Problem statement: ¶

"Predict Possibility of Patient Readmission within 60days from discharge"

To do:

- 1. Exploratory Data Analysis
- 2. Assumptions (if any)
- 3. Feature Engineering & Feature Selection
- 4. Model Building
- 5. Predicted output
- 6. Model Evaluation Metrics

Code:

Import Libraries

```
In [1]: import numpy as np
    import pandas as pd
    import warnings
    warnings.simplefilter('ignore')
```

Read Data

```
In [2]: admissions = pd.read_csv('readmission_data//ADMISSIONS.csv')
    services = pd.read_csv('readmission_data//SERVICES.csv')
    icd = pd.read_csv('readmission_data//DIAGNOSES_ICD.csv')
    d_icd = pd.read_csv('readmission_data//D_ICD_DIAGNOSES.csv')
    icu_stays = pd.read_csv("readmission_data//ICUSTAYS.csv")
    patients = pd.read_csv('readmission_data//PATIENTS.csv')
    drg_codes = pd.read_csv('readmission_data//DRGCODES.csv')
```

Prepare a KEY

```
In [3]:
        admissions['SUBJECT ID'] = admissions['SUBJECT ID'].astype(str)
        admissions['HADM ID'] = admissions['HADM ID'].astype(str)
        admissions['key'] = admissions[['SUBJECT ID', 'HADM ID']].apply(lambda x: ''.j
        oin(x), axis=1)
        services['SUBJECT ID'] = services['SUBJECT ID'].astype(str)
        services['HADM ID'] = services['HADM ID'].astype(str)
        services['key'] = services[['SUBJECT ID', 'HADM ID']].apply(lambda x: ''.join(
        x), axis=1)
        icd['SUBJECT ID'] = icd['SUBJECT ID'].astype(str)
        icd['HADM_ID'] = icd['HADM_ID'].astype(str)
        icd['key'] = icd[['SUBJECT_ID', 'HADM_ID']].apply(lambda x: ''.join(x), axis=1
        icu_stays['SUBJECT_ID'] = icu_stays['SUBJECT_ID'].astype(str)
        icu_stays['HADM_ID'] = icu_stays['HADM_ID'].astype(str)
        icu_stays['key'] = icu_stays[['SUBJECT_ID', 'HADM_ID']].apply(lambda x: ''.joi
        n(x), axis=1)
        drg codes['SUBJECT ID'] = drg codes['SUBJECT ID'].astype(str)
        drg_codes['HADM_ID'] = drg_codes['HADM_ID'].astype(str)
        drg_codes['key'] = drg_codes[['SUBJECT_ID', 'HADM_ID']].apply(lambda x: ''.joi
        n(x), axis=1)
```

Drop un-wanted columns

· Column: Reason

ROW_ID: Unique

HADM ID: Already formed a key

DESCRIPTION: Text

DRG SEVERITY: 46% Missing

DRG MORTALITY: 46% Missing

DOB: Invalid dates, Cant use to get age

DOD: Already we have expire flag

• DOD HOSP: 79% Blanks

DOD SSN: 72% Blanks

ICUSTAY_ID: Unique

• INTIME: Already we have LOS

· OUTTIME: Already we have LOS

SHORT TITLE: text

LONG TITLE: text

SEQ NUM: Key

· TRANSFERTIME: Not found useful

· EDREGTIME: Not found useful

EDOUTTIME: Not found useful

· DIAGNOSIS: Not found useful

```
In [5]: admissions.drop(admission_drop, axis=1, inplace=True)
    services.drop(services_drop, axis=1, inplace=True)
    icd.drop(icd_drop, axis=1, inplace=True)
    d_icd.drop(d_icd_drop, axis=1, inplace=True)
    icu_stays.drop(icu_stays_drop, axis=1, inplace=True)
    patients.drop(patients_drop, axis=1, inplace=True)
    drg_codes.drop(drg_codes_drop, axis=1, inplace=True)
```

Drop duplicates

```
In [6]: admissions = admissions.drop_duplicates()
    services = services.drop_duplicates()
    icd = icd.drop_duplicates()
    d_icd = d_icd.drop_duplicates()
    icu_stays = icu_stays.drop_duplicates()
    patients = patients.drop_duplicates()
    drg_codes = drg_codes.drop_duplicates()
In [7]: # # delete unwanted objects
    del admission_drop
    del services_drop
    del icd drop
```

Clean tables for preparing Master table

del d_icd_drop
del icu_stays_drop
del patients_drop
del drg_codes_drop

```
In [9]: drg_codes['APR'] = [1 if i == 'APR ' else 0 for i in drg_codes.DRG_TYPE]
    drg_codes['HCFA'] = [1 if i == 'HCFA' else 0 for i in drg_codes.DRG_TYPE]
    drg_codes['MS'] = [1 if i == 'MS ' else 0 for i in drg_codes.DRG_TYPE]
    drg_codes.drop(['DRG_TYPE', 'DRG_CODE'], axis = 1, inplace=True)
    drg_codes = drg_codes.groupby('key').agg('sum').reset_index()
    drg_codes = drg_codes.drop_duplicates()
```

