Roll no: 225229113

# **Lab6.Predictive Analysis for Hospitals**

In [1]: import pandas as pd

## Step1. Import dataset

In [2]: df=pd.read\_csv('diabetes.csv')
 df

Out[2]:

:		Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	ВМІ	DiabetesPedigreeFunction
	0	6	148	72	35	0	33.6	0.627
	1	1	85	66	29	0	26.6	0.351
	2	8	183	64	0	0	23.3	0.672
	3	1	89	66	23	94	28.1	0.167
	4	0	137	40	35	168	43.1	2.288
	763	10	101	76	48	180	32.9	0.171
	764	2	122	70	27	0	36.8	0.340
	765	5	121	72	23	112	26.2	0.245
	766	1	126	60	0	0	30.1	0.349
	767	1	93	70	31	0	30.4	0.315

768 rows × 9 columns

In [3]: #head
 df.head(5)

Out[3]:

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	DiabetesPedigreeFunction	1
0	6	148	72	35	0	33.6	0.627	
1	1	85	66	29	0	26.6	0.351	
2	8	183	64	0	0	23.3	0.672	
3	1	89	66	23	94	28.1	0.167	
4	0	137	40	35	168	43.1	2.288	
4								<b>•</b>

```
In [4]: #shape
        df.shape
Out[4]: (768, 9)
In [5]: #columns
        df.columns
Out[5]: Index(['Pregnancies', 'Glucose', 'BloodPressure', 'SkinThickness', 'Insulin',
                'BMI', 'DiabetesPedigreeFunction', 'Age', 'Outcome'],
              dtype='object')
In [6]: #dtype
        df.dtypes
Out[6]: Pregnancies
                                       int64
        Glucose
                                       int64
        BloodPressure
                                       int64
        SkinThickness
                                       int64
        Insulin
                                       int64
        BMI
                                     float64
        DiabetesPedigreeFunction
                                     float64
        Age
                                       int64
        Outcome
                                       int64
        dtype: object
```

```
In [7]:
         #info
         df.info
Out[7]: <bound method DataFrame.info of</pre>
                                                   Pregnancies Glucose BloodPressure
                                                                                            SkinT
         hickness Insulin
                                BMI
                                  148
                                                    72
                          6
                                                                     35
                                                                                0
                                                                                   33.6
         1
                          1
                                   85
                                                    66
                                                                     29
                                                                                0
                                                                                    26.6
         2
                          8
                                  183
                                                    64
                                                                      0
                                                                                0
                                                                                   23.3
         3
                          1
                                   89
                                                    66
                                                                     23
                                                                               94
                                                                                   28.1
         4
                          0
                                  137
                                                    40
                                                                     35
                                                                              168
                                                                                   43.1
         . .
                                  . . .
                                                   . . .
                                                                    . . .
                                                                              . . .
                        . . .
                                                    76
                                                                              180
                                                                                   32.9
         763
                         10
                                  101
                                                                     48
         764
                          2
                                  122
                                                    70
                                                                     27
                                                                                0 36.8
                                                                              112 26.2
         765
                          5
                                                    72
                                                                     23
                                  121
         766
                          1
                                  126
                                                    60
                                                                      0
                                                                                0
                                                                                   30.1
         767
                          1
                                   93
                                                    70
                                                                     31
                                                                                0
                                                                                   30.4
               DiabetesPedigreeFunction
                                            Age
                                                 Outcome
         0
                                    0.627
                                             50
                                                         1
         1
                                    0.351
                                             31
                                                         0
         2
                                    0.672
                                             32
                                                         1
         3
                                    0.167
                                                         0
                                             21
                                    2.288
                                                         1
         4
                                             33
                                             . . .
         763
                                    0.171
                                             63
                                                        0
         764
                                    0.340
                                             27
                                                        0
         765
                                    0.245
                                             30
                                                        0
         766
                                    0.349
                                                         1
                                             47
                                    0.315
                                             23
                                                        0
         767
         [768 rows x 9 columns]>
In [8]:
         #value_counts
         df.Glucose.value_counts
Out[8]: <bound method IndexOpsMixin.value_counts of 0</pre>
                                                                   148
                  85
         2
                 183
                  89
         3
         4
                 137
         763
                 101
         764
                 122
         765
                 121
         766
                 126
         767
                  93
```

Step2. Identify relationships between feature

Name: Glucose, Length: 768, dtype: int64>

In [9]:	<pre>: df.style.background_gradient(cmap = 'GnBu')</pre>								
Out[9]:		Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	ВМІ	DiabetesPedigree	
	0	6	148	72	35	0	33.600000		
	1	1	85	66	29	0	26.600000		
	2	8	183	64	0	0	23.300000		
	3	1	89	66	23	94	28.100000		
	4	0	137	40	35	168	43.100000		
	5	5	116	74	0	0	25.600000		
	6	3	78	50	32	88	31.000000		
	7	10	115	0	0	0	35.300000		
	8	2	197	70	45	543	30.500000		
	9	8	125	96	0	0	0.000000		
	10	4	110	92	0	0	37.600000	•	
								<b>&gt;</b>	

## Step3. Prediction using one feature

```
In [14]: logr.predict(X test)
Out[14]: array([0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0,
             0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 1, 0, 0, 0, 0, 1, 0,
             0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 1, 0, 0,
             1, 0, 1, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 1, 0, 0, 0, 0,
             0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 1, 1, 0, 0, 0, 0,
             0, 1, 0, 0, 0, 0, 1, 0, 1, 0, 0, 0, 0, 1, 0], dtype=int64)
In [15]: print("Coef :",logr.coef_)
       print("Intercept :",logr.intercept_)
       Coef : [[0.04278121]]
       Intercept : [-2.06807344]
In [16]: logr.predict([[60]])
       C:\Users\Harsmitha\anaconda3\lib\site-packages\sklearn\base.py:450: UserWarnin
       g: X does not have valid feature names, but LogisticRegression was fitted with
       feature names
         warnings.warn(
Out[16]: array([1], dtype=int64)
In [17]: | lrf=logr.coef *60+logr.intercept
       from scipy.special import expit
       d = expit(lrf)
In [18]: if d>0.5:
           print('YES he will become diabetic')
       else:
           print('NO he will not be diabetic')
       YES he will become diabetic
```

## Step4.Prediction using many features

```
In [19]: X1=df[['Glucose','BMI','Age']]
X1_train,X1_test,Y1_train,Y1_test = train_test_split(X1,Y,test_size=.25,random_st
logr1= LogisticRegression()
logr1.fit(X1_train,Y1_train)

C:\Users\Harsmitha\anaconda3\lib\site-packages\sklearn\utils\validation.py:993:
    DataConversionWarning: A column-vector y was passed when a 1d array was expecte
    d. Please change the shape of y to (n_samples, ), for example using ravel().
        y = column_or_1d(y, warn=True)
Out[19]: LogisticRegression()
```

```
In [20]: logr1.predict(X1 test)
Out[20]: array([0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 1, 0, 0, 0, 1, 1, 0, 0, 0, 1, 1, 0, 0,
                1, 0, 0, 0, 0, 0, 1, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
                0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 1, 1, 0, 1, 0, 1, 0, 0, 0,
                0, 0, 0, 0, 0, 1, 1, 0, 0, 0, 0, 1, 0, 0, 1, 0, 0, 0, 0, 0, 0,
                1, 1, 1, 1, 0, 1, 0, 0, 1, 1, 0, 0, 0, 0, 1, 0, 0, 1, 0, 0, 1,
                0, 1, 0, 1, 0, 0, 0, 1, 0, 1, 0, 0, 0, 0, 0, 0, 0, 1, 0, 1, 0, 0,
                1, 0, 0, 0, 0, 0, 1, 1, 1, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0,
                0, 0, 1, 1, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0,
                0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0], dtype=int64)
In [21]: print("Coef :",logr1.coef_)
         print("Intercept :",logr1.intercept_)
         Coef: [[0.03358049 0.07889299 0.02722911]]
         Intercept : [-8.37441801]
In [22]: |lrf1=logr1.coef *150*40*30+logr1.intercept
         from scipy.special import expit
         expit(lrf1)
Out[22]: array([[1., 1., 1.]])
In [23]: logr1.predict([[150,40,30]])
         C:\Users\Harsmitha\anaconda3\lib\site-packages\sklearn\base.py:450: UserWarnin
         g: X does not have valid feature names, but LogisticRegression was fitted with
         feature names
           warnings.warn(
Out[23]: array([1], dtype=int64)
In [24]: logr1.predict_proba([[150,40,30]])
         C:\Users\Harsmitha\anaconda3\lib\site-packages\sklearn\base.py:450: UserWarnin
         g: X does not have valid feature names, but LogisticRegression was fitted with
         feature names
           warnings.warn(
Out[24]: array([[0.34632141, 0.65367859]])
```

Step5. Build LoR model with all features

```
In [25]: X2=df.drop(['Outcome'],axis=1)
         X2 train, X2_test, Y2_train, Y2_test = train_test_split(X2, Y, test_size=.25, random_st
         logr2=LogisticRegression()
         logr2.fit(X2 train,Y2 train)
         logr2.predict(X2_test)
         C:\Users\Harsmitha\anaconda3\lib\site-packages\sklearn\utils\validation.py:993:
         DataConversionWarning: A column-vector y was passed when a 1d array was expecte
         d. Please change the shape of y to (n_samples, ), for example using ravel().
           y = column or 1d(y, warn=True)
         C:\Users\Harsmitha\anaconda3\lib\site-packages\sklearn\linear model\ logistic.p
         y:814: ConvergenceWarning: lbfgs failed to converge (status=1):
         STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.
         Increase the number of iterations (max_iter) or scale the data as shown in:
             https://scikit-learn.org/stable/modules/preprocessing.html (https://scikit-
         learn.org/stable/modules/preprocessing.html)
         Please also refer to the documentation for alternative solver options:
             https://scikit-learn.org/stable/modules/linear_model.html#logistic-regressi
         on (https://scikit-learn.org/stable/modules/linear_model.html#logistic-regressi
         on)
           n iter i = check optimize result(
Out[25]: array([0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 1, 1, 0, 0,
                1, 0, 0, 0, 0, 0, 1, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0,
                0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 1, 1, 0, 1, 0, 1, 0, 0, 0,
                0, 0, 0, 0, 0, 1, 1, 0, 0, 0, 0, 1, 0, 0, 1, 0, 0, 0, 0, 0, 0,
                0, 1, 1, 1, 0, 1, 0, 0, 0, 0, 0, 1, 0, 0, 1, 0, 0, 1, 0, 0, 1,
                0, 1, 0, 1, 0, 0, 0, 1, 0, 1, 0, 0, 0, 0, 0, 0, 0, 1, 0, 1, 0, 0,
                1, 0, 0, 0, 0, 0, 1, 1, 0, 0, 0, 1, 0, 0, 1, 0, 0, 1, 0, 0, 0,
                0, 0, 1, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 1, 0, 0, 0, 0, 0, 0,
                0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0], dtype=int64)
In [26]: Y2_pred=logr2.predict(X2_test)
         Y2_pred
Out[26]: array([0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 1, 1, 0, 0,
                1, 0, 0, 0, 0, 0, 1, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0,
                0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 1, 1, 0, 1, 0, 1, 0, 0, 0,
                0, 0, 0, 0, 0, 1, 1, 0, 0, 0, 0, 1, 0, 0, 1, 0, 0, 0, 0, 0, 0,
                0, 1, 1, 1, 0, 1, 0, 0, 0, 0, 1, 0, 0, 1, 0, 0, 1, 0, 0, 1,
                0, 1, 0, 1, 0, 0, 0, 1, 0, 1, 0, 0, 0, 0, 0, 0, 0, 1, 0, 1, 0, 0,
                1, 0, 0, 0, 0, 0, 1, 1, 0, 0, 0, 0, 1, 0, 0, 1, 0, 0, 1, 0, 0,
                0, 0, 1, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 1, 0, 0, 0, 0, 0,
                0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0], dtype=int64)
In [27]: from sklearn.metrics import roc auc score
         lor_auc=roc_auc_score(Y2_test,Y2_pred)
         print("Auc:",lor_auc)
```

Auc: 0.7142857142857143

#### **Step6. Forward selection prodcure**

```
In [28]:
         #get auc
         def get auc(var,tar,df):
             fx = df[var]
             fy = df[tar]
             logr4=LogisticRegression()
             logr4.fit(fx,fy)
             pred=logr4.predict_proba(fx)[:,1]
             auc_val = roc_auc_score(Y,pred)
             return auc val
         get_auc(['Glucose',"BMI"],['Outcome'],df)
         C:\Users\Harsmitha\anaconda3\lib\site-packages\sklearn\utils\validation.py:993:
         DataConversionWarning: A column-vector y was passed when a 1d array was expecte
         d. Please change the shape of y to (n_samples, ), for example using ravel().
           y = column_or_1d(y, warn=True)
Out[28]: 0.8109328358208956
In [29]: | get_auc(['Pregnancies','BloodPressure','SkinThickness'],['Outcome'],df)
         C:\Users\Harsmitha\anaconda3\lib\site-packages\sklearn\utils\validation.py:993:
         DataConversionWarning: A column-vector y was passed when a 1d array was expecte
         d. Please change the shape of y to (n_samples, ), for example using ravel().
           y = column or 1d(y, warn=True)
Out[29]: 0.6444962686567164
```

```
In [30]: #fuction of best next
         def best next(current,cand,tar,df):
             best auc=-1
             best var=None
             for i in cand:
                 auc_v = get_auc(current+[i],tar,df)
                 if auc_v>=best_auc:
                     best_auc=auc_v
                     best_var=i
             return best var
         current=['Insulin','BMI','DiabetesPedigreeFunction','Age']
         cand=['Pregnancies','Glucose','BloodPressure','SkinThickness']
         tar=['Outcome']
         next_var = best_next(current,cand,tar,df)
         next_var
         C:\Users\Harsmitha\anaconda3\lib\site-packages\sklearn\utils\validation.py:993:
         DataConversionWarning: A column-vector y was passed when a 1d array was expecte
```

C:\Users\Harsmitha\anaconda3\lib\site-packages\sklearn\utils\validation.py:993:
DataConversionWarning: A column-vector y was passed when a 1d array was expecte
d. Please change the shape of y to (n\_samples, ), for example using ravel().
 y = column\_or\_1d(y, warn=True)
C:\Users\Harsmitha\anaconda3\lib\site-packages\sklearn\utils\validation.py:993:
DataConversionWarning: A column-vector y was passed when a 1d array was expecte
d. Please change the shape of y to (n\_samples, ), for example using ravel().
 y = column\_or\_1d(y, warn=True)
C:\Users\Harsmitha\anaconda3\lib\site-packages\sklearn\utils\validation.py:993:
DataConversionWarning: A column-vector y was passed when a 1d array was expecte
d. Please change the shape of y to (n\_samples, ), for example using ravel().
 y = column\_or\_1d(y, warn=True)
C:\Users\Harsmitha\anaconda3\lib\site-packages\sklearn\utils\validation.py:993:
DataConversionWarning: A column-vector y was passed when a 1d array was expecte
d. Please change the shape of y to (n\_samples, ), for example using ravel().
 y = column\_or\_1d(y, warn=True)

Out[30]: 'Glucose'

```
In [31]: | tar =['Outcome']
         current=[]
         cand=['Pregnancies','Glucose','BloodPressure','SkinThickness','Insulin','BMI','Di
         max num=7
         num_it = min(max_num,len(cand))
         for i in range(0,num_it):
             next var = best next(current,cand,tar,df)
             current += [next_var]
             cand.remove(next var)
             print("variable added in step "+str(i+1)+' is '+ next_var +" .")
         C:\Users\Harsmitha\anaconda3\lib\site-packages\sklearn\utils\validation.py:99
         3: DataConversionWarning: A column-vector y was passed when a 1d array was ex
         pected. Please change the shape of y to (n_samples, ), for example using rave
         1().
           y = column or 1d(y, warn=True)
         C:\Users\Harsmitha\anaconda3\lib\site-packages\sklearn\utils\validation.py:99
         3: DataConversionWarning: A column-vector y was passed when a 1d array was ex
         pected. Please change the shape of y to (n_samples, ), for example using rave
         1().
           y = column_or_1d(y, warn=True)
         C:\Users\Harsmitha\anaconda3\lib\site-packages\sklearn\utils\validation.py:99
         3: DataConversionWarning: A column-vector y was passed when a 1d array was ex
         pected. Please change the shape of y to (n_samples, ), for example using rave
           y = column or 1d(y, warn=True)
         C:\Users\Harsmitha\anaconda3\lib\site-packages\sklearn\utils\validation.py:99
         3: DataConversionWarning: A column-vector y was passed when a 1d array was ex
         pected. Please change the shape of y to (n samples, ), for example using rave
         1().
                          4 1/
In [32]:
         current
Out[32]: ['Glucose',
           'BMI',
           'Pregnancies',
           'DiabetesPedigreeFunction',
           'BloodPressure',
           'Age',
           'SkinThickness']
         step-7 Plot line graph of AUC values and select cut-off
```

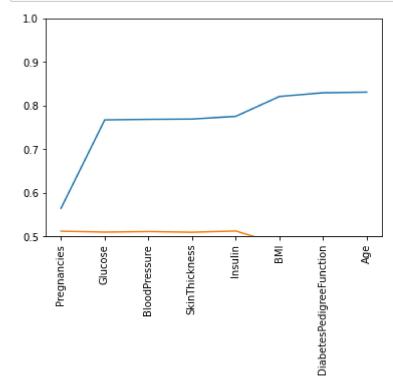
```
In [33]: X2_train,X2_test,Y2_train,Y2_test = train_test_split(X2,Y,stratify=Y,test_size=.5
In [34]: prediction=logr2.predict_proba(X2_test)
```

```
In [36]: #plot auc values
         train = pd.concat([X2_train,Y2_train],axis =1)
         test = pd.concat([X2_test,Y2_test],axis =1)
         def auc_train_test (variables, target, train, test):
             X_train = train[variables]
             X_test = test[variables]
             Y_train =train[target]
             Y_test = test[target]
             LoR=LogisticRegression()
             LoR.fit(X_train,Y_train)
             prediction_train = LoR.predict_proba(X_train)[:,1]
             prediction_test = LoR.predict_proba(X_test)[:,1]
             auc_train = roc_auc_score(Y_train, prediction_train)
             auc_test = roc_auc_score(Y_train,prediction_test)
             return (auc_train,auc_test)
         auc_values_train=[]
         auc_values_test=[]
         variable_evaluate=[]
         for v in X2.columns:
             variable_evaluate.append(v)
             auc_train,auc_test = auc_train_test(variable_evaluate,['Outcome'],train,test)
             auc_values_train.append(auc_train)
             auc values test.append(auc test)
         C:\Users\Harsmitha\anaconda3\lib\site-packages\sklearn\utils\validation.py:99
         3: DataConversionWarning: A column-vector y was passed when a 1d array was ex
         pected. Please change the shape of y to (n samples, ), for example using rave
           y = column_or_1d(y, warn=True)
         C:\Users\Harsmitha\anaconda3\lib\site-packages\sklearn\utils\validation.py:99
         3: DataConversionWarning: A column-vector y was passed when a 1d array was ex
         pected. Please change the shape of y to (n_samples, ), for example using rave
           y = column_or_1d(y, warn=True)
         C:\Users\Harsmitha\anaconda3\lib\site-packages\sklearn\utils\validation.py:99
         3: DataConversionWarning: A column-vector y was passed when a 1d array was ex
         pected. Please change the shape of y to (n samples, ), for example using rave
           y = column_or_1d(y, warn=True)
         C:\Users\Harsmitha\anaconda3\lib\site-packages\sklearn\utils\validation.py:99
         3: DataConversionWarning: A column-vector y was passed when a 1d array was ex
         pected. Please change the shape of y to (n_samples, ), for example using rave
           y = column_or_1d(y, warn=True)
         C:\Users\Harsmitha\anaconda3\lib\site-packages\sklearn\utils\validation.py:99
         3: DataConversionWarning: A column-vector y was passed when a 1d array was ex
         pected. Please change the shape of y to (n samples, ), for example using rave
           y = column_or_1d(y, warn=True)
         C:\Users\Harsmitha\anaconda3\lib\site-packages\sklearn\utils\validation.py:99
         3: DataConversionWarning: A column-vector y was passed when a 1d array was ex
         pected. Please change the shape of y to (n samples, ), for example using rave
           y = column_or_1d(y, warn=True)
         C:\Users\Harsmitha\anaconda3\lib\site-packages\sklearn\utils\validation.py:99
         3: DataConversionWarning: A column-vector y was passed when a 1d array was ex
```

```
Lab sheet - 6 - Jupyter Notebook
pected. Please change the shape of y to (n_samples, ), for example using rave
1().
 y = column_or_1d(y, warn=True)
C:\Users\Harsmitha\anaconda3\lib\site-packages\sklearn\linear model\ logisti
c.py:814: ConvergenceWarning: lbfgs failed to converge (status=1):
STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.
Increase the number of iterations (max_iter) or scale the data as shown in:
    https://scikit-learn.org/stable/modules/preprocessing.html (https://sciki
t-learn.org/stable/modules/preprocessing.html)
Please also refer to the documentation for alternative solver options:
    https://scikit-learn.org/stable/modules/linear_model.html#logistic-regres
sion (https://scikit-learn.org/stable/modules/linear model.html#logistic-regr
ession)
  n_iter_i = _check_optimize_result(
C:\Users\Harsmitha\anaconda3\lib\site-packages\sklearn\utils\validation.py:99
3: DataConversionWarning: A column-vector y was passed when a 1d array was ex
pected. Please change the shape of y to (n_samples, ), for example using rave
1().
  y = column or 1d(y, warn=True)
C:\Users\Harsmitha\anaconda3\lib\site-packages\sklearn\linear_model\_logisti
c.py:814: ConvergenceWarning: lbfgs failed to converge (status=1):
STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.
Increase the number of iterations (max_iter) or scale the data as shown in:
    https://scikit-learn.org/stable/modules/preprocessing.html (https://sciki
t-learn.org/stable/modules/preprocessing.html)
Please also refer to the documentation for alternative solver options:
    https://scikit-learn.org/stable/modules/linear model.html#logistic-regres
sion (https://scikit-learn.org/stable/modules/linear model.html#logistic-regr
ession)
  n_iter_i = _check_optimize_result(
```

In [37]: import matplotlib.pyplot as plt

```
In [38]: import numpy as np
    x =np.array(range(0,len(auc_values_train)))
    my_train = np.array(auc_values_train)
    my_test = np.array(auc_values_test)
    plt.xticks(x,X2.columns,rotation=90)
    plt.plot(x,my_train)
    plt.plot(x,my_test)
    plt.ylim(0.5,1)
    plt.show()
```



### In [41]: pip install scikit.plot

#### Collecting scikit.plot

Downloading scikit plot-0.3.7-py3-none-any.whl (33 kB)

Requirement already satisfied: scikit-learn>=0.18 in c:\users\harsmitha\anacond a3\lib\site-packages (from scikit.plot) (1.0.2)

Requirement already satisfied: scipy>=0.9 in c:\users\harsmitha\anaconda3\lib\s ite-packages (from scikit.plot) (1.7.3)

Requirement already satisfied: matplotlib>=1.4.0 in c:\users\harsmitha\anaconda 3\lib\site-packages (from scikit.plot) (3.5.1)

Requirement already satisfied: joblib>=0.10 in c:\users\harsmitha\anaconda3\lib \site-packages (from scikit.plot) (1.1.0)

Requirement already satisfied: numpy>=1.17 in c:\users\harsmitha\anaconda3\lib\site-packages (from matplotlib>=1.4.0->scikit.plot) (1.21.5)

Requirement already satisfied: fonttools>=4.22.0 in c:\users\harsmitha\anaconda 3\lib\site-packages (from matplotlib>=1.4.0->scikit.plot) (4.25.0)

Requirement already satisfied: packaging>=20.0 in c:\users\harsmitha\anaconda3 \lib\site-packages (from matplotlib>=1.4.0->scikit.plot) (21.3)

Requirement already satisfied: pyparsing>=2.2.1 in c:\users\harsmitha\anaconda3 \lib\site-packages (from matplotlib>=1.4.0->scikit.plot) (3.0.4)

Requirement already satisfied: cycler>=0.10 in c:\users\harsmitha\anaconda3\lib \site-packages (from matplotlib>=1.4.0->scikit.plot) (0.11.0)

Requirement already satisfied: python-dateutil>=2.7 in c:\users\harsmitha\anaco nda3\lib\site-packages (from matplotlib>=1.4.0->scikit.plot) (2.8.2)

Requirement already satisfied: kiwisolver>=1.0.1 in c:\users\harsmitha\anaconda 3\lib\site-packages (from matplotlib>=1.4.0->scikit.plot) (1.3.2)

Requirement already satisfied: pillow>=6.2.0 in c:\users\harsmitha\anaconda3\lib\site-packages (from matplotlib>=1.4.0->scikit.plot) (9.0.1)

Requirement already satisfied: six>=1.5 in c:\users\harsmitha\anaconda3\lib\sit e-packages (from python-dateutil>=2.7->matplotlib>=1.4.0->scikit.plot) (1.16.0) Requirement already satisfied: threadpoolctl>=2.0.0 in c:\users\harsmitha\anaconda3\lib\site-packages (from scikit-learn>=0.18->scikit.plot) (2.2.0)

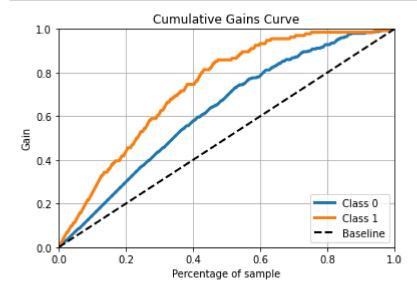
Installing collected packages: scikit.plot

Successfully installed scikit.plot-0.3.7

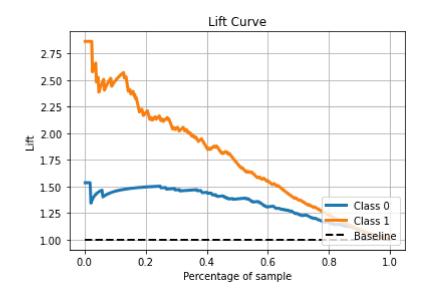
Note: you may need to restart the kernel to use updated packages.

#### In [42]: import scikitplot as skplt

```
In [45]: skplt.metrics.plot_cumulative_gain(Y2_test,prediction)
    plt.show()
    plt.figure(figsize=(7,7))
    skplt.metrics.plot_lift_curve(Y2_test, prediction)
    plt.show()
```



<Figure size 504x504 with 0 Axes>



In [ ]: