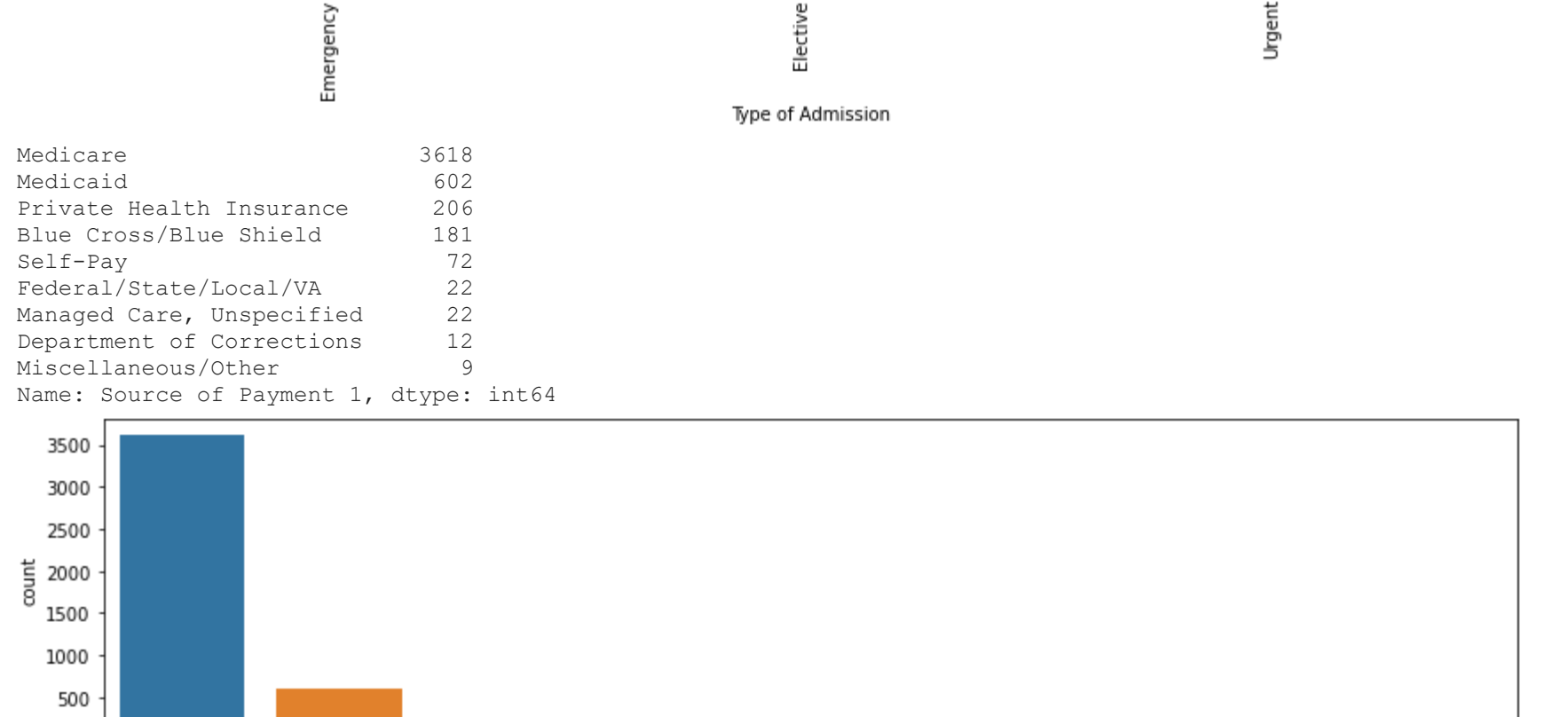
```
70 or Older    3082
50 to 69      1378
30 to 49       237
0 to 19        28
18 to 29       19
Name: Age Group, dtype: int64
```



```
White          2554
Other Race     2190
Name: Gender, dtype: int64
```



```
White          2875
Other Race     999
Black/African American    778
Multi-racial    92
Name: Race, dtype: int64
```



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1383  1
1384  1
1385  1
1386  1
1387  1
1388  1
1389  1
1390  1
1391  1
1392 
```


Total NaN Percent of NaN Nunique Dtype					
Facility ID	0	0.0	8	int64	
Age Group	0	0.0	5	object	
Total Charges	0	0.0	4733	float64	
Emergency Department Indicator	0	0.0	2	object	
Source of Payment 1	0	0.0	9	object	
APR Risk of Mortality	0	0.0	4	object	
APR Severity of Illness Description	0	0.0	4	object	
APR Severity of Illness Code	0	0.0	4	int64	
APR MDC Description	0	0.0	1	object	
APR DRG Code	0	0.0	1	int64	
CCS Procedure Description	0	0.0	75	object	
CCS Procedure Code	0	0.0	75	int64	
CCS Diagnosis Description	0	0.0	4	object	
CCS Diagnosis Code	0	0.0	4	int64	
Patient Disposition	0	0.0	17	object	
Type of Admission	0	0.0	3	object	
Length of Stay	0	0.0	47	float64	
Ethnicity	0	0.0	4	object	
Race	0	0.0	4	object	
Gender	0	0.0	2	object	
Zip Code - 3 digits	0	0.0	47	object	
Total Costs	0	0.0	4733	float64	

```
In [ ]:

In [102]: #selecting all categorical columns
fields = []
for col in train.columns:
    if train[col].dtype == 'object':
        fields.append(col)
print(fields)

['Age Group', 'Zip Code - 3 digits', 'Gender', 'Race', 'Ethnicity', 'Type of Admission', 'Patient Disposition',
'CCS Diagnosis Description', 'CCS Procedure Description', 'APR MDC Description', 'APR Severity of Illness Description',
'APR Risk of Mortality', 'Source of Payment 1', 'Emergency Department Indicator']
```

Label Encoding

```
In [103]: train['Length_of_Stay'] = train['Length of Stay']
train.drop('Length of Stay',axis = 1,inplace = True)
train['Total_Costs'] = train['Total Costs']
train.drop('Total Costs',axis = 1,inplace = True)

In [103]: train['Total_Charges'] = train['Total Charges']
train.drop('Total Charges',axis = 1,inplace = True)

In [103]: train['Total_Costs'][:17]
```

```
Out[103]: 7337.06
```

```
In [103]: #transforming the 'timeline' columns into gaussian distribution - NOT NEEDED FOR TREE BASED MODELS
#train['Timeline'] = np.log(train['Timeline']).values
#test['Timeline'] = np.log(test['Timeline']).values
#label encoding the target variables: 'Total Cost'

Length of Stay map = {'>12','10-12','6-10','<6'}
Total_Cost_map = {'>19000','17,000-19,000','11,000-17,000','<11,000'}
#manual mapping
#fields = ['Length of Stay', 'Total Costs']
for index in list(train.index):
    if (train['Total_Costs'][index] <= 10000):
        train['Total_Costs'][index] = 1
    elif (train['Total_Costs'][index] >= 11000) & (train['Total_Costs'][index] <= 17000):
        train['Total_Costs'][index] = 2
    elif (train['Total_Costs'][index] >= 17000) & (train['Total_Costs'][index] <= 19000):
        train['Total_Costs'][index] = 3
    elif (train['Total_Costs'][index] >= 19000):
        train['Total_Costs'][index] = 4
    else:
        index+=1
```

```
In [103]: #label encoding the target variables: 'Length of Stay'
Length_of_Stay_map = {'>12','10-12','6-10','<6'}
#Total_Cost_map = {'>19000','17,000-19,000','11,000-17,000','<11,000'}
#manual mapping
#fields = ['Length of Stay', 'Total Costs']
for index in list(train.index):
    if (train['Length_of_Stay'][index] <= 6):
        train['Length_of_Stay'][index] = 1
    elif (train['Length_of_Stay'][index] >= 6) & (train['Length_of_Stay'][index] <= 10):
        train['Length_of_Stay'][index] = 2
    elif (train['Length_of_Stay'][index] >= 10) & (train['Length_of_Stay'][index] <= 12):
        train['Length_of_Stay'][index] = 3
    elif (train['Length_of_Stay'][index] >= 12):
        train['Length_of_Stay'][index] = 4
    else:
        index+=1
```

```
In [103]: #representation of the four categories of Total Costs
train['Total_Costs'].value_counts()
```

```
Out[103]: 1.0    2283
         4.0    1281
         2.0     947
         3.0     227
         Name: Total_Costs, dtype: int64
```

```
In [103]: #representing the four categories of Total Costs
train['Length_of_Stay'].value_counts()
```

```
Out[103]: 1.0    3193
         4.0     426
         3.0    214
         2.0     227
         Name: Length_of_Stay, dtype: int64
```

```
In [103]: quality_report(train)
```

Total NaN Percent of NaN Nunique Dtype					
Facility ID	0	0.0	8	int64	
Age Group	0	0.0	5	object	
Total Costs	0	0.0	4	float64	
Length of Stay	0	0.0	4	float64	
Emergency Department Indicator	0	0.0	2	object	
Source of Payment 1	0	0.0	9	float64	
APR Risk of Mortality	0	0.0	4	float64	
APR Severity of Illness Description	0	0.0	4	float64	
APR Severity of Illness Code	0	0.0	4	int64	
APR MDC Description	0	0.0	1	float64	
APR DRG Code	0	0.0	1	int64	
CCS Procedure Description	0	0.0	75	float64	
CCS Procedure Code	0	0.0	75	int64	
CCS Diagnosis Description	0	0.0	4	float64	
CCS Diagnosis Code	0	0.0	4	int64	
Patient Disposition	0	0.0	17	float64	
Type of Admission	0	0.0	3	float64	
Ethnicity	0	0.0	4	float64	
Race	0	0.0	4	float64	
Zip Code - 3 digits	0	0.0	47	float64	
Total Charges	0	0.0	4733	float64	

```
In [104]: #renaming some columns
train['Total_1']=train['Total Costs']
train.drop('Total Costs',axis =1,inplace= True)
train['Total_Charges']=train['Total Charges']
train.drop('Total Charges',axis =1,inplace= True)
```

```
In [104]: train.head(6)
```

Facility ID	Age Group	Zip Code - 3 digits	Gender	Race	Ethnicity	Type of Admission	Patient Disposition	CCS Diagnosis Code	CCS Diagnosis Description	CCS Procedure Code	CCS Procedure Description	CCS Procedure Code	CCS Procedure Description	APR DRG Code	APR MDC Description
0	66	40	430	1.0	3.0	1.0	1.0	6.0	108	0.0	58	28.0	194	0.0	
1	66	30	430	0.0	3.0	1.0	1.0	11.0	108	0.0	0	36.0	194	0.0	
2	66	30	430	1.0	3.0	1.0	1.0	16.0	108	0.0	0	36.0	194	0.0	
3	66	40	430	1.0	3.0	1.0	1.0	6.0	108	0.0	201	7.0	194	0.0	
4	66	40	430	1.0	3.0	1.0	1.0	11.0	108	0.0	0	36.0	194	0.0	
5	66	40	430	0.0	3.0	1.0	1.0	16.0	108	0.0	0	36.0	194	0.0	

6.Feature Engineering

Polynomial features

This generates extra features based on perceived interactions between attributes

```
In [104]: poly_feature_1 = ['Gender', 'Race', 'Ethnicity',
                    'CCS Diagnosis Description', 'CCS Procedure Code', 'CCS Procedure Description',
                    'APR Severity of Illness Code', 'APR Severity of Illness Description',
                    'APR Risk of Mortality', 'Source of Payment 1',
                    'Emergency Department Indicator', 'Length of Stay', 'Total_Costs',
                    'Total_Charges', 'poly1_0', 'poly1_1', 'poly1_2', 'poly1_3', 'poly1_4',
                    'poly1_5', 'poly1_6', 'poly1_7', 'poly1_8', 'poly1_9', 'poly1_10', 'poly1_11', 'poly1_12',
                    'poly1_13', 'poly1_14', 'geography', 'CCS Profile', 'APR Profile',
                    'Total_Charges_binned'], dtype='object')

In [104]: train.drop('Total Costs',axis =1,inplace= True)
```

```
In [104]: #from sklearn.preprocessing import PolynomialFeatures
poly = PolynomialFeatures(degree=2, interaction_only=True, include_bias=False)
poly1 = poly.fit_transform(train[poly_feature_1])
poly2 = poly.fit_transform(train[poly_feature_2])
poly3 = poly.fit_transform(train[poly_feature_3])
```

```
In [104]: df_poly1 = pd.DataFrame(poly1, columns=["poly1_{}".format(i) for i in range(poly1.shape[1])])
df_poly2 = pd.DataFrame(poly2, columns=["poly2_{}".format(i) for i in range(poly2.shape[1])])
df_poly3 = pd.DataFrame(poly3, columns=["poly3_{}".format(i) for i in range(poly3.shape[1])])
```

```
In [104]: train.columns
```

```
Out[104]: Index(['Facility ID', 'Age Group', 'Zip Code - 3 digits', 'Gender', 'Race',
        'Ethnicity', 'Type of Admission', 'Patient Disposition',
        'CCS Diagnosis Code', 'CCS Diagnosis Description', 'CCS Procedure Code',
        'CCS Procedure Description', 'APR DRG Code', 'APR MDC Description',
        'APR Severity of Illness Code', 'APR Severity of Illness Description',
        'APR Risk of Mortality', 'Source of Payment 1',
        'Emergency Department Indicator', 'Length of Stay', 'Total_Costs',
        'Total_Charges', 'poly1_0', 'poly1_1', 'poly1_2', 'poly1_3', 'poly1_4',
        'poly1_5', 'poly1_6', 'poly1_7', 'poly1_8', 'poly1_9', 'poly1_10', 'poly1_11', 'poly1_12',
        'poly1_13', 'poly1_14', 'geography', 'CCS Profile', 'APR Profile',
        'Total_Charges_binned'],
        dtype='object')
```

```
In [104]: new_data = pd.concat([train, df_poly1], axis = 1)
new_data = pd.concat([new_data, df_poly2], axis = 1)
new_data = pd.concat([new_data, df_poly3], axis = 1)
```

```
In [104]: new_data.columns
```

```
Out[104]: Index(['Facility ID', 'Age Group', 'Zip Code - 3 digits', 'Gender', 'Race',
        'Ethnicity', 'Type of Admission', 'Patient Disposition',
        'CCS Diagnosis Code', 'CCS Diagnosis Description', 'CCS Procedure Code',
        'CCS Procedure Description', 'APR DRG Code', 'APR MDC Description',
        'APR Severity of Illness Code', 'APR Severity of Illness Description',
        'APR Risk of Mortality', 'Source of Payment 1',
        'Emergency Department Indicator', 'Length of Stay', 'Total_Costs',
        'Total_Charges', 'poly1_0', 'poly1_1', 'poly1_2', 'poly1_3', 'poly1_4',
        'poly1_5', 'poly1_6', 'poly1_7', 'poly1_8', 'poly1_9', 'poly1_10', 'poly1_11', 'poly1_12',
        'poly1_13', 'poly1_14', 'geography', 'CCS Profile', 'APR Profile',
        'Total_Charges_binned'],
        dtype='object')
```

```
In [104]: new_data['geography'] = new_data[poly_feature_1].sum(axis = 1)
new_data['CCS Profile'] = new_data[poly_feature_2].sum(axis = 1)
new_data['APR Profile'] = new_data[poly_feature_3].sum(axis = 1)
```

```
In [104]: # bin_label = [1,2,3,4,5]
new_data['Total_Charges_binned'] = pd.qcut(new_data.Total_Charges, q = [0, .2, .4, .6, .8, 1], duplicates="drop",
#new_data.drop('Total_Charges',axis =1,inplace= True)
```

```
In [104]: new_data.columns
```

```
Out[104]: Index(['Facility ID', 'Age Group', 'Zip Code - 3 digits', 'Gender', 'Race',
        'Ethnicity', 'Type of Admission', 'Patient Disposition',
        'CCS Diagnosis Code', 'CCS Diagnosis Description', 'CCS Procedure Code',
        'CCS Procedure Description', 'APR DRG Code', 'APR MDC Description',
        'APR Severity of Illness Code', 'APR Severity of Illness Description',
        'APR Risk of Mortality', 'Source of Payment 1',
        'Emergency Department Indicator', 'Length of Stay', 'Total_Costs',
        'Total_Charges', 'poly1_0', 'poly1_1', 'poly1_2', 'poly1_3', 'poly1_4',
        'poly1_5', 'poly1_6', 'poly1_7', 'poly1_8', 'poly1_9', 'poly1_10', 'poly1_11', 'poly1_12',
        'poly1_13', 'poly1_14', 'geography', 'CCS Profile', 'APR Profile',
        'Total_Charges_binned'],
        dtype='object')
```

```
In [ ]:
```

```
In [111]: #splitting the data into train and test features
# randomly predicting on 10% of the data set
test = new_data.sample(frac = 0.10)
```

```
In [111]: #dropping the prediction set from the data to ensure unique training data
train = new_data.drop(list(test.index),axis = 0)
```

```
In [111]: new_data.shape
```

```
Out[111]: (4749, 57)
```

```
In [111]: test.shape
```

```
Out[111]: (475, 57)
```

```
In [111]: train.shape
```

```
Out[111]: (4274, 57)
```

```
In [111]: features = train.select_dtypes(include = 'number').columns
features
```

```
Out[111]: Index(['Facility ID', 'Age Group', 'Zip Code - 3 digits', 'Gender', 'Race',
        'Ethnicity', 'Type of Admission', 'Patient Disposition',
        'CCS Diagnosis Code', 'CCS Diagnosis Description', 'CCS Procedure Code',
        'CCS Procedure Description', 'APR DRG Code', 'APR MDC Description',
        'APR Severity of Illness Code', 'APR Severity of Illness Description',
        'APR Risk of Mortality', 'Source of Payment 1',
        'Emergency Department Indicator', 'Length of Stay', 'Total_Costs',
        'Total_Charges', 'poly1_0', 'poly1_1', 'poly1_2', 'poly1_3', 'poly1_4',
        'poly1_5', 'poly1_6', 'poly1_7', 'poly1_8', 'poly1_9', 'poly1_10', 'poly1_11', 'poly1_12',
        'poly1_13', 'poly1_14', 'geography', 'CCS Profile', 'APR Profile',
        'Total_Charges_binned'],
        dtype='object')
```

```
In [111]: train.head(7)
```

Facility ID	Age Group	Zip Code - 3 digits	Gender	Race	Ethnicity	Type of Admission	Patient Disposition	CCS Diagnosis Code	CCS Diagnosis Description	CCS Procedure Code	CCS Procedure Description	CCS Procedure Code	CCS Procedure Description	APR DRG Code	APR MDC Description
0	66.0	4.0	430	1.0	3.0	1.0	1.0	6.0	108.0	0.0	58.0	28.0	194.0	0.0	
1	66.0	3.0	430	0.0	3.0	1.0	1.0	11.0	108.0	0.0	0.0	36.0	194.0	0.0	
2	66.0	4.0	430	1.0	3.0	1.0	1.0	16.0	108.0	0.0	0.0	36.0	194.0	0.0	
3	66.0	3.0	430	1.0	3.0	1.0	1.0	6.0	108.0	0.0	201.0	7.0	194.0	0.0	
4	66.0	4.0	430	1.0	3.0	1.0	1.0	11.0	108.0	0.0	0.0	36.0	194.0	0.0	
5	66.0	4.0	430	0.0	3.0	1.0	1.0	16.0	108.0	0.0	0.0	36.0	194.0	0.0	
6	66.0	3.0	430	1.0	3.0	1.0	1.0	6.0	108.0	0.0	0.0	36.0	194.0	0.0	

```
In [112]: #dropping missing values
train.dropna(axis = 0,inplace= True)
train.shape
```

```
Out[112]: (4255, 57)
```

6. Modeling

```
In [105]: from catboost import CatBoostClassifier
from sklearn.model_selection import StratifiedFold
from sklearn.metrics import log_loss, f1_score
from xgboost import XGBClassifier
from lightgbm import LGBMClassifier
from sklearn.model_selection import train_test_split, cross_val_score
```

```
In [106]: #defining both length of stay and total costs
X = train[features]
X.drop('Length of Stay',axis =1,inplace=True)
#X.drop('Total Costs',axis =1,inplace=True)
target = 'Length of Stay'
#target = 'Total_Costs'
y = train[target]
```

```
In [106]: X.columns
```

```
Out[106]: Index(['Facility ID', 'Age Group', 'Zip Code - 3 digits', 'Gender', 'Race',
        'Ethnicity', 'Type of Admission', 'Patient Disposition',
        'CCS Diagnosis Code', 'CCS Diagnosis Description', 'CCS Procedure Code',
        'CCS Procedure Description', 'APR DRG Code', 'APR MDC Description',
        'APR Severity of Illness Code', 'APR Severity of Illness Description',
        'APR Risk of Mortality', 'Source of Payment 1',
        'Emergency Department Indicator', 'Length of Stay', 'Total_Costs',
        'Total_Charges', 'poly1_0', 'poly1_1', 'poly1_2', 'poly1_3', 'poly1_4',
        'poly1_5', 'poly1_6', 'poly1_7', 'poly1_8', 'poly1_9', 'poly1_10', 'poly1_11', 'poly1_12',
        'poly1_13', 'poly1_14', 'geography', 'CCS Profile', 'APR Profile',
        'Total_Charges_binned'],
        dtype='object')
```

```
In [106]: #splitting with train_test_split
```

```
In [106]: X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
```

```
In [106]: X_train.columns
```

```
Out[106]: Index(['Facility ID', 'Age Group', 'Zip Code - 3 digits', 'Gender', 'Race',
        'Ethnicity', 'Type of Admission', 'Patient Disposition',
        'CCS Diagnosis Code', 'CCS Diagnosis Description', 'CCS Procedure Code',
        'CCS Procedure Description', 'APR DRG Code', 'APR MDC Description',
        'APR Severity of Illness Code', 'APR Severity of Illness Description',
        'APR Risk of Mortality', 'Source of Payment 1',
        'Emergency Department Indicator', 'Length of Stay', 'Total_Costs',
        'Total_Charges', 'poly1_0', 'poly1_1', 'poly1_2', 'poly1_3', 'poly1_4',
        'poly1_5', 'poly1_6', 'poly1_7', 'poly1_8', 'poly1_9', 'poly1_10', 'poly1_11', 'poly1_12',
        'poly1_13', 'poly1_14', 'geography', 'CCS Profile', 'APR Profile',
        'Total_Charges_binned'],
        dtype='object')
```

```
In [106]: #oversampling
from imblearn.over_sampling import SMOTE
sm = SMOTE(random_state=42)
X_train_res, y_train_res = sm.fit_resample(X, y)
```

```
Out[106]: Index(['Facility ID', 'Age Group', 'Zip Code - 3 digits', 'Gender', 'Race',
        'Ethnicity', 'Type of Admission', 'Patient Disposition',
        'CCS Diagnosis Code', 'CCS Diagnosis Description', 'CCS Procedure Code',
        'CCS Procedure Description', 'APR DRG Code', 'APR MDC Description',
        'APR Severity of Illness Code', 'APR Severity of Illness Description',
        'APR Risk of Mortality', 'Source of Payment 1',
        'Emergency Department Indicator', 'Length of Stay', 'Total_Costs',
        'Total_Charges', 'poly1_0', 'poly1_1', 'poly1_2', 'poly1_3', 'poly1_4',
        'poly1_5', 'poly1_6', 'poly1_7', 'poly1_8', 'poly1_9', 'poly1_10', 'poly1_11', 'poly1_12',
        'poly1_13', 'poly1_14', 'geography', 'CCS Profile', 'APR Profile',
        'Total_Charges_binned'],
        dtype='object')
```

```
In [107]: #MODEL 1 XGBClassifier #Benchmark f1 score 0.450828729281768
#MODEL 1 XGBClassifier(max_depth = 8, n_estimators = 500,num_class=4,objective='multi:softmax',learning_rate=0.01,
model = XGBClassifier(max_depth=8, n_estimators=1000,
max_depth=5,
min_child_weight=1,
gamma=0,
average='micro',
subsample=0.8,
colsample_bytree=0.8,
objective='multi:softmax',
nthread=4,
num_class=4,
seed=27)
```

```
xgb1.fit(X_train, y_train)
prediction = xgb1.predict(X_test)
```

```
[12:41:53] WARNING: C:\Users\Administrator\workspace\xgboost-win64_release_1.4.0\src\learner.cc:573:
Parameters: { "average" } might not be used.

This may not be accurate due to some parameters are only used in language bindings but
passed down to XGboost core. Or some parameters are not used but slip through this
verification. Please open an issue if you find above cases.
```

```
[12:41:53] WARNING: C:\Users\Administrator\workspace\xgboost-win64_release_1.4.0\src\learner.cc:1095: Starting
in XGboost 1.3.0, the default evaluation metric used with the objective 'multi:softmax' was changed from 'merr
or' to 'mlogloss', explicitly set eval_metric if you'd like to restore the old behavior.

Accuracy: 85.33%
```

```
In [106]: from sklearn.metrics import accuracy_score

#MODEL 1 XGBClassifier #Benchmark f1 score 0.450828729281768
model = XGBClassifier(max_depth = 8, n_estimators = 500,num_class=4,objective='multi:softmax',learning_rate=0.01,
model.fit(X_train, y_train)
prediction = model.predict(X_test)
```

```
# evaluate predictions
accuracy = accuracy_score(y_test, prediction)
print("Accuracy: %.2f%%" % (accuracy * 100.0))

[12:42:17] WARNING: C:\Users\Administrator\workspace\xgboost-win64_release_1.4.0\src\learner.cc:1095: Starting
in XGboost 1.3.0, the default evaluation metric used with the objective 'multi:softmax' was changed from 'merr
or' to 'mlogloss', explicitly set eval_metric if you'd like to restore the old behavior.

Accuracy: 84.86%
```

```
In [106]: # evaluate predictions
#print(y_test)

print("Accuracy: %.2f%%" % (accuracy * 100.0))

Accuracy: 84.86%
```

MODEL 2 LGBM Classifier with oversampled inputs #Benchmark Accuracy: 95.77%

```
In [106]: #defining both length of stay and total costs
X = train[features]
X.drop('Length of Stay',axis =1,inplace=True)
#X.drop('Total Costs',axis =1,inplace=True)
target = 'Length of Stay'
#target = 'Total_Costs'
y = train[target]
```

```
In [107]: #undersampling
from imblearn.under_sampling import RandomUnderSampler
#rus = RandomUnderSampler(random_state = 0)
#X_train_res, y_train_res = rus.fit_resample(X, y)
```

```
In [107]: X.columns
```

```
Out[107]: Index(['Facility ID', 'Age Group', 'Zip Code - 3 digits', 'Gender', 'Race',
        'Ethnicity', 'Type of Admission', 'Patient Disposition',
        'CCS Diagnosis Code', 'CCS Diagnosis Description', 'CCS Procedure Code',
        'CCS Procedure Description', 'APR DRG Code', 'APR MDC Description',
        'APR Severity of Illness Code', 'APR Severity of Illness Description',
        'APR Risk of Mortality', 'Source of Payment 1',
        'Emergency Department Indicator', 'Length of Stay', 'Total_Costs',
        'Total_Charges', 'poly1_0', 'poly1_1', 'poly1_2', 'poly1_3', 'poly1_4',
        'poly1_5', 'poly1_6', 'poly1_7', 'poly1_8', 'poly1_9', 'poly1_10', 'poly1_11', 'poly1_12',
        'poly1_13', 'poly1_14', 'geography', 'CCS Profile', 'APR Profile',
        'Total_Charges_binned'],
        dtype='object')
```

```
In [107]: #oversampling
from imblearn.over_sampling import SMOTE
sm = SMOTE(random_state=42)
X_train_res, y_train_res = sm.fit_resample(X, y)
```

```
Out[107]: Index(['Facility ID', 'Age Group', 'Zip Code - 3 digits', 'Gender', 'Race',
        'Ethnicity', 'Type of Admission', 'Patient Disposition',
        'CCS Diagnosis Code', 'CCS Diagnosis Description', 'CCS Procedure Code',
        'CCS Procedure Description', 'APR DRG Code', 'APR MDC Description',
        'APR Severity of Illness Code', 'APR Severity of Illness Description',
        'APR Risk of Mortality', 'Source of Payment 1',
        'Emergency Department Indicator', 'Length of Stay', 'Total_Costs',
        'Total_Charges', 'poly1_0', 'poly1_1', 'poly1_2', 'poly1_3', 'poly1_4',
        'poly1_5', 'poly1_6', 'poly1_7', 'poly1_8', 'poly1_9', 'poly1_10', 'poly1_11', 'poly1_12',
        'poly1_13', 'poly1_14', 'geography', 'CCS Profile', 'APR Profile',
        'Total_Charges_binned'],
        dtype='object')
```

```
In [107]: #MODEL 3 CatBoostClassifier no sampling #Benchmark f1 score 0.3827314238273142
#MODEL 3 CatBoostClassifier(loss_function='MultiClass', class_weights=[.15, .35, .35, .15], iterations=1000, des
model.fit(X_train, y_train)
prediction = model.predict(X_test)
```

```
print("Accuracy: %.2f%%" % (accuracy_score(prediction, y_test)*100.0))
```


[illegible]

	Facility ID	Age Group	Zip Code	Gender	Race	Ethnicity	Type of Admission	Patient Disposition	CCS Code	Diagnosis Description	CCS Procedure Code	CCS Procedure Description	APR DRG Code	APR DRG Description	
	4464	14640	40	30	10	30	20	10	70	1080	00	1930	200	1940	
	4545	5410	30	120	10	20	10	00	150	1080	00	2130	600	1940	
	2770	5410	30	160	10	30	10	10	60	1080	00	00	360	1940	
	995	4130	30	410	10	30	20	10	70	990	00	00	360	1940	
	1609	5410	30	160	10	30	20	10	70	990	10	2220	40	1940	
	3113	14380	40	110	10	20	20	10	70	1080	00	1790	130	1940	
	1631	5410	40	140	00	30	10	10	70	1080	00	2160	670	1940	
	818	1810	30	230	10	30	10	10	60	1080	00	2280	650	1940	
	3173	14380	30	110	10	20	20	10	60	1080	00	2310	620	1940	
	2365	5410	40	90	00	30	10	10	70	1080	00	2170	610	1940	
	717	1810	30	230	00	30	10	10	70	1080	10	00	360	1940	
	2972	10050	40	260	10	30	10	10	60	990	10	00	360	1940	
	798	1810	20	240	10	20	30	10	60	1080	00	00	360	1940	
	4198	14640	40	30	00	20	20	10	70	990	10	00	360	1940	
	3076	14380	30	110	10	00	10	10	120	1080	00	2310	620	1940	
	2049	5410	40	110	10	30	10	10	60	1080	00	2170	610	1940	
	1440	4130	40	440	10	30	10	20	160	1080	00	00	360	1940	
	1728	5410	30	140	00	20	20	10	60	1080	00	580	280	1940	
	2828	10050	40	260	10	20	10	10	70	1080	00	00	360	1940	
	2920	10050	40	260	00	30	10	10	160	1080	00	00	360	1940	
	4628	14640	40	00	00	30	10	10	160	1080	00	00	360	1940	
	2346	5410	40	160	00	20	10	10	60	1080	00	00	360	1940	
	4589	14640	40	00	00	20	10	10	30	990	10	2160	670	1940	
	1540	4130	30	440	10	30	10	10	90	3000	30	410	470	1940	
	1464	14640	40	00	10	20	20	10	60	1080	00	00	360	1940	
	1522	4130	40	420	10	30	10	10	100	1080	00	00	360	1940	
	228	1810	40	240	00	30	10	10	160	990	10	00	360	1940	
	1066	4130	40	410	00	30	10	10	90	1080	00	00	360	1940	
	4131	14640	30	20	10	30	10	10	160	1080	00	260	540	1940	
	3090	14380	30	00	00	20	10	10	120	1080	00	1930	200	1940	
	4317	14640	20	30	00	10	20	10	70	1080	00	00	360	1940	
	2904	10050	40	260	10	30	10	10	70	990	10	00	360	1940	
	1084	4130	40	400	10	30	10	10	70	1080	00	00	360	1940	
	3219	14380	20	00	10	00	10	10	60	1080	00	2310	620	1940	
	4475	14640	30	00	10	10	00	10	60	1080	00	00	360	1940	
	2462	5410	40	140	10	00	10	10	60	1080	00	00	360	1940	
	1895	5410	40	120	10	30	10	10	160	1080	00	00	360	1940	
	4090	14640	40	30	00	20	10	10	70	1080	00	1930	200	1940	
	3059	14380	20	30	10	00	10	10	60	1080	00	2310	620	1940	
	302	1810	30	230	10	30	10	10	70	1080	00	00	360	1940	
	2443	5410	40	160	10	20	10	10	70	1080	00	410	470	1940	
	155	660	40	370	00	30	10	10	160	1080	00	00	360	1940	
	2283	5410	40	130	00	30	20	10	70	1080	00	00	360	1940	
	3089	14380	40	130	00	20	10	10	90	1080	00	2310	620	1940	
	1100	4130	30	380	10	30	10	20	30	1080	00	610	370	1940	
	460	1810	40	230	00	20	10	10	60	1080	00	00	360	1940	
	1302	4130	30	420	10	00	10	10	60	1080	00	00	360	1940	
	886	4130	40	420	00	30	10	10	60	1080	00	00	360	1940	
	1343	14630	40	00	10	10	30	10	160	1080	00	00	360	1940	
	3070	4130	40	420	00	30	10	10	260	60	00	00	360	1940	
	858	1810	40	240	00	20	30	10	60	990	10	00	360	1940	
	3401	14630	40	110	10	20	30	10	00	70	1080	00	2130	640	1940
	2366	5410	40	120	00	30	10	10	60	1080	00	1930	200	1940	
	519	1810	30	240	10	00	10	10	70	1080	00	00	360	1940	
	1282	4130	40	420	00	30	10	10	70	1080	00	00	360	1940	
	4688	14640	40	00	00	20	10	10	60	1080	00	00	360	1940	
	2930	10050	40	260	00	30	10	10	70	990	10	00	360	1940	
	1037	4130	40	360	10	30	10	10	70	1080	00	00	360	1940	
	4446	14640	30	30	10	10	00	10	70	1080	00	1930	200	1940	
	425	1810	40	240	00	30	10	10	70	1080	00	00	360	1940	
	1014	14380	30	120	10	30	10	10	60	1080	00	1960	330	1940	
	3494	4130	40	410	10	30	10	10	70	1080	00	2180	670	1940	
	304	1810	40	240	10	30	10	10	110	1080	00	00	360	1940	
	929	4130	30	290	10	30	10	10	60	1080	00	00	360	1940	
	4667	14640	20	30	00	10	20	10	60	1330	20	2160	670	1940	
	2696	10050	30	260	10	00	10	10	60	1080	00	00	360	1940	
	4666	14640	30	00	00	00	20	10	70	1080	00	2310	620	1940	
	4700	14640	40	00	00	00	10	10	60	990	10	1930	200	1940	
	102	660	40	430	00	30	10	10	160	1080	00	00	360	1940	
	202	1810	30	230	00	20	10	10	70	1080	00	1930	200	1940	
	1550	4130	30	310	00	30	10	20	60	1080	00	00	360	1940	
	2715	10050	30	260	00	30	10	10	70	990	10	910	630	1940	
	8896	14630	30	00	00	20	10	10	70	1080	00	1930	200	1940	
	2441	5410	40	140	00	30	10	10	160	1080	00	2160	670	1940	
	1574	4130	30	420	10	00	10	10	70	990	10	00	360	1940	
	421	14640	30	420	10	00	20	10	70	990	10	2310	620	1940	
	1926	5410	40	140	00	30	10	10	60	1080	00	400	400	1940	
	2673	10050	40	260	10	30	10	20	30	1080	00	00	360	1940	
	4716	14640	30	460	10	20	10	20	70	1080	00	650	50	1940	
	4675	14640	30	460	10	20	10	10	60	3000	30	410	470	1940	
	4252	14640	20	460	00	30	10	00	60	1080	00	00	360	1940	
	323	14380	30	110	00	00	10	10	60	1080	00	2310	620	1940	
	3229	14380	30	110	00	00	20	10	60	1080	00	2310	620	1940	
	2589	5410	40	140	00	30	10	10	60	1080	00	00	360	1940	
	1660	14640	40	00	00	20	10	10	60	990	10	2310	620	1940	
	343	1810	30	240	10	30	10	10	60	990	10	00	360	1940	
	4232	5410	30	230	10	30	10	10	60	3000	30	410	470	1940	
	2034	1810	30	130	10	30	10	10	160	1080	00	00	360	1940	
	1828	5410	40	130	00	00	10	10	110	1080	00	00	360	1940	
	2821	10050	30	260	00	30	10	10	70	1080	00	00	360	1940	
	3699	14630	20	00	00	20	20	10	60	990	10	00	360	1940	
	688	5410	40	230	10	30	10	10	60	1080	00	00	360	1940	
	2088	1810	30	140	10	00	10	10	60	1080	00	700	730	1940	
	2319	5410	40	140	00	30	10	10	60	1080	00	2170	610	1940	
	436	1810	30	240	10	00	10	10	160	1080	00	00	360	1940	
	819	1810	30	240	10	00	10	10	60	1080	00	00	360	1940	
	1659	5410	30	130	10	00	20	10	60	1080	00	9260	670	1940	
	4424	14640	30	00	10	20	20	10	60	1080	00	2160	670	1940	
	1680	5410	30	130	10	30	10	10	70	1080	00	2160	670	1940	
	304	1810	40	230	10	00	20	10	60	1080	00	00	360	1940	
	4629	14640	40	40	10	30	10	00	30	1080	00	2160	670	1940	
	3361	14630	30	460	00	00	10	10	70	1080	00	00	360	1940	
	4128	14640	30	460	10	20	10	10	60	1080	00	2310	620	1940	
	2258	5410	40	120	00	30	10	10	60	1080	00	2220			