Prepare Master

```
In [11]: # # prepare master
    admissions['SUBJECT_ID'] = admissions['SUBJECT_ID'].astype('int')
    admissions = admissions.merge(patients, on = 'SUBJECT_ID', how = 'left')
    admissions = admissions.merge(services, on = 'key', how = 'left')
    admissions = admissions.merge(icd, on = 'key', how = 'left')
    admissions = admissions.merge(icu_stays, on = 'key', how = 'left')
    admissions = admissions.merge(drg_codes, on = 'key', how = 'left')
```

Prepare Target

```
In [12]: admissions['ADMITTIME'] = pd.to_datetime(admissions.ADMITTIME)
    admissions['DISCHTIME'] = pd.to_datetime(admissions.DISCHTIME)
    admissions['DEATHTIME'] = pd.to_datetime(admissions.DEATHTIME)

In [13]: admissions['SUBJECT_ID'] = admissions['SUBJECT_ID'].astype('int')

In [14]: admissions = admissions.sort_values(['SUBJECT_ID', 'ADMITTIME'], ascending=[True, True])

In [15]: admissions['target'] = 0
    admissions = admissions.reset_index()
    del admissions['index']
```

```
In [16]: n = admissions.shape[0]
In [17]: def prepare_target(admissions, n):
             i = 0
             while i < n:
                  temp = admissions.SUBJECT_ID[i]
                  count = admissions[admissions.SUBJECT ID == temp].shape[0]
                  if count == 1:
                      i = i + 1
                  else:
                      while count > 1:
                          tempadmitdate = admissions.ADMITTIME[i + 1]
                          tempdichargedate = admissions.DISCHTIME[i]
                          if (tempdichargedate - tempadmitdate).days < 60:</pre>
                              admissions['target'][i] = 1
                          i = i + 1
                          count = count - 1
                      i = i + 1
             return(admissions)
In [18]:
         temp_admissions = admissions.copy() # take a backup !
         admissions = prepare_target(admissions = admissions, n = n)
In [19]:
         # admissions = pd.read_csv('..//data//full_data.csv')
In [20]: # del admissions['daystoadmit']
In [21]:
         ## check target class
         admissions['target'].value_counts(normalize=True) # only 6.2% of 1's
Out[21]: 0
              0.788795
              0.211205
         Name: target, dtype: float64
```

Check and handle missing values

In [22]: | admissions.isnull().sum(axis=0)

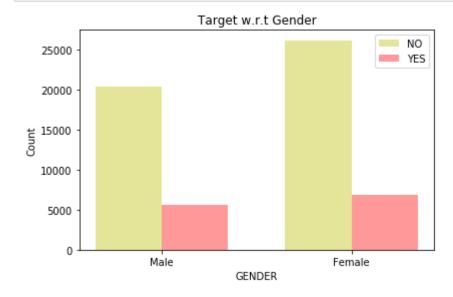
```
Out[22]: SUBJECT ID
                                      0
         ADMITTIME
                                      0
         DISCHTIME
                                      0
         DEATHTIME
                                  53122
         ADMISSION TYPE
                                      0
         ADMISSION LOCATION
                                      0
         DISCHARGE_LOCATION
                                      0
         INSURANCE
                                      0
         LANGUAGE
                                  25332
                                    458
         RELIGION
         MARITAL STATUS
                                  10128
         ETHNICITY
                                      0
                                      0
         HOSPITAL_EXPIRE_FLAG
         HAS CHARTEVENTS DATA
                                      0
         key
                                      0
         GENDER
                                      0
         EXPIRE FLAG
                                      0
                                     50
         service ct
         icd_ct
                                      0
         Avg LOS
                                   1200
         N CAREUNITS
                                   1190
         DBSOURCE
                                   1190
         APR
                                     86
         HCFA
                                     86
         MS
                                     86
         target
                                      0
         dtype: int64
         drop_cols = ['DEATHTIME', 'LANGUAGE', 'MARITAL_STATUS'] # many missing !
In [23]:
         admissions.drop(drop cols, axis = 1, inplace = True)
         admissions['Avg_LOS'].fillna((admissions['Avg_LOS'].mean()), inplace=True)
In [24]:
         admissions['N_CAREUNITS'].fillna((admissions['N_CAREUNITS'].mean()), inplace=T
         admissions['HCFA'].fillna((admissions['HCFA'].mode()[0]), inplace=True)
         admissions['APR'].fillna((admissions['APR'].mode()[0]), inplace=True)
         admissions['MS'].fillna((admissions['MS'].mode()[0]), inplace=True)
         admissions['service_ct'].fillna((admissions['service_ct'].mean()), inplace=Tru
         e)
         admissions['DBSOURCE'].fillna('carevue', inplace = True)
In [25]:
          admissions['RELIGION'].fillna('CATHOLIC', inplace = True)
```

EDA

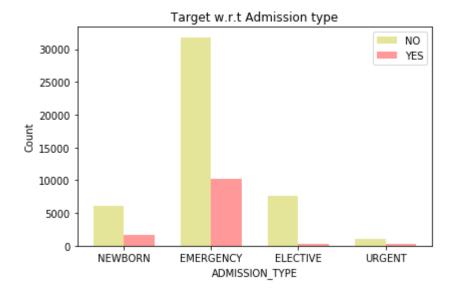
```
In [26]: import matplotlib.pyplot as plt
%matplotlib inline
```

```
def group_bar(data, column, n_groups, x_label, title, labels):
In [27]:
             no = np.array(data[[column, 'target']][data.target == 0].groupby(column)
         .agg('count').reset_index()['target'])
             yes = np.array(data[[column, 'target']][data.target == 1].groupby(column
         ).agg('count').reset_index()['target'])
             fig, ax = plt.subplots()
             index = np.arange(n groups)
             bar width = 0.35
             rects1 = plt.bar(index, no, bar_width, alpha=0.4, color='y', label='NO')
             rects2 = plt.bar(index + bar_width, yes, bar_width, alpha=0.4, color='r'
         , label='YES')
             plt.xlabel(x_label)
             plt.ylabel('Count')
             plt.title(title)
             plt.xticks(index + bar_width / 2, labels)
             plt.legend()
             plt.tight_layout()
             return plt
```

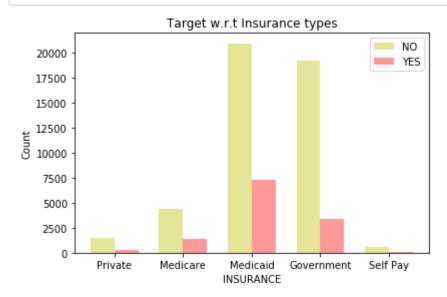
In [28]: p1 = group_bar(admissions, 'GENDER', 2, 'GENDER', 'Target w.r.t Gender', ['Mal
e', 'Female'])



In [29]: p2 = group_bar(admissions, 'ADMISSION_TYPE', 4, 'ADMISSION_TYPE', 'Target w.r.
t Admission type', ["NEWBORN", "EMERGENCY", "ELECTIVE", "URGENT"])



In [30]: types = ["Private", "Medicare", "Medicaid", "Government", "Self Pay"]
 p3 = group_bar(admissions, 'INSURANCE', 5, 'INSURANCE', 'Target w.r.t Insurance types', types)



Prepare Model data

In [36]: from sklearn.tree import DecisionTreeClassifier

```
In [33]: t_admissions = admissions.copy()
In [34]: le = LabelEncoder()
    for eachCol in cat_cols:
        admissions[eachCol] = le.fit_transform(admissions[eachCol])
In [35]: admissions.drop(drop_cols, axis = 1, inplace = True)
```

Build Model

```
from sklearn.ensemble import RandomForestClassifier
         from sklearn.model selection import train test split
         from sklearn.model selection import cross val score
         from sklearn.metrics import f1 score
         # conda install -c conda-forge imbalanced-learn
         from imblearn.under_sampling import RandomUnderSampler
         from imblearn.over sampling import SMOTE
In [37]:
         def data_sampling(X, y, method = 'rus', kind = 'regular', seed = 0):
             Function to balance classes using random under sampling or smote
             Parameters:
                 X: the independent variable values in a numpy array / pandas dataframe
                 y: the target variable valeus in a numpy array / pandas dataframe
                 method: currently accepts: 'rus' or 'smote'
                 kind: parameter for smote where kind = ['regular', 'borderline1', 'bor
         derline2', 'svm']
                 seed: A random seed to maintain reproducability
             if method == 'smote':
                 sm = SMOTE(kind = kind)
                 X_res, y_res = sm.fit_sample(X, y)
```

```
In [38]: feature_names = [x for x in admissions.columns if x not in ['target']]
    target = admissions['target']
```

X res, y res, idx res = rus.fit sample(X, y)

return {'X' : X_res, 'y' : y_res}

rus = RandomUnderSampler(return indices = True, random state = seed)

```
In [39]: X_train, X_valid, y_train, y_valid = train_test_split(admissions[feature_names
], target, train_size = 0.7, stratify = target, random_state = 0)
```

Handling Imbalance with RUS & SMOTE

1) Random Forest

```
In [42]: clf = RandomForestClassifier(random_state=0)
In [43]: cross_val_score(clf, X_SMOTE_train, y_SMOTE_train, cv=5)
Out[43]: array([0.68922018, 0.68176606, 0.67029817, 0.67230505, 0.68559954])
In [44]: clf.fit(X_SMOTE_train, y_SMOTE_train)
Out[44]: RandomForestClassifier(bootstrap=True, class_weight=None, criterion='gini', max_depth=None, max_features='auto', max_leaf_nodes=None, min_impurity_decrease=0.0, min_impurity_split=None, min_samples_leaf=1, min_samples_split=2, min_weight_fraction_leaf=0.0, n_estimators=10, n_jobs=1, oob_score=False, random_state=0, verbose=0, warm_start=False)
In [45]: f1_score(y_valid, clf.predict(X_valid))
Out[45]: 0.476584836833284
In []: from xgboost import XGBClassifier
